



UNIVERSITY of the
WESTERN CAPE

**THE EFFECTS OF SELECTIVE INHIBITORS OF N-GLYCOSYLATION
AND ENDOPLASMIC RETICULUM STRESS INDUCERS ON THE
EXPRESSION OF NEUROBLASTOMA DRUG RESISTANCE**

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DECLARATION

I, **Wejdan A.B. Mahmud Husein**, declare that “**The Effects of Selective Inhibitors of N-Glycosylation and Endoplasmic Reticulum Stress Inducers on the Expression of Neuroblastoma Drug Resistance**” is my original work and that all the sources that I have used or cited have been indicated and acknowledged by means of complete references, and that this document has not been submitted for degree purposes at any other academic institution.

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: 13 July 2017

DEDICATION

This study is dedicated to my family.



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ABSTRACT

Neuroblastoma (NB) represents 8-10% of all childhood tumours and accounts for approximately 15% of all cancer-related deaths in the paediatric population. Approximately half of newly diagnosed children with this tumour will present with metastatic disease or histologically aggressive large tumours that are at high risk for treatment failure. Since NBs are often widely disseminated and the tumours genetically heterogeneous in terms of their growth and metastatic behaviour, it is challenging to pinpoint their origin and predict disease prognosis. Several risk factors have been identified to play a role in disease progression, including age at the time of initial presentation, tumour stage, histology and ploidy of the tumour, and cytogenetic aberrations such as *MYCN* amplification, anaplastic lymphoma kinase (ALK), loss of heterozygosity of 11q and gain of 17q chromosomes.

Heredity is an important risk factor in about 1% to 2% of all NBs as children inherit an increased risk of developing NB from a parent. The stages and risk groups for NB are complex and can be perplexing. Advances in our knowledge of the biology and genetic basis of NB have led to the development of targeted and potentially useful therapeutic modalities. Many aggressive NBs exhibit multidrug resistance (MDR), attributable to p53 mutations and/or a loss of p53 function acquired during chemotherapy, which escalates the likelihood of relapse and thus presents a major obstacle to effective tumour eradication. Most metastatic drug-resistant NBs derive from the selection of clones (side population cells) that express the *MDR1* (*ABCB1*), *MRP1/ABCC1* and *MRP4/ABCC4* gene family, which may or may not correlate with *MYCN* amplification and poor outcome.

In NB, ganglioside signatures may influence tumour behaviour and clinical outcome. Thus, NB glycobiology impact on tumour growth and antitumour therapy. Targeted immunotherapy of NB with antibodies directed against disialoganglioside (GD2) has been amply documented. Direct and coordinate transcriptional targets of *MYCN* include several of the ATP-binding

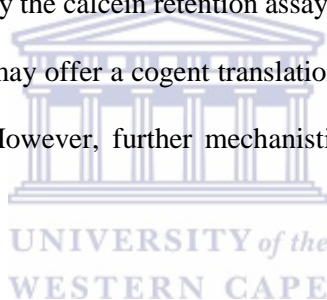
cassette (ABC) transporters—ABCB1 (P-glycoprotein/P-gp/MDR1). The expression of these MDR transporters are strongly prognostic of NB outcome since they extrude a wide array of structurally- and functionally-related or -unrelated chemotherapeutic drugs. The ABC transporters are thus promising candidates for therapeutic suppression in high-risk NB (HR-NB), the rationale behind increasing drug bioavailability (therapeutic efficacy) in refractory tumours which overexpress these glycoproteins.

P-gp is known to be overexpressed in NB, including the SK-N-BE(2) cell line selected for this study. Glycosylation of P-gp is critical for its location and function as a drug efflux pump to mediate MDR. In this study, the effects of aspirin (acetyl salicylic acid, a non-steroidal anti-inflammatory drug known to activate PERK and upregulate pro-apoptotic transcription factor CHOP (GADD153) which, together with cleavage of caspase-12, are hallmarks of ERS-mediated responses); bacitracin (an antibiotic that ablates glycoprotein synthesis at its first stage and interferes with P-glycoprotein (P-gp) expression and localization); castanospermine (a plant alkaloid that specifically inhibits α -glucosidases I and II, thus blocking elongation of glycan chains and formation of mature glycoproteins); brefeldin A (a metabolic inhibitor of N-glycosylation and disruptor of microtubule and actin cytoskeleton organization) and thapsigargin (a potent inducer of GRP78 expression and ERS, and activator of the UPR through non-competitive inhibition of the sarcoplasmic/endoplasmic reticulum calcium ATPase/SERCA) on SK-N-BE(2) cells were investigated.

The methods used to determine these effects were dose-response analysis, triplex-based fluorescence and luminescence cell cytotoxicity, viability and apoptosis assays, Annexin-V Cy3 fluorescence microscopy for apoptosis visualization and measurement of P-gp-mediated calcein efflux function. In this study, aspirin produced cytotoxicity towards SK-N-BE(2) cells, but viability was not affected. Aspirin had no effect on cell apoptosis at low concentrations, but at higher concentrations it decreased apoptosis induction. Bacitracin was shown to exert concentration-dependent effects on apoptosis in SK-N-BE(2) cells, i.e., at low concentrations

it increased caspase-dependent apoptosis, but at higher concentrations it reduced apoptosis. Such duality of effects is difficult to explain in the absence of mechanistic studies, especially since it was observed that bacitracin also decreased cytotoxicity commensurate with increased viability, but had no impact on P-gp efflux function. Results obtained in this study showed that castanospermine at all concentrations tested produced no cytotoxic effects, but at high concentrations resulted in contrasting effects, viz, increased viability and apoptosis, but no effect on calcein retention.

Brefeldin A, regarded as an inhibitor of P-gp, induced cytotoxicity and apoptosis in SK-N-BE(2) cells, but no inhibition of P-gp function was evident in the concentration range tested. Thapsigargin increased cytotoxicity and apoptosis in SK-N-BE(2) cells, but had no effect on P-gp function as measured by the calcein retention assay. It is concluded that efficacy of these ERS aggravators (ERSAs) may offer a cogent translational targeted cancer chemotherapeutic approach to treating NB. However, further mechanistic studies are needed to explain the responses observed.



Keywords: childhood cancer, neuroblastoma, inhibitors of N-glycosylation, endoplasmic reticulum stress inducers, neuroblastoma drug resistance, cytotoxicity, P-glycoprotein-mediated drug efflux

LIST OF ABBREVIATIONS

ABMT	Autologous Bone Marrow Transplantation
ABC	ATP-Binding Cassette
ADCC	Antibody Mediated Cell Cytotoxicity
ALK	Anaplastic Lymphoma Kinase
AnnCy3	Annexin-Cy3
ANOVA	One-Way Analysis of Variance
ANS	Autonomic Nervous System
ASCT	Autologous Stem-Cell Transplantation
ATCC	American Type Culture Collection
ATRA	All-Trans Retinoic Acid
AURKA	Aurora A Kinase
BAC	Bacitracin
BDNF	Brain-Derived Neurotrophic Factor
BFA	Brefeldin A
BMP	Bone Morphogenetic Protein
BMT	Bone Marrow Transplantation
Calcein-AM	Calcein–Acetoxymethylester
CAMs	Cell Adhesion Molecules
CBC	Complete Blood Count
CCG	Children’s Cancer Group
CCK-8	Cell Counting Kit-8
CD102	Cluster of Differentiation 102
6-CF	6-Carboxyfluorescein
6-CFDA	6-Carboxyfluorescein Diacetate
95%CI	95% Confidence Interval
COG	Children’s Oncology Group
COX	Cyclooxygenase
CRA	13-Cis-Retinoic Acid (Isotretinoin)
CSCs	Cancer Stem Cells

CST	Castanospermine
CT	Computed Tomography
DBH	Dopamine Beta-Hydroxylase Promoter
DEVD	Aspartic Acid, Glutamic Acid, Valine, Aspartic Acid
2dGlc/2-DG	2-Deoxyglucose
DLTs	Dose-Limiting Toxicities
DMEM	Dulbecco's Modified Eagle Medium
DMSO	Dimethylsulfoxide
DNJ/DMJ	Deoxynojirimycin/Deoxymannojirimycin
Dol-PP	Dolichol Pyrophosphate
DON	6-Diazo-5-Oxo-L-Norleucine
GD2	Diganglioside (Disialoganglioside)
EC	Epidural Compression
ECM	Extracellular Matrix
EFS	Event-Free Survival
EMT	Epithelial-to-Mesenchymal Transition
ER	Endoplasmic Reticulum
ERAD	Endoplasmic Reticulum-Associated Degradation
ERQC	Endoplasmic Reticulum Protein Quality Control System
ERS	Endoplasmic Reticulum Stress
ERSAs	Endoplasmic Reticulum Stress Aggravators
FDA	Food and Drug Administration
FDG	18-Fluorodeoxy-Glucose (18F-FDG)
FDG-PET	Fluorodeoxyglucose Positron Emission Tomography
FGF	Fibroblast Growth Factor
FISH	Fluorescence <i>In Situ</i> Hybridization
FKBP12	FK-Binding Protein 12
GAGs	Glycosaminoglycans
GATA	GATA transcription factors are a family of transcription factors characterized by their ability to bind to the DNA sequence "GATA".
GBPs	Glycan-Binding Proteins
GM-CSF	Granulocyte Macrophage Colony-Stimulating Factor

GN	Ganglioneuroma
GPI	Glycosylphosphatidylinositol
GRP/ GRPR	Gastrin-Releasing Peptide/Gastrin-Releasing Peptide Receptor
HAND2	Heart- and Neural Crest Derivatives-Expressed Protein
hESCs	Human Embryonic Stem Cells
HIF	Hypoxia-Inducing Factor
HIFBS	Heat-Inactivated Foetal Bovine Serum
HPR/4-HPR	N-(4-Hydroxyphenyl) Retinamide
HR-NB	High-Risk Neuroblastoma
HSCR	Hirschsprung's Disease
HSCT	Haematopoietic Stem Cell Transplantation
HSR	Homogeneously Staining Region(s)
hNCSCs	Human Neural Crest Stem Cells
HVA	Homovanillic Acid
IC₅₀	The Half Maximal Inhibitory Concentration of a Drug
ICAM-2	Intercellular Adhesion Molecule-2
IDRFs	Image-Defined Risk Factors
IGF2BP1	Insulin-Like Growth Factor-2 mRNA-Binding Protein 1
IMT	Inflammatory Myofibroblastic Tumour
INRG	International Neuroblastoma Risk Group
INRGSS	International Neuroblastoma Risk Group Staging System
INSS	International Neuroblastoma Staging System
INPC	International Neuroblastoma Pathology Classification
IUPAC	International Union of Pure and Applied Chemistry
IV	Intravenous
JNK	c-Jun Amino N-Terminal Kinase
LNN	Large Nucleolar Neuroblastoma
LOH	Loss of Heterozygosity
67LR	67-kDa Laminin Receptor
mAbs	Monoclonal Antibodies
MAPK	Mitogen-Activated Protein Kinase
MAT	Myeloablative Therapy

MASH1	Mammalian Achaete Scute Homologue-1
MDM2	Mouse Double Minute 2 Homologue
MDR	Multidrug Resistance
MEM	Modified Eagles Medium
MIBG	Meta-Iodobenzylguanidine
MKI	Mitosis-Karyorrhexis Index
MMPs	Matrix Metalloproteases
MRD	Minimal Residual Disease
MRI	Magnetic Resonance Imaging
MRP1	Multidrug Resistance-Associated Protein-1
MTDs	Maximal-Tolerated Doses
mTOR	Mammalian Target of Rapamycin
MSCs	Mesenchymal Stromal Cells
MYCL	Myelocytomatosis Viral Oncogene
MYCN	The <i>v-myc</i> avian myelocytomatosis viral oncogene neuroblastoma-derived homolog. <i>MYCN</i> remains the best-characterized genetic marker of risk in neuroblastoma.
NB(s)	Neuroblastoma(s)
NC	Neural Crest
NCAM	Neural Cell Adhesion Molecule
NGF	Nerve Growth Factor
NKT	Natural Killer T Cells
NMP	Nucleophosmin
NTRK	Neurotrophic Tyrosine Receptor Kinase(S)
NSAIDs	Nonsteroidal Anti-Inflammatory Drugs
NSCLC	Non-Small-Cell Lung Carcinoma
NT3	Neurotrophin-3 Growth Factor
ODC1	Ornithine Decarboxylase 1
OMS	Opsoclonus-Myoclonus Syndrome
OS	Overall Survival
PAH	Pulmonary Arterial Hypertension
PBS	Phosphate Buffered Saline
PBSCT	Peripheral Blood Stem Cell Transplantation

PCD	Programmed Cell Death
PET	Positron Emission Tomography
PFS	Progression-Free Survival
PI3K	Phosphoinositide-3-Kinase
Pgp	P-glycoprotein
PHOX2B	Paired-Like Homeobox 2b
PLC	Phospholipid C
pNTs	Peripheral Neuroblastic Tumours
POG	Paediatric Oncology Group
PPM1D	Protein Phosphatase Magnesium-Dependent 1 Delta
PS	Phosphatidylserine
PSA	Polysialic Acid
PTMs	Post-Translational Modifications
RA	Retinoic Acid
RARs/RXR s	Retinoic Acid Receptors/Retinoic Acid X (Rexinoid) Receptors
RAREs	Retinoic Acid Response Elements
RH	Relative Humidity
RTK	Receptor Tyrosine Kinase(s)
SEER	Surveillance, Epidemiology, and End Results Programme
SERCA	Sarcoplasmic/Endoplasmic Reticulum Calcium ATPase
SNS	Sympathetic Nervous System
SPECT	Single-Photon Emission Computed Tomography
SWSN	Swainsonine
RET	Rearranged During Transfection
TAFs	Tumour-Associated Fibroblasts
TAMs	Tumour-Associated Macrophages
TGFB	Transforming Growth Factor Beta
TH	Tyrosine Hydroxylase
TM	Tunicamycin
TME	Tumour Microenvironment
TICs	Tumour-Initiating Cells
TRK/Trk	Tyrosine Receptor Kinase

TSG(s)	Tumour Suppressor Gene(s)
UPR	Unfolded Protein Response
VEGF	Vascular Endothelial Growth Factor
VIP	Vasoactive Intestinal Peptide
VMA	Vanillylmandelic Acid
Wnt	Wingless/Integrated Proto-Oncogene
β-D-Xyl	β -D-Xyloside



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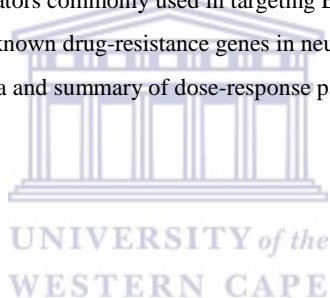
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CHAPTER 1

INTRODUCTION AND LITERATURE REVIEW

SECTION A: NEUROBLASTOMA

A1. Introduction

Neuroblastoma (NB) is the most common extracranial and deadly solid tumour in children and its origin has been clearly linked to the development of the sympathetic nervous system (SNS) because it originates from sympathetic precursor neuroblasts derived from the neural crest. Neuroblastoma represents 8-10% of all childhood tumours and accounts for approximately 15% of all cancer-related deaths in the paediatric population. The incidence of neuroblastoma is 10.2 cases per million children under 15 years of age, and nearly 500 new cases are reported annually. While 90% of cases are diagnosed before the age of 5, 30% of those are within the first year. The median age of diagnosis is 22 months. Rarely does it present in adolescence and adulthood, but outcomes are much poorer in this age group.

There does not appear to be an increased prevalence among races, but there is a slight predilection for males (1.2:1). With a family history noted in 1-2% of diagnoses, there are several reports of autosomal dominant patterns of inheritance. In such pedigrees, patients are frequently diagnosed at an earlier age (median age of 9 months) than those with sporadic disease and are more likely to have associated multiple primary cancers. Neuroblastoma has also been diagnosed in conjunction with other congenital conditions such as Hirschsprung's disease, congenital hypoventilation disorder and neurofibromatosis type 1. There was early interest in the co-occurrence of neuroblastoma and neurofibromatosis, as they are both disorders of neural crest cells. However, this may represent coincidence rather than a true association.^{1,2}

Approximately half of newly diagnosed children with this tumour will present with metastatic disease or histologically aggressive large tumours that are at high risk for treatment failure. Neuroblastomas found in patients older than 1 year are usually aggressive and eventually kill the patients despite intensive therapy, whereas those in patients younger than 1 year often regress spontaneously or mature, resulting in a more favourable prognosis. Because of the unique biological features of NBs, the neoplasm shows a wide range of clinical hallmarks, including the highest rate of spontaneous regression of any human malignancy, a potential for undergoing both induced and spontaneous maturation and also aggressive clinical courses with poor survival outcomes. The biological characteristics of NB are complex aneusomies, aneuploidies and ploidy shifts acquired by the tumour cells, and some of these chromosomal changes are known to be associated with clinical behaviour. The most commonly observed aneusomies include gains of 17q and deletions or allelic losses of 1p and 11q.

Combinations of these and other less prevalent genetic changes are detected in different genetic and clinical subsets of NB, and shown to be associated with tumour phenotype. Notwithstanding intensive investigation to map the shortest region of overlaps at deleted segments of 1p and 11q, a consistently involved candidate tumour suppressor gene has not yet been identified. Similarly, although PPM1D has been reported to be the most likely target genes with a recurrent oncogenic role at the minimal common region of gains at 17q remains to be identified. For years, *MYCN* was the only oncogene known to be involved recurrently in approximately 22% of tumours, and the *MYCN* protein is overexpressed via high copy number gains of the gene in tumours with advanced stages and aggressive clinical behaviour. The sections that follow will describe the global disease landscape of neuroblastoma in greater detail.

A2. What is Neuroblastoma?

Neuroblastoma (NB) is a paediatric cancer that originates from undifferentiated migratory neural crest (NC) progenitor or stem cells of the developing sympathetic nervous system

present in an embryo or foetus.³⁻⁶ These early precursor nerve cells are called neuroblasts—the term *neuro* signifies nerves, while *blastoma* denotes a cancer that affects immature or developing cells. Thus, neuroblastoma is a solid tumour that stems from the developing sympathetic nervous system (SNS) and can be found anywhere along this system.⁷ Neuroblastoma is prevalent in infants and young children—it rarely occurs in children older than 10 years.⁸ About 1 out of 3 neuroblastomas start in the adrenal glands and 1 out of 4 begin in sympathetic nerve ganglia in the abdomen, whereas the rest start in sympathetic ganglia near the spine in the chest or neck, or in the pelvis.⁶

Since NBs are often widely disseminated and the tumours genetically heterogeneous in terms of their growth and metastatic behaviour—some grow and spread quickly, while others grow slowly—it is challenging to pinpoint their origin and predict disease prognosis.⁹⁻¹¹ Sometimes in very young children, the cancer cells die for no reason and the tumour regresses spontaneously.¹² In other cases, the cells sometimes mature on their own into normal ganglion cells and stop dividing. This terminal differentiation makes the tumour a ganglioneuroma (GN).⁸ In order to understand neuroblastoma, it is essential to reflect on how the SNS functions.¹³ Therefore, the SNS is described briefly in the subsection that follows.

A3. The Sympathetic Nervous System

The nervous system consists of the brain, spinal cord and the nerves that reach out from them to all areas of the body. The nervous system is essential for cognitive function, sensation, movement and many sensory and motor functions that we are hardly ever aware of, including heart rate, breathing, blood pressure, and digestion. This division of the nervous system is known as the autonomic nervous system (ANS). Figure 1.1 shows the organization of the nervous system.

The SNS is a subdivision of the ANS. The SNS includes nerve fibres that run along either side the spinal cord. Clusters of nerve cells called ganglia (plural of ganglion) occur at certain points

along the path of the nerve fibres. Nerve-like cells are found in the medulla (centre) of the adrenal glands located superiorly to each kidney. These glands produce hormones (such as adrenaline (epinephrine) that help control heart rate, blood pressure, blood glucose and how the body reacts to stress. The main cells that make up the nervous system are called nerve cells or neurons. These cells interact with other types of cells in the body by releasing small amounts of chemical messengers (hormones). This is important, because neuroblastoma cells often release certain hormones that can cause symptoms of the tumour.

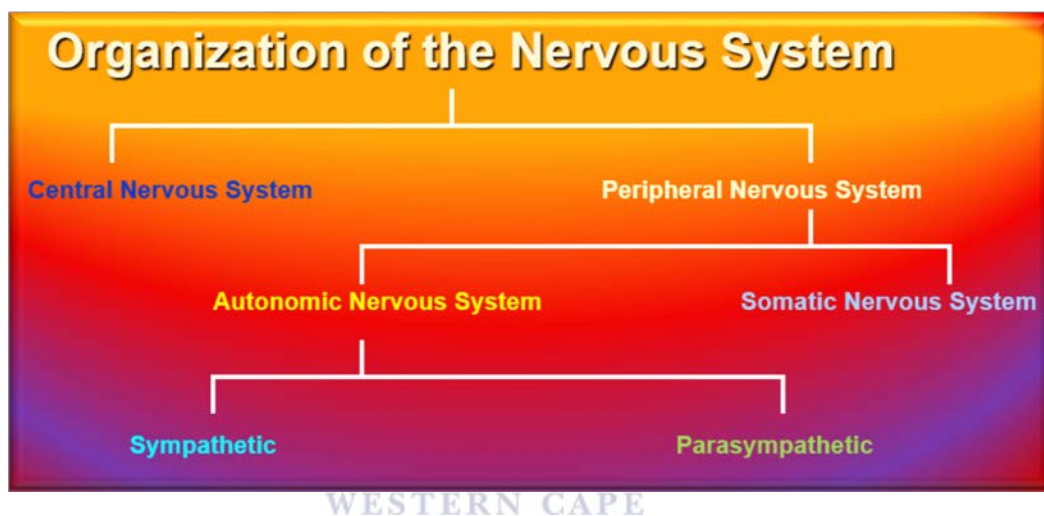
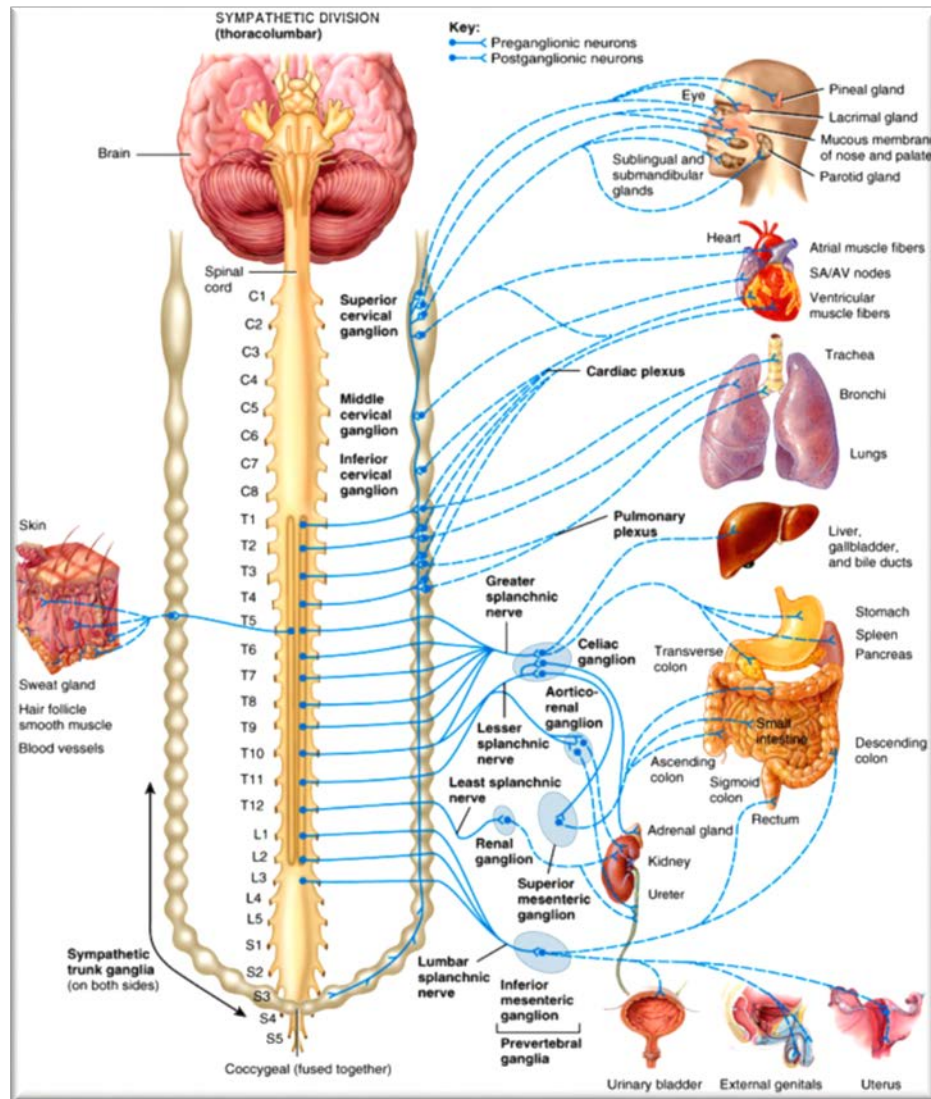


Figure 1.1: Organization of the nervous system

Figure 1.2 shows the greater structure of the SNS and thirty-one pairs of spinal nerves connected to the spinal cord, namely, 8 cervical nerve pairs (C1 through C8), 12 thoracic nerve pairs (T1 through T12), 5 lumbar nerve pairs (L1 through L5), 5 sacral nerve pairs (S1 through S5) and 1 coccygeal fused nerve pair.

A4. Epidemiology of Neuroblastoma: Incidence and Mortality Statistics

Neuroblastoma is by far the most common cancer in infants (less than 1 year old).¹⁴ Neuroblastoma accounts for about 6% of all cancers in children, with about 700 new cases reported each year in the United States. This number has remained constant for many years. The average age of children when they are diagnosed is about 1 to 2 years.



Source:¹⁵ Solid lines represent preganglionic axons; dashed lines represent postganglionic axons. Although the innervated structures are shown for only one side of the body for diagrammatic purposes, the sympathetic division actually innervates tissues and organs on both sides. Reproduced from Tortora GJ, Derrickson B. *Principles of Anatomy & Physiology*. 14 ed. Hoboken, NJ: John Wiley & Sons; 2014, with clearance from Wiley Global Permissions (Appendix 1)

Figure 1.2: Structure and pathways of the sympathetic division of the nervous system

In rare cases, NB is detected by ultrasound even before birth, but most cases (about 90%) are diagnosed by age 5. The malignancy is very rare in individuals over the age of 10 years. In about 2 of 3 cases, the disease has already spread to the lymph nodes or to other parts of the body when it is diagnosed. Table 1.1 summarizes USA survey and epidemiological data on childhood and adolescent cancers retrieved from *Cancer Facts & Figures 2014—Special Section: Childhood and Adolescent Cancers* at cancer.org/statistics. Table 1.2 outlines 5-year

survival rates for childhood cancers.¹⁶ Cancer is the second most common cause of death among children aged 1 to 14 years in the United States, exceeded only by accidents. In 2016, an estimated 10,380 children (birth to 14 years) will be diagnosed with cancer.

Table 1.1: Epidemiological information on childhood and adolescent cancers 2016

New Cases	An estimated 10,380 new cases of childhood cancers (ages 0-14 years) are expected to occur in 2016.
Incidence Trends	Childhood cancer incidence rates have gradually increased by 0.6% per year since 1975, when population-based cancer registration began in the US.
Deaths	An estimated 1,250 cancer deaths are expected to occur among children in 2016. Cancer is the second leading cause of death in children aged 1-14 years, eclipsed only by accidents.
Mortality Trends	Childhood cancer death rates dropped by 66% from 1969 (6.5 per 100,000) to 2012 (2.2 per 100,000), mainly as a result of improved treatment and high rates of participation in clinical trials. From 2003 to 2012, the childhood cancer death rate declined by 1.3% per year.
Survival	Survival for all invasive childhood cancers combined has improved markedly over the last 30 years due to novel and improved treatment strategies. The five-year relative survival rate increased from 58% in the mid-1970s to 83% in most recent times (2005-2011). However, rates differ considerably according to cancer type, patient age and other variables (see also Tables 1.1 and 1.2).
Some paediatric cancer patients experience treatment-induced side effects long after treatment, including impairment of function of specific organs (e.g., cognitive defects) and secondary cancers. ¹⁷ The Children's Oncology Group (COG) has developed guidelines for screening and managing of late effect survivors of childhood cancer (see COG website at survivorshipguidelines.org). The <i>Childhood Cancer Survivor Study</i> has followed more than 14,000 long-term childhood cancer survivors and posted valuable information on ccss.stjude.org .	

Approximately 1,250 children will die from the disease. Benign and borderline brain tumours are not included in the 2016 case estimates because the calculation method requires historical data and these tumours were not required to be reported until 2004. Leukaemia (76% of which are lymphoid leukaemias) accounts for 30% of all childhood cancers (including benign brain tumours). Cancers of the brain and other nervous system are the second most common cancer type (26%), followed by soft tissue sarcomas (7%, almost one-half of which are rhabdomyosarcoma), neuroblastoma (6%), non-Hodgkin lymphomas, including Burkitt's lymphoma (6%), renal (Wilms) tumours (5%), and Hodgkin lymphomas (3%).¹⁶

The 5-year survival rate refers to the percentage of children who live at least 5 years after their cancer is diagnosed. Many children may live much longer than 5 years (and many are even cured). In order to obtain 5-year survival rates, doctors have to look at children who were treated at least 5 years ago. Improvements in treatment since then may result in a better outlook for children now being diagnosed with NB.⁸

A5. Risk Factors for Neuroblastoma

Neuroblastoma is one of the most common childhood (age 0-14 years) cancers, being only surpassed in the paediatric age group by leukaemia and brain tumours.^{1-3,14,18} A risk factor is any factor that affects an individual's chance of getting a disease such as cancer. Different cancers have different risk factors. Lifestyle-related risk factors such as body weight, physical activity, diet and smoking play a major role in many adult cancers. However, these factors usually take many years to influence cancer risk, and are not usually associated with childhood cancers, including NBs. Also, no environmental factors (such as exposures during the mother's pregnancy or in early childhood) are known to increase the child's chance of getting NB.

Neuroblastoma is a very heterogeneous disease with features ranging from spontaneous regression during the foetal period to disseminated metastasis at the time of diagnosis. Several risk factors have been identified to play a role in disease progression, including age at the time of initial presentation, tumour stage, histology and ploidy of tumour, and cytogenetic aberrations such *MYCN* amplification, loss of heterozygosity of 11q and gain of 17q.^{4,9,11,19-24} According to the International Neuroblastoma Risk Group (INRG) task force report, age-range between 18 and 60 months is considered as a high risk group.^{25,26}

Heredity is an important risk factor in about 1% to 2% of all NBs as children inherit an increased risk of developing NB from a parent. Children with the familial form of NB (those with an inherited tendency to develop this cancer) usually come from families with one or more members who had NB as infants.

Table 1.2: Trends in 5-year relative survival rates for children (birth to 14 years) by year of diagnosis

Childhood cancer	1975-1977	1978-1980	1981-1983	1984-1986	1987-1989	1990-1992	1993-1995	1996-1998	1999-2001	2002-2004	2005-2011
All sites	58	62	67	68	72	76	77	79	81	83	83[†]
Acute lymphocytic leukaemia	57	66	71	72	78	83	84	87	89	92	91 [†]
Acute myeloid leukaemia	19	26	27 [‡]	31 [‡]	37 [‡]	42	41 [‡]	49	58	61	67 [†]
Bones and joints	50 [‡]	48	57 [‡]	57 [‡]	67 [‡]	67	74	70	70	78	77 [†]
Brain & other nervous system	57	58	57	62	64	64	71	75	74	75	74 [†]
Hodgkin lymphoma	81	87	88	90	87	97	95	96	94	98	98 [†]
Neuroblastoma	53	57	55	52	63	76	67	66	72	73	74[†]
Non-Hodgkin lymphoma	43	53	67	70	71	77	81	83	90	85	88 [†]
Soft tissue	61	74	69	73	66	80	77	71	77	85	79 [†]
Wilms tumour	73	79	87	91	92	92	92	92	94	89	94 [†]

Data are for surveys conducted in the United States, 1975 to 2011.

*Survival rates are adjusted for normal life expectancy and are based on follow-up of patients through 2012.

[†]The difference in rates between 1975 to 1977 and 2005 to 2011 is statistically significant ($p < 0.05$).

[‡]The standard error of the survival rate is between 5 and 10 percentage points.

Source¹⁶

Many other large epidemiological studies focusing on the incidence, prognostic factors, and the treatment outcomes in their patients have been carried out in different parts of the world, including Turkey,²⁷ Norway,²⁸ The European Neuroblastoma Study Group,²⁹ The Surveillance, Epidemiology, and End Results Programme (SEER, <http://seer.cancer.gov>),³⁰⁻³² Iran,^{33,34} Mexico³⁵ Australia, Europe, Japan, North America²⁶ and Italy.³⁶

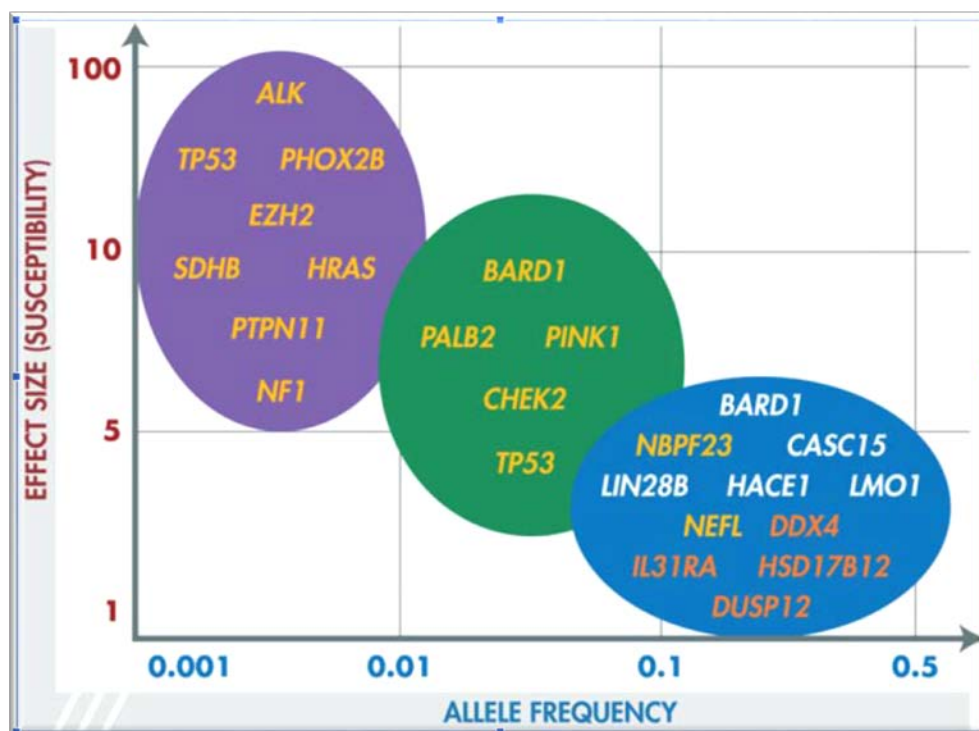
The average age at diagnosis of familial cases is younger than the age for sporadic (not inherited) cases. Children with familial NB sometimes develop 2 or more of these cancers in different organs (for example, in both adrenal glands or in more than one sympathetic ganglion). It's important to distinguish NBs that originate in more than one organ from those that have metastasized (spread) from one primary organ to secondary sites or organs (metastatic NBs). Frequently, tumours that have developed in several places at once implies a familial form. Both familial and sporadic NB can spread to other organs.

A recent review encapsulates the current state of knowledge about NB genetics and genomics, highlighting the improved prognosis and potential therapeutic opportunities that have arisen from recent advances in understanding germline predisposition, recurrent segmental chromosomal alterations, somatic point mutations and translocations and clonal evolution in relapsed NB.³⁷ Figure 1.3 shows rare and common genomic variants that predispose to NB.

A6. Staging of Neuroblastoma

The stages and risk groups for NB are complex and can be perplexing. A staging system is a standard approach used by a multidisciplinary cancer care team to classify the extent and burden of the cancer. Since the mid-1990s, most cancer centres have adopted the International Neuroblastoma Staging System (INSS) to stage NB.²⁵ This postsurgical staging system takes into account the results of surgery to remove the tumour.

Neuroblastoma belongs to a group collectively known as peripheral neuroblastic tumours, which also includes intermixed ganglioneuroblastoma, ganglioneuroma, and nodular ganglioneuroblastoma.³⁸⁻⁴⁰ Neuroblastoma can further be divided based on the degree of neuroblastic differentiation (undifferentiated, poorly differentiated, and differentiating) and the mitosis-karyorrhexis index (MKI) (low, intermediate, or high).⁴¹ Histologically, it has limited Schwannian cell production, is stroma-poor, and has abundant neuroblasts.



Source:³⁷ Reproduced from Bosse KR, Maris JM. Advances in the translational genomics of neuroblastoma: From improving risk stratification and revealing novel biology to identifying actionable genomic alterations. *Cancer* 2016;122(1):20-33, Copyright, American Cancer Society, with permission from John Wiley and Sons. See Appendix 2 for copyright clearance.

Figure 1.3: Rare and common genomic variants that predispose to neuroblastoma

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Left: In addition to anaplastic lymphoma kinase (ALK)-associated and paired-like homeobox 2B (PHOX2B)-associated familial neuroblastoma, neuroblastoma can also arise in the setting of genetic syndromes with underlying rat sarcoma oncogene-mitogen activated protein kinase (RAS-MAPK) pathway germline mutations, such as neurofibromin 1 (NF1) in neurofibromatosis type 1,50 protein tyrosine phosphatase, nonreceptor type 11 (PTPN11) in Noonan syndrome,^{51,52} and Harvey rat sarcoma viral oncogene homolog (HRAS) in Costello syndrome. Tumour protein 53 (TP53) mutations associated with Li Fraumeni syndrome, 53 enhancer of zeste 2 polycomb repressive complex 2 subunit (EZH2) mutations associated with Weaver syndrome, and succinate dehydrogenase complex, Subunit B, iron sulphur (Ip) (SDHB) mutations in familial paraganglioma/pheochromocytoma (PGL/PCC) are also rarely associated with neuroblastoma genesis. **Middle:** Low-frequency alleles in multiple DNA damage-response genes (BRCA1-associated ring domain 1 [BARD1], checkpoint kinase 2 [CHEK2], partner and localizer of BRCA2 [PALB2], and TP53) with an intermediate effect size also contribute to neuroblastoma predisposition. PINK1 indicates phosphatase and tensin homolog-induced putative kinase 1. **Right:** More common alleles with a modest effect size discovered using a genome-wide association study approach also collectively contribute to neuroblastoma genesis and, at times, specifically to a high-risk (white) or low-risk (orange) neuroblastoma phenotype. CASC15 indicates cancer susceptibility candidate 15; DDX4, DEAD (Asp-Glu-Ala-Asp) box polypeptide 4; DUSP12, dual specificity phosphatase 12; HACE1, HECT domain and ankyrin repeat containing E3 ubiquitin protein ligase 1; HSD17B12, hydroxysteroid (17b) dehydrogenase 12; IL31RA, interleukin 31 receptor A; LIN28B, lin-28 homologue B; LMO1, LIM domain only 1; NBPF23, neuroblastoma breakpoint family, member 23; NEFL, neurofilament, light polypeptide.

The International Neuroblastoma Pathology Classification has been used to predict prognosis based on the histopathology of the tumour and age of the patient. This system takes into account the degree of cell differentiation, MKI, and the presence of Schwann cells. Following these guidelines, the unfavourable group encompasses patients with any tumour over 60 months; undifferentiated tumours with a high MKI at any age; and undifferentiated or poorly

differentiated tumours with intermediate or high MKI in children older than 18 months. In simplified form, the stages are summarized in Table 1.3. A risk-group staging system now coming into use is known as the International Neuroblastoma Risk Group Staging System (INRGSS).²⁵ It is analogous to the INSS, but it does not use the results of surgery to help define the stage. This lets doctors determine a stage before surgery, based on the results of imaging tests, usually a computed tomography (CT) or magnetic resonance imaging (MRI) scan, and a meta-iodobenzylguanidine (MIBG) scan, as well as examinations and biopsies. The stage can then be used to help predict how resectable the tumour is—that is how much of it can be removed with surgery. The INRGSS uses image-defined risk factors (IDRFs) which are seen on imaging tests that might mean the tumour will be harder to remove. This includes features like the tumour growing into a nearby vital organ or growing around important blood vessels. The INRGSS divides NBs into 4 stages (Table 1.4).

The Children's Oncology Group (COG) uses major prognostic factors (section A7) combined with the INSS stage of the disease, to place children into 3 different risk groups: low, intermediate, and high (Table 1.5). These risk groups are used to help predict how likely it is that a child can be cured. For example, a child in a low-risk group can often be cured with limited treatment, such as surgery alone. With children in higher risk groups, the chance of cure is not as great, so more intensive treatment is often needed. A newer risk group classification system, the International Neuroblastoma Risk Group (INRG) classification (Table 1.6), is now being studied and may soon replace the COG system above. This system is based on the newer INRGSS staging system, which includes the image-defined risk factors (IDRFs), as well as many of the prognostic factors discussed in section A7 below.

A7. Prognostic Markers for Neuroblastoma

Prognostic markers are features that help predict whether the child's outlook for cure is better or worse than would be predicted by the stage alone. Markers used to help determine a child's prognosis are summarized in Table 1.7.

Table 1.3: The International Neuroblastoma Staging System (INSS)

Stage	Description
1	Localized tumour with complete gross excision with or without microscopic residual disease; ipsilateral and contralateral lymph node (LN) negative for tumour microscopically. The cancer is still in the area where it started. It is on one side of the body (right or left). All visible tumour has been removed completely by surgery (although looking at the tumour's edges under the microscope after surgery may show some cancer cells). Lymph nodes outside the tumour are free of cancer (although nodes enclosed within the tumour may contain neuroblastoma cells).
2A	Unilateral tumour with incomplete gross resection; ipsilateral and contralateral LN negative for tumour microscopically. The cancer is still in the area where it started and on one side of the body, but not all of the visible tumour could be removed by surgery. Lymph nodes outside the tumour are free of cancer (although nodes enclosed within the tumour may contain neuroblastoma cells).
2B	Unilateral tumour with or without complete gross excision with ipsilateral LN positive for tumour; contralateral LN negative microscopically. The cancer is on one side of the body, and may or may not have been removed completely by surgery. Nearby lymph nodes outside the tumour contain neuroblastoma cells, but the cancer has not spread to lymph nodes on the other side of the body or elsewhere.
3	The cancer has not spread to distant parts of the body, but one of the following is true of the cancer: <ul style="list-style-type: none"> • Cannot be removed completely by surgery and it has crossed the midline (defined as the spine) to the other side of the body. It may or may not have spread to nearby lymph nodes. • Is still in the area where it started and is on one side of the body. It has spread to lymph nodes that are relatively nearby but on the other side of the body. • Is in the middle of the body and is growing toward both sides (either directly or by spreading to nearby lymph nodes) and cannot be removed completely by surgery.
4A	Any primary tumour with dissemination to distant LN, bone, bone marrow, liver, skin, or other organs (except as defined for stage 4S). Tumour infiltrating across the midline with or without regional LN involvement, localized unilateral tumour with contralateral regional LN involvement, or midline tumour with bilateral extension by infiltration (unresectable) or by LN involvement. The cancer has spread to distant sites such as distant lymph nodes, bone, liver, skin, bone marrow, or other organs (but the child does not meet the criteria for stage 4S)
4S	Localized primary tumor (as defined for stage 1 or 2) with dissemination limited to skin, liver, or bone marrow (limited to infants <1 yr of age). Also called "special" neuroblastoma. The child is younger than 1 year. The cancer is on one side of the body. It might have spread to lymph nodes on the same side of the body but not to nodes on the other side. The neuroblastoma has spread to the liver, skin, and/or the bone marrow. However, no more than 10% of marrow cells are cancerous, and imaging tests such as a meta-iodobenzylguanidine (MIBG) scan do not show that the cancer has spread to the bones or the bone marrow.
Recurrent	While not formally part of the staging system, this term is used to describe cancer that has come back (recurred) after it has been treated. The cancer might come back in the area where it first started or in another part of the body.

Table 1.4: The International Neuroblastoma Risk Group Staging System (INRGSS)

Stage	Description
L1	Localized disease that does not involve vital structures and is confined to one body compartment, i.e., a tumour that has not spread from where it started and has not grown into vital structures as defined by the list of IDRFs. It is confined to one body compartment, such as the neck, chest, or abdomen.
L2	Localized disease with image-defined risk factors, i.e., a tumour that has not spread far from where it started (for example, it may have grown from the left side of the abdomen into the left side of the chest), but that has at least one IDRF.
M	Distant metastatic disease, i.e., a tumour that has spread (metastasized) to a distant part of the body (except tumours that are stage MS).
MS	Metastatic disease in children younger than 18 months with cancer spread only to skin, liver, and/or bone marrow. No more than 10% of marrow cells are cancerous, and an MIBG scan does not show spread to the bones and/or the bone marrow.
<p>IDRFs: image-defined risk factors</p> <p>MIBG: meta-iodobenzylguanidine</p>	

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Table 1.5: Children’s Oncology Group (COG) risk groups

Risk group	Description
Low risk	<ul style="list-style-type: none"> • All children who are Stage 1 • Any child who is Stage 2A or 2B and younger than age 1 • Any child who is Stage 2A or 2B, older than age 1, whose cancer has <i>no</i> extra copies of the <i>MYCN</i> gene • Any child who is Stage 4S (younger than age 1), whose cancer has favorable histology, is hyperdiploid (excess DNA) and has no extra copies of the <i>MYCN</i> gene
Intermediate risk	<ul style="list-style-type: none"> • Any child who is Stage 3, younger than age 1, whose cancer has no extra copies of the <i>MYCN</i> gene • Any child who is Stage 3, older than age 1, whose cancer has no extra copies of the <i>MYCN</i> gene and has favorable histology (appearance under the microscope) • Any child who is Stage 4, younger than age 1, whose cancer has no extra copies of the <i>MYCN</i> gene • Any child who is Stage 4S (younger than age 1), whose cancer has no extra copies of the <i>MYCN</i> gene and has normal DNA ploidy (number of chromosomes) and/or has unfavorable histology
High risk	<ul style="list-style-type: none"> • Any child who is Stage 2A or 2B, older than age 1, whose cancer has extra copies of the <i>MYCN</i> gene • Any child who is Stage 3, younger than age 1, whose cancer has extra copies of the <i>MYCN</i> gene • Any child who is Stage 3, older than age 1, whose cancer has extra copies of the <i>MYCN</i> gene • Any child who is Stage 3, older than 18 months of age, whose cancer has unfavorable histology • Any child who is Stage 4, whose cancer has extra copies of the <i>MYCN</i> gene regardless of age • Any child who is Stage 4 and older than 18 months • Any child who is Stage 4 and between 12 and 18 months old whose cancer has extra copies of the <i>MYCN</i> gene, unfavorable histology, and/or normal DNA ploidy (a DNA index of 1) • Any child who is Stage 4S (younger than age 1), whose cancer has extra copies of the <i>MYCN</i> gene

Table 1.6: International Neuroblastoma Risk Group (INRG) classification

Classification
<ul style="list-style-type: none"> • The child's age • Tumour histology • The presence or absence of <i>MYCN</i> gene amplification • Certain changes in chromosome 11 (known as an 11q aberration) • DNA ploidy (the total number of chromosomes in the tumour cells)
<p>The INRG classification uses these factors to put children into 16 different pre-treatment groups (lettered A through R). Each of these pretreatment groups falls into 1 of 4 overall risk groups:</p> <ul style="list-style-type: none"> • Very low risk • Low risk • Intermediate risk • High risk <p>This system has not yet been widely adopted, but it is being researched in new treatment protocols.</p>

Table 1.8 indicates the Children's Oncology Group (COG) survival outlook according to disease risk. Risk-based treatment approaches for NB have been used for many years. However, the criteria employed to delimit risk in various institutional and cooperative groups were incongruent, limiting the evaluation of clinical trial results. To alleviate this drawback and boost collaborative research, homogenous pretreatment patient cohorts have been defined by the INRG classification system. This treatment rationale has yielded improved outcomes, even though survival for high-risk patients remains poor, underscoring the dire need to develop more effective treatment strategies.

Advances in our knowledge of the biology and genetic basis of NB have led to the development of targeted and potentially useful therapeutic modalities.^{10,42,43} The collaborative ventures of institutions and international cooperative groups have refined risk classification and stratified treatment strategies, resulting in improved survival rates for NB patients.^{2,44,45}

A8. Other Autonomic Nervous System Tumours in Children

Not all childhood ANS tumours are malignant. Features of two of the more common ANS tumours are summarized in Table 1.9.

Table 1.7: Prognostic markers for neuroblastoma

Marker	Clinical Significance
Age	Younger children (under 12-18 months) are more likely to be cured than older children.
Tumour histology	Tumour histology is based on how the neuroblastoma cells look under the microscope. Tumours that contain more normal-looking cells and tissues tend to have a better prognosis and are said to have a favourable histology. Tumours whose cells and tissues look more abnormal under a microscope tend to have a poorer prognosis and are said to have an unfavourable histology.
DNA ploidy	The amount of DNA in each cell, known as ploidy or the DNA index, can be measured using special lab tests, such as flow cytometry or imaging cytometry. Neuroblastoma cells with about the same amount of DNA as normal cells (a DNA index of 1) are classified as diploid. Cells with increased amounts of DNA (a DNA index higher than 1) are termed hyperdiploid. In infants, hyperdiploid cells tend to be associated with earlier stages of disease, respond better to chemotherapy, and usually predict a more favourable prognosis (outcome) than diploid cells. Ploidy is not as useful a factor in older children.
MYCN gene amplifications	<i>MYCN</i> is an oncogene, a gene that helps regulate cell growth. Changes in oncogenes can make cells grow and divide too quickly, as with cancer cells. Neuroblastomas with too many copies (amplification) of the <i>MYCN</i> oncogene tend to grow quickly and are less likely to mature. Children whose neuroblastomas have this feature tend to have a worse prognosis than other children with neuroblastoma.
Chromosome changes	Tumour cells that are missing certain parts of chromosomes 1 or 11 (known as 1p deletions or 11q deletions) may predict a less favorable prognosis. It is thought that these chromosome parts, which are missing in many neuroblastomas, may contain important tumour suppressor genes (TSGs), but more studies are needed to verify this. Having an extra part of chromosome 17 (17q gain) is also linked with a worse prognosis. This probably means that there is an oncogene in this part of chromosome 17.
Neurotrophin (nerve growth factor) receptors	These are substances on the surface of normal nerve cells and on some neuroblastoma cells. They normally allow the cells to recognize neurotrophins—hormone-like chemicals that help the nerve cells mature. Neuroblastomas that have more of certain neurotrophin receptors, especially the nerve growth factor receptor called TrkA, may have a better prognosis.
Serum markers	Serum (blood) levels of certain substances can be used to help predict prognosis. Neuroblastoma cells release ferritin, an important regulator of the body's normal iron metabolism, into the blood. Patients with high ferritin levels tend to have a worse prognosis. Neuron-specific enolase (NSE) and lactate dehydrogenase (LDH) are synthesized by normal cells as well as by NB cells. Increased levels of NSE and LDH in the blood are often linked with a worse outlook in children with NB. A substance on the surface of many nerve cells known as ganglioside GD2 is often increased in the blood of NB patients. Although the usefulness of GD2 in predicting prognosis is unknown, it may turn out to be more important in treating NB.

