

Investigations on bendamustine esters as new antitumor agents and the role of ABCG2 as a surrogate marker of breast cancer initiating cells



Dissertation

Zur Erlangung des Doktorgrades der Naturwissenschaften (Dr. rer. nat.)
an der Fakultät Chemie und Pharmazie der Universität Regensburg.

vorgelegt von

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2015

Die vorliegende Arbeit entstand in der Zeit vom März 2011 bis November 2014 unter der Anleitung von Herrn Prof. Dr. Armin Buschauer am Institut für Pharmazie der Naturwissenschaftlichen Fakultät IV – Chemie und Pharmazie – der Universität Regensburg.

Das Promotionsgesuch wurde eingereicht im Februar 2015.

Tag der mündlichen Prüfung: 06.03.2015

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Prof. Dr. A. Buschauer (Erstgutachter)
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Prof. Dr. J. Wegener (Drittprüfer)

Für meine Familie

Danksagung:

An dieser Stelle möchte ich mich herzlich bei allen die zum Gelingen dieser Arbeit beigetragen und mich während der Promotion begleitet haben bedanken. Besonderer Dank geht an:

Herrn Prof. Dr. Armin Buschauer für die Möglichkeit interessante und vielseitige Projekte zu bearbeiten, für die stete Ermöglichung Ideen umsetzen zu können und die gewährte forschersische Freiheit, für die zu jedem Zeitpunkt gesicherte Finanzierung und nicht zuletzt für die konstruktive Kritik und die vielen investierten Stunden bei der Durchsicht der Publikationen und dieser Arbeit.

Prof. Dr. Gero Brockhoff (Abteilung für Frauenheilkunde und Geburtshilfe, Universität Regensburg) für die Unterstützung und das stete, große Interesse am Fortgang meiner Arbeit, für die kompetente Hilfe bei verschiedenen Fragestellungen und für die konstruktive Kritik und investierte Zeit bei der Durchsicht der Publikation.

Prof. Dr. Günther Bernhardt für seine Hilfsbereitschaft, sein breites Wissen, die konstruktive Kritik bei der Durchsicht der Publikationen und für das hervorragende Sommerfest.

Der Arevi Pharma GmbH sowie Dr. Helmut Schickaneder und Christian Schickaneder für die Synthese und Bereitstellung der Bendamustinester.

Prof. Dr. Jörg König (Abteilung klinische Pharmakologie, Institut für Pharmakologie und Toxikologie, Friedrich-Alexander Universität, Erlangen) für die zur Verfügung gestellten HEK-OCT Zelllinien und die Durchführung der qPCR auf OCT-Expression in humanen Krebszelllinien sowie die Durchsicht der Publikation.

Dr. Anja Wege (Abteilung für Frauenheilkunde und Geburtshilfe, Universität Regensburg) für die hohe Motivation, außerordentliche Freundlichkeit und stete Hilfsbereitschaft bei zytometrischen und tierexperimentellen Arbeiten, sowie für die Durchführung der Versuche im NSG-Mausmodell.

Johannes Hüttner (Institut für Pharmakologie und Toxikologie, Universität Regensburg) für viele zum einen kritische, fachlich inspirierende und zum anderen freundschaftliche und unterhaltsame Gespräche inner- und außerhalb der Universität sowie für die tatkräftige Unterstützung bei der Konfokalmikroskopie.

Frauke Antoni für die gute Zusammenarbeit und ihr großes Engagement und Interesse während ihrer Masterarbeit sowie für die Synthese und Testung der Bendamustinester die Teil dieser Arbeit wurden.

PD Dr. Petra Hoffmann (Klinik und Poliklinik für Innere Medizin III, Universität Regensburg) und ihren Mitarbeitern Dr. Monique Germerodt und Rüdiger Eder für die Möglichkeit den FACSARIA™ Ili zur Zellsortierung zu nutzen und die große Hilfe bei der praktischen Durchführung.

Dr. Thilo Spruss und Franz Wiesenmayer für die Betreuung und Hilfe bei der Durchführung der tierexperimentellen Arbeiten.

