Aus dem Fachbereich Medizin der Johann Wolfgang Goethe-Universität Frankfurt am Main

betreut am

Georg-Speyer-Haus

Institut für Tumorbiologie und Experimentelle Therapie

Direktor: Prof. Dr. Florian Greten

Exploration of Nilotinib enhancing the anti-leukaemic Effects of Gas6 Deletion in Ph⁺ B-ALL by inducing Immunogenic Cell Death

Dissertation
zur Erlangung des Doktorgrades der Medizin
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vorgelegt von
Carolin Wachtel

aus Heppenheim (Bergstraße)

Frankfurt am Main, 2020

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In Dank und Liebe meinen Eltern,

Ute und Ralf Wachtel,

gewidmet

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List of Abbreviations

ALL	Acute lymphoblastic leukaemia
ANXA1	Annexin A1
AnxV	Annexin V
ATP	Adenosine triphosphate
B-ALL	B-cell acute lymphoblastic leukaemia
BMDM	Bone marrow-derived macrophages
CAFs	Cancer-associated fibroblasts
CALR	Calreticulin
CD	Cluster of differentiation
CDKN2A	Cyclin-dependent kinase inhibitor 2A
CXCL	CXC motif chemokine ligand
DAMPs	Damage-associated molecular patterns
DMSO	Dimethyl sulfoxide
DPBS	Dulbecco's Phosphate Buffered Saline
ECM	Extracellular matrix
ELISA	Enzyme-linked immunosorbent assay
ER	Endoplasmic reticulum
FACS	Fluorescence-activated cell sorting
FBS	Foetal Bovine Serum
FS	Forward scatter
GFP	Green fluorescent protein
Gas6	Growth-arrest-specific gene-6
Gas6KO	Gas6 knockout
HD-TKI	High-dose TKI
HE	Haematoxylin-Eosin
HMGB1	High-mobility-group-protein B1
IC ₅₀	Half maximal inhibitory concentration
ICD	Immunogenic cell death
IHC	Immunohistochemistry

IL	Interleukin
INF	Interferon
IRES	Internal ribosomal entry site
LC3	Light chain 3
LPS	Lipopolysaccharide
M-CSF	Macrophage colony-stimulating factor
MDSCs	Myeloid-derived suppressor cells
MHC	Major histocompatibility complex
M-MDSCs	Monocyte-related myeloid-derived suppressor cells
MTX	Mitoxantrone
NCCD	Nomenclature Committee on Cell Death
NK	Natural killer
PAGE	Polyacrylamide gel electrophoresis
Pi	Propidium iodide
PRRs	Pattern recognition receptors
PVDF	Polyvinylidene difluoride
RIPK	Receptor-interacting protein kinases
rL/L-reagent	Recombinant Luciferin/Luciferase-reagent
RLU	Relative light units
rpm	Revolutions per minute
RTKs	Receptor tyrosine kinases
SD	Standard deviation
SDS	Sodium dodecyl sulphate
SDS-PAGE	Sodium dodecyl sulphate polyacrylamide gel electrophoresis
SS	Side scatter
TAM	Tyro3, Axl, Mer (collective term)
TAMs	Tumour-associated macrophages
TGF	Transforming growth factor
TIICs	Tumour-infiltrating immune cells
TIME	Tumour immune microenvironment
TKI	Tyrosine kinase inhibitor
TLR	Toll-like receptor
TME	Tumour microenvironment

TNF	Tumour necrosis factor
Tregs	Regulatory T cells
wt	Wild-type

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1 Abstract

Cancer is the major cause of death besides cardiovascular disease. Leukaemia represents the most prevalent malignancy in children with a frequency of 30 % and is one of the ten leading types of cancer in adults. Philadelphia Chromosome-positive B-ALL (Ph+ B-ALL) is driven by the cytogenetic aberration of the reciprocal chromosomal translocation t(9;22)(q34;q11) leading to the formation of the Philadelphia chromosome with a BCR-ABL1 fusion gene. This fusion gene encodes a BCR-ABL1 oncoprotein which is characterized by a constitutively enhanced tyrosine kinase activity promoting amplified proliferation, differentiation arrest and resistance to cell death. Ph+ B-ALL is considered the most aggressive ALL subtype with a long-term survival rate in the range of only 30 % despite intensive standard of care including chemotherapy in combination with a tyrosine kinase inhibitor (TKI) followed by allogeneic stem cell transplantation after remission for clinically fit patients.

The efficacy of chemotherapy has long been mainly attributed to tumour cell toxicity while immune modulating effects have been overlooked, especially in light of known immunosuppressive properties. Accumulative evidence, however, emphasizes the ability of chemotherapeutic agents, including TKIs, to normalise or re-educate a dysfunctional tumour microenvironment (TME) resulting in enhanced anti-tumour immunity. One of the underlying mechanisms of immune modulation is the induction of immunogenic cell death (ICD). ICD is an antitumour agent-induced cell death modality determined by the capacity to convert cancer cells into anti-cancer vaccines. The induction of ICD relies on the release of damage-associated molecular patterns (DAMPs) from dying tumour cells succumbing to ICD. Translocation of CALR to the cell surface, extracellular secretion of ATP and release of HMGB1 from the nucleus are key hallmarks of ICD that mediate anti-tumour immunity upon binding to antigen presenting cells resulting in a tumour antigen-specific immune response. Besides these molecular determinants, ICD is functionally defined by the inhibition of tumour growth in a vaccination assay in which mice are injected with tumour cells exposed to the potential ICD inducer in-vitro and then re-challenged with live tumour cells of the same cancer type. Both molecular and functional criteria determine the gold

standard approach to assess ICD. By increasing the immunogenicity of cancer cells, ICD contributes to the restoration of immunosurveillance as an essential feature of tumour rejection, which is clinically reflected by improved therapeutic efficacy and disease outcome in patients. Therefore, identifying novel ICD inducers is an objective of interest in the context of cancer therapy.

In respect of these considerations, the aim addressed in the present work is the examination of the second-generation TKI Nilotinib for the ability to induce ICD. The thesis is set in the context of the group's research on the role of Gas6/TAM signalling within the TME regarding the pathogenesis of acute leukaemia. In in-vivo experiments of our research group it has been consistently observed that the use of Nilotinib enhances the anti-leukaemic immunity mediated by a deletion of Gas6. Against the background of increasing importance of chemotherapeutic agents as potent modulators of a dysregulated TME, it was hypothesized that Nilotinib may synergize with a Gas6-deficient environment by inducing ICD in Ph+ B-ALL cells.

In growth inhibition and Annexin V/Propidium iodide cell death assays Nilotinib was shown to induce cell death in concentration-dependent manner that occurs bimodally in terms of cell death modality ranging between apoptosis and necrosis. By ICD marker analysis, comprising flow-cytometric detection of CALR exposure, chemoluminescence-based ATP measurement and immunoblotting for HMGB1, it was found that Nilotinib-induced cell death is not accompanied by CALR exposure and ATP secretion, but is associated with the release of HMGB1. In macrophages co-culture experiments with Nilotinib-treated leukaemic cells, no relevant shift in terms of macrophages activation and polarisation was observed in either a juxtacrine or paracrine setup. In consistency with the results obtained in the in-vitro experiments, Nilotinib was not potent to elicit a protective immune response in mice within a vaccination assay.

Conclusively, Nilotinib was identified to not qualify as bona fide ICD inducer. The role of Nilotinib-induced cell death and HMGB1 release are proposed as objective for further investigation concerning the synergistic interplay between Nilotinib and a Gas6-deficient environment. Efforts addressing exploration and optimisation of the immunological potential of chemotherapeutic agents are a promising approach aimed at providing cancer patients with the best possible treatment in future.

2 Zusammenfassung

Krebs stellt neben Herz-Kreislauf-Erkrankungen eine der Haupttodesursachen da. Leukämie ist mit einer Häufigkeit von 30 % die häufigste bösartige Erkrankung bei Kindern und gehört zu den zehn führenden Krebsarten bei Erwachsenen. Die Philadelphia-Chromosom-positive B-ALL (Ph+ B-ALL) wird durch die zytogenetische Aberration der reziproken chromosomalen Translokation t(9;22)(q34;q11) hervorgerufen, die zur Bildung des Philadelphia-Chromosoms mit einem BCR-ABL1-Fusionsgen führt. Dieses Fusionsgen kodiert ein BCR-ABL1-Onkoprotein, das durch eine konstitutiv erhöhte Tyrosinkinase-Aktivität zu verstärkter Proliferation, Differenzierungsblock und Resistenz gegenüber Zelltod führt. Ph+ B-ALL gilt als die aggressivste Form der ALL mit einer Langzeitüberlebensrate von lediglich 30 % trotz intensiver Chemotherapie, die in Kombination mit einem Tyrosinkinase Inhibitor (TKI) eingesetzt wird und welcher sich bei klinisch geeigneten Patienten nach Remission eine allogene Stammzelltransplantation anschließt.

Die Wirksamkeit von Chemotherapien wurde lange Zeit hauptsächlich der Toxizität gegenüber Tumorzellen zugeschreiben, wohingegen immunmodulierende Effekte insbesondere angesichts der bekannten immunsuppressiven Eigenschaften weitgehend unberücksichtigt blieben. Zunehmend zeigt sich jedoch, dass Chemotherapeutika, einschließlich TKI, eine dysfunktionale Tumormikroumgebung (TME) normalisieren oder umstrukturieren und somit eine verstärkte Antitumorimmunität bewirken können. Einer der zugrunde liegenden Mechanismen der Immunmodulation stellt die Induktion von immunogenem Zelltod (ICD) da. ICD ist eine Form des Zelltods, die dadurch charakterisiert ist, dass innerhalb der Tumortherapie abgetötete Krebszellen in Vakzinen gegen Krebs transformiert werden. Durch die Freisetzung von Damage Associated Molecular Patterns (DAMPs) aus sterbenden Tumorzellen wird ICD induziert, wobei die Translokation von CALR zur Zelloberfläche, die extrazelluläre Sekretion von ATP und die Freisetzung von HMGB1 aus dem Zellkern die zentralen Elemente des ICD darstellen und durch Bindung an antigenpräsentierende Zellen eine tumorantigenspezifische Immunantwort hervorrufen. Neben diesen molekularen Bestimmungsgrößen wird ICD funktionell über eine

Inhibition des Tumorwachstums im Rahmen eines Impftests definiert, in welchem Mäuse lebenden Tumorzellen ausgesetzte werden, nachdem sie zuvor mit den als Vakzine fungierenden Krebszellen geimpft wurden. Zusammen bilden molekulare und funktionelle Kriterien den Goldstandard zur Beurteilung von ICD. Durch die Erhöhung der Immunogenität von Krebszellen trägt ICD zur Wiederherstellung der Immunüberwachung bei, die ein wesentliches Merkmal der Tumorabwehr ist und sich klinisch in einem verbesserten Behandlungsergebnis äußert. Daher stellt die Identifizierung von neuen ICD-Induktoren ein Ziel von Interesse im Hinblick auf Behandlungsstrategien von Krebs dar.

Unter Berücksichtigung dieser Aspekte, ist die Untersuchung des TKI Nilotinib hinsichtlich seiner Fähigkeit, ICD zu induzieren, das Anliegen der vorliegenden Arbeit. Diese steht im Kontext der Forschungsarbeiten der Arbeitsgruppe zur Rolle der Gas6/TAM-Achse innerhalb des TME bei der Entstehung von akuten Leukämien. In den in-vivo-Experimenten unserer Forschungsgruppe wurde konsistent beobachtet, dass der Einsatz von Nilotinib die durch die Deletion von Gas6 vermittelte antileukämische Immunität verstärkt. Vor dem Hintergrund der zunehmenden Bedeutung von Chemotherapeutika als potente Modulatoren eines dysregulierten TME wurde daher die Hypothese aufgestellt, dass Nilotinib mit einer Gas6-defizienten Umgebung durch die Induktion von ICD in Ph+ B-ALL-Zellen synergistisch wirken könnte.

In Wachstumshemmungs- und Annexin V/Propidiumiodid-Zelltod-Assays konnte gezeigt werden, dass Nilotinib konzentrationsabhängig Zelltod induziert, welcher in Bezug auf die Zelltodmodalität bimodal apoptotisch und nekrotisch verläuft. Mittels ICD-Markeranalyse, die den durchflusszytometrischen Nachweis der Translokation von CALR, die Chemolumineszenz basierte ATP-Messung und das Immunoblotting für HMGB1 umfasst, wurde festgestellt, dass der durch Nilotinib induzierte Zelltod nicht mit einer CALR-Translokation oder ATP-Sekretion einhergeht, aber mit der Freisetzung von HMGB1 assoziiert ist. In Makrophagen-Co-Kultur-Experimenten mit Nilotinib behandelten leukämischen Zellen wurde weder in einem juxtakrinen noch in einem parakrinen Versuchsaufbau eine relevante Veränderung bezüglich der Makrophagenaktivierung und -polarisation beobachtet. In Übereinstimmung mit den Ergebnissen der in-vitro Experimente konnte Nilotinib im Rahmen des Impftests keine schützende Immunantwort in Mäusen hervorrufen.

Zusammenfassend wurde herausgearbeitet, dass Nilotinib nicht als Induktor von ICD eingestuft werden kann. Aspekte des Nilotinib induzierten Zelltods und der HMGB1-Freisetzung werden als Gegenstand weiterer Untersuchungen bezüglich des synergistischen Zusammenspiels zwischen Nilotinib und einer Gas6-defizienten Umgebung vorgeschlagen. Bemühungen um die Erforschung und Optimierung des immunologischen Potenzials von Chemotherapeutika stellen einen vielversprechenden Ansatz dar, um Krebspatienten in Zukunft die bestmögliche Behandlung zukommen zu lassen.

3.1 Philadelphia Chromosome-positive ALL (Ph⁺ ALL)

Cancer is the second leading cause of death after cardiovascular disease.1 Regarding cancer type, leukaemia is the tenth most common cancer in adults with a frequency of 3.5 % and the most common one in children amounting 30 %.1,2 The classification of leukaemia is based on the course of the disease and the affected cell lineage so that a distinction is made between acute and chronic leukaemia on the one hand and between myeloid and lymphoblastic leukaemia on the other.³ Thereof, acute lymphoblastic leukaemia (ALL) is the most common subtype in children, representing up to 80 % of all cases and 20 % of leukaemia in adults.^{4,5} Pathogenetically, ALL is based on an uncontrolled clonal proliferation of aberrant lymphatic cells of haematopoiesis in the bone marrow with blocked cell maturation at a certain level of differentiation.^{6,5,6} The determination of this differentiation stage and the allocation to B- or T-cell lineage determined according to the antigen expression pattern immunophenotyping.⁷ Clinically, the increasing suppression of normal haematopoiesis results in progressive bone marrow insufficiency with symptoms typical of anaemia, susceptibility to infection due to leukocytopenia and tendency to bleed as a consequence of thrombocytopenia in addition to accompanying general symptoms such as fever, night sweats and fatigue.8 ALL is often characterised by various cytogenetic aberrations accounting for the heterogenicity in clinical disease course and prognosis, notably when comparing paediatric and adult ALL.9,10 One of those cytogenetic abnormalities is the chromosomal translocation t(9;22)(q34;q11) leading to the formation of the Philadelphia chromosome with a BCR-ABL1 fusion gene via the reciprocal fusion of the ABL1-tyrosine kinase gene on chromosome 9 with the BCR-gene on chromosome 22 (Figure 1). 11 This fusion gene encodes a BCR-ABL1 oncoprotein which is characterized by a constitutively enhanced tyrosine kinase activity resulting in amplified proliferation, differentiation arrest and resistance to cell death. 12,13 Philadelphia chromosome-positive ALL (Ph+ ALL) presents 25 % of adult ALL, and about 4 % of paediatric ALL cases. 14,15 The standard treatment regimen includes intensive chemotherapy in combination with a tyrosine kinase inhibitor (TKI) followed by allogeneic stem cell transplantation after remission for clinically fit patients. ¹⁶ The significant improvement in outcome compared to the very poor prognosis of Ph+ ALL prior to the introduction of TKI is due to both the fact that more patients are qualified to receive stem cell transplantation and to a reduced dependence on potentially highly toxic chemotherapy. ^{16,17} Nevertheless, despite an initial remission rate of up to 95 %, the 5-year survival rate remains 33 % to 46 % because of emerging resistance to TKI and relapses. ^{18,19} Various acquired and intrinsic mechanisms of resistance have been revealed, notably including mutations that prevent TKI binding, like T315I, or reduce TKI binding affinity by an altered protein confirmation, such as L248V. ²⁰⁻²² This has led to the development of second (Nilotinib, Dasatinib, Bosutinib) and third (Pronatinib) generation TKIs overcoming the resistance observed in the first generation TKI Imatinib. ²³

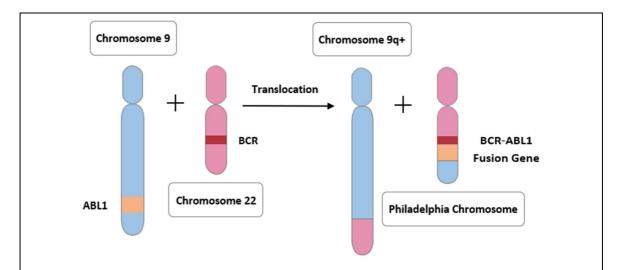


Figure 1. Philadelphia Chromosome Formation. Acute lymphoblastic leukaemia (ALL) is often driven by genetic aberrations, which are of relevance for the diagnostic classification of ALL into different subtypes. Moreover, these alterations highly influence leukaemic development, risk of treatment failure and disease outcome. Chromosomal rearrangements are one key type of alterations resulting in the expression of fusion genes. Thus, in Philadelphia Chromosome-positive ALL (Ph+ ALL) the reciprocal translocation of the ABL1-tyrosine kinase gene on chromosome 9 with the BCR-gene on chromosome 22 leads to the formation of the Philadelphia chromosome with a BCR-ABL1 fusion gene. The BCR-ABL1 oncoprotein encoded by this fusion gene displays a constitutively enhanced tyrosine kinase activity promoting amplified proliferation, differentiation arrest and resistance to cell death.

3.2 Immunotherapy and Tumour Microenvironment

However, despite great efforts and continuous improvement, the treatment of cancer with conventional therapy concepts such as tumour surgery, chemo- and radiotherapy remains suboptimal.²⁴ This gave rise to the investigation of alternative approaches such as immunotherapy, that has become an established part of cancer treatment improving prognosis and outcome in a broad field of malignant diseases.^{25,26} The functional principle of immunotherapy involves the enhancement of immune response in a general way and, in special, the promotion of the immune system in order to attack cancer cells.²⁷ More detailed, immunotherapy operates by (1) targeting immunosuppressive and tumourpromoting metabolism, (2) altering tumour microenvironment, (3) stimulating innate and adaptive anti-tumour immune response, (4) advancing detection of tumour cells that have escaped the immune system and (5) inhibiting oncogenic inflammation.²⁸ To address those objectives, several strategies are under investigation as illustrated in Figure 2, which are categorized in clinical use as antibody therapy, cellular immunotherapy, immune adjuvants therapy and combination immunotherapy. 28,29 Although immunotherapy has proven to be a beneficial option for cancer treatment especially when used in synergistic combination with conventional tumour therapy, there is growing evidence that differences in the response to cancer immunotherapy are related to the heterogeneity of the tumour microenvironment (TME).³⁰⁻³² The tumour microenvironment is composed of tumour cells, tumour-infiltrating immune cells (TIICs), cancer-associated fibroblasts (CAFs), the tumour vasculature and the extracellular matrix (ECM).33,34 Strikingly, the tumour microenvironment is characterized by the contrary ability to provoke both beneficial and adverse conditions for tumorigenesis in dependency of the interactions within the TME underlining the important role in either suppressing or enhancing anti-tumour immune response (Figure 3).35-38 A tumour-promoting, immune-suppressive tumour microenvironment exhibits a variety of mechanisms such as (1) supporting neoplastic transformation and sustaining tumour growth, (2) inducing angiogenesis, (3) enabling immortality by resistance to cell death, (4) altering and escaping the immune system of the host, (5) activating invasion and metastasis, (6) providing niches for dormancy and (7) fostering therapeutic resistance. 39-41

As a consequence, targeting the complexity of the tumour microenvironment is a promising approach in the field of cancer therapy.^{42,43}

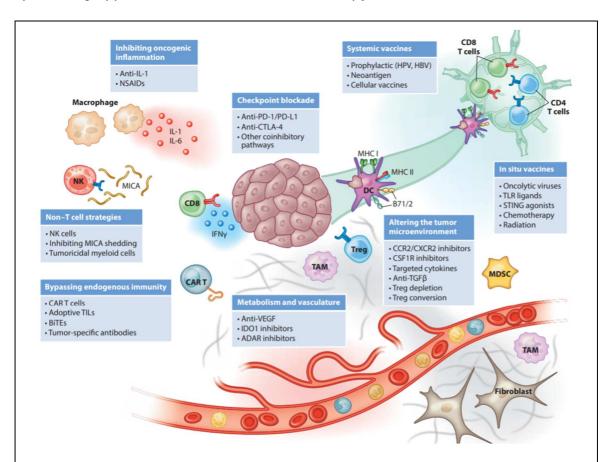


Figure 2. Investigational Strategies of Immunotherapy. Multiple investigational strategies aim to overcome immunosuppressive features and resistance to anticancer immunosurveillance that protect tumour cells from productive anti-tumour immunity. These strategies not only address the tumour microenvironment, oncogenic inflammation as well as tumour metabolism and vasculature, but also stimulate innate and adaptive anti-tumour immune response. Several therapeutic options including targeted antibodies, checkpoint inhibitors, T cell therapy, cytokines and both in situ and systemic vaccines are already in clinical use or under evaluation. From Dougan *et al.*, 2019 (28).