Table 1.8: Survival of neuroblastoma patients by Children’s Oncology Group (COG) risk group

Risk Group	5-Year Survival Rate
Low	Higher than 95%
Intermediate	90% to 95%
High	40% to 50%
Source: <i>Neuroblastoma Summary</i> , American Cancer Society (http://www.cancer.org/)	

Table 1.9: Other autonomic nervous system tumours in children

Tumour	Features
Ganglioneuroma	A benign (non-cancerous) tumour made up of mature ganglion and nerve sheath cells.
Ganglioneuroblastoma	<p>A tumour that has both malignant and benign parts. It contains neuroblasts (immature nerve cells) that can grow and spread abnormally, similar to neuroblastoma, as well as areas of more mature tissue that are similar to ganglioneuroma.</p> <p>Ganglioneuromas are usually removed by surgery and looked at carefully under a microscope to be sure they don’t have areas of malignant cells (which would generate a ganglioneuroblastoma). If the final diagnosis is ganglioneuroma, no other treatment is needed. If it’s found to be a ganglioneuroblastoma, it’s treated the same as a neuroblastoma.</p>
Source: <i>Neuroblastoma Summary</i> , American Cancer Society (http://www.cancer.org/)	

A9. Detection, Diagnosis and Prognosis of Neuroblastoma

A9.1 Imaging and Laboratory Tests

Neuroblastomas are customarily suspected when a child presents with signs or symptoms, but a definite diagnosis is made after correlating physical examination with laboratory tests.^{44,46-52}

Table 1.10 outlines contemporary procedures and approaches used in confirming a diagnosis of neuroblastoma.^{44,46-51,53-61}

Table 1.10: Contemporary procedures and approaches used in confirming a diagnosis of neuroblastoma

Procedure/Approach	Diagnostic Significance/Prognostic Value
Medical history and physical examination	If the child presents with signs or symptoms that might suggest neuroblastoma, a complete medical history, as well as a family history of any type of cancer, is indispensable. Possible signs of a neuroblastoma are an abnormal mass or swelling in the body, lumps or bumps under the skin or high blood pressure. Neuroblastomas that grow adjacent to the spinal cord can affect the movement and strength in the child's arms and legs, so particular attention has to be paid to these. Some signs that could be caused by neuroblastoma, such as fever and enlarged lymph nodes, are much more likely to be caused by an infection, so it is prudent to look for other signs of infection at first. If the history and examination imply a child might have a neuroblastoma (or another type of tumour), other specialized tests will be mandatory, including blood and urine tests, imaging tests, and biopsies. These tests are important because many of the symptoms and signs of neuroblastoma can also be caused by other diseases, such as infections, or even other types of cancer.
Blood and urine catecholamine tests	Sympathetic nerve cells normally release hormones called catecholamines, such as epinephrine (adrenaline) and norepinephrine (noradrenaline) into the blood. Eventually the body degrades these into metabolites which is secreted in urine. Neuroblastoma cells can also synthesize these hormones. In most cases, neuroblastoma cells make enough catecholamines to be detected by blood or urine tests. The 2 catecholamine metabolites most often measured are homovanillic acid (HVA) and vanillylmandelic acid (VMA).
Other lab tests	If neuroblastoma is suspected or has been found in a child, certain blood tests will be requested to check blood cell counts, liver and kidney function, and the balance of salts (electrolytes) in the body. A urinalysis (urine test) may also be done to further check kidney function.
Imaging tests	<p>Imaging tests use X-rays, magnetic fields, sound waves, or radioactive substances to create pictures of the inside of the body. Imaging tests can be performed for a number of rationales, including to help find out if a suspicious area might be cancerous, to learn how far cancer has spread, to help determine if treatment has been effective. Most children who have or might have neuroblastoma will have one or more of these tests. Children with neuroblastoma are often very young, so it can be hard to perform some of these tests.</p> <p>Ultrasound</p> <p>Ultrasound is often one of the first tests done in small children if a tumour is suspected, because it is fairly quick and easy, it does not use radiation, and it can often give the doctor a good view inside the body, especially in the abdomen (belly). This test uses sound waves to create pictures of organs or masses inside the body. For this test, the child lies on a table (or sits) while a small wand called a transducer is placed on the skin over the belly (which is first lubricated with gel).</p>

Continued/...

Table 1.10: Contemporary procedures and approaches used in confirming a diagnosis of neuroblastoma (continued)

Procedure/Approach	Diagnostic Significance/Prognostic Value
Imaging tests (continued)	<p>Ultrasound (continued)</p> <p>The wand gives off sound waves and picks up the echoes as they bounce off organs. The echoes are converted by a computer into a black and white image on a screen. The test is not usually painful, but it might cause some discomfort if the transducer is pressed down hard on the belly. Ultrasound is used most often to look for tumors in the abdomen. It's not used to look in the chest because the ribs block the sound waves. Ultrasound can detect if kidneys have become swollen because the outflow of urine has been blocked by enlarged lymph nodes or a mass. It can also be used to help guide a biopsy needle into a suspected tumour to get a sample for testing. It is particularly useful in checking to see if tumours in the abdomen are shrinking. The pictures from ultrasound are not as detailed as those from some other tests, so even if a tumour is found, computed tomography (CT) or magnetic resonance imaging (MRI) scans (described below) might still be needed.</p> <p>X-rays</p> <p>The doctor may also order an X-ray of the chest or another part of the body as an early test if a child is having symptoms but it is not clear what might be causing them. But the images might not always be detailed enough to spot tumours. If neuroblastoma has already been diagnosed, X-rays can be useful to see if cancer has spread to certain bones. An X-ray of the head may be done to see if cancer has spread to the skull bones. A meta-iodobenzylguanidine (MIBG) scan or a bone scan (described below) is usually better for looking at the bones in the rest of the body, but X-rays may be used in infants, where these scans might not be possible. A standard chest X-ray may be done if doctors suspect that the tumour has invaded the lungs, but a CT or MRI scan of the chest can show the area in more detail.</p> <p>Computed tomography (CT or CAT) scan</p> <p>CT scans are often used to look for neuroblastoma in the abdomen, pelvis and chest. The CT scan is an X-ray test that produces detailed cross-sectional images of parts of the body. Instead of taking one picture, like a regular X-ray, a CT scanner takes many pictures as it rotates around the child while s/he lies on a table. A computer then combines these pictures into images showing slices of the part of the body being studied. Unlike a regular X-ray, a CT scan creates detailed images of the soft tissues in the body. Before the test, the child may be asked to drink a contrast solution and/or get an intravenous (IV) injection of a contrast dye. This helps better outline structures in the body. The contrast may cause some flushing (a feeling of warmth, especially in the face). Some people are allergic and get hives. Rarely, more serious reactions like laboured breathing or low blood pressure can occur. The doctor needs to ascertain if the child has any allergies or has ever had a reaction to any contrast material used for X-rays. CT scans take longer than regular X-rays. Younger children may be sedated before the test to reduce movement and help make sure the pictures come out well.</p>

Continued/...

Table 1.10: Contemporary procedures and approaches used in confirming a diagnosis of neuroblastoma (continued)

Procedure/Approach	Diagnostic Significance/Prognostic Value
Imaging tests (continued)	<p>CT-guided needle biopsy</p> <p>CT scans can also be used to help guide a biopsy needle into a tumour. For this procedure, the child lies on the CT scanning table while a radiologist advances a biopsy needle through the skin and toward the mass. CT scans are repeated until the needle is within the mass. A biopsy sample is then removed and looked at under a microscope. In children, this procedure is always done under general anaesthesia.</p> <p>Magnetic resonance imaging (MRI) scan</p> <p>MRI scans provide detailed images of soft tissues in the body. These scans are very helpful in looking at the brain and spinal cord. They may be slightly better than CT scans for seeing the extent of a neuroblastoma tumour, especially around the spine, but this test can be harder to do in small children. MRI scans use radio waves and strong magnets to create the images instead of x-rays, so there is no radiation. A contrast material called gadolinium may be injected into a vein before the scan to better see details, but this is needed less often than with a CT scan. It usually does not cause allergic reactions, but it can cause other problems in children with kidney disease, so doctors are careful when they use it. MRI scans take longer than CT scans, often up to an hour. For most MRI machines, the child has to lie inside a narrow tube, which is confining and can be distressing. Newer, more open MRI machines may be an option in some cases, but they still require the child to stay still for long periods of time. The MRI machine also makes loud buzzing and clicking noises that may be disturbing. Younger children are often given medicine to help keep them calm or even asleep during the test.</p> <p>Meta-iodobenzylguanidine (MIBG) scan</p> <p>This scan uses a form of the chemical meta-iodobenzylguanidine (MIBG) that contains a small amount of radioactive iodine. MIBG is similar to norepinephrine, a hormone produced by sympathetic nerve cells. It is injected into a vein and travels through the blood, and in most patients it will attach to neuroblastoma cells anywhere in the body. Several hours or days later, the body is scanned with a special camera to look for areas that incorporated the radioactivity. This helps doctors tell where the neuroblastoma is and whether it has spread to the bones and/or other parts of the body. This test is preferred by many doctors as a standard test in children with neuroblastoma. It can be repeated after treatment to see if it has been effective. It is also good to know if the tumour takes up the MIBG because in some cases, this radioactive molecule can be used at higher doses to treat the neuroblastoma.</p>

Continued/...

Table 1.10: Contemporary procedures and approaches used in confirming a diagnosis of neuroblastoma (continued)

Procedure/Approach	Diagnostic Significance/Prognostic Value
Imaging tests (continued)	<p>Positron emission tomography (PET) scan</p> <p>For a PET scan, a radioactive substance, usually a glucose analogue, known as fluorine-18-fluorodeoxy-glucose (18F-FDG) is injected into the blood. The amount of radioactivity used is very low and will pass out of the body within a day or so. Because cancer cells in the body are growing quickly, they absorb large amounts of the radioactive sugar. After about an hour, your child will be moved onto a table in the PET scanner. He or she will lie on the table for about 30 minutes while a special camera creates a picture of areas of radioactivity in the body. Younger children may be given medicine to help keep them calm or even asleep during the test. The picture from a PET scan is not as detailed as a CT or MRI scan, but it can provide helpful information about the whole body. Some newer machines can do a PET and CT scan at the same time (PET/CT scan). This lets the doctor compare areas of higher radioactivity on the PET scan with the more detailed appearance of that area on the CT scan.</p> <p>Bone scan</p> <p>A bone scan can help show if a cancer has spread to the bones, and can provide a picture of the entire skeleton at once. Neuroblastoma often causes bone damage, which a bone scan can find. This test used to be done routinely, but in some centres it has been replaced by use of MIBG or PET scans. For this test, a small amount of low-level radioactive material (technetium-99) is injected into a vein. (The amount of radioactivity used is very low and will pass out of the body within a day or so.) The substance settles in areas of damaged bone throughout the skeleton over the course of a couple of hours. Your child then lies on a table for about 30 minutes while a special camera detects the radioactivity and creates a picture of the skeleton. Younger children may be given medicine to help keep them calm or even asleep during the test. Areas of active bone changes attract the radioactivity and appear as “hot spots” on the skeleton. These areas may suggest cancer, but other bone diseases can also cause the same pattern. To help tell these apart, other imaging tests such as plain x-rays or MRI scans, or even a bone biopsy might be needed.</p>
Biopsies	<p>Examinations and tests might strongly suggest a child has neuroblastoma, but a biopsy (removing some of the tumour for viewing under a microscope and other lab testing) is often done to be sure. During a biopsy, the doctor removes a sample of the tumour mass. In adults, biopsies are sometimes done using local anaesthetic (numbing medicine), but in children they are more often done while the child is under general anaesthesia. There are 2 main types of biopsies:</p> <p>Incisional (open or surgical) biopsy</p> <p>This type of biopsy is done by removing a piece of the tumour through an incision (cut) in the skin. For tumours deep in the body this may be done laparoscopically using long, thin surgical tools inserted through small cuts in the skin.</p>

Continued/...

Table 1.10: Contemporary procedures and approaches used in confirming a diagnosis of neuroblastoma (continued)

Procedure/Approach	Diagnostic Significance/Prognostic Value
Biopsies (continued)	<p data-bbox="862 416 1093 440">Needle (closed) biopsy</p> <p data-bbox="862 461 2022 651">For this type of biopsy, a thin, hollow needle is placed through the skin and into the tumour to remove a small sample. If the tumour is deep within the body, CT scans or ultrasound can be used to help guide the needle into the tumour. The biopsy samples are sent to a lab, where they are viewed under a microscope by a pathologist (a doctor with special training in identifying cancer cells). Some neuroblastomas are easily recognized when looked at by experienced doctors. But some may be hard to tell apart from other types of child cancers. In these cases, special lab tests must be done to show the tumour is a neuroblastoma. Other lab tests may also be done on neuroblastoma samples to help determine how quickly the tumour is likely to grow.</p> <p data-bbox="862 671 1234 695">Bone marrow aspiration and biopsy</p> <p data-bbox="862 716 2022 1015">Neuroblastoma often spreads to the bone marrow (the soft inner parts of certain bones). If blood or urine levels of catecholamines are increased, then finding cancer cells in a bone marrow sample is enough to diagnose neuroblastoma (without getting a biopsy of the main tumour). If neuroblastoma has already been diagnosed by a biopsy done elsewhere in the body, bone marrow tests are done to help determine the extent of the disease. A bone marrow aspiration and biopsy are usually done at the same time. In most cases the samples are taken from the back of both of the pelvic (hip) bones. Even when the area is numbed with local anaesthetics, these tests can be painful, so in most cases the child is also given other medicines to reduce pain or even be asleep during the procedure. For a bone marrow aspiration, a thin, hollow needle is inserted into the bone and a syringe is used to suck out a small amount of liquid bone marrow. A bone marrow biopsy is usually done just after the aspiration. A small piece of bone and marrow is removed with a slightly larger needle that is pushed down into the bone. Once the biopsy is done, pressure is applied to the site to help stop any bleeding. Samples from the bone marrow are sent to a lab, where they are looked at and tested for the presence of cancer cells.</p>
<p data-bbox="197 1050 2022 1150">Source: <i>Neuroblastoma Summary</i>, American Cancer Society (http://www.cancer.org/). For detailed references, see text. Examples of more general references have been published previously.^{47,52,62} See also PDQ Screening and Prevention Editorial Board. Neuroblastoma Screening (PDQ®): Health Professional Version. 2014 Feb 6. In: PDQ Cancer Information Summaries [Internet]. Bethesda (MD): National Cancer Institute (US); 2002-. Available from: http://europepmc.org/books/NBK66025 and PDQ Pediatric Treatment Editorial Board. Neuroblastoma Treatment (PDQ®): Health Professional Version. 2016 Jan 14. In: PDQ Cancer Information Summaries [Internet]. Bethesda (MD): National Cancer Institute (US); 2002-. Available from: http://europepmc.org/books/NBK65747.</p>	

Depending on the clinical presentation, the index of suspicion must be high. Initial diagnostic testing should include complete blood count (CBC), serum electrolytes, liver function tests and a chest radiograph, which may reveal calcifications or a posterior mediastinal mass. Complementary diagnostic findings will encompass increased levels of urine or serum catecholamines or catecholamine metabolites such as dopamine, vanillylmandelic acid (VMA) and homovanillic acid (HVMA).^{63,64} Elevated levels of non-specific biomarkers such as lactate dehydrogenase (>1500 U/ml),⁶⁵ ferritin (>142 ng/ml),⁶⁶ and neuron-specific enolase (>100 ng/ml) may be correlated with advanced stage neuroblastoma and/or relapse.^{54,65,67-71}

A computed tomography (CT) and functional single-photon emission computed tomography (SPECT) or positron emission tomography (PET) scan of the neck, chest and abdomen is the gold standard for diagnostic imaging as it can concurrently focus the tumour and determine the degree of disease progression.^{48,50,59} Ultrasound may be used primarily to differentiate the tumour.^{48,72} Magnetic resonance imaging (MRI) may be beneficial if there is concern for spinal extension, and imaging of the brain is only necessary in the setting of neurological symptoms.^{54,55,73-76}

While not routinely used, a ^{123/131}I-radiolabelled meta-iodobenzylguanidine (MIBG) scan is valuable in both the detection of primary tumours and metastases since MIBG is a norepinephrine analogue that is selectively concentrated in sympathetic nervous tissue. MIBG has also proven exceptionally practical in surveillance of patient treatment responses and disease recurrence.^{54,55,57,76-80} While MIBG is generally more sensitive for the detection of lesions, fluorodeoxyglucose positron emission tomography (FDG-PET) may be better at localizing soft tissue metastases.^{50,56,57}

Despite advances in diagnostic medicine, the diagnosis of neuroblastoma can only be confirmed pathologically with tissue obtained from tumour or bone marrow. Specimens can be obtained either during resection of the primary tumour or as an open biopsy for unresectable

disease. Bilateral posterior iliac crest marrow aspirates are required to exclude metastatic disease. Molecular studies, such as fluorescence *in situ* hybridization (FISH), can be performed on tissue samples to note ploidy and other chromosomal aberrations.^{81,82} Recently, expression of the insulin-like growth factor-2 mRNA-binding protein 1 (*IGF2BP1*) gene has been found to be associated with more advanced tumours and decreased patient survival in neuroblastoma, suggesting its prognostic value.^{83,84}

A9.2 Histopathology of Neuroblastoma

Neuroblastoma, ganglioneuroblastoma and ganglioneuroma are classified as peripheral neuroblastic tumours (pNTs) which constitute a clinically and genomically complex disease. The pNTs represent significant disease models for analyzing the biologic and prognostic relationships between molecular/genomic alterations and accompanying morphological appearances. The International Neuroblastoma Pathology Classification (INPC) is particularly useful for patient stratification and protocol assignment in clinical trials of the Children's Oncology Group.⁸⁵⁻⁹¹ Table 1.11 summarizes the INPC classification of neuroblastic tumours and Table 1.12 shows a comparison between the categories and subtypes used in the INPC and the original Shimada classification.^{92,93}

Joshi and co-authors⁹⁴ advocated minor modifications to the terminology of the Shimada classification to include “borderline” ganglioneuroblastoma for the “stroma-rich, well-differentiated” subtype which is now called “ganglioneuroma (Schwannian stroma-dominant), maturing” subtype in the INPC. Morphologic confirmation of the Schwannian stroma-poor, stroma-rich, and stroma-dominant categories, as well as among subtypes in each category, may be challenging since the subtypes may express stages of a biologic and morphologic continuum. Likewise, macroscopic categorizing a nodular lesion of the composite tumour may be arduous.^{90,91,93} The INPC further differentiates between ‘favourable’ and ‘unfavourable’ histology groups⁹⁵ based on the age-linked morphological changes⁹⁶ (cut-offs of 18 and 60 months at diagnosis) by three major biologic/molecular mechanisms:

1. **Cross-talk between neuroblastic cells and Schwann cells** essential for tumour maturation; three categories, i.e., neuroblastoma (Schwannian stroma-poor), ganglioneuroblastoma-intermixed (Schwannian stroma-rich), and ganglioneuroma (Schwannian stroma-dominant), are defined;
2. **High-affinity nerve growth factor (NGF; TrkA/II)** expression critical for neuroblastic differentiation; 3 subtypes, i.e., undifferentiated, poorly differentiated, and differentiating, are defined in the neuroblastoma category; and
3. **MYCN amplification** as the powerful driving force for preventing neuroblastic differentiation and promoting mitotic and karyorrhectic activities; 3 classes of MKI (mitosis-karyorrhexis index), i.e., low <100/5000 cells, intermediate 100–200/5000 cells, and high >200/5000 cells, are defined in the neuroblastoma category. The INPC also includes the fourth category – ganglioneuroblastoma, nodular (composite, Schwannian stroma-dominant/stroma-rich and stroma-poor).^{85,97}

A new subtype of large nucleolar neuroblastoma (LNN) in the NB category have a characteristic nucleus containing large and prominent nucleoli, but do not show cytoplasmic enlargement/maturation. LCN can well be included in this group of LNN, as a large cell variant of undifferentiated/poorly differentiated NB.⁹⁸ Representative images of the histology of peripheral neuroblastic tumours are shown in Figures 1.4 and 1.5. Neuroblastoma predominantly comprises neuroblasts at different stages of differentiation and a varying amount of Schwannian-like stroma. The proportion of both cell types fluctuates according to the degree of tumour maturation and a correlation exists between the degree of differentiation of the neuroblastic subtype, the proportion of the Schwannian-like stroma and disease prognosis. Undifferentiated stroma-poor NB is the most malignant and the stroma rich ganglioneuroma (GN) is a benign form. The relationship between Schwannian-like stromal cells and neuroblastic cells needs further clarification.

Table 1.11: INPC classification of neuroblastic tumours

International Neuroblastoma Pathology Classification				
Category	Favorable Histology		Unfavorable Histology	
Neuroblastoma (Schwannian stroma-poor)	Grade of Differentiation	MKI	Grade of Differentiation	MKI
Age <18 months	poorly differentiate differentiating	low or intermediate low or intermediate	undifferentiated poorly differentiate differentiating	any high high
Age 18 – 60 months	differentiating	low	undifferentiating poorly differentiated differentiating	any any intermediate or high
Age ≥60 months	–	–	any	any
Ganglioneuroblastoma Intermixed (Schwannian stroma-rich)	all cases		–	–
Ganglioneuroma (Schwannian stroma-dominant)	all cases		–	–
Ganglioneuroblastoma, Nodular (composite, Schwannian stroma-rich/ stroma-dominant and stroma-poor)	favorable subset		unfavorable subset	

MKI: Mitosis-Karyorrhexis Index, Low <100/5,000 cells, Intermediate 100-200/5,000 cells, High >200/5,000 cells
Ganglioneuroblastoma, Nodular: Distinction of Favorable and Unfavorable subsets by the same criteria of age-appropriate evaluation of Grade of Differentiation and MKI as used for the Neuroblastoma category.

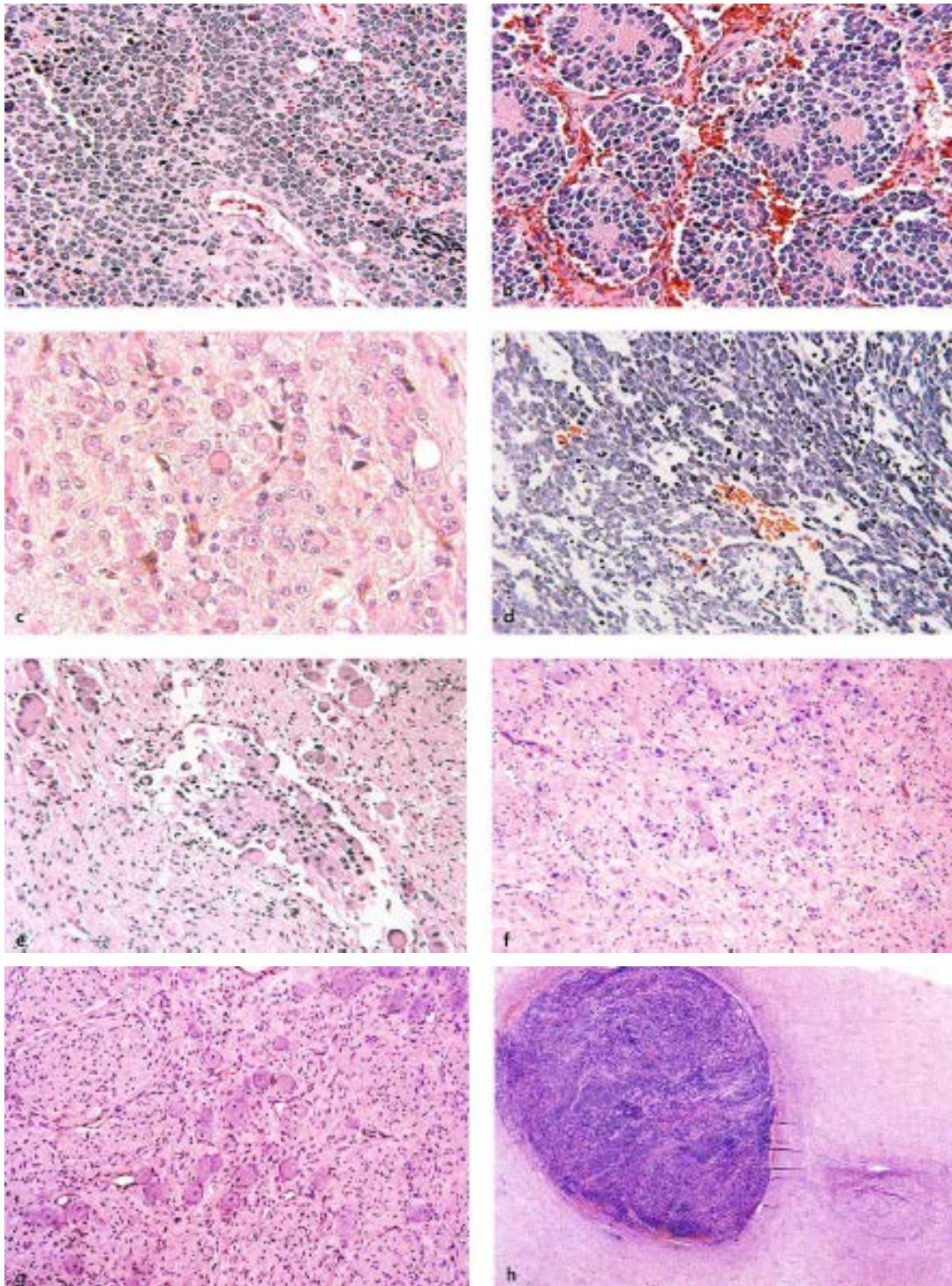
Source:⁸⁶

Table 1.12: Comparison of the INPC and the Shimada classification of neuroblastic tumours

International neuroblastoma pathology classification		
Schwannian (ganglioneuromatous) development	Category and subtype	Shimada classification
None to minimal	Neuroblastoma (Schwannian stroma-poor)	Stroma-poor
	Undifferentiated Poorly differentiated	Undifferentiated
None to minimal to <50% of the tumor tissue	Differentiating	Differentiating
	Ganglioneuroblastoma, intermixed (Schwannian stroma-rich) Ganglioneuroma, (Schwannian stroma-dominant)	Stroma-rich Intermixed
Dominant	Maturing ^a	Stroma-rich Well differentiated
	Mature	Ganglioneuroma
Proportion of Schwannian stroma-rich/stroma dominant and stroma-poor (nodular) areas, variable	Ganglioneuroblastoma, nodular (Composite Schwannian stroma-rich/stroma-dominant and stroma-poor)	Stroma-rich Nodular

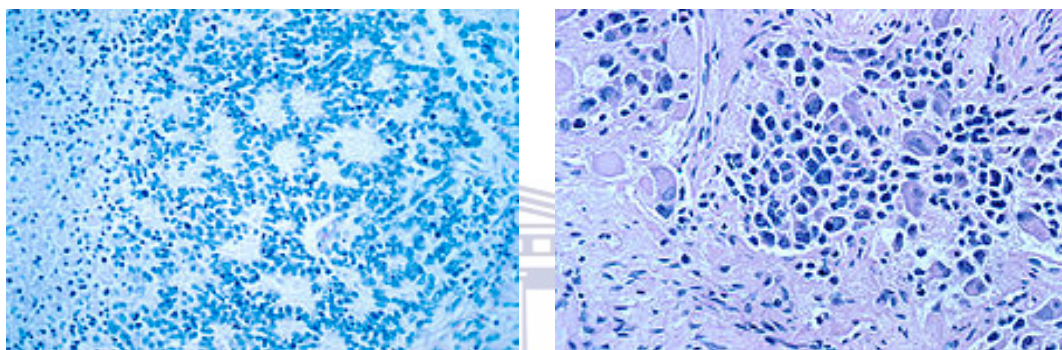
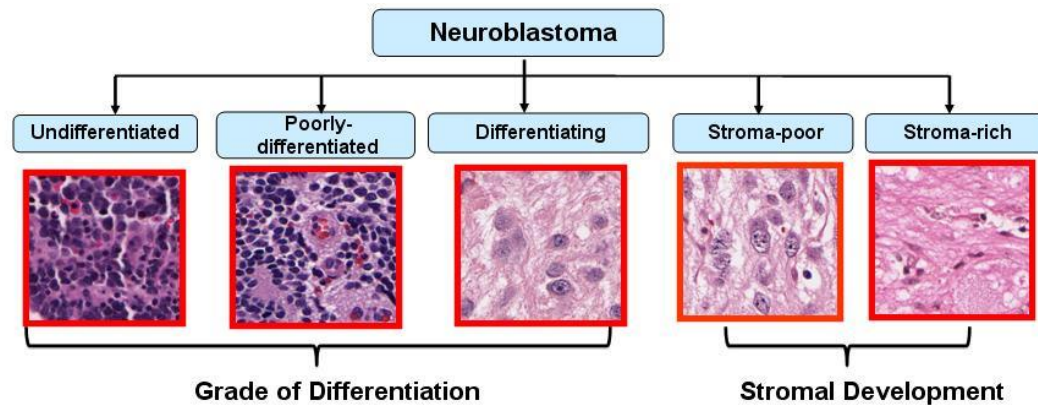
^a This subtype is termed "borderline" ganglioneuroblastoma, according to the Joshi classification.

Source:⁹⁰



Source:⁹¹ a Neuroblastoma (Schwannian stroma-poor), undifferentiated subtype. b Neuroblastoma (Schwannian stroma-poor), poorly differentiated subtype. c Neuroblastoma (Schwannian stroma-poor), differentiating subtype. d Neuroblastoma (Schwannian stroma-poor) with a high mitosis–karyorrhexis index. e Ganglioneuroblastoma, intermixed (Schwannian stroma-rich). f Ganglioneuroma (Schwannian stroma-dominant), maturing subtype. g Ganglioneuroma (Schwannian stroma-dominant), mature subtype. h Ganglioneuroblastoma, nodular (composite, Schwannian stroma-rich/stroma-dominant and stroma-poor).

Figure 1.4: Histology of peripheral neuroblastic tumours



Neuroblastoma with rosette formation (x40)

Stroma-rich ganglioneuroblastoma (x40)

Sources: <https://visualsonline.cancer.gov/details.cfm?imageid=2593>; On microscopy, the tumour cells are typically described as small, round and blue, and rosette patterns (Homer-Wright rosettes) may be seen. Homer-Wright rosettes are tumour cells around the neuropil, not to be confused with pseudorosettes, which are tumour cells around a blood vessel. Two typical true rosettes in retinoblastoma occur in the form of Flexner-Wintersteiner and Homer-Wright rosettes (<http://www.pathologystudent.com/?p=5400>; <https://en.wikipedia.org/wiki/Neuroblastoma#Histology>).^{99,100}

Figure 1.5: Microscopic views of typical neuroblastoma histopathology

The amount of Schwannian-like stroma in the tumour is associated with better prognosis since it is thought that Schwannian-like cells may downregulate tumour growth signal transduction pathways by secreting antiproliferative and/or antiangiogenic factors. In this regard, experimental evidence shows that when co-cultured *in vitro*, neuroblasts derived from neuroblastoma tumours can enhance the proliferation of Schwann cells. Schwannian-like stromal cells in neuroblastic tumours are likely to be reactive in nature and may have been recruited from normal tissue.^{95,101}

A10. Clinical Presentation, Signs and Symptoms of Neuroblastoma

The signs and symptoms of neuroblastoma vary widely, depending on the size of the tumour, where it is localized, how far it has spread, and if the tumour cells secrete hormones.⁶² Many of the signs and symptoms below are more likely to be caused by factors other than neuroblastoma. For example, a neuroblastoma may arise from sympathetic nervous tissue anywhere in the body, but most often develops in the abdomen. The presentation depends on the local effects of the solid tumour and any metastases. An abdominal mass in a child may also be due to Wilms' tumour (also known as nephroblastoma).⁷³ This neoplasm may present with renal signs and symptoms, such as hypertension,¹⁰² haematuria and abdominal pain.^{11,103-107} Multiple factors play a role in a patient's clinical presentation since it depends largely on tumour location, size, degree of invasion, effects from catecholamine secretion,¹⁰⁸ and symptoms due to paraneoplastic syndromes.^{109,110}

Nearly 65% of tumours arise in the abdomen with half of those localized to the medulla of the adrenal gland.^{59,111-113} However, they can occur in the neck (5%), chest (20%), or pelvis (5%), and 1% of patients have no detectable primary tumours.¹¹⁴ Many patients are asymptomatic, yet some may present with constitutional symptoms (malaise, fevers, and weight loss), an enlarging mass, pain, abdominal distension, lymphadenopathy, or respiratory distress secondary to compression or hepatomegaly. Pelvic masses may cause constipation or difficulty urinating, while thoracic involvement can cause dysphagia, dyspnoea, or rarely, thoracic outlet syndrome. For cervical tumours, a patient develop Horner's syndrome,⁵² and in up to 15% of patients, epidural extension may result in neurological deficits such as progressive paralysis.^{1,11,37,46,115-117}

At the time of diagnosis, 50% of patients present with localized disease while 35% already have regional lymph node spread. Metastasis can occur by haematogenous and/or lymphatic route, seeding bone marrow,¹¹⁸ liver, and bone. Neuroblastoma originating from cells of the primitive neural crest eventually populates the sympathetic ganglia and the inner adrenal gland.

In approximately one-half of cases, the primary tumour arises at the level of the paravertebral ganglia and may infiltrate the adjacent intervertebral foramina and compress the intraspinal structures. Although modern imaging studies document the infiltration of the intervertebral foramina by the tumour in at least one-third of neuroblastomas, only 5–7% of the cases develop symptoms related to epidural compression (EC).¹¹⁹

In these instances, various neurological deficits can ensue which may progressively worsen, and could end in paraplegia.⁷⁴ The signs of EC are difficult to detect in an early phase, especially among the youngest children, and this may account for the frequent delay in diagnosis and thus lead to development of permanent neurological impairment. Several studies have analyzed the outcome of various types of treatment for EC. Neurosurgical decompression, chemotherapy and radiation therapy have all proven to be effective in relieving the symptoms. Although the occurrence of short-term sequelae were reported in some publications, only one study addressed the issue of the authors found that the majority of children actually did recover normal neurological function, but they developed an excess of spinal deformities, in particular when treatment included laminectomy.¹¹⁹⁻¹²¹

Commonly, the orbits are involved, which manifests as periorbital swelling and proptosis (“raccoon eyes”).⁶⁴ When dissemination occurs to the skin, patients develop blue subcutaneous nodules known as blueberry muffin syndrome. Surprisingly, this is associated with a favourable prognosis with likely spontaneous tumour regression. Because of its neuroendocrine properties, neuroblastoma has the potential to secrete catecholamines, which results in early-onset hypertension and tachycardia.^{108,122} Patients may also experience paraneoplastic syndromes.^{110,122} Examples include intractable diarrhoea with electrolyte disturbances due to release of vasoactive intestinal peptide (VIP),¹²³ encephalomyelitis, or sensory neuropathy. There have been reports of the development of opsoclonus-myoclonus syndrome (OMS),^{115,124} which occurs when antibodies cross-react with cerebellar tissue.^{1,125-127} The characteristic symptoms and signs of OMS include rapid, conjugate eye nystagmus

with involuntary spasms of the limbs. Interestingly, the patients with intractable diarrhoea due to VIP secretion or OMS generally tend to present with less aggressive neuroblastomas. Thus far, symptomatic paraneoplastic syndromes are rarely diagnosed (prevalence < 0.01% of all cancer), but they may indicate early signs of disease relapse.^{1,128}

A10.1 Signs or Symptoms Caused by the Main Tumour

A10.1.1 Tumours in the Abdomen or Pelvis

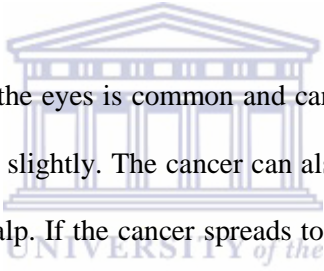
One of the most common signs of a neuroblastoma is a large lump or swelling in the child's abdomen.¹²⁹ The child might not want to eat (which can lead to weight loss). If the child is old enough, s/he may complain of feeling full or having abdominal pain. But the lump itself is usually not painful to the touch. Sometimes, a tumour in the abdomen or pelvis can affect other parts of the body. For example, tumours that press against or grow into the blood and lymph vessels in the abdomen or pelvis can stop fluids from getting back to the heart. This can sometimes lead to swelling in the legs and, in boys, the scrotum. In some cases, the pressure from a growing tumour can affect the child's bladder or bowel, which can cause problems urinating or having bowel movements.

A10.1.2 Tumours in the Chest or Neck

Tumours in the neck can often be seen or felt as a hard, painless lump. If the tumour is in the chest, it might press on the superior vena cava. This can cause swelling in the face, neck, arms, and upper chest (sometimes with a bluish-red skin colour). It can also cause headaches, dizziness, and a change in consciousness if it affects the brain. The tumour might also press on the throat or windpipe, which can cause coughing and troubled breathing (dyspnoea) or swallowing. Neuroblastomas that press on certain nerves in the chest or neck can sometimes cause other symptoms, such as a drooping eyelid and a small pupil (the black area in the centre of the eye). Pressure on other nerves near the spine might affect the child's ability to feel or move their arms or legs.

A10.2 Signs or Symptoms Caused by Metastatic Spread of the Cancer

About 2 out of 3 neuroblastomas have already spread to the lymph nodes or other parts of the body by the time they are found. Lymph nodes are bean-sized collections of immune cells found throughout the body. Cancer that has spread to the lymph nodes can cause them to swell. These nodes can sometimes be felt as lumps under the skin, especially in the neck, above the collarbone, under the arm, or in the groin. Enlarged lymph nodes in children are much more likely to be a sign of infection rather than cancer, but they should be checked by a doctor. Neuroblastoma often spreads to bones. A child who can talk may complain of bone pain. The pain may be so bad that the child limps or refuses to walk. If it spreads to the bones in the spine, tumours can press on the spinal cord and cause weakness, numbness, or paralysis in the arms or legs.



Spread to the bones around the eyes is common and can lead to bruising around the eyes or cause an eyeball to stick out slightly. The cancer can also spread to other bones in the skull, causing bumps under the scalp. If the cancer spreads to the bone marrow, the child may not have enough red blood cells, white blood cells, or platelets. These shortages of blood cells can result in tiredness, irritability, weakness, frequent infections, and excessive bruising or bleeding from small cuts or scrapes. Rarely, large tumours can start to break down, leading to a loss of clotting factors in the blood. This can result in a high risk of serious bleeding, which is known as a consumption coagulopathy and can be life threatening.

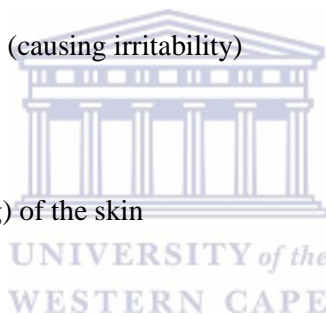
A special widespread form of neuroblastoma (known as stage 4S)⁹⁷ occurs only during the first few months of life. In this special form, the neuroblastoma has spread to the liver, to the skin, and/or to the bone marrow (in small amounts). Blue or purple bumps that look like small blueberries may be a sign of spread to the skin. The liver can become very large and can be felt as a mass on the right side of the belly. Sometimes it can grow large enough to push up on the lungs, which can make it hard for the child to breathe. Despite the fact that the cancer is already widespread when it is found, stage 4S neuroblastoma is very treatable, and often

shrinks or regresses spontaneously. Almost all children with this form of neuroblastoma can be cured.^{12,97,130}

A10.3 Signs or Symptoms Caused by Hormones Secreted by the Tumour

Neuroblastomas sometimes release hormones that can cause problems with tissues and organs in other parts of the body, even though the cancer has not spread to those tissues or organs. These problems are called paraneoplastic syndromes as described above. Symptoms of paraneoplastic syndromes can include:

- ❧ Constant diarrhoea
- ❧ Fever
- ❧ High blood pressure (causing irritability)
- ❧ Rapid heartbeat
- ❧ Reddening (flushing) of the skin
- ❧ Sweating



An uncommon set of symptoms is called the opsoclonus-myoclonus-ataxia syndrome (OMS) or “dancing eyes, dancing feet.”^{52,64} The child has irregular, rapid eye movements (opsoclonus), twitch-like muscle spasms (myoclonus), and appears uncoordinated when standing or walking (ataxia). S/he may also have trouble speaking. For reasons that are not clear, neuroblastomas that cause this syndrome tend to be less life-threatening than other forms of the disease.

A11. Molecular Pathogenesis, Genetics and Genomics of Neuroblastoma

Two major causes have been identified in the origin of NB, namely, (i) *familial origin* which is identified in the loss-of-function mutation in the *PHOX2B* gene and (ii) *sporadic origin* which results in chromosomal losses.¹³¹ These will be explained in the subsections.

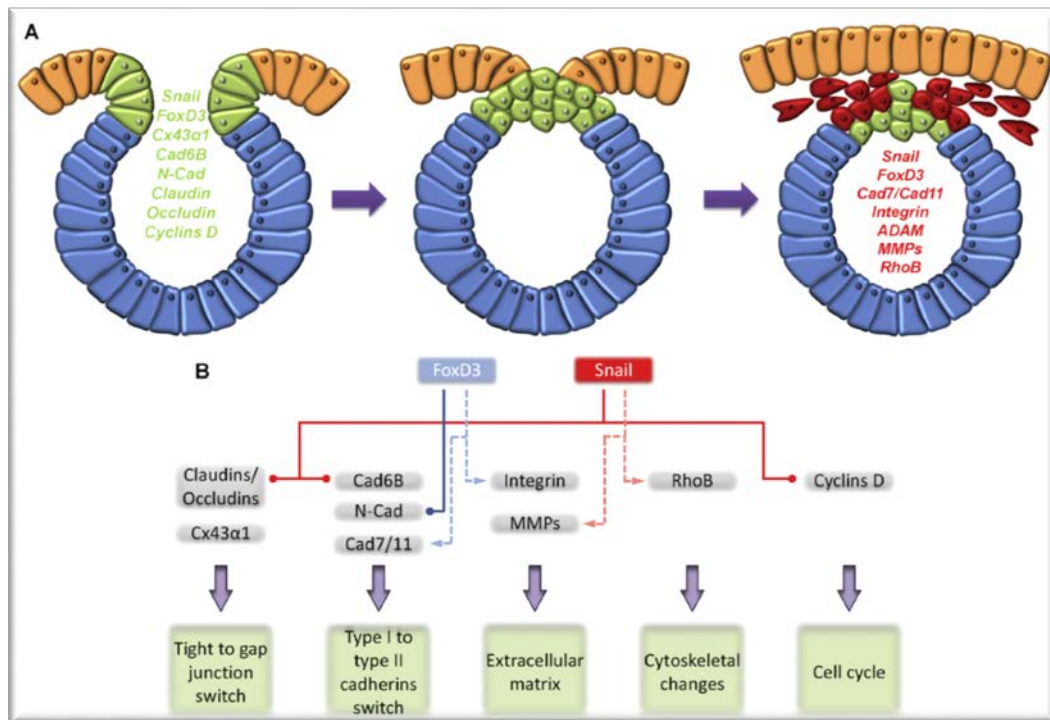
A11.1 Neural Development and Neuroblastoma

In considering neural development, the contribution of the neural crest to sympathetic ganglia and the adrenal gland is important. The majority of NB tumours appear to arise from neural crest-derived cells in the abdomen adjacent to the aorta in the region of the kidney or in the medullary region of the adrenal gland.^{6,132,133} Thus, NB is a sympathoadrenal lineage neural crest-derived tumour.¹³⁴ The neural crest arises from the dorsal region of the closing neural tube beneath the ectoderm.¹³⁵ This transient population of cells produces multipotential progenitor cells that give rise to the peripheral nervous system, the enteric nervous system, pigment cells, Schwann cells, adrenal medullary cells, and cells of the craniofacial skeleton.¹³⁵

This process is regulated by both extrinsic and intrinsic factors. The Hedgehog and Wnt signalling pathways are especially crucial for proper neural crest development.^{5,135,136} Lineage studies in the developing embryo have shown that neural crest cells within the trunk region generate multiple neural crest derivatives such as melanocytes, Schwann cells, glia, and neurons of the dorsal root ganglia. A subset of these trunk crest cells, commonly referred to as the sympathoadrenal lineage, contributes to the sympathetic ganglia and medullary region of the adrenal gland. This lineage of cells is thought to be the origin of NB.^{132,133} However, given the fact that NB can develop anywhere along the sympathetic axis, it is likely that NB can also arise from earlier crest derivatives, before development of the sympathoadrenal lineage but after the initial fate specification. This could contribute to the heterogeneous histology and pathology of NB.¹³⁷

A11.2 EMT and MET Transitions in the Neural Crest

During maturation, the neural crest undergoes programmed epithelial-to-mesenchymal transition (EMT).^{138,139} Figure 1.6 is a schematic representation of this process. The progression of NC EMT is synchronized by (i) the coordinated activity of transcription factors and molecular signaling pathways, (ii) changes in cell junctions and polarity, (iii) changes in adhesion properties, and (iv) changes in the extracellular matrix (ECM).



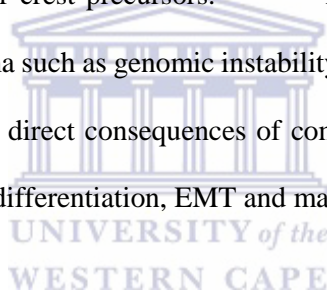
Source:¹³⁹ (A) Genes expressed on neural crest cells—prior (green) and after (red)—epithelial-to-mesenchymal transition. (B) Neural crest epithelial-to-mesenchymal transition regulation. NC specifiers, FoxD3 and Snail down-regulate expression of molecules that are associated with epithelial static cell populations, such as N-Cad and E-Cad (or Cad6B in chick and mouse), respectively, to relinquish space to the upregulation of mesenchymal migratory proteins, such as Cad7. Similarly, Snail down-regulates tight junction claudins/occludins to permit the upregulation of gap junction protein connexin-43 α 1 (Cx43 α 1), which may also depend on Snail expression. Gene regulation in which the repressors Snail or FoxD3 up-regulate the expression of matrix metalloproteases (MMPs), integrins, Cad7 or RhoB may denote indirect regulatory interactions, possibly mediated by other repressors (denoted by dotted lines). Reproduced from Strobl-Mazzulla PH, Bronner ME. Epithelial to mesenchymal transition: New and old insights from the classical neural crest model. *Seminars in Cancer Biology* 2012;22(5-6):411-416, with permission from Elsevier®. See Appendix 3 for copyright clearance.

Figure 1.6: Neural crest programmed epithelial-to-mesenchymal transition

Signalling pathways activated during the course of EMT in the NC are triggered by the integration of ECM signalling molecules and any number of secreted ligands such as members of transforming growth factor beta (TGF β), wingless/integrated proto-oncogene (Wnt) and fibroblast growth factor (FGF) families. These early cellular changes during EMT are essential for the switch from neuroepithelial precursors into migratory NC cells through activation and coordination of several transcriptional regulators, including the zinc finger transcription factors, Snail1 and Snail2 (formerly known as Slug) and the winged-helix transcription factor FoxD3.^{139,140} During embryonic development, mesenchymal transformation involves, among other processes, loss of E-cadherins, loss of cell contacts, activation of matrix

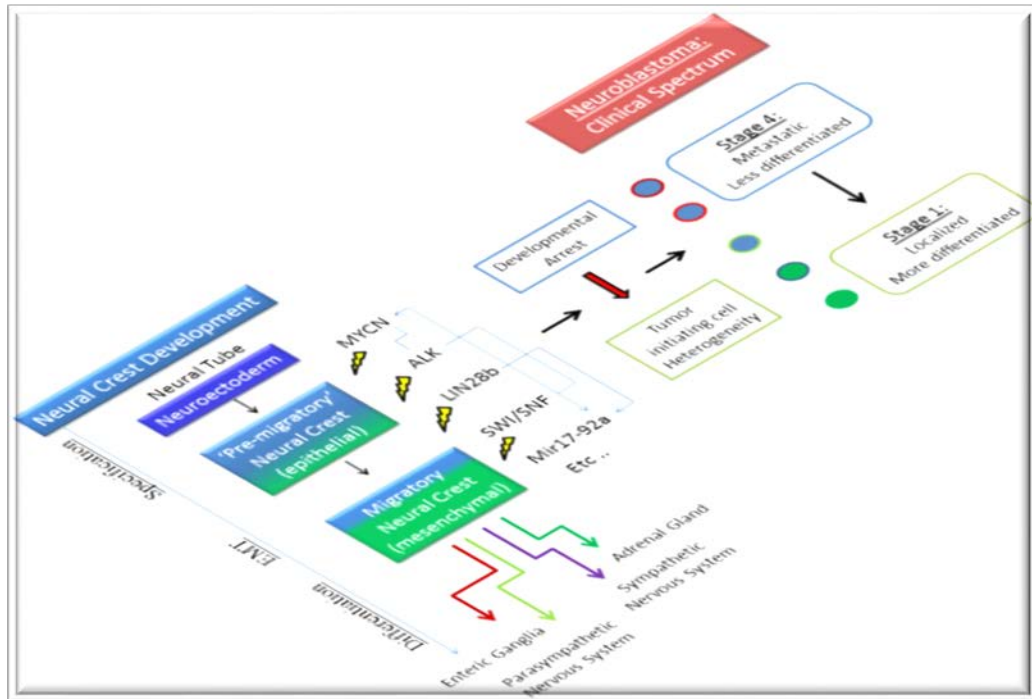
metalloproteinases (MMPs). Bone morphogenetic proteins (BMP, multi-functional growth factors that belong to the TGF β superfamily), Wnt and FGF signalling within the microenvironment further drive differentiation of these mesenchymal migratory NC cells. The early neural crest is similar to other pluripotent cell populations with regard to their committed self-renewal capacity and selective propensity to generate many different tissue types. Expression of pro-survival and pluripotency factors such as SOX10, FOXD3, C-Myc and MYCN confer on these cells an increased proliferative advantage coupled to an aggressive apoptosis evasive potential.^{138,141}

Neuroblastoma tumour-initiating cells (TICs) or cancer stem cells (CSCs) derived from diverse environments may direct the clonal evolution of distinct tumour phenotypes according to the developmental stage of their crest precursors.^{6,19,142-144} It is likely that the clinicopathologic correlations of neuroblastoma such as genomic instability, tumour heterogeneity and disparate treatment outcomes may be direct consequences of complex molecular signalling pathways that coordinate neural crest differentiation, EMT and maturation/specialization (Figure 1.7).



A11.3 Hallmarks of the Neuroblastoma Tumour Microenvironment

Drawing on the hallmarks of cancer expounded by Douglas Hanahan and Robert A. Weinberg,^{145,146} a recent article reviewed how an integrated biological systems repertoire, encompassed by the tumour microenvironment (TME), regulates tumour progression and metastasis in NB. The authors views converge on the respective contributions of innate [TAMs, neutrophils, natural killer cells (NK), dendritic cells (DC)] and adaptive (T- and B-lymphocytes, and natural killer T cells (NKT)] immune cells, tumour-associated fibroblasts (TAFs), bone marrow-derived mesenchymal stromal cells (MSCs), endothelial cells, Schwann cells, and the extracellular matrix (ECM). Neuroblastoma cells exploit the cell-cell and cell-ECM communication apparatus to “instruct” the TME and TME cells to activate neuroblastoma signalling pathways to express and maintain their neoplastic behaviour (Figures 1.8 and 1.9).¹⁴⁷



Neuroblastoma is a spectrum of diseases with a wide range of clinical behaviours. Disruption of the normal maturation progression with different genetic drivers at different times leads to heterogeneity of tumour-initiating cells. Interaction between different epigenetic and genetic factors complicates the task of defining a primary oncogenic driver or pathway for this disease. This results in a wide range of pathologies with highly variable responses to treatment.

Source:⁶ Reproduced and adapted from Louis CU, Shohet JM. Neuroblastoma: Molecular pathogenesis and therapy. *Annual Review of Medicine* 2015;66:49-63, permission not required as stipulated by the Annual Review of Medicine. See Appendix 4 for copyright clearance.

Figure 1.7: Clinicopathologic correlations of neuroblastoma

The ten hallmarks of cancer¹⁴⁵⁻¹⁴⁷ are the ability of cancer cells to:

- ☞ Sustain proliferative signals
- ☞ Evade growth-suppressors
- ☞ Invade and metastasize
- ☞ Enable replicative immortality
- ☞ Induce angiogenesis
- ☞ Resist cell death or apoptosis
- ☞ Escape immune destruction

- ☞ Deregulate cellular metabolism
- ☞ Confer and express genomic instability
- ☞ Induce tumour-promoting inflammation

Since these hallmarks of NB have been discussed thoroughly in the cited references in the context of their contribution to the neuroblastoma malignant phenotype and current clinical trials that target the TME in neuroblastoma patients,^{147,148} no further consideration will be accorded to them in this section. An essential description of neuroblastoma oncogenic drivers and transcriptional networks is provided in the next section.