Maria Beer-Krön für die tatkräftige Unterstützung bei verschiedensten Laborarbeiten sowie für ihr stetes Interesse, ihre Fröhlichkeit und Hilfsbereitschaft bei jedwedem Problem.

Petra Pistor und Rosa Kromas für die Anfertigung der histologischen Färbungen der Maustumore.

Peter Richthammer für die Unterstützung bei technischen Problemen und der Wartung von Geräten.

Brigitte Wenzl für die Vorbereitung des Biochemiepraktikums und die Durchführung der Mycoplasmentests.

Meinen Bürokollegen Dr. Johannes Felixberger, Dr. Uwe Nordemann und Nicole Plank für viele unterhaltsame Stunden, anregende Diskussionen fachlicher und anderweitiger Natur und die stets hervorragende Stimmung.

Allen aktuellen und ehemaligen Mitarbeitern des Lehrstuhls, insbesondere Dr. Paul Baumeister, Dr. Tobias Birnkammer, Dr. Miriam Ertel, Dr. Roland Geyer, Dr. Tobias Holzammer, Dr. Melanie Kaske, Dr. Carolin Meyer, Dr. Nikola Pluym und Dr. Stefanie Rodler für die herzliche Aufnahme im Arbeitskreis, viele unterhaltsame Gespräche und die schöne gemeinsame Zeit in Regensburg.

Zuletzt denjenigen die den Weg bis zur Promotion erst ermöglicht haben und derer Unterstützung ich mir auch währenddessen immer bewusst sein konnte: meinen Eltern, meinen Brüdern und meiner Freundin Maria.

Patents and Publications:

Patents:

1. Schickaneder H., Schickaneder C., Buschauer A., Bernhardt G., Huber S., Limmert M., Bendamustine derivatives and related compounds, and medical use thereof in cancer therapy, International Patent Application, Int. Pub. Nr.: WO2013/189847 A1, **2013**
2. Schickaneder H., Buschauer A., Bernhardt G., Schickaneder C., Limmert M., Huber S., Esters of bendamustine and related compounds, and medical use thereof, European Patent Application, Appl. Nr.: 12165728.2, **2013**

Publications:

1. Huber S, Antoni F, Schickaneder C, Schickaneder H, Bernhardt G, Buschauer A, Stabilities of neutral and basic esters of bendamustine in plasma compared to the parent compound: Kinetic investigations by HPLC, J Pharm Biomed Anal. 2015;104: 137-143
2. Huber S, Wege A K, Bernhardt G, Buschauer A, Brockhoff G, Topotecan-induced ABCG2 expression in MCF7-cells is associated with decreased CD24 and EpCAM expression and a loss of tumorigenicity, Cytometry A. 2014, under revision
3. Huber S, Huettner J P, Hacker K, Bernhardt G, König J, Buschauer A, Esters of bendamustine are by far more potent cytotoxic agents than the parent compound against human sarcoma and carcinoma cells, PloS One. 2015, submitted

Poster presentations:

- 10/2014 Annual conference of the DGFZ, Dresden.
Huber S., Wege A. K., Bernhardt G., Buschauer A., Brockhoff G.
„Topotecan-induced decrease in CD24 expression leads to a loss of tumorigenicity of human MCF-7 breast cancer cells.”
- 10/2013 Annual meeting of the DPhG, Freiburg.
Huber S., Bernhardt G., Buschauer A.
„Bendamustine esters versus bendamustine: cytotoxicity and stability“
- 10/2013 Annual meeting of the DPhG, Freiburg.
Huber S., Bernhardt G., Buschauer A.
„Esters of bendamustine are superior to the parent compound in terms of induction of p53 expression and cellular accumulation“

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1 Chapter I

General introduction

1.1 Cancer - definition and significance

The National Cancer Institute defines cancer as a “term used for diseases in which abnormal cells divide without control and are able to invade other tissues”. More than 100 different identified types of cancer underline the variability of this disease (c.f. <http://www.cancer.gov/cancertopics/cancerlibrary/what-is-cancer>). Malignant neoplasms, termed cancer, differ from benign neoplasms in the ability to infiltrate the surrounding tissues, resulting in the destruction of the latter, as well as the ability to spread to regional lymph nodes and distant organs via blood vessels and lymphatic channels (metastasis).