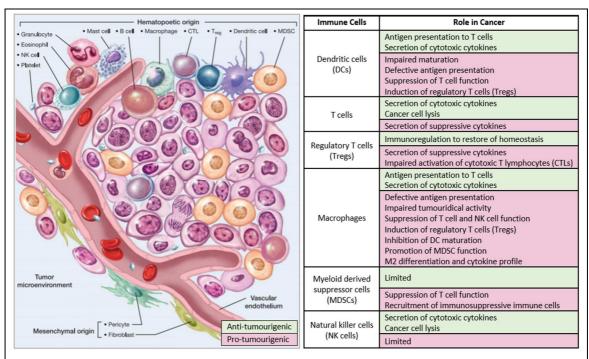


Figure 3. Plasticity of the Tumour Microenvironment. The tumour microenvironment (TME) forms a heterogeneous environment around the tumour, including tumour cells, tumour-infiltrating immune cells (TIICs), cancer-associated fibroblasts (CAFs), the tumour vasculature and the extracellular matrix (ECM). Depending on the different interactions and signalling within the TME, either a tumour-suppressive or a tumour-promoting environment is established. Taking advantage of this plasticity of the TME, cancer cells modulate the TME by various mechanisms in order to build up beneficial conditions. The adverse nature of the TME is particularly evident with respect to tumour-infiltrating immune cells, which collectively form the tumour immune microenvironment (TIME). Anti- and pro-tumourigenic functional features of different immune cells are outlined in this figure underlining the complexity of the TME. Adopted from Kerkar *et al.*, 2012 (41).

3.3 Research Background

The research work of our group is aimed to address the tumour microenvironment in the context of the pathogenesis of malignant diseases in the haematopoietic system. The work is focused on the mechanisms contributing to the establishment of an immune-suppressive environment in acute leukaemia and, in particular, on the role of the Gas6/TAM axis in anti-leukaemic immunity.

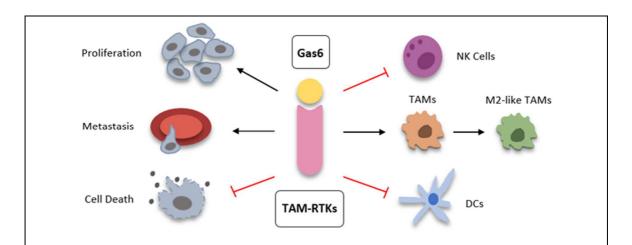


Figure 4. The Role of Gas6/TAM in Cancer Cells and the Tumour Microenvironment. Growth arrest-specific 6 (Gas6) displays its effects by binding to the receptor tyrosine kinases (RTKs) Axl, Tyro3 and Mer, that are collectively named TAM. In cancer cells Gas6/TAM signalling not only promotes cancer cell proliferation and metastasis but also inhibits cell death. Moreover, Gas6/TAM targets diverse immune cell functions resulting in the establishment of an immune-suppressive tumour microenvironment (TME). These transformations include both the inhibition of natural killer (NK) and dendritic cell function as well as the polarisation of tumour-associated macrophages (TAMs) towards a M2-like phenotype.

Gas6, also known as Growth arrest-specific 6, is a member of the vitamin K-dependent family of proteins binding with decreasing affinity to the receptor tyrosine kinases (RTKs) Axl, Tyro3 and Mer, that are collectively named TAM.44 Next to diverse processes including proliferation, differentiation, apoptosis, leucocyte migration and platelet aggregation, 45,46 Gas6 has been found to contribute to the development of several cancers, 47-52 including acute leukaemia, 53-55 in which overexpression of Gas6/TAM is associated with an unfavourable prognosis.54,55 In the context of cancer development and progression, Gas6 has been described to promote cancer cell proliferation, sustain cancer cell survival by inhibiting apoptosis and induce cancer cell migration.⁵⁶⁻⁵⁸ Moreover, Gas6/TAM signalling has been found to modulate the tumour microenvironment resulting in an immune-suppressive, inflammatory, tissue-repair phenotype.^{59,60} The Gas6/TAM axis affects the tumour microenvironment by targeting diverse components and functions, including those of immune cells.61 The effects on the tumour immune microenvironment (TIME) result from the instruction of stromal cells and macrophages to secrete Gas6 into the TME altering cellular immune response by binding to TAM receptors on immune cells. 61,62 The changes in cellular immune response comprise (1) the inhibition of tumour eradicating activity of natural killer cells (NK cells), (2) the impairment of cytokine production and antigen presentation of dendritic cells (DCs) and (3) the polarisation of tumour-associated macrophages (TAMs) towards a M2-like phenotype (Figure 4). 63-66 With regard to their immunogenic function, cytokine profile and surface marker expression, macrophages can be classified into classically activated (M1) and alternatively activated (M2) macrophages.⁶⁷ While M1 macrophages exhibit pro-inflammatory, tissue-damaging, immune-stimulating and tumour-suppressive attributes, M2 macrophages display anti-inflammatory, tissue-remodelling, immune-regulatory and tumour-promoting properties (Figure 5).68-70 However, typical M1 and M2 phenotypes represent extremes of the heterogeneity in macrophage functional stage, emphasising the dynamic plasticity in macrophage polarisation as a reaction to different microenvironmental stimuli. 70 Remarkably, tumourassociated macrophages cannot be explicitly assigned to a M1 or M2 phenotype, but they are often associated with a tumour-promoting M2-like phenotype (Supplementary Figure S1, page 88).71-74

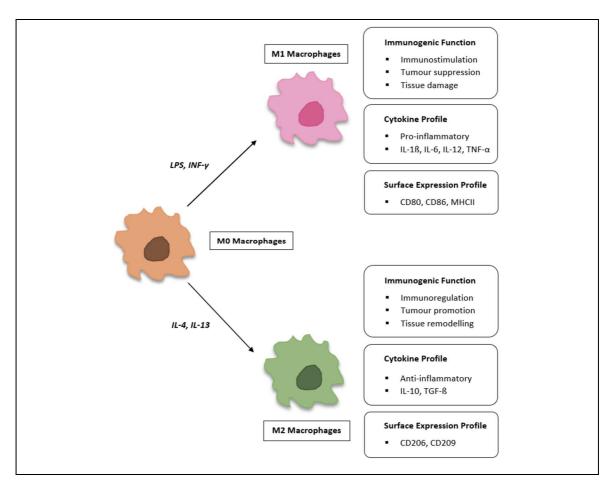


Figure 5. Macrophage Polarisation. Macrophage polarisation describes the process of macrophage differentiation to adopt a phenotype in response to different environmental stimuli. According to the binary polarisation concept, macrophages are classified into classically activated (M1) and alternatively activated (M2) macrophages regarding their immunogenic function, cytokine profile and surface marker expression. Thus, M1 macrophages exhibit immunestimulating, tumour-suppressive, tissue-damaging and pro-inflammatory attributes, whereas M2 macrophages display immune-regulatory, tumour-promoting, tissue-remodelling and anti-inflammatory properties. However, typical M1 and M2 phenotypes represent extremes of heterogeneity in macrophage functional stage, emphasising the diverse spectrum of macrophage subtypes and the dynamic plasticity of polarisation.

3.4 Immunogenic Cell Death (ICD)

Targeting Gas6/AxI axis is a complex but promising approach in cancer treatment. The findings of our group demonstrate that addressing Gas6/AxI signalling, by ablation of Gas6 in the TME or AxI in macrophages, promotes both anti-leukaemic immune response and memory resulting in leukaemic clearance and prolonged survival in pre-clinical mouse models of acute leukaemia when combined with standard of treatment.

The combination of immunotherapy with conventional chemotherapy is a conductive strategy to improve cancer treatment, but little is known about the effects of chemotherapeutic agents on the tumour microenvironment. Given the concerns about the various immunosuppressive effects, chemotherapy has not been considered an immunological modality. However, in different studies certain chemotherapeutic agents have been shown to improve anti-tumour immune response and even reverse immune tolerance by modulating the TME. The underlying mechanisms of immune modulation include (1) the eradication of immune-suppressing regulatory T cells (Tregs) and myeloid-derived suppressor cells (MDSCs), (2) the activation and maturation of DCs, (3) the promotion of TAM accumulation and reprogramming towards an immune-stimulating activity, (4) the facilitation of tumour infiltration by NK cells and cytotoxic T lymphocytes (CTLs), (5) the upregulation of class I major histocompatibility complex (MHCI) molecules on cancer cells and (6) the induction of immunogenic cell death (ICD). Tr, T9,80

Immunosurveillance is an essential feature of cancer rejection.⁸¹ Resistance to host anticancer immunosurveillance and anti-tumour immunity is acquired through several mechanisms, involving exhibition of low immunogenicity, induction of immunotolerance or active immunosuppression and evasion of immune cell-mediated lysis.^{82,83} To reinforce immunosurveillance, immunogenicity of dying or dead cancer cells can be increased by immunogenic cell death.⁸⁴

ICD is a cell death modality determined by the capacity to convert dying or dead cancer cells into vaccines leading to an antigen-specific anti-cancer immune response.85 So far, a variety of agents have been identified as potent inducers of ICD, including chemotherapeutics, radiotherapy, biological therapeutics and physio-chemical or mechanical stress-based therapies.86 Molecularly, ICD is characterised by spatiotemporal release of damage-associated molecular patterns (DAMPs) via endoplasmic reticulum (ER) stress triggering immunestimulating effects upon binding to pattern recognition receptors (PRRs) expressed by immune cells.87 Thus, DAMPs act as danger signals mediating the immunogenicity and anti-cancer vaccine effect of ICD.88,89 With regard to the mechanism of ICD induction, ICD is classified in type I and type II ICD.90 Type I ICD is induced through collateral ER stress effect, indirectly evoking ICDassociated danger signalling, whereas, in contrast, type II ICD induction selectively targets the ER, promoting focused ER stress effect. 90 Next to diverse DAMPs contributing to the immunogenicity of cell death in a limited amount, translocation of ER chaperon calreticulin (CALR) to the cell surface, extracellular secretion of adenosine triphosphate (ATP) and release of endogenous Toll-like receptor (TLR) agonist high-mobility-group-protein B1 (HMGB1) from the nucleus are attributed a key role.91-94 CALR, ATP and HMGB1 released by dying tumour cells succumbing to ICD bind to CD91, P2RX7 and TLR4, respectively, on macrophages and dendritic cells to mediate anti-tumour immune effects.95 These effects comprise (1) the engulfment and presentation of tumour antigens operated by CALR as eat-me signal, (2) the recruitment and maturation of antigen presenting cells facilitated by ATP as find-me signal and (3) the release of proinflammatory cytokines as well as T cell priming stimulated by HMGB1, finally resulting in (4) the eradication of tumour cells by tumour-specific cytotoxic CD8+ T lymphocytes (Figure 6).84,95,96 Accordingly, CALR exposure, ATP secretion and

HMGB1 release are considered as references for predicting the ICD-inducing capacity of anti-cancer agents.⁹⁷

However, besides these molecular determinants of immunogenic cell death, ICD is defined by tumour-rejecting immunity in a vaccination context. ⁹⁶ In fact, gold-standard approach to asses bona fide ICD is based on in-vivo vaccination assays. ⁹⁸ In this setting, cancer cells are exposed to a potential ICD inducer in-vitro and then injected into immunocompetent mice, followed by a re-challenge with live cancer cells of the same cancer type. ⁹⁹ Hence, bona fide ICD is induced if resulting in inhibition of tumour growth, that is based on the activation of an effective anti-tumour immune response (Figure 7). ⁸⁰

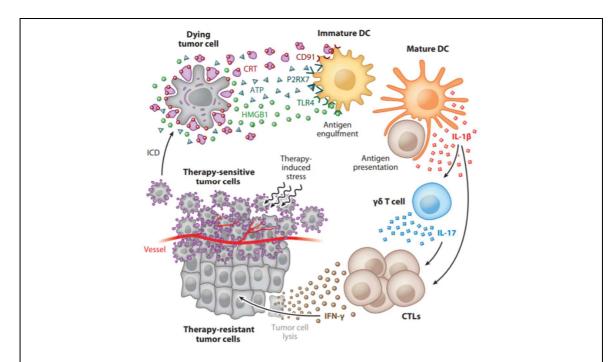


Figure 6. Molecular Definition of Immunogenic Cell Death. Dying tumour cells succumbing to immunogenic cell death (ICD) express damage-associated molecular pattern (DAMPs) hallmarks, including calreticulin (CALR), adenosine triphosphate (ATP) and high-mobility-group-protein B1 (HMGB1), that promote anti-tumour immune response by binding to respective pattern recognition receptors (PRRs) on antigen presenting cells, such as macrophages and dendritic cells (DCs). CALR engages CD91 receptors and provides an eat-me signal stimulating tumour antigen phagocytosis and presentation. ATP/P2RX7 regarded as find-me signal triggers the recruitment and maturation of antigen presenting cells. The binding of HMGB1 to Toll-like receptor 4 (TLR4) induces the release of proinflammatory cytokines and T cell priming. Consequently, these processes result in a potent Interleukin-1ß (IL-1ß)- and Interleukin-17 (IL-17)-dependent, Interferon-γ (INF-γ)-mediated tumour antigen-specific T cell response involving both γδ T cells and cytotoxic CD8+ T lymphocytes (CTLs), which lead to the eradication of tumour cells. From Kroemer *et al.*, 2013 (92).

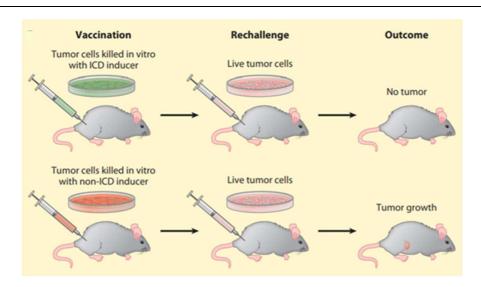


Figure 7. Functional Definition of Immunogenic Cell Death. From an operational point of view, immunogenic cell death (ICD) is defined by tumour-rejecting immunity in an in-vivo vaccination assay. Therefore, tumour cells are exposed to a potential ICD inducer in-vitro and then injected into immunocompetent mice, followed by a re-challenge with live tumour cells of the same cancer type. The operational definition of a bona fide ICD-inducer is fulfilled if a protective tumour-specific immune response is elicited resulting in the inhibition of tumour growth. From Kroemer *et al.*, 2013 (92).

3.5 Objective and Prospect

Addressing immunogenic cell death in the tumour microenvironment provides a strategy to simulate a dysfunctional anti-tumour immune system.⁹⁷ The induction of ICD contributes to the restoration of immunosurveillance and the establishment of immunological memory with long-lasting protective anti-tumour immunity, resulting in improved therapeutic efficacy and disease outcome in patients.^{100,101} Therefore, identifying novel ICD inducers is an objective of interest in the context of cancer immunotherapy.¹⁰¹

In respect of these considerations, the aim addressed in the present work is to examine the second-generation tyrosine kinase inhibitor Nilotinib for its ability to induce immunogenic cell death. The work is in the context of the group's research on the role of Gas6/TAM signalling within the tumour microenvironment regarding the pathogenesis of acute leukaemia. Hence, the question arises whether Nilotinib synergizes with a Gas6-deficient environment by inducing immunogenic cell death in Ph+ ALL to promote anti-tumour immunity.

In accordance with the criteria for assessing an anti-cancer agent for its capacity to induce immunogenic cell death, Nilotinib was investigated for the molecular and functional determinants of ICD in in-vitro and in-vivo experiments. The in-vitro experiments were performed on Ph⁺ ALL cell lines, whereby those were exposed to Nilotinib as a potential ICD inducer as well as to the commonly known ICD inducer Mitoxantrone^{85,80,85} in the sense of a positive control. In addition, untreated leukemic cells were used as negative control in the experiments. For in-vivo evaluation, a vaccination assay was performed on a Gas6 knockout (Gas6KO) mouse model on a C57BL/6 background.

The underlying methods as well as the obtained results and their evaluation are displayed hereinafter.

4 Material and Methods

4.1 Material

4.1.1 Consumable Material

The list of the consumable material is attached in the appendix (Table 1, page 89).

4.1.2 Reagents and Chemicals

Reagents and chemicals used in the experiments are listed in the appendix (Tables 2-7, page 90-96).

4.1.3 Antibodies

Table 8: Antibodies for Flow Cytometry

Antigen (Catalogue Number)	Conjugate	Dilution	Manufacturer
Annexin V (550474)	APC	1:20	BD Bioscience, Franklin Lakes, NJ, US
Calreticulin		0.25 μg/10 ⁶	R&D Systems,
(MAB38981)	-	cells, 500 µg/mL	Minneapolis, MN, US
CD11b	PE-CF594	1:1000	BD Bioscience, Franklin
(562317)	PE-0F394	1.1000	Lakes, NJ, US
CD16/CD32			BD Bioscience, Franklin
Fc receptor block	-	1:1000	,
(553141)			Lakes, NJ, US
CD45	DV/700	1.200	BD Bioscience, Franklin
(564225)	BV786	1:300	Lakes, NJ, US

CD45R/B220	APC	1:1000	BD Bioscience, Franklin
(553092)			Lakes, NJ, US
CD45R/B220	BV786	1:1000	BD Bioscience, Franklin
(563894)			Lakes, NJ, US
CD80	V450	1:300	BD Bioscience, Franklin
(560523)			Lakes, NJ, US
CD86	PE-Cy7	1:1000	BD Bioscience, Franklin
(560582)			Lakes, NJ, US
CD206	APC	1:100	BD Bioscience, Franklin
(565250)			Lakes, NJ, US
F(ab') ₂ Fragment	Alexa Fluor 594	1:500	Jackson
(715-586-150)			ImmunoResearch,
(713-300-130)	1 1001 334		Bar Harbor, ME, US
F4/80	PE	1:300	BD Bioscience, Franklin
(565410)	PE		Lakes, NJ, US
MHCII	BV650	1:1000	BD Bioscience, Franklin
(563415)			Lakes, NJ, US
Propidium iodide			Thermo Fisher
(P1304MP)	PerCP	1:5000	Scientific,
(1 1004IVII)			Waltham, MA, US

Table 9: Antibodies for Western Blot

Antigen	Conjugate	Dilution	Manufacturer
(Catalogue Number)	Conjugato	Bhation	Manaraotaror
HMGB1	_	1:5000	R&D Systems,
(MAB16901)		1.5000	Minneapolis, MN, US
IgG			Amersham
(NA935)	HRP	1:10000	Biosciences,
(144933)			Little Chalfont, UK
α-Tubulin	_	1:300	Abcam,
(ab4074)		1.000	Cambridge, UK
IgG	HRP	1:10000	Abcam,
(ab6721)	THU	1.10000	Cambridge, UK

Table 10: Antibodies for Immunohistochemistry

Antigen	Conjugate	Dilution	Manufacturer
(Catalogue Number)			- Wandiacturer
GFP	-	1:1000	Abcam,
(ab6673)			Cambridge, UK
Granzyme B		1:500	Abcam,
(ab4059)	_		Cambridge, UK

4.1.4 Chemotherapeutic Agents

Mitoxantrone was purchased from Sigma-Aldrich and supplied as crystalline solid of Mitoxantrone hydrochloride. A 2 mM stock solution was made by dissolving 10 mg Mitoxantrone hydrochloride in 9.7 mL ethanol while sonicating to be then aliquoted and stored at -20 °C.

Nilotinib was purchased from APExBIO and supplied as lyophilized powder. To prepare a 10 mM stock solution 100 mg of Nilotinib was dissolved in 18.9 mL Dimethyl sulfoxide (DMSO). Aliquots were stored at -20 °C.