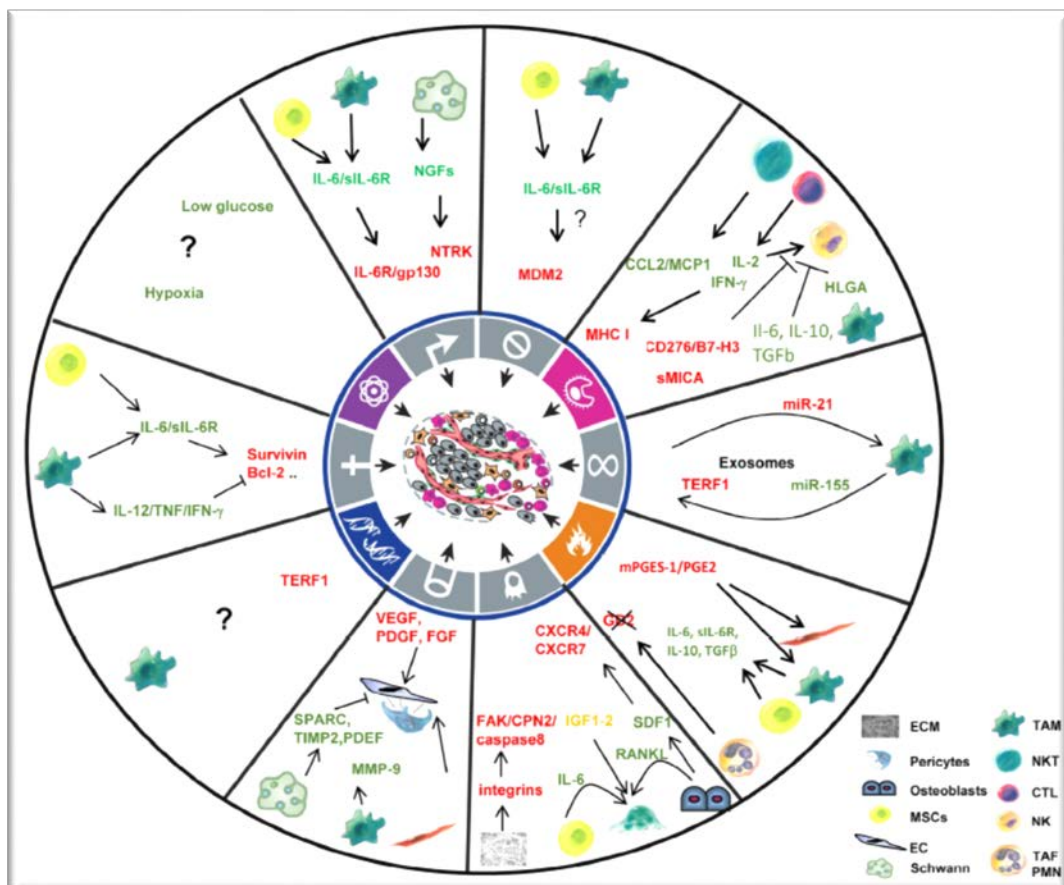
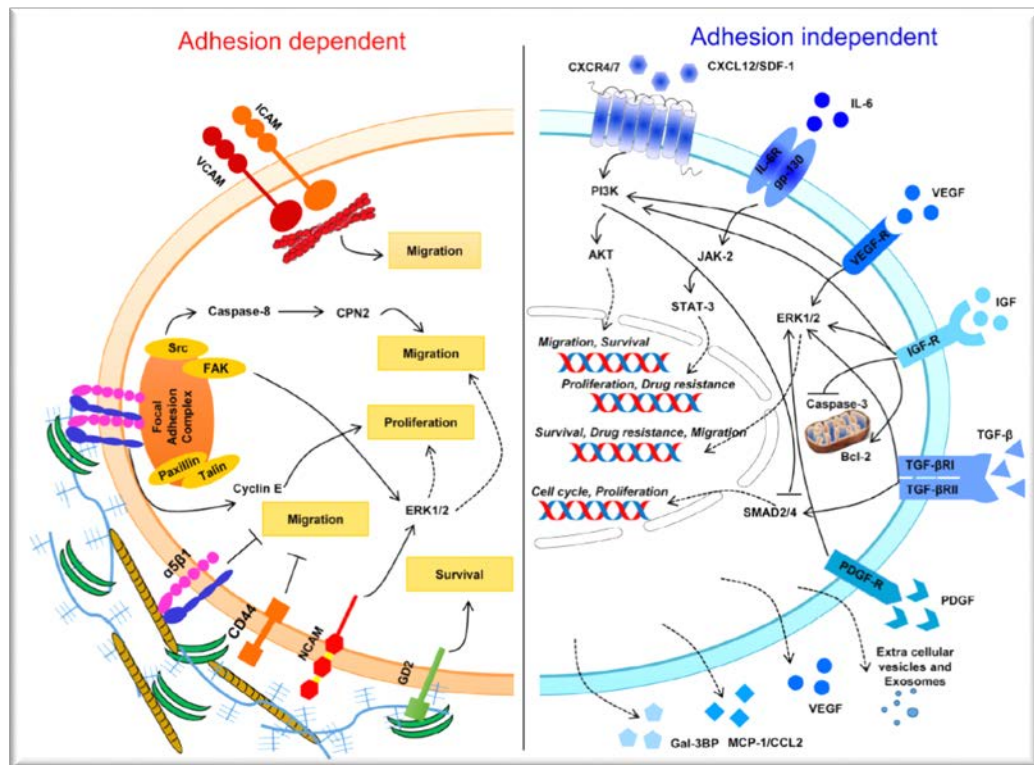


Diagram summarizing the contribution of the cells and ECM in the TME to the ten hallmarks of cancer shown at the centre of the wheel. The central graph was reproduced from Hanahan and Weinberg.¹⁴⁶

Source:¹⁴⁷ Reproduced from Borriello L, Seeger RC, Asgharzadeh S, DeClerck YA. More than the genes, the tumour microenvironment in neuroblastoma. *Cancer Letters* 2015;doi: 10.1016/j.canlet.2015.11.017, with permission from Cancer Letters, Elsevier Ireland Ltd. See Appendix 5 for copyright clearance.

Figure 1.8: Contribution of the cells and ECM in the TME to the ten hallmarks of neuroblastoma



Source:¹⁴⁷ Reproduced from Borriello L, Seeger RC, Asgharzadeh S, DeClerck YA. More than the genes, the tumour microenvironment in neuroblastoma. *Cancer Letters* 2015;doi: 10.1016/j.canlet.2015.11.017, with permission from Cancer Letters, Elsevier Ireland Ltd. See Appendix 5 for copyright clearance.

Figure 1.9: Pathways activated via communication between neuroblastoma and TME cells in the ECM

WESTERN CAPE

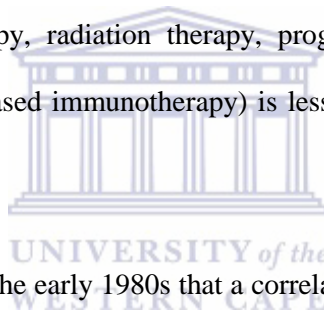
A11.4 Genetic Lesions, Transcriptional Networks and Oncogenic Drivers in Neuroblastoma

Somatic alterations, including mutations, gain of alleles, loss of alleles, or conversions in tumour-cell ploidy, have long been regarded as critical factors in the development and progression of NB. Many of these chromosomal aberrations are strong prognostic markers that can be used separate from clinical traits in risk stratification and treatment of NB patients.^{93,137,149,150} Chromosome regions and genes known to be involved in NB oncogenesis is outlined schematically in Figure 1.10. Some of these gene expression profiles of NB are described in the subsections that follow.

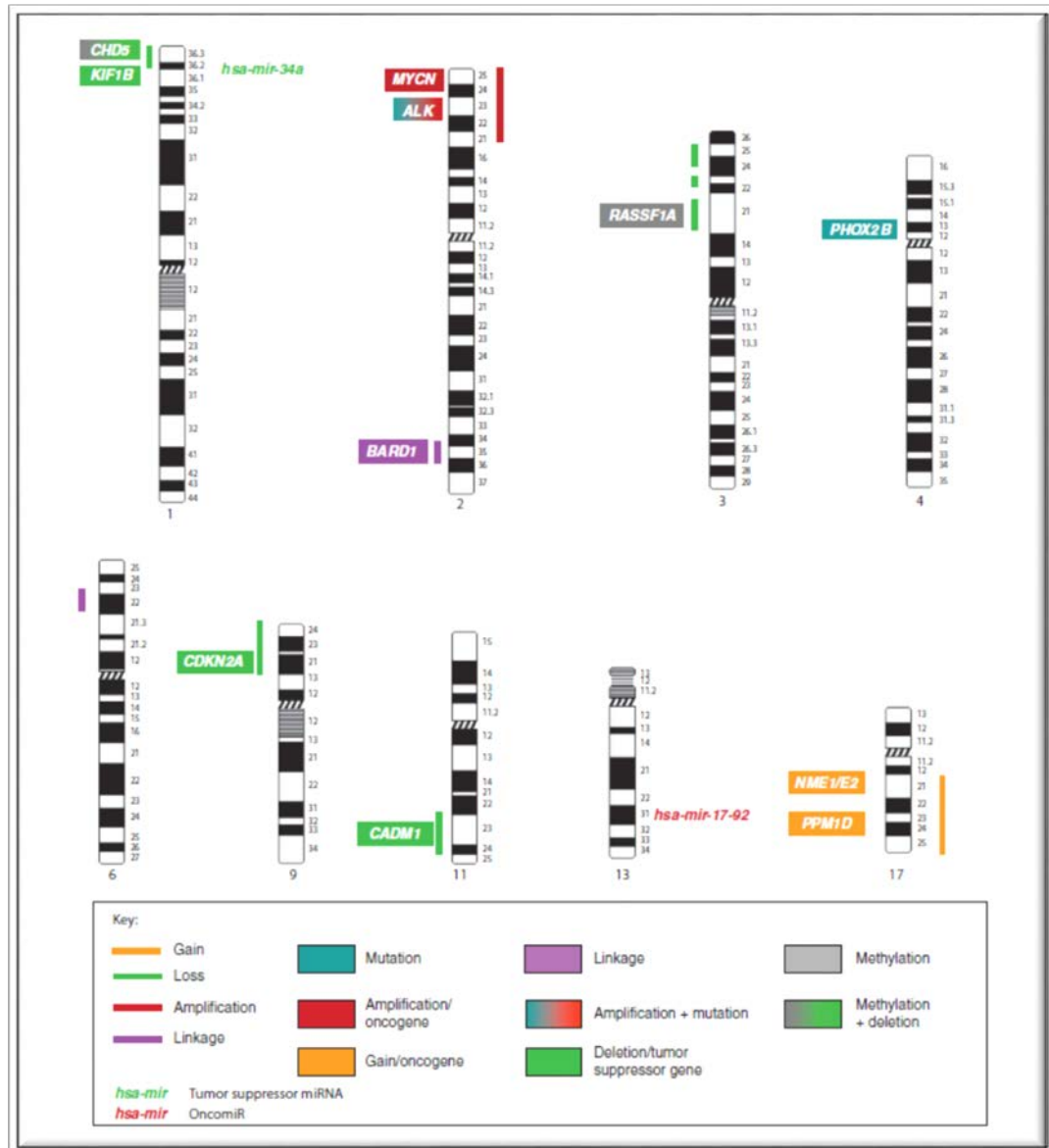
A11.4.1 Familial Genetic Lesions

Neuroblastoma originates from neuroepithelial cells that migrate from the neural crest to form

the sympathetic nervous system.¹⁵¹ Even though intratumour heterogeneity represents the predictable stage of oncogenesis,¹⁴³ it is well-established that neuroblastoma is not exclusively initiated by gene signatures.¹⁴⁷ As such, hereditary NB is both rare and heterogeneous, accounting for less than 5% of all NBs. Hereditary NB predisposition loci have been mapped to chromosomes 16p12–13 and 4p16, indicating that other familial predisposition mutations may exist, but hitherto no specific genes have been unequivocally shown to be inactivated or mutated in these chromosomal regions.^{133,137} Neuroblastoma presents as a locoregional tumour with no detectable amplification of the *MYCN* oncogene in 50% of children, but it correlates with extraordinary prognosis—overall survival (OS)>90%.¹⁴⁷ By contrast, in children older than 18 months of age diagnosed with NB, with or without *MYCN* amplification and metastatic disease, the chance of event-free long term survival despite an intensive combination therapy (myeloablative chemotherapy, radiation therapy, progenitor cell transplantation, surgery, isotretinoin and antibody-based immunotherapy) is less than 50%, and is thus indicative of high-risk disease.^{152,153}



The landmark discovery in the early 1980s that a correlation exists between *MYCN* oncogene amplification and advanced stage NB raised hopes that other similar genetic associations may be identified.¹⁵⁴ A family history of NB occurs in 1–2% of patients and 2 mutated genes have been distinguished as promising cancer biomarkers in this regard—anaplastic lymphoma kinase (ALK, exemplifying a gain of function) and the paired-like homeobox 2B (PHOX2B, denoting a loss of function)—in 80% of the familial cases.^{155,156} The advent of genome-wide association studies yielded additional gene polymorphisms with a low, but significant risk of NB, including *BARD1*, *LMO1* and *LIN28B*.¹⁵⁷ Genomic analysis of over 200 NBs showed, unpredictably, low levels of recurrent-driver mutations, most notably activation mutation and amplification of *ALK* (8% of the cases), activation mutations in *PTPN11* (a tyrosine phosphatase), inactivating mutations in chromatin remodelling genes (*ATRX* and *ARID1A*) and activating mutations in *NRAS*, in addition to amplification and activation mutations of *MYCN*.^{147,158-160}



A schematic overview of chromosome regions and genes known to be involved in neuroblastoma oncogenesis. This overview is not comprehensive, and only those regions and genes mentioned in the article are indicated. Gene abbreviations: ALK, anaplastic lymphoma receptor tyrosine kinase; BARD1, BRCA1 associated RING domain 1; CADM1, cell adhesion molecule 1; CDKN2A, cyclin dependent kinase inhibitor 2A; CHD5, chromodomain helicase DNA binding protein 5; KIF1B, kinesin family member 1B; MYCN, v-myc myelocytomatosis viral related oncogene, neuroblastoma derived; NME1/E2, non-metastatic cells 1, protein (NM23A) expressed in/non-metastatic cells 2, protein (NM23A) expressed in; PHOX2B, paired-like homeobox 2b; PPM1D, protein phosphatase 1D magnesium-dependent, delta isoform; RASSF1A, Ras association (RalGDS/AF-6) domain family member 1.

Source:¹⁵⁰ Van Roy N, De Preter K, Hoebeek J, Van Maerken T, Pattyn F, Mestdagh P, Vermeulen J, Vandesompele J, Speleman F. The emerging molecular pathogenesis of neuroblastoma: Implications for improved risk assessment and targeted therapy. *Genome Medicine* 2009;1(7):74, with permission from BioMed Central (BMC) Reprints and Permissions (<http://www.biomedcentral.com/about/policies/reprints-and-permissions> [14/06/2016 16:35:15]). See Appendix 6 for copyright clearance.

Figure 1.10: Chromosome regions and genes known to be involved in neuroblastoma oncogenesis

Evaluation of a subset of patients with MYCN non-amplified NB showed that infiltration of tumour-associated macrophages (TAMs) was significantly higher in metastatic NBs than their locoregional equivalents. In addition, metastatic tumours diagnosed in patients at age ≥ 18 months had higher expression of inflammation-related genes than those in patients diagnosed at age < 18 months. Expression of genes representing TAMs (CD33/CD16/IL-10/FCGR3) in addition to IL-6 receptor (IL-6R) influenced 25% of the accuracy of a novel 14-gene tumour classification score.¹⁶¹ Moreover, infiltration with Th2-driven macrophages expressing CD163 and CD206 was also recently observed in a subset of high-risk neuroblastoma tumours with deletion of chromosome 11q and high levels of prostaglandin-synthase and elevated levels of PGE2.¹⁶²

A11.4.2 PHOX2B Germline Mutations

Germline mutations in the paired-like homeobox 2B (PHOX2B) gene on chromosome 4p13 are the first predisposition mutations identified in NB.^{156,163,164} PHOX2B, a *master regulator* of sympathetic neuronal development and mainly expressed in sympathetic neural progenitors,¹⁶⁵ as well as mammalian achaete scute homolog-1 (MASH1), are expressed early in the developing sympathoadrenal progenitors. Shortly after expression of MASH1 and PHOX2B in the sympathoadrenal lineage, heart- and neural crest derivatives-expressed protein 2 (HAND2), PHOX2A, and GATA2/3 appear. PHOX2B has also been shown to be essential for the expression of the glial family ligand tyrosine kinase coreceptor RET (rearranged during transfection) and for the specification of noradrenergic fates, particularly the biosynthetic enzymes tyrosine hydroxylase (TH) and dopamine beta-hydroxylase promoter (DBH).¹³⁷

NB patients with PHOX2B mutations also have familial disorders of the neural crest such as Hirschsprung's disease (HSCR) and congenital hypoventilation syndrome.^{156,163} It is not clear whether the mutations in PHOX2B found in familial NB result in gain or loss of function, although many PHOX2B mutations stabilize the PHOX2B protein and decrease or eliminate the ability of PHOX2B to transactivate the DBH promoter.^{166,167} The findings that PHOX2B is

necessary for the differentiation of autonomic neurons and overexpression of PHOX2B inhibits proliferation in neuron progenitors and cell lines suggests PHOX2B is a tumour suppressor.^{166,167} However, the absence of tumours with loss of heterozygosity (LOH) or mutation in second allele suggests gain-of-function, dominant negative effect, or haploinsufficiency.¹⁶⁴

A11.4.3 Anaplastic Lymphoma Kinase

Anaplastic lymphoma kinase (ALK) is a member of receptor tyrosine kinases (RTKs) and was first identified as a part of the fusion gene nucleophosmin (NMP)–ALK in anaplastic large cell lymphoma via chromosome translocation of t(2;5)(p23;q25).¹⁶⁸ ALK is thought to play a role in the normal development of the central and peripheral nervous system since ALK mRNA is expressed throughout the nervous system in mouse and rat, but is not present in normal haematopoietic cells.^{169,170} Similar patterns of expression are observed in humans although additional ALK transcripts of differing size, most likely due to alternative splicing, have been observed in colon, prostate, testis, small intestine, and brain of adults.^{137,171} Full-length ALK protein is comprised of an extracellular region and an intracellular region containing a RTK domain, linked by a transmembrane (TM)-spanning segment, whereas the NMP–ALK fusion protein generated as a result of the t(2;5)(p23;q25) translocation contains the N-terminal of NMP and C-terminal kinase domain of ALK. Translocation of the gene is also evident in other tumours, such as inflammatory myofibroblastic tumour (IMT), and non-small-cell lung carcinoma (NSCLC), but not in NB.¹⁷¹

Overexpression of wild-type ALK has also been observed in thyroid carcinoma, breast cancer, NB, melanoma, small cell lung carcinoma, glioblastoma, astrocytoma, retinoblastoma, Ewing sarcoma, and rhabdomyosarcomas NB.¹⁷¹⁻¹⁷³ During 2008, several reports focused attention on ALK point mutations in 8–12% of all NB patients (both hereditary and sporadic) and some NB cell lines.¹⁷⁴⁻¹⁷⁷ Almost all the point mutations identified occurred in the kinase domain and resulted in the constitutive activation of ALK. Two of these activating ALK mutants were able

to transform NIH3T3 fibroblasts and induce tumour formation in nude mice.¹⁷⁸ In addition, knockdown of ALK or small molecular ALK inhibitors could reduce cell proliferation and induce apoptosis.^{175,176} Amplification of the ALK gene and/or overexpression of the ALK protein is seen in as many as 77% of all NB tumours, suggesting that overexpression of the ALK protein may also contribute to NB.¹⁷⁹ The downstream effects of ALK in NB need to be elucidated. Current data suggest that ALK may function through the Shc and MAP kinase pathways.^{180,181} More recent studies also suggest that activation of ALK enhances RAP1 activity via interaction with C3G, a Crk-binding protein and Crk-like protein (CRKL), and that this complex contributes to NB tumour cell growth and neurite outgrowth.¹⁸²

A11.4.4 Chromosome Gain and Oncogene Activation

Many genetic abnormalities have been identified in non-familial NB tumours, including amplification of the *MYCN* proto-oncogene (25–33% of patients) and consistent areas of chromosomal deletion and rearrangement that result in loss of 1p36 (25–35%), 11q23 (35–45%), and 14q23 (16–27%), as well as unbalanced gain of 17q22 (~50%).^{132,133} In contrast, known tumour suppressor genes (TSGs) such as p16^{INK4a}, pRb, p53 and p14^{ARF} are not frequently deleted or mutated in NB, although the nuclear localization of the p16^{INK4a} and p53 proteins has been reported to be altered in some tumour cell lines.^{132,133,183} Many of these abnormalities are convincing prognostic markers and are highly related to clinical outcome. For example, amplification of *MYCN* in NB patients is correlated with chromosome 1p36 LOH. NB tumours which harbour 1p36 LOH and *MYCN* amplification are usually advanced-stage (stages 3 and 4) aggressive tumours that are frequently metastatic and generally respond poorly to chemotherapy/irradiation.^{132,133} In recent years, clinical trials are increasingly based on such tumour genetic markers.

A11.4.5 Amplification of MYCN and the 2p24 Locus

MYCN gene amplification is a hallmark of aggressive NB.¹⁸⁴ In 1983, Schwab found that a novel *myc* homologue gene was amplified in several NB cell lines and one NB tumour.¹⁸⁵

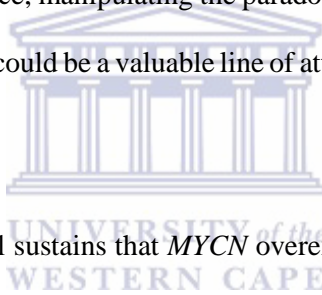
Later, several papers termed this gene as *MYCN* based on homology to *c-myc* and expression pattern in the developing nervous system, and identified its location at chromosome 2p24.^{185,186} Additional studies have shown that N-myc protein is a nuclear phosphoprotein that is a member of the myc family of helix loop-helix transcription factors.¹⁸⁷ Amplification of the *MYCN* gene in patient tumours ranges from 10-fold to more than 500-fold, although the majority of tumours exhibit 50- to 100-fold *MYCN* gene amplification levels. The amplified DNA typically contains a large region of chromosome 2 ranging from 100 kb to 1 Mb which includes the entire *MYCN* gene and varying amounts of adjacent DNA.¹⁸⁸

Although other genes may be co-amplified with *MYCN*, it is the only consistent amplified gene from this region.¹⁸⁹ *MYCN* amplification is rarely observed on chromosome 2p24 in primary tumours, but is found to be at homogeneously staining regions (HSRs) on different chromosomes or, more frequently, as double minutes (DMs; which are small fragments of extrachromosomal DNA).^{185,190} In cell culture, the amplification unit frequently integrates into chromosomes to become HSRs. The reason for the differences in the location of the amplicon in primary tumours and cultured cells remains unclear. Amplification of *MYCN* is highly associated with aggressive NB tumours and poor outcome. The precise role of *MYCN* in NB is still sketchy, however, amplification of the gene is frequently associated with the overexpression of the N-myc protein. Studies on *MYCN* regulation suggest that the transcription factor and signalling pathways controlling the upregulation of *MYCN* are dependent on cell type.¹⁹¹ These factors include IL-7 and Pax-5, NF- κ B in pre-B cells, and insulin-like growth factors I and II (IGFI and IGFII) in NB cells.¹⁹²

In contrast, *MYCN* transcription is repressed by retinoic acid (RA) in association with E2F binding, nerve growth factor (NGF) binding to TrkA receptor, the iron chelator deferoxamine mesylate and transforming growth factor-beta 1 (TGF- β 1).¹⁹² Myc proteins form heterodimers with the Max protein. These heterodimers bind to E-box elements (CACGTG) to activate transcription. However, Myc–Max dimers can also associate with other transcription factors

such as Miz-1 and Smad and bind to Inr (initiator) elements to repress transcription. Max can also form homodimers or heterodimers with Mad to compete or suppress Myc-Max binding.^{137,187,193,194}

The targets of Myc–Max are involved in various cellular processes, including cell growth, proliferation, loss of differentiation, and apoptosis, and include proteins such as MASH1 and important molecules in the normal development of sympathoadrenal lineage cells, such as the multidrug resistance protein 1 (MRP1) and MDM2.^{187,193,195} *MDM2*, which negatively regulates p53, is a direct transcriptional target of *MYCN* in NB and modulates cell cycle and transcriptional events as demonstrated by targeted inhibition of *MYCN* in a *MYCN*-amplified neuroblastoma cell line which concomitantly decreased *MDM2* expression, stabilized p53 and induced apoptosis.^{196,197} Hence, manipulating the paradoxical apoptosis-promoting function of *MYCN* amplification in NB could be a valuable line of attack in the high-risk, *MYCN*-amplified subset of neuroblastoma.¹⁹⁸



The transgenic mouse model sustains that *MYCN* overexpression is a primordial stage in NB tumourigenesis. In this model, overexpression of the human *MYCN*, followed by NB tumour formation, is driven by the rat TH promoter, which is expressed in migrating cells of the neural crest early in development.¹⁹⁹ However, other factors are also likely to be involved in the early stages of tumour formation since amplification of the *MYCN* oncogene occurs in only about one-third of NBs. Moreover, the tumours in these transgenic mice rarely exhibit significant metastasis despite the presence of high levels of N-myc protein suggesting that the other genetic alterations and/or epigenetic changes are critical for tumour formation and metastasis. The pioneering of a unique Cre-conditional human *MYCN*-driven mouse model for NB that robustly recapitulates features of the human disease, including tumour localization, histology, marker expression and genomic profile sets a significant benchmark for advance translational approaches to preclinical and molecularly targeted therapies for NBs.²⁰⁰

Likewise, it has recently been proposed that human neural crest stem cells (hNCSCs) isolated from *in vitro*-differentiating human embryonic stem cells (hESCs) may be an invaluable model system to study human neural crest development and diseases, since suppression of *MYCN* in hNCSCs not only arrests cell growth and cell cycle progression, but its knockdown induces the expression of *Cdkn1a*, *Cdkn2a* and *Cdkn2b*, which encodes the cyclin-dependent kinases p21^{CIP1}, p16^{INK4a} and p15^{INK4b}, highlighting its critical function in stem cell growth and cell cycle progression. Remarkably also, *MYCN* is involved in the regulation of human sympathetic neurogenesis, as knockdown of *MYCN* augments the expression of key transcription factors involved in sympathetic neuron differentiation, including PHOX2A, PHOX2B, MASH1, HAND2 and GATA3, which may have implications for targeted therapy of NB.¹⁶⁵

Several other genetically engineered mouse models (GEMMs) of NB have been reviewed recently: tyrosine hydroxylase (TH)-*MYCN*, TH-*MYCN*/Trp53(+/-), TH-*MYCN*/TH-Cre/Casp8(flox/flox), TH-*MYCN*/TH-ALK(F1174L) and DBH-iCre/CAG-LSL-Lin28b.^{111,201} Correspondingly, studies focusing on *MYCN*-amplified neuroblastoma patient-derived (human) xenograft models demonstrated sensitivity to the BCL-2 inhibitor ABT-199 which was partly the result of low anti-apoptotic BCL-xL expression, high pro-apoptotic NOXA expression, paradoxical *MYCN*-driven upregulation of NOXA and widespread induced apoptosis mediated by Aurora kinase A inhibitor MLN8237 combined with ABT-199, which further led to tumour shrinkage and in several instances complete tumour regression.¹⁹⁸

A11.4.6 Gain of Chromosome Arm 17q

Gain of genetic material from chromosome arm 17q (gain of segment 17q21–qter) is the most frequent cytogenetic abnormality of neuroblastoma cells. This gain has been linked with progressive disease, infants ≥ 1 year old, deletion of chromosome arm 1p, and amplification of the *N-myc* oncogene, all of which predict an adverse outcome.²³ Gain of chromosome arm 17q was originally detected by G-banded cytogenetic analysis in early 1980s. However, this observation was regarded as trivial in comparison to *MYCN* amplification and 1p loss of

heterozygosity (LOH). In the middle 1990s, the significance of 17q abnormalities in NB became increasingly discernible because fluorescence *in situ* hybridization (FISH) technology revealed that translocation of this chromosome arm was prevalent in 50% of primary NB tumours, resulting in an unbalanced gain of one to three copies of 17q which may confer a selective survival advantage for NB tumour cells.^{132,133}

It is approximated that multiples of the 17q chromosome fragment (ca. 20 Mb) accounting for more than 200 genes can be translocated in NB tumours, thus making it difficult to spot the genes responsible for the selective persistence advantage. Several genes in this region have been deemed as good candidate oncogenes or tumour suppressors based on correlations between expression levels and unbalanced gain of 17q. These include survivin, PPM1D and NM23A.¹³⁷

Overexpression of *survivin*, an anti-apoptosis gene, mapped to 17q25, is significantly associated with poor prognosis and promotes cell survival in human neuroblastoma.^{202,203} Wip1 (wild-type p53-inducible phosphatase 1) or PPM1D (protein phosphatase magnesium-dependent 1delta) is a p53-inducible Ser/Thr protein phosphatase which negatively regulates the DNA damage response through the dephosphorylation and inactivation of p53, ATM, p38 and Chk1/2, and hence drives oncogenesis.²⁰⁴⁻²⁰⁷

Low expression of NM23A (Nm23/NDP kinase), a metastasis suppressor, has been correlated with poor patient prognosis and survival, lymph node infiltration, and histopathological indicators of high metastatic potential in a number of cancer types, including NB.^{149,208-210}

Unbalanced gain of 17q correlates with other chromosomal deletions. The most frequent deletion site is the short arm of chromosome 1, followed by 11q. At least 30 translocation sites on 20 different chromosomes have been detected in patient samples and cell lines²¹¹⁻²¹³ Nevertheless, NB tumours harbouring unbalanced gain of 17q exhibit a more aggressive phenotype and a poorer prognosis than those without this abnormality.¹³⁷

A11.4.7 Amplification and Chromosome Gain of Other Loci

Besides the amplification of the *MYCN* gene, numerous other regions of gene amplifications have been distinguished in small groups of NB cases. These include amplification of the *MDM2* gene at 12q13, the *DDX* gene at 2p24, the *MYCL* gene at 1p32, and unexplained DNA from chromosome 2p22 and 2p13.^{214,215} The mouse double minute 2 homologue (*MDM2*) gene (*MDM2*) is amplified in various NB cell lines and primary tumours. Like the *MYCN* gene amplification, *MDM2* amplification unit first developed within DMs and then integrates into a different chromosome to form HSRs.²¹⁵

The *DDXI* gene, which encodes a RNA helicase, was found to be co-amplified with *MYCN* in 4/6 NB cell lines and 6/16 tumours with *MYCN* amplification, however, *DDXI* amplification was not detected in the absence of *MYCN* amplification.²¹⁶ Moreover, the *MYCL* gene is co-amplified with *MYCN* in NB cell lines. *MYCL*, another member of myc gene family, is commonly overexpressed in small cell lung carcinoma.²¹⁷ In addition to gain of 17q, other chromosome gains have been seen on 1q, 4q, 5q, 6p, 7q, 18q using comparative genomic hybridization (CGH) methodology, although their biological and clinical significance have yet to be elucidated.¹³⁷

A12.4 Chromosome Loss and Tumour Suppressor Genes

In addition to mutation, gene amplification and increased chromosome copy number, NB tumours also experience loss of genetic material and deletion of putative tumour suppressor genes (TSGs).¹³⁷

A12.4.1 Loss of Heterozygosity of Chromosome 1p and *CHD5*, *miR-34*, *KIF1B β*

Loss of the short arm of chromosome 1 occurs in about 25–35% NB tumours. 1p LOH is correlated with amplification of *MYCN* in NB patients. Loss of 1p correlates with and may stem from unbalanced gain of 17q, but the precise process that underscore these dualistic

outcomes is not well-defined. The significance of 1p LOH is borne out by research in which transferring chromosome 1p material into human NB cells caused differentiation and suppression of tumorigenicity.²¹⁸ Whereas patients with 1p36 abnormalities without *MYCN* amplification have been identified, the reverse situation virtually never occurs suggesting either that 1p36 LOH provides a permissive environment for *MYCN* amplification or that tumours with these two associated genetic defects have a high degree of genomic instability.¹³²

Remarkably, NB tumours with 1p36 LOH and *MYCN* amplification are frequently aggressive with high metastatic potential and generally resistant to chemotherapy/irradiation. Even though the chromosomal regions defined above are crucial in NB, the TSGs that reside within these regions have not been sufficiently characterized. Nevertheless, contemporary studies have classified three new putative tumour suppressors on chromosome 1p36: the chromodomain helicase DNA-binding domain 5 (CHD5), microRNA-34a (mir-34a), and the kinesin superfamily protein 1B beta (KIF1B β).²¹⁹⁻²²¹ These tumour suppressors proteins mediate their effects through cell growth dynamics, e.g., the effects of CHD5 on cell growth were shown to be dependent on p53 and CDH5 positively regulates p53 via p19ARF expression.²¹⁹

Thus, overexpression of CHD5 results in enhanced apoptosis and cellular senescence, increased p53 and p19ARF levels, and sequestration of MDM2, the negative regulator of p53, by p19ARF. On the other hand, cells lacking CHD5 exhibit decreased p16 and p19ARF expression, the latter paralleled a decrease in p53 levels and enhanced cellular proliferation. Therefore, CHD5 acts as a tumour suppressor that controls proliferation, apoptosis, and senescence via effects on the p19ARF/p53 pathway. These effects are largely attributable to changes in the accessibility of the p16/p19ARF gene locus resulting from the chromatin remodelling function of CHD5.²¹⁹ By analogy, mir-34a was found to be expressed at very low levels in unfavourable primary tumours and NB cell lines.²²² Introduction of this microRNA (miRNA) into cell lines diminished cell proliferation and enhanced caspase-dependent apoptosis, by targeting E2F3 mRNA and suppressing its expression.²²¹ E2F3 a transcription

factor that upregulates the expression of countless genes associated with cellular proliferation. Overexpression of KIF1B β induced cell death while decreased KIF1B β levels correlated with cell proliferation and enhanced tumour development in nude mice, implying that KIF1B β is also a prospective TSG candidate.²²⁰ Moreover, KIF1B β is a downstream target of prolyl hydroxylase EglN3 and an inducer of apoptosis in neuronal progenitor cells or NB cells when NGF is deficient. Missense mutations of KIF1B β in inherited NBs and pheochromocytomas strongly support the hypothesis that KIF1B β is a conceivable TSG candidate.²²³

A12.4.2 Loss of Heterozygosity of 11q and TSLC1

Loss of the long arm of chromosome 11 occurs in 35–45% NB primary tumours with a single copy *MYCN* gene. Two large patient studies that analyzed 295 NB primary tumours observed loss of 11q in 44% cases, and common regions of LOH located at 11q23, signifying that putative TSGs reside in this region.^{224,225} Loss of 11q correlated with adverse clinical features such as late stage disease, older age of disease onset and unfavourable histology, although it is strongly inversely correlated with *MYNC* amplification and 1p loss. Hence, 11q loss is a valuable and principal marker for verifying the clinical prognosis for those advanced stage tumours without *MYCN* amplification. Transfer of chromosome 11 induced differentiation in NB cell lines supporting the importance of loss of 11q in tumorigenesis.²¹⁸

Another putative tumour suppressor, the *IGSF4* (immunoglobulin superfamily 4) gene, was originally localized to the common 11q23 LOH region in 1999.²²⁶ *IGSF4*, also known as TSLC1/CADM1 (tumour suppressor in lung cancer 1/cell adhesion molecule 1), is a plausible TSG for lung cancers. A recent CGH study which examined 236 primary tumour samples found the TSLC1 LOH locus in 35% tumours. Notably, the level of TSLC1 expression correlated with tumour stage, histological classification, *MYCN* and *TrkA* expression levels. Reduced expression of TSLC1 was found in unfavourable tumours. Furthermore, introduction of TSLC1 decreased cell proliferation in NB cell lines and thus a representative NB tumour suppressor candidate.²²⁷ Interestingly, a recent study indicates that expression of both KIF1B β

and TSLC1 is controlled by the polycomb protein Bmi1, whose expression is regulated by N-myc.^{137,228}

A12.4.3 Loss of Heterozygosity of 14q

Loss of the long arm of chromosome 14 is also commonly found in NB primary tumours (~16–27% of the patients).¹³⁷ LOH on chromosome 14q was first identified in 1989 using a polymorphic DNA marker which detected allelic deletion at specific 14q23 loci.²²⁹ LOH analysis of 14q in a large number of primary tumours using 11 polymorphic DNA markers found 14q LOH in 83 of 372 tumours (22%).^{230,231} 14q LOH was highly correlated with 11q loss and had an inverse relationship with 1p loss and *MYCN* amplification.²³⁰ However, LOH for 14q was present in tumours from all clinical stages, suggesting this abnormality may be a universal early event during tumour development.¹³⁷

A13. Treatment and Management of Neuroblastoma

A13.1 Overall Therapeutic Landscape of Neuroblastoma

Neuroblastoma is the most frequent extracranial solid cancer in paediatric patients and has long puzzled scientists and oncologists alike since its biological and clinical behaviour vary between resistance to multimodal cancer therapies and complete spontaneous regression.⁶⁴ Most children diagnosed with NB are classified as high-risk cases with disseminated metastases and a mortality rate of more than 50%.³³ The cornerstone of treatment of NB consists of chemotherapy, surgical resection and/or radiotherapy. Novel personalized and molecular-guided therapy for the treatment of patients with relapsed or refractory NB are therefore a dire need to stem the tide of NB-related deaths in infants.^{19,232,233} Currently, various efforts are being pursued through innovations in basic medical sciences and translation of promising novel and molecular NB drug targets into successful clinical therapeutic practice.^{8,9,37,151,234} Treating NB is complex and frequently entails a multidisciplinary team of health professionals, including a paediatric cancer surgeon, a paediatric oncologist, a paediatric radiation oncologist. However,

many other specialists may be involved in the care of children with NB such as physician assistants, nurse practitioners, nurses, psychologists, social workers and rehabilitation specialists. Treatment of NB depends on the stage of the cancer and the child's age. Current treatment modalities can include any one or several combinations of surgery, chemotherapy, radiation therapy, high-dose chemotherapy/radiation therapy and stem cell transplantation, retinoid therapy and immunotherapy. Complementary and alternative methods to treat cancer or relieve symptoms are optional. These methods can include vitamins, herbs, and special diets, or other methods such as acupuncture or massage. Even though some of these methods might be helpful in relieving symptoms of NB, many have no proven efficacy and might even cause unwanted side effects (*Neuroblastoma*, American Cancer Society, <http://www.cancer.org/>).

A13.1.1 Spontaneous Regression and Stage 4S Disease

Spontaneous regression of cancer is not a new concept and has been defined in the 1950s as “the partial or complete disappearance of a malignant tumour in the absence of all treatment, or in the presence of therapy which is considered inadequate to exert a significant influence on neoplastic disease.”^{235,236} However, one of the first allusions to regression of malignant NB was published in 1927.²³⁷ A significant feature of NB is that occasionally it undergoes spontaneous regression.²³⁸⁻²⁴⁰ This propensity is consistent with the notion that NB is largely caused by aberrations in the embryonic progressions of the neural crest and thus the sympathetic nervous system.¹¹¹

Overexpression of the *MYCN/c-MYC* target gene is a hallmark of malignant NB progression—a process primarily driven by *c-MYC* in stage 4-non-amplified tumours.²⁴¹ It has been proposed that moderate gain of *MYCN* function in stage 4S-non-amplified tumours induces a number of target genes that retain their ability to trigger spontaneous regression.^{241,242} More importantly, the fifth stage of NB tumours (stage 4S) is said to undergo spontaneous regression with minimum treatment or even without medical intervention.^{238,243}

Recent genomic studies of NB corroborated the striking heterogeneity in the clinical behaviour of this disease, which encompasses spontaneous regression or differentiation in some patients, to relentless disease progression in others, in the face of intensive multimodality therapy.¹² Thus, some NBs regress spontaneously without therapy while others progress with a fatal outcome despite therapy. In one such study of infants younger than 12 months, spontaneous regression was noted in almost 50% of the study population within three years of follow-up.²⁴⁴ Several conceivable mechanisms may account for the spontaneous regression observed in NBs, including HOX gene expression (HOXC9 expression is downregulated in advanced-stage NB and is involved in cell cycle control and the processes of NB cell differentiation),^{245,246} neurotrophin signalling (especially those through nerve growth factor and its receptor, TrkA),²⁴⁷⁻²⁵⁰ activation of developmentally programmed apoptosis, humoral or cellular immunity, loss of human telomerase (h-Tert) activity, epigenetic changes in gene expression controlled by DNA methylation, histone modification, or alterations in chromatin remodelling.¹² A better understanding of the mechanisms of spontaneous regression might help to identify optimal therapeutic approaches for patients with these tumours.¹¹¹

A13.1.2 Surgery

The goal of surgical resection of NB is to attain macroscopic tumour resection with minimal residual disease (MRD). Surgery in ‘low’ and ‘intermediate risk’ groups is aimed at complete resection—wherever possible—with minimal injury to adjacent structures which are frequently adherent to, if not encased by the tumour mass.^{251,252} NB is a highly infiltrative neoplasm and poses several challenges for the paediatric cancer surgeon, e.g., difficulty in obtaining microscopically negative resection margins wherein gross total resection (GTR) or subtotal tumour resection (STR) is desirable.²⁵³ Much controversy and debate subsists on the defining role of surgery in advanced stage 4 disease. Moreover, significant overall survival advantage for radical surgical clearance in stage 4 disease has never been clearly demonstrated.²⁵²⁻²⁵⁸

The first recorded successful excision of a NB occurred in 1916, but for many years there was no other form of treatment, and the outlook remained dismal. The use of radiotherapy (1928) and subsequently combination chemotherapy (1965) had a modest impact. Recent advances in accurate disease imaging and staging has allowed a more coherent approach to diagnosis and treatment of NB.^{47,251,257} Generally, surgery in NB can be performed safely, but moderate to serious intra- and postoperative surgical complications have been experienced, including massive haemorrhage, major vascular injury, respiratory failure requiring mechanical ventilation after major surgery, cardiac arrest, tumoural rupture, nephrectomies, Bernard-Horner syndrome and pleural effusions.²⁵¹ It has been suggested that presurgical chemotherapy may lead to a more extensive and safer removal of locally advanced tumours.²⁵⁶

A recent analysis of the SEER database (The Surveillance, Epidemiology, and End Results Programme (SEER, <http://seer.cancer.gov>), after accounting for selection bias, indicated improved survival following surgery and radiation therapy for olfactory neuroblastoma. However, the efficacy, timing, and optimum approach for combining chemotherapy with surgery and radiotherapy could not be established.²⁵⁹ Also, in intensively treated patients with stage 4 neuroblastoma age 18 months or older at diagnosis, surgery of the primary tumour site had no impact on local control rate and outcome.²⁵⁵ Similarly, no substantial survival benefit had been noted in stage IV neuroblastoma patients undergoing complete tumour resection, organ preservation and minimalization of morbidity.²⁵⁴ Despite these challenges, surgical resection remains a cornerstone of therapy in paediatric patients suffering from this clinically and biologically heterogeneous and complex disease.²⁵³

A13.1.3 Chemotherapy

About 80% of patients with high-risk NB often survive their primary tumour, i.e., attain remission through high-dose chemotherapy, surgery, radiation and stem cell transplantation. However, those with relapsed metastatic disease after treatment have a discouraging long-term survival outcome.²⁶⁰ As a paediatric extracranial solid tumour paradigm, NB correlates with a

disproportionately high mortality rate, i.e., 15% of all cancer-related deaths in children.¹⁶ In spite of 5-year event-free survival (EFS) and overall survival (OS) rates >90% for low- and intermediate-risk NB groups, the survival rate in children with high-risk NB remains 40%–50%.^{3,251,259,261} Also, the manifestation of minimal residual disease (MRD) persists to be a substantial obstacle to bettering the prognosis of patients with high-risk NB.²⁶²⁻²⁶⁴

Chemotherapy for high-risk NB entails three empirical stages, viz., (i) induction of remission, (ii) consolidation of remission and ultimately (iii) maintenance phase focused on the eradication of MRD.^{64,265} Generally, induction regimens utilize various combinations of anthracyclines, platinum-based compounds, etoposide, microtubule disruptors and alkylating agents. NB combination chemotherapy encompasses dose-intensive cycles of cisplatin and etoposide alternating with vincristine, doxorubicin, and cyclophosphamide.²⁶⁶ Very low and low-risk patients may require only observation or surgical resection, except in cases of life- or organ-threatening symptoms at diagnosis. Intermediate risk group patients whose tumours are not compatible for primary resection are put on chemotherapy designed to selectively destroy rapidly dividing tumour cells, eliminate life-threatening symptoms or make tumour resectability easier.

High-risk patients undergo chemotherapy protocols combining carboplatin, etoposide, cyclophosphamide, doxorubicin, and vincristine. Furthermore, patients in the high-risk group also receive myeloablative chemotherapy to impede tumour infiltration (even though normal bone marrow is also suppressed), followed by bone marrow transplantation and granulocyte macrophage colony-stimulating factor (GM-CSF) induction. This sequential therapy regimen has proved beneficial in enhancing EFS.²⁶⁵ Mild chemotherapy regimens, especially those presently indicated for intermediate-risk NB are deemed ineffective for preventing evolution into advanced-stage (high-risk) disease. There is a disinclination to expose a clinically disease-free infant or child to aggressive, highly toxic multimodal therapy that is only partially effective against advanced-stage disease. Hence, close clinical monitoring of such patients is

warranted.²⁵¹ However, chemotherapy is invaluable in the early treatment of patients with stage-2 tumours who present with spinal cord susceptibility from a paraspinal mass or airway weakness from a tumour in the superior mediastinum. Biologic findings strengthen contentions surrounding the use of cytotoxic therapy in localized disease. The salient disparities in chromosomal features of lethal vs low-risk forms of NB lend biologic support for the radical dichotomy in prognosis. In this regard, progression of non-stage-4 NB with low-risk biologic features (triploidy, unamplified *MYCN*) to lethal stage-4 disease is a rare event. Several lines of evidence reinforced the hypothesis that non-stage-4 NB without *MYCN* amplification rarely, if ever, evolves into lethal disease.²⁵¹

Biologically favourable (low-risk) stage-4S disease resolves spontaneously in the majority of cases and surgical resection of primary tumours at diagnosis is no longer recommended since these tumours are likely to regress.²⁶⁷ Some stage-4S tumours with low-risk prognostic markers, e.g., non-amplified *MYCN*, hyperdiploidy and favourable histopathology, can cause dire life-threatening cardiopulmonary complications and coagulopathies in the neonatal period. Such medical emergencies may abate after treatment with one to two cycles of low-dose chemotherapy and/or modest doses of radiotherapy in order to spare the kidneys and spine. In the face of persistence of liver lesions, once clinical adjustment has been achieved, supplementary cytotoxic therapy may not be needed as it might pose some risk and the residual disease, even if extensive, is likely to regress.²⁵¹

Infants with stage-3 NB lacking *MYCN* amplification have survival rates close to 100%. In multi-institution studies in North America and Europe, these patients have received various modest dose-combinations of platinum compounds, etoposide, cyclophosphamide, doxorubicin and/or vincristine.²⁶⁸⁻²⁷⁰ The French Society of Paediatric Oncology achieved similar success using alternating cycles of carboplatin/etoposide and cyclophosphamide/doxorubicin/vincristine, in moderate doses.^{251,271} In a large Children's Cancer Group (CCG) study, a regimen involved 9 months of combination chemotherapy

incorporating cisplatin, etoposide, cyclophosphamide, and doxorubicin.²⁷⁰ In another large Paediatric Oncology Group (POG) study, cycles of high-dose cisplatin/etoposide alternated with low-dose cyclophosphamide/doxorubicin, and in a follow-up POG study, patients received cycles of cyclophosphamide, etoposide, vincristine, plus either cisplatin or carboplatin.²⁷² Similar results described in clinical trials referred to above have been reported for treatment of Infant Stage-4 NB, at lower but improving cure rates from 10 to 50 to >70%.²⁶⁸

Neonatal NB constitutes less than 5% of all cases of the neoplasm and frequently correlates with a good prognosis provided that patients are stratified into low- or intermediate-risk groups for disease recurrence. In neonates less than or older than 30 days, NB has the unusual potential to undergo spontaneous regression and this characteristic is used as a paradigm by several paediatric oncology groups (CCG, COG, POG and INRG) to moderate therapy given to neonates with low-risk NB.^{8,44} These groups also strive for the agency of reduced cytotoxic chemotherapy therapy and surgical tumour ablation for certain low- and intermediate-risk patients, but advocate observation approaches for such favourable subsets. By analogy, high-risk patients should receive aggressive chemotherapy, radiation, surgery and myeloablative and immunotherapies.^{18,45,273}

A13.1.4 Radiotherapy

While NB responds favourably to radiotherapy, the efficacy of total body irradiation in paediatric patients remains debateable in the face of long-lasting adverse events. Currently, COG promotes the concept that high-risk patients receive radiation to the primary tumour site irrespective of the coverage of surgical resection and to metastatic sites that display persistent MIBG avidity on pre-transplantation scans.^{54,55,260} Radiotherapy is not indicated for low-risk NB, even with local residual disease as risks outweighing potential benefits. In low- and intermediate-risk groups, radiation therapy is reserved for patients with progressive clinical relapse despite chemotherapy and surgery. Infants with stage 4S disease are usually excluded from radiotherapy, except those with severe respiratory distress or abdominal compartment

syndrome with precipitous hepatomegaly. Synergistic, additive and antagonistic acute and long-term tumour responses as well as side effects may be produced by parallel use of radiosensitizing agents.^{260,274} Radiosensitizers include cisplatin, topotecan and irinotecan²⁶⁰ which are usually safe to give with radiation therapy. Currently, spinal cord compression is managed by chemotherapy, radiotherapy or surgical resection with or without laminectomy, but is contingent on a case by case basis. Radiation therapy is contraindicated in intraspinal tumours because it can trigger gross vertebral impairment and growth arrest resulting in severe scoliosis, but it may be used selectively as an emergency therapy for patients with symptomatic spinal cord compression.⁶⁴

A13.1.5 Haematopoietic / Peripheral Blood Stem Cell Transplantation

Almost 56% of NB patients present with disseminated disease at the time of diagnosis. The bone, bone marrow, liver, non-contiguous lymph nodes and central nervous system (including the choroid plexus) are the most frequent metastatic foci.^{118,133,275-279} The low 5-year survival rate (40–45%) of NB patients with secondary tumours (metastases) underscores the therapeutic hurdles faced by paediatric oncologists in the face of advanced treatment options.²⁸⁰ Children with bone metastasis have a dismal outcome with survival rates below 7%.²⁸¹ Moreover, 40–50% of patients relapse (presenting with occult NB cells in peripheral blood), often after total remission following multi-modal treatment (surgery, chemotherapy and radiation).²⁸²

The bone marrow is a chief metastatic site in stage IV NB and therefore assessment of MRD in the bone marrow is implicit in disease prognosis. Since high-risk NB correlates with a worse prognosis, autologous bone marrow transplantation (ABMT)²⁸³ and autologous peripheral blood stem cell transplantation (PBSCT) have become a therapeutic mainstay to enhance the prognosis of such patients, in particular to support haematopoietic rescue following high-dose chemotherapy.^{264,279,284-287} However, re-infusion of PBSC contaminated with tumour at the time of autologous transplantation may play a significant role in the high proportion of relapse in children with NB who eventually succumb to the disease.^{264,285,286,288,289} Recently, pulmonary

arterial hypertension (PAH) has been acknowledged as a rare condition with high mortality rate after paediatric haematopoietic stem cell transplantation (HSCT). Generally, there is a propensity to overlook PAH in the differential diagnosis of cardiorespiratory failure after HSCT as the clinical presentation is non-specific and may mimic other aetiologies.²⁹⁰ Accordingly, paediatricians overseeing HSCT recipients should be cognizant of this serious post-transplant complication as appropriate diagnosis and treatment may improve clinical outcomes.²⁹¹

A13.1.6 Management of Minimal Residual Disease and Relapse

Minimal residual disease (MRD) is a major barrier to the obliteration of malignant neoplasms. Even with the high sensitivity of various cancers to therapy, fractions of residual tumour cells persist and give to tumour recurrence and treatment failure.^{292,293} The detection of minimal amounts of tumour cells in bone marrow, peripheral blood, putative metastatic sites, lymph nodes or in other tissues, compartments or body fluids has become a major goal in cancer diagnostics. For NB patients, the clinical significance of minimal residual disease (MRD) in bone marrow after induction of chemotherapy or within stem cell harvests prior to autologous transplantation has become a critical clinical context.^{262,263} In paediatric patients, relapse is a frequent occurrence after autologous BMT, indicating the presence of malignant stem cells that are resistant to dose-intensive myeloblastic chemotherapy.^{7,118,131,294}

MRD status in PBSCs might be crucial in high-risk NB because PBSCs contaminated with tumour cells are thought to contribute to relapse and increased mortality rate. Detection of tyrosine hydroxylase (TH) transcripts by RT-PCR is one way to assess whether PBSCs are contaminated with tumour cells.²⁶⁴ Re-infusion of PBSC contaminated with tumour at the time of autologous BMT may play a significant role in this relapse.²⁸⁸ Early observations that retinoids and other agents are useful to induce differentiation of NB, and, hence improve survival outcomes, have raised interest in these compounds as biological response modifiers against MRD in the bone and bone marrow.^{152,262,295} Immunocytology and quantitative RT-

PCR for tumour specific markers such as diganglioside (GD2)²⁹⁶ and TH are currently used to detect and quantify MRD. Therapeutic regimens presently integrate novel biological therapies, including retinoic acid post-consolidation therapy for high-risk NB, aimed at eradicating MRD.²⁹⁵ Recently, a MRD model has been conceptualized as a novel approach to testing preclinical therapies and interpreting mechanisms of MRD and metastatic disease in experimental NB.²⁹⁷

Currently, patients receive six courses of 13-cis-retinoic acid (CRA) to eradicate residual disease that may still be present despite meeting imaging criteria for complete remission. This treatment is guided by the observation that high-dose CRA administered after chemoradiation significantly improved EFS in high-risk NB.¹⁵² Side effects such as skin dryness and cheilitis are the dose-limiting factors, and consequently, CRA therapy involves of 2-week courses alternating with 2 weeks for mucocutaneous recovery.^{1,298} CRA appears to be most suited in the setting of MRD.⁶² High dose CRA and pulse schedules of other retinoids show therapeutic and chemopreventive efficacy in NB, but low-dose, chronic retinoid administration may not be as effective to treat MRD.²⁹⁹

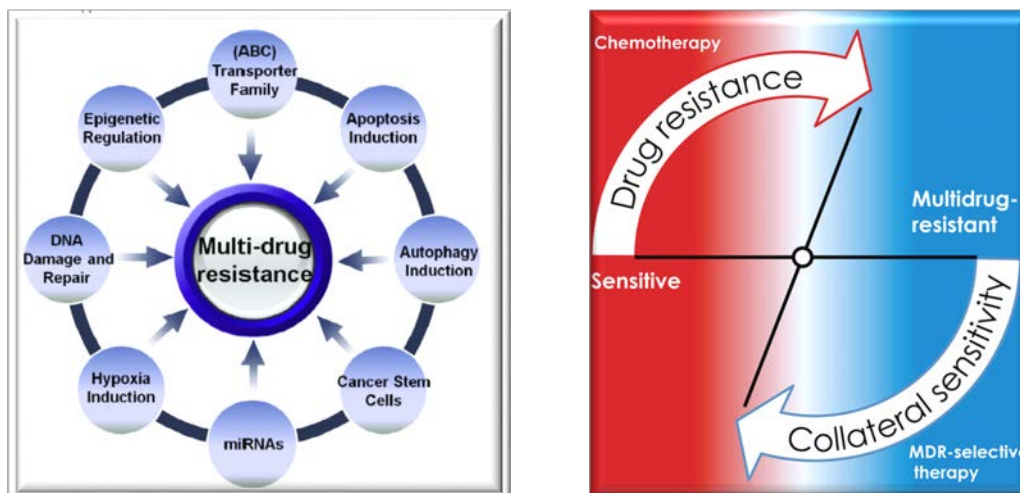
Clinical trials involving myeloablative chemotherapy and ¹³¹I-MIBG have been undertaken in an effort to minimize adverse side effects and rationalize more targeted therapies.³⁰⁰⁻³⁰² ¹³¹I-MIBG exhibits activity against refractory NB with response rates ranging from 10-50%. In a phase I trial of ¹³¹I-MIBG therapy for relapsed NB, myelosuppression was the most significant toxicity at doses >15 mCi/kg, as nearly half of the patients enrolled required haematopoietic cell transfusion. Despite this, the response rate (36%), EFS (18% at 1 year), and OS (49% at 1 year; 29% at 2 years) were found to be significantly higher in patients older than 12 years and who had fewer than three prior treatment regimens.³⁰³ Subsequently, a phase I dose escalation study of ¹³¹I-MIBG with myeloablative chemotherapy and stem cell rescue showed a significant response rate of 25% in patients with primary refractory disease. These observations, together with recent research, indicate that ¹³¹I-MIBG may prove useful in

conjunction with other treatment modalities,^{1,49,80,300,302,304,305} as corroborated also by a recent study that autologous stem-cell transplantation (ASCT) may not be indispensable for better outcomes when anti-GD2 immunotherapy is used for consolidation after dose-intensive conventional chemotherapy.³⁰⁶ However, in certain cases, even after 12 months of high-dose chemotherapy, treatment failures due to MRD are still widespread because of the acquisition of drug resistance and most patients who relapse eventually die from disease progression. Also, those patients who achieve a cure with initial therapy persist to be at risk for developing long-term complications related to treatment, including blindness, hearing loss, infertility, and secondary malignancies.^{307,308}

A13.1.7 Multidrug Resistance and Monitoring Response to Treatment

The term multidrug resistance (MDR), as applied to cancer, is a phenomenon in which tumour cells have developed decreased sensitivity to a wide variety of unrelated drugs (cross-resistance) with different modes of pharmacological activity.³⁰⁹⁻³¹⁴ Many of the mechanisms that decrease cell sensitivity to chemotherapeutic agents are caused by well-defined genotypic and phenotypic alterations, including overexpression of the ATP-binding cassette (ABC) transporter family, apoptosis induction, autophagy induction, cancer stem cell (CSC) regulation, miRNA regulation, hypoxia induction, DNA damage and repair, and epigenetic regulation (Figure 1.11).^{312,315,316}

The molecular basis of MDR generally involves specific members of the ABC drug transporters or efflux pumps such as ABCB1 (P-glycoprotein/P-gp/MDR1), ABCG2 (breast cancer resistance protein/BCRP) and ABCC1 (multidrug resistance protein 1/MRP1) and ABCC4/MRP4.^{311,313,317,318} In paediatric malignancies, different CSC phenotypes have been detected—including those that overexpress ABC transporters^{293,319-323}—not only accounting for the tumour heterogeneity associated with NB, but also raising hopes that further insight into the mechanisms that control the traits of CSCs may aid in the design of novel strategies to overcome chemoresistance.^{7,144,294,324}



A: Source:³¹⁵ Wu Q, Yang Z, Nie Y, Shi Y, Fan D. Multi-drug resistance in cancer chemotherapeutics: Mechanisms and lab approaches. *Cancer Letters* 2014;347(2):159-166, with permission from Cancer Letters, Elsevier Ireland Ltd.) See Appendix 7 for copyright clearance.