Cancer is among the leading causes of death worldwide, with estimated 8.2 million deaths in 2012 [1] and additional 32.6 million people living with cancer. Among the multitude of different cancer types, lung (1.6 million), liver (0.75 million), stomach, colorectal (both 0.7 million) and breast cancers (0.5 million) cause the highest number of deaths each year. The frequency of cancer types and related deaths is sex-dependent, which becomes most obvious in the case of breast cancer which had the highest mortality among women in 2008 [2].

1.2 Classification of cancer

The variety of cancer types can be described according to different criteria. Based on the cells of origin, cancer is usually distinguished in carcinomas, and sarcomas, the latter including hematological neoplasms (leukemia and lymphoma), all of which are divided into subgroups [3]. Carcinomas, which are divided into the major subtypes adenocarcinoma and squamous cell carcinoma, arise from epithelial cells and represent the vast majority of all cancer types (85 %), including most neoplasms of lung, breast, prostate, colon and pancreas. Sarcomas are defined as cancer arising from mesenchymal origin, e.g. bone, muscle or connective tissues. The hematological neoplasms leukemia and lymphoma arise from the hematopoietic tissue. Leukemia develops in the bone marrow and is characterized by an overproduction of immature white blood cells. Lymphomas (e. g. Hodgkin and non-Hodgkin lymphoma) and myelomas develop in the primary and secondary lymphatic tissues.

1.3 Development of cancer

Cancer development is most commonly regarded as a clonal process, derived from a single cell [4], comprising different stages, termed initiation, promotion and progression [5]. Initiation is characterized by alterations (mutations) in the genome, whereby additional spontaneous mutations and external effectors are considered as triggers. The external carcinogenic factors comprise physical (e.g. ionizing radiation), chemical (e.g. polycyclic aromatic hydrocarbons,

alkylating agents) and biological (e.g. viral infections [6]) damages. Epigenetic alterations such as changes in DNA methylation [7] as well as a variety of possible mutations, for example alterations in single nucleotides [8, 9] or chromosomal translocation [10-12], have been associated with cancer development.

Especially mutations in tumor suppressor genes (usually accompanied by a loss of function) and proto-oncogenes (usually gain of function) are associated with cancer [13]. For instance, cells with mutations in the tumor suppressor gene p53, i.e. decrease/loss of function of the p53 protein, can escape apoptosis [14]. However, "Gain of function" such as enhanced cell proliferation or the stimulation of epithelial-to-mesenchymal transition, cell motility and invasion are associated with p53 mutations as well [14]. In contrast, mutated or abnormally expressed proto-oncogenes (then referred to as activated oncogenes) and the related oncoproteins (e.g. platelet derived growth factor, epidermal growth factor receptor) commonly trigger proliferation or restrict apoptosis [15]. Besides mutations in tumor suppressor or proto-oncogenes, mutations resulting in reduced DNA repair contribute to cancer development [16, 17].

It is assumed that promoting agents (e.g. phorbol esters, reactive oxygen species [18-21]), acting as mitogens, trigger clonal expansion and proliferation of the initiated tumor cell [5, 22, 23]. The genetic instability, acquired during initiation, in combination with the proliferation and clonal expansion in the promotion phase leads to varying malignant phenotypes during the progression phase [24], a process which can take many years, increasing the heterogeneity of the developing tumor [5]. However, it should be noted that carcinogenesis is considered as a multicausal, multigenic and multistep process, involving genetic as well as epigenetic factors [25]. Besides spontaneous mutations and external factors, stromal cells, forming a permissive microenvironment (niche) surrounding the initiated cell, are assumed to contribute to carcinogenesis [26].

1.4 Tumor heterogeneity and cancer stem cells

The phenotypic heterogeneity of cancer cells [27, 28] is not fully understood. Two models the “stochastic” and the “cancer stem cell model”, were proposed [29-31].