Table 11: Chemotherapeutic Agents

Product Name	Manufacturer	
(Catalogue Number)	Manufacturer	
Mitoxantrone dihydrochloride	Sigma-Aldrich, St. Louis, MO, US	
(M6545)		
Nilotinib, AMN-107	ADEVALO Houston TV LIC	
(A8232)	APExBIO, Houston, TX, US	

4.1.5 Leukaemic Cell Lines (BCR-ABL B-ALL)

BCR-ABL B-ALL leukaemic cell lines were generated by Jacques Ghysdael, Research Director of Cellular Signaling and Oncogenesis at the Institut Curie in Paris. For this, whole bone marrow from Arf knockout (Arf-/-) mice (CDKN2A) was transduced with BCR-ABL-IRES-GFP retrovirus and subsequently transplanted

intravenously into sub-lethally irradiated (9 Gy) recipient wild-type mice. BCR-ABL-induced B-ALL developed individually in mice as primary leukaemias. Then primary leukaemias were cell culture adapted to generate successively stable leukaemic cell lines, such as the leukaemic cell lines L1-1 and L3 that are used in the experiments.

4.1.6 Mouse Stock

C57BL/6 wild-type mice were obtained from Charles River Laboratories. Gas6 knockout mice on a C57BL/6 background were generated by UCDAVIS KOMP Repository at the University of California Davis and Children's Hospital Oakland Research Institute.

Keeping of animals was performed according to the guidelines of Regierungspräsidium Darmstadt, Dezernat V 54 für Veterinärwesen und Verbraucherschutz. All experiments involving mice procedures were approved by the IACUC.

4.1.7 Devices and Software

The list of the devices and software is attached in the appendix (Table 12, page 96).

4.2 Methods

4.2.1 Cell Culture

L1-1 and L3 leukaemic suspension cells were routinely cultured in tissue culture flasks in leukaemic cell line medium containing RPMI 1640 Medium supplemented with 15 % heat-inactivated Foetal Bovine Serum (FBS), 1 % L-Glutamine (200 mM) and 1 % Pen Strep (10000 U/mL Penicillin, 10000 μ g/nL Streptomycin) at 37 °C in a 5 % CO₂ humidified incubator. In order to keep growth and colony conditions constant, leukaemic cells were passaged every second or third day according to their cell density. As suitable cell density 0.2 x 10⁶ to 0.5 x 10⁶ leukaemic cells per 1 mL medium were considered. Leukaemic cells were cultured no more than 12 passages before used in in-vitro or in-vivo experiments.

4.2.2 Growth Curve and Viability Count

To study the dynamics of cell growth and viability under the treatment with Nilotinib and Mitoxantrone, 0.2×10^6 L1-1 and L3 leukaemic cells were seeded in triplicate in 12-well plates in 1 mL leukaemic cell line medium and treated with increasing concentrations of Nilotinib using 10 nM, 50 nM, 100 nM, 500 nM and 1000 nM Nilotinib or 1 μ M Mitoxantrone. An untreated triplicate of both leukaemic cell lines was performed as untreated control. Number and viability of leukaemic cells per well were both monitored using the light microscope and counted by ViCell XR Cell Counter every 24 hours up to five days. To maintain appropriate culture conditions as well as to keep concentrations of Nilotinib and Mitoxantrone effective, every second day leukaemic cells were split at a 1:2 ratio, medium was exchanged and retreatment with Nilotinib or Mitoxantrone was performed at indicated concentrations in the respective wells.

4.2.3 Flow Cytometry

Flow cytometry is a method to detect the expression of cell surface and intracellular molecules.

The functional principle is based on the detection of scattered light emitted by cells passing through a light beam. Hydrodynamic focusing by a sheath fluid allows the cells in suspension to pass through the laser beam aligned one cell at a time. Light emitted by a laser light source and scattered by the passing cells is sensed by distinct detectors. A detector in front of the light beam facilitates the measurement of forward scatter (FS) and several detectors to the side of side scatter (SS) (Figure 8). As FS is proportional to cell size and SS correlates with cell granularity, cell populations can be distinguished by differences in their size and granularity in this manner (Figure 9). Next to determine cells based on FS and SS, cells can be analysed by their expression of particular proteins. For this, a fluorochrome-conjugated antibody stains the cell by binding to the referring protein of interest. The fluorescent stained cells are able to be detected individually by the emitted light of the fluorochrome when excited by a laser with the corresponding excitation wavelength. This signal of emission is converted to a voltage pulse, known as event, and amplified linearly or logarithmically. The total event height and area is measured by the flow cytometer whereby the measured event area correlates directly to the intensity of fluorescence (Figure 10).

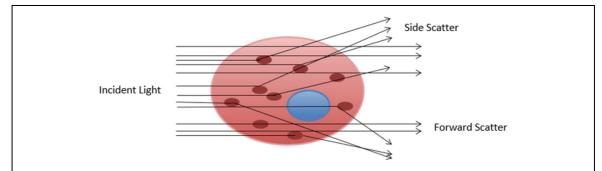


Figure 8. Light Scattering by the Cell. The direction of light scattered by the cell correlates to cell size and granularity, which is detected as forward scatter (FS) and side scatter (SS), respectively.

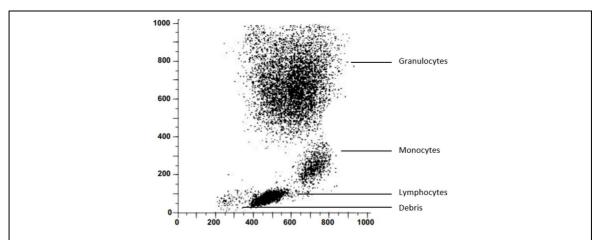


Figure 9. Determination of the Cell Population based on FS and SS in a flow cytometric Data Analysis. Dot plot of forward scatter versus side scatter of a bone marrow sample showing the characteristic position of different cell populations. Each dot represents a single cell analysed by the flow cytometer. Adapted from Riley R. S. Principles and Applications of Flow Cytometry. http://www.flowlab-childrens-harvard.com. Accessed January 10, 2020.

The process of using fluorochrome-labelled antibodies is called direct staining. In contrast to this, in indirect staining an antibody without fluorochrome conjugation binds to the target protein and is then detected by a fluorochrome-labelled secondary antibody. Flow cytometry can be used to perform extracellular staining of surface proteins as well as to run intracellular staining of proteins within the cell after fixation and permeabilization with specific agents.

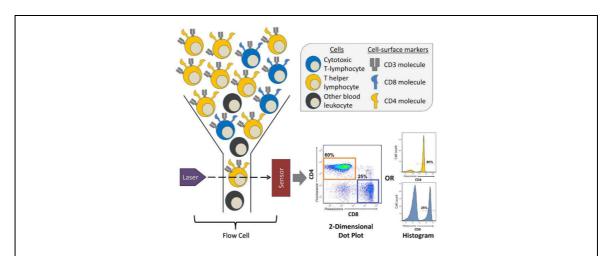


Figure 10. Set-up and functional Principle of Flow Cytometry and Data Analysis. Cells pass through the laser beam focused by a sheath fluid one cell at a time. Both forward and side scattered light as well as fluorescence emitted from stained cells are detected by a sensor. The derived information is visualised digitally as two-dimensional dot plots or one-dimensional histograms allowing the quantification of frequency of the analysed cells. From Verschoor *et al.*, 2015 (1).

Taking into account the mentioned aspects about flow cytometry, this method was used to analyse different cell types in heterogenous cell populations of different experiments. Thereby, cells were classified within a specific gating strategy regarding not only FS and SS but also their fluorescence. In order to correct the signal overlap between the channels of the emission spectra of different fluorochromes, compensation was performed for all experiments. To avoid non-specific binding all samples were incubated with an Fc receptor blocking antibody. Antibodies with respective dilutions used in the experiments are listed in Table 8. All antibody-mixes were prepared on ice using DPBS containing 2 % FBS as staining buffer. To each sample 50 µL of antibody-mix were applied for staining, that was performed at room temperature in the dark. Before and after staining as well as between different staining steps, cells were washed twice with DPBS by spinning down for 5 minutes at 1500 revolutions per minute (rpm) at 4 °C and discarding the supernatant. For flow cytometry analysis cells were then re-suspended in FACS buffer using 300 µL of DPBS supplemented with 2 % FBS. Flow cytometry data were acquired on a BD LSRFortessa Flow Cytometer and analysed using FlowJo software. At least 10000 events were acquired per sample.

4.2.4 Western Blot

Western Blot is a technique to detect a protein of interest in a complex protein mixture by specific antibody-antigen interactions that allows a qualitative and semi-quantitative analysis of the target protein.

The first step in a western blotting procedure is the sample preparation including the lysis in an appropriate lysis buffer as well as the reduction in sample buffer and the following heat denaturation. Thereafter, proteins are separated by gel electrophoresis commonly using polyacrylamide gel electrophoresis (PAGE). While native PAGE separates proteins according to their mass-charge ratios, sodium dodecyl sulphate-PAGE (SDS-PAGE) separates proteins based on their mass only by masking the endogenous protein charge with a negative charge. Thus, upon application of an electric field, proteins of lower molecular weight run faster towards the positive cathode.

Following electrophoresis, proteins are transferred from the gel onto a membrane generally made of nitrocellulose or polyvinylidene difluoride (PVDF). Electrophoretic transfer, performed as wet or semi-dry transfer, is the most used transfer method allowing the negatively charged proteins to move out of the polyacrylamide gel towards the positive cathode and onto the membrane when an electric field is applied. In order to prevent non-specific binding and to reduce background interference, the free sites on the membrane are blocked with a blocking buffer before antibody staining. The process of immunodetection is performed by probing the blocked membrane in a first step with a primary antibody that binds to the protein of interest and in a second step with an enzyme-conjugated secondary antibody. Finally, adding chromogenic, fluorogenic or chemiluminescent substrates allows the visualisation of signal development that is based on the enzyme-substrate reaction and detected with a suitable detection system.

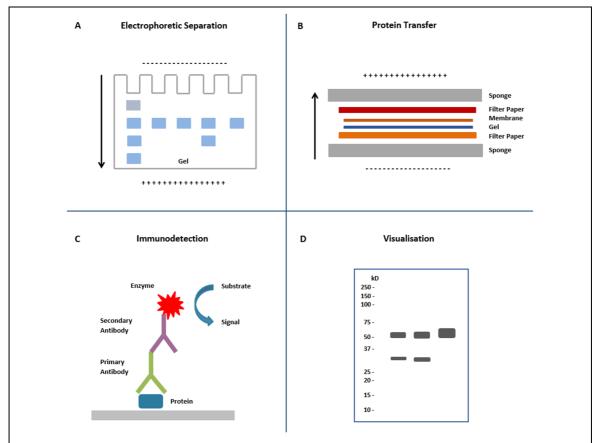


Figure 11. The Technique of Western Blotting. Western Blot is performed in four main steps: Protein separation by gel electrophoresis (A), protein transfer from the gel onto a membrane (B), immunodetection of the protein by antibody staining (C) and visualisation of the protein detection (D).

To check for equal loading of the gel and a correct transfer of the proteins onto the membrane, ideally, a loading control is done when running a Western Blot. Hence, antibodies used for loading controls target housekeeping proteins showing a ubiquitous and consistently high level of expression. The technique of Western blotting is illustrated in Figure 11.

4.2.5 Cell Death Assay

In order to assess the modality of Nilotinib-induced cell death in leukaemic cell lines, 1 x 10 6 L1-1 and L3 leukaemic cells were plated in 12-well plates in 2 mL leukaemic cell line medium and treated with Nilotinib at increasing concentrations of 10 nM, 50 nM or 100 nM for 4, 8, 12 and 48 hours, respectively. To compare with a positive and negative control, same conditions were tested using 1 μ M Mitoxantrone or leaving cells untreated. After incubation at 37 °C in a 5 % CO₂ incubator at indicated concentrations and time, leukaemic cells were harvested and stained for cell death markers Annexin V and Propidium iodide together with B cell surface marker B220. After 30 minutes of incubation at room temperature in the dark, cells were analysed by flow cytometry.

4.2.6 Determination of ICD Markers

4.2.6.1 Calreticulin Exposure

To determine CALR exposure under treatment with Nilotinib, 1 x 10^6 L1-1 and L3 leukaemic cells were seeded in duplicate in 12-well plates in 1 mL leukaemic cell line medium and treated with 10 nM or 50 nM of Nilotinib for 4 and 6 hours, respectively. Under same conditions an untreated negative control as well as a positive control treated with 1 μ M Mitoxantrone were performed. After incubation, cells were first stained with an anti-CALR primary antibody for 30 minutes and then, in a second step, with an Alexa Fluor 594-conjugated IgG secondary antibody for 15 minutes at room temperature in the dark. In order to detect non-specific background staining, secondary controls were carried out for all conditions. Finally, samples were analysed using flow cytometry.

4.2.6.2 Extracellular ATP Release

Nilotinib-induced release of extracellular ATP was quantified using an ENLITEN ATP Assay System Bioluminescence Detection Kit for ATP Measurement according to the manufacturer's instructions.

In the experiment 2.5×10^3 L1-1 leukaemic cells were plated in triplicate in a 96-well plate in 50 µL leukaemic cell line medium. Nilotinib was added at concentrations of 10 nM or 50 nM and untreated triplicates as negative control as well as triplicates treated with 1 µM Mitoxantrone as positive control were performed. To examine the dynamics of extracellular ATP release over the time during treatment, conditions mentioned above were tested for 4, 6, 8, 12 and 24 hours. In order to determine possible effects of the sample buffer (= leukaemic cell line medium) on the luciferase reaction, an ATP standard curve in the range of 10^{-18} M ATP to 10^{-10} M ATP was run in duplicate. Furthermore, both a triplicate with leukaemic cell line medium only and a triplicate with ATP-free water were carried out to quantify background luminescence.

ATP-derived chemoluminescence was detected on a LUMIstar OPTIMA Microplate Luminometer with an automatic reagent delivery system applying $50~\mu L$ of recombinant Luciferin/Luciferase-reagent (rL/L-reagent) to each sample. The measurement was run with a 2-second delay time after reagent injection and a 10-second luminescence signal integration.

4.2.6.3 HMGB1 Release

Aiming to investigate the release of HMGB1 under the treatment with Nilotinib, Western Blotting was performed. Therefore, 10 x 10 6 L1-1 leukaemic cells were seeded into tissue culture flasks in 20 mL leukaemic cell line medium. Nilotinib at the concentrations of 10 nM or 50 nM was added and cells were incubated for 8 hours. Next to that, a positive control treated with 1 μ M Mitoxantrone as well as an untreated negative control were performed under the same conditions. To determine cytosolic HMGB1, whole-cell extracts were prepared by lysing cells with RIPA lysis buffer (10 μ L/1 x 10 6 cells) supplemented with protease inhibitor (10 μ L/1 mL lysis buffer) and phosphatase inhibitor (10 μ L/1 mL lysis buffer). In order to assess extracellular HMGB1 release, 5 mL of cell culture supernatants

were collected, and proteins precipitated using the method of acetone precipitation. Then, samples were reduced and denatured by adding 4x Laemli sample buffer and heating at 95 °C for 5 minutes. After loading of 20 µL of cytosolic extracts and 10 µL of precipitated cell culture supernatants together with a molecular weight marker, proteins were separated by 15 % SDS-PAGE running 30 minutes at 80 V and 1 hour at 100 V in running buffer. Thereafter, proteins were transferred from the gel onto a 0.2 µm nitrocellulose blotting membrane in a wet transfer system in transfer buffer at 250 mA for 1.5 hours at 4 °C on ice. To reduce non-specific binding, the membrane was placed in blocking buffer containing 5 % skimmed milk overnight at 4 °C before immunodetection. The latter was performed by probing the blocked membrane with an anti-HMGB1 primary antibody overnight at 4 °C followed by the incubation with an HRPconjugated secondary IgG antibody for 1 hour at room temperature with both antibodies diluted in blocking buffer containing 2.5 % skimmed milk. Before and after every step of antibody incubation, the membrane was washed with TBS-T buffer three times for 5 minutes. For visualization, the chemiluminescent ECL Prime Western Blotting Detection Reagent with a final volume of 0.1 mL/cm² membrane was added onto the probed membrane, incubating for 5 minutes. Then, the membrane was exposed to an X-ray film in a dark room for 30 seconds to detect cytosolic HMGB1 protein bands and for 90 seconds to visualise extracellular HMGB1 from cell culture supernatants. Finally, the film was developed immediately in the X-ray film developer machine XR24 Pro to acquire the western blot images. An α-Tubulin loading control was performed for cytosolic HMGB1.

4.2.7 Macrophages Co-Culture

In the purpose of examining the effects of Nilotinib-treated leukaemic cells on macrophages activation and polarisation in an in-vitro setting, bone marrow cells were harvested from the femurs of a 12-week-old female C57BL/6 wild-type mouse and cultured in 30 mL of macrophages medium in Teflon-coated cell culture bags for 7 days at 37 °C in a 5 % CO2 humidified incubator to generate mature CD11b+F4/80+ M0-polarised macrophages. Macrophages medium containing DMEM Medium supplemented with 10 % FBS, 1 % L-Glutamine (200 mM) and 1 % Pen Strep (10000 U/mL Penicillin, 10000 μ g/nL Streptomycin) in the presence of recombinant mouse macrophage colony-stimulating factor (M-CSF, 100 μ g/mL) was exchanged every second day.

On day 6, 0.5 x 10⁶ macrophages were plated into 12-well UpCell MultiDish cell culture dishes in 1 mL macrophages medium for optimal attachment. Next to that, 1 x 10⁶ L1-1 and L3 leukaemic cells were seeded in 6-well plates in 2 mL leukaemic cell line medium and treated with Nilotinib at concentrations of 10 nM or 50 nM for 24 hours. For a positive and negative control, same conditions were performed using 1 µM Mitoxantrone or leaving leukaemic cells untreated for 24 hours, respectively. On day 7, bone marrow-derived macrophages (BMDM) were then co-cultured with 0.5 x 10⁶ treated leukaemic cells in 1 mL macrophages medium (1:1 ratio) or with 500 µL of treated leukaemic cells medium (= supernatant) in 500 μL macrophages medium (1:1 ratio) for additional 24 hours, respectively. Moreover, bone marrow-derived macrophages without co-culturing were cultured in duplicate for control. Finally, after incubation with 30 µL of Brilliant Stain Buffer for 15 minutes at room temperature in the dark, cell cultures were stained with fluorescence-labelled antibodies against CD45, CD11b, F4/80, CD80, CD86, CD206 and MHCII and analysed by flow cytometry (Figure 12).

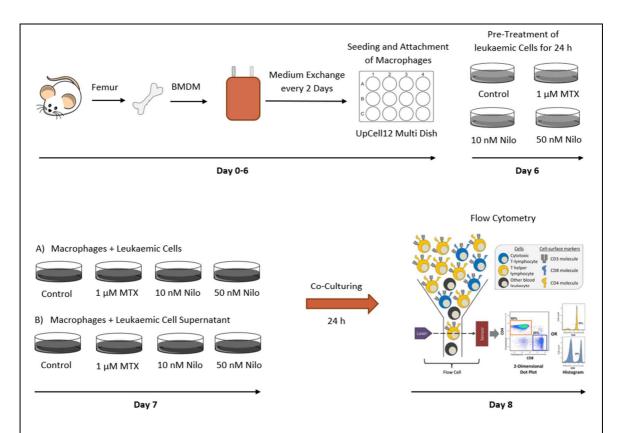


Figure 12. Experimental Setup of Macrophages Co-Culture. Bone marrow cells of two murine femurs were cultivated in macrophages medium in Teflon-coated cell culture bags and seeded into 12-well UpCell MultiDish cell culture dishes in order to generate mature bone marrow-derived macrophages (BMDM) with M0-polarisation. Leukaemic cells were treated with Nilotinib (Nilo) or Mitoxantrone (MTX) at indicated concentrations or left untreated for 24 hours, respectively. Finally, bone marrow-derived macrophages were co-cultured with treated leukaemic cells or with treated leukaemic cells supernatant at a 1:1 ratio. After additional 24 hours, cell cultures were stained and flow cytometrically analysed for markers of macrophages activation and polarisation.