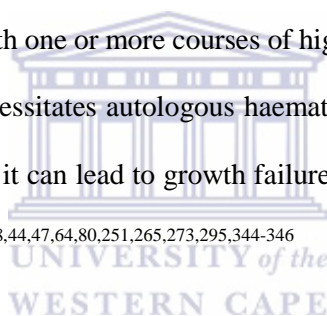
B: Source:³²⁵ Szakacs G, Hall MD, Gottesman MM, Boumendjel A, Kachadourian R, Day BJ, Baubichon-Cortay H, Di Pietro A. Targeting the Achilles heel of multidrug-resistant cancer by exploiting the fitness cost of resistance. *Chemical Reviews* 2014;114(11):5753-5774, with permission from the American Chemical Society (ACS; <http://pubs.acs.org/copyright/permissions.html>).

Figure 1.11: Mechanisms of MDR and the concept of MDR targeting based on collateral sensitivity

Overexpression of ABCG2/BCRP and ABCC4/MRP4 in subpopulations of stem cells was demonstrated in NBs.^{311,326,327} ABC transporters such as ABCC1, ABCC3 and ABCC4 are subject to direct transcriptional control of MYCN,³²⁸ and their expression compellingly correlates with poor prognosis.^{327,329,330} Many aggressive NBs exhibit MDR,³¹¹ attributable to p53 mutations and/or a loss of p53 function acquired during chemotherapy,^{331,332} which escalates the likelihood of relapse and thus presents a major obstacle to effective tumour eradication.^{333,334}

Most metastatic drug-resistant NBs derive from the selection of clones (side population cells) that express the *MDR1* (*ABCB1*), *MRP1/ABCC1* and *MRP4/ABCC4* gene family, which may or may not correlate with *MYCN* amplification and poor outcome.^{7,294,323,327,328,330,335} Moreover, MRD, the major cause of tumour recurrence (relapse) and metastasis, is enriched in CSCs with an increased drug efflux capacity mediated through overexpression of ABC transporters.^{336,337}

Hence, in the case of refractory NBs, the treatment algorithm is determined by the patient's disease stage and risk stratification.¹ The objective of induction chemotherapy is to promote remission by alleviating the tumour burden, which then simplifies complete resection when indicated—such as stage 1-2B tumours, but in the case of more advanced-stage NB (stages 3 and 4), surgical intervention is limited to an open biopsy, and for infants who are stage 4S, surgical resection is not recommended since these tumours tend to spontaneously differentiate and regress.^{338,339} Induction chemotherapy consists of various combinations of cyclophosphamide, doxorubicin, cisplatin, melphalan, carboplatin, etoposide, topotecan, ifosfamide and vincristine, an example of the chemotherapeutic platform of collateral sensitivity (the hypersensitivity of resistant cancer cells to other drugs) that aims to kill MDR cells selectively over the parental cells from which they were derived.^{325,340-343} After induction, treatment is consolidated with one or more courses of high-dose chemotherapy to induce bone marrow ablation, which necessitates autologous haematopoietic stem cell support. Rescue is not without complication as it can lead to growth failure, endocrinopathy, and the occurrence of secondary metastases.^{6,8,18,44,47,64,80,251,265,273,295,344-346}



A13.1.8 Alleviating the Burden of Late Effects

A standard component of NB management is follow-up of survivors to monitor and treat adverse events that may debilitate their quality of life and increase the rate of early mortality.^{8,62,73,347} NB patients invariably are subjected to intensive therapy which are increasingly associated with a number of complications, including: hearing loss linked to platinum compounds and ototoxic antibiotics indicated for neutropaenic infections (may result in learning and speech impediments); compression of renal vessels resulting in hypertension (renal toxicity caused by platinum agents, myeloablative therapy (MAT) with ASCT or HSCT, radiotherapy, nephrectomy, nephrotoxic antibiotics); secondary metastases initiated by alkylating agents or radiotherapy; leukaemias and myelodysplastic syndromes reportedly related to high doses of etoposide and cyclophosphamide (a reduction of these was achieved by limiting the number of cycles; thyroid cancers and other solid tumours; hypothyroidism;

growth impairment; musculoskeletal abnormalities (scoliosis, osteoporosis and bony and soft tissue hypoplasia may result from surgery and/or radiotherapy); cardiopulmonary sequelae induced by anthracyclines or thoracic radiotherapy (endocrine complications and reduced fertility).^{62,347-353} It is therefore imperative to develop new therapeutic regimens that will improve the survival and quality of life of NB patients.⁶⁴

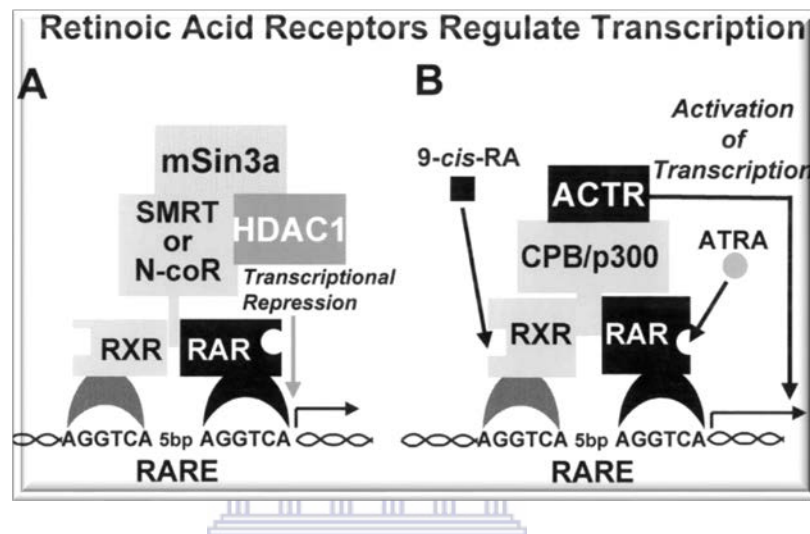
A13.2 Current Research Milestones and Proposed Novel Therapies

A13.2.1 Differentiation and Retinoids

Retinoids, including 13-cis-retinoic acid (CRA, isotretinoin), all-trans retinoic acid (ATRA) and N-(4-hydroxyphenyl) retinamide (HPR, fenretinide) are endogenous, lipophilic vitamin A derivatives that arrest cell growth and induce differentiation of cell lines derived from tumours that are resistant to anticancer drugs.^{152,298,354-361} In the late 1980s, ATRA has been hailed as a therapeutic advancement for acute promyelocytic leukaemia and high-risk neuroblastoma (HR-NB).³⁵⁴ Retinoic Acid (RA) is included in multimodal therapies because it stimulates differentiation of NB cells *in vitro* and decreases the risk of tumour recurrence.^{362,363} CRA is given at completion of cytotoxic therapy to control MRD in NB.^{262,264,295,364-368}

ATRA or CRA have been shown to inhibit cell proliferation and induce morphological differentiation of human NB cell lines,^{355,356} whereas HPR (an apoptosis inducer) is highly active against retinoic-acid-resistant NB cell lines by deregulating *MYC* expression and removing its transcriptional repression of NB cell differentiation.^{369,370} HPR inhibits NB-induced angiogenesis and may be applied usefully in aggressive HR-NB.^{62,371} HPR is also well-tolerated in clinical trials, but has a poor solubility and may decrease night vision.³⁷² Interestingly, downregulation of the acylated and glycosylated 67-kDa laminin receptor (67LR) by RA has been shown to correlate with reduced metastatic aggressiveness of human NB cells, rendering 67LR as a molecular target in NB.³⁷³ The retinoids bind to members of the nuclear receptor family of proteins (transcription factors), i.e., the retinoic acid receptors (RARs—multiple isoforms of RAR α , β , and γ) and the retinoid X (rexinoid) receptors RXRs—

multiple isoforms of RXR α , β , and γ)—which alter interactions of transcription complexes with numerous cancer stem cell (CSC) genes, thus triggering an exit from the self-renewing, pluripotent CSC reserve into a differentiated mature cell niche.^{354,374} The retinoids are activated by RARs and the rexinoids by RXRs—RAR/RXR heterodimers constitutively associate with RA response elements (RAREs) in promoters of target genes (Figure 1.11).^{298,354,357,374,375}



The mechanism of action of retinoids is mediated via zinc-finger transcriptional regulators which function as heterodimers to regulate promoter activity of certain target genes. The RAR and RXR proteins bind to specific direct repeat DNA sequences (AGGTCA are separated by either 2 or 5 nucleotides) in gene promoters, known as RA response elements. (A) In the absence of ligand, the RAR/RXR heterodimers interact with nuclear co-repressors including N-CoR and SMRT, which in turn bind to a common adapter protein mSin3 that complexes to proteins with histone deacetylase activity to repress transcription. (B) RA binds to the RAR portion of the complex causing a conformational change in the RAR and RXR proteins which releases the co-repressor complex and facilitates binding of 9-cis-RA to the RXR protein (the latter enhances the activation response). The transcriptional co-regulator CBP/p300 then binds to the receptor complex and recruits the coactivator protein ACTR, which contains histone acetyltransferase activity, that promotes transcription; RARE= retinoic acid response elements.

Source:²⁹⁸ Reynolds CP, Matthyay KK, Villablanca JG, Maurer BJ. Retinoid therapy of high-risk neuroblastoma. *Cancer Letters* 2003;197(1-2):185-192, with permission from Cancer Letters, Elsevier Ireland Ltd.) See Appendix 8 for copyright clearance.

Figure 1.12: The mechanism of action of retinoids and rexinoids

The zinc-finger cluster of ZNF423 (also known as Ebfaz, OAZ, or Zfp423) has been shown to be indispensable for retinoid-induced differentiation. ZNF423 combines with the RAR α /RXR α nuclear receptor complex and is critical for transactivation in response to retinoids. Blockade of ZNF423 expression by RNA interference in NB cells confers a growth advantage and acquired resistance to RA-induced differentiation whereas its overexpression triggers growth

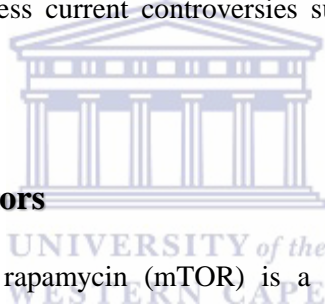
arrest and amplified differentiation. Correspondingly, deregulation of ZNF423 expression correlates with poor disease outcome in HR-NB patients.³⁵⁹ Several clinical trials have been conducted to establish the maximal-tolerated doses (MTDs), pharmacokinetics, efficacies and dose-limiting toxicities (DLT) of the retinoids/rexinoids.³⁵⁷ Current selected clinical trials on various interventions in NB have recently been published.³⁷⁶ In a phase I trial in children 2 to 12 years of age with NB, treatment with CRA (isotretinoin) doses escalated from 100 to 200 mg/m²/day following BMT, DLTs in 6 of 9 patients were observed, including hypercalcaemia (n=3), rash (n=2), and anaemia/thrombocytopenia/emesis/rash (n=1). All toxicities resolved after CRA withdrawal. Three complete responses were observed in bone marrow metastases.³⁷⁷

Another study was performed on eligible patients which included children and adolescents (1 to 18 years of age) with newly diagnosed HR-NB to assess whether MAT in conjunction with ABMT improved EFS as compared with chemotherapy alone, and whether subsequent treatment with CRA further improves EFS. In this study, 434 patients had Evans stage IV NB; 72 had stage III disease with one or more of the following: amplification of the *MYCN* oncogene, a serum ferritin level of at least 143 ng/ml and unfavourable histopathological findings; 1 had stage II disease with amplification of *MYCN* (age > 1 year); 13 had stage I or II disease with bone metastases before therapy other than surgery; and 19 had had stage IV disease with *MYCN* amplification for less than one year. The conclusion was that treatment with MAT and ABMT improved EFS among children with HR-NB and, significantly, that CRA administered successively to chemotherapy or transplantation had a favourable outcome for patients without progressive disease.¹⁵²

However, a recent phase 2 trial of ATRA, administered orally at a dose of 90 mg/m²/day in three divided doses for 3 consecutive days per week—and IFN- α 2a administered subcutaneously daily at a dose of 3×10^6 U/m²/day for 5 consecutive days per week, in 4 week cycles—was inactive in children with relapsed or refractory NB and Wilms tumour.³⁷⁸ By comparison, assessment of the long-term outcome of HR-NB patients enrolled on the CCG-

3891 study in which patients were randomly assigned to undergo ABMT or to receive chemotherapy and subsequent treatment with CRA indicated that MAT and ABMT, significantly improved 5-year EFS than non-MAT chemotherapy and neither MAT with ABMT nor CRA given after consolidation therapy significantly improved OS.³⁷⁹ The aforementioned differences may be ascribed to pharmacogenetic variation on CRA disposition in patients with HR-NB and emphasize the need for personalized therapies.^{357,380}

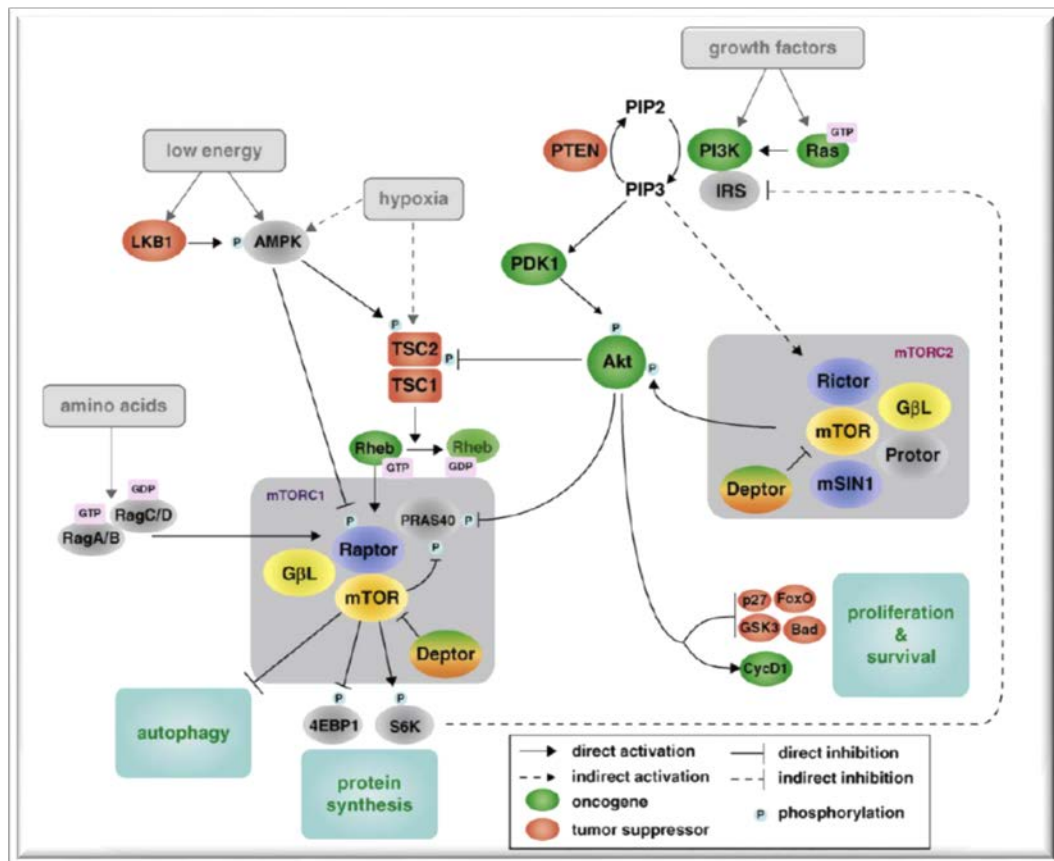
In a retrospective cohort design to verify if intensive chemoradiotherapy with purged ABMT and/or CRA improved outcome for HR-NB patients with no metastatic distant sites, it was deduced that patients with high-risk INSS Stage 3 NB have an overall poor prognosis despite aggressive chemoradiotherapy.³⁵⁸ Clearly, further research into the efficacies and DLTs of the retinoids is needed to address current controversies surrounding their beneficial status in patients with HR-NB.²⁹⁵



A13.2.2 mTOR Inhibitors

The mammalian target of rapamycin (mTOR) is a serine/threonine protein kinase and downstream effector of the PI3K/AKT pathway and master regulator that orchestrates, via a network of regulatory loops, various signals from nutrient and energy sensors with cell growth and proliferation, survival and motility to ensure that they are activated exclusively during favourable conditions within different organs (Figure 1.12).³⁸¹

The mTOR pathway can be activated by various exogenous stimuli such as growth factors, nutrients, energy and stress signals, and essential signalling pathways (e.g., PI3K, MAPK and AMPK) in order to mediate temporal control of physiological processes.³⁸² Deregulation of the mTOR pathway, including PI3K amplification or mutation, PTEN loss of function, AKT gain of function, and S6K1, 4EBP1 and eIF4E overexpression, has been correlated with oncogenesis, tumour progression and metastases of many cancers, including NB.^{381,383-387}



mTOR signaling pathway. One branch of mTORC1 activation is mediated by the class I phosphoinositide-3-kinase (PI3K), Akt (also known as Pkb) and the tuberous sclerosis complex (TSC). TSC is formed by TSC1 and TSC2, and inhibits a direct activator of mTORC1, the GTPase Ras-homolog enriched in brain (Rheb) by hydrolyzing its GTP into GDP. TSC2 is activated by phosphorylation by AMP-activated protein kinase (AMPK), which is directly activated by a high AMP versus ATP ratio. AMPK also directly phosphorylates and inactivates Raptor, so it inhibits mTORC1 by TSC-dependent and TSC-independent manners. The activity of AMPK is regulated by phosphorylation by the tumour suppressor LKB1. This protein, like TSC1/2, was found mutated in the germline of patients with different hamartomatous syndromes. Akt is a serine/threonine kinase and an important player in regulating mTORC1 activity. Akt positively regulates mTORC1 by acting at different levels. First, Akt inactivates TSC1/2 by phosphorylating TSC2.

Second, Akt inhibits PRAS40, negative regulator of mTORC1 that counteracts Rheb function. Akt is activated by PI3K, which responds to a variety of growth factors. When activated by insulin or insulin-like growth factors (IGFs), as well as other growth factors, class I PI3K catalyzes the formation of the lipidic second messenger phosphoinositide-3,4,5-tri-phosphate (PIP3) from the bi-phosphate form PIP2. PIP3 triggers the relocation of Akt to the inner surface of the plasma membrane, where it is activated by phosphoinositide-dependent kinase 1 (PDK1) and transduces the signal as described above.

Opposing Akt function is the tumour suppressor phosphatase and tensin homolog deleted on chromosome ten (PTEN), a lipid phosphatase that converts PIP3 to PIP2, thus shutting off signaling from PI3K. PTEN deficiency causes a series of hamartomatous syndromes collectively classified as PTEN hamartoma tumour syndrome (reviewed in. Amino acids activate mTORC1 by an independent route mediated by the Rag family of proteins. The activation of mTORC2 is not well understood, but this complex directly activates Akt (and Akt-related kinases) by phosphorylation. Akt, in addition, regulates many proteins involved in cell survival and cell-cycle progression.

Source:³⁸¹ Efeyan A, Sabatini DM. mTOR and cancer: Many loops in one pathway. *Current Opinion in Cell Biology* 2010;22(2):169-176, with permission from Current Opinion in Cell Biology, Copyright Clearance Center's RightsLink service, Elsevier) See Appendix 9 for copyright clearance.

Figure 1.13: Overview of the mTOR signalling pathway in cancer

In a benchmark study, AKT and mTOR were found to be overexpressed in primary NB tissue samples, but in non-malignant adrenal medullas this pattern could not be demonstrated. mTOR inhibitors (rapamycin and CCI-779) arrested the growth of NB cells in culture, particularly cell lines with a high *MYCN* gene expression signature. *In vivo*, mTOR inhibitors increased apoptosis, decreased cell proliferation and blocked angiogenesis in established NB tumours. Significantly also, mTOR inhibitors induced downregulation of VEGF-A secretion, cyclin D1 and *MYCN* protein expression *in vitro* and *in vivo*. Even though mTOR inhibitors may inhibit proliferation of human NB cells without suppression of the *MYCN* oncoprotein,³⁸⁸ the above findings underscore the therapeutic efficacy of mTOR inhibitors in aggressive *MYCN* amplified NBs and corroborate similar observations in NB tumours with 1p36 aberrations, advanced stage disease at diagnosis and unfavourable histology in which AKT and *MYCN* are co-amplified.^{389,390} The mTOR pathway and VEGF signalling are implicated in the regulation of clonal proliferation, angiogenesis and metastasis. Collateral inhibition, either in a concurrent or successive design, of mTOR and VEGF signalling exemplifies an interesting therapeutic rationale to overcome MDR and optimize efficacious tumour ablation and also to identify prognostic biomarkers for neuroendocrine neoplasms (NENs).³⁹¹

Novel drugs targeting the PI3K/AKT/mTOR cascade in various malignancies, including NB, are currently being refined.^{384-387,392-399} The mTOR inhibitor rapamycin (also termed sirolimus) and its analogues (rapalogs), including temsirolimus, everolimus, and ridaforolimus) form a complex with the cytosolic protein FK-binding protein 12 (FKBP12) which attaches directly to mTOR, impeding its function and activating downstream effectors, such as cyclin D1, p21, and HIF1a/b.^{197,397} Of concern, however, is the ability of rapamycin to induce the anti-apoptotic protein, survivin which, in NB, may favour clonal proliferation of resistant cells.²⁰³

Rapamycin and some rapalogs have been approved for clinical trials because of their propensity to inhibit NB cell proliferation.^{388,389} Evaluation of the efficacy of temsirolimus in a phase II trial of children with relapsed or refractory high-grade NB did not produce

encouraging results despite the observation that disease stabilization occurred which makes it a candidate for combination therapy.^{197,400} Similar clinical outcomes have been reported for everolimus in refractory solid NB in paediatric patients.⁴⁰⁰⁻⁴⁰² Preclinical evaluation of mTOR inhibitors that mimic ATP-competitive inhibitors (e.g., INK128/MLN0128, AZD2014, and OSI027) have shown limited potential as inhibitors of NB growth,⁴⁰³ but compounds that target the feedback loops in mTOR/AKT signalling (e.g., MK-2206, an AKT inhibitor) show promise in suppressing tumour growth and increasing survival in mice bearing xenograft NB tumours.⁴⁰⁴ Recent efforts focusing on the development and validation of pharmacodynamic biomarkers to evaluate both the mechanism of action of and proof of concept for drugs that block MYCN and PI3K/AKT/mTOR pathways in children with NB may prove useful in future clinical trials.^{395,399}

A13.2.3 Aurora A Kinase and MDM2 as MYCN Targets

Aurora A kinase (AURKA, a serine/threonine kinase), along with p53 and MDM2, are downstream effectors of *MYCN* that regulate cell cycle progression (particularly during the G2 to M phase transition), the DNA damage response, differentiation and apoptosis in NB^{405,406} AURKA has been implicated in centrosome maturation, spindle assembly and orientation, meiotic maturation and cytokinesis. Targeted inhibition of AURKA leads to deregulation of autophosphorylation and p53 phosphorylation, monopolar spindles and G2-M arrest. Overexpression of AURKA is widespread in solid tumours and associated with resistance to apoptosis, making it a significant focus for the development of anticancer agents, some of which are currently in early-phase NB clinical trials.^{197,407,408}

For example, MLN8237 (alisertib), a reversible AURKA inhibitor, is being investigated in phase I clinical trials by the COG for patients who have experienced relapse. *In vitro* and *in vivo* effects of MLN8237 include apoptosis induction, upregulation of p53 and the tumour suppressor genes p21 and p27.^{37,198,409,410} MLN8054 inhibits N-Myc-dependent transcription, correlating with tumour regression and prolonged survival in a mouse model of MYCN-driven

NB.⁴¹¹ AURKA also increases VEGF secretion and NB angiogenesis.⁴¹² Thus, AURKA is a negative prognostic factor in human NB.⁴¹³ Dose-escalation and combination therapy studies are currently in progress.^{1,11} In a phase 1 trial, alisertib, an oral AURKA inhibitor, in combination with irinotecan and temozolomide, showed promising response and progression-free survival (PFS) rates in patients with advanced NB.⁴¹⁴ Alisertib and a novel pan-AURKA inhibitor, BPR1K653, show potential for the management of patients with MDR1 (ABCB1)-related drug resistance after prolonged chemotherapeutic treatments.⁴¹⁵⁻⁴¹⁷

A13.2.4 Tyrosine Receptor Kinase Neurotrophin Receptor Inhibitors

The tyrosine receptor kinase (Trk) neurotrophin receptors (also referred to as neurotrophic tyrosine receptor kinases)—TrkA/NTRK1, TrkB/ NTRK2 and TrkC/NTRK3 (3 isoforms) are crucial modulators of normal central and peripheral nervous system developmental outcomes (e.g., neuronal differentiation and survival) and NB pathogenesis. Their respective ligands are nerve growth factor (NGF), brain-derived neurotropic factor (BDNF) and neurotrophin-3 (NT3) growth factor.^{12,247,418-421} Overexpression of TrkA correlates with favourable prognosis and suppressed MYCN amplification,^{422,423} whereas upregulation of TrkB and its ligand, BDNF, is associated with aggressive NB—invasion, metastasis, angiogenesis and drug resistance—and unfavourable clinical outcomes.⁴²⁴

Tumours isolated from patients with low-stage and 4S disease frequently express elevated levels of TrkA.¹² It is therefore not surprising that the induction of apoptosis and tumour regression in NBs by targeting neurotrophin receptor pathways such as TrkA and p75^{NTR}—a member of the tumour necrosis factor (TNF) receptor superfamily—is considered a promising therapeutic paradigm. This is supported by observations that TrkA-expressing tumour cells in primary culture will survive and even differentiate in the presence of NGF, but undergo apoptosis in its absence.⁴²⁵⁻⁴²⁷ Noteworthy also, lestaurtinib (CEP-701), has shown proof-of-principle as a small molecule inhibitor of Trk neurotrophin receptors (TrkA, TrkB and TrkC) against TrkB-expressing NB xenografts,⁴²⁸⁻⁴³¹ and in a phase I trial in children with recurrent

and/or refractory neuroblastoma.⁴³² Several second-generation Trk inhibitors are currently in phase I clinical trials or in preclinical development.⁴³³ Interestingly, oncogenic TRK gene fusions are found across multiple tumour types, and those involving NTRK1, NTRK2 and NTRK3, and in-frame deletions or splice variants of NTRK1 signify newer rational drug targets in cancer and are likely to be actionable oncogenes based on preclinical data.⁴³⁴

A13.2.5 Targeted Immunotherapy and Disialoganglioside

Immunotherapy of NB is gaining momentum as a treatment elective to enhance the survival of patients suffering from this challenging paediatric cancer.⁴³⁵⁻⁴³⁸ Targeted immunotherapy of MDR microscopic NB offers an approach which exploits tumour selectivity and minimizes cross-resistance or overlapping side effects (toxicities) with chemotherapy.^{62,436} Disialoganglioside, GD2, is expressed on the surface of tumours of neuroectodermal origin, including NB.^{296,439} Anti-GD2 monoclonal antibodies (mAbs) ablate tumour cells through both complement- and cell-mediated lysis (antibody mediated cell cytotoxicity or ADCC), and are therefore exceptional candidates for targeted immunotherapy since they have specificity, high affinity and are relatively nontoxic.²⁹⁶

A number of anti-GD2 mAbs ± GM-CSF ± CRA have been tested in clinical trials and favourable therapeutic outcomes were reported.^{11,151,440,441} However, a recent trial inferred a lack of survival advantage with ASCT in HR-NB consolidated by anti-GD2 immunotherapy and CRA, adding to the complexity of developing targeted immunotherapies for NB.³⁰⁶ Targeting NB immune escape pathways, intrinsic NB cell defects such as impaired expression of the human leukocyte antigen (HLA) class I related antigen processing machinery and functional alterations of the tumour microenvironment (TM) induced by NB cell-derived immunosuppressive molecules such as human major histocompatibility complex (MHC) class I chain-related gene A (MICA) and HLA-G are critical considerations of such therapeutic interventions.⁴⁴²

A13.2.6 Angiogenesis and VEGF Signalling Inhibitors

Angiogenesis or neovascularization—one of the hallmarks of cancer—encompasses the sprouting of new blood vessels in tumours that enable them to grow, survive and metastasize before their metabolic demands are restricted due to diffusion limits of oxygen and nutrients in the tumour microenvironment.^{145-148,443} Vascular endothelial cell growth factor (VEGF) is the most potent activator of angiogenesis and comprises six members (VEGF-A, VEGF-B, VEGF-C, VEGF-D, VEGF-E and placental growth factor) that bind differentially with three cell surface RTKs, the VEGFRs, or a second class of non-signalling co-receptors, the neuropilins.⁴⁴³ In NB, elevated expression of pro-angiogenic factors correlates with advanced stage disease while low vascular tumour density is associated with non-metastatic localized disease and favourable prognosis.^{62,444,445} Accordingly, inhibition of angiogenesis has long been regarded as a promising line of attack in the management HR-NB.⁴⁴⁶⁻⁴⁵⁰ Current anti-vascular NB therapies combining multimodal antiangiogenic, anti-vasculogenic mimicry and anti-lymphangiogenic strategies may yield increased efficacy.^{278,391,446,448,449,451-462}

A13.2.7 The PI-3 Kinase-Akt-MDM2-Survivin Signalling Axis in High-Risk Neuroblastoma

The phosphatidylinositol-3 (PI-3) kinase-Akt pathway is a central convergent molecular conduit, and PI-3 kinase is a multiplex signalling hub downstream of various growth factor receptors, including TrkB, VEGFR, PDGFR and EGFR.⁴⁶³ Activation of the PI3K/Akt pathway in NB correlates with poor patient prognosis, and the forkhead transcription factor, FOXO3a, is a key target of the PI3K/AKT pathway in NB. FOXO3a expression was shown to be upregulated in low-stage NB and normal embryonal neuroblasts, but ablated in late-stage NB. Thus, inactivation of FOXO3a by AKT is essential for neuroblastoma cell survival.⁴⁶⁴ The mammalian target of rapamycin (mTOR) protein (section A13.2.2 on mTOR inhibitors) is currently being regarded as a potential therapeutic target in NB patients.^{203,391,463,465-467} Well-defined adjustments are associated with mitochondrial proteins, e.g., VDAC1/Porin protein, an integral part of the mitochondrial permeability transition pore complex, during loss of

mitochondrial membrane potential with subsequent cytochrome c release and caspase-3 activation. VDAC1 is negatively regulated by the PI3K/Akt pathway via GSK3beta and inhibition of GSK3beta is activated when Akt is blocked.⁴⁶⁸ Similarly, a recent study showed that guanosine offers protection against mitochondrial oxidative stress by a signalling pathway that implicates PI3K/Akt/GSK-3beta proteins and induction of the antioxidant enzyme, haem oxygenase-1.⁴⁶⁹ Also, the PI3K/Akt pathway is obligatory for RA-induced NB cell differentiation, and may be exploited as a novel therapeutic strategy against poorly differentiated NB.^{394,395,467,470}

A13.2.8 Gastrin-Releasing Peptide Receptors

Gastrin-releasing peptide (GRP) receptors (GRPR), a member of G-protein coupled receptor family, are overexpressed in undifferentiated NB.^{471,472} The decreased expression of the tumour suppressor protein PTEN in aggressive undifferentiated NB is associated with an increase in GRP binding capacity, as a result of GRP-R overexpression.⁴⁷³⁻⁴⁷⁵ It has been suggested that inhibition of the PTEN tumour suppressor gene may be an important regulatory mechanism involved in GRP-induced cell proliferation in NB which offers promising scenarios for the use of radiolabelled and cytotoxic GRP analogues and antagonists for cancer diagnosis and therapy.⁴⁷⁶⁻⁴⁷⁹ GRPR transactivates the epidermal growth factor receptor (EGFR) and may thus modulate therapeutic responses to EGFR inhibitors, e.g., gefitinib.⁴⁸⁰ GRP upregulates proangiogenic IL-8 expression in an Ets1-dependent manner, implying a key role during GRP-induced NB angiogenesis and metastasis,^{481,482} and its promise as a NB biomarker and gene silencing therapeutic target.⁴⁸³⁻⁴⁸⁶

A13.2.9 Anaplastic Lymphoma Kinase

Germline ALK activating mutations have been implicated in the majority of hereditary NB and somatic ALK activating mutations have also frequently been observed in sporadic cases of advanced NB. Accordingly, gain of function mutations in the gene encoding the anaplastic lymphoma kinase (ALK) is currently deemed the most frequent druggable mutations identified

in NB. Preclinical studies warrant the notion of an oncogene addiction of NB cells to mutated ALK and corroborate that ALK inhibitory therapy effectively blocks tumour models. Recently, a paediatric phase I trial for the first approved ALK inhibitor, crizotinib, illustrated significant antitumoural efficacy in NB patients. A successive international phase I study with the second generation ALK inhibitor, LDK-378, has been launched that makes ALK inhibitory therapy accessible to paediatric patients.⁴⁸⁷

However, crizotinib is not as effective in blocking the activity of ALK when activating mutations are present within its kinase domain, as with the F1174L mutation. A new ALK inhibitor, AZD3463, effectively suppresses the proliferation of NB cell lines with wild type ALK (WT) as well as ALK activating mutations (F1174L and D1091N) by blocking the ALK-mediated PI3K/AKT/mTOR pathway and induces apoptosis and autophagy. Moreover, AZD3463 synergistically enhances the cytotoxicity of doxorubicin on NB cells and shows significant therapeutic efficacy on the growth of NB tumours with WT and F1174L activating mutation ALK in orthotopic xenograft mouse models. These results indicate that AZD3463 is a promising therapeutic agent in the treatment of NB.⁴⁸⁸

Interestingly, the ALK/ROS1 inhibitor, PF-06463922, ablates primary resistance to crizotinib in ALK-driven NB. PF-06463922 has high potency across ALK variants and inhibits ALK more effectively than crizotinib *in vitro*. Essentially, PF-06463922 causes complete tumour regression in both crizotinib-resistant and crizotinib-sensitive xenograft mouse models of NB, as well as in patient-derived xenografts harbouring the crizotinib-resistant F1174L or F1245C mutations. Hence, PF-06463922 shows potential to reverse crizotinib resistance and exert significant activity as a single targeted agent against F1174L and F1245C ALK-mutated xenograft tumours, while also inducing responses in an R1275Q xenograft model. These results provide the reasoning to advance PF-06463922 into clinical trials for treatment of patients with ALK-mutated NB.^{489,490}

A13.2.10 Future Therapeutic Perspective

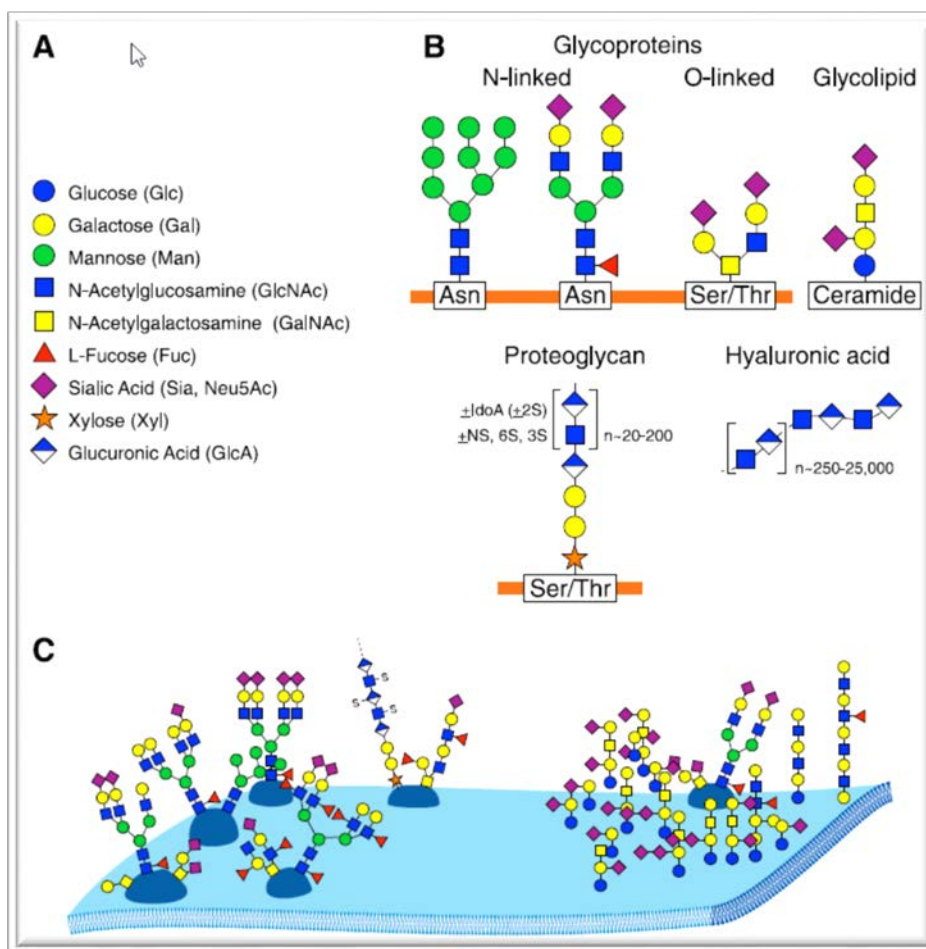
In spite of the knowledge provided in the preceding sections, NB remains a therapeutic enigma. As we are driven to improve outcomes and survival, the ideal therapy also remains elusive. However, there are many fronts on which to attack, and it seems unquestionable that the cure will require a multimodal approach. Part of the solution is to effectively eradicate MRD, as this appears to put patients at highest risk for relapse and progression. Selected clinical trials on current interventions in NB have recently been collated, but most are recruiting and results may not be available yet (www.clinicaltrials.gov).

SECTION B: GLYCOBIOLOGY AND GLYCOMICS OF NEUROBLASTOMA

B1. Orientation to Glycans

Glycans are defined by the International Union of Pure and Applied Chemistry (IUPAC, <https://iupac.org/> and <http://www.chem.qmul.ac.uk/iupac/>) as compounds that consist of monosaccharides or oligosaccharides linked by N- or O glycosidic bonds). Glycans and complementary glycan-binding proteins (GBPs) are indispensable metabolic, structural and modulatory components of various cell functions, including cell-cell communication, cell-matrix interactions, immunity, cancer pathogenesis and progression.⁴⁹¹ The term glycan may also be used to refer to the carbohydrate portion of a glycoconjugate, such as a glycoprotein, glycolipid, or a proteoglycan (Figure 1.14).

The human glycome stems from 9 building blocks that are merged by enzymes (*writers*: glycosyltransferases, glycosidases and glycan modifying enzymes) with precise and regulated biosynthetic functions into a wide diversity of glycan patterns (Figure 1.15) that are functionally read by various human GBPs (*readers*).⁴⁹²⁻⁴⁹⁵ The importance of glycan recognition, for example, in infection and immunity, and advances in our understanding and technologies in the field of glycobiology,^{496,497} have already led to the design and use of glycan mimetic anti-infective and anti-inflammatory drugs.^{492,498-501}



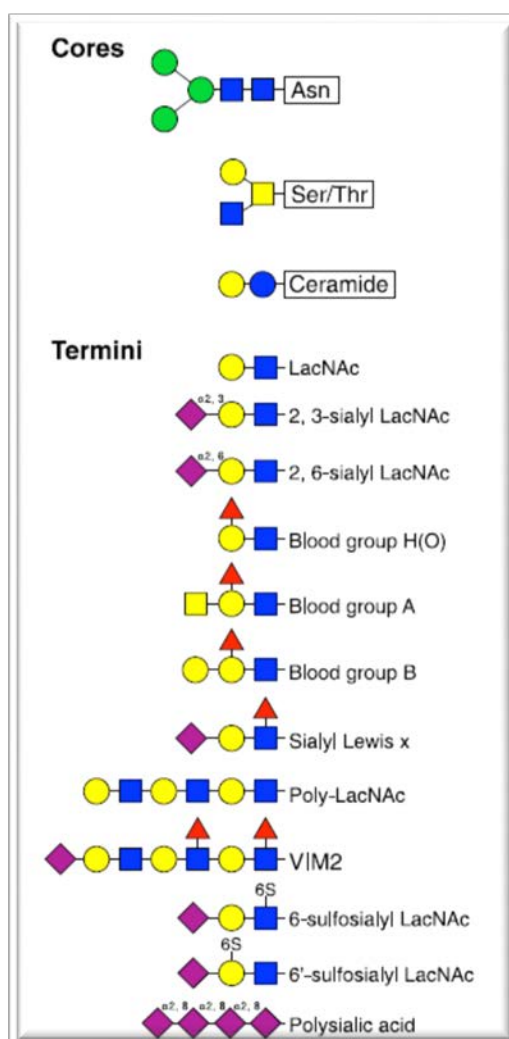
Major human glycans. (A) The 9 sugars that comprise most of the human glycome, with their broadly accepted symbol representations.⁵⁰² (B) Major classes of human glycans. Linkage details (hydroxyl attachment sites and anomeric configurations at each glycosidic bond) that are keys to structural recognition are omitted here for simplicity. Representative asparagine (N-linked) and serine or threonine (O-linked) glycoprotein structures, a glycosphingolipid (ceramide-linked), a proteoglycan (most frequently O-linked), and hyaluronic acid (HA) (unlinked) are shown. (C) A schematic representation of glycans on a cell surface. Notable features important for understanding glycan recognition include varied glycan branching patterns, variations in terminal glycan structures, and the tendency of glycans to form distinctive, lateral glycan patches.

Source:⁴⁹² Schnaar RL. Glycobiology simplified: Diverse roles of glycan recognition in inflammation. *Journal of Leukocyte Biology* 2016;99(6):825-838, with permission from *Journal of Leukocyte Biology*. See Appendix 10 for copyright clearance.

Figure 1.14: Major human glycans

Glycans are significant regulators of biological homeostasis,⁵⁰³ playing pivotal roles in protein folding, trafficking and stability,^{504,505} and in vertebrate development, morphogenesis and organogenesis,⁵⁰⁶⁻⁵⁰⁸ and cellular senescence and human aging.⁵⁰⁹ Inside cells, protein glycosylation, conceivably in unison with protein phosphorylation, regulates key signal transduction cascades,⁵⁰³ intercellular communication,⁵¹⁰ pathogen recognition and

immunological differentiation of self from non-self.^{492,511,512} Moreover, the glycosylation state of both cell-surface proteins and lipids are altered in response to external stimuli and internal cellular dysfunction.⁵¹³ Thus, the dynamics and profiles of glycoproteins and glycolipids reflect the cell's physiological and pathological (disease) status.⁵¹⁴⁻⁵¹⁸



Theme and variation in the human glycome. Glycan recognition often involves variations of terminal glycan structures that are attached to core structures (linkage details are omitted for simplicity). The upper panel provides examples of the invariant, N-linked glycoprotein pentasaccharide core, one of several serine/ threonine-linked glycoprotein cores, and a common glycosphingolipid (ceramide-linked) core. The lower panel provides a sampling of terminal structures. A representation of how these might be grouped on a cell surface is shown in Figure 1.14-C.

Source:⁴⁹² Schnaar RL. Glycobiology simplified: Diverse roles of glycan recognition in inflammation. *Journal of Leukocyte Biology* 2016;99(6):825-838, with permission from *Journal of Leukocyte Biology*. See Appendix 10 for copyright clearance.

Figure 1.15: Theme and variation in the human glycome

B2. Protein Glycosylation in Neuroblastoma

B2.1 General Principles of Glycosylation

Cell-surface and soluble proteins of the secretory pathway are post-translationally glycosylated in the ER.⁵¹⁹ Generally, glycans on membranes, extra cellular matrix (ECM) and secreted proteins are found covalently attached to a protein core at asparagine Asn (N-glycosylation) or at serine/threonine residues (O-glycosylation) (Figure 1.14).^{520,521} Glycosaminoglycans (GAGs) are O-linked glycans initiated by a highly conserved tetrasaccharide (GlcA- β 1,3-Gal- β 1,3-Gal- β 1,4 Xyl- β) and classified by the configuration of their disaccharide repeats that consist of either sulfated or non-sulfated monosaccharides.^{499,500,522} Typical GAGs are chondroitin sulfate, keratan sulfate, dermatan sulfate and heparan sulfate.

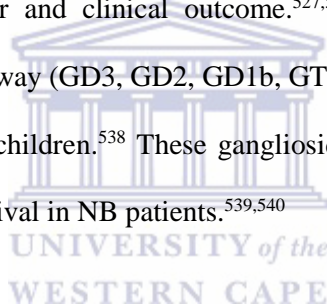
A glycoprotein with one or more GAG chains extending from its protein core is called a proteoglycan which exists as secreted, transmembrane or glycosylphosphatidylinositol (GPI)-anchored units.^{523,524} Hyaluronic acid, a GAG-like polysaccharide of the ECM, is the only glycan that is not linked to protein or lipid.^{525,526} N-linked glycosylation has been correlated with several physiological and pathological processes such as protein folding and conformation, oligomerization, cell-cell interactions, targeting proteins to sub-cellular or extracellular sites (<http://themedicalbiochemistrypage.org/glycoproteins.php#nglycans> for a quick glance at glycans, glycoproteins and glycosylation).

B2.2 Gangliosides

Most tumour cells, including those of neuroectodermal cell origin, have upregulated levels of gangliosides.^{527,528} Gangliosides also accumulate in activated glia in the developing brain,⁵²⁹ and may have neuroprotective roles via activation of microglia and astrocytes in response to acute ethanol concentrations in the neonatal brain. However, chronic ethanol exposure can induce an inappropriate proinflammatory glial reaction and neurotoxicity.⁵³⁰ Gangliosides are glycosphingolipids which comprise of a carbohydrate chain bearing one or several sialic acid

(N-acetyl-neuraminic acid) residues and a lipid portion (ceramide backbone), which attaches (anchors) the ganglioside molecule to the cell membrane.^{531,532} Figure 1.16 illustrates the consecutive glycosylation steps in ganglioside biosynthesis which involves two primary pathways indicated as “a” (GM2, GM1a, and GD1a) and “b” (GD3, GD2, GD1b, GT1b and GQ1b), from a common precursor (GM3). GM1a/GD1b synthase (UDP-Gal:betaGlcNAc-beta-1,3-galactosyltransferase) is the key enzyme in ganglioside biosynthesis and has been implicated in human NB.⁵³³

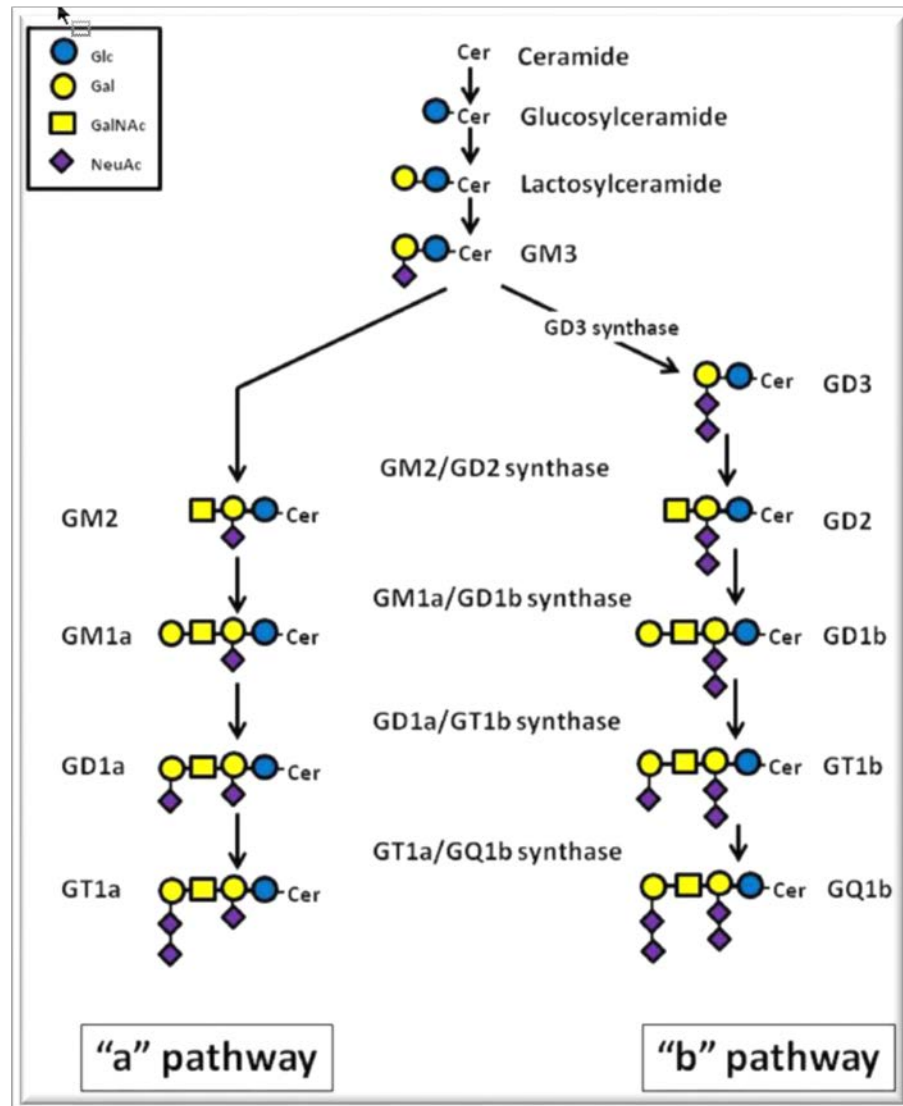
Even though gangliosides are predominantly expressed on tumour cell surfaces, they may be shed into the tumour microenvironment and ultimately appear in the patient’s plasma to induce the production of anti-glycan antibodies.^{527,534-536} In NB, ganglioside signatures or arrays may influence tumour behaviour and clinical outcome.^{527,537} For instance, elevated levels of gangliosides of the “b” pathway (GD3, GD2, GD1b, GT1b, GQ1b) are prevalent in infant NB in contrast to NB in older children.⁵³⁸ These gangliosides correlate with an aggressive NB phenotype and reduced survival in NB patients.^{539,540}



Understandably, complex gangliosides have stimulated interest as diagnostic biomarkers to predict clinical outcome, to stratify NB patients for targeted anticancer therapy and to monitor efficacy of treatment.⁵⁴¹ Thus far, retinoic acid has proved useful for maintenance therapy of disseminated NB as it induces a remarkable shift from synthesis of simple gangliosides toward predominant expression of structurally complex “a” and “b” pathway ganglioside molecules.^{527,542}

Targeted immunotherapy of NB with antibodies directed against disialoganglioside (GD2) has been discussed in detail in section A13.2.5. Various other glycans such as polysialic acid (PSA), galectin-1 (Gal-1), and other related processes such as N- and O-protein glycosylation, glycosyltransferases and glycosidases, have been amply documented for their respective involvement in NB glycopathobiology and, therefore, share parallel platforms and themes with

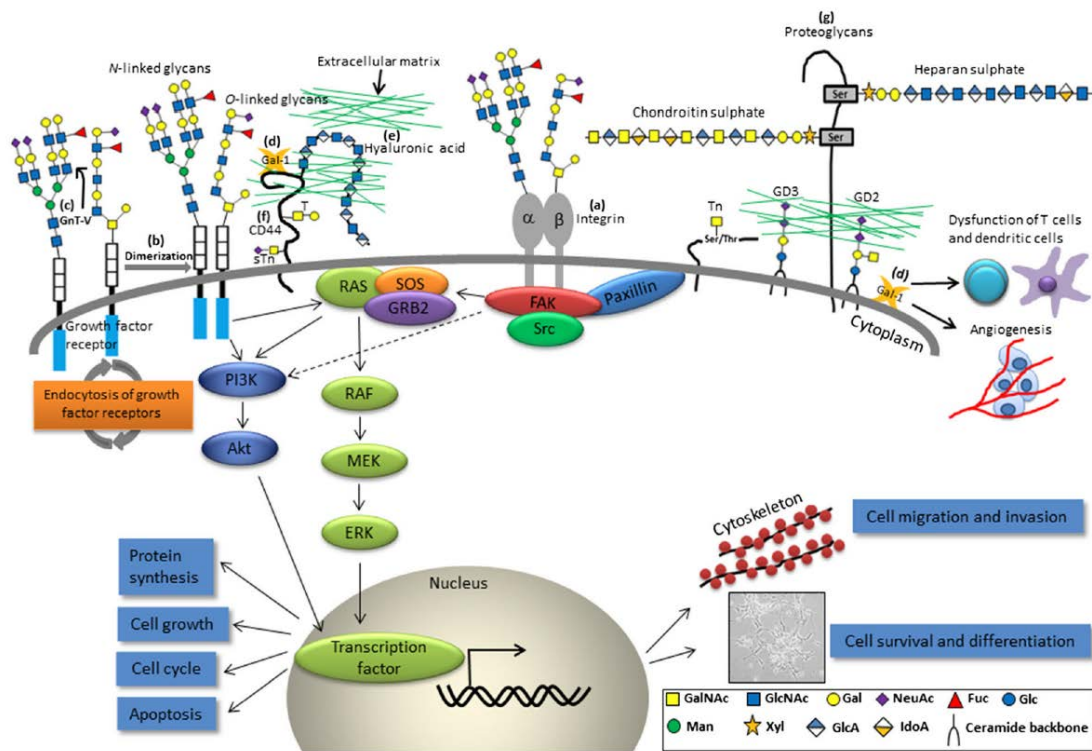
the gangliosides.^{491,492,515,516,527,529,543-546} For the sake of objectivity and some degree of inclusiveness, some of these highlights are encapsulated in Figures 1.17 and 1.18.