4.2.8 Vaccination Assay

In order to evaluate Nilotinib-induced immunogenic cell death in-vivo, L1-1 leukaemic cells were incubated with 100 nM Nilotinib for 72 hours or with 1 μ M Mitoxantrone for 24 hours. Then 10 x 10³ treated leukaemic cells were inoculated subcutaneously into the left flanks of eight-week-old female C57BL/6 wild-type and Gas6 knockout mice. After two weeks, mice were challenged with 1 x 10³ living L1-1 leukaemic cells injected with a viability of 96 % into the tail vein. Thereafter, mice were monitored routinely for the incidence of leukaemic burden. When both clinical symptoms of progressing leukaemia and high leukaemic

burden of the peripheral blood in flow cytometry analysis were displayed, mice were sacrificed for a final analysis (Figure 13). Extent and procedure of the final analysis are descripted in Table 13.

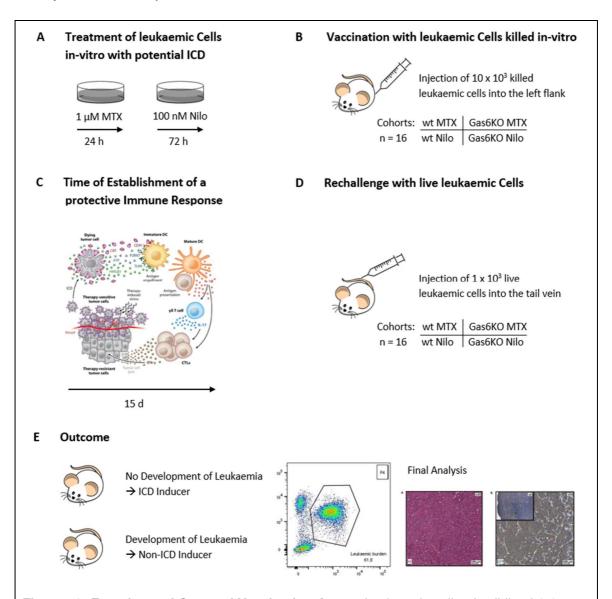


Figure 13. Experimental Setup of Vaccination Assay. Leukaemic cells of cell line L1-1 were incubated with 1 μ M Mitoxantrone for 24 hours or with 100 nM Nilotinib for 72 hours (A). For vaccination, 10 x 10³ killed leukaemic cells were injected subcutaneously into the left flank of the experimental mice (B). After 15 days considered for the establishment of a protective immune response (C), mice were re-challenged with 1 x 10³ live leukaemic cells applied into the tail vein (D). Then, mice were monitored routinely for the incidence of leukaemic burden. When both leukaemia-typical symptoms and high leukaemic burden in flow cytometry analysis were displayed, mice were sacrificed for a final analysis (E). The cohort of experimental mice included 8 wild-type and 8 Gas6 knockout mice, half of which were vaccinated and re-challenged with Nilotinib treated cells and the other half with Mitoxantrone treated cells.

Table 13: Extent and Procedure of Final Analysis

Nilotinib co	<u>ohort</u> : n = 8, i	ncluding 4 wt and 4 Gas6KO mice	
		nalysis on day 8	
$\underline{\text{Mitoxantrone cohort}}: \qquad \qquad n = 8, \text{ in}$		ncluding 4 wt and 4 Gas6KO mice nalysis on day 13	
	i iiai a	Bones were harvested. Femur was fixed	
Femur + Tibia	 IHC (GFP) Flow cytometry analysis of leukaemic burden 	for Immunohistochemistry (IHC) and tibia was crushed to stain cells with PI for dead cell exclusion and B-cell surface marker B220. Leukaemic burden was quantified by the frequency of GFP+B220+ cells using flow cytometry analysis.	
Blood	 White Blood Count Flow cytometry analysis of leukaemic burden 	Blood was taken from the submandibular vein. White Blood Count was performed using scil Vet abc hematology analyzer. Blood lysed with lysing buffer was stained with PI for dead cell exclusion and B-cell surface marker B220. Leukaemic burden was quantified by the frequency of GFP+B220+ cells using flow cytometry analysis.	
Spleen	 Spleen weight IHC (GFP) Flow cytometry analysis of leukaemic burden 	Spleen was harvested and weighted. Half of the spleen was fixed for Immunohistochemistry (IHC). The remaining half of the spleen was smashed to stain cells with PI for dead cell exclusion and B-cell surface marker B220. Leukaemic burden was quantified by the frequency of GFP+B220+cells using flow cytometry analysis.	
Tumour	HE stain and IHC (GFP)	All mice of the Nilotinib cohort developed subcutaneous tumours at the injection site. Tumours were harvested and a piece was fixed for Haematoxylin-Eosin (HE) stain and Immunohistochemistry (IHC).	

4.2.9 Haematoxylin-Eosin Stain and Immunohistochemistry

Collected tissues from the vaccination experiment were prepared for Immunohistochemistry. Therefore, tissues were fixed with Roti-Histofix for 24 hours and then washed with DPBS for another 24 hours. In addition to this, femurs were decalcified in decalcification buffer for 14 days. To permit sectioning, samples were dehydrated and paraffin embedded. Dehydration was done in a Leica ASP300 S Tissue Processor by passing the samples through a series of alcohol solutions in ascending concentrations (70 %, 80 %, 96 %, 96 %, 100 %, 100 %, 100 % ethanol) and by clearing with Xylene to remove the ethanol prior to paraffin-embedding. After immersion and blocking in paraffin wax, tissues were sectioned at 3 µm with a Leica RM2235 Manual Rotary Microtome and then transferred onto glass slides. Haematoxylin-Eosin stain was performed using Leica ST5010 Autostainer XL. Leica BOND-MAX was used for fully automated GFP-immunostaining applying 150 µL of antibody diluted with Bond Primary Antibody Diluent per slide. After staining, slides were dehydrated in Leica ST5010 Autostainer XL through an ascending series of alcohol (50 %, 70 % and 96 % ethanol, 100 % isopropanol) and cleared with Xylene to then get cover-slipped with Leica CV5030 Glass Coverslipper using CellTexx mounting medium. Finally, slides were examined under an Axio Imager 2 Research Microscope and imaged using the AxioVision SE64 Microscopy software.

4.2.10 Statistical Analysis

Statistical analysis was performed using GraphPad Prism software. Results are expressed as mean \pm standard deviation (SD). Significant difference between two groups was analysed by unpaired, two-tailed Student t-tests. Statistical significance was determined by a value of p < 0.05 for all analyses. P values are presented in the figures as * p < 0.05, ** p < 0.01, *** p < 0.001, **** p < 0.0001.

5.1 Effects of Nilotinib on Tumour Cell Growth and Viability

To study the effects of Nilotinib on both growth trends and viability of leukaemic tumour cells in concentration-dependent manner, cells of the leukaemic cell lines L1-1 and L3 were treated in-vitro with increasing doses of Nilotinib between 10 nM and 1000 nM. The number and viability of cells were monitored every 24 hours for 5 days.

While untreated tumour cells showed unrestrained positive accelerated growth, Nilotinib significantly inhibited growth at all concentrations (Figure 14A and 14B). Here, a higher efficacy was seen at the concentrations greater than 10 nM Nilotinib, as at those concentrations there was an overall decreasing cell count after an initial increase, whereas leukaemic cells treated with 10 nM Nilotinib showed a constantly increasing number of cells despite inhibited growth. However, there were no significant differences in efficacy between the concentrations of Nilotinib greater than 10 nM (Figure 14C and 14D). Next to that, the two leukaemic cell lines exhibited no statistically different sensitivities to Nilotinib exposure.

As Nilotinib, Mitoxantrone was potent to inhibit tumour cell growth significantly compared to the untreated control (Figure 15A and 15B). In contrast, the growth-inhibiting effect of Mitoxantrone was not delayed as with Nilotinib, resulting in an immediately decreasing number of cells (Figure 15C and 15D). Likewise, the leukaemic cell lines L1-1 and L3 were similar in their sensitivity referring to the treatment with Mitoxantrone.

In terms of viability, effects were seen with concentration of Nilotinib greater than 10 nM for leukaemic cell line L1-1 and above 50 nM for L3, while untreated cells maintained a high viability of in average 97 % (L1-1) or 94 % (L3). In comparison to Mitoxantrone, that exposed high und prompt cytotoxicity, cell viability was gradually decreasing with Nilotinib over the time (Figure 16A and 16B).

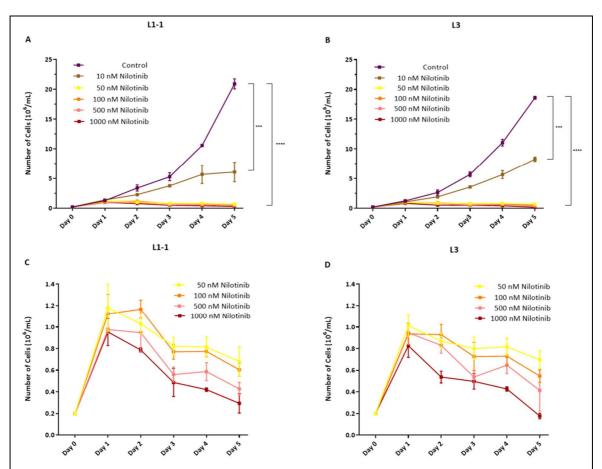


Figure 14. Effects of Nilotinib on Tumour Cell Growth. Tumour cells of leukaemic cell lines L1-1 (A, C) and L3 (B, D) were treated with 10, 50, 100, 500 or 1000 nM Nilotinib or left untreated as control. Number of cells was measured using a ViCell XR Cell Counter every 24 hours for 5 days. Data presented as mean \pm SD. Statistical was significance analysed by unpaired, two-tailed Student t-test. Significance values: * p < 0.05, ** p < 0.01, *** p < 0.001, **** p < 0.0001.

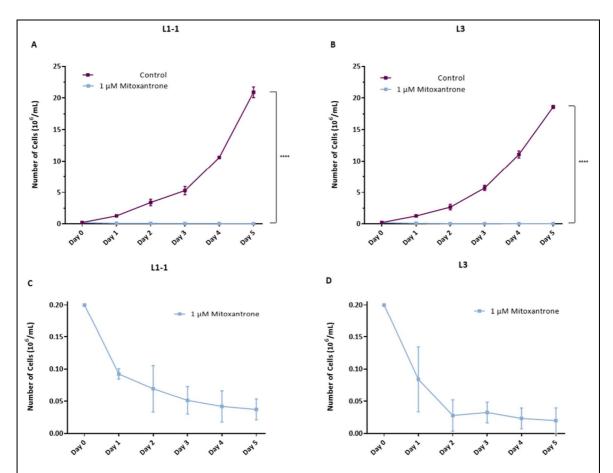


Figure 15. Effects of Mitoxantrone on Tumour Cell Growth. Tumour cells of leukaemic cell lines L1-1 (A, C) and L3 (B, D) were treated with 1 μ M Mitoxantrone or left untreated as control. Number of cells was quantified using a ViCell XR Cell Counter every 24 hours for 5 days. Data presented as mean \pm SD. Statistical significance was analysed by unpaired, two-tailed Student t-test. Significance values: * p < 0.05, ** p < 0.01, *** p < 0.001, **** p < 0.0001.

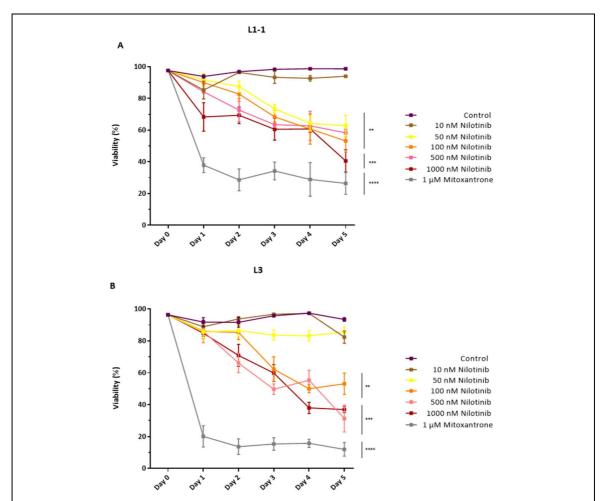


Figure 16. Effects of Nilotinib on Tumour Cell Viability. Tumour cells of leukaemic cell lines L1-1 (A) and L3 (B) were treated with 10, 50, 100, 500 or 1000 nM Nilotinib. For comparison, leukaemic cells were treated with 1 μ M Mitoxantrone or left untreated as control. Cell viability was measured using a ViCell XR Cell Counter every 24 hours for 5 days. Viability is defined as the percentage of viable cells of the total number of cells. Data presented as mean \pm SD. Statistical significance was analysed by unpaired, two-tailed Student t-test. Significance values: * p < 0.05, ** p < 0.01, *** p < 0.001, **** p < 0.0001.

5.2 Characterisation of Nilotinib-induced Cell Death

To further determine the process of Nilotinib-induced cell death, an in-vitro Annexin V/ Propidium iodide cell death assay was performed. This assay allows not only to quantify the extent of cell death but also to distinguish between the cell death modality ranging from apoptosis to necrosis. Therefore, Nilotinib-treated leukaemic cells of both cell lines were stained with Annexin V (AnxV) and Propidium iodide (Pi) in order to qualify cell death by a flow cytometry-based

analysis. With the help of the a gating strategy (Supplementary Figure S2, page 98), cells were finally categorized on the one hand into viable cells (AnxV-Pi-) and on the other hand into dead apoptotic cells (AnxV+Pi- and AnxV+Pi+) as well as dead necrotic cells (AnxV-Pi+). Flow cytometry data with all concentrations and different times of incubations are presented in Figure 17 (L1-1) and Figure 18 (L3). In addition to that, data are figured as bar diagrams in Figure 19.

Both types of figure show that after 4 and 8 hours of Nilotinib treatment leukaemic cells were mainly viable and displayed a distribution similar to the untreated control. Depending on the dose, slightly more dead cells compared to the control were found after 24 hours. While Nilotinib at a concentration of 10 nM had no more considerable effect on cell death after another 24 hours, 48 % of the L1-1 and 34 % of the L3 leukaemic cells treated with 50 nM Nilotinib and 60 % of the L1-1 and 40 % of the L3 leukaemic cells treated with 100 nM Nilotinib were dead after 48 hours. Notably, referring to the cell death modality, Nilotinib induced a bimodal cell death, that was predominantly necrotic in L1-1 and apoptotic in L3 leukaemic cells.

In comparison, Mitoxantrone-treated cells of both leukaemic cell lines displayed a nearly complete apoptotic cell death within 4 hours.

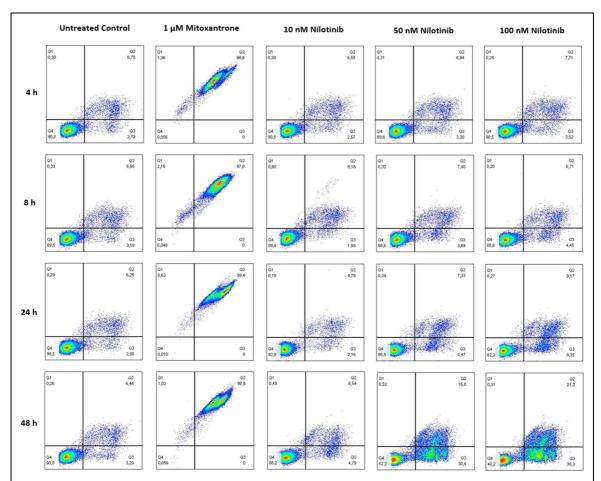


Figure 17. Flow Cytometry based Analysis of Nilotinib-induced Cell Death. Cell Line 1-1. Tumour cells of leukaemic cell lines L1-1 were exposed to 10, 50 or 100 nM Nilotinib for 4, 8, 24 or 48 hours, respectively. Comparatively, same conditions were performed with an untreated control and leukaemic cells treated with 1 μ M Mitoxantrone. Cell death was evaluated by flow cytometry analysis using a staining with Annexin V (AnxV) and Propidium iodide (Pi) to discriminate between apoptotic (Q1 + Q2: AnxV+Pi- and AnxV+Pi+), necrotic (Q3: AnxV-Pi+) and viable (Q4: AnxV-Pi-) cells.

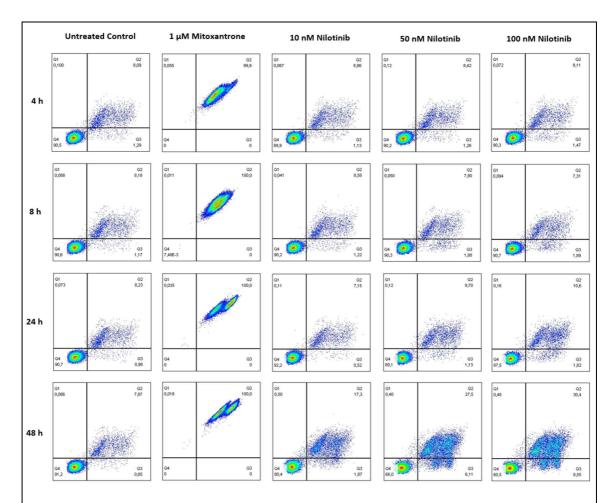


Figure 18. Flow Cytometry based Analysis of Nilotinib-induced Cell Death. Cell Line 3. Tumour cells of leukaemic cell lines L3 were exposed to 10, 50 or 100 nM Nilotinib for 4, 8, 24 or 48 hours, respectively. Comparatively, same conditions were performed with an untreated control and leukaemic cells treated with 1 μM Mitoxantrone. Cell death was evaluated by flow cytometry analysis using a staining with Annexin V (AnxV) and Propidium iodide (Pi) to discriminate between apoptotic (Q1 + Q2: AnxV+Pi- and AnxV+Pi+), necrotic (Q3: AnxV-Pi+) and viable (Q4: AnxV-Pi-) cells.

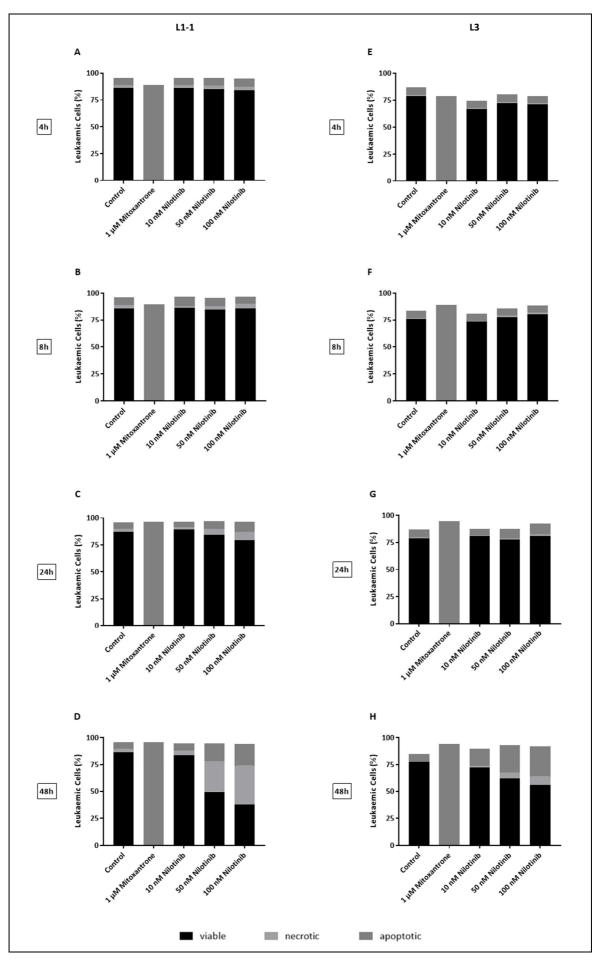


Figure 19. Cell Death Assay. Tumour cells of leukaemic cell lines L1-1 (A-D) and L3 (E-H) were exposed to 10, 50 or 100 nM Nilotinib for 4, 8, 24 or 48 hours, respectively. Comparatively, same conditions were performed with an untreated control and leukaemic cells treated with 1 μ M Mitoxantrone. Cell death was evaluated by flow cytometry analysis using a staining with Annexin V and Propidium iodide to discriminate between viable, apoptotic and necrotic cells. Note: In the bar diagrams 100 % of leukaemic cells were not reached in total due to the exclusion of debris and doublets from the cell population.

5.3 Molecular Determinants of Immunogenic Cell Death. ICD Marker Analysis

5.3.1 Calreticulin Exposure

In order to investigate a Nilotinib-triggered cell surface exposure of Calreticulin, leukaemic cells of both cell lines were stained for Calreticulin by flow cytometry after treatment with Nilotinib. The flow cytometry-based analysis was performed using a gating strategy, in which the positive gating region for the expression of Calreticulin was constituted with the aid of unstained controls (Supplementary Figure S3, page 99).

The results are illustrated in Figure 20, where Calreticulin expressing cells are given as frequency of total. Here, only the percentage increase in Calreticulin expression compared to the untreated control was evaluated as drug-induced expression, considering the background staining of the target antibody, since the unstimulated control should not be expressing the target.

Accordingly, cell line L1-1 exhibited no Calreticulin exposure after 4 hours and only 2 % after 6 hours, while an exposure of Calreticulin in cell line L3 was detected neither after 4 hours nor after 6 hours of Nilotinib treatment. In contrast, a Mitoxantrone-induced Calreticulin exposure of 8 % in cell line L1-1 after 6 hours and of 6 % in cell line L3 after 4 hours was noted.

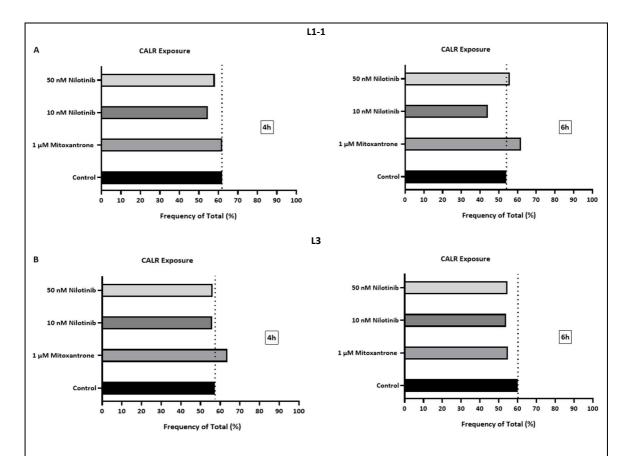


Figure 20. ICD Marker Analysis of Calreticulin Exposure. Tumour cells of leukaemic cell lines L1-1 (A) and L3 (B) were incubated with 10 or 50 nM Nilotinib for 4 or 6 hours, respectively. For comparison, same conditions were performed with an untreated control and leukaemic cells treated with 1 μ M Mitoxantrone. Subsequently, cells were stained by flow cytometry for cell surface expression of Calreticulin (CALR). Results are given as frequency of total. The dashed line sets the limit to the drug-induced exposure of Calreticulin alone with regard to the untreated control.

5.3.2 Extracellular ATP Release

For the quantification of Nilotinib-induced release of extracellular ATP a bioluminescence-based ATP assay was used. Therefore, leukaemic cells of cell line L1-1 were exposed to Nilotinib and ATP-derived luminescence was measured at different points of time on a luminometer.

The principle of quantifying extracellular ATP by the measurement of luminescence is based on the assumption that the intensity of the emitted light is proportional to the ATP concentration, since ATP is the limiting component of the luciferase reaction (Figure 21B). To take into account possible effects of the sample buffer, here leukaemic cell line medium, on the luciferase reaction, an

ATP standard curve was prepared (Figure 21A). Thus, the concentration of ATP was determined by comparing to the standard curve. In fact, luminescence was generally lower in the sample buffer as FBS-supplemented media is known to inhibit ATP availability for luciferase reaction.¹⁰² Consequently, the buffer inhibition of light output was corrected by a correction factor.

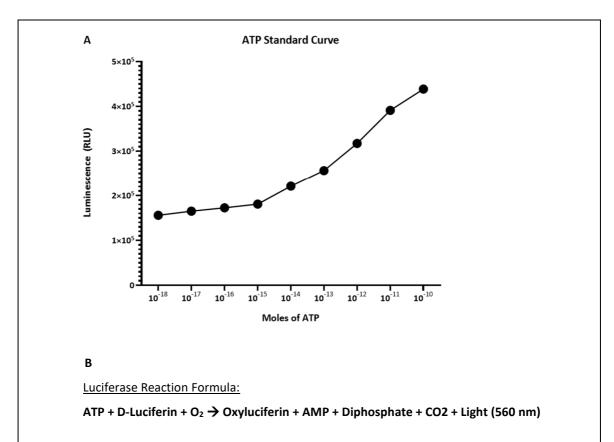


Figure 21. Bioluminescence based ATP Assay for the Detection of Extracellular ATP. ATP Standard Curve and Luciferase Reaction Formula. In order to determine possible effects of the sample buffer (= leukaemic cell line medium) on the luciferase reaction, an ATP standard curve in the range of 10⁻¹⁸ M ATP to 10⁻¹⁰ M ATP was constructed by diluting the ATP Standard (10⁻⁷ M ATP) with sample buffer to the proper concentrations (A). The quantification of extracellular ATP by the measurement of luminescence as light intensity (RLU = relative light units) is based on the assumption that the intensity of the emitted light is proportional to the ATP concentration, since ATP is the limiting component of the luciferase reaction. The underlying formula of the luciferase reaction is shown in (B).

Data of the analysis of extracellular ATP release are presented in Figure 22. At any time point, there was effectively no release of ATP under the exposure of either 10 nM or 50 nM Nilotinib. In contrast, a significant release of ATP from leukaemic cells under treatment with Mitoxantrone was measured with absolute

peak levels of 500 x 10^{-15} M ATP after 4 and 6 hours and, in relation to the untreated control, a relative peak level of 400×10^{-15} M ATP after 8 hours, which were gradually decreasing within 24 hours. Results were evaluated considering both the baseline and the untreated control. The baseline indicates the baseline ATP concentration and was set with respect to ATP concentrations measured in ATP-free water, which is why only values above the baseline were assessed as specific.

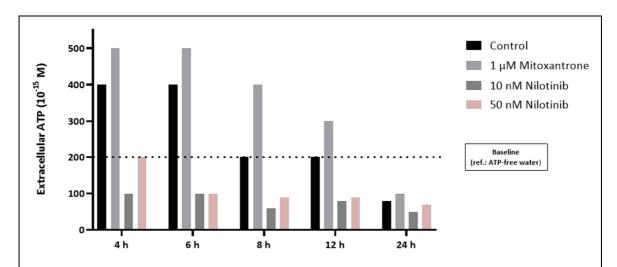


Figure 22. ICD Marker Analysis of Extracellular ATP Release. Tumour cells of leukaemic cell line L1-1 were treated with 10 or 50 nM Nilotinib. Using an ENLITEN ATP Assay System Bioluminescence Detection Kit for ATP Measurement, extracellular ATP release over the time was quantified after 4, 6, 8, 12 and 24 hours of exposure to Nilotinib based on ATP-derived chemoluminescence detected with a luminometer. Comparatively, conditions mentioned above were performed with an untreated control and leukaemic cells incubated with 1 μ M Mitoxantrone. Baseline ATP concentration is indicated as dashed line.

5.3.3 **HMGB1 Release**

Nilotinib-induced release of HMGB1 from leukaemic cells was assessed by Western blotting. Therefore, HMGB1 in both cytosolic extracts and cell culture supernatants of Nilotinib-exposed leukaemic cells of cell line L1-1 was determined after 12 hours of incubation. For cytosolic HMGB1 release, a loading control using α -Tubulin was performed.

Graphical data of Western blotting indicated a dose-dependent release of cytosolic HMGB1 from leukaemic cells challenged with 10 nM or 50 nM Nilotinib.

Likewise, cytosolic HMGB1 release under treatment with Mitoxantrone was found to be between the two levels of Nilotinib. Hence, in comparison to the untreated control, a drug-specific release of cytosolic HMGB1 was approved for Nilotinib and Mitoxantrone (Figure 23A).

Taking into account that the release of HMGB1 is the process of HMGB1 translocation from the nucleus via the cytosol to the extracellular space during cell death, there was not yet an extracellular HMGB1 release into cell culture supernatants from leukaemic cells exposed to Nilotinib. In contrast to that, a Mitoxantrone-induced release of extracellular HMGB1 was already detected (Figure 23B).

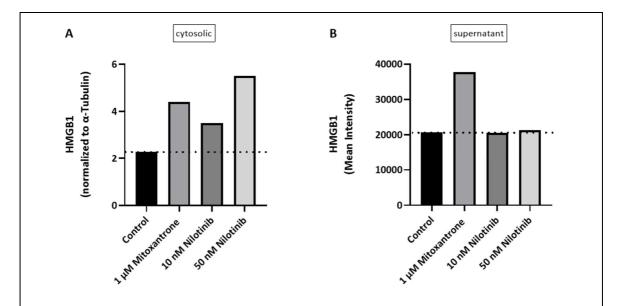


Figure 23. ICD Marker Analysis of HMGB1 Release. Tumour cells of leukaemic cell line L1-1 were exposed to 10 or 50 nM Nilotinib for 12 hours. Comparatively, same conditions were performed with an untreated control and leukaemic cells treated with 1 μ M Mitoxantrone. Then, HMGB1 in cytosolic extracts (A) and cell culture supernatants (B) were determined by Western blotting. Cytosolic HMGB1 release is given normalized to α -Tubulin that was used as loading control. Extracellular HMGB1 release into cell culture supernatants is expressed as mean intensity. Dashed line is set with regard to the untreated control displaying drug-specific release of HMGB1.

5.4 Immunological Determinants of Immunogenic Cell Death

5.4.1 Macrophages Co-Culture

In order to examine the immunological effects of Nilotinib on macrophages activation and polarisation, macrophages were co-cultured on the one hand side with Nilotinib pre-treated leukaemic cells in a juxtacrine setup and on the other hand side with leukaemic cells supernatant of Nilotinib pre-treated leukaemic cells in a paracrine setup. Therefore, leukaemic cells of both leukaemic cell lines L1-1 and L3 were pre-treated with 10 nM or 50 nM Nilotinib for 24 hours and subsequently co-culturing with bone marrow-derived macrophages was performed for another 24 hours. Finally, macrophages were analysed by flow cytometry with the help of a gating strategy (Supplementary Figure S4, page 100) investigating macrophages activation and polarisation based on their surface expression of CD80, CD86, CD206 and MHCII.

Data of the flow cytometry-based analysis of macrophages co-culture presented in Figure 24 did not reveal a relevant shift in macrophages activation and polarisation in the juxta- and paracrine co-culture conditions with Nilotinib pretreated leukaemic cells of both cell lines compared to the corresponding co-culture conditions with untreated leukaemic cells. More precisely, the shift of the surface expression profile regarding CD80, CD86, CD206 and MHCII was less than 5 % on total average and with partly opposing changes among the juxta-and paracrine co-culture conditions as well as between the two cell lines L1-1 and L3 (Table 14 and 15, page 101-102).

In contrast to that, there were effects on macrophages activation and polarisation in the co-culture conditions with Mitoxantrone pre-treated leukaemic cells in comparison with the corresponding co-culture conditions of untreated leukaemic cells involving an average upregulation of CD80 by 4.4 %, of CD86 by 26.5 % and of CD206 of 24.1 % besides a downregulation of MHCII by 9.0 %. Here, the changes in the surface expression profile of the two cell lines were similar, but more pronounced by an average of 58.0 % in the paracrine setup comparing the two co-culture conditions (Table 16, page 103).

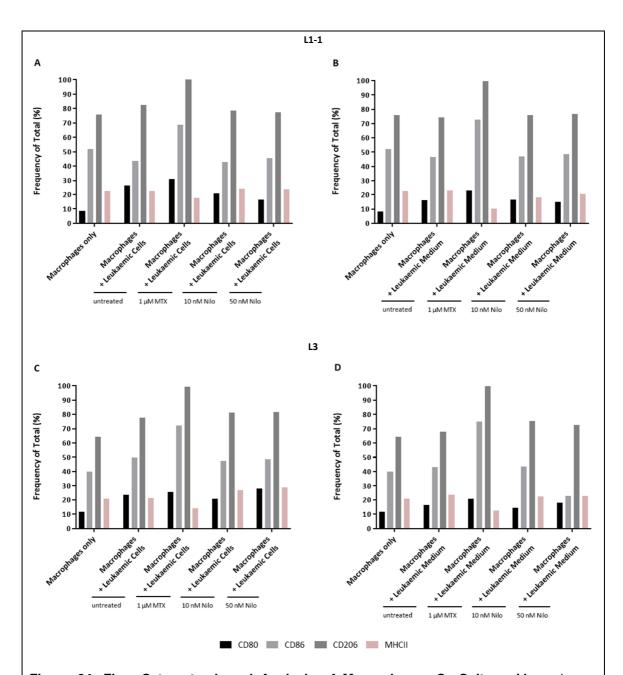


Figure 24. Flow Cytometry based Analysis of Macrophages Co-Culture. Macrophages co-culture was performed with both leukaemic cell lines L1-1 and L3 in two conditions: a juxtacrine setup (A, C) and a paracrine setup (B, D). For the juxtacrine co-culture condition, bone marrow-derived macrophages (BMDM) were cultured with leukaemic cells pre-treated for 24 hours with 10 nM or 50 nM Nilotinib (Nilo). In contrast to that, bone marrow-derived macrophages (BMDM) were cultured with leukaemic cells supernatant (= leukaemic cells medium) of leukaemic cells pre-treated with 10 nM or 50 nM Nilotinib for the paracrine co-culture condition. Comparatively, all conditions were run with leukaemic cells either pre-treated with 1 μ M Mitoxantrone (MTX) or left untreated. After 24 hours of co-culturing, macrophages were analysed by flow cytometry in order to investigate macrophages activation and polarisation based on their surface expression profile of CD80, CD68, CD206 and MHCII. In addition to that, macrophages only without co-culturing were analysed equally. Results are given as frequency of total.

5.4.2 Vaccination Assay

A vaccination assay was performed to examine the potential of Nilotinib to induce immunogenic cell death in-vivo. The first step within the experiment involved the vaccination of mice with killed leukaemic cells functioning as vaccine in order to elicit a protective immune system. For this purpose, leukaemic cells of the cell line L1-1 were incubated with 100 nM Nilotinib for 72 hours or with 1 µM Mitoxantrone for 24 hours and then injected subcutaneously into the left flank of mice. In a second step, after 15 days those mice were re-challenged with live leukaemic cells injected into the tail vein. Thereafter, mice were regularly checked for the development of leukaemia manifesting both as clinical symptoms and as high leukaemic burden of the peripheral blood in flow cytometry (Supplementary Figure S5, page 104). The cohort of experimental mice included 8 wild-type and 8 Gas6 knockout mice, half of which were vaccinated and re-challenged with Nilotinib treated cells and the other half with Mitoxantrone treated cells. The injection of live leukaemic cells failed in a wild-type and a Gas6 knockout mouse from the Nilotinib cohort, which is why these 2 mice were excluded from the final analysis.

As shown in Figure 25, there was no development of leukaemia in the mice after vaccination, confirmed by the lack of leukaemia-typical symptoms and the absent detection of leukaemic cells in the peripheral blood by flow cytometry.

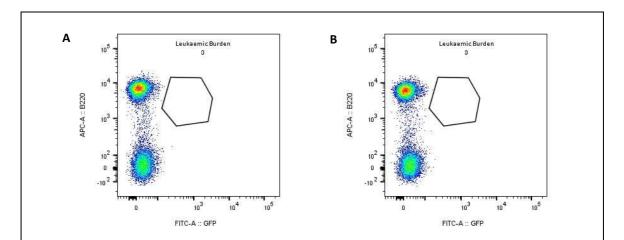


Figure 25. Monitoring the Occurrence of Leukaemia after Vaccination. The development of leukaemia in mice after vaccination was monitored with the help of flow cytometric analysis of the peripheral blood. The leukaemic burden was quantified by the amount of GFP+B220+ leukaemic cells. Exemplary analyses are depicted here illustrating no occurrence of leukaemia in the experimental mice vaccinated with Nilotinib (A) or Mitoxantrone (B) killed leukaemic cells.

Thus, mice were re-challenged with live leukaemic cells in a second step of the vaccination assay. Under regular clinical and flow cytometric monitoring, a development of leukaemia was observed in both cohorts, notably, with leukaemia manifesting more rapidly in the Nilotinib cohort within 8 days than in the Mitoxantrone cohort within 13 days. More detailed, while there were no leukaemic cells detected in the peripheral blood of the Mitoxantrone cohort on day 8, a leukaemic burden of 45.3 % (wt) and 41.9 % (Gas6KO) was found in the peripheral blood of the Nilotinib cohort. In accordance with that, the Nilotinib cohort displayed a distinctly elevated white blood count of 57.7 x 10³/mm³ (wt) and 47.7 x 10³/mm³ (Gas6KO) leukocytes, whereas in the Mitoxantrone cohort a normal white blood count of 7.8 x 10³/mm³ (wt) and 9.2 x 10³/mm³ (Gas6KO) leukocytes was obtained (Figure 26). So once leukaemia occurred in a cohort, all mice of that cohort were sacrificed for a final analysis comprising (1) the flow cytometric assessment of leukaemic burden in bone marrow, peripheral blood and spleen, (2) immunohistochemistry of femur, spleen and subcutaneous tumours, (3) white blood count of peripheral blood and (4) spleen weight.

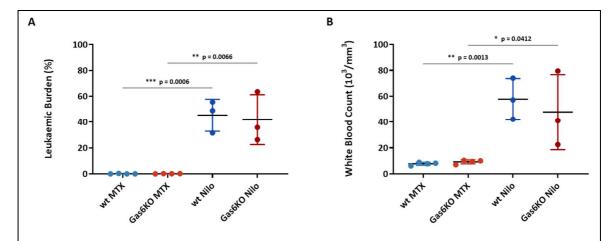


Figure 26. Time of Manifestation of Leukaemia after Re-challenge. The leukaemic burden of peripheral blood was assessed by flow cytometry and expressed as frequency of total (A). The number of leukocytes was measured using white blood count and indicated per cubic millimetre (B). Statistical significance was analysed by unpaired, two-tailed Student t-test. Significance values: * p < 0.05, ** p < 0.01, *** p < 0.001, **** p < 0.001.

The final analysis revealed lower infiltration of leukaemic cells in the Nilotinib cohort as opposed to the Mitoxantrone cohort with respect to the leukaemic burden in all three compartments, bone marrow (wt MTX: 36.6 %, wt Nilo:

32.5 %; Gas6KO MTX: 40.2 %, Gas6KO Nilo: 34.5 %), peripheral blood (wt MTX: 49.3 %, wt Nilo: 45.3 %; Gas6KO MTX: 45.5 %, Gas6KO Nilo: 41.9 %) and spleen (wt MTX: 56.1 %, wt Nilo: 47.5 %; Gas6KO MTX: 31.1 %, Gas6KO Nilo: 11.7 %), which was significant in terms of leukaemic burden of the spleen in the Gas6KO group (Figure 27-29). In terms of the white blood count, a higher number of leukocytes was found in the Nilotinib cohort compared to the Mitoxantrone cohort, as depicted in Figure 28 (wt MTX: 42.7 x 10³/mm³, wt Nilo: 57.7 x 10³/mm³; Gas6KO MTX: 28.0 x 10³/mm³, Gas6KO Nilo: 47.7 x 10³/mm³). Considering the lower infiltration of peripheral blood with leukaemic cells as described above, this indicated an increased fraction of normal leukocytes. In a comparison of the two cohorts regarding spleen weight as a correlate of leukaemic spread, lower spleen weights were measured in the Nilotinib cohort (wt MTX: 0.50 g, wt Nilo: 0.44 g; Gas6KO MTX: 0.47 g, Gas6KO Nilo: 0.34 g), which was significant in the Gas6KO group (Figure 29). Furthermore, all mice of the Nilotinib cohort developed subcutaneous tumours at the injection site in the left flank. To further investigate the consistency of those tumours, histological samples were prepared and stained for HE and GFP revealing a tumour mass of densely packed GFP-positive leukaemic cells, as illustrated in Figure 30.

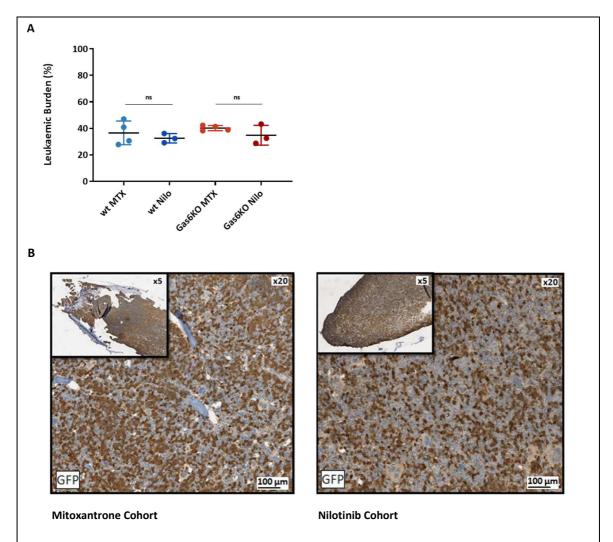


Figure 27. Final Analysis of Bone Marrow. The leukaemic burden of bone marrow was assessed by flow cytometry and given as frequency of total (A). Immunohistochemistry was used to visualise the leukaemic infiltration of bone marrow. Therefore, histological samples of femur were prepared and stained for GFP. Exemplary images of both cohorts are shown here (B). Statistical significance was analysed by unpaired, two-tailed Student t-test. Significance values: $^*p < 0.05$, $^{**}p < 0.01$, $^{***}p < 0.001$, $^{****}p < 0.0001$.