Each ganglioside is structurally more complex than its precursor molecule, and the stepwise addition of monosaccharide or sialic acid (N-acetyl-neuraminic acid) residues in the Golgi apparatus is catalyzed by the same specific membrane-bound glycosyltransferases in both pathways.⁵⁴⁷ Gangliosides can also be grouped into structurally simple (SG) and complex (CG) molecules. The enzyme GM1a/GD1b synthase (UDP-Gal:betaGlcNAc-beta-1,3-galactosyltransferase) converts its substrates, the simple gangliosides, GM2 and GD2, into the corresponding initial complex ganglioside products, GM1a and GD1b.

Source:⁵²⁷ Berois N, Osinaga E. Glycobiology of neuroblastoma: Impact on tumour behavior, prognosis, and therapeutic strategies. *Frontiers in Oncology* 2014;4:114, with permission from *Frontiers in Oncology*. See Appendix 11 for copyright clearance.

Figure 1.16: Schematic representation of the major ganglioside biosynthesis pathways



(a) β 1,4-N-acetylgalactosaminyltransferase 3 (B4GALNT3) and β 1,4-galactosyltransferase 3 (B4GALT3) exhibit differential effects on malignant phenotypes by modification of β 1 integrin in NB cells; (b) N-acetylgalactosaminyltransferase 2 (GALNT2) modifies O-glycans on IGF-1R, thereby suppressing IGF-1-induced IGF-1R dimerization and downstream signaling; (c) N-acetylglucosaminyltransferase V (GnT-V) modulates the sensitivity of NB to apoptosis; (d) Gal-1 promotes attachment of NB cells to the extracellular matrix (ECM) and endothelial cells through binding to CD44. Besides, Gal-1 may dampen the function of T cells and dendritic cells (DC) as well. Glycosaminoglycans present as (e) free polysaccharides (hyaluronic acid), a major counterreceptor for (f) CD44, or (g) as part of proteoglycans (heparan sulfate and chondroitin sulfate). *GalNAc* N-acetylgalactosamine, *GlcNAc* N-acetylglucosamine, *Gal* galactose, *NeuAc*, N-acetylneuraminic acid, *Fuc* fucose, *Glc* glucose, *Man* mannose, *Xyl* xylose, *GlcA* glucuronic acid, *IdoA* iduronic acid.

Source:⁴⁹⁴ Ho WL, Hsu WM, Huang MC, Kadomatsu K, Nakagawara A. Protein glycosylation in cancers and its potential therapeutic applications in neuroblastoma. *Journal of Hematology & Oncology* 2016;9(1):100.4, with permission from *BioMed Central* in terms of the Creative Commons Attribution 4.0 International License (<http://creativecommons.org/licenses/by/4.0/>), which permits unrestricted use, distribution, and reproduction in any medium, provided you give appropriate credit to the original author(s) and the source, provide a link to the Creative Commons license, and indicate if changes were made.

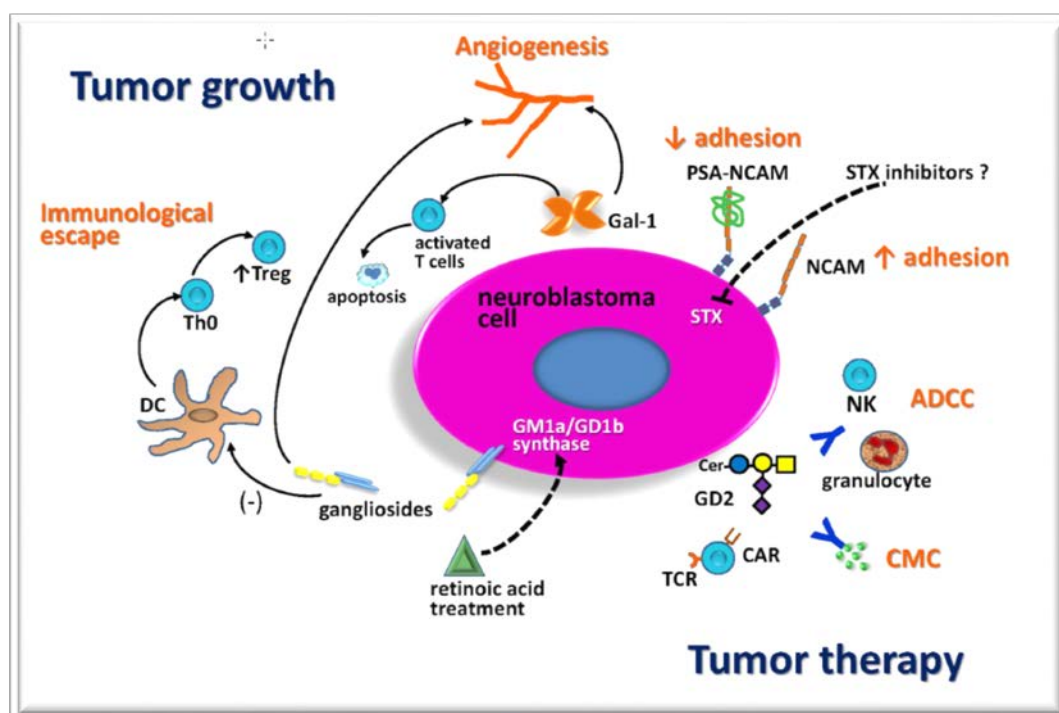
Figure 1.17: Altered glycans and related pathophysiological events involved in NB progression

B2.3 Intercellular Adhesion Molecule-2

Cell adhesion molecules (CAMs, e.g., selectins, integrins, cadherins and immunoglobulin-like CAMs) comprise four ubiquitously occurring families of glycosylated, membrane-bound proteins involved in multiple cellular processes, including cell-cell communication, cell motility, inside-out and outside-in signalling, tumourigenesis, angiogenesis and metastasis.⁵⁴⁸⁻

⁵⁵¹ In the context of NB, it was found that intercellular adhesion molecule-2 (ICAM-2, a 55-60

kDa transmembrane glycoprotein), also known as CD102 (Cluster of Differentiation 102), which has six N-linked glycosylation sites at Asn47, Asn82, Asn105, Asn153, Asn178 and Asn87, suppressed tumour cell motility and dissemination, but not tumourigenic potential, *in vivo* in a murine model of metastatic NB.^{527,552,553} N-glycosylation of ICAM-2 is critical for these effects,⁵⁵³ and ICAM-2 confers a non-metastatic phenotype in NB cells by interaction with α -actinin.⁵⁵⁴



Treatment with retinoic acid markedly enhances the activity of GD1b/GM1a synthase, resulting in increased expression of complex gangliosides, associated with less-aggressive tumours. NB gangliosides promote dendritic cells (DC) to develop with decreased costimulatory signals and IL-12 production. These DC promote differentiation of human T-helper type 0 (Th0) cells toward regulatory T-cells (Treg). Galectin-1 (Gal-1) secreted by NB also contributes to the immunosuppressive tumour microenvironment, limiting T-cell survival and impairing DC function. Both, gangliosides and Gal-1 contribute to tumour angiogenesis. The presence of polysialic acid (PSA) on neural cell-adhesion molecule (NCAM) reduces NCAM-mediated adhesion processes promoting NB cell migration. The fact that sialyltransferase (STX) is the dominant polysialyltransferase for PSA biosynthesis in NB suggests that this enzyme could be a good therapeutic target. Disialoganglioside (GD2) is a relevant antigen for NB immunotherapy. Anti-tumour activity of anti-GD2 antibodies is mediated by antibody-dependent cell-mediated cytotoxicity (ADCC) in the presence of human natural killer (NK) cells and granulocytes, as well as by complement-mediated cytotoxicity (CMC). Anti-GD2 chimeric antigen receptor T-cells (CAR-T-cells) activity could induce NB tumour regression.

Source:⁵²⁷ Berois N, Osinaga E. Glycobiology of neuroblastoma: Impact on tumour behavior, prognosis, and therapeutic strategies. *Frontiers in Oncology* 2014;4:114, with permission from *Frontiers in Oncology*. See Appendix 11 for copyright clearance.

Figure 1.18: Neuroblastoma glyco-biology impact on tumour growth and antitumour therapy

B2.4 Anaplastic Lymphoma Kinase

The role of ALK as a predisposing gene in NB has been discussed in detail in section A11.4.3. ALK has 16 highly conserved putative sites of N-linked glycosylation in its extracellular domain.⁵²⁷ Previous studies have observed that perturbation of N-linked glycosylation ablates ALK phosphorylation and blocks downstream pro-survival signalling and cell viability in NB cell lines selected for mutated or amplified ALK,⁵⁵⁵ raising hopes that inhibition of this post-translational modification could be applied usefully in NB targeted therapy.

B2.5 Cell-Surface Mucin-Type O-Glycans

Cell-surface mucins are glycoproteins with large branches of O-linked oligosaccharides. The most abundant mucin-type glycoproteins typically contain an α -N-acetylgalactosamine residue (GalNAc) covalently linked to the alpha hydroxyl group of Ser/Thr residues.^{494,517} Such linkages are catalyzed by UDP-GalNAc:polypeptide-N-acetyl-galactosaminyl-transferases (GalNAc-T). GalNAc-T is a multigene family of 20 or more isoenzymes (<http://www.cazy.org>).⁵⁵⁶ Several carcinomas express truncated O-glycosylated tumour-associated glycan antigens (terminal structures arising from sialylation and fucosylation) such as (Tn, sTn, T, and sLe^{a/x} and Thomsen–Friedenreich antigen (TF) which correlate with adverse outcome and poor prognosis in cancer patients, thus making them candidate therapeutic targets.^{494,514,527,557-559}

In the case of NB, recent evidence suggests that the expression of enzymes encoded by the GALNT [UDP-N-acetyl- α -D-galactosamine:polypeptide N-acetylgalactosaminyltransferase (GALNAC-T)] gene family which catalyze the first step in O-glycosylation, correlates with improved overall survival in low- and high-risk groups and improved clinical outcome (overall and disease-free survival) in low-risk NB patients. Hence, GALNT9 expression may be a valuable prognostic marker for personalized therapy.⁵⁶⁰ Likewise, elevated expression of β 1,3-N-acetylglucosaminyltransferase-3 (B3GNT3), the enzyme responsible for adding GlcNAc to core 1 (T antigen), predicted a favourable prognosis in NB patients distinct from other

prognostic markers. B3GNT3 overexpression also interfered with T antigen production and malignant signatures such as migration and invasion of SK-N-SH cells, while B3GNT3 knockdown enhanced these phenotypes of SK-N-SH cells. Additionally, B3GNT3 expression abolished phosphorylation of focal adhesion kinase (FAK), Src, paxillin, Akt and ERK1/2 (Figure 1.17). Thus, B3GNT3 as a modulator of mucin-type O-glycosylation and signalling in NB cells, may be a precise clinical predictor of NB behaviour and therapeutic outcome.⁵⁶¹

B2.6 Polysialic Acid

Polysialic acid (PSA) exemplifies a distinctive post-translational modification of the neural cell adhesion molecule (NCAM).^{527,562} PSA assembly involves extended linear homopolymerization⁵⁶³ of 150-200 α 2,8-linked sialic acids on N-glycans of the fifth immunoglobulin-like domain of NCAM. During normal development, PSA mediates cell migration and axonal growth, but in undifferentiated NB, it promotes NB cell proliferation and metastatic potential.⁵⁶⁴ PSA expression is upregulated in high-risk NB. In the Golgi apparatus, two homologous polysialyltransferases, ST8SiaII (STX) and ST8SiaIV (PST), catalyze the synthesis of variable amounts of PSA in tumours.⁵⁶⁵

The ST8SiaII gene is expressed predominantly during embryonic development and thought to be silent in normal tissue, but is highly expressed in metastatic NB.^{566,567} ST8SiaIV is the major polysialyltransferase in the adult brain.⁵²⁷ STX has attracted considerable interest as a molecular marker and therapeutic target for metastatic NB⁵⁶⁸ as borne out by recent efforts aimed at reducing STX-mediated polysialylation of NCAM using cytidine monophosphate (CMP)⁵⁶⁹ and inhibiting migration of IMR-32 NB cells with the sialic acid precursor, ManNProp.⁵⁶⁷

B2.7 Lectins (Glycan-Binding Proteins)

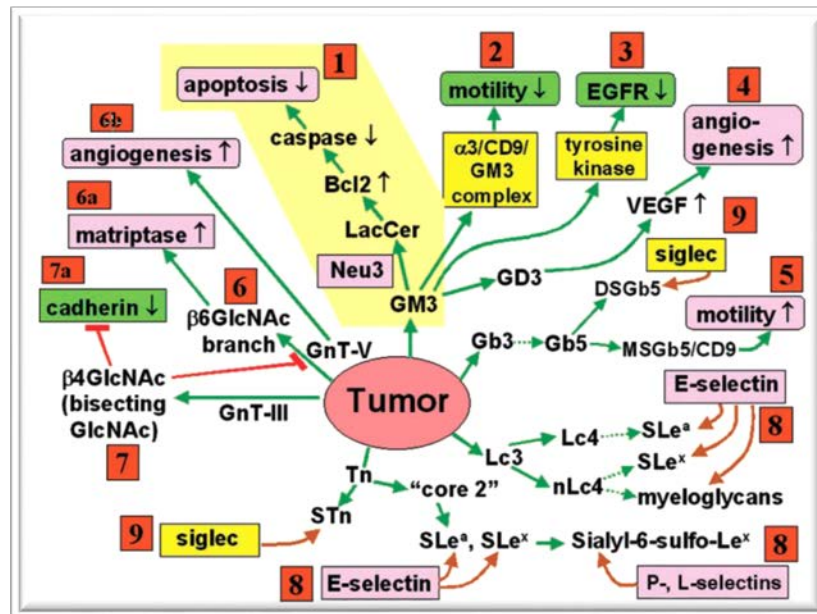
Three main categories of lectins, namely, siglecs (sialic acid binding Ig-like lectins), galectins and selectins are GBPs that have a high specificity for sugar moieties. Endogenous lectins are

involved in processes such as cell-cell recognition, cell adhesion and motility, pathogen-host recognition, and tumour progression and metastasis. Many lectins are expressed on the surface of immune and endothelial cells or exist as ECM components and cytoplasmic adhesion molecules.⁵⁷⁰ Normal glycans of colonic epithelial cells, for example, suppress cyclooxygenase-2 expression by resident macrophages, thus maintaining immunological homeostasis in mucosal membranes, whereas loss of immunosuppressive glycans by impaired glycosylation during colonic carcinogenesis triggers inflammatory destruction of colonic mucosa.

Siglec-7 and -9, expressed on resident macrophages in the colonic lamina propriae bind to ligands di-sLe^a and 6-sulfo sLe^x, but loss of this function occurs during malignant transformation coupled with a gain of expression of sLe^a and sLe^x which have no siglec ligand activity.⁵⁷¹ Siglec-7 is expressed mainly on natural killer (NK) cells and suppresses NK cell-mediated cytotoxicity towards target cells overexpressing α 2,8-disialic acid-bearing ganglioside, GD3 (see Figure 1.16).^{572,573} Malignant melanoma and NB overexpress GD3, a cancer signature that may confer on these tumours the ability to evade immunosurveillance and elimination by NK cells. This concept may indeed be exploited in the targeted inhibition of siglec-7 and NB metastasis.⁵⁷³

In recent years, galectin-1 (Gal-1) has gained prominence as a burgeoning target in NB translational therapeutics.^{494,527} Gal-1 is an adaptable regulator of multiplex signalling pathways such as tumour–host interaction,⁵⁷⁴ angiogenesis,⁵⁷⁵ promotion of immunosuppression by T cell apoptosis and impairment of dendritic cell (DC) function in numerous cancers, including NB.⁵⁷⁶⁻⁵⁷⁸ Aggressive NB tumours express high levels of neurotrophin receptor TrkB (section A13.2.4) and Gal-1 which are not only coupled with invasive behaviour and high metastatic potential, but also associated with therapy resistance and thus poor prognosis⁵⁷⁹. These Gal-1 phenotypic features of NB are meticulously being probed for their glycan-based therapeutic potential.^{494,527,546,578,580}

Figure 1.19 summarizes how glycosylation/glycans, in particular, the selectins cast the hallmarks of the cancer phenotype.



Tumour cell malignancy is defined by several key phenotypes: apoptosis (route 1), motility (routes 2 and 5), EGF receptor tyrosine kinase (route 3), angiogenesis (routes 4 and 6b), matriptase (matrix-destroying enzyme) activity (route 6a), self-adhesion (through cadherin) (route 7a), adhesion to ECM (through integrin), adhesion to ECs and platelets (through E- or P-selectin) (route 8), adhesion to blood cells and other parenchymatous cells (through siglecs) (route 9). Each phenotype is up- or down-regulated (1, 2) by different status of glycosylation. Phenotypes with 1 or 2 and green color inhibit tumour invasiveness. Those with 1 or 2 and pink color promote invasiveness. Glycosyl epitopes capable of binding to specific ligands (pink color without arrow) promote invasiveness. Ligands with yellow color have variable or unclear effect on invasiveness. Note that a given phenotype is produced by different glycosylations, and a given glycosylation produces different phenotypes. Phenotypic changes have cooperative effects on malignancy. For example, GM3 inhibits motility through $\alpha 3/CD9$ complex and also inhibits EGF receptor tyrosine kinase (routes 2 and 3). Reduction of GM3 inhibits apoptosis (route 1), but promotes motility and proliferation (negative route 2 and 3 effect). Essentially all glycosylation pathways catalyzed by multiple glycosyltransferases and their genes are well established (for review see 2). However, the mechanism by which each type of glycosylation affects the various phenotypes remains to be studied. Structures of GSLs are abbreviated according to International Union of Pure and Applied Chemistry–International Union of Biochemistry nomenclature recommendations. S, sialyl; MS, monosialyl; DS, disialyl.

Source:⁵⁸¹ Hakomori S. Glycosylation defining cancer malignancy: New wine in an old bottle. *Proceedings of the National Academy of Sciences of the United States of America* 2002;99(16):10231-10233, with permission from PNAS, <http://www.pnas.org/site/aboutpnas/rightperm.xhtml>, accessed 4 December 2016.

Figure 1.19: Glycosylation defining malignancy—invasive and metastatic phenotype of tumours

B2.8 Glycosyltransferases

Tumour cells exhibit striking changes in cell-surface glycosylation as a consequence of dysregulated glycosyltransferases and glycosidases.⁵⁸² Aberrant glycosylation is a cancer hallmark which correlates with differential expression of cell-surface and cytosolic glycans

and tumour-associated antigens.^{527,560} Glycosyltransferases have clinical relevance as cancer biomarkers for different tumours, markers for minimal residual disease (MRD) detection, risk group assignment and as prognostic predictors (<https://pob.abcc.ncifcrf.gov/cgi-bin/JK>).^{494,527,566,583}

Elevated β 1,6-N-acetylglucosaminyltransferase V (GnT-V) expression, for example, predicts a favourable prognosis and treatment outcome in NB.⁵⁸⁴ The expression, clinical relevance and functional significance of several glycosyltransferases, including β 1,4-N-acetylgalactosaminyltransferase (GD2synthase), sialyltransferase (STX or ST8SiaII), β 1,3-N-acetylglucosaminyltransferase 3 (B3GNT3), UDP-polypeptide GalNAc-transferase 13 (GalNAc-T13, GALNT13), β 1,4-galactosyl-transferase 3 (B4GALT3), and N-acetylgalactosaminyltransferase 2 (GALNT2), in NB and other cancers, have been widely documented.^{494,527,561,585-587}

B2.9 ATP-Binding Cassette Multidrug Transporters

Direct and coordinate transcriptional targets of MYCN (section A11.4.5) include several of the ATP-binding cassette (ABC) transporters—ABCB1 (P-glycoprotein/P-gp/MDR1), ABCG2 (breast cancer resistance protein/BCRP) and ABCC1 (multidrug resistance protein 1/MRP1), ABCC3 (MRP3) and ABCC4/MRP4.^{311,328,588} The expression of these multidrug resistance (MDR) transporters are strongly prognostic of NB outcome (section A13.1.7) since they extrude a wide array of structurally- and functionally-related or -unrelated chemotherapeutic drugs.^{311,329,330}

Moreover, endogenous substrates of MDR transporters such as bioactive lipid mediators (e.g., prostaglandins and leukotrienes) may modify normal neural development by switching on processes (angiogenesis, cell signalling, inflammation, proliferation, and migration and invasion) that promote NB initiation and progression. The ABC transporters are thus promising candidates for therapeutic suppression in HR-NB, the rationale behind increasing drug

bioavailability (therapeutic efficacy) in refractory tumours which overexpress these glycans.³²⁶ The glycosylation of ABCB1 (P-glycoprotein/P-gp) has been studied widely and will thus be considered as the representative drug transporter. Previous studies have shown that the most strongly upregulated genes associated with acquired drug resistance and an important cause of NB treatment failure was *GALNT13*, followed by *ABCB1 (MDR1)*.^{527,589,590} *GALNT13* encodes the UDP-GalNAc:polypeptide GalNAc-transferase-13 (GalNAc-T13), constitutively expressed in neural tissue.^{583,591} It has been demonstrated unequivocally that inhibition of protein glycosylation reverses the MDR phenotype of cancer cell lines.⁵⁹² Likewise, inhibition of N-linked glycosylation impairs ALK phosphorylation and perturbs pro-survival signalling in NB cell lines.⁵⁵⁵ Glycosylation of P-gp corresponds to the *en bloc* transfer of the oligosaccharide portion of a lipid-linked oligosaccharide onto the acceptor asparagine of nascent proteins, typical for all N-glycans.⁵¹³

P-gp is synthesized as a 140–150 kDa precursor protein which is escorted by chaperones (calnexin and Hsp70) in the ER lumen to the Golgi. P-gp is modified post-translationally by N-glycosylation encompassing various sugar moieties—a process essential for its destination docking (dynamic integration into the membrane) and, ultimately, mature functioning (drug efflux pump activity), as inferred from experiments with cDNA encoding N-glycosylation-deficient P-gp showing that the immature or non-glycosylated protein is trapped in subcellular compartments.^{593,594} Tunicamycin (one of the prototype inhibitors of glycosylation) suppresses P-gp activity thereby triggering the accumulation of cytostatic drugs in the cells, and thus providing evidence that inhibitors of glycosylation ablate the P-gp-mediated MDR phenotype.^{595,596} By contrast, some researchers assert that while N-glycosylation may stabilize correct folding of P-gp, guiding its proper subcellular localization and protecting it from luminal protease degradation, its precise role in P-gp function remains open-ended as tunicamycin treatment neither altered P-gp cellular localization to the plasma membrane nor the P-gp drug efflux activity.⁵⁹⁷⁻⁵⁹⁹

B2.10 Inhibitors of N-Linked Glycosylation

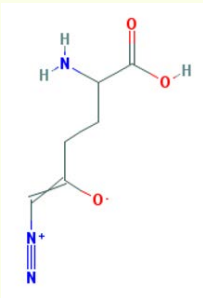
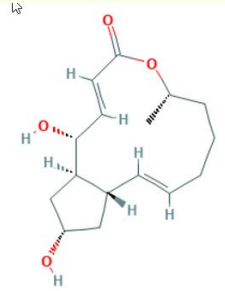
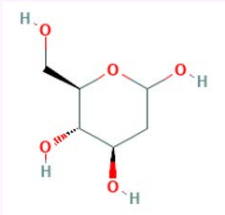
The myriad types of N-linked oligosaccharides are formed by two sequential reactions: 1) the formation of the lipid-linked saccharide precursor, $\text{Glc}_3\text{Man}_9(\text{GlcNAc})$ -2-pyrophosphoryl-dolichol, by the stepwise addition of GlcNAc, mannose and glucose to dolichyl-P, and 2) the removal of glucose and mannose by membrane-bound glycosidases and the addition of GlcNAc, galactose, sialic acid, and fucose by Golgi-localized glycosyltransferases to generate diverse complex oligosaccharide structures. Many glycoproteins contain more than one N-linked oligosaccharide structure—one oligosaccharide may be of the high-mannose type whereas another may be a complex chain.

Various methodologies are used to establish the role of specific structures in glycoprotein function, including inhibitors that hinder the different modification steps, resulting in the production of aberrant glycoproteins with altered carbohydrate structures. Several alkaloid-mimetic/specific inhibitors of the glucosidases and mannosidases involved in glycoprotein processing have been characterized. These inhibitors trigger the assembly of glycoproteins with glucose-containing high mannose structures, or various high-mannose or hybrid chains, depending on the site of inhibition. These inhibitors have also been useful for studying the glycan processing pathways and for comparing processing enzymes from different organisms.^{544,593,600-603} N-linked glycosylation inhibitors are categorized according to their mechanism of action and target.^{593,604-606} They may prevent N-glycosylation through:

1. Interference with the turnover of the process precursors;
2. Inhibition of glycosyltransferases and glycosylases;
3. Inhibition of transport of modified proteins between cellular compartments engaged in N-glycosylation, endoplasmic reticulum, Golgi apparatus; and
4. Functioning as substrate analogues.

Table 1.13 summarizes some of the known inhibitor classes of N-glycosylation.

Table 1.13: Specific classes and examples of N-glycosylation inhibitors

Inhibitor Class/Example(s)	Structure	Mode of Action	References
<p><i>Metabolic Inhibitors</i></p> <p>6-Diazo-5-Oxo-L-Norleucine (DON)</p>		<p>Affects turnover of glycosylation precursors, primarily at the stage of their formation. DON blocks glutamine:fructose-6-phosphate aminotransferase which catalyzes the synthesis of glucosamine from glutamine and fructose. Effects include disruption of mitochondrial internal membrane, permeabilization and dilation of the endoplasmic reticulum and induction of apoptosis. DON exhibits antitumour effects.</p>	593,606-608
<p>Brefeldin A (BFA)</p>		<p>A macrocyclic lactone synthesized from palmitate by various fungi. Inhibits the early transport of proteins from the endoplasmic reticulum to the Golgi apparatus. BFA also disrupts organization of the microtubule and actin cytoskeletons.</p>	609,610
<p><i>Sugar Analogues</i></p> <p>2-Deoxyglucose (2dGlc/2DG)</p>		<p>Inhibits glycosyltransferases so that saccharides are not transferred to the nascent glycoprotein and extended branching of the sugar core is obliterated. 2dGlc impacts gene expression, protein phosphorylation and signalling pathways and it blocks the cell cycle progression, DNA repair which culminates in apoptosis.</p>	593,611-613

Continued/...

Table 1.13: Specific classes and examples of N-glycosylation inhibitors (continued)

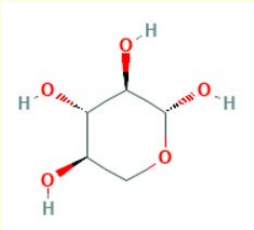

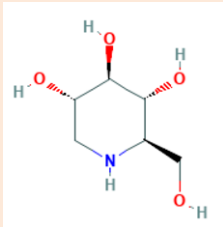
Inhibitor Class/Example(s)	Structure	Mode of Action	References
<p><i>Glycoside Primers</i></p> <p>β-D-Xyloside (β-D-Xyl)</p>		<p>β-D-Xyl blocks synthesis of glycosaminoglycans on growing proteoglycans. Disrupts glycoprotein assembly by elongating the oligosaccharide chains with exogenous primers instead of the endogenous glycoprotein core with specific sugar moieties which causes premature inhibition of glycan synthesis. β-D-Xyl also reversibly inhibits cell proliferation by arresting cells in the G₁ phase of the cell cycle.</p>	606,614-616
<p><i>Plant Alkaloids</i></p> <p>Castanospermine (CST)</p>		<p>CST specifically inhibits α-glycosidases I and II, thus impeding elongation of saccharide chains. Inhibition occurs particularly at the stage of glycosylation, after formation of a 14 monomer-long chain (Glc₃Man₉GlcNAc₂). Hence, plant alkaloids block formation of mature glycoproteins.</p>	593,601,603,606,617-619
<p>Deoxynojirimycin (DNJ)/ Deoxymannojirimycin (DMJ)</p>		<p>Castanospermine and deoxynojirimycin both obstruct angiogenesis <i>in vitro</i>. 1-Deoxymannojirimycin predominantly inhibits Golgi mannosidase I. 1-Deoxynojirimycin (1-DNJ) has been shown to possess antimetastatic potential.</p>	601,606,620-622

Table 1.13: Specific classes and examples of N-glycosylation inhibitors (continued)

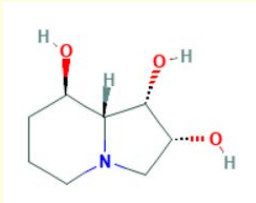
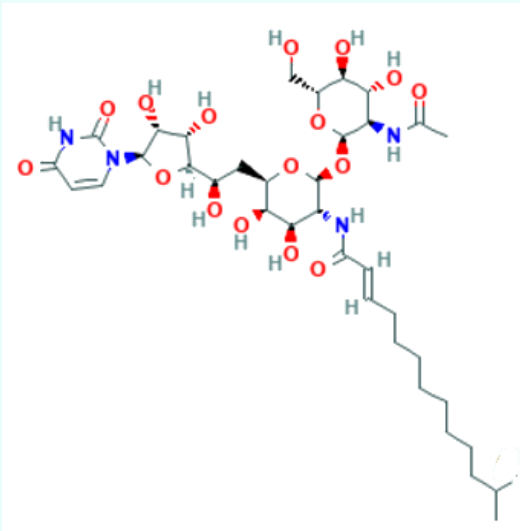
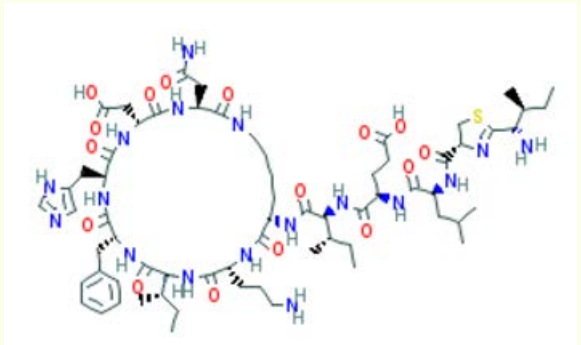
Inhibitor Class/Example(s)	Structure	Mechanism of Action	References
<p><i>Plant Alkaloids</i></p> <p>Swainsonine (SWSN)</p>		<p>Blocks Golgi α-mannosidase II and lysosomal α-mannosidase.</p> <p>It has a neurotoxic effect (“syndrome loco”) and intoxication with it occasions in accumulation of glycoproteins in lymph nodes. It also abrogates metastasis of melanoma cells by triggering natural killer cell proliferation or their anti-tumour activity. The drug is the first glycosidase inhibitor to have undergone anticancer clinical testing.</p>	601,603,606,617,623
<p><i>Antibiotics</i></p> <p>Tunicamycin (TM)</p>		<p>It is a nucleoside antibiotic which blocks the enzyme responsible for the transfer of 1-phospho-N-acetylglucosamine from UDP-N-acetylglucosamine to dolichol phosphate, i.e., the first glycosylation step. Tunicamycin induces E-cadherin-mediated cell–cell interactions and apoptosis in neoplastic cells. The antibiotics also exerts synergistic effects in combination with doxorubicin, cisplatin and vincristine.</p>	593,596,614,620,624,625

Table 1.13: Specific classes and examples of N-glycosylation inhibitors (continued)

Inhibitor Class/Example(s)	Structure	Mechanism of Action	References
<p><i>Antibiotics</i></p> <p>Bacitracin (BAC)</p>		<p>Prevents dolichol pyrophosphate (Dol-PP) hydrolysis to dolichol phosphate which, the after-effect being ablation of glycoprotein synthesis at its first stage. Bacitracin interferes with P-gp expression and localization. Bacitracin is a competitive inhibitor of protein disulfide isomerase and reduces phosphorylated focal adhesion kinase (p-FAK) and secreted matrix metalloproteinase-2 (MMP-2), which are the downstream of integrin and play a major role in cell migration and invasion, considered a therapeutic target for glioblastoma. Bacitracin has significant effects on both non-catalyzed protein folding and on other molecular chaperones. Bacitracin further disrupts ER function and causes higher ERS-mediated apoptosis in melanoma cells as demonstrated by the upregulation of ER chaperones.</p>	626-630
<p>2-D structures of the N-glycosylation inhibitors were downloaded from https://pubchem.ncbi.nlm.nih.gov/.</p>			

SECTION C: PROTEIN GLYCOSYLATION, ENDOPLASMIC RETICULUM STRESS AND THE UNFOLDED PROTEIN RESPONSE

C1. Introduction

In eukaryotic cells, many proteins are covalently modified during or immediately after translation. These modifications (e.g., phosphorylation, acetylation, glycosylation, methylation, sumoylation, sulfation, nitrosylation and ubiquitylation) collectively referred to as post-translational modifications (PTMs), regulate protein maturation, stability, dynamics, assembly, translocation, molecular interactions and cellular functions.⁶³¹⁻⁶³⁵ Modifications of a protein at asparagine Asn (N-glycosylation) or at serine/threonine residues (O-glycosylation) (Figure 1.14) are arguably the most prevalent PTMs which impact protein folding, maturation and activity.^{491,636} Protein glycosylation occurs in the eukaryotic secretory pathway and encompasses discrete biosynthetic transitions between the endoplasmic reticulum (ER) and Golgi apparatus.^{491,605} Unlike nucleic acids and proteins, glycan structures are not directly determined by genes or synthesized from a template, and may be linear or branched, and even undergo additional modification by acetylation, sulfation or phosphorylation.⁶³¹

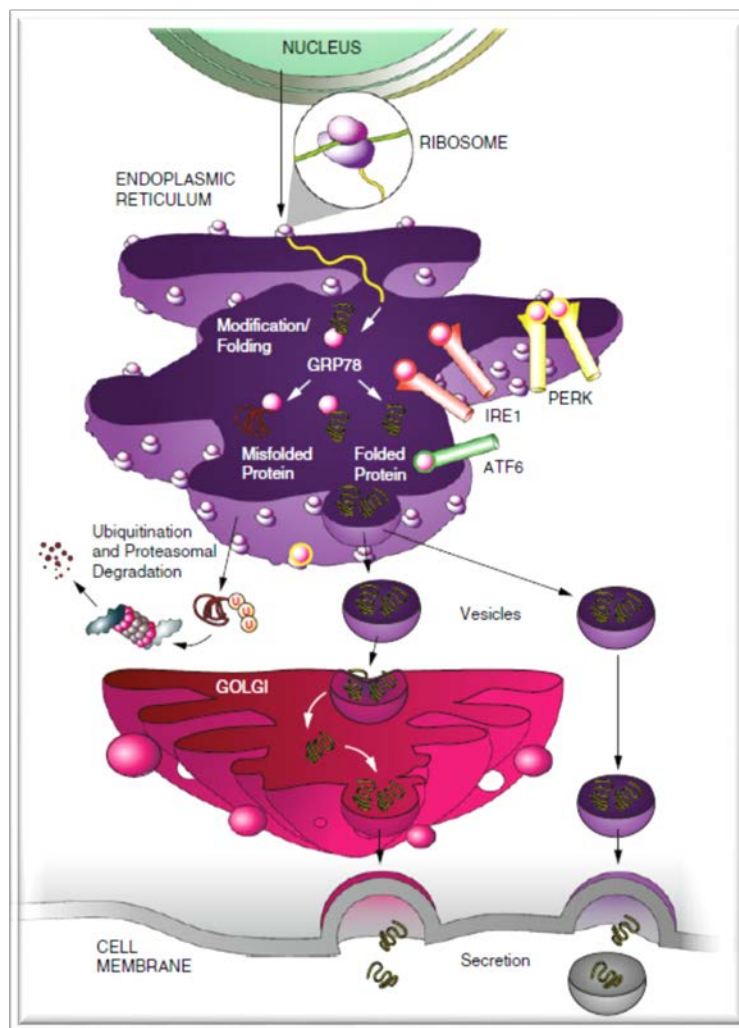
Glycosylation is an important component of the ER protein quality control system (ERQC) which precisely sorts and corrects misfolded proteins for reprocessing.⁶³⁷ The ERQC, starting in the ER and ending at the trans-Golgi, direct the dispatch, transit, secretion and fate (final localization) of properly folded and glycosylated proteins to the cell surface or external environment (terminal compartments). This progression is imperative in the development and homeostasis, as well as cell-to-cell communication in complex multicellular organisms.^{631,636,638,639} The glycan moieties of glycoproteins play crucial roles in intricate processes ranging from protein solubility, stability, conformation and function—and thus their half-life in the blood (circulation)—to their altered expression in most chronic or acquired infectious diseases, ERS and cancer.^{491,497,605,640-645}

C2. Endoplasmic Reticulum Stress and the Unfolded Protein Response

The ER is a major cellular compartment for protein synthesis, assembly and trafficking. Within the lumen and membranous network of the ER, a robust protein quality control system (ERQC) verifies whether secretory and membrane proteins are properly folded and modified before they are dispatched to their final destinations (cytosol, membrane and extracellular milieu).^{639,646-648} Incorrectly or misfolded proteins cannot assume their final conformation and functionally active structures, and are, therefore, retained in the lumen of ER until they are reconfigured into their proper conformations.⁶⁴⁹⁻⁶⁵¹

If the mature tertiary structure cannot be synthesized, misfolded proteins are then redirected to the cytoplasm to undergo ubiquitination and proteasome-mediated degradation (Figure 1.20), a process referred to as ER-associated degradation (ERAD).⁶⁵⁰⁻⁶⁵³ ERAD is indispensable in cells that cannot constitutively induce the unfolded protein response (UPR). Equally, loss of ERAD function leads to constitutive UPR induction. Ultimately, concurrent loss of ERAD and the UPR significantly decreases cell viability, suggesting that the UPR and ERAD are dynamic responses vital for the synchronized clearance of misfolded proteins, even in the absence of acute stress.^{654,655}

Under normal physiological conditions, the ERQC can cope with cellular demands, but extreme conditions of intracellular and extracellular stress (e.g., increased protein synthesis, genetic mutations that cause defects in folding, alteration in calcium homeostasis, and nutrient starvation such as glucose deprivation),^{649,650} neurodegenerative disorders, heart disease, smoking, diabetes and malignancy may overwhelm the ERQC capacity, leading to ERS.^{637,656-659} Thus, perturbations of ER homeostasis, in particular, protein homeostasis (proteostasis), results in the accumulation of unfolded proteins which then activates the ERAD and the UPR—integrated transcriptional and translational systems for transmitting information about the status of protein folding to the cytosol and nucleus (Figure 1.21).⁶⁶⁰



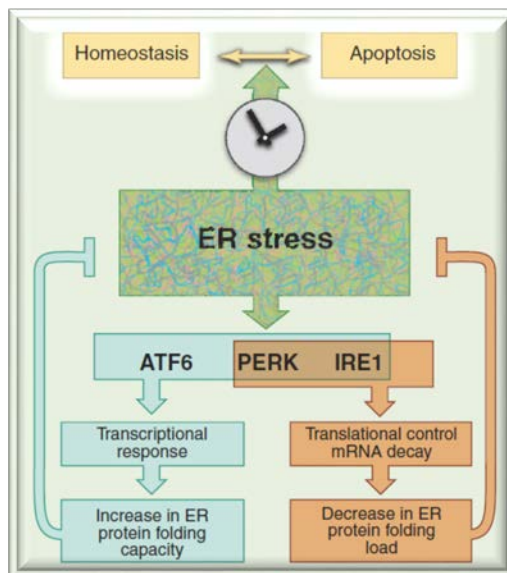
Normal state. Proteins that enter the ER are folded and transported to the Golgi apparatus or other destinations. GRP78 is bound to the luminal domains of PERK, IRE1, and domains of PERK, IRE1 and ATF6.

Source:⁶⁶¹ Park SW, Ozcan U. Potential for therapeutic manipulation of the UPR in disease. *Seminars in Immunopathology* 2013;35(3):351-373, permission granted by *Springer*, under the terms of the Creative Commons Attribution License which permits any use, distribution, and reproduction in any medium, provided the original author(s) and the source are credited; <http://link.springer.com/article/10.1007%2Fs00281-013-0370-z#copyrightInformation>; accessed 8 January 2017.

Figure 1.20: Endoplasmic reticulum protein folding function under normal physiological conditions

In eukaryotic cells, three ER transmembrane components (Figure 1.22)—inositol-requiring enzyme 1 (IRE1), protein kinase RNA-like ER kinase (PERK) and activating transcription factor 6 (ATF6), initiate distinct UPR signalling arms.⁶⁶²⁻⁶⁶⁴ The UPR triggers upregulation of genes encoding ER chaperones [e.g., heat shock protein 90 (HSP90), HSP70, CAAT/enhancer binding protein- α homologous protein (CHOP)/ growth arrest and DNA damage-inducible protein (GADD153), XBP1 (X-box binding protein 1),⁶⁶⁵ calreticulin (CRT),⁶⁶⁶ ER HSP40

ERdj3/DNAJB11,^{667,668} and ERdj5,⁶⁶⁹ and calnexin],⁶⁷⁰ attenuation of translation, and initiation of the ERQC to reinstate ER homeostasis.^{671,672}

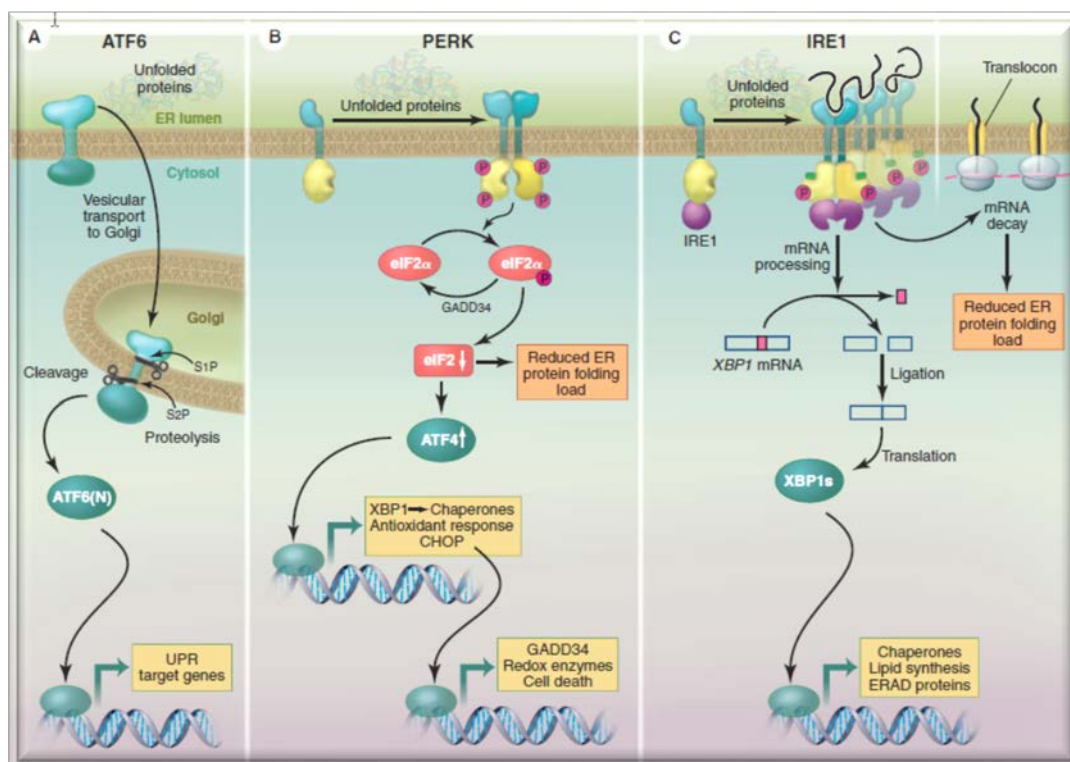


ERS activates the stress sensors ATF6, IRE1, and PERK, representing the three branches of the UPR. Activation of each sensor produces a transcription factor [ATF6(N), XBP1, and ATF4, respectively] that activates genes to increase the protein-folding capacity in the ER. IRE1 (via RIDD) and PERK (via eIF2a phosphorylation) also decrease the load of proteins entering the ER. Both outcomes work as feedback loops that mitigate ERS. If cells cannot reestablish homeostasis, but continue to experience prolonged and unmitigated ERS (depicted by the timer), they apoptose.

Source:⁶⁵⁰ Walter P, Ron D. The unfolded protein response: From stress pathway to homeostatic regulation. *Science* 2011;334(6059):1081-1086. Permission granted by *Science* (American Association for the Advancement of Science), <http://www.sciencemag.org/help/reprints-and-permissions>; accessed 8 January 2017.

Figure 1.21: Core elements of the UPR signalling network

The ER luminal binding protein—immunoglobulin heavy chain-binding protein (BiP)—also called glucose-regulated protein, 78kDa (GRP78), a member of the heat shock protein 70 (HSP70) family, is the most abundant ER-chaperone, and a central regulator of the ERQC machinery. BiP binds to and suppresses the activity of the mammalian ERS sensors, PERK, IRE1, and ATF6.^{673,674} Pro-survival (yang) GRP78 and pro-apoptotic (yin) CHOP are quintessential antagonistic mediators of the ERS response.⁶⁷⁵ When the UPR is insufficient to restore the steady state in the ER, programmed cell death (PCD) or apoptosis ensues, but chronic ERS can lead to pathological states.^{521,646,676-679} ERS is also a strong inducer of autophagy, a self-degradative process that has an adaptive function.⁶⁸⁰⁻⁶⁸²



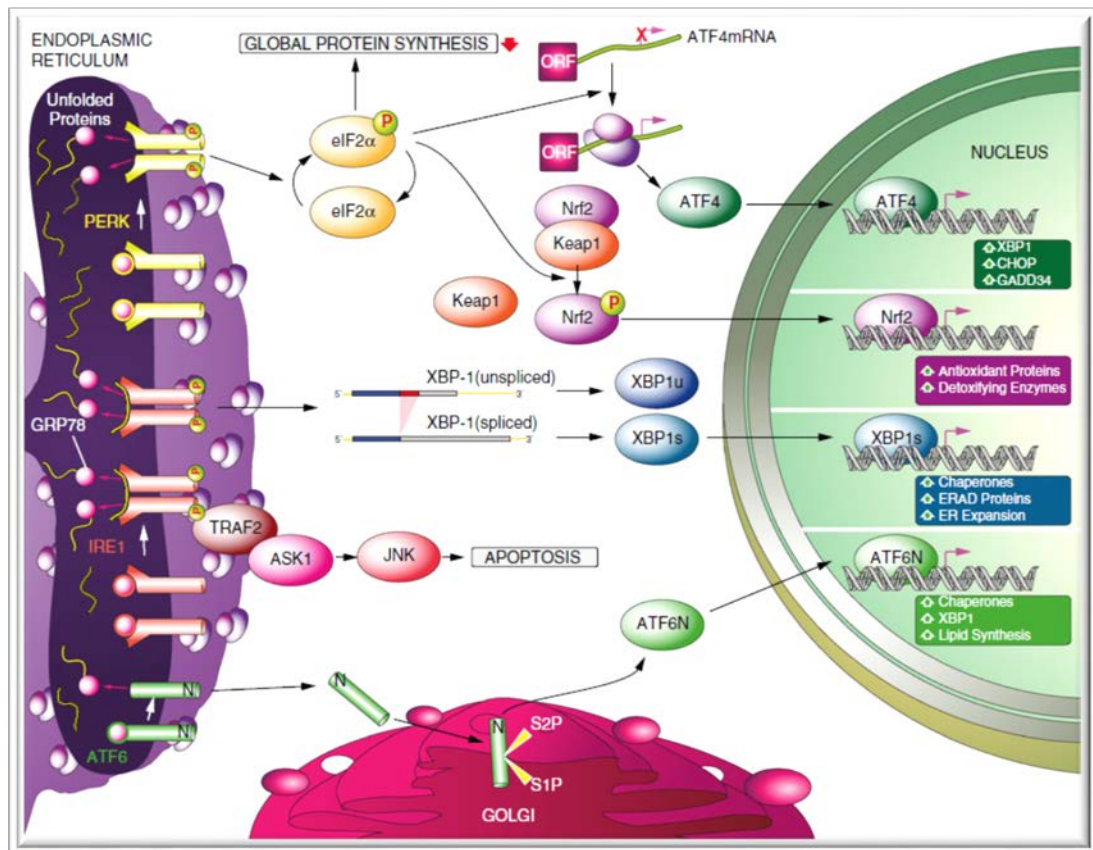
Three families of signal transducers (ATF6, PERK, and IRE1) sense the protein-folding conditions in the ER lumen and transmit that information, resulting in production of bZIP transcription regulators that enter the nucleus to drive transcription of UPR target genes. Each pathway uses a different mechanism of signal transduction: ATF6 by regulated proteolysis, PERK by translational control, and IRE1 by non-conventional mRNA splicing. In addition to the transcriptional responses that largely serve to increase the protein-folding capacity in the ER, both PERK and IRE1 reduce the ER folding load by down-tuning translation and degrading ER bound mRNAs, respectively.

Source:⁶⁵⁰ Walter P, Ron D. The unfolded protein response: From stress pathway to homeostatic regulation. *Science* 2011;334(6059):1081-1086. Permission granted by *Science (American Association for the Advancement of Science)*, <http://www.sciencemag.org/help/reprints-and-permissions>; accessed 8 January 2017.

Figure 1.22: The three branches of the UPR

C3. ER Stress and the UPR in Cancer

Most of functions mediated by the UPR in cellular homeostasis are also displayed in the role of ERS response in diseases that include cancer,⁶⁸³ diabetes, and metabolic, genetic, inflammatory, and neurodegenerative disorders.^{671,682,684} Several lines of evidence suggest that all branches of the UPR either promote or impede cancer initiation and progression, implicating various hallmarks of cancer,^{145,146,685,686} including sustaining proliferative signalling, evading growth suppressors, resisting cell death, enabling replicative immortality, inducing angiogenesis and activating invasion and metastasis (Figure 1.24).

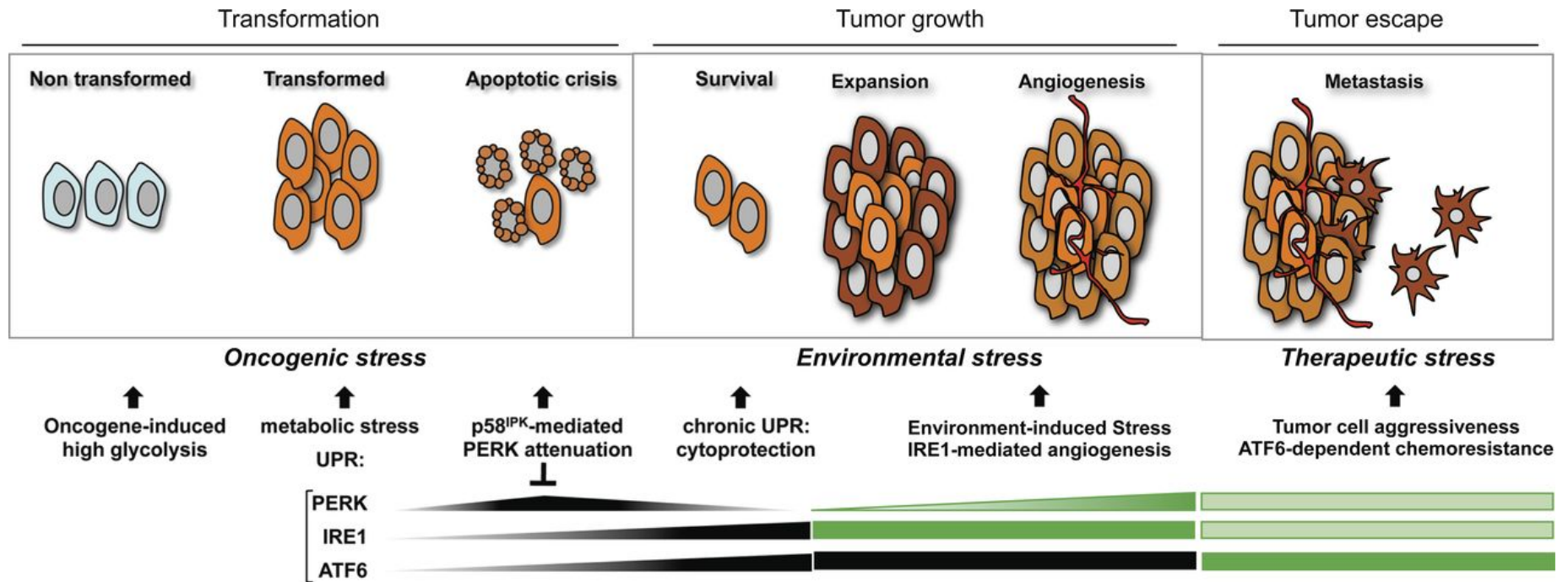


ERS state. GRP78 dissociates from PERK, IRE1, and ATF6. PERK and IRE1 oligomerize, forming a dimeric structure with a deep groove where peptide can bind. Upon oligomerization, PERK and IRE1 are auto-phosphorylated. PERK phosphorylates eIF2 α , leading to attenuation in global protein synthesis. Phosphorylated eIF2 α leads to translation and nuclear translocation of ATF4 and Nrf2. Activated IRE1 mediates unconventional mRNA splicing of XBP1 to generate XBP1s. IRE1 also recruits TRAF2 and ASK1 and leads to activation of JNK. ATF6 translocates to the Golgi apparatus and the cytoplasmic tail of ATF6 acts as a transcription factor to regulate UPR target genes.

Source:⁶⁶¹ Park SW, Ozcan U. Potential for therapeutic manipulation of the UPR in disease. *Seminars in Immunopathology* 2013;35(3):351-373, permission granted by *Springer*, under the terms of the Creative Commons Attribution License which permits any use, distribution, and reproduction in any medium, provided the original author(s) and the source are credited; <http://link.springer.com/article/10.1007%2Fs00281-013-0370-z#copyrightInformation>; accessed 8 January 2017.

Figure 1.23: Endoplasmic reticulum protein folding function under ERS conditions

Both intrinsic and extrinsic factors can activate the UPR in cancer cells, including hyperactivation of oncogenes and loss-of-function mutations in tumour suppressor genes, which may inappropriately amplify protein synthesis and translocation into the ER in response to excessive metabolic demands. Furthermore, mutations in oncogenes and tumour suppressor genes are known to inhibit ERS-induced apoptosis.^{675,683,687} Cancer cells exploit the ERS responses to promote survival and growth.



The activation status of the three UPR arms is shown at the bottom of the scheme. Green indicates a predominant role of the arm concerned in the tumourigenic process indicated at the top of the scheme (the gradient in green indicates the relative contribution of each arm).

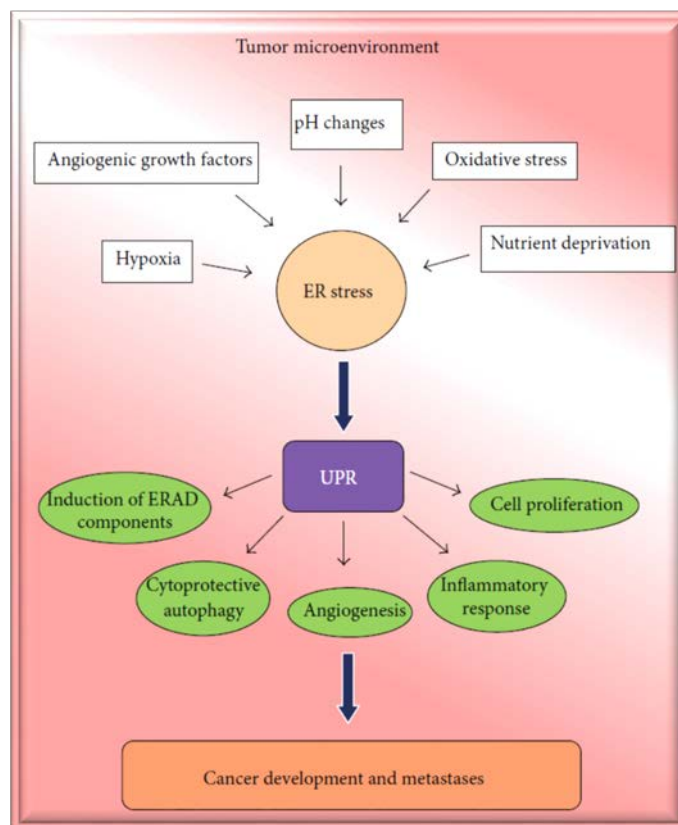
Source:⁶⁸⁴ Manie SN, Lebeau J, Chevet E. Cellular mechanisms of endoplasmic reticulum stress signaling in health and disease. 3. Orchestrating the unfolded protein response in oncogenesis: An update. *American Journal of Physiology Cell Physiology* 2014;307(10):C901-907, permission granted by *American Journal of Physiology Cell Physiology, The American Physiological Society*, <https://s100.copyright.com/AppDispatchServlet#formTop>; accessed 8 January 2017.

Figure 1.24: Involvement of UPR signaling during cell transformation and tumour growth

For example, the ER protein chaperone BiP is commonly overexpressed in breast cancer, lung cancer, prostate cancer, melanoma, and other malignancies to mediate the prosurvival response of cancer cells to major environmental stress.^{665,688,689} Surprisingly, unremitting ERS and UPR activation may interchange the cytoprotective functions of UPR into cell death programmes, a principle that can be exploited as a line of attack against cancer cells.^{661,675,690,691}

C4. Targeting ER Stress and the UPR

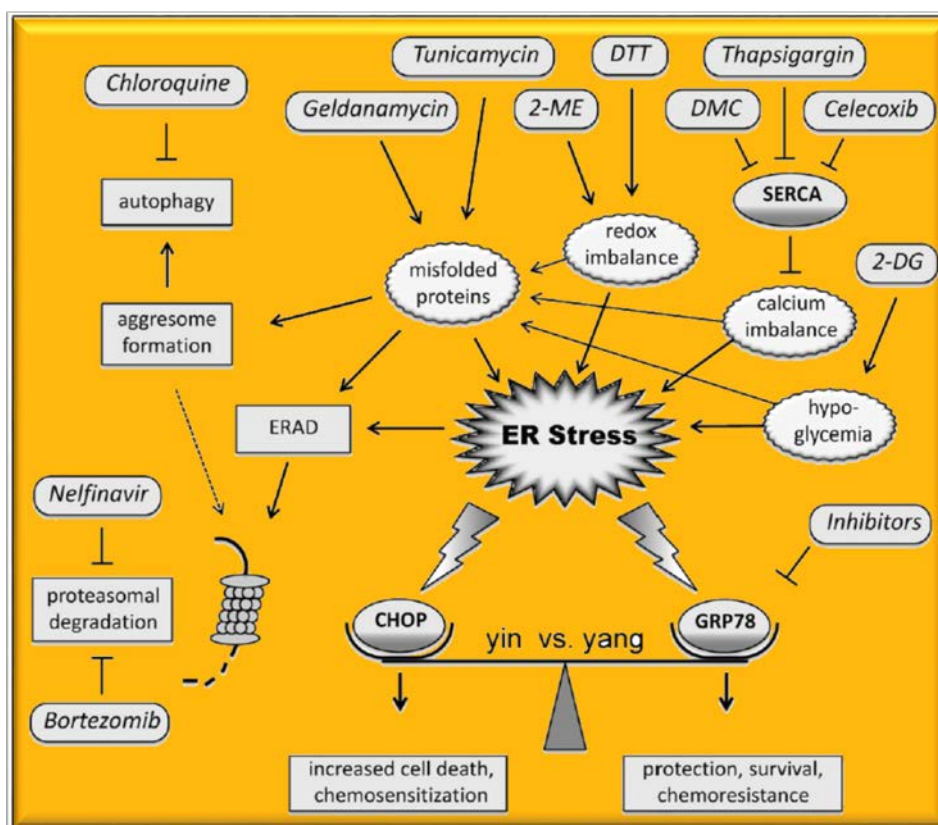
Cancer cells are resistant to extreme environmental stress conditions that induce ERS and UPR responses leading to cancer initiation and metastases (Figure 1.25).⁶⁸³



Source:⁶⁸³ Giampietri C, Petrunaro S, Conti S, Facchiano A, Filippini A, Ziparo E. Cancer microenvironment and endoplasmic reticulum stress response. *Mediators of Inflammation* 2015;2015:417281, permission granted (Open Access) by *Hindawi Publishing Corporation*; <https://www.hindawi.com/oa/>; accessed 9 January 2017.

Figure 1.25: Tumour microenvironment and activation of ERS and UPR responses in cancer

Therefore, the various molecular interconnections defining ERS and UPR in cancer, neurodegenerative and metabolic diseases offer a promising targeted therapeutic raison d'être. Several excellent reviews have recently described the merits of such pharmacological targeting of the UPR and the literature on this topic is expanding at an alarming rate.^{660,661,663,675,692-697} Selected examples of ERS aggravators (ERSAs) and UPR responses are indicated in Figures 1.26 and 1.27, and Table 1.14.^{675,686}



Pharmacological agents (shown in rounded rectangles) cause physiological imbalances (shown in undulated ovals) or directly block SERCA, autophagy, or the proteasome, thus causing the accumulation of misfolded proteins and resulting in aggravated ERS. ERS can be further exacerbated via inhibition of GRP78 with specific inhibitors (left, bottom; see text for details). Activation of the ERS response system/UPR involves GRP78 as key pro-survival and CHOP as key pro-apoptotic components, and these two proteins are representatives of the antagonistic cellular struggle for survival vs. cell death. Shifting this yin–yang balance towards dominance of CHOP will ensure cell death and abort the cell. However, if the yin module (in particular, GRP78) prevails, cell survival and, in the case of tumour cells, increased chemoresistance, will be favoured. CHOP, C/EBP homology protein; 2-DG, 2-deoxy-d-glucose; DMC, 2,5-dimethyl-celecoxib; 2-ME, 2-mercaptoethanol; DTT, dithiothreitol; SERCA, sarcoplasmic/endoplasmic reticulum calcium ATPase; ERAD, endoplasmic reticulum-associated degradation.

Source:⁶⁷⁵ Schönthal AH. Pharmacological targeting of endoplasmic reticulum stress signaling in cancer. *Biochemical Pharmacology* 2013;85(5):653-666, permission granted by *Biochemical Pharmacology, Elsevier* (Appendix 12).

Figure 1.26: Cellular impact of ERS aggravators that weigh on the yin vs yang balance

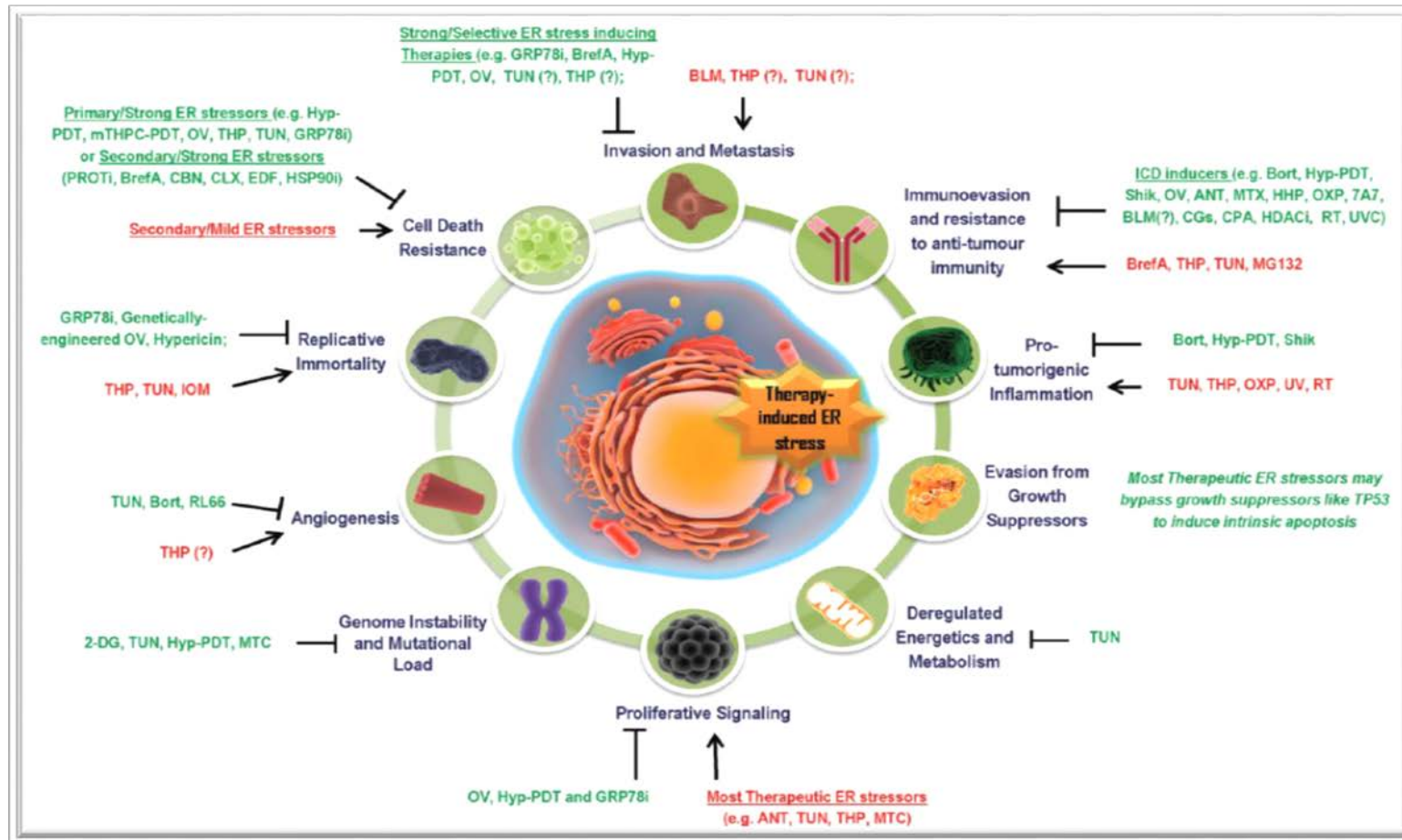


Figure 1.27: An overview of therapeutic ERS-based targeting of the main hallmarks of cancer

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Figure 1.27: An overview of therapeutic ERS-based targeting of the main hallmarks of cancer (continued)

Respective therapy-based ERS inducers have been segregated into 2 categories (wherever possible) based on their ability to target each of the hallmarks of cancer; such that therapies or drugs labeled with green inhibit the hallmark (thereby inhibiting or suppressing tumorigenesis) whereas those labeled with red support the hallmark (thereby enabling or supporting tumorigenesis). The question mark in parenthesis (?) indicates that data supporting the ability of the given therapy or drug to target or support a hallmark of cancer are not conclusive but are evidenced by either contradictory or incomplete observations. Please see the text for further details. 2-DG, 2-deoxyglucose; 7A7, murine anti-EGFR antibody; ANT, anthracycline; BLM, bleomycin; Bort, bortezomib; BrefA, brefeldin A; CBN, cannabinoids; CG, cardiac glycoside; CLX, celecoxib; CPA, cyclophosphamide; EDF, edelfosine; GRP78i, BiP/GRP78 inhibitor; HDACi, HDAC inhibitor; HHP, high hydrostatic pressure; HSP90i, HSP90 inhibitor; Hyp-PDT, hypericin-based photodynamic therapy; ICD, immunogenic cell death; IOM, ionomycin; MTC, microtubule-targeting chemotherapy; MTX, mitoxantrone; OV, oncolytic viruses; OXP, oxaliplatin; PROTi, proteasome inhibitor; RL66, an analog of curcumin; RT, radiotherapy; Shik, shikonin; THP, thapsigargin; TUN, tunicamycin; UVC, UV irradiation of C-band wavelength.

Source:⁶⁸⁶ Garg AD, Maes H, van Vliet AR, Agostinis P. Targeting the hallmarks of cancer with therapy-induced endoplasmic reticulum (ER) stress. *Molecular and Cellular Oncology* 2015;2(1):e975089; permission granted (Open Access, under the terms of the Creative Commons Attribution-Non-Commercial License (<http://creativecommons.org/licenses/by-nc/3.0/>)).



Table 1.14: Pharmacologic modulators commonly used in targeting ERS and UPR signalling

Biologic or Pharmacologic Modulator	Endoplasmic Reticulum Stress Mechanistic Principles and Therapeutic Targeting Strategy	References
Thapsigargin	<p>Thapsigargin, derived from <i>Thapsia garganica</i>, is a potent inducer of GRP78 expression and endoplasmic reticulum stress (ERS) and activator of the UPR through non-competitive inhibition of SERCA (sarcoplasmic/endoplasmic reticulum calcium ATPase). SERCA inhibition causes extensive efflux of calcium from ER stores into the cytosol and is thus a strong inducer of ERS—a potent trigger for autophagy, a self-degradative process that has an adaptive function. ERS mediated by thapsigargin promotes CHOP and death receptor 5 (DR5, also referred to as Apo2) synthesis, thus sensitizing TRAIL treatment, which induces oesophageal squamous cell carcinoma cell (ESCC) apoptosis. Thapsigargin synergistically enhances the anticancer activity of drugs against human ESCCs, including inhibition of tumour cell proliferation, invasion and metastasis, and induction of apoptosis. However, thapsigargin causes systemic toxicity, including potent tumour promoter, causes histamine release, and stimulates arachidonic acid metabolism. Combination of bortezomib with SERCA inhibitors, such as thapsigargin, celecoxib, or 2,5-dimethyl-celecoxib (DMC), aggravates ERS and greatly increases glioblastoma cell death <i>in vitro</i> and <i>in vivo</i>, pointing to the potential of thapsigargin as a combination agent in therapeutic regimens.</p>	675,680,686,698-702
Tunicamycin	<p>Tunicamycin is an asparagine-linked (N-linked) glycosylation inhibitor which causes impairment of protein folding and thus ERS. Tunicamycin blocks cell surface receptor tyrosine kinases (RTKs), thereby interrupting mitogenic and pro-survival signalling pathways and sensitizing tumour cells to cytotoxic therapies.</p>	596,703-709

Continued/...

Table 1.14: Pharmacologic modulators commonly used in targeting ERS and UPR signalling (continued)

Biologic or Pharmacologic Modulator	Endoplasmic Reticulum Stress Mechanistic Principles and Therapeutic Targeting Strategy	References
Tunicamycin	Tunicamycin inhibits angiogenesis <i>in vitro</i> and <i>in vivo</i> by arresting cells in the G ₁ phase of the cell cycle. It prevents the progression of a double- and a triple-negative breast tumour in athymic nude mice by inducing ERS, followed by apoptosis. Tunicamycin is efficacious alone or in combination with radiation/radiotherapy.	686
Brefeldin A	Brefeldin A (BFA) is an inhibitor of protein transport from ER to Golgi, and thus the secretory pathway. It is also an ADP-ribosylation factor (ARF) inhibitor. BFA is also a known perturbant of P-glycoprotein (P-gp), an ATP-dependent efflux pump encoded by the <i>MDR1</i> gene which mediates multidrug resistance of tumour cells to cancer therapy. BFA induces caspase activation and apoptosis and triggers GRP78 upregulation and ER dilation, markers of ERS.	609,675,685,686,710,711
GRP78/BiP Inhibitors	The ER chaperone, GRP78, is one of the most dynamic components of cancer cells. Overexpression of GRP78 correlates with apoptosis, angiogenesis, proliferation, tumourigenesis, invasion/metastasis, inflammation, immunity and drug resistance. GRP78 inhibitors are not generally available for clinical testing, but are developed under screening platform licences (https://nuevolution.com/wp-content/uploads/2016/08/EFMC-ISMC-2016_Poster_Final-PDF.pdf ; https://nuevolution.com/pipeline/1097/ ;). GRP78 levels can be reduced using a GRP78-specific small interfering RNA (siRNA). In NB cells, Akt increases the accumulation of GRP78 in response to 2-DG. Inhibition of GRP78 is of therapeutic utility for cancer and for bacterial and viral infections.	685,686,712-717

Continued/...

Table 1.14: Pharmacologic modulators commonly used in targeting ERS and UPR signalling (continued)

Biologic or Pharmacologic Modulator	Endoplasmic Reticulum Stress Mechanistic Principles and Therapeutic Targeting Strategy	References
<p>Celecoxib (CEL) 2,5-Dimethyl-Celecoxib (DMC)</p>	<p>Cyclooxygenase 2 inhibitor. Postulated to impede SERCA and perturb intracellular calcium homeostasis. Relentless Ca²⁺ dysregulation can induce ERS-mediated apoptosis. Celecoxib induces apoptosis independently from its COX-2 inhibitory action via a mitochondrial apoptosis pathway. It also prevents neuroblastoma tumour initiation and progression and potentiates the effect of chemotherapeutic drugs <i>in vitro</i> and <i>in vivo</i>. ERS inducer DMC augments TRAIL-induced apoptosis in glioblastoma and inhibits cell cycle progression and induces apoptosis in human leukaemia cells.</p>	<p>675,708,718-722</p>
<p>NSAIDs (Aspirin, Salicylates and Diclofenac)</p>	<p>Some non-steroidal anti-inflammatory drugs (NSAIDs) like aspirin (acetyl salicylic acid) and its metabolite, sodium salicylate, have profound effects on cellular functions and survival. Aspirin activates PERK and upregulates expression of the pro-apoptotic transcription factor CHOP (GADD153), a downstream event to eIF2α phosphorylation which, together with cleavage of caspase-12, are hallmarks of ERS-mediated responses. Salicylates inhibit prostaglandin H synthase (cyclooxygenase/COX) activity. By contrast, diclofenac (another NSAID) has been reported to exert protective effects against ER-stress-induced apoptosis in human neuroblastoma SH-SY5Y cells, by suppressing the activation of caspases in the intrinsic apoptotic pathway. NSAIDs perturb ER homeostasis by upregulating the expression of GRP78 and CHOP, and the activation of PERK and ATF6, but rarely the other UPR arm, viz. IRE-1. Inhibition of PGE₂ production with diclofenac, resulted in reduced tumour growth in an <i>in vivo</i> model of 11q-deleted neuroblastoma.</p>	<p>162,723-727</p>

Continued/...

Table 1.14: Pharmacologic modulators commonly used in targeting ERS and UPR signalling (continued)

Biologic or Pharmacologic Modulator	Endoplasmic Reticulum Stress Mechanistic Principles and Therapeutic Targeting Strategy	References
Geldanamycin	HSP90 and GRP94 inhibitor	233,665,727,728
Irestatin	Inhibits IRE-1 α activity	665,694,729
Bortezomib	Reversible inhibitor of the 26S proteasome. Bortezomib exerts synergistic cytotoxic effects in cancer cells by turning off the prosurvival ER chaperone BIP/Grp78 and turning on the pro-apoptotic NF-kappaB. Both nelfinavir and bortezomib lead to autophagy-dependent growth arrest and the radiosensitization of cancer cells.	688,730-732
Ritonavir and Nelfinavir	HIV protease inhibitors that activate certain UPR components such as CHOP and GRP78 and hence induce accumulation of misfolded proteins. Nelfinavir, in combination with various NSAIDs, causes reduction in cell survival and an increase in apoptosis. Also, HIV protease inhibitors significantly impede ABC transporters, including P-gp.	310,643,686,725,733-736
Resveratrol	Induces GRP78 and CHOP, p-eIF2 α and XBP1 splicing. Downregulate P-gp expression via inhibiting PI3K/Akt/mTOR pathway. Augments ER stress and the cytotoxic effects of glycolytic inhibition in neuroblastoma by downregulating Akt. Enhances mitochondrial biogenesis and riggers UPR.	657,713,737,738
Epidermal Growth Factor (EGF)-SubA	Targets GRP78, impeding its function and affecting proteostasis in ER.	686
Oncolytic Viruses	Stress the ER through viral protein overload.	686
Protein Disulphide Isomerase (PDI) Inhibitors	Protein disulfide isomerase (PDI) is an essential enzyme of disulphide bond formation in the ER. PDI inhibitors cause rapid accumulation of misfolded or unfolded proteins in the ER. There is an interrelation of ER stress and ROS with redox signalling mediators such as PDI-ER oxidoreductin (ERO)-1, glutathione (GSH)/glutathione disulphide (GSSG), NADPH oxidase 4 (Nox4), NADPH-P450 reductase (NPR) and calcium.	686,739-741

Continued/...

Table 1.14: Pharmacologic modulators commonly used in targeting ERS and UPR signalling (continued)

Biologic or Pharmacologic Modulator	Endoplasmic Reticulum Stress Mechanistic Principles and Therapeutic Targeting Strategy	References
Versipelostatin/Epigallocatechin Gallate	Derived from green tea extract and causes inhibition of GRP78.	675,685,686,742
Anthracyclines/Mitoxantrone/Carboplatin	Cause ROS production leading to ROS-based ERS.	686,694,743
Chloroquine	Lysosomotropic agent and inhibitor of autophagy. Combination of nelfinavir and chloroquine significantly increased ER stress and caused selective cell death in multiple cell line models with hyperactive mTORC1.	744-746
BRAF Inhibitor	Interferes with cytosolic Ca ²⁺ homeostasis causing ERS. Predominantly causes activation of the PERK–eIF2 α –ATF4/ATF3 pathway, which in turn promotes cytoprotective autophagy. Combined BRAF and autophagy inhibition promotes tumour regression in BRAFi-resistant xenografts.	312,686,745,747-749
Cannabinoids	Cause ERS through ceramide accumulation and eIF2 α phosphorylation. Plant-derived cannabinoids are moderately effective in reversing MDR in CEM/VLB100 cells by decreasing P-gp expression.	686,750
Curcumin (Turmeric)	Plant polyphenols that have been identified to possess proteasome-inhibitory activity include (-)-epigallocatechins-3-gallate (EGCG), genistein, luteolin, apigenin, chrysin, quercetin, curcumin and tannic acid. SERCA inhibition causes ER Ca ²⁺ imbalance and ERS. Down regulates calreticulin. Liposome-encapsulated curcumin suppresses neuroblastoma growth through nuclear factor-kappa B inhibition. Curcumin down-regulates transcription factors important for cell growth and survival, through modulation of the NF-kB and PI3K/AKT pathways.	661,686,742,751
HDAC inhibitors (HDACi) e.g. Vorinostat	Cause GRP78 acetylation, inhibiting GRP78 function and compromises ER protein folding, causing ERS.	
Ceapins	Selectively targets the ATF6 α branch.	752
Chemotherapeutic Agents	Induce ERS through various mechanisms.	676,686,687,692,699,730-732

C5. The ERS and UPR in Perspective

Pro-survival (yang) GRP78 and pro-apoptotic (yin) CHOP are quintessential opposing regulators of the ERS response. Whereas suprabasal levels of GRP78 are commonly found in many tumour cell lines and primary tumour tissues (and levitate upon ERS), CHOP is predominantly infrabasal, but intensifies in response to short-term, acute ERS. Factoring in the differential in baseline ERS levels in tumour vs. normal cells, manipulated pharmacological aggravation of pre-existing ERS in tumour cells can be exploited to ‘‘overload’’ this already burdened system to eclipse the tumour cells’ capacity for adaptation. During this process, the ERS system’s pro-apoptotic module would surpass the pro-survival module, leading to increased chemosensitivity of the tumour. By analogy, normal cells should be reasonably safeguarded from the toxic outcomes of pharmacologically increased ERS. Their ERS response system, which had not been subjected to chronic activation, would be triggered from significantly lower baseline levels and therefore would have the competence to stem increased ERS levels. Generally, the protective module of normal cells would presumably dominate and shield the cell from stress-induced toxicity significantly longer than is the case in tumour cells. This is the sine qua non of probing ERS and UPR dynamics and hence targeting the hallmarks of cancer with therapy-induced ERS.

SECTION D: RESEARCH CONTEXT

D1. Problem Statement and Research Questions

Despite recent advances in cancer cell glycomics and the arsenal of investigational and approved drugs against NB therapeutic targets, successful treatment of high-risk neuroblastoma (HR-NB) remains a challenging task since 40 % of patients still relapse during or after glycan-based immunotherapy following standard therapy.^{153,494} Therefore, a dire need exists to develop novel treatment modalities that target the NB glycome, proteome and transcriptome. Neuroblastoma (NB) is the most common paediatric cancer and accounts for

15% of all oncology deaths in infants. NB continues to perplex scientists and oncologists alike because its biological and clinical behaviour fluctuate between complete spontaneous regression and clinical multidrug resistance, strongly indicating that besides genetic events, the tumour microenvironment (TME) significantly influences these characteristics of NB.^{64,147} Survival rates of HR-NB remain less than 50%, with amplification of the *MYCN* oncogene being the most significant hallmark associated with aggressive NB and poor survival outcome.^{184,185}

Transcriptionally, the ABC transporters are directly and coordinately regulated by *MYCN*³²⁸ and, correspondingly, their overexpression correlates with poor prognosis.^{327,329} Most aggressive NBs exhibit MDR,³¹¹ attributable to p53 mutations and/or a loss of p53 function induced during chemotherapy,⁷⁵³ which further exacerbates the probability of relapse.^{333,334} While the prognostic merit of the ABC transporters in childhood NB is generally ascribed to their role in cytotoxic drug efflux, several reports claim that they might promote the malignant phenotype independent of this function, thus unlocking their potential as therapeutic targets,³³⁰ and strengthening the less well understood, but evolving theme of the drug efflux-independent contributions of ABC transporters to cancer biology and treatment failure.³¹¹ Similarly, some metastatic MDR NBs derive from the clonal selection of side population cells that constitutively express the *MDR1* (*P-gp*, *ABCB1*), *MRP1/ABCC1* and *MRP4/ABCC4* gene family, which may or may not correlate with *MYCN* amplification and poor outcome.^{7,294,323,327,328,330,335} Moreover, minimal residual disease (MRD), the major cause of tumour recurrence (relapse) and metastasis, is enriched in cancer stem cells (CSCs) with an increased drug efflux capacity mediated through overexpression of ABC transporters.^{336,337}

Charging into the fray are elevated levels of different types of gangliosides that profoundly contribute to aggressive NB behaviour and poor patient survival.^{527,537,539,540} By contrast, overexpression of both complex “a” gangliosides (CaG and CbG), eradicates aggressive tumour-cell behaviour *in vitro* (e.g., cellular proliferation and migration) and promotes

differentiation.⁷⁵⁴ In recent years, interest in gangliosides has been revived, mainly as prognostic biomarkers to stratify NB patients for targeted anticancer immunotherapy and to monitor efficacy of treatment.⁵⁴¹ Diverse molecules involved in NB glycobiology play key roles in tumour growth and are therefore potential targets for anti-tumour therapy.⁵²⁷

As mentioned above, the overexpression of MDR transporters strongly correlates with poor NB therapeutic outcome since they efflux a wide array of endogenous compounds and anticancer drugs from cancer cells.^{311,329,330} Theoretical and contextual issues emerging from the literature point to a multitude of unclarified roles of endogenous compounds and anticancer drugs at the intersection of ABC transporters and NB behaviour and cancer cell responses to chemotherapy. Perturbation of ABC transporters may provide insightful options for therapeutic repression of HR-NB, and proof of concept for increasing drug bioavailability (therapeutic efficacy) in refractory tumours which overexpress these glycan.³²⁶

The MDR transporter, P-glycoprotein (ABCB1, P-gp), has been shown to be one of the most strongly upregulated genes associated with acquired drug resistance and NB treatment failure.^{527,589,590} Inhibition of protein glycosylation reverses the MDR phenotype of several cancer cell lines.⁵⁹² Equally, inhibition of N-linked glycosylation hampers ALK phosphorylation and pro-survival signalling in NB cell lines.⁵⁵⁵ Inhibitors of N-glycosylation, e.g., tunicamycin, hinder P-gp-mediated MDR phenotype.^{595,596} The precise role of N-glycosylation in P-gp function remains to be fully unravelled.⁵⁹⁷⁻⁵⁹⁹ Several classes of N-linked glycosylation inhibitors are available that need to be evaluated further for their potential to alter NB behaviour.⁵⁹³

The glycan moieties of glycoproteins are critical for various cellular processes such as protein solubility, stability, conformation and function. Thus, altered expression of glycans has been implicated in chronic or acquired infectious diseases, endoplasmic reticulum stress (ERS) and cancer.^{491,497,605,640-645} Perturbation of N-linked glycosylation can also result in the

accumulation of unfolded/misfolded proteins which, in turn, may trigger ERS, the unfolded protein response (UPR) and, ultimately, decreased cell viability and apoptosis.^{650,661} Severe ERS may as well induce autophagy, a self-degradative process that has a life-saving adaptive function.⁶⁸⁰⁻⁶⁸² The TME is the arena for ERS and UPR responses that sustain various hallmarks of cancer.^{683,684,686}

It is clear from the above considerations that NB glycopathobiology, particularly the diverse cancer landscape exemplified by the N-glycoproteome in eukaryotic cells (N-linked protein glycosylation in the ER coupled with MDR, ERS, UPR activation, apoptosis and autophagy), offers an emerging theme in the therapeutic targeting of cancers, including NB.^{326,493,494,519,527,544,546,569,588,593,599,645,755-763} To this end, we have set out to explore the effects of various glycosylation inhibitors and ERS inducers on SK-N-BE(2) NB cell survival and ability to efflux calcein-AM, a P-gp substrate.

D2. Purpose of the Study

The purpose of the study was to investigate the effects of various glycosylation inhibitors and ERS inducers on the behaviour of NB cells in culture. For this study, we have selected the continuous SK-N-BE(2) cell line as representative of human NB cells *in vitro* that overexpress readily detectable levels of P-glycoprotein (P-gp, ABCB1) and other ABC transporters.^{764,765} The SK-N-BE(2) cell line was derived from a bone marrow metastases in a patient refractory to chemotherapy.⁷⁶⁶⁻⁷⁶⁹ Figure 2.1 in Chapter 2 shows the experimental design of the project presented in this thesis.

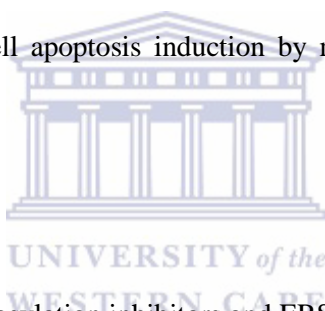
D3. Aims of the Study

The aims of the study were to determine the effects of N-glycosylation inhibition and ERS induction on SK-N-BE(2) cell proliferation and viability, apoptosis and P-glycoprotein drug efflux function.

D4. Objectives of the Study

In this study, the following N-glycosylation inhibitors and ERS inducers— aspirin, bacitracin, castanospermine, brefeldin A and thapsigargin— have been evaluated for their effects on:

1. SK-N-BE(2) NB cell growth and viability, using the CCK-8 assay.
2. P-glycoprotein (P-gp, ABCB1)-mediated cellular drug efflux function in SK-N-BE(2) NB cells, using Cayman's Calcein-AM multidrug resistance assay.
3. SK-N-BE(2) NB cell viability, cytotoxicity and apoptosis induction by caspase-3 activation, using the Apotox-Glo Triplex assay.
4. SK-N-BE(2) NB cell apoptosis induction by morphological staining of cells with Annexin-FITC.



D5. Hypothesis

We hypothesize that N-glycosylation inhibitors and ERS inducers will alter the manifestation of SK-N-BE(2) cancer cell hallmarks evaluated, namely, cell survival (proliferation, viability and apoptosis) and P-glycoprotein-mediated drug efflux function.

SECTION E: SUMMARY

This chapter provided the introduction and literature review on NB encompassing the epidemiology of the disease, risk factors and staging, prognostic markers, histopathological characteristics, detection, diagnosis and prognosis, clinical presentation, signs and symptoms, molecular pathogenesis, genetics and genomics and therapeutic landscape. In addition, the chapter underscored the significance of glycans and protein glycosylation in NB and the targeting strategies for ER stress and the UPR. Finally, the chapter outlined the research context of the study in terms of problem statement, aims and objectives and hypothesis.

CHAPTER 2

RESEARCH METHODOLOGY

2.1 Experimental Design

The focus of this chapter is to outline and describe the research methodology and experimental design that have been chosen for the study. It summarizes the materials and methods used such as chemicals required, drugs tested and the maintenance of the parental SK-N-BE(2) (American Type Culture Collection / ATCC® CRL2271™) neuroblastoma cell line. Analyses of the SK-N-BE(2) neuroblastoma cells exposed to brefeldin A (BFA), thapsigargin (TG), aspirin (AS), castanospermin (CST) and bacitracin a (BAC) included growth curves, cell viability and cytotoxicity assays by means of the Cell Counting Kit-8 (CCK-8), measurement of P-glycoprotein (P-gp, ABCB1) cellular drug efflux pump function using the Calcein-AM (Cayman's multidrug resistance) assay kit, the Apotox-Glo™ triplex cell viability, cytotoxicity and apoptosis assays, caspase-3 activation and morphological staining of apoptotic cells using the Annexin V-FITC kit (Figure 2.1). Further details of the experimental design are covered in the subsections that follow. The statistical methods used for data analysis are also described.

2.2 Drugs and Chemicals

Drugs and chemicals used in this study included thapsigargin from plant *Thapsia garganica* (CAS 67526-95-8, Sigma-Aldrich, St Louis, MO, USA), brefeldin A from *Penicillium brefeldianum* (CAS 20350-15-6, Sigma-Aldrich, St Louis, MO, USA), castanospermin from *Castanospermum australe* seeds (CAS 79831-76-8, Sigma-Aldrich, St Louis, MO, USA), bacitracin (CAS 1405-87-4, Sigma-Aldrich, St Louis, MO, USA), aspirin (CAS 50.78.2, Sigma-Aldrich, St Louis, MO, USA), heat inactivated foetal bovine serum (Biochrome, The Scientific Group), phosphate buffered saline (PBS) (Gibco, Life Technologies), Dulbecco's

Modified Eagles Medium supplemented with F-12 glutamax (DMEM F-12 glutamax) (Gibco Life Technologies), penicillin/streptomycin (Invitrogen or Gibco, Life Technologies), trypsin-EDTA (Gibco, Life Technologies), dimethyl sulfoxide (DMSO) (Sigma-Aldrich, St Louis, MO, USA), trypan blue (CAS 72-57-1, Sigma-Aldrich, St Louis, MO, USA), Cell Countin Kit-8 (CCK-8)(Item no 06041406, Enzo Life Sciences) Cayman Chemicals Multi Drug Resistance kit (Calcein-AM) (Item no. 600370), Annexin V-CY3 (Cat: APOAC; Sigma-Aldrich, St Louis, MO, USA), ApoTox-Glo[®] triplex assay kit (Cat: G6320, Promega).

2.3 Culture and Maintenance of SK-N-BE(2) Neuroblastoma Cells

The continuous human neuroblastoma (NB) SK-N-BE(2) cell line, originally purchased from the American Type Culture Collection (ATCC, Rockville, MA), was kindly provided by Dr AM Serafin, Radiobiology Laboratory, Department of Medical Imaging and Clinical Oncology, Faculty of Medicine and Health Sciences, University of Stellenbosch, South Africa. The SK-N-BE(2) cell line is known to overexpress readily detectable levels of P-glycoprotein (P-gp, ABCB1) and other ABC transporters.^{764,765} The SK-N-BE(2) cell line was established from a bone marrow biopsy of a metastases in November 1972 of a patient refractory to chemotherapy.⁷⁶⁶⁻⁷⁶⁹

All tissue culture operations were carried out in a model NU-5510E NuAire DHD autoflow automatic CO₂ air-jacketed incubator and an AireGard NU-201-430E horizontal laminar airflow cabinet with a HEPA-filtered clean work area (NuAire). SK-N-BE(2) cells were maintained in Dulbecco's Modified Eagle Medium (DMEM) and supplemented with 10% heat-inactivated foetal bovine serum (HIFBS), 1% penicillin/streptomycin (100 µg/ml penicillin and 10 µg/ml streptomycin) and grown as monolayer cultures at 37°C in relative humidity (RH) of 80%) in an atmosphere of 5% CO₂:95% air. Routinely, cryovials containing frozen SK-N-BE(2) cells in 40% HIFB, 50% DMEM, 10% DMSO were removed from -80°C freezer and thawed in a 37°C water bath.

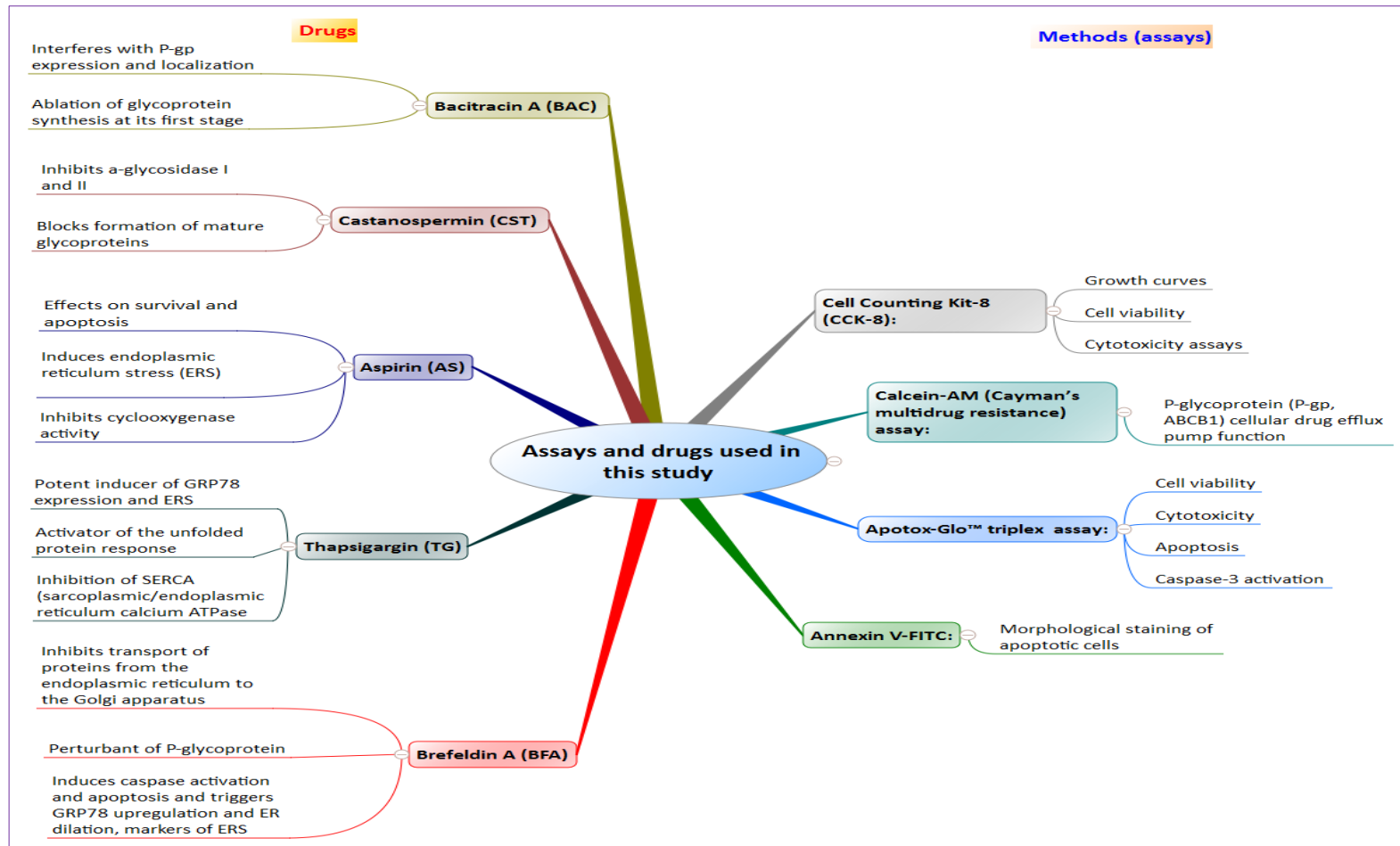
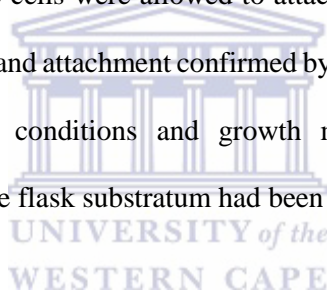


Figure 2.1: Experimental design: Assays and drugs used in this study

The caps were wiped with 70% ethanol and the contents of the vial transferred aseptically to a 15-ml conical centrifuge tube containing 1 ml of Modified Eagles Medium (MEM)/F-12 supplemented with 1% penicillin-streptomycin and 10% HIFBS, centrifuged for 5 minutes at 2500 rpm. After centrifugation, the supernatant was discarded and the cell pellet resuspended in 2 ml of complete medium. The cells were mixed thoroughly to ensure a homogeneous cell suspension, 1 ml of which was transferred to T-25 culture flask (surface area 2500 mm²) containing 5 ml complete medium to maintain stock cultures.

The flask was placed on a PrimoVert phase-contrast microscope to visualize the presence of suspended cells, and then placed in a 37°C incubator at 5% CO₂ and 80% RH, the incubation specifications were kept constant throughout for cells to acclimatize and attach to the substratum of the flask. The cells were allowed to attach for 24 h, after which the flask was removed from the incubator and attachment confirmed by microscopy. The flask was incubated under ideal tissue culture conditions and growth medium periodically changed until approximately 80-90% of the flask substratum had been occupied by SK-N-BE(2) cells.



Once confluency had been reached, cells were gently trypsinized. The medium was aspirated and the cells rinsed with 2 ml PBS. After 1 minute, the PBS was aspirated and replaced with 2 ml of 0.25% Trypsin-EDTA and placed in the incubator for 5-15 minutes in order for detachment of the cell monolayer to be achieved. The flask was then removed from the incubator and placed in a laminar flow cabinet. Thereafter, 4 ml complete medium was added to the flask to deactivate the trypsin. The cells were gently mixed using an electronic pipette aid and detached cells aspirated and transferred to a 15 -ml conical centrifuge tube, centrifuged 5 minutes at 2500 rpm to separate the cells from the medium-trypsin solution. After centrifugation, the supernatant was discarded and the cell pellet resuspended in 5 ml of complete medium. The cells were mixed to ensure a homogeneous cell suspension, 1 ml of which was transferred to T-25 culture flask (surface area 2500 mm²) containing 5 ml complete medium to maintain stock cultures or for use in experiments.

2.4 Growth Curve Analysis of SK-N-BE(2) Neuroblastoma Cells

SK-N-BE(2) neuroblastoma cells were trypsinized and transferred to a 15-ml centrifuge tube and spun at 2500 rpm for 5 min. The supernatant was removed and the pellet resuspended in 5 ml of complete fresh medium. Cells were seeded into 24-well plates at density of 1×10^5 cells/ml per well (100 μ l of suspension was mixed with 1.9 ml of fresh media to obtain a final volume of 2 ml per well. The plates were incubated for 24 hours (24h), 48h, 72h and 96h, respectively. After the incubation period, cells were harvested with 1 ml trypsin-EDTA from wells every 24h for the duration of the experiment. Viable cells were counted using the Bio-Rad TC-20 cell counter at a ratio 1:1 cell suspension: 0.4 μ M trypan blue. The experiments were conducted in quadruplicate and the results pooled.

2.5 Cell Counting Kit-8 (CCK-8) Cell Viability Assays

The Cell Counting Kit-8 (CCK-8, CCK-8; Dojindo Laboratories, Japan) permits precise assays by utilizing Dojindo's highly water-soluble tetrazolium salt. WST-8 [2-(2-methoxy-4-nitrophenyl)-3-(4-nitrophenyl)-5-(2,4-disulfophenyl)-2H-tetrazolium monosodium salt] produces a water-soluble formazan dye upon reduction in the presence of an electron carrier.⁷⁷⁰⁻
⁷⁷² CCK-8, being non-radioactive, allows sensitive colorimetric assays for the determination of the number of viable cells in cell proliferation and cytotoxicity assays. WST-8 is reduced by dehydrogenases in cells to give a yellow-coloured product (formazan), which is soluble in tissue culture medium. The amount of the formazan dye generated by the activity of dehydrogenases in cells is directly proportional to the number of living cells. Cell viability was measured using the CCK-8 kit, according to manufacturer's protocol.

All CCK-8 assays were carried out in 96-well flat bottom tissue culture plates. SK-N-BE(2) NB cells were seeded at density of 5×10^4 cells/ml. A 100 μ l of cell suspension was added to each well and cells were allowed to attach for 24h under normal incubation conditions. After 24h, the media was aspirated from all wells, first and second columns were replaced with 100 μ l of complete media alone while the other columns were replaced with increasing \log_{10}

concentrations of test compounds: bacitracin (0.001, 0.01, 0.1, 1, 10, 100 mM), castanospermine (0.00001, 0.0001, 0.001, 0.01, 0.1, 1 mM), aspirin (0.0001, 0.001, 0.01, 0.1, 1, 10 mM), thapsigargin (0.01, 0.1, 1, 10, 100, 1000 nM) and brefeldin A (0.002, 0.02, 0.2, 2, 20, 200 µM) in quadruplicate wells. Following incubation with the compounds for various time periods, 100 µl of CCK-8 solution was added to each well of the plate, and then plates were placed in a 37°C incubator at 5% CO₂ and RH 80%. The optical density (OD, absorbance) was obtained at 450 nm using a Promega GloMax™ Multiscan plate reader. The mean blank-corrected absorbance (MBCA) was derived from the following equation:

$$MBCA = \frac{1}{4} \sum_{i=1}^4 (A_i - A_0)$$

where A_i represents the absorbance reading of well i and A_0 is the absorbance reading of the blank well (inoculated cells without test compound=untreated controls with variable molar concentrations of vehicle approximating final concentrations present in the test wells).

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2.6 Apotox-Glo™ Triplex Cell Cytotoxicity, Viability and Apoptosis Assays

2.6.1 Principle of the Apotox-Glo™ Triplex Assay

A number of 96-well assays are available for high throughput screening of cytotoxicity of drugs (<https://www.promega.com/-/media/files/promega-worldwide/north-america/promega-us/webinars-and-events/assessmentcellhealthwebinfo0412.pdf>).⁷⁷³⁻⁷⁷⁷ The Promega ApoTox-Glo™ Triplex Assay combines three assay chemistries to assess viability, cytotoxicity and caspase activation events within a single assay well. The first part of the assay simultaneously measures two protease activities; one is a marker of cell viability and the other is a marker of cytotoxicity. The live-cell protease activity is restricted to intact viable cells and is measured using a fluorogenic, cell-permeant, peptide substrate (glycylphenylalanyl aminofluorocoumarin, GF-AFC). The substrate enters intact cells where it is cleaved by the live-cell protease activity to generate a fluorescent signal proportional to the number of living cells.

This live-cell protease becomes inactive upon loss of cell membrane integrity and leakage into the surrounding culture medium. A second, fluorogenic cell-impermeant peptide substrate (bis-alanylalanyl-phenylalanyl-rhodamine 110; bis-AAF-R110) is used to measure dead-cell protease activity, which is released from cells that have lost membrane integrity. Because bis-AAF-R110 is not cell-permeant, essentially no signal from this substrate is generated by intact, viable cells. The live- and dead-cell proteases produce different products, AFC and R110, which have different excitation and emission spectra, allowing them to be detected simultaneously.

The second part of the assay uses a luminogenic caspase-3/7 substrate, which contains the tetrapeptide sequence DEVD (aspartic acid, glutamic acid, valine, aspartic acid), in a reagent optimized for caspase activity, luciferase activity and cell lysis. Adding the Caspase-Glo® 3/7 reagent in an “add-mix-measure” format results in cell lysis, followed by caspase cleavage of the substrate and generation of a “glow-type” luminescent signal produced by luciferase. Luminescence is proportional to the amount of caspase activity present. The Caspase-Glo® 3/7 reagent relies on the properties of a proprietary thermostable luciferase (Ultra-Glo™ Recombinant Luciferase), which is formulated to generate a stable “glow-type” luminescent signal and improve performance across a wide range of assay conditions.