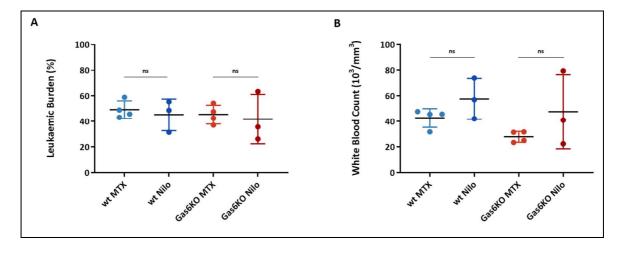


Figure 28. Final Analysis of Peripheral Blood. The leukaemic burden of peripheral blood was assessed by flow cytometry and expressed as frequency of total (A). The number of leukocytes was measured using white blood count and indicated per cubic millimetre (B). Statistical significance was analysed by unpaired, two-tailed Student t-test. Significance values: * p < 0.05, ** p < 0.01, *** p < 0.001, **** p < 0.001, **** p < 0.0001.

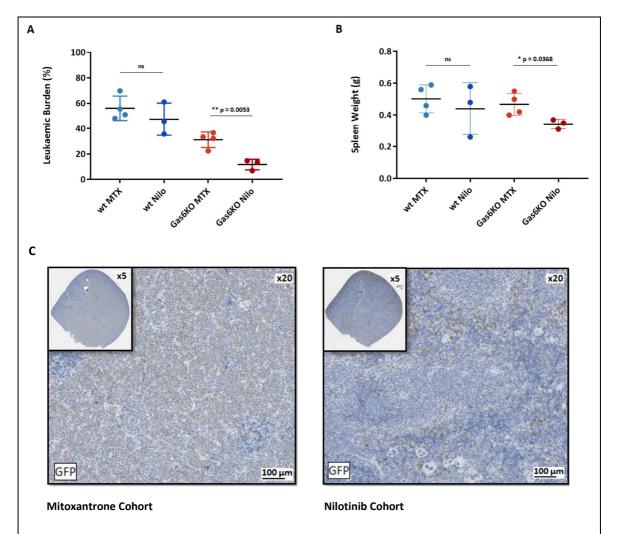


Figure 29. Final Analysis of Spleen. The leukaemic burden of spleen was assessed by flow cytometry and quoted as frequency of total (A). Spleen weight was determined and given in grams (B). To visualise the leukaemic infiltration of spleen, histological samples were prepared and stained for GFP. Representative images of both cohorts are presented here (C). Statistical significance was analysed by unpaired, two-tailed Student t-test. Significance values: * p < 0.05, ** p < 0.01, *** p < 0.001, *** p < 0.001, **** p < 0.0001.

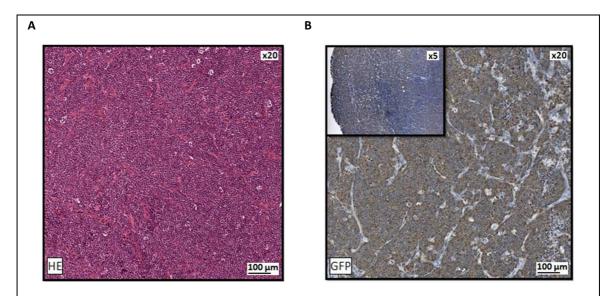


Figure 30. Histological Examination of subcutaneous Tumours of the Nilotinib Cohort. For the histological examination of the subcutaneous tumours that all mice of the Nilotinib cohort developed at the injection site in the left flank, samples were prepared and stained. Representative images of both HE (A) and GFP stain (B) are given here.

6.1 Results on Interpretation and Discussion

6.1.1 Growth Inhibition and Viability Assay

At first, the effects of Nilotinib on cell growth and viability of Ph+ B-ALL cells were studied in a concentration-dependent manner. It was shown that Nilotinib inhibited cell growth in all tested concentrations ranging between 10 nM and 1000 nM, whereby a concentration of 10 nM appeared to be a threshold dose, since at this concentration the cell number continued to increase constantly despite decelerated growth. This was consistent with the effects on cell viability, which dropped significantly above a concentration of 10 nM (L1-1) and 50 nM (L3). However, when exposed to the highest concentration of Nilotinib, neither cell count was below baseline nor viability declined below 37 % implying, especially in comparison with Mitoxantrone, which showed strong and prompt cytotoxicity, that Nilotinib exerts more cytostatic than cytotoxic activity on the leukaemic cells. These results are in accordance with the known pharmacological properties of both drugs. Nilotinib exhibits anti-proliferative properties by competitively binding to the ATP binding site of the BCR-ABL1 kinase¹⁰³, thereby inhibiting cell proliferation¹⁰⁴, whereas Mitoxantrone displays cytotoxic effects by DNA intercalation and inhibition of topoisomerase II. 105,106 In other studies investigating the effects of Nilotinib on BCR-ABL1+ cells, both similar concentrations were tested, and consistent results were obtained. 107-109 Additionally, in other Ph+ B-ALL models, including the BV-173 cell line, and in Ba/F3, that is a popular tool in kinase drug discovery¹¹⁰, a half maximal inhibitory concentration (IC₅₀) of \leq 12 nM Nilotinib was evaluated, ¹⁰⁸ corresponding to the concentration of 10 nM Nilotinib determined in the experiments of this work.

6.1.2 Cell Death Assay

Although Nilotinib predominantly exhibited cytostatic properties, it also appeared to have cytotoxic potential to some extent. To both quantify the extent of cell death and to assess the cell death modality ranging between apoptosis and

necrosis, an Annexin V/Propidium iodide cell death assay was performed. The Annexin V/PI protocol is a commonly used approach for investigating cell death, 111 which allows to distinguish between viable, apoptotic and necrotic cells based on the differences in plasma membrane integrity and permeability. 112,113 Annexin V detects apoptotic cells by its ability to bind to phosphatidylserine, which is translocated from the intracellular side of the plasma membrane to the surface during apoptosis. 114 PI is a nucleic acid intercalator that can enter and stain only permeable cells, that is why it is used for the detection of late apoptotic and necrotic cells with decreased membrane integrity. 115 The cell death assay showed that after 24 hours Nilotinib treated leukaemic cells began to undergo cell death, which was increasing as a function of concentration and exposure time. In terms of cell death modality, Nilotinib induced a bimodal cell death, which was predominantly necrotic in cell line L1-1 and apoptotic in cell line L3. In contrast, apoptosis was triggered in nearly all cells within 4 hours in response to Mitoxantrone exposure. Taken together, these findings are in line with the results observed in the cell growth inhibition and viability assay. Other studies have also demonstrated that inhibition of cell growth by Nilotinib is associated with the induction of cell death, with varying degrees within different BCR-ABL+ cell lines. 108,116 As one aspect of the potential causes for that, cell maturity is cited. 117 In this context, it has been reported that CD34+ hematopoietic stem and progenitor cells do not respond with cell death even at very high concentrations of 5 µM Nilotinib, whereas more mature cells start to undergo cell death. 117,118 Of note, this fact is considered a contributing factor to disease persistence, minimal residual disease and relapse. 119,120 So far, classical mechanisms of resistance, such as point mutations, inadequate active drug uptake or enhanced drug efflux have been disproved to explain the different sensitivity of Ph+ cell lines to Nilotinib and other TKIs. 117,118 Interestingly, studies suggested an interrelation between the expression level of mouse double minute 2 homolog (MDM2) and drug sensitivity, with Nilotinib displaying stronger cytotoxic effects in Ph+ cells with MDM2 overexpression. At this, it is reported that both the tumour suppressor p53 is activated and the anti-apoptotic X-linked inhibitor of apoptosis protein (XIAP) is suppressed by Nilotinib-induced inhibition of MDM2, leading to the induction of apoptotic cell death in BRC-ABL+ cells. 116,121 Moreover, high-dose TKI (HD-TKI) pulse-exposure represents a further aspect regarding Nilotinib-induced cell death in leukaemic cells. Cytotoxic activity of TKIs was found to be enhanced by HD-TKI pulse-exposure due to prolonged intracellular TKI accumulation and retention resulting in an increased fraction of cancer cells initiating apoptosis. 122,123

Besides the aspect of the initiation of cell death per se, which is an obligatory requirement for the induction of ICD, considerations in terms of cell death modality have been controversially discussed. Morphologically, apoptosis is characterized by progressive cell condensation and nuclear fragmentation resulting in the formation of apoptotic bodies surrounded by membranes, whereas necrosis is defined by rupture of the plasma membrane as a result of cell swelling without displaying hallmarks of apoptosis. 124 According to the Nomenclature Committee on Cell Death (NCCD), the definitions of cell death modalities were adjusted from a morphologic to a molecular one. 125 Thereby, apoptosis represents a programmed cell death modality mediated by caspase activation in contrast to necrosis, which is an unregulated, caspase independent form of cell death. 125 Moreover, it was found that necrosis can also occur in a regulated manner, termed necroptosis, involving active disintegration of cellular membranes with limited compartmentalisation of organelles and requiring the activation of receptor-interacting protein kinases 1 (RIPK1) and 3 (RIPK3). 126,127 Initially, when efforts were made to determine the characteristics of ICD, the occurrence of an apoptotic cell death was assumed to be necessary for the immunogenic potential of ICD.91 This might seem peculiar, since on the one hand side apoptotic cell death has long been hypothesised to be of low immunogenicity, 128 and on the other hand side, necrosis is known to be accompanied by the release of DAMPs¹²⁷, which have been identified to actually evoke the immunogenic effects of ICD. 129 Current data indicate that both regulated necrosis and apoptosis play an important role in immunological processes⁸⁸ and may promote analogous effect in the tumour microenvironment including in terms of ICD.¹²⁷ Therefore, it is suggested that different cell death modalities are capable to induce ICD.84,97,101,127 Notably, this assumption seems to be plausible considering that cell death modalities can both occur simultaneously within a cell and merge into each other, as it is known in the form of apoptosis-necrosis switch. 130,131 Even though the exact signalling pathways that determine immunogenic and non-immunogenic cell death have not yet been fully explored, the capacity to elicit ICD seems to be based on the induction of ER stress response.^{97,127} However, it remains to be investigated which modality of cell death maximises the immunogenicity of dying cancer cells within the TME.¹²⁷

6.1.3 ICD Marker Analysis

Next, Nilotinib-treated leukaemic cells were examined for the exhibition of molecular ICD hallmarks. According to the criteria for the assessment of ICD, those comprise translocation of CALR to the cell surface, extracellular secretion of ATP and release of HMGB1 from the nucleus. 91,95,96 The ICD marker analysis was performed with concentrations of Nilotinib at 10 nM and 50 nM. The former to study a possible influence on the level of ICD marker expression at half maximal inhibition, and the latter to investigate potential ICD marker expression at the lowest concentration tested as fully inhibitory. Regarding CALR cell surface exposure, no exposure in Nilotinib-treated leukaemic cells could be detected by flow cytometry. In contrast to that, CALR surface exposure was observed in leukaemic cells exposed to Mitoxantrone after 4 (L3) and 6 (L1-1) hours. Also, in terms of ATP, Nilotinib did not induce extracellular ATP release measured in a bioluminescence-based ATP assay. On the other hand, extracellular ATP release was observed under exposure to Mitoxantrone with absolute peak levels after 4 and 6 hours and, in relation to the untreated control, with relative peak levels after 8 hours. Considering that ATP is released both in the absence of external stimuli and in response to physiological stimuli, 102 this is most likely to give the reason that ATP release was also measured in the untreated controls at early measurement time points. ATP release is seen during mechanical stress, as is the case in experimental manoeuvres. 132 Therefore, conclusions regarding the effective ATP release under exposure to a drug have to be made both in respect to an evaluated baseline (ATP-free water) and in comparison to an untreated negative control, as performed in the ATP assay of this work. In addition, the fact that the levels of ATP release of Nilotinib-treated leukaemic cells were below baseline may not to be considered significant but rather stochastically random in the scope of technical inaccuracy, since the quantified values lay within the 95 % confidence interval. In order to asses HMGB1 release from the nucleus, HMGB1 was determined in both cytosolic extracts and cell culture supernatants of Nilotinib-exposed leukaemic cells by Immunoblotting. Thereby a concentration-dependent release of HMGB1 into the cytosol but not into the cell culture supernatant was found. In the case of leukaemic cells exposed to Mitoxantrone, HMGB1 release was not only observed in the cytosol but also in the supernatant. This is consistent with the results from the cell death assay taking into account that the release of HMGB1 is the process of passive translocation from the nucleus via the cytosol into the extracellular space in response to several forms of cell death, including apoptosis, necrosis and necroptosis by dying or dead cells. 133-135 It seems that HMGB1 release from Nilotinib exposed cells has not yet progressed as far after 12 hours as in those cells the initiation of cell death was just beginning, which is why HMGB1 was only detected in the cytosol and not yet extracellularly. In contrast, the leukaemic cells treated with Mitoxantrone were already completely succumbed to cell death at this time, so that HMGB1 was found in both the cytosol and the supernatant.

Taken together, the results support the assumption that Nilotinib is not capable of inducing immunogenic cell death in Ph+ leukaemic cells in-vitro considering the use of techniques referred as standard⁹¹ as well as by comparison with Mitoxantrone as a positive control. In fact, the orchestration of all three molecular determinants together is considered a required condition for the eligibility of ICD, whereas the exposure of only one or two determinants is viewed as insufficient. 91,95,96 However, the different requirements for the immunogenicity of cell death have not been fully investigated. Although CALR, ATP and HMGB1 are considered key hallmarks, several other processes have been linked to the immunogenicity of cell death, involving release of the heat shock proteins (HSP) 70 and 90, unfolded protein response, secretion of CXC motif chemokine ligand 10 (CXCL10), autophagy induction, release of annexin A1 (ANXA1) as well as inflammasome and TLR3 signalling.84 Besides, the exact mechanisms that underlie the release of the immunogenic hallmarks remain to be completely elucidated.⁹⁷ The process of CALR exposure to the cell surface is suggested to take place in dependence of the stage of cell death, via secretory pathway, phosphatidylserine-based or as a result of a general exposure of chaperons from the ER. 136 In terms of ATP, the release is reported to occur by secretory pathway, pannexin 1- or autophagy-based secretion or as passive release due to plasma membrane permeabilisation. 136 The release of HMGB1 is revealed least so far

and is considered to be operated in autophagy- or caspase activity-dependent manner next to a passive release during plasma membrane permeabilisation.¹³⁷ Moreover, the full spectrum of aspects regarding the effects exhibited by the molecular ICD determinants within ICD are still under investigation, especially in terms of the role of HMGB1.^{136,137} HMGB1 shows cytokine, chemokine and growth factor activity that acts in multiple biological processes, thus making HMGB1 an important target in inflammatory diseases, cancer and immune response.¹³⁸⁻¹⁴⁰ Remarkably, HMGB1 has been found to cause different and even adverse effects within the immune system depending on the redox state induced by the cell or the extracellular environment.^{141,142} Referring to this, the activity of HMGB1 shifts between chemoattractant (reduced redox state), cytokine-inducing (intermediate redox sate) and nonimmune activity (oxidised redox state).¹⁴³

6.1.4 Macrophages Co-Culture

To further assess Nilotinib for its functional potential in inducing ICD, a macrophage co-culture was performed, as macrophages play a key role in mediating the immunological effects of ICD by antigen presentation and T cell activation. The prerequisite for this is the activation and maturation of macrophages towards an immune-stimulating phenotype. Therefore, bone marrow-derived macrophages were investigated for their activation and polarisation in response to co-cultivation with Nilotinib pre-treated leukaemic cells in both a juxtacrine and paracrine setup. The experiment was set on the background that macrophages display polarisation in reaction to environmental stimuli towards an M1 or M2 phenotype, that differ in immunogenic function and is characterised by a specific cytokine and surface expression profile. Referring to this, M1 macrophages represent an immune-stimulating, tumour-suppressive phenotype with pro-inflammatory cytokine production, while M2 macrophages exhibit immune-suppressive, tumour-promoting features associated with an anti-inflammatory cytokine profile. Referois

Regarding macrophages co-culture with Nilotinib-treated leukaemic cells, no relevant shift in terms of macrophages activation and polarisation was observed in either the juxtacrine or paracrine setup compared to the corresponding co-culture conditions of untreated leukaemic cells. In contrast to that, macrophages

co-cultured with Mitoxantrone-incubated leukaemic cells were found to exhibit an average upregulation of CD80, CD86 and of CD206 next to a downregulation of MHCII. The findings of this more functional approach are in line with the results of the molecular ICD marker analysis suggesting that leukaemic cells exposed to Mitoxantrone but not to Nilotinib exhibit increased immunogenicity indicated by the upregulated expression of CD80 and CD86 in macrophages co-cultured with Mitoxantrone-treated leukaemic cells, as CD80 and CD86 represent costimulatory molecules on activated macrophages mediating activation of cytotoxic T cells. 144 However, with regard to macrophage polarisation, the results are not as distinct as in terms of macrophage activation. In the context of macrophage polarisation, CD80 and CD86 are related to a M1 phenotype, while CD206 is considered a typical marker for M2 macrophages.⁶⁸ The simultaneous upregulation of CD206 might therefore initially seem surprising but may be explained by the fact that tumour-associated macrophages are assumed to be not explicitly attributable to either phenotype by expressing overlapping M1/M2 properties.^{71-74,145} This fact becomes evident with respect to MHCII, which is not only a marker of macrophage activation but also of macrophage polarisation, where it is attributed to both the M1 and M2 phenotype. 72,145 Furthermore, in the context of macrophage polarisation, not only the coexistence of M1 and M2 macrophages within the TME but also the shift in macrophages polarisation status over the time in response to the TME have been observed. 145,146 Moreover, TAMs are found to exist in distinct TAM subsets residing in different tumour compartments, exerting specialised functions and displaying a certain surface expression profile.147 Taken together, these aspects underline both the heterogeneity and the plasticity of TAMs emphasising the approach of classifying different TAMs subsets according to their function within the TME rather than in terms of their M1/M2 phenotype. 148-150 As an example, the observation that hypoxia modifies macrophages polarisation and function can be cited. TAMs in hypoxic areas, such as the tumour-stroma border or the tumour centre, are sessile and pro-angiogenic macrophages that are CD206+MHCIIlow and in general more M2-like orientated, while TAMs in oxygen-rich, perivascular areas exhibit migratory and pro-metastatic features associated with an CD206-MHCII^{high} expression profile and a rather M1-like phenotype. 147,151

In terms of the co-culture setup, effects on activation and polarisation of macrophages co-cultured with Mitoxantrone-treated leukaemic cells were found both in the juxtacrine and the paracrine co-culture condition. Juxtracrine signalling represents a direct signalling mechanism via close cell-cell interaction, whereas paracrine signalling is an indirect signalling mechanism that is mediated by the secretion of signalling molecules form neighbouring cells reaching the target cells via diffusion. 152 The paracrine co-culture condition was performed with the aim to demonstrate that the effects on macrophage activation and polarisation are not only based on unspecific cell-cell contact between leukaemic cells and macrophages but in fact are due to the release of DAMPs from dying tumour cells succumbing to ICD. Interestingly, the effects were more pronounced in the paracrine than in the juxtacrine setup. Concerning this observation, there are almost no data available. In the study of Chang et al. 153 it was seen that the effects on macrophage activation were more distinct in the paracrine setup at early time points of co-culture, while juxtacrine signalling dominated at later time suggesting that the predominant mechanism of signalling might be in relation to the duration of co-cultivation. This may appear plausible considering that once the process of cell death is completed the release of DAMPs dries up.

6.1.5 Vaccination Assay

Finally, to evaluate the potential of Nilotinib to induce immunogenic cell death in-vivo, a vaccination assay was performed that is defined as gold-standard approach to asses bona fide ICD.⁹⁸ Therefore, leukaemic cells were treated with the drug in-vitro and then injected into the left flank of mice for vaccination. After 15 days, mice were re-challenged with live leukaemic cells injected into the tail vein. In the vaccination assay a vaccination in the sense of an active immunisation is aimed to be evoked. Thereby, in response to immortalised pathogens, the immune system is stimulated to build up a pathogen-specific immunocompetence with an immune memory so that a subsequent exposure to living pathogens leads to an effective elimination based on the established immune response.¹⁵⁴

In order to make leukaemic cells immortalised and potentially immunogenic by the induction of ICD, they were exposed to Nilotinib and Mitoxantrone, respectively. After vaccination, leukaemia did not occur in mice as expected, but all mice of the Nilotinib cohort developed subcutaneous tumours at the injection site. Histological staining revealed tumours of densely packed leukaemic cells, indicating that Nilotinib was not sufficient to completely kill the leukaemic cells. This was to be assumed considering that Nilotinib predominantly exerts antiproliferative rather than cytotoxic effects unlike Mitoxantrone.