2.6.2 Assay Conditions for the ApoTox-Glo™ Triplex Assay

All Apoptox-Glo™ Triplex assays were carried out in white opaque bottom 96-well plates. 1×10^5 cells/ml were seeded into each well in a final volume of 100 μ l per well and allowed to attach for 24h. After the attachment period, culture medium was removed, first and second columns were replaced with 100 μ l medium containing vehicle and vehicle control (untreated cells and vehicle), while the other columns were replaced with increasing concentrations of test compounds in four replicate wells as described for the CCK-8 assay. Plates were placed in a 37°C incubator at 5% CO₂ and RH 80%. After exposure of SK-N-BE(2) cells for 24h, 20 μ l of viability/cytotoxicity reagent containing both GF-AFC substrate and bis-AFF-R110

substrate was added to all wells, plates were covered in foil and briefly mixed by orbital shaking at 300 rpm for 30 seconds(s). After the reagents were added and allowed to mix, plates were placed in the CO₂ incubator at 37°C for 1 hour. Following the incubation period, plates were removed from the incubator, the foil removed and the fluorescence measured at excitation (Ex) wavelength of 400 nm (λ_{ex} 400 nm) and emission wavelength of 505 nm (λ_{em} 505 nm) and λ_{ex} 485 nm / λ_{em} 520 nm for viability and cytotoxicity, respectively using the Promega GloMax™ Multiscan plate reader. To determine apoptosis, 25 μ l of Caspase-Glo® 3/7 reagent was added to each well, the plate was covered in foil, and briefly mixed by orbital shaking at 300 rpm for 30 seconds. Thereafter, the plate was incubated for 1 hour and luminescence was measured (caspase-3 activation, a hallmark of apoptosis) using the Promega GloMax™ Multiscan plate reader.

2.7 Measurement of P-Glycoprotein-Mediated Efflux Function

A number of *in vitro* assays have been used to identify compounds as MDR protein modulators, either as a substrate or as inhibitors of P-glycoprotein (Pgp; ABCB1), a member of the ATP-binding cassette (ABC) superfamily which actively exports structurally diverse hydrophobic compounds from the cell by ATP hydrolysis. Of these, the calcein-acetoxymethylester (Calcein-AM) assay has been shown to identify both substrates and inhibitors of MDR proteins, and therefore offer an advantage over other assays.⁷⁷⁸⁻⁷⁸⁴ Calcein-AM is cell-permeable non-fluorescent dye. Upon transport into live cells, its acetomethoxy group is removed by intracellular esterases, thereby trapping the compound inside the cell where it exhibits strong green fluorescence. As an MDR protein substrate, calcein-AM is rapidly excluded from cells expressing MDR protein, thus reducing fluorescent calcein in the cytosol. This property makes calcein-AM an ideal probe for identifying MDR protein overexpressing cells. Cayman's Multi-Drug Resistance Assay Kit provides a convenient tool for studying MDR protein modulators. The kit employs calcein-AM, a substrate for MDR proteins, including P-gp and MRP, as a probe for the detection of chemical compounds interacting with MDR proteins. Cyclosporin A, a competitive inhibitor, and verapamil, a non-competitive

inhibitor of P-gp, are included as positive controls. All experiments using the calcein-AM kit were carried out in black, clear bottom 96-well tissue culture treated plates. SK-N-BE(2) cells were seeded at a density of 5×10^5 cells/well in 100 μ l of complete DMEM cell culture medium and incubated allowing for cells to attach and grow overnight (24h). On the day of the experiment, the plate was centrifuged for 5 min at 400 x g at room temperature, the medium aspirated from all the wells and replaced with 100 μ l concentrations of test compounds: thapsigargin at (0.5, 1, 2 nM), brefeldin A (0.001, 0.01, 0.1 μ M), bacitracin (0.2, 0.8, 1.6 mM), aspirin (1, 8, 16 mM) and castanospermine (0.5, 1, 2 mM). Included in the kit was cyclosporin A and verapamil which were used as positive diluted 1:1000 and 1:2000, respectively, into culture medium.

The plates were incubated for 24h in CO₂ incubator at 37°C. It is recommended for positive controls to be incubated for 30 min. At the end of the specified treatment interval, 100 μ l of the prepared calcein-AM solution (2X) was added to each of the sample wells and incubated for additional 30 min in in CO₂ incubator at 37°C. Then, the plates were centrifuged for 5 min at 400 x g at room temperature. The supernatants were aspirated and another 100 μ l of the prepared calcein-AM solution (2X) added to each of sample wells and incubated for an additional 30 min in a CO₂ incubator at 37°C. The plates were again centrifuged for 5 min at 400 x g at room temperature, supernatants aspirated and finally 200 μ l of ice cold medium added to each well. The plates were analyzed immediately with a fluorescent plate reader (Promega GloMax™ Multiscan). Cells that have taken up calcein-AM display strong fluorescence intensity with excitation and emission wavelengths of 485 nm and 535 nm, respectively.

2.8 Annexin-V Cy3™ Apoptosis Assay

2.8.1 Principle of Annexin-V Cy3™ Apoptosis Assay

The annexins are a group of homologous proteins that bind phospholipids in the presence of calcium. Apoptosis, or programmed cell death (PCD), is an important mechanism that most

cells use to negatively select cells deleterious to the host. Many cells of the immune system such as thymocytes, self-reactive B- and T-cells undergo apoptosis as a result of normal cell selection processes. The cellular changes involved in the process include loss of cell membrane phospholipid asymmetry during early stages of apoptosis. In living cells, phosphatidylserine (PS) is transported to the inner plasma membrane leaflet by the enzyme Mg-ATP dependent aminophospholipid translocase. However, during the onset of apoptosis, PS is transported to the external leaflet of the plasma membrane. PS is then available for binding to annexin-V and any of its conjugates in the presence of Ca^{2+} ions. Apoptotic cells can be differentiated from necrotic cells in several ways. The method employed by this kit involves the use of two labels: Annexin-Cy3 (AnnCy3) binds to PS present in the outer leaflet of the plasma membrane of cells starting the apoptotic process. The binding is observed as red fluorescence. 6-Carboxyfluorescein diacetate (6-CFDA) is used to measure viability. When this non-fluorescent compound enters living cells, esterases present hydrolyze it, producing the fluorescent compound, 6-carboxyfluorescein (6-CF). This appears as green fluorescence. Cells can be incubated either with AnnCy3 or 6-CFDA separately, or with the two compounds simultaneously. After labelling at room temperature, the cells are immediately observed by fluorescence microscopy. Live cells will be labelled only with 6-CF (green), while necrotic cells will label only with AnnCy3 (red). Cells in the early stage of apoptosis, however, will be labelled with both AnnCy3 (red) and 6-CF (green).

2.8.2 Assay Conditions for Annexin-V Cy3™ Apoptosis Assay

SK-N-BE(2) neuroblastoma cells were seeded into into 24-well plates, at density of 5×10^5 cells per well in 1 ml of culture medium. Cells were then incubated and allowed to attach for 24h. After 24h, cells were exposed to 100 μl of relative concentrations of test compounds: aspirin (1, 8, 16 mM), bacitracin (0.2, 0.8, 16 mM), castanospermine (0.5, 1, 2 mM), brefeldin A (0.001, 0.01, 0.1 μM) and thapsigargin (0.5, 1, 2 nM). Staurosporine (1 $\mu\text{g}/\text{ml}$) was used as positive control. After inducing apoptosis using the specified concentrations of ER stress inducers, cells were washed in PBS. Thereafter, cells were trypsinized (500 $\mu\text{l}/\text{well}$) and

detached cells were transferred to a 15-ml conical centrifuge tube and centrifuged for 5 min. The cell pellet was resuspended in PBS (1 ml). A 2-mm tip PAP pen (Sigma-Aldrich), a special marking pen that delivers a thin film-like green-tinged hydrophobic barrier when a circle is drawn around a specimen on a slide, was used to draw circles of 1 cm diameter on poly-prep-poly-L-lysine-coated slides to restrict movement of cell suspension to the slide. A droplet (50 μ l) of cell suspension was deposited inside the circle and cells were allowed to attach to the slide by incubating at room temperature.

The cells were washed twice with 50 μ l of binding buffer (10 mM HEPES/NaOH, pH 7.5, containing 150 mM NaCl, 5 mM KCl and 2.5 mM CaCl_2) and 50 μ l of a double label staining solution (Sigma-Aldrich; Annexin-V Cy3.18 and 6-CFDA) added onto each circle and covered with foil. Cells were incubated at room temperature for 10 min. Slides were washed three times with 50 μ l 1X binding buffer in order to remove excess unbound staining solution. A drop of binding buffer (35 μ l) was added to the centre of each slide. A cover slip (24 X 50 mm) was placed onto the slide and results were viewed and recorded using a Nikon Eclipse 50i fluorescence microscope (IMP, Cape Town, South Africa, <http://www.imp.co.za/>).

2.9 Statistical Analysis

The Apotox-Glo™ Triplex cell viability, cytotoxicity and apoptosis assays were analyzed by One-way analysis of variance (ANOVA) followed by Dunnett's multiple comparisons test, using GraphPad Prism version 7.02 for Windows, GraphPad Software, La Jolla California USA, www.graphpad.com. Results are expressed as the mean \pm SEM (n=4), from three independent experiments. A difference of $P < 0.05$ was considered to be significant as compared to untreated SK-N-BE(2) cells (vehicle-treated controls). Transformed raw data of P-gp ATPase activity (Calcein-AM assay) were analyzed by ANOVA followed by Tukey's multiple comparisons test with the significance criterion set a priori at 0.05. Calcein-AM assay data are presented as mean \pm 95% CI (n=4).

CHAPTER 3

RESULTS AND DISCUSSION

3.1 Introduction

In this study, specific classes of N-glycosylation inhibitors (Table 1.13, Chapter 1) and pharmacologic modulators commonly used in targeting endoplasmic reticulum stress (ERS) and the unfolded protein response (UPR) signalling (Table 1.14, Chapter 1) were used to evaluate their effects on SK-N-BE(2) neuroblastoma cell proliferation, viability and induction of apoptosis.

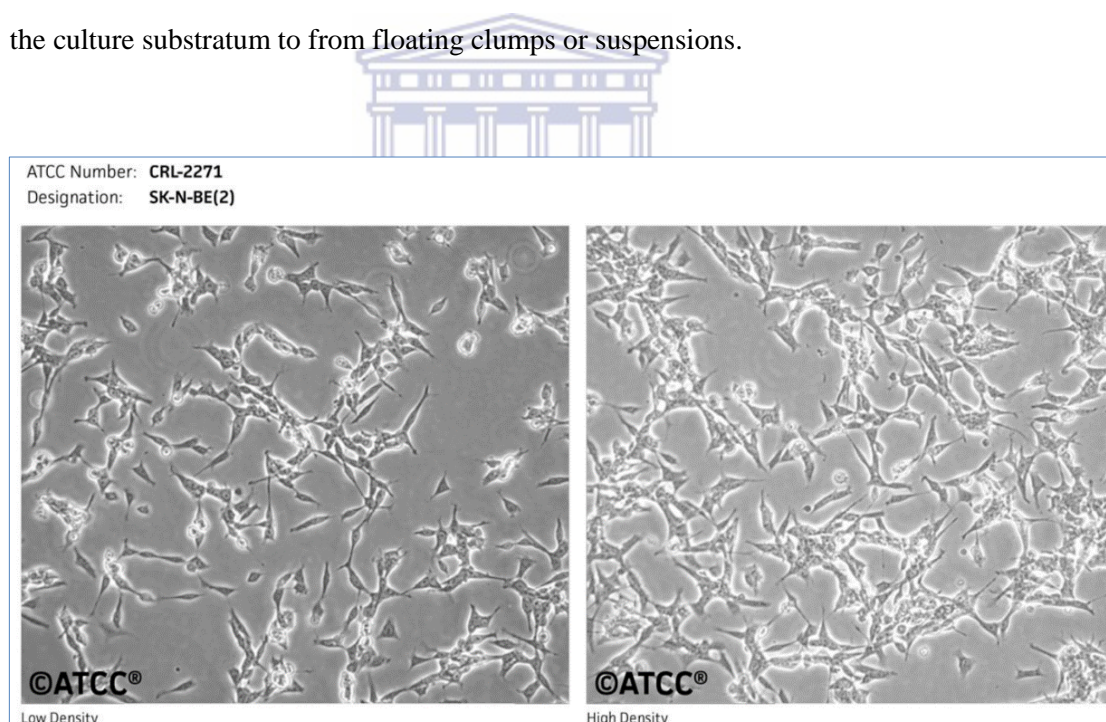
The compounds are: aspirin (acetyl salicylic acid, a non-steroidal anti-inflammatory drug known to activate PERK and upregulate pro-apoptotic transcription factor CHOP (GADD153) which, together with cleavage of caspase-12, are hallmarks of ERS-mediated responses); bacitracin (an antibiotic that ablates glycoprotein synthesis at its first stage and interferes with P-glycoprotein (P-gp) expression and localization); castanospermine (a plant alkaloid that specifically inhibits α -glycosidases I and II, thus blocking elongation of glycan chains and formation of mature glycoproteins); brefeldin A (a metabolic inhibitor of N-glycosylation and disruptor of microtubule and actin cytoskeleton organization) and thapsigargin (a potent inducer of GRP78 expression and ERS, and activator of the UPR through non-competitive inhibition of the sarcoplasmic/endoplasmic reticulum calcium ATPase/SERCA).

To evaluate the effects of these compounds on SK-N-BE(2) cells, distinctive characteristics previously reported on the cells such as morphological characteristics and their expression of P-gp were retrieved and integrated with experimental data obtained in this study, using tissue methodologies to evaluate cell growth and proliferation, multiplex fluorescence and luminescence assays for cell proliferation, viability and apoptosis, microscopic visualization

of annexin-based fluorescence apoptosis and the Calcein-AM P-gp drug efflux assay. Statistical analyses were performed using GraphPad Prism version 7.02 for Windows (GraphPad Software, La Jolla California USA, www.graphpad.com).

3.2 Morphology of SK-N-BE(2) Neuroblastoma Cells

The continuous SK-N-BE(2) neuroblastoma (NB) cell line was derived from a bone marrow biopsy taken from a 2-year old boy with disseminated (metastatic) NB after repeated courses of chemotherapy and radiotherapy.⁷⁶⁶⁻⁷⁶⁹ SK-N-BE(2) cells can achieve a saturation density (confluence) in excess of 1×10^6 cells/cm². The morphology of the cells fluctuates with some cells exhibiting long processes and others assuming an epithelioid organization (Figure 3.1). In culture, the cells often aggregate and form dense clusters or, at high density, detach from the culture substratum to form floating clumps or suspensions.



Source: <https://www.lgcstandards-atcc.org/~media/45C32586E3974C539A8D3F5579EC5920.ashx>

Figure 3.1: Morphology of SK-N-BE(2) neuroblastoma cells

3.3 Expression of P-Glycoprotein in SK-N-BE(2) Neuroblastoma Cells

SK-N-BE(2) NB cells are known to overexpress the multidrug transporter, P-glycoprotein (P-gp) (Figure 3.2A).⁷⁸⁵ Therefore, of particular relevance to the work presented in this thesis is

that P-gp is indeed expressed in SK-N-BE(2) NB cells.⁷⁸⁶ A recent study compared the expression of various members of the ATP-Binding Cassette (ABC) family of drug transporters, including ABCC1 (*MRP1*), ABCC2 (*MPR2*), ABCC6 (*mrp6*), ABCC8 (*mrp8*), ABCC10 (*mrp10*), ABCC11 (*mrp11*), ABCC12 (*mrp12*) and ABCC13 (*mrp13*), and found that these were expressed either at similar or elevated levels in drug-resistant NB cell lines compared to parental controls (Table 3.1).⁷⁸⁶ Western immunoblotting demonstrated significantly greater upregulation of P-gp in the SK-N-BE(2) subset doxorubicin-resistant (DoxR) SK-N-BE(2)C cells than the vorinostat-treated doxorubicin-resistant (DoxR-v) cells, relative to wild-type (WT) parental cells (Figure 3.2B).⁷⁶⁴

Table 3.1: Relative expression of known drug-resistance genes in neuroblastoma cell lines

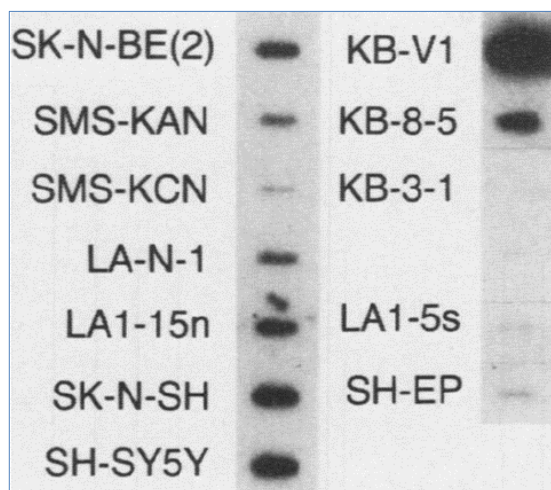
Gene Symbol	Gene Entrez	DoxR		DoxR-v	
		SK-N-SH	SK-N-Be(2)C	SK-N-SH	SK-N-Be(2)C
ABCB1 (<i>mdr1</i>)	5243	4.63	4.23	1.98	2.04
ABCB6 (<i>prp</i>)	10058	1.68	1.81	1.57	1.81
ABCC3 (<i>mrp3</i>)	8714	N.D.	-2.39	N.D.	N.D.
ABCC4 (<i>mrp4</i>)	10257	2.17	2.12	N.D.	N.D.
ABCC5 (<i>mrp5</i>)	10057	-1.63	N.D.	-2.26	-2.68
ABCC9 (<i>mrp9</i>)	10060	-2.76	-4.15	-2.22	-2.89
BCL-2	596	1.53	N.D.	N.D.	N.D.
SIRT1	23411	1.75	2.02	N.D.	N.D.
BDNF	627	3.69	5.17	2.56	3.60
TH	7054	-1.87	-2.47	25.5	31.0

Relative expression of known drug-resistance genes in doxorubicin resistant (DoxR) and vorinostat-treated doxorubicin-resistant (DoxR-v) cells compared to the parental lines. Results are expressed as a fold-change (all $p < 0.1$). N.D. indicates no difference in gene expression (fold-change, 1.5 and/or $p < 0.1$). The following genes had no significant difference in any comparison: ABCC1 (*MRP1*), ABCC2 (*MPR2*), ABCC6 (*mrp6*), ABCC8 (*mrp8*), ABCC10 (*mrp10*), ABCC11 (*mrp11*), ABCC12 (*mrp12*), ABCC13 (*mrp13*), MGMT, SOD, HDAC1-8. doi:10.1371/journal.pone.0040816.t001.

Source:⁷⁶⁴ Lautz TB, Jie C, Clark S, Naiditch JA, Jafari N, Qiu YY, Zheng X, Chu F, Madonna MB. The effect of vorinostat on the development of resistance to doxorubicin in neuroblastoma. *PloS One* 2012;7(7):e40816, with permission (This is an open-access article distributed under the terms of the Creative Commons Attribution License, which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited, <https://www.ncbi.nlm.nih.gov/pmc/articles/PMC3400660/>).

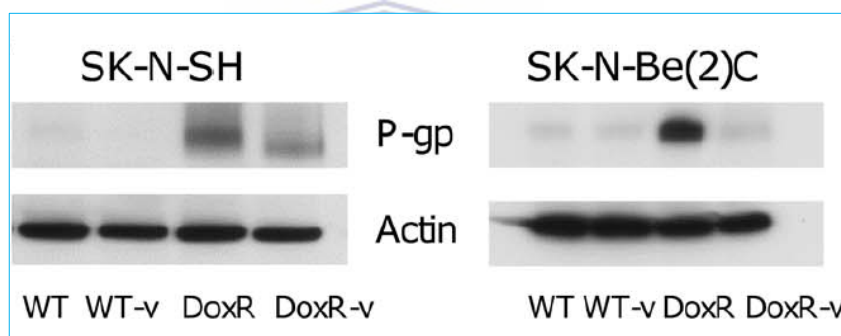
3.4 Growth Curve Analysis of SK-N-BE(2) Neuroblastoma Cells

A cell viability growth curve for SK-N-BE(2) cells is shown in Figure 3.3. SK-N-BE(2) NB cells were seeded at a density of 100,000 cells per well onto 96-well plates and subsequently monitored over a 4-day period.



A. Expression of *mdrl/Pgp* in nine human neuroblastoma cell lines.

Source:⁷⁸⁶ Bates SE, Mickley LA, Chen YN, Richert N, Rudick J, Biedler JL, Fojo AT. Expression of a drug resistance gene in human neuroblastoma cell lines: Modulation by retinoic acid-induced differentiation. *Molecular and Cellular Biology* 1989;9(10):4337-4344, with permission from the American Society for Microbiology (ASM), under the Creative Commons license and ASM Journals Public Access Policy (http://journals.asm.org/site/misc/index_compliance.xhtml).

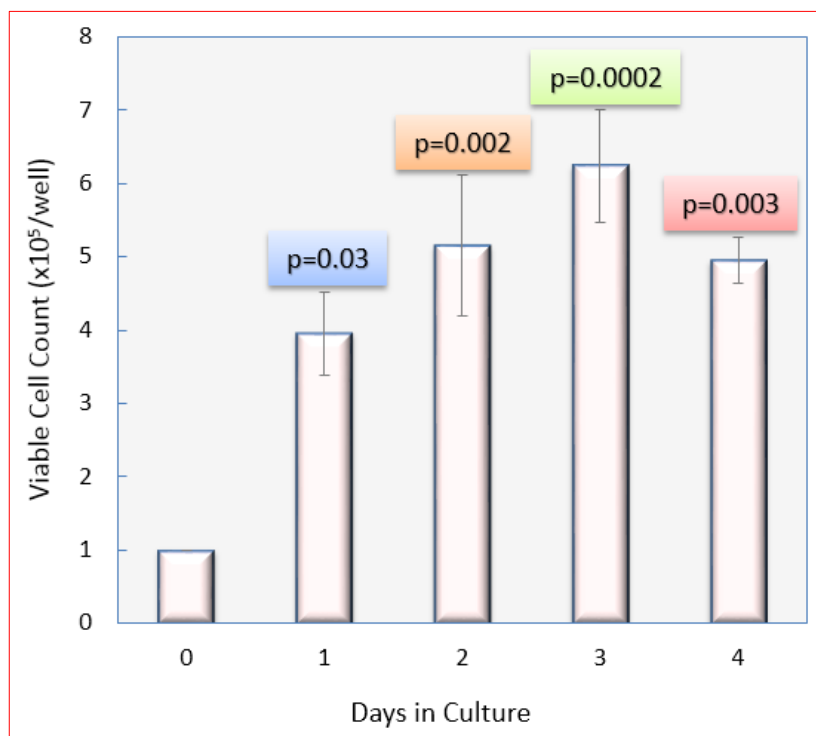


B. Western immunoblotting of P-gp in doxorubicin-resistant (DoxR) and vorinostat-treated doxorubicin-resistant (DoxR-v) cells compared to wild-type (WT) cells.

Source:⁷⁶⁴ Lautz TB, Jie C, Clark S, Naiditch JA, Jafari N, Qiu YY, Zheng X, Chu F, Madonna MB. The effect of vorinostat on the development of resistance to doxorubicin in neuroblastoma. *PloS One* 2012;7(7):e40816, with permission (This is an open-access article distributed under the terms of the Creative Commons Attribution License, which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited, <https://www.ncbi.nlm.nih.gov/pmc/articles/PMC3400660/>).

Figure 3.2: Upregulation of P-glycoprotein expression in neuroblastoma cell lines

Cell growth progressed through an initial log phase and then reached a plateau by day 3, after which time the viability began to decline. Viable cell counts were compared by one-way ANOVA with Tukey's multiple comparisons post-hoc test using GraphPad Prism 7 (www.graphpad.com). The one-way ANOVA tests yielded p values as indicated for control vs days 1, 2, 3 and 4, respectively. On all days, the viable cell count of SK-N-BE(2) NB cells was significantly greater ($p < 0.05$) than that of day 0.

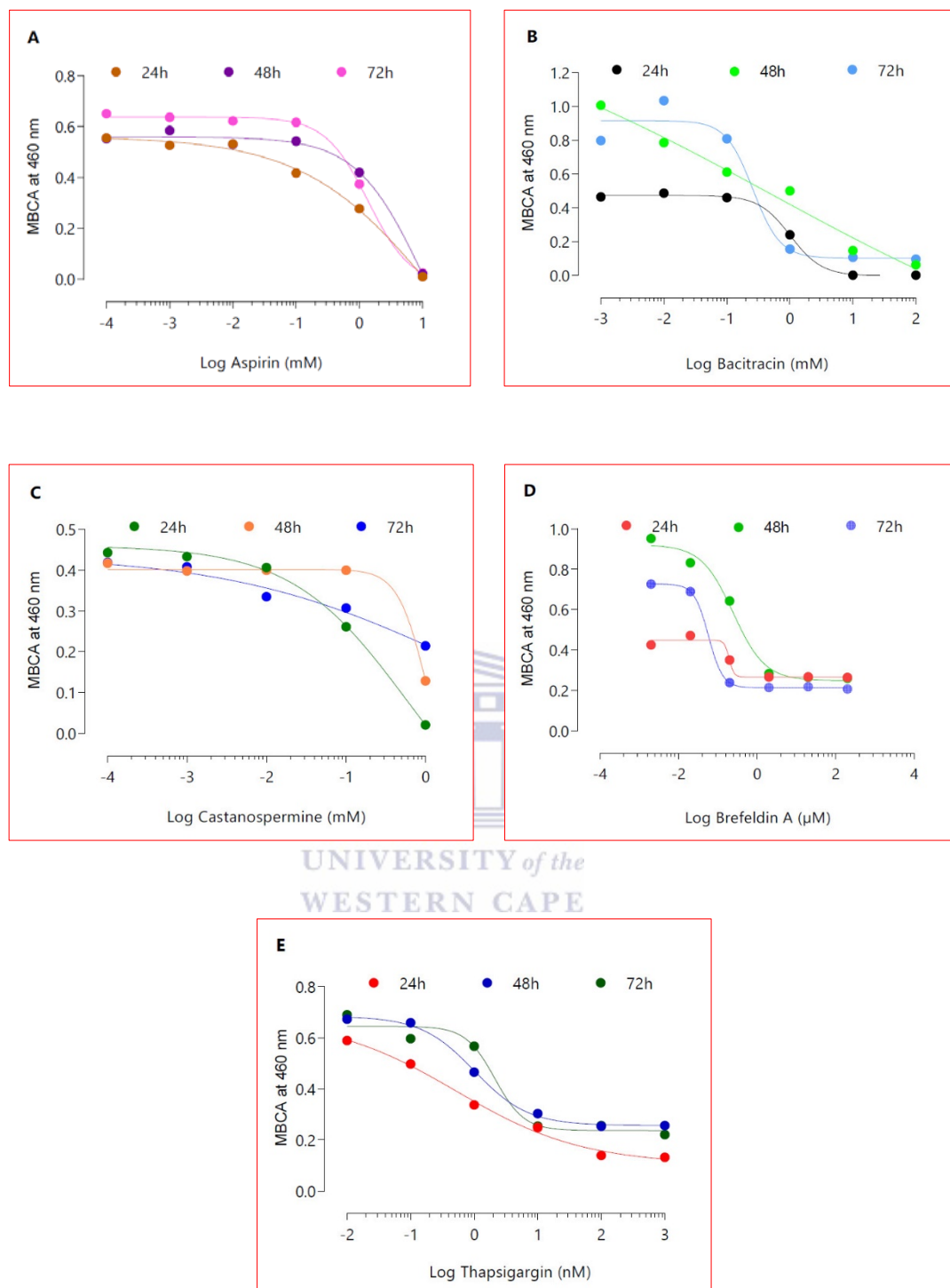


Values are means \pm SEM (n=4) of a representative experiment. One-way ANOVA (Tukey's multiple comparisons test) yielded p values as indicated for control vs days 1, 2, 3 and 4, respectively.

Figure 3.3: Growth curve analysis of SK-N-BE(2) neuroblastoma cells in culture

3.5 Cell Counting Kit-8 (CCK-8) Cell Viability Assays

SK-N-BE(2) NB cell proliferation was assessed by the CCK-8 cell viability assay as described in the research methodology (Chapter 2). Cells (5×10^4 cells/ml) were seeded in 96-well plates and incubated for 24h, 48h and 72h with incremental \log_{10} concentrations of aspirin, bacitracin, castanospermine, brefeldin A and thapsigargin. Cell viability in the presence of different concentrations of these ERS inducers or glycoprotein processing inhibitors was determined by comparison with untreated control cells, i.e., the data points represent blank-corrected absorbances at 460 nm. Figure 3.4 shows the CCK-8 dose-response curves for test compounds assessed at varying concentrations using the GraphPad Prism 7 (www.graphpad.com) four-parameter non-linear regression model with variable Hill slope. Table 3.2 summarizes the non-linear regression analysis data and validation parameters of the respective dose-response curves for the test compounds.



SK-N-BE(2) neuroblastoma cells were exposed to \log_{10} increments of ERS inducer or glycoprotein processing inhibitor for the times indicated. Data points (blank-corrected absorbance at 460 nm) are the means of quadruplicate measurements ($n=4$), representing one of 3 independent experiments. The non-linear regression model used does not assume a standard slope, but rather fits the Hill slope from the data, and so is called a Variable slope model or a four-parameter dose-response curve, or four-parameter logistic curve, abbreviated as 4PL. MBCA, mean blank-corrected absorbance.

Figure 3.4: CCK-8 dose-response curves for test compounds assessed at varying concentrations

Table 3.2: Regression analysis data and summary of dose-response parameters

Drug (μM)	Exposure Time (Hours)	IC ₅₀ (μM)	95% CI (μM)	R ²
Aspirin (mM)	24	16.16	6.74e-007 to 3.87e008	0.99
	48	9.95	1.58e-008 to 6.27e009	0.99
	72	1.39	0.60 to 3.22	0.99
Bacitracin (mM)	24	1.02	0.75 to 1.38	0.99
	48	0.50	2.22e-013 to 1.1e012	0.98
	72	0.26	0.009 to 7.81	0.96
Castanospermine (mM)	24	0.58	0.00017 to 1903	0.99
	48	1.04	0 to ∞	0.99
	72	2.22	0 to 4.13e026	0.97
Brefeldin A (μM)	24	0.20	0 to ∞	0.97
	48	0.25	0.06 to 1.02	0.99
	72	0.06	0.042 to 0.08	0.99
Thapsigargin (nM)	24	0.52	0.02 to 15.56	0.99
	48	1.01	0.54 to 1.89	0.99
	72	2.12	0.14 to 32.79	0.98

IC₅₀, half maximal inhibitory concentration of a drug estimated by the non-linear four-parameter logistic regression model; ∞ , infinity symbol; 95% CI, 95 percent confidence interval; R², regression coefficient (goodness of fit). All regression plots passed the test for homoscedasticity (equal variances, homogeneity of variance, i.e., same scatter across the independent variable around the regression line).

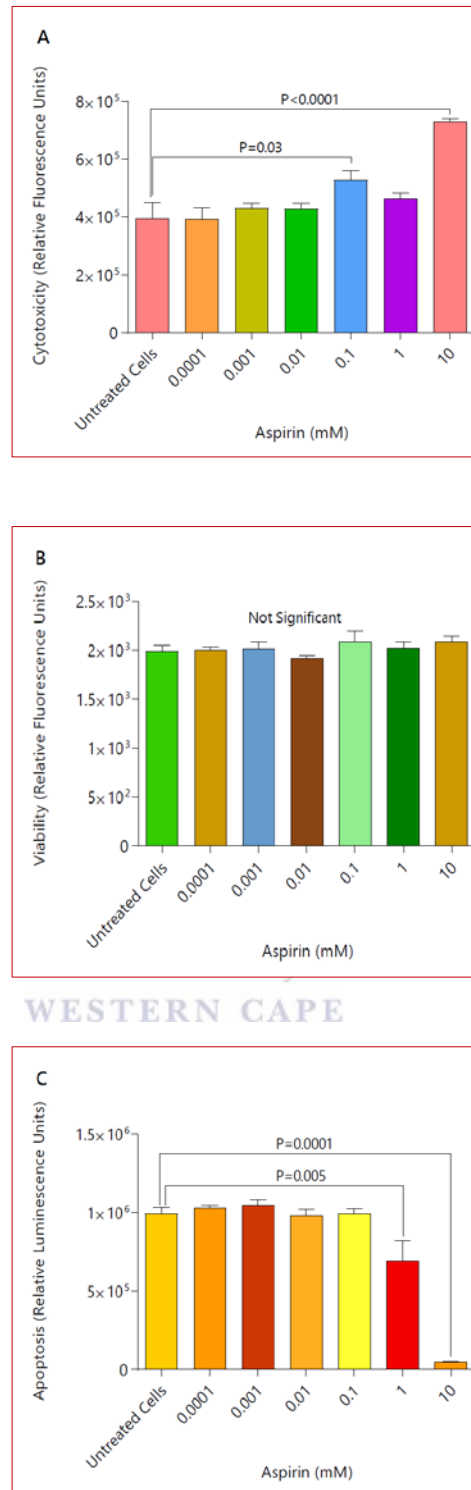
The half maximal inhibitory concentrations (IC₅₀) of aspirin for SK-N-BE(2) NB cells after 24h, 48h and 72h exposure times were 16.16 μM (95% CI: 6.74e-007 to 3.87e008; R²=0.99), 9.95 μM (95% CI: 1.58e-008 to 6.27e009; R²=0.99) and 1.39 μM (95% CI: 0.60 to 3.22; R²=0.99), respectively. Thus, aspirin exhibited the greatest potency after 72h exposure (Figure 3.4A and Table 3.2). In the case of bacitracin, a similar pattern to that of aspirin was observed, with a peak potency after 72h, i.e., the lowest IC₅₀ of 0.26 μM (95% CI: 0.009 to 7.81; R²=0.96), representing a 4-fold and 2-fold increased potency over values estimated for 24h (IC₅₀ of 1.02 μM ; 95% CI: 0.75 to 1.38; R²=0.99) and 48h (IC₅₀ of 0.5 μM ; 95% CI: 2.22e-013 to 1.1e012; R²=0.98), respectively (Figure 3.4B and Table 3.2).

Castanospermine showed an inverse potency pattern (decreased IC_{50} equates to increased potency) compared to bacitracin, i.e., its IC_{50} of 2.22 μM (95% CI: 0 to 4.13e026; $R^2=0.97$) increased 4-fold and 2-fold over values obtained for 24h (IC_{50} of 0.58 μM ; 95% CI: 0.00017 to 1903; $R^2=0.99$) and 48h (IC_{50} of 0.5 μM ; 95% CI: 0 to ∞ ; $R^2=0.99$), respectively (Figure 3.4C and Table 3.2). A 95% CI of 0 to ∞ , in the case of the 48h exposure of SK-N-BE(2) NB cells to castanospermine signifies that the non-linear dose-response curve did not entirely fit the four-parameter logistic model, despite a regression coefficient of 0.99 (Figure 3.4C and Table 3.2).

Almost identical potencies were obtained for brefeldin A, following exposure of SK-N-BE(2) NB cells to this plant alkaloid inhibitor of glycosidases for 24h (IC_{50} of 0.20 μM ; 95% CI: 0 to ∞ ; $R^2=0.97$) and 48h (IC_{50} of 0.25 μM ; 95% CI: 0.06 to 1.02; $R^2=0.99$). However, brefeldin A exerted the greatest potency after 72h (IC_{50} of 0.06 μM ; 95% CI: 0.042 to 0.08; $R^2=0.99$), representing a 3-fold and 4-fold increase in potency over the 24h and 48h exposure periods, respectively (Figure 3.4D and Table 3.2). Thapsigargin's potency profile (Figure 3.4E and Table 3.2) mimics closely that of castanospermine (Figure 3.4C and Table 3.2).

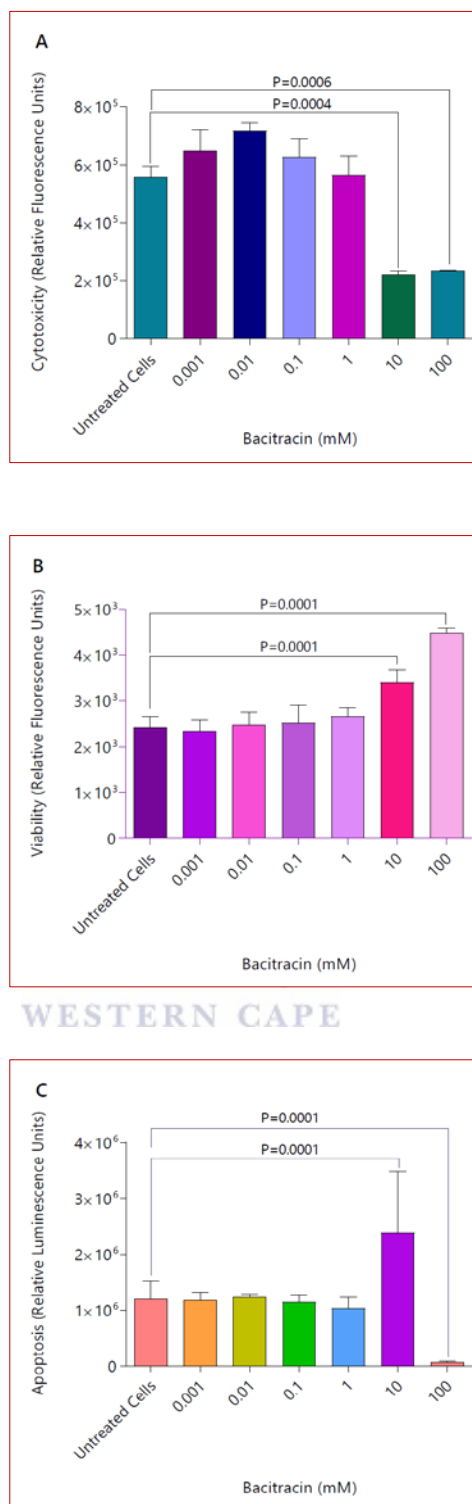
3.6 Apotox-Glo™ Triplex Cell Cytotoxicity, Viability and Apoptosis Assays

The effects of aspirin, bacitracin, castanospermine, brefeldin A and thapsigargin on SK-N-BE(2) neuroblastoma cell cytotoxicity, viability and apoptosis were measured using the Promega MultiTox-Fluor™ and Caspase-Glo® 3/7 Triplex assay kit according to the manufacturer's protocol. The results are presented in Figures 3.5 to 3.9 below and interpreted as follows: The Triplex cell-based assay simultaneously measures three parameters—cell viability, cytotoxicity, and apoptosis. The method combines two fluorescent and one luminescent assay chemistries offered by Promega (Caspase-Glo® 3/7 and MultiTox-Fluor™ Assays) in the same assay well to extract information about viability, cytotoxicity and caspase activation events. These parameters are particularly useful to define mechanisms associated with a cytotoxic profile of an investigative compound.



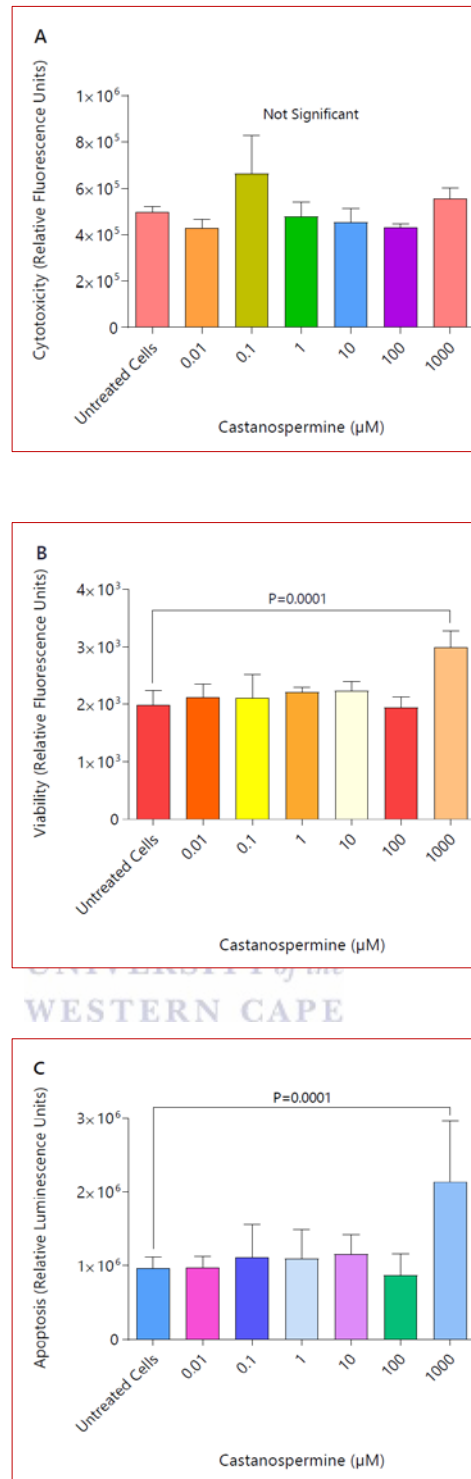
Values are means \pm SEM (n=4) of a representative experiment. One-way ANOVA followed by Dunnett's multiple comparisons test yielded P values for comparison of treated vs untreated SK-N-BE(2) cells (vehicle-treated controls).

Figure 3.5: Effects of aspirin on SK-N-BE(2) NB cell cytotoxicity, viability and apoptosis



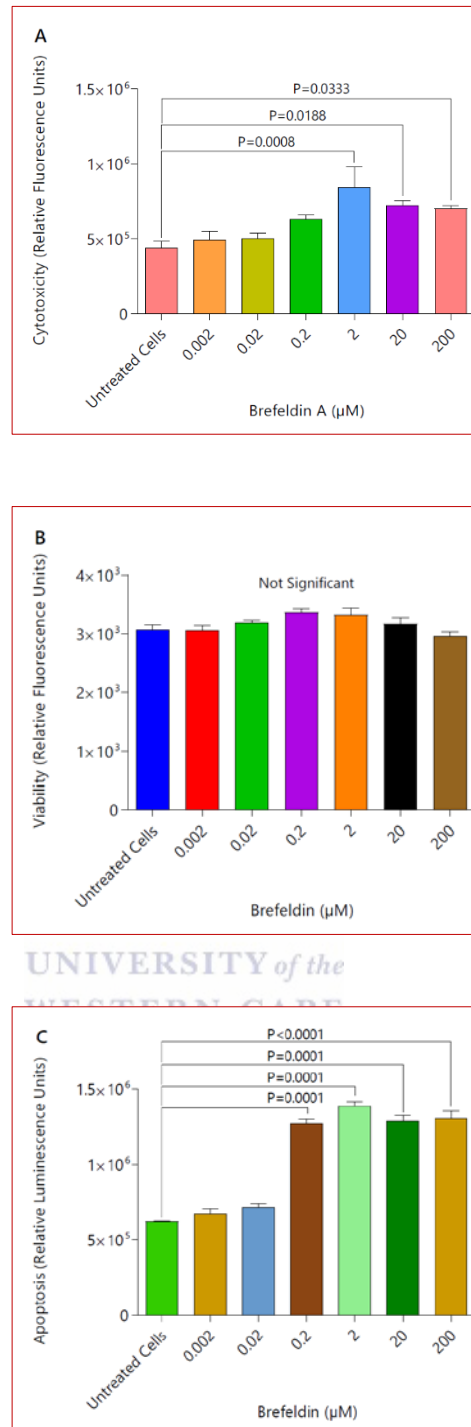
Values are means \pm SEM (n=4) of a representative experiment. One-way ANOVA followed by Dunnett's multiple comparisons test yielded P values for comparison of treated vs untreated SK-N-BE(2) cells (vehicle-treated controls).

Figure 3.6: Effects of bacitracin on SK-N-BE(2) NB cell cytotoxicity, viability and apoptosis



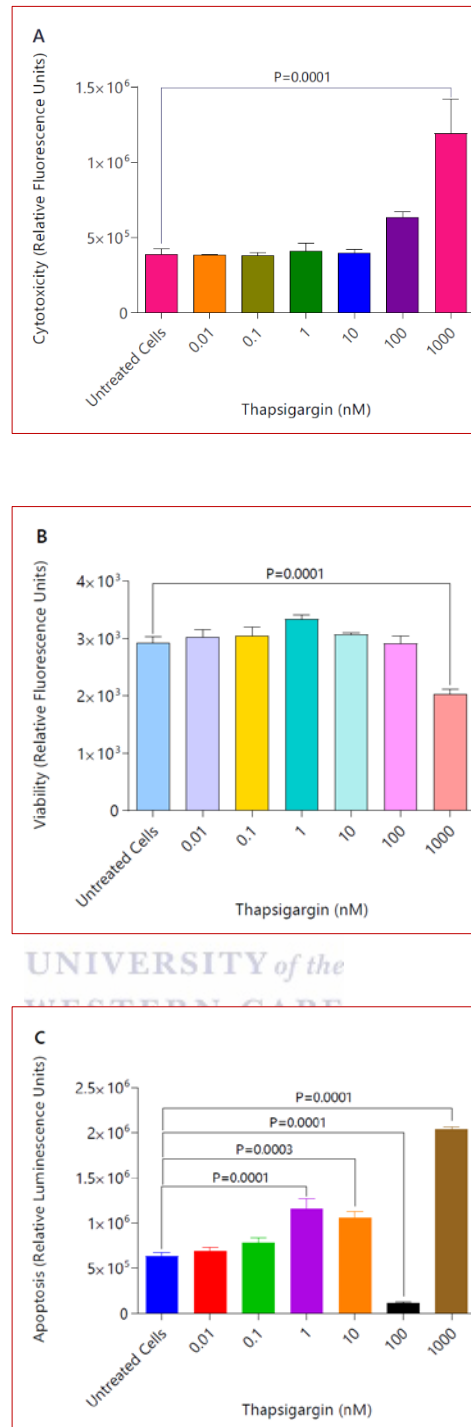
Values are means \pm SEM (n=4) of a representative experiment. One-way ANOVA followed by Dunnett's multiple comparisons test yielded P values for comparison of treated vs untreated SK-N-BE(2) cells (vehicle-treated controls).

Figure 3.7: Effects of castanospermine on SK-N-BE(2) NB cell cytotoxicity, viability and apoptosis



Values are means \pm SEM ($n=4$) of a representative experiment. One-way ANOVA followed by Dunnett's multiple comparisons test yielded P values for comparison of treated vs untreated SK-N-BE(2) cells (vehicle-treated controls).

Figure 3.8: Effects of brefeldin A on SK-N-BE(2) NB cell cytotoxicity, viability and apoptosis



Values are means \pm SEM (n=4) of a representative experiment. One-way ANOVA followed by Dunnett's multiple comparisons test yielded P values for comparison of treated vs untreated SK-N-BE(2) cells (vehicle-treated controls).

Figure 3.9: Effects of thapsigargin on SK-N-BE(2) NB cell cytotoxicity, viability and apoptosis

The Triplex assay is comprised of the Promega MultiTox-Fluor™ and Caspase-Glo® 3/7 Assays. The MultiTox-Fluor™ Assay is a non-lytic chemistry that allows measurement of live and dead cells in a single sample well. Specifically, for live cell assessment, live-cell protease activity is measured by the fluorogenic, cell-permeant peptide substrate Gly-Phe-7-amino-4-trifluoromethyl coumarin (GFAFC). This live-cell protease activity marker labels only live cells because it becomes inactive upon loss of membrane integrity and leakage into the surrounding culture medium, and thus does not contribute to the dead cell measurement.

For dead cell assessment, a second protease activity marker, the cell-impermeant peptide substrate bis-(Ala-Ala-Phe)-rhodamine 110 (bis-AAFR110), is used to measure the activity of a dead-cell protease from cells that have lost membrane integrity and leaked the biomarker into the surrounding culture medium.

The Caspase-Glo® 3/7 Assay is a luminescent assay that measures caspase-3 and -7 activities in cultures of cells, which are indicative of apoptosis. The assay provides a proluminescent caspase-3/7 substrate which contains the tetrapeptide sequence DEVD (aspartic acid, glutamic acid, valine, aspartic acid). This substrate is cleaved to release aminoluciferin, a substrate of luciferase used in the production of light. The amount of light produced correlates with caspase-3/7 activity.

Together, these assays provide a researcher with three data parameters per well (cell viability, cytotoxicity, and caspase activity) which can be used to more accurately profile compound effects on cells. The following convention pertains: cytotoxicity (MultiTox-Fluor, *dead cells*), viability (Multi-Tox-Fluor, *live cells*) and Caspase-Glo 3/7 (*apoptosis-caspase activity*).^{773,776}

Compared to the untreated cell control, the cytotoxicity fluorescence of 0.1 and 10 mM aspirin increased significantly, $p=0.003$ and $p<0.0001$, respectively. The other concentrations did not produce any cytotoxic effects on SK-N-BE(2) neuroblastoma cells (Figure 3.5A). Aspirin in the \log_{10} increment concentration range of 0.0001 to 10 mM had no effect on the viability

fluorescence of SK-N-BE(2) cells (Figure 3.5B), but at 1 mM ($p=0.005$) and 10 mM ($p=0.0001$) caused significant reduction in apoptosis luminescence (Figure 3.5C) compared with control.

Bacitracin at concentrations of 10 ($p=0.0004$) and 100 mM ($p=0.0006$) produced a reduction in cytotoxicity with a commensurate increase in viability (Figures 3.6A and B). However, at 10 mM ($p=0.0001$), bacitracin significantly increased caspase-dependent apoptosis whereas at 100 mM ($p=0.0001$), apoptosis was significantly reduced in SK-N-BE(2) cells relative to untreated controls (Figure 3.6C).

Castanospermine in the \log_{10} concentration range of 0.01 to 1000 μM had no cytotoxic effect on SK-N-BE(2) cells (Figure 3.7A), whereas castanospermine concentrations of 0.01 to 100 μM did not affect cell viability (Figure 3.7B), but somewhat perplexing at 1000 μM , viability (Figure 3.7B) and apoptosis (Figure 3.7C) in these cells were significantly increased ($p=0.0001$) over that of untreated control.

Brefeldin A concentrations of 2 μM ($p=0.0008$), 20 μM ($p=0.0188$) and 200 μM ($p=0.0333$) exerted significant cytotoxic effects on SK-N-BE(2) cells (Figure 3.8A), but in the \log_{10} concentration range of 0.002 to 200 μM did not affect cell viability (Figure 3.8B). The cytotoxicity of brefeldin A (Figure 3.8A) was commensurate with an increase in apoptosis after treatment of SK-N-BE(2) cells with 0.2, 2, 20 and 200 μM concentrations ($p\leq 0.0001$ in all cases, Figure 3.8C).

Thapsigargin only significantly increased cytotoxicity fluorescence in SK-N-BE(2) cells at a concentration of 1000 nM ($p=0.0001$, Figure 3.9A), commensurate with a decrease in viability fluorescence at the same concentration ($p=0.0001$, Figure 3.9B). Apoptosis luminescence in SK-N-BE(2) cells was increased at thapsigargin concentrations of 1 nM ($p=0.0001$), 10 nM ($p=0.0003$) and 1000 nM ($p=0.0001$), but decreased at 100 nM ($p=0.0001$, Figure 3.9C).

3.7 Annexin-V Cy3™ Apoptosis Assays

The Annexin-V Cy3™ Apoptosis Assay kit provides a rapid and convenient assay for apoptosis in cells. Annexin-Cy3 binds to phosphatidylserine (PS) in apoptotic cells and is visualized as red fluorescence. 6-Carboxyfluorescein diacetate (6-CFDA) is used to measure cell viability. Upon entering of 6-CFDA (non-fluorescent compound) into living cells, esterases cleave it, producing the fluorescent compound, 6-carboxyfluorescein (6-CF). This appears as green fluorescence. Cells are incubated either with Annexin-Cy3 or 6-CFDA separately or simultaneously. After labelling at room temperature, the cells are immediately observed by fluorescence microscopy. Live cells will be labelled only with 6-CF (*green*), while necrotic cells will label only with Annexin-Cy3 (*red*). Cells in the early stage of apoptosis, however, will be labelled with both Annexin-Cy3 (*red*) and 6-CF (*green*).

To induce apoptosis, SK-N-BE(2) cells were exposed for 24h to selected concentrations of aspirin (1, 8, 16 mM), bacitracin (0.2, 0.8, 16 mM), castanospermine (0.5, 1, 2 mM), brefeldin A (0.001, 0.01, 0.1 μ M) and thapsigargin (0.5, 1, 2 nM), using staurosporine as a positive control for apoptosis, as described in the research methodology (Chapter 2). Morphological observation of apoptosis induced by these compounds in SK-N-BE(2) cells was done by fluorescence microscopy and images were processed using Nikon Eclipse 50i software (IMP, Cape Town, South Africa, <http://www.imp.co.za/>).

Figure 3.10 shows the fluorescence micrographs of the effects of aspirin on SK-N-BE(2) cell apoptosis. Untreated SK-N-BE(2) cells displayed live cells (green fluorescence) as well as canonical apoptotic transformation as evidenced by various morphological changes, including apoptotic bodies, nuclear condensation and cell shrinkage, which were also observed under fluorescence microscopy, but very few necrotic cells (red fluorescence, Figures 3.10A to 3.14A). The staurosporine apoptosis-positive control micrograph shows mostly live cells and very few green-and-red fluorescing cells (cells in the process of undergoing apoptosis) and some fully apoptosed cells (Figure 3.10B to 3.14B).