After rechallenge with live leukaemic cells, leukaemia occurred in both cohorts manifesting more rapidly in the Nilotinib cohort within 8 days than in the Mitoxantrone cohort within 13 days. This suggests that Nilotinib is not potent in eliciting an anti-leukaemic immune response by the induction of ICD in-vivo reinforcing the observations of the in-vitro experiments in which Nilotinib was shown to not meet the criteria of a bona fide ICD inducer. With regard to Mitoxantrone as a well-known inductor of ICD, the following considerations may be mentioned to explain why leukaemia developed in mice after rechallenge. One aspect is that Ph+ B-ALL is a highly aggressive leukaemia model, which is clinically emphasised by the fact that Philadelphia chromosome positive B-ALL is the most aggressive ALL subtype with a long-term survival rate in the range of only 30 %.155 Therefore, the effects of ICD induction by Mitoxantrone might not be sufficient to prevent but at least delay the occurrence of leukaemia as seen in the vaccination assay. Furthermore, the in-vivo vaccination assay is designed for the application in solid tumours and has not yet been investigated in the context of non-solid leukaemia according to current data, rendering its implementation and interpretation challenging.

Notably, lower leukaemic burden was observed in the Nilotinib cohort compared to the Mitoxantrone cohort with respect to all three compartments involving bone marrow, peripheral blood and spleen. Additionally, in accordance with this, lower spleen weights were found in the Nilotinib cohort. These findings regarding leukaemic spread were most pronounced in spleen and here were found significant in the Gas6KO but not in the wild-type group. This may support the observations consistently seen in the in-vivo experiments of our research group, that Nilotinib synergizes with a Gas6-deficient environment. Moreover, it was observed by our group that the anti-leukaemic effects of Gas6 deletion in the microenvironment diminish with progression of leukaemia, initially at the site of origin in the bone marrow and finally in peripheral organs such as the spleen.

This might be the reason why in the vaccination assay the synergistic impact between Gas6KO and Nilotinib was found to be significant only in spleen, considering that leukaemia was already quite progressed in the mice with an infiltration of 35 % in bone marrow and 42 % in peripheral blood.

Besides, in white blood count a higher number of leukocytes was measured in the Nilotinib cohort compared to the Mitoxantrone cohort, indicating an increased fraction of normal leukocytes taking into account the lower leukaemic burden in peripheral blood. This finding could be interesting to be further investigated, taking into consideration that, as mentioned before, in several studies chemotherapeutic agents including TKIs have been reported to improve antitumour immune response by modulating the tumour immune microenvironment, such as by increasing the infiltration of cytotoxic CD8+ T cells.⁷⁸

6.1.6 Conclusion

In the in-vivo experiments of our research group it has been consistently observed that the use of Nilotinib but not of other chemotherapeutic agents enhances the anti-leukaemic immunity mediated by a deletion of Gas6 in the TME, which is why I was assigned to examine this observation in the present work. Against the background of increasing importance of chemotherapeutic agents as potent modulators of a dysregulated TME, it was hypothesized that Nilotinib may synergize with a Gas6-deficient environment by inducing ICD. Thus, according to the criteria for the assessment of a potential ICD inducer, Nilotinib was evaluated for the molecular and functional determinants of ICD in in-vitro and in-vivo experiments.

So far, the induction of ICD in leukaemic cells has been found in several studies. 156-162 However, there is no data available investigating the induction of ICD in leukaemic cells by Nilotinib. In the present work it was shown that Nilotinib induces cell death in leukaemic cells in concentration-dependent manner that occurs bimodally in terms of cell death modality ranging between apoptosis and necrosis. Furthermore, it was found that Nilotinib-induced cell death is not accompanied by CALR exposure and ATP release, but is associated with the release of HMGB1. No relevant shift in terms of macrophages activation and polarisation was observed in either a juxtacrine or paracrine setup of a

macrophages co-culture experiment with Nilotinib-treated leukaemic cells. In consistency with the results obtained in the in-vitro experiments, Nilotinib was not potent to elicit a protective immune response in the in-vivo vaccination assay. Conclusively, Nilotinib was identified to not qualify as a bona fide ICD inducer. This gives rise to the question by which mechanism Nilotinib instead synergistically contributes the anti-leukaemic effects of a Gas6-deficient environment. In this context, the role of Nilotinib-induced cell death and HMGB1 release may be of relevance for further investigation.

6.2 Limitations and Options

A critical review of the used methods and the obtained results is crucial to ensure quality and reliability of scientific work. Therefore, it is necessary to examine one's own work with respect to this.

By growth inhibition and viability assay it was shown that Nilotinib displays concentration- and exposure time-dependent efficacy. This relation to concentration and exposure time was further affirmed in the cell death assay. To assess Nilotinib as a potential ICD inducer under strict experimental conditions, the experiments were performed using Nilotinib at a concentration of 50 nM, which gives the lowest concentration tested with full inhibitory activity. Additionally, experiments were carried out using 10 nM Nilotinib to study a possible influence on the level of ICD hallmark expression at half maximal inhibition. As Nilotinib could not be identified as a bona fide inducer of ICD, this should be confirmed for both higher concentrations and longer exposure times in order to validate this finding.

To examine the capacity of Nilotinib to induce ICD in-vitro, ICD marker analysis was performed using techniques referred as standard according to the consensus guidelines for the detection of immunogenic cell death. Nonetheless, the use of additional methods allows the results obtained to be substantiated. In this respect, the evaluation of CALR exposure and HMGB1 release can be complemented by fluorescence microscopy. Moreover, extracellular HMGB1 can be assessed using enzyme-linked immunosorbent assay (ELISA). Regarding CALR, exposure to the cell surface in leukaemic cells incubated with

Mitoxantrone as positive control was not pronounced in flow cytometry indicating that the staining protocol may be optimized. In this context, modifications in terms of antibody dilution and exposure time, as well as the testing of other secondary antibodies, with which fewer non-specific background staining is obtained, could improve the experimental design.

Antigen presenting cells, including dendritic cells and macrophages, play a key role in ICD, as CALR, ATP and HMGB1 released by dying tumour cells succumbing to ICD bind to those cells mediating anti-tumour immune response. 95 The studies of our research group suggested that TAM-expressing macrophages are the essential mediators of Gas6 immune effects. For this reason, in the coculture experiment the effects of Nilotinib in the context of ICD were assessed on macrophages and not, as primarily common, on dendritic cells. Based on the surface expression profile macrophages were investigated for their activation and polarisation. As discussed above, an evaluation concerning macrophage polarisation by surface expression markers alone is not sufficient, as in this way TAMs cannot be assigned to an M1/M2 phenotype due to overlapping characteristics.⁷¹⁻⁷⁴ Therefore, a cytokine profile assay including IL-1B, IL-6, TNF- α , IL-10 and TGF- β performed by ELISA may provide a helpful contribution. Besides, both phagocytosis and antigen presentation capacity of macrophages could be studied by flow cytometry using DiO-labelled or OVA peptide SIINFEKLexpressing leukaemic cells, respectively. In addition, methods and addressed aspects of the macrophage co-culture experiment can be transferred to the investigation of dendritic cells for completing the picture.

Finally, the potential of Nilotinib to induce ICD in-vivo was evaluated in a vaccination assay. Unlike in the other experiments, no negative control with mice vaccinated and re-challenged using untreated leukaemic cells was performed from questions of rational resource management. With regards to the intention to elicit a protective immune response by vaccination with killed leukaemic cells that have succumbed to ICD, the question arises about how many cells have to be applied. Moreover, the efficacy of vaccination needs to be examined for a dependence on the time point of rechallenge in order to ensure optimal conditions. As outlined above, the in-vivo vaccination assay is designed for the application in solid tumours rendering its implementation and interpretation challenging in terms of leukaemia. In addition to that, all mice of the Nilotinib

cohort developed tumours at the injection site because Nilotinib was insufficient to completely kill leukemic cells due to displaying predominantly antiproliferative rather than cytotoxic activity. In order to circumvent these issues, an ex-vivo immune stimulation approach may be suggested. In this case, instead of the vaccination with leukaemic cells killed by the potential ICD inducer, bone marrow-derived macrophages and/or dendritic cells including their supernatant would be applied, which have been co-cultured with leukaemic cells pre-treated with the drug of interest. In the final analysis of the vaccination assay leukaemic burden was assessed. Considering that chemotherapeutic agents have been shown to promote anti-tumour immune response by modulating the tumour immune microenvironment, ⁷⁶⁻⁷⁸ studying the impact on the immune cell infiltrate could be a contributing addition to the final analysis. Since the induction of ICD results in a tumour antigen-specific T cell immune response, the frequency of both total and INF-γ producing T cells as well as the CD8+/CD4+ and CD8+/Treg ratio might be an investigation of interest.

6.3 Perspectives

Besides cardiovascular disease, cancer is one of the major causes of death.¹ Therefore, highest efforts are required concerning effective treatment of cancer. Since conventional therapy concepts remain suboptimal, alternative approaches need to be investigated. Tumour cells display various mechanisms to promote tumour growth and metastasis including tumour immune escape. In this way cancer evades immunosurveillance and cannot be detected and eliminated by the immune system.⁸² The efficacy of chemotherapy has long been mainly attributed to tumour cell toxicity while immune modulating effects were neglected. But in fact, recent advances emphasise the use of chemotherapeutic agents, including TKIs, for normalising or re-educating the dysfunctional TME in the context of cancer treatment.⁷⁸ Accumulating evidence suggests that reactivation of comprised immunosurveillance in the tumour microenvironment is a critical approach for improved disease outcome and patient survival that can be achieved by the use of agents inducing immunogenic cell death in tumour cells.¹⁰⁰ However, the majority of chemotherapeutic agents are incapable of inducing ICD.

In order to restore immunogenicity of cell death, the co-administration of chemicals is conceived that convert non-ICD inducer into bona fide inducer of ICD by boosting, eliciting or substituting hallmarks of ICD, such as tumour cell death, CALR exposure, ATP secretion, HMGB1 release and Type I IFN production (Figure 31).96,101 Optimising the immunological potential of conventional chemotherapy is of particular importance in view of the acquired resistance to TKIs that represent the major clinical challenge of TKI treatment, as rapid but not durable tumour response is achieved.⁷⁸

In conclusion, emerging evidence supports the concept that conventional chemotherapeutic agents can display immunomodulatory effects within the TME and synergise with immunotherapeutic approaches. Testing anticancer drugs for their capability to induce ICD and establishing optimized combinatorial therapeutic regimes is a promising strategy to enhance anti-tumour immunity and to improve therapeutic efficacy with the aim of providing cancer patients with the best possible treatment in the future.

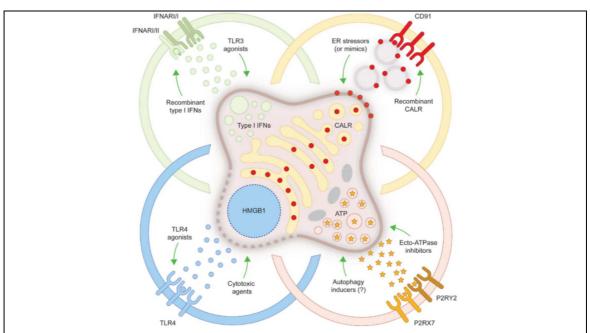


Figure 31. Strategies for the Induction of ICD. The induction of ICD relies on the release of DAMPs form dying tumour cells succumbing to ICD. CALR exposure, ATP secretion and HMGB1 release are considered key hallmarks of ICD mediating anti-tumour immunity upon binding to antigen presenting immune cells, which results in an INF-γ-mediated tumour antigen-specific T cell response. If any of these DAMPs is not released, dying cancer cells cannot be perceived as immunogenic by the immune system. Several strategies have been conceived aiming to convert non-immunogenic cell death into bona fide ICD in order to restore immunogenicity. From Bezu *et al.*, 2015 (101).

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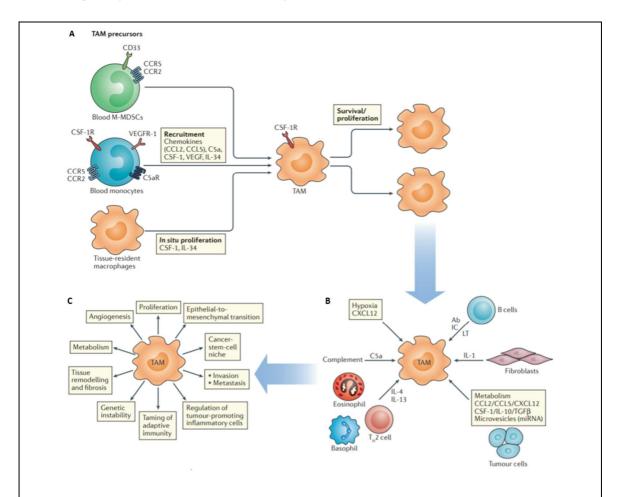
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8.1 Supplementary Figure S1: The Role of Tumour-associated Macrophages in Tumour Progression



Supplementary Figure S1. The Role of Tumour-associated Macrophages in Tumour Progression. Macrophages populating in the microenvironment around the tumour are termed as tumour-associated macrophages (TAMs). Both circulating monocytes and monocyte-related myeloid-derived suppressor cells (M-MDSCs) derived from bone marrow as well as tissue-resident macrophages of embryonic origin represent TAM precursors contributing to the TAM population, that are recruited or stimulated to in situ proliferation, respectively, in response to diverse chemoattractants (A). A variety of stimuli within the tumour microenvironment (TME), including those of tumour and immune cells, fibroblast and metabolism, have impact on TAM polarisation (B). Although TAMs cannot be explicitly assigned to a M1 or M2 phenotype, they are often associated with a tumour-promoting M2-like phenotype. The effects of TAM in the context of tumour progression address all aspects of tumour cell biology and function as depicted in (C). Hence, anti-cancer strategies are aimed to target macrophage plasticity for re-educating TAMs towards a tumour-suppressive phenotype. From Mantovani *et al.*, 2017 (72).

8.2 Table 1: Consumable Material

Product Name	Manufacturer
BD Microfine+ U100 Insulin Syringes,	BD Bioscience, Franklin Lakes,
0.30 mm x 8 mm, 0.5 mL	NJ, US
Cell Culture Multiwell Plate, 6 Well	Greiner Bio-One,
	Kremsmünster, AT
Cell Culture Multiwell Plate, 12 Well	Greiner Bio-One,
	Kremsmünster, AT
Cell Culture Multiwell Plate, 96 Well, white	Greiner Bio-One,
	Kremsmünster, AT
Cellstar Serologic Pipettes, 5 mL, bulk,	Greiner Bio-One,
sterile	Kremsmünster, AT
Cellstar Serologic Pipettes, 5 mL, single	Greiner Bio-One,
packed, sterile	Kremsmünster, AT
Cellstar Serologic Pipettes, 10 mL, bulk,	Greiner Bio-One,
sterile	Kremsmünster, AT
Cellstar Serologic Pipettes, 10 mL, single	Greiner Bio-One,
packed, sterile	Kremsmünster, AT
Cellstar Serologic Pipettes, 25 mL, bulk,	Greiner Bio-One,
sterile	Kremsmünster, AT
Cellstar Serologic Pipettes, 25 mL, single	Greiner Bio-One,
packed, sterile	Kremsmünster, AT
Falcon Conical Centrifuge Tubes, 15 mL	Corning Life Sciences,
	Amsterdam, NL
Falcon Conical Centrifuge Tubes, 50 mL	Corning Life Sciences,
	Amsterdam, NL
Filter Cap Cell Culture Flasks, 50 mL,	Greiner Bio-One,
25 cm ²	Kremsmünster, AT
Filter Cap Cell Culture Flasks, 250 mL,	Greiner Bio-One,
75 cm ²	Kremsmünster, AT
Filter Cap Cell Culture Flasks, 550 mL,	Greiner Bio-One,
175 cm ²	Kremsmünster, AT

Teflon Bag, Teflon-Coated Cell Culture	American Durafilm, Holliston,
Bag	MA, US
TipOne 10 μl Pipette Tips, red	STARLAB, Hamburg, DE
TipOne 200 μl Pipette Tips, yellow	STARLAB, Hamburg, DE
TipOne 1000 μl Pipette Tips, blue	STARLAB, Hamburg, DE
Tubes, 5 ml, 75 mm x 13 mm, PS	Sarstedt, Nümbrecht, DE
UpCell MultiDish Cell Culture Dish, 12	Thermo Fisher Scientific,
Well	Waltham, MA, US

8.3 Table 2: Common Reagents and Chemicals

Manufacturer
Manuacturer
Carl Roth, Karlsruhe, DE
US
Sigma-Aldrich, St. Louis, MO,
US
Sigma-Aldrich, St. Louis, MO,
US
Carl Roth, Karlsruhe, DE
US
Sigma-Aldrich, St. Louis, MO,
US
Carl Roth, Karlsruhe, DE

Rotiphorese Gel 30, 30 % acrylamide and	
bisacrylamide stock solution (ratio 37.5:1)	Carl Roth, Karlsruhe, DE
(3029)	
Sodium chloride ≥ 99 %	Sigma-Aldrich, St. Louis, MO,
(S9888)	US
Sodium dihydrogen phosphate ≥ 99.0 %	Sigma-Aldrich, St. Louis, MO,
(S8282)	US
Sodium dodecyl sulphate 98 %	Sigma-Aldrich, St. Louis, MO,
(862010)	US
Tetramethylethylenediamine ~ 99 %	Sigma-Aldrich, St. Louis, MO,
(T9281)	US
Tris base	Sigma-Aldrich, St. Louis, MO,
(TRIS-RO)	US
Tween 20, Polyoxyethylene (20) sorbitan	Sigma-Aldrich, St. Louis, MO,
monolaurate	US
(P2287)	

8.4 Table 3: Reagents and Chemicals for Cell Culture

Product Name (Catalogue Number)	Manufacturer
Dulbecco's Modified Eagle Medium, DMEM Medium (11960044)	Thermo Fisher Scientific, Waltham, MA, US
Foetal Bovine Serum, FBS (10437028)	Thermo Fisher Scientific, Waltham, MA, US

Leukaemic Cell Line Medium

- RPMI 1640 Medium
- 15 % Foetal Bovine Serum (FBS)
- 1 % L-Glutamine (200 mM)
- 1 % Pen Strep

L-Glutamine (200 mM), 100x	Thermo Fisher Scientific,
(25030081)	Waltham, MA, US
Macrophages Medium	
DMEM Medium	
 10 % Foetal Bovine Serum (FBS) 	
• 1 % L-Glutamine (200 mM)	
1 % Pen Strep	
• 100 μg/mL Recombinant Mouse M-C	SF
Pen Strep, Penicillin (10000 U/mL)-	Thermo Fisher Scientific,
Streptomycin (10000 μg/nL)	Waltham, MA, US
(15140122)	Waltham, WA, OO
Recombinant Mouse M-CSF	BioLegend, San Diego, CA, US
(576406)	BioLegena, oan biego, o/t, oo
RPMI 1640 Medium	Thermo Fisher Scientific,
(21870076)	Waltham, MA, US
Trypan Blue Solution	Sigma-Aldrich, St. Louis, MO,
(T8154)	US

8.5 Table 4: Reagents and Chemicals for Flow Cytometry

Product Name (Catalogue Number)	Manufacturer
BD Horizon Brilliant Stain Buffer	BD Bioscience, Franklin Lakes,
(563794)	NJ, US
BD Pharm Lyse Lysing Buffer	BD Bioscience, Franklin Lakes,
(555899)	NJ, US
Dulbecco's Phosphate Buffered Saline, 1x,	Thermo Fisher Scientific,
DPBS	·
(14190094)	Waltham, MA, US
Dulbecco's Phosphate Buffered Saline,	Thermo Fisher Scientific,
10x, DPBS	·
(14200075)	Waltham, MA, US

FACS Buffer

- Dulbecco's Phosphate Buffered Saline (DPBS)
- 2 % Foetal Bovine Serum (FBS)

Staining Buffer

- Dulbecco's Phosphate Buffered Saline (DPBS)
- 2 % Foetal Bovine Serum (FBS)

8.6 Table 5: Reagents and Chemicals for Western Blot

Product Name (Catalogue Number)	Manufacturer
Amersham Hyperfilm ECL High performance chemiluminescence film (28906836)	GE Healthcare Life science, Chalfont St Giles, UK
Amersham Protran Premium 0.2 µm NC Nitrocellulose Blotting Membrane (10600001)	GE Healthcare Life science, Chalfont St Giles, UK

Blocking Buffer

- TBS-T Buffer
- 2.5 % or 5 % Skim milk powder

Laemmli Sample Buffer, 4x	Bio-Rad Laboratories, Hercules,
(1610747)	CA, US
Phosphatase Inhibitor Cocktail	Thermo Fisher Scientific,
(78428)	Waltham, MA, US

Polyacrylamide Gel

- a) 15 % Resolving Gel (10 mL, 1 gel)
 - 5 mL 30 % Acrylamide mix
 - 2.5 mL 1.5 M Tris base (pH 8.8)
 - 100µL 10 % Sodium dodecyl sulphate (SDS)
 - 100 μL 10 % Ammonium persulfate (APS)
 - 4 μL Tetramethylethylenediamine (TEMED)
 - 2.3 mL Distilled water

b) 5 % Stacking Gel (3 mL, 1 gel)

- 500 μL 30 % Acrylamide mix
- 380 μL 12.5 % 1.0 M Tris base (pH 6.8)
- 30 μL 10 % Sodium dodecyl sulphate (SDS)
- 30 µL 10 % Ammonium persulfate (APS)
- 3 µL Tetramethylethylenediamine (TEMED)
- 2.1 mL Distilled water

Precision Plus Protein WesternC Blotting Standards (1610376)	Bio-Rad Laboratories, Hercules, CA, US
Protease Inhibitor Cocktail, EDTA-free	Thermo Fisher Scientific,
(78425)	Waltham, MA, US
RIPA Lysis and Extraction Buffer	Thermo Fisher Scientific,
(89900)	Waltham, MA, US

Running Buffer (pH 8.3)

- 25 mM Tris base
- 192 mM Glycine
- 0.1 % Sodium dodecyl sulphate (SDS)
- Distilled water

Skim milk powder	Sigma-Aldrich, St. Louis, MO,
(1.15363)	US

TBS-T Buffer (pH 7.2 - 7.6)

- 20 mM Tris base
- 150 mM Sodium chloride (NaCl)
- 0.05 % Tween 20
- Distilled water

Wet Transfer Buffer (pH 8.3)

- 25 mM Tris base
- 190 mM Glycine
- 20% Methanol
- Distilled water

Western Blotting Filter Paper	Thermo Fisher Scientific,
(84784)	Waltham, MA, US

8.7 Table 6: Reagents and Chemicals for Haematoxylin-Eosin Stain and Immunohistochemistry

Product Name (Catalogue Number)	Manufacturer
BOND Primary Antibody Diluent (AR9352)	Leica Biosystems, Wetzlar, DE
Cover glass (3800145ACS)	Leica Biosystems, Wetzlar, DE
CellTexx Mounting Medium (19360)	Englebrecht Medizin- und Labortechnik, Edermünde, DE

Decalcification Buffer

- 900 mL 0.5 M Ethylenediaminetetraacetic acid (EDTA, pH 8.0)
- 100 mL 10x NaPBS (pH 7.4)
 - o 150 mM Sodium chloride (NaCl)
 - o 10.8 mM Disodium hydrogen phosphate (Na₂HPO₄)
 - o 1.6 mM Sodium dihydrogen phosphate (NaH₂PO₄)
 - o 100 mL Distilled water

EM-400 Embedding Medium Paraffin (3801320)	Leica Biosystems, Wetzlar, DE
IP Cassette III (38440210)	Leica Biosystems, Wetzlar, DE
Roti-Histofix 4 % (P087)	Carl Roth, Karlsruhe, DE
Snowcoat Pearl Microscope Slides (3809299)	Leica Biosystems, Wetzlar, DE

8.8 Table 7: Detection Kit for Extracellular ATP Release

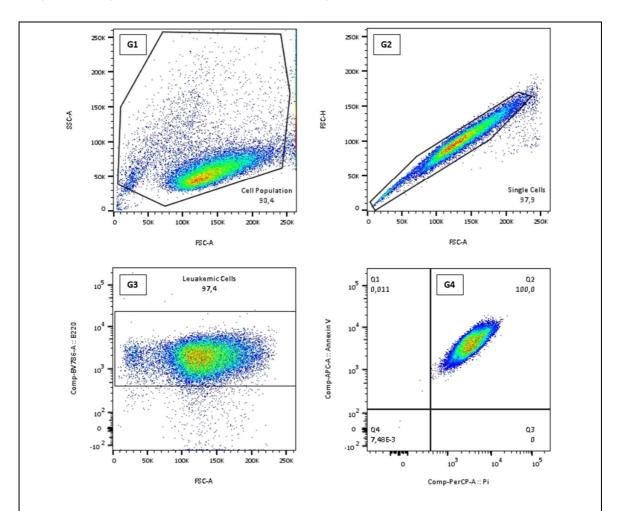
Product Name	Manufacturer
(Catalogue Number)	
ENLITEN ATP Assay System	
Bioluminescence Detection Kit for ATP	Promega, Mannheim, DE
Measurement	r Tomega, Marinielin, DE
(FF2000)	

8.9 Table 12: Devices and Software

Product Name	Manufacturer	
Axio Imager 2 Research Microscope	Carl Zeiss, Oberkochen, DE	
PD I SPEartages Flow Cytomotor	BD Bioscience, Franklin Lakes,	
BD LSRFortessa Flow Cytometer	NJ, US	
BINDER CO ₂ Incubator C170	BINDER, Tuttlingen, DE	
Heraeus Megafuge 40R Centrifuge	Thermo Fisher Scientific,	
Tieraeus Megaluge 4011 Gerilliuge	Waltham, MA, US	
Heraeus Pico 17 Centrifuge	Thermo Fisher Scientific,	
Tieraeus i ico i / Gentinuge	Waltham, MA, US	
Leica ASP300 S Fully Enclosed Tissue	Leica Biosystems, Wetzlar, DE	
Processor		
Leica BOND-MAX Fully Automated IHC	Leica Biosystems, Wetzlar, DE	
and ISH		
Leica CV5030 Fully Automated Glass	Leica Biosystems, Wetzlar, DE	
Coverslipper		
Leica RM2235 Manual Rotary Microtome	Leica Biosystems, Wetzlar, DE	
Leica ST5010 Autostainer XL	Leica Biosystems, Wetzlar, DE	
LUMIstar OPTIMA Microplate	BMG Labtech, Ortenberg, DE	
Luminometer	Divid Labledii, Orleiberg, DL	
scil Vet abc hematology analyzer	scil animal care company,	
Son vot abo nematology analyzer	Gurnee, IL, US	
ViCell XR Cell Counter	Beckman Coulter, Brea, CA, US	

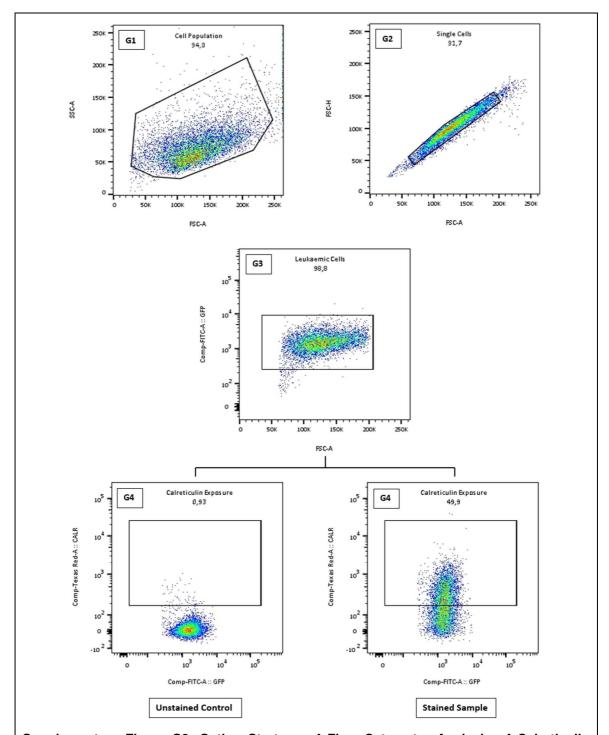
X-ray film developer machine XR24 Pro	Dürr Dental, Bietigheim- Bissingen, DE
AxioVision Microscopy software Vision SE64 4.9	Carl Zeiss, Oberkochen, DE
BD FACSDiva software Version 8.0.1	BD Bioscience, Franklin Lakes, NJ, US
FlowJo software Version 10.5.3	BD Bioscience, Franklin Lakes, NJ, US
GraphPad Prism software Version 8	GraphPad Software, San Diego, CA, US
Lumistar 32 software Version 1.7	BMG Labtech, Ortenberg, DE
Vi-Cell XR software Version 2.04	Beckman Coulter, Brea, CA, US

8.10 Supplementary Figure S2: Gating Strategy of Flow Cytometry based Cell Death Assay



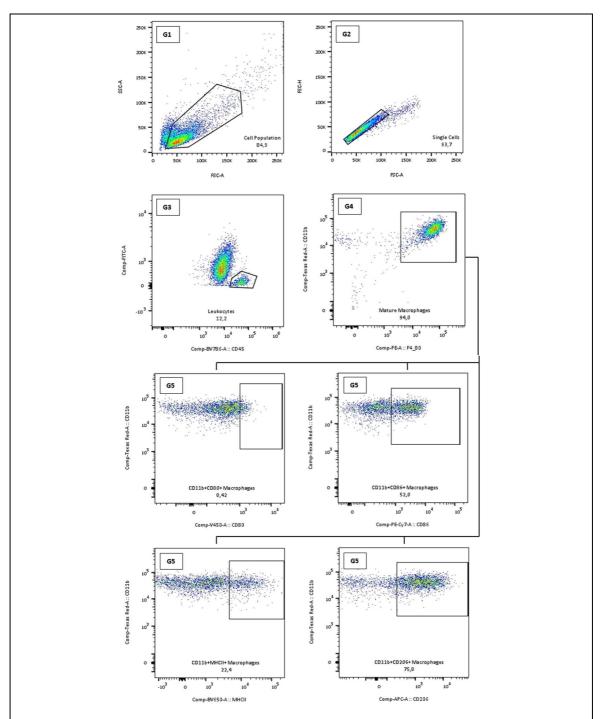
Supplementary Figure S2. Gating Strategy of Flow Cytometry based Cell Death Assay. First, the whole cell population was gathered excluding debris (G1). Next, single cells were selected by doublet discrimination (G2). Then, leukaemic cells, indicated by their expression of B cell lineage marker B220 (G3), were characterised upon their Annexin V/Propidium iodide profile (G4). According to this, apoptotic cells were determined as AV+Pi- (Q1, early apoptotic) and AV+Pi+ (Q2, late apoptotic), necrotic cells as AV-Pi+ (Q3) and viable cells as AV-Pi- (Q4).

8.11 Supplementary Figure S3: Gating Strategy of Flow Cytometry Analysis of Calreticulin Exposure



Supplementary Figure S3. Gating Strategy of Flow Cytometry Analysis of Calreticulin Exposure. After the exclusion of debris (G1) and doublets (G2), leukaemic cells identified as GFP+ (G3) were examined for their expression of CALR on the cell surface. In order to discriminate positive from negative cells, unstained controls were performed allowing the setup of the positive gating region (G4). Thus, a gate with a frequency of less than 1 % of unstained cells was defined as positive gating region.

8.12 Supplementary Figure S4: Gating Strategy of Flow Cytometry Analysis of Macrophages Co-Culture



Supplementary Figure S4. Gating Strategy of Flow Cytometry Analysis of Macrophages Co-Culture. After the exclusion of debris (G1) and doublets (G2), macrophages were differentiated from leukaemic cells by their expression of leucocyte common antigen CD45 (G3). Mature macrophages characterized by their double positivity for CD11b and F4/80 (G4) were finally examined for their expression profile of macrophages activation and polarisation marker CD80, CD86, CD206 and MHCII (G5).

8.13 Table 14: Shift in Macrophages Activation and Polarisation. Co-Culture Conditions with 10 nM Nilotinib

	CD80	CD86	CD206	MHCII
L1-1 (juxtacrine)	- 5.5 %	- 0.7 %	- 3.9 %	+ 1.2 %
L1-1 (paracrine)	+ 0.2 %	+ 0.6 %	+ 1.5 %	- 4.8 %
L3 (juxtacrine)	- 2.6 %	- 2.6 %	+ 3.5 %	+ 5.6 %
L3 (paracrine)	- 2.1 %	+ 0.6 %	+ 7.5 %	- 1.2 %
L1-1 (average)	- 2.7 %	- 0.1 %	- 1.2 %	- 1.8 %
L3 (average)	- 2.4 %	- 1.0 %	+ 5.5 %	+ 2.2 %
Juxtacrine (average)	- 4.1 %	- 1.7 %	- 0.2 %	+ 3.4 %
Paracrine (average)	- 1.0 %	+ 0.6 %	+ 4.5 %	- 3.0 %
Total (average)	- 2.5 %	- 0.5 %	+ 2.2 %	+ 0.2 %

Shift in macrophages activation and polarisation in the co-culture conditions with 10 nM Nilotinib pre-treated leukaemic cells compared to the corresponding co-culture conditions with untreated leukaemic cells.

8.14 Table 15: Shift in Macrophages Activation and Polarisation. Co-Culture Conditions with 50 nM Nilotinib

	CD80	CD86	CD206	MHCII
L1-1	- 10.0 %	+ 2.0 %	- 5.4 %	+ 1.1 %
(juxtacrine)				
L1-1 (paracrine)	- 1.5 %	+ 2.2 %	+ 2.5 %	- 2.7 %
L3				
(juxtacrine)	+ 4.8 %	- 1.3 %	+ 3.9 %	+ 7.9 %
L3	+ 1.7 %	- 20.1 %	+ 4.8 %	- 0.6 %
(paracrine)	+ 1.7 %	20.1 70	7 4.0 70	0.0 70
L1-1	- 5.8 %	+ 2.1 %	- 1.5 %	- 0.8 %
(average)		7 2.1 70	1.5 %	- 0.0 78
L3	+ 3.3 %	- 10.7 %	+ 4.4 %	+ 3.7 %
(average)		- 10.7 /8	T 4.4 /6	+ 3.7 /6
Juxtacrine	- 2.6 %	+ 0.4 %	- 0.8 %	+ 4.5 %
(average)		+ 0.4 /8	- 0.0 /8	7 7.5 76
Paracrine	+ 0.1 %	- 9.0 %	+ 3.7 %	- 1.7 %
(average)		- 3.0 /0	T U.1 /0	- 1.7 /0
Total	- 1.3 %	- 4.3 %	+ 1.5 %	+ 1.4 %
(average)		- 4.0 /0	+ 1.5 /0	+ 1. 4 /0

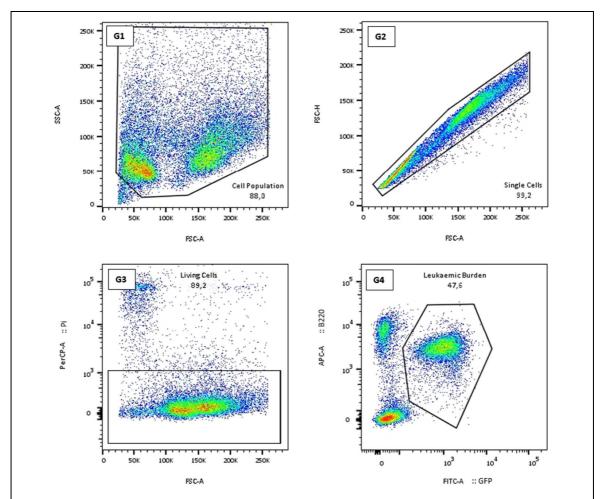
Shift in macrophages activation and polarisation in the co-culture conditions with 50 nM Nilotinib pre-treated leukaemic cells compared to the corresponding co-culture conditions with untreated leukaemic cells.

8.15 Table 16: Shift in Macrophages Activation and Polarisation. Co-Culture Conditions with 1 μ M Mitoxantrone

	CD80	CD86	CD206	MHCII
L1-1 (juxtacrine)	+ 4.5 %	+ 25.4 %	+ 17.4 %	- 5.0 %
L1-1 (paracrine)	+ 6.6 %	+ 26.1 %	+ 25.5 %	- 12.9 %
L3 (juxtacrine)	+ 2.1 %	+ 22.5 %	+ 21.6 %	- 7.1 %
L3 (paracrine)	+ 4.4 %	+ 31.9 %	+ 31.9 %	- 11.0 %
L1-1 (average)	+ 5.6 %	+ 25.8 %	+ 21.5 %	- 9.0 %
L3 (average)	+ 3.3 %	+ 27.2 %	+ 26.8 %	- 9.1 %
Juxtacrine (average)	+ 3.3 %	+ 24.0%	+ 19.5 %	- 6.1 %
Paracrine (average)	+ 5.5 %	+ 29.0 %	+ 28.7 %	- 12.0 %
Total (average)	+ 4.4 %	+ 26.5 %	+ 24.1 %	- 9.0 %

Shift in macrophages activation and polarisation in the co-culture conditions with 1 μ M Mitoxantrone pre-treated leukaemic cells compared to the corresponding co-culture conditions with untreated leukaemic cells.

8.16 Supplementary Figure S5: Gating Strategy of Flow Cytometry Analysis of Vaccination Assay. Quantifying Leukaemic Burden



Supplementary Figure S5. Gating Strategy of Flow Cytometry Analysis of Vaccination Assay. Quantifying Leukaemic Burden. After the exclusion of debris (G1) and doublets (G2), dead cells positive for Propidium iodide (Pi) were excluded from living cells (G3). Finally, leukaemic burden was assessed by the frequency of cells double positive for GFP and B cell lineage marker B220 (GFP+B220+) that represent the amount of BCR-ABL B-ALL leukaemic cells correlating with the progression of leukaemia (G4).

9 Publication

9.1 Publication 2019

Tirado-Gonzalez I, Nevmerzhitskaya A, Descot A, Devona Soetopo, Ewelina Czlonka, Maresa Weitmann, Carolin Wachtel, Julia Slotta-Huspenina, Christine Tran-Quang, Katharina Götze, Emily Alberto, Carla Vanina Rothlin, Jacques Ghysdael and Hind Medyouf. Abstract A092: TAM receptors targeting unleashes antileukemic immunity and enables checkpoint blockade leading to eradication of leukemic cells. In: Tackling the Tumor Microenvironment: Beyond T-cells: American Association for Cancer Research; 022019:A092-A092.

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- Translationale Forschung im Bereich Hämatoonkologie und Immunologie
- Doktorvater Herr Prof. Dr. Florian Greten, Betreuerin Frau Dr. Hind Medyouf

Publikation

02/2019

TAM receptors targeting unleashes antileukemic immunity and enables checkpoint blockade leading to eradication of leukemic cells

Irene Tirado-Gonzalez, Aleksandra Nevmerzhitskaya, Arnaud Descot, Devona Soetopo, Ewelina Czlonka, Maresa Weitmann, Carolin Wachtel, Julia Slotta-Huspenina, Christine Tran-Quang, Katharina Götze, Emily Alberto, Carla Vanina Rothlin, Jacques Ghysdael and Hind Medyouf

International Cancer Immunotherapy Conference: Translating Science into Survival September 30 - October 3, 2018; New York, NY

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Sprachen

Deutsch: Muttersprache

Englisch: fließend in Schrift und Sprache
 Spanisch: fortgeschrittene Kenntnisse
 Französisch: Grundkenntnisse

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12 Schriftliche Erklärung

Ich erkläre ehrenwörtlich, dass ich die dem Fachbereich Medizin der Johann Wolfgang Goethe-Universität Frankfurt am Main zur Promotionsprüfung eingereichte Dissertation mit dem Titel

Exploration of Nilotinib enhancing the anti-leukaemic Effects of Gas6 Deletion in Ph+ B-ALL by inducing Immunogenic Cell Death

am Georg-Speyer-Haus, Institut für Tumorbiologie und Experimentelle Therapie, unter Betreuung und Anleitung von Prof. Dr. Florian Greten mit Unterstützung durch Dr. Hind Medyouf ohne sonstige Hilfe selbst durchgeführt und bei der Abfassung der Arbeit keine anderen als die in der Dissertation angeführten Hilfsmittel benutzt habe. Darüber hinaus versichere ich, nicht die Hilfe einer kommerziellen Promotionsvermittlung in Anspruch genommen zu haben.

Ich habe bisher an keiner in- oder ausländischen Universität ein Gesuch um Zulassung zur Promotion eingereicht. Die vorliegende Arbeit wurde bisher nicht als Dissertation eingereicht.

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