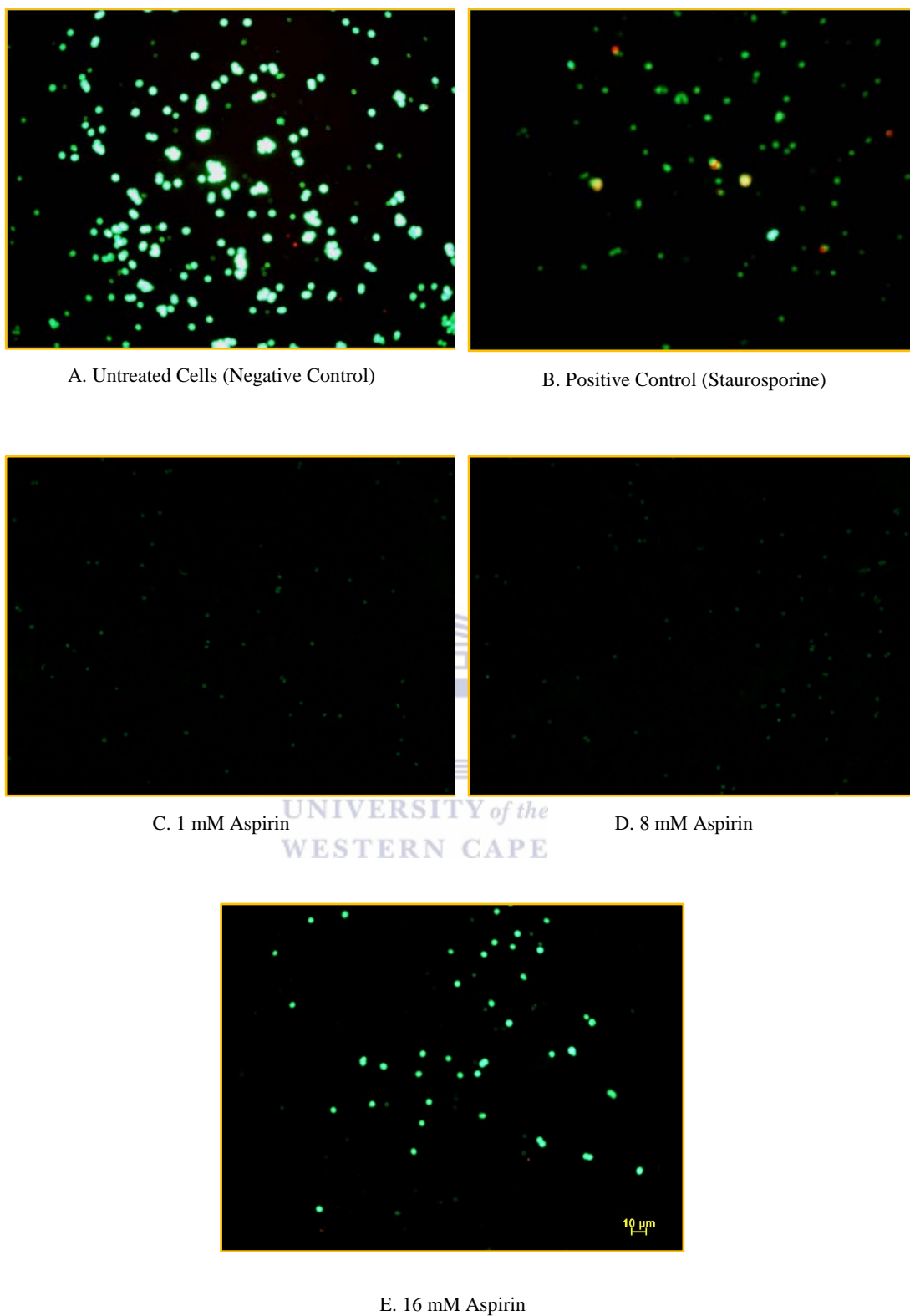


Figure 3.10: Fluorescence micrographs of the effects of aspirin on SK-N-BE(2) cell apoptosis

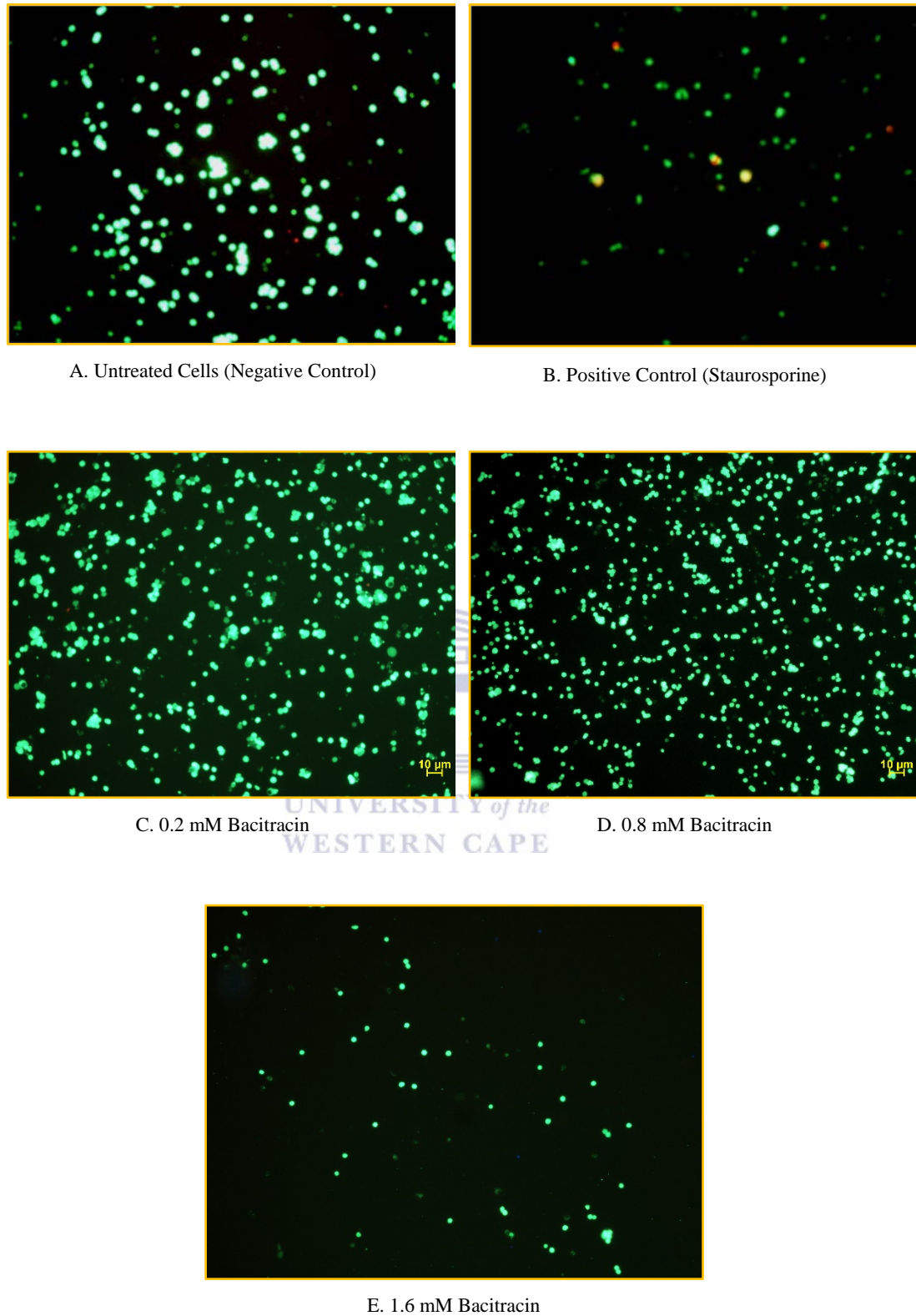


Figure 3.11: Fluorescence micrographs of the effects of bacitracin on SK-N-BE(2) cell apoptosis

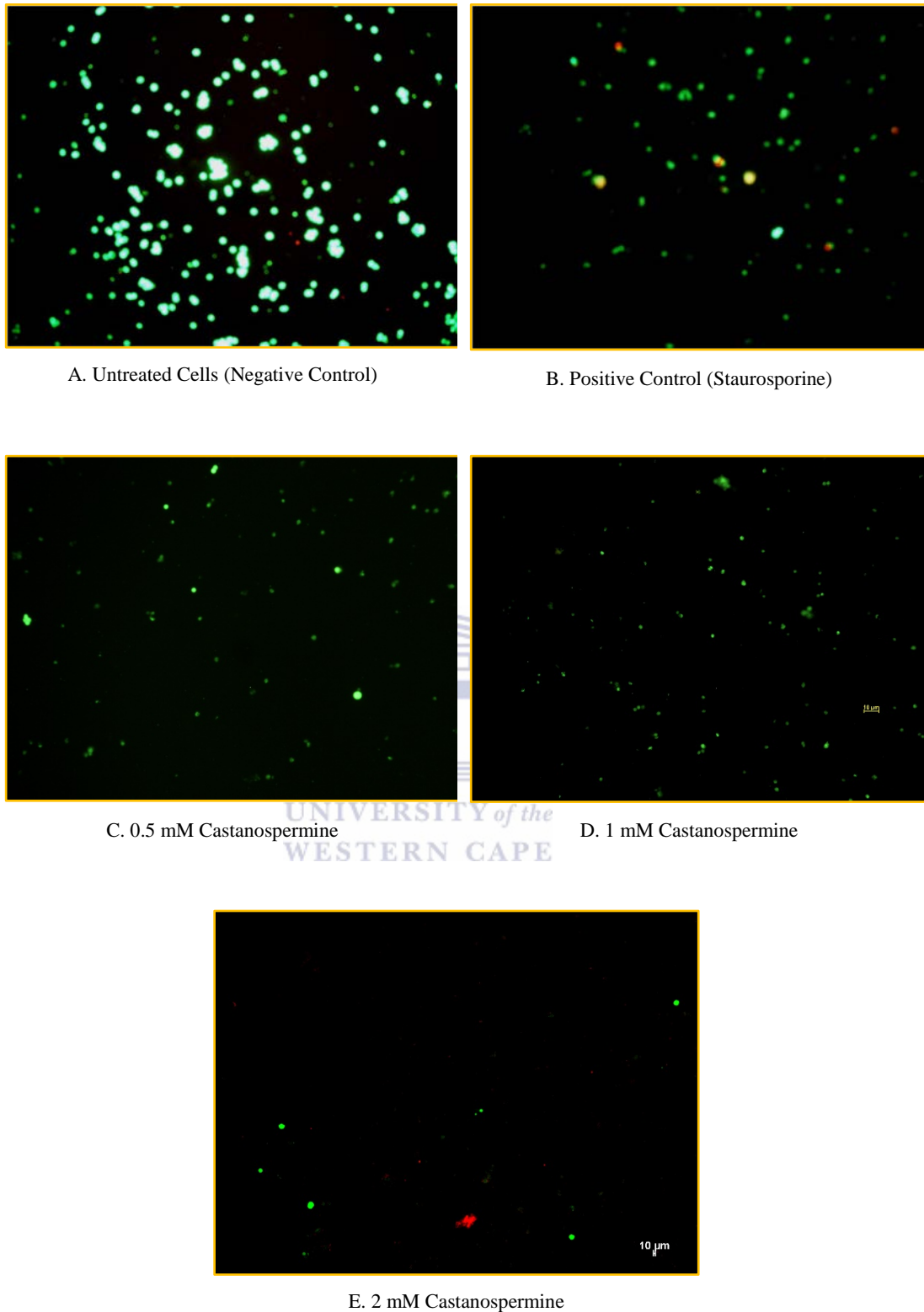


Figure 3.12: Fluorescence micrographs of the effects of castanospermine on SK-N-BE(2) cell apoptosis

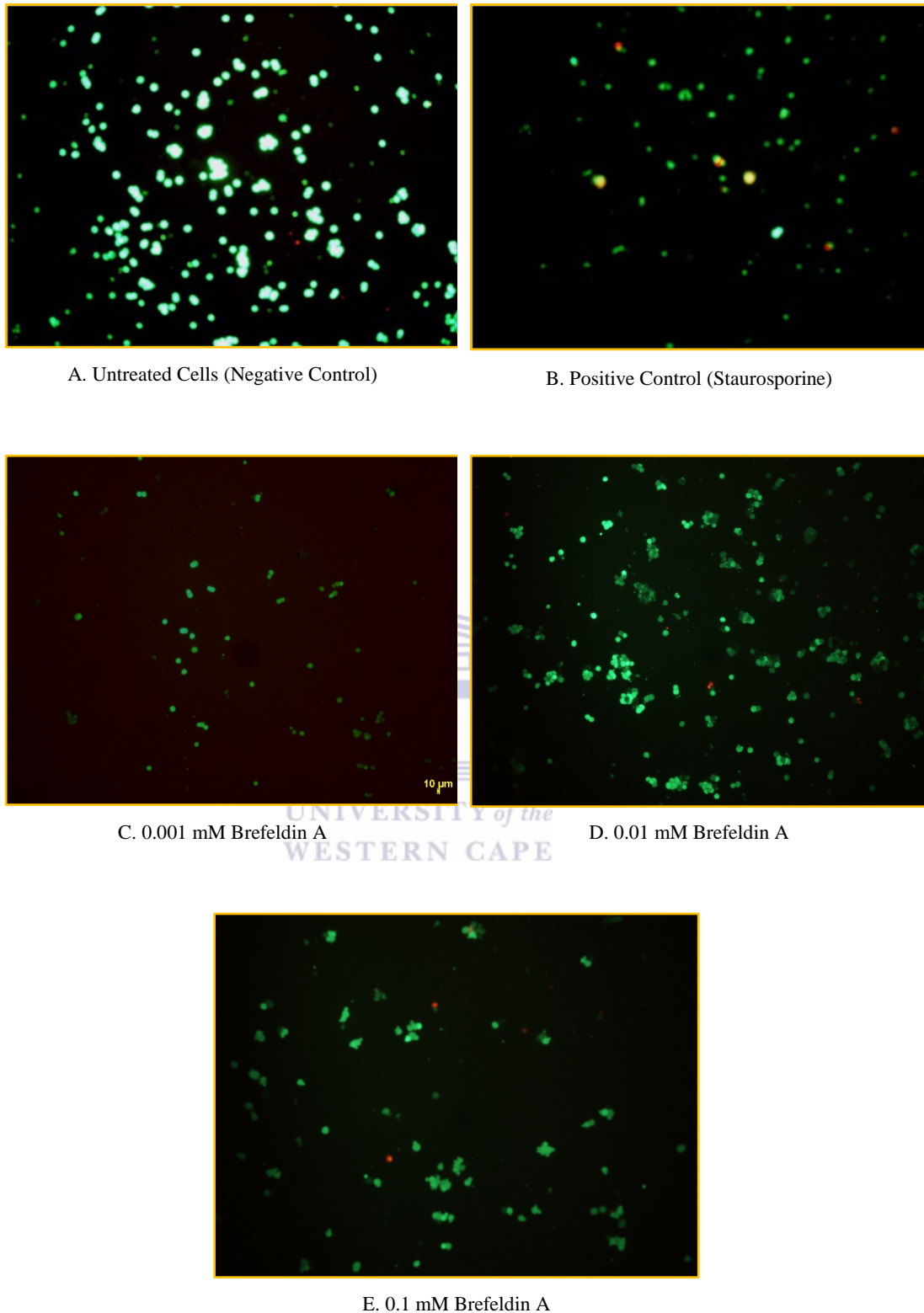


Figure 3.13: Fluorescence micrographs of the effects of brefeldin A on SK-N-BE(2) cell apoptosis

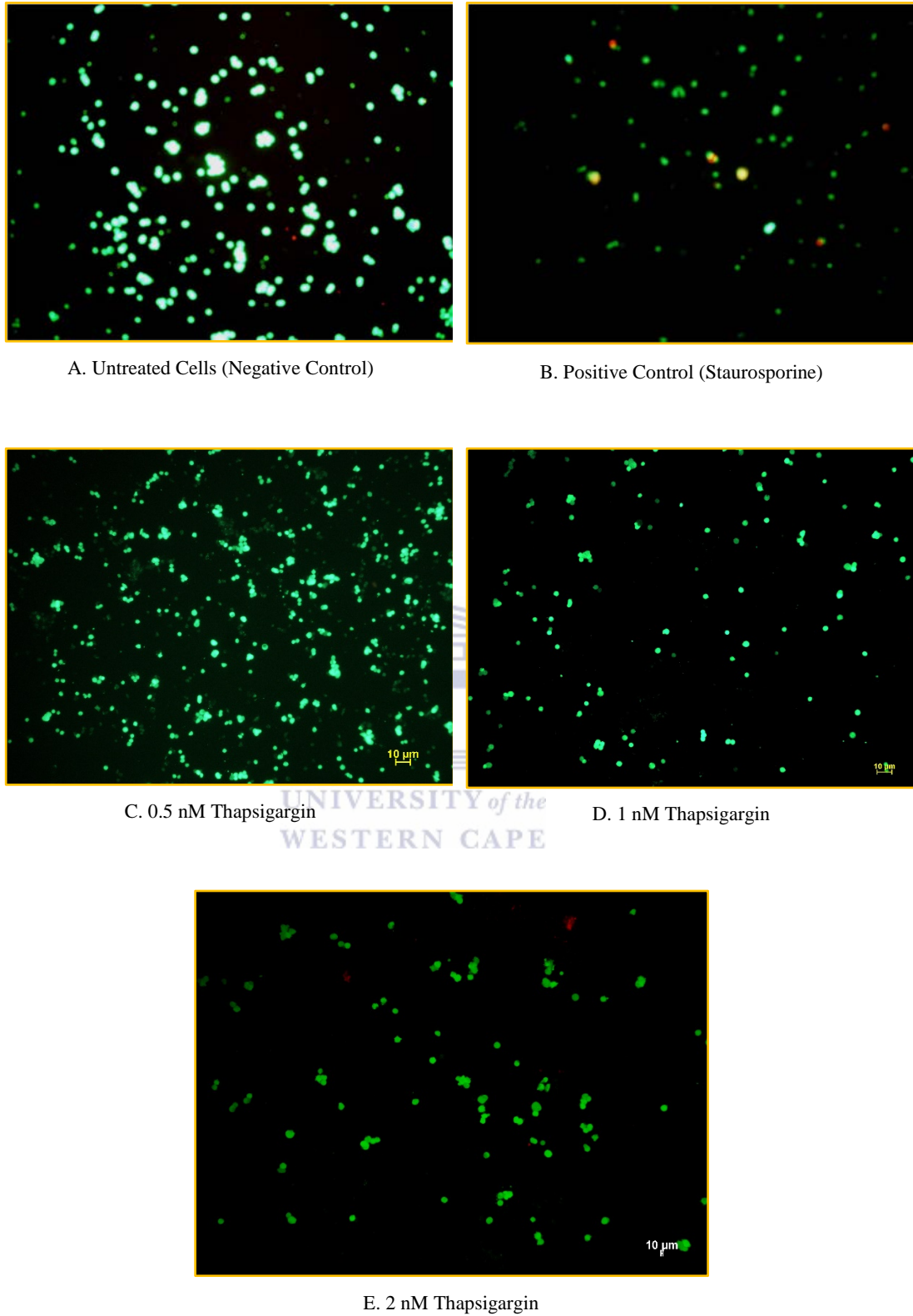


Figure 3.14: Fluorescence micrographs of the effects of thapsigargin on SK-N-BE(2) cell apoptosis

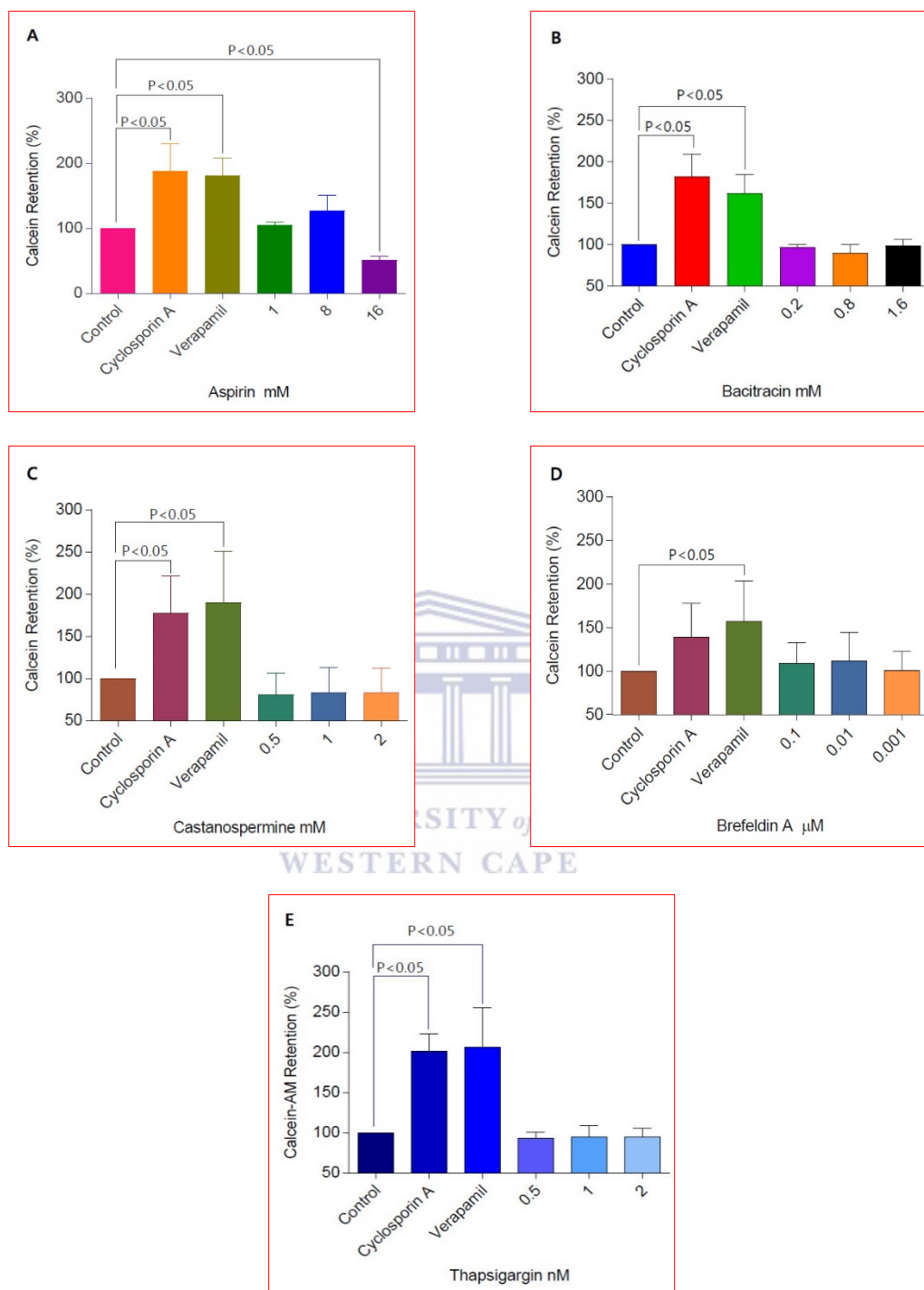
SK-N-BE(2) cells treated with 1, 8 and 16 mM aspirin showed exclusively live cells (Figure 3.10C-E). Exposure of cells to 0.2 and 0.8 mM bacitracin yielded equal proportions of live and mitotic cells (Figure 3.11C and D), whereas 1.6 mM bacitracin showed only live non-dividing cells (Figure 3.11E). Castanospermine at 0.5, 1 and 5 mM also did not produce any observable apoptosis luminescence in SK-N-BE(2) cells (Figure 3.12C-E) as did 0.001 mM brefeldin A (Figure 3.13C), but a negligible percentage of apoptosis-positive cells was noted in cells treated with 0.01 and 0.1 mM brefeldin A (Figure 3.13D and E). Similarly, thapsigargin at 0.5, 1 and 2 nM did not produce any significant apoptosis in SK-N-BE(2) cells (Figure 3.14 C to E).

3.8 Measurement of P-Glycoprotein-Mediated Efflux Function

Calcein–acetoxymethylester (calcein-AM) was used as a neutral substrate to determine ABC transporter activity, i.e., P-glycoprotein (P-gp) efflux function on the basis of fluorescence.^{778-781,787} The Cayman's Multi-Drug Resistance Assay Kit (Calcein-AM) was used as described in the research methodology (Chapter 2). The rate of calcein accumulation in human MDR1-expressing cells is significantly lower relative to control cells, while various drug-resistance reversing agents (verapamil, vinblastine, oligomycin, cyclosporin A and MDR1-specific monoclonal antibodies) greatly increase calcein trapping only in the MDR1-expressing cells.⁷⁷⁹

In this study, cyclosporin A, a competitive inhibitor, and verapamil, a non-competitive inhibitor of P-gp, were included as positive controls. Representative results of calcein retention in SK-N-BE(2) cells treated for 24h with aspirin, bacitracin, castanospermine, brefeldin A and thapsigargin, relative to control, are summarized in Figure 3.15A to E.

In all cases (except for cyclosporin A, Figure 3.15D) cyclosporine A and verapamil (inhibitors of P-gp-mediated efflux function) significantly ($p < 0.05$) increased calcein retention in SK-N-BE(2) cells whereas the concentrations of aspirin, bacitracin, castanospermine, brefeldin A and thapsigargin tested had a reducing effect (Figure 3.15A-C and E).



The extent of calcein-AM retention was determined by fluorescence measurements using a plate reader. Data were transformed to percentages, with the control set as 100%. Values are means \pm 95% CI (n=4) of a representative experiment. One-way ANOVA followed by Tukey's multiple comparisons test yielded P values for comparison of treated vs control SK-N-BE(2) cells (vehicle-treated controls). Cyclosporin A, a competitive inhibitor, and verapamil, a non-competitive inhibitor of P-gp, were included as positive controls.

Figure 3.15: Effects of test compounds on P-glycoprotein function in SK-N-BE(2) neuroblastoma cells

3.9 Summary

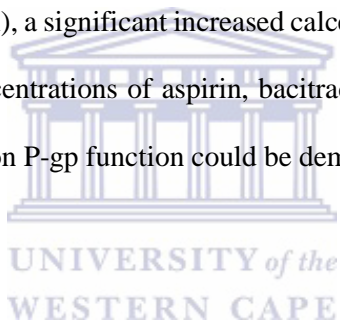
This chapter presented the results obtained in this study. The SK-N-BE(2) is a model for a continuous neuroblastoma (NB) cell line. In culture, these cells grow as epithelial-like monolayers exhibiting cellular elongation processes. The cells form dense aggregates and when overgrown or post-confluent, they lift from the culture substratum to form floating clusters. SK-N-BE(2) NB cells, like most drug-resistant NB cell lines, overexpress the multidrug transporter, P-glycoprotein (P-gp) and various other members of the ATP-Binding Cassette (ABC) family of drug efflux pumps. Growth curve analysis of SK-N-BE(2) cells indicated that the cells conformed to canonical log-plateau-decline cell proliferation kinetics.

CCK-8 cell proliferation \log_{10} incremental dose-response assays for the N-glycosylation inhibitors and pharmacologic modulators of ERS and UPR signalling, were used to estimate their time-dependent (24, 48, 72h) half maximal inhibitory concentrations (IC_{50}) or potencies according to the non-linear four-parameter logistic regression model for variable Hill slopes. These IC_{50} values were used to intuitively select concentrations of the test compounds for their further analyses of Annexin-V Cy3 apoptosis and measurement of P-gp-mediated efflux function in SK-N-BE(2) NB cells.

The Apotox-Glo™ Triplex (cell cytotoxicity, viability and apoptosis) assays showed that aspirin produced significant cytotoxicity fluorescence at concentrations of 0.1 and 10 mM, but at all concentrations of aspirin tested no effect on viability fluorescence was observed whereas at 1 and 10 mM it reduced apoptosis luminescence, as compared with control. Bacitracin at concentrations of 10 and 100 mM decreased cytotoxicity fluorescence with a commensurate increase in viability at these concentrations, however, at 10 mM, bacitracin significantly increased caspase-dependent apoptosis whereas, at 100 mM, apoptosis was significantly reduced in SK-N-BE(2) cells relative to untreated controls.

Generally, castanospermine in the \log_{10} concentration range of 0.01 to 1000 μM did not produce any cytotoxic effects, but cell viability and apoptosis was increased at 1000 μM . Brefeldin A at 2, 20 and 200 μM exerted significant cytotoxic effects with a parallel increase in apoptosis luminescence at the entire \log_{10} concentration range of 0.002 to 200 μM , but viability remained essentially unchanged. Thapsigargin only significantly increased cytotoxicity and viability fluorescence at 1000 nM, but apoptosis luminescence was increased at 1, 10 and 1000 nM, while at 100 nM a decreased apoptosis was observed.

Annexin-V Cy3 apoptosis assays revealed mostly live cells and occasional apoptotic cells for all the test compounds. In all cases, except of course for cyclosporin A and verapamil (inhibitors of P-gp-mediated efflux function), a significant increased calcein retention in SK-N-BE(2) cells were observed, but for all the concentrations of aspirin, bacitracin, castanospermine, brefeldin A and thapsigargin tested no effect on P-gp function could be demonstrated.



CHAPTER 4

CONCLUSIONS AND FUTURE PERSPECTIVES

4.1 Introduction

Neuroblastoma (NB) accounts for about 10% of all childhood cancers and is responsible for close to 15% of cancer-related deaths in the paediatric population.^{14,16} Newly diagnosed children invariably present with metastatic disease or aggressive multidrug resistant (MDR) tumours and are therefore at high risk of treatment failure, associated with poor survival outcomes and high mortality rates. In paediatric malignancies, including NB, various cancer stem cell (CSC) phenotypes overexpress multidrug-resistance (MDR) or ABC transporters^{293,294,319-323} which have attracted interest in their therapeutic targeting to overcome chemoresistance.^{7,144,294,315,324,325} Recent evidence suggests that alterations in glycolipids and protein glycosylation pathways are associated with NB biological behaviour.⁵²⁷ Current efforts are increasingly being directed at defining the molecular features of the tumour microenvironment (TME), particularly with regard to changes in the expression of glycan-related genes, as well as enzymes such as glycosyltransferases and glycosidases.⁵⁴⁶ The role of protein glycosylation in cancers⁷⁵⁹ and its potential therapeutic applications in NB have also become focal points in recent years.⁴⁹⁴

4.2 Research Hypothesis and Objectives of the Study

In this study, the N-glycosylation inhibitors and ERS inducers—aspirin, bacitracin, castanospermine, brefeldin A and thapsigargin—were evaluated for their effects on SK-N-BE(2) NB cell growth, viability, apoptosis and P-gp function. It was hypothesized that N-glycosylation inhibitors and ERS inducers will alter the expression of SK-N-BE(2) cancer cell hallmarks evaluated, namely, cell survival (proliferation, viability and apoptosis) and P-gp-

mediated drug efflux function. Based on the results of this study it can be accepted that, indeed, these investigative compounds altered the proliferation, viability, apoptosis and P-gp function in SK-N-BE(2) NB cells, albeit somewhat unconvincingly because of the conflicting and duality of responses observed.

4.3 Context and Significance of the Study

4.3.1 P-Glycoprotein, Endoplasmic Reticulum Stress and Glycosylation

The expression of the transmembrane multidrug resistance (MDR)-associated drug efflux pump, P-glycoprotein (P-gp), by cancer cells is one of the principal reasons for failure of cancer chemotherapy.³¹⁴ Elevated expression of the ABC transporter genes confers both clinical and *in vitro* drug resistance and correlates with poor prognosis in NB.^{327,788} MDR is mediated by the enhanced efflux of drugs (and thus reduced intracellular retention and cytotoxicity) by transmembrane ABC transporters, of which P-gp is a member.³¹¹ The SK-N-BE(2) NB cell line used in this study is known to overexpress P-gp.^{785,786} P-gp is modified post-translationally by N-glycosylation which is thought to play a significant role in its maturation, location and activity as a drug transporter. Accordingly, inhibitors of glycosylation have been shown to perturb P-gp in various ways.⁵⁹³

The endoplasmic reticulum (ER) regulates the synthesis, folding and aggregation of intracellular proteins.^{637,643,789,790} Relentless aberrant protein glycosylation within the ER may induce ER stress (ERS) and dysregulation of signal transduction pathways coupled to the unfolded protein response (UPR) which culminate in apoptosis or programmed cell death.⁶⁸³ The wide use of glycosylation inhibitors and oligosaccharide-processing reinforces the significance of glycosylation patterns of cell surface glycoproteins and glycolipids in the malignant phenotype.^{593,637} Thus, efforts devoted to pharmacological targeting of the ERS and UPR are intensifying at a startling rate.^{660,661,663,675,692-697} In the present study, various glycosylation inhibitors and ERS inducers (aspirin, bacitracin, castanospermine, brefeldin A and thapsigargin) were tested for their efficacy to induce apoptosis in the SK-N-BE(2) NB cell

line and inhibition of P-gp-mediated drug transport function and drug resistance. Overall, the significance of the study relates to the tenet that, transcriptionally, MYCN (a hallmark of NB) directly regulates the ABC transporters,³²⁸ and their overexpression correlates with poor prognosis.^{327,329} Most aggressive NBs exhibit MDR,³¹¹ attributable to p53 mutations and/or a loss of function induced during chemotherapy,³³¹ which further worsens the probability of relapse.^{333,334}

4.3.2 Aspirin

In this study, aspirin (acetylsalicylic acid) produced cytotoxicity towards SK-N-BE(2) cells, but viability was not affected. Cytotoxicity does not necessarily imply cell killing. Also, aspirin had no effect on cell apoptosis at low concentrations, but at higher concentrations it decreased apoptosis induction. Aspirin, at the concentrations studied, did not interfere with P-gp function as measured by the calcein retention assay. These findings may be significant from the perspective that the Wnt/beta-catenin pathway is a key modulator of aspirin-induced apoptosis in mesenchymal stem cells (MSCs) via regulation of mitochondrial/caspase-3 function.⁷⁹¹ However, it has been shown that aspirin has dual effects (it can either enhance or decrease) on cyclo-oxygenase-2 (COX-2) expression mediated via the Wnt/beta-catenin pathway. Remarkably, low-dose aspirin has been found to impede inflammatory tumour progression *in vivo* in a transgenic mouse model of neuroblastoma.⁷⁹²

Aspirin affects the activity and expression of several molecules implicated in ERS, triggering a variety of cellular processes, including transcriptional activation of ERS responsive genes.⁷²⁵ Aspirin was also found to induce *in vitro* P-gp expression and to suppress proliferation (contrasting effects) in LNCaP prostate cancer cells.⁷⁹³ Aspirin also enhances *MDR1* expression in human Molt-4 T lymphoma cells.⁷⁹⁴ Recently, the post-diagnosis use of aspirin in patients with gastrointestinal tract cancer was demonstrated to correlate with increased survival, hence lending support to the hypothesis that the anticancer effects of aspirin are not tumour-site specific and may be modulated through the TME.⁷⁹⁵ Interestingly, among women

living at least 1 year after a breast cancer diagnosis, aspirin use was associated with a decreased risk of distant recurrence and breast cancer death.⁷⁹⁶ Thus, regular aspirin use after a cancer diagnosis may improve survival outcomes in the adjuvant setting where the risk:benefit ratio will tip the scales away from the known adverse effects of this non-steroidal anti-inflammatory drug.^{726,797}

4.3.3 Bacitracin

Bacitracin is a peptide antibiotic widely used as an inhibitor of protein disulfide isomerase (PDI) to validate the role of the protein-folding catalyst in a variety of molecular pathways.⁶²⁸ Bacitracin interferes with ER function and enhances ER stress-mediated apoptosis in melanoma cells via up-regulation of ER chaperones.⁷⁹⁸ In this study, bacitracin, was shown to exert concentration-dependent effects on apoptosis in SK-N-BE(2) cells, i.e., at low concentrations it increased caspase-dependent apoptosis, but at higher concentrations it reduced apoptosis. Such duality of effects is difficult to explain in the absence of mechanistic studies, especially since it was observed that bacitracin also decreased cytotoxicity commensurate with increased viability, but had no impact on P-gp efflux function. Recently, bacitracin was reported to decrease phosphorylated focal adhesion kinase (p-FAK) and secreted matrix metalloproteinase-2 (MMP-2), which are downstream of integrin and play a major role in cell migration and invasion, and thus offering a rational therapeutic strategy for targeting malignant glioblastoma.⁶²⁷ Bacitracin may also have purported application in delineating the interrelationships of ulcerative colitis, expression of ABC drug transporters, inflammation and the pathogenesis of colorectal cancer.⁷⁹⁹

4.3.4 Castanospermine

Castanospermine is a plant alkaloid and natural inhibitor of glycosidases and thus blocks elongation of glycan chains.⁸⁰⁰ Cells exposed to castanospermine express altered levels of cell surface glycoprotein receptors.^{593,801,802} Results obtained in this study showed that castanospermine at concentrations tested produced no cytotoxic effects, but at a high

concentration (1000 μM) resulted in contrasting effects, viz, increased viability and apoptosis, but no effect on calcein retention. Previous studies have shown that castanospermine effectively altered endothelial cell glycosylation, blocked angiogenesis, and reduced tumour growth.⁸⁰³

4.3.5 Brefeldin A

Brefeldin A is an inhibitor of the secretory protein traffic pathway, i.e., it blocks translocation of proteins from the ER to the Golgi complex, that causes accumulation of secretory proteins in the ER, and hence ERS.^{665,675,685,804} Brefeldin A is a regulator of the ER resident chaperone GRP78 (a master regulator of ERS and the UPR) gene expression in mammalian cells.^{609,805} In this study, brefeldin A induced cytotoxicity and apoptosis in SK-N-BE(2) cells, but no inhibition of P-gp function was evident in the concentration range tested. Brefeldin A is regarded as an inhibitor of P-gp.⁷¹⁰ Induction of ER stress and inhibition of ARF activity are central to the proof of concept of the anticancer potential of brefeldin A.⁶⁷⁵ In growth inhibition assays using human breast carcinoma MDA-MB-435 cells, brefeldin A showed synergism in combination with taxol and tiazofurin.⁸⁰⁶ A water-soluble pro-drug analogue of brefeldin, called breflate, has been developed to facilitate parenteral administration of brefeldin as an investigational antineoplastic in clinical trials.⁸⁰⁷ However, clarification of the complex signalling pathways and associated ERS that stem from the Golgi complex in response to brefeldin A is needed.^{668,675,686}

4.3.6 Thapsigargin

Thapsigargin is a high affinity and widely-used inhibitor of ER Ca^{2+} transport ATPases.⁸⁰⁸ Thapsigargin is a potent inducer of GRP78 expression and ERS and activator of the UPR through non-competitive inhibition of SERCA (sarcoplasmic/endoplasmic reticulum calcium ATPase) pump.^{665,675} In this study, thapsigargin increased cytotoxicity and apoptosis in SK-N-BE(2) cells, but had no effect on P-gp function as measured by the calcein retention assay. Recent studies have shown that thapsigargin and other well-known ER homeostasis modifiers

induce ERS and epithelial-mesenchymal transition (EMT) in lung adenocarcinoma cells.⁸⁰⁹ Thapsigargin has been demonstrated to induce apoptosis when autophagy (both processes are regulated by the ROS-dependent pathway) is inhibited in hepatoma (HepG2) cells.⁷⁴⁴ Thapsigargin also sensitizes human oesophageal cancer to TRAIL-induced apoptosis via AMPK activation.⁶⁹⁸ It has been reported that inhibition of caspase activity significantly reduced cell death in both tunicamycin- or thapsigargin-treated cells and that caspases are crucial mediators in inducing cell death in response to ERS.⁸¹⁰

This well-documented efficacy of ERS aggravators (ERSAs) such as thapsigargin, tunicamycin and nelfinavir offers a cogent targeted cancer chemotherapeutic approach.⁶⁷⁵ A novel thapsigargin-based targeted prodrug, mipsagargin, has shown promise in a phase I clinical trial in patients with refractory, advanced or metastatic solid tumours.⁸¹¹ Intracellular Ca^{2+} is a key signalling pathway modulator of NB pathophysiology and thus presents a conceivable drug target for the treatment of NB, particularly using thapsigargin to monitor NB events such as differentiation, proliferation, drug resistance, apoptosis and autophagy.^{680,686,732,740,812} Related work drawing attention to the cross-talk between autophagy and ER homeostasis illustrated that induction of ERS by thapsigargin involves impairment of autophagosome-lysosome fusion.^{732,813,814}

4.4 Limitations of the Study

This study is limited to SK-N-BE(2) NB cells. Considerable effort was made to study a range of concentrations of the investigative compounds in all assays, but funding constraints hindered such objectives.

4.5 Conclusions and Future Outlook

Neuroblastoma (NB) is the most frequent type of solid extra-cranial tumour in children associated with approximately 15% of paediatric cancer-related deaths.^{261,358,815} NB is predominantly heterogeneous with biological and clinical behaviour fluctuating between

complete spontaneous regression and aggressive clinical multidrug resistance, suggesting that apart from genetic events, the tumour microenvironment (TME) strongly influences these characteristics of NB.^{64,147} *MYCN* oncogene expression is the most significant cancer signature associated with aggressive or HR-NB and poor survival outcome.^{184,816} Pharmacologic targeting of *MYCN* is not straightforward as this NB oncogenic driver is not very amenable to direct preclinical and clinical targeting and efforts should thus be centred on indirectly targeting *MYCN*.¹⁹⁸ Therefore, the development of innovative rational targeted therapies based on druggable pathways specifically activated in NB with *MYCN* amplification should be encouraged to diversify more efficacious treatment modalities.

MYCN has pro-growth and pro-survival functions, but can switch to an apoptosis initiating mode via p53.⁸¹⁷⁻⁸¹⁹ Thus, the paradoxical apoptosis-promoting function of *MYCN* amplification in NB could be a valuable line of attack in the high-risk, *MYCN*-amplified subset of NB.¹⁹⁸ Moreover, aggressive NBs express MDR,³¹¹ ascribed to p53 mutations and/or a loss of p53 function acquired during chemotherapy,^{331,332} which exacerbates the odds for relapse and thus treatment efficacy.^{331,333,334} The MDR transporter, P-gp, has been shown to be one of the most strongly upregulated genes associated with acquired drug resistance and NB treatment failure.^{527,589,590} This, together with the emerging themes of NB glycobiology (glycomics),⁵²⁷ glycosylation in cancer,^{544,593,820-822} and challenges of 40% relapse among HR-NB patients associated with glycan-based immunotherapy following standard therapy,^{153,494} underscores the need for targeting P-gp.^{311,313,315,326,823,824} {Silva, 2015 #2890; Garg, 2015 #6059; Wang, 2014 #5845} Also, ER stress and the UPR pathways have consistently been regarded as promising targets for developing drugs for several cancers, which may be further explored.^{661,695,825-829}

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
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Author: Kristopher R. Bosse, John M. Maris

Publication: Cancer

Publisher: John Wiley and Sons

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Author: Pablo H. Strobl-Mazzulla, Marianne E. Bronner

Publication: Seminars in Cancer Biology

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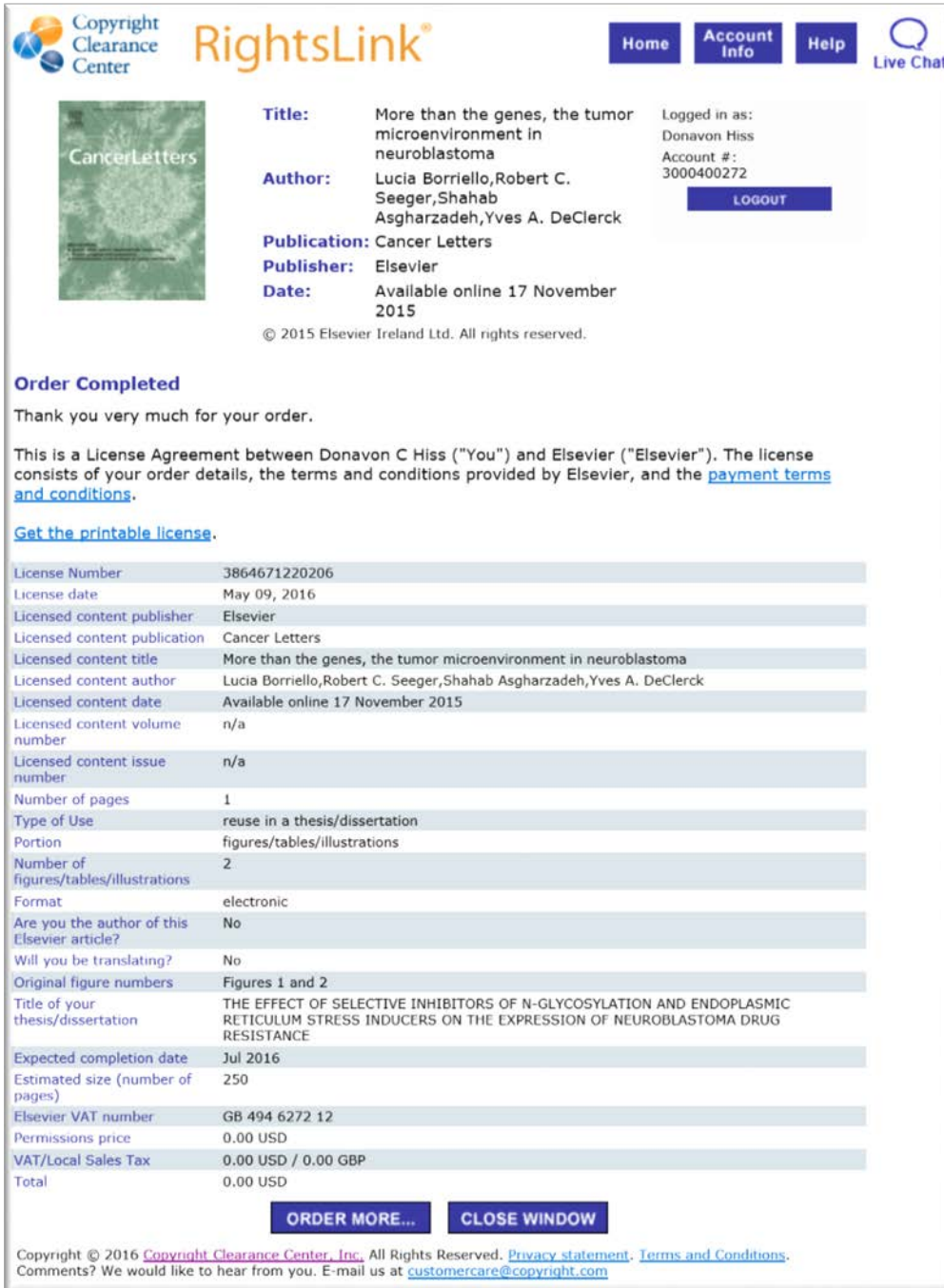
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Title: More than the genes, the tumor microenvironment in neuroblastoma

Author: Lucia Borriello, Robert C. Seeger, Shahab Asgharzadeh, Yves A. DeClerck

Publication: Cancer Letters

Publisher: Elsevier

Date: Available online 17 November 2015

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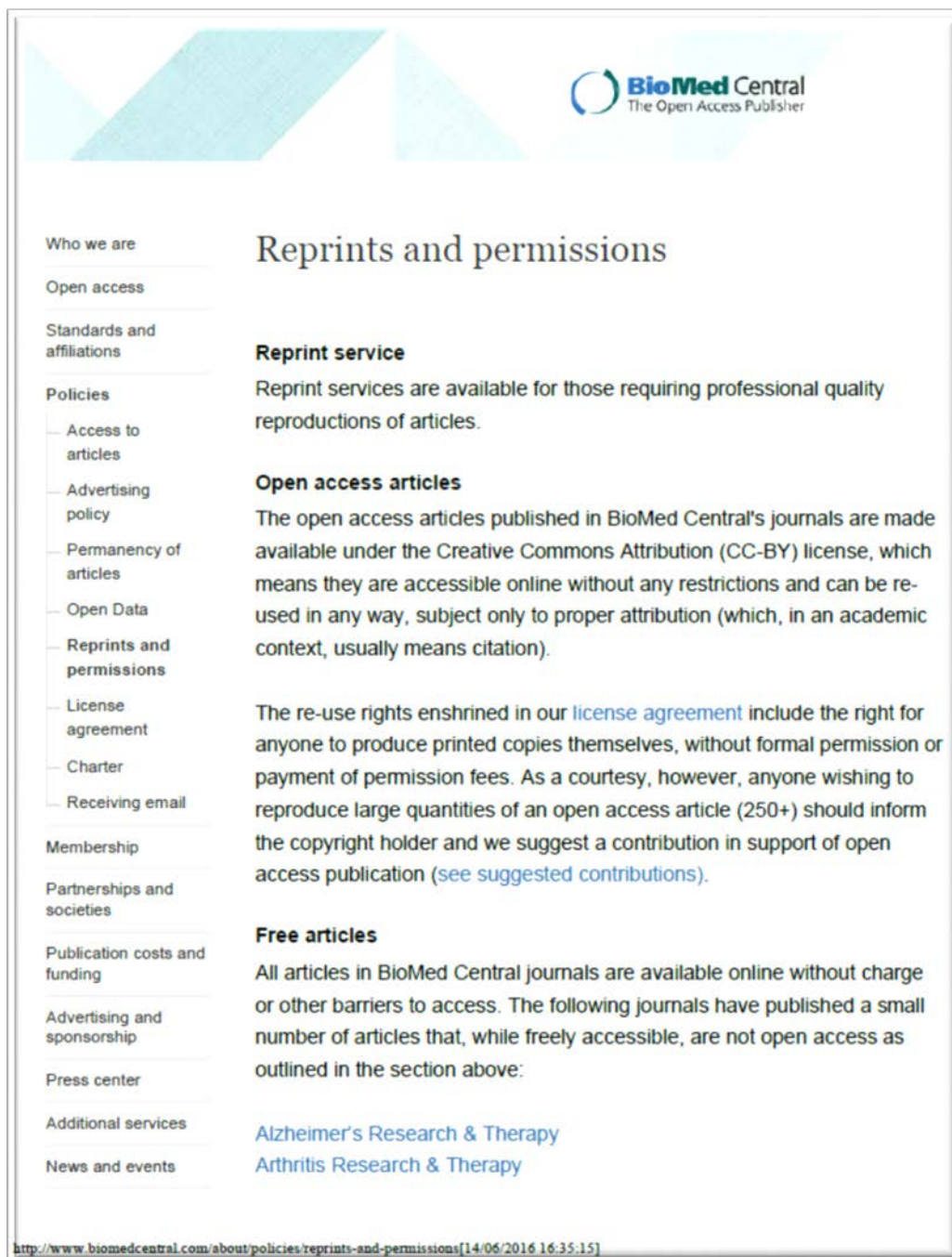
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Title: Multi-drug resistance in cancer chemotherapeutics: Mechanisms and lab approaches

Author: Qiong Wu,Zhiping Yang,Yongzhan Nie,Yongquan Shi,Daiming Fan

Publication: Cancer Letters

Publisher: Elsevier

Date: 1 June 2014

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Title: Retinoid therapy of high-risk neuroblastoma

Author: C.Patrick Reynolds,Katherine K. Matthay,Judith G. Villablanca,Barry J. Maurer

Publication: Cancer Letters

Publisher: Elsevier

Date: 18 July 2003

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Title: mTOR and cancer: many loops in one pathway

Author: Alejo Efeyan, David M Sabatini

Publication: [Current Opinion in Cell Biology](#)

Publisher: Elsevier

Date: April 2010

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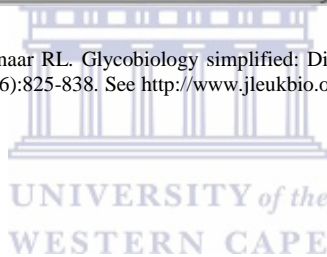
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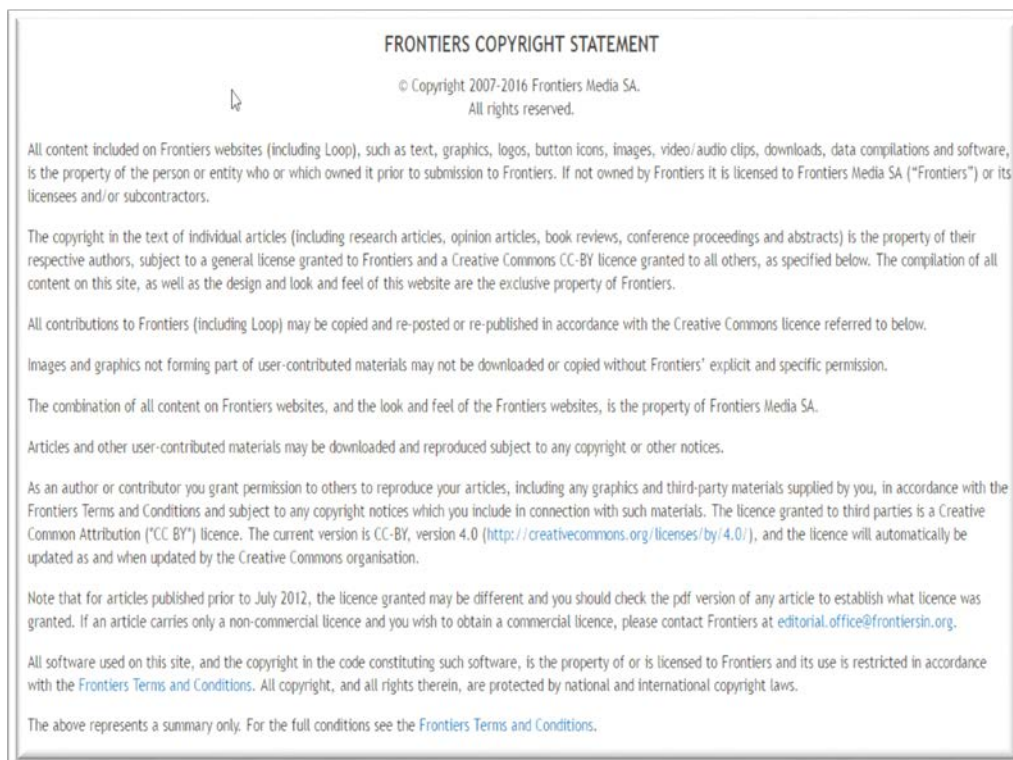



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