The “classical” model assumes that random (stochastic) events change the behavior of single tumor cells, resulting in heterogeneity. These events can either be intrinsic (e.g. transcription factors) or extrinsic (e.g. microenvironment, immune response). In this model, all cancer cells have the potential of self-renewal and proliferation.

In contrast, the stem cell model assumes the similarity of tumors to healthy tissues (e.g. colon epithelium and blood cells), which are commonly organized in hierarchies with stem cells, possessing self-renewing capacity, on top. In this model, so-called cancer stem cells give rise to progenitor cells, which in turn differentiate or proliferate resulting in heterogeneous populations of cancer cells.

However, high plasticity (ability to change the phenotype) between different “states” of cancer cells was recently reported for breast cancer. Sorted cells of three distinct differentiation states, described as stem-like ($CD24^-/CD44^{high}/EpCAM^{low}$), basal ($CD24^-/CD44^{high}/EpCAM^-$) and luminal ($CD24^{high}/CD44^{low}/EpCAM^{high}$), were able to give rise to all three phenotypes by interconversion, resulting in initial equilibrium proportions [32]. It has to be stated that the authors of the cited study did not claim that their data would contradict the presence of “cancer stem cells”. In contrast, they concluded that cells in different “states” are able to give rise to “cancer stem cells”. The increasing knowledge about plasticity of cancer cells challenges cancer hierarchy, requiring more complex models to explain cancer cell heterogeneity. One approach, addressing this issue is for example the “CSC plasticity model” [33], which considers the majority of the tumor cell mass as cells with stem cell properties. The stem cell properties are supposed to vary depending, for example, on the microenvironment.

1.4.1 Cancer stem cells or cancer initiating cells?

Uni- or multipotent adult stem cells in healthy tissues share properties with cancer cells, such as self-renewal and asymmetric division, giving rise to new tissue [34]. This fueled speculations about the existence of cancer stem cells (CSC) and their involvement in carcinogenesis and relapse. The self-renewing capacity of both cell types is considered to be regulated by similar signaling pathways (e.g. Wnt [35, 36], Sonic Hedgehog [36, 37] and Notch [38, 39]). According to McCulloch and Siminovitch [40, 41], the proof of stem cell properties is achieved by demonstrating the self-renewing capacity, illustrated by colony forming ability *in vitro* [42] and tumorigenicity *in vivo*. Lapidot et al. identified “cancer stem cells” in acute myeloid leukemia (AML) [43], and suggested the formation of colonies or tumor spheres *in vitro* [44, 45] and the

tumorigenicity in immunodeficient mice as the state-of-the-art methods to define “cancer stem cell” properties [29, 46]. In accordance to the methods used for the actual proof of stem cell-like properties by tumor initiation respectively sphere formation, Beck et al. suggested the use of the terms “tumor propagating” or “cancer initiating cell” (TPC, CIC) rather than cancer stem cell [29]. Apart from self-renewal and asymmetric cell division, adult stem cells and CICs share characteristics such as the ability to secrete growth factors, leading to independent growth control of cells and induction of angiogenesis [47], the expression of ABC transporters [29] and other cell surface proteins referred to as stem cell markers (discussed in section 1.4.3).

1.4.2 Origin of cancer initiating cells

The origin of CICs has not unambiguously been clarified. Different hypotheses especially for hematological diseases, have been discussed in the last years [43]. CICs may result from transforming mutations of adult stem cells leading to abnormal growth and differentiation [34, 48-50]. This is supported by the fact that tissue specific stem cell markers are also expressed on the corresponding CICs [51]. The influence of the surrounding microenvironment (niche), especially in endothelial tissue, may also play a crucial role in the development of CICs originating from adult stem cells [47, 52, 53].

A different concept considers mutations in progenitor cells as initial step in the development of CICs [34, 54], since only few mutations affecting mechanisms which regulate cell death would be necessary to gain stem cell properties accompanied by uncontrolled cell growth [43, 55, 56]. Additionally, de-differentiation of progenitor or differentiated cells as well as the fusion of tissue-specific stem cells with circulating bone marrow stem cells are discussed [51, 57].

1.4.3 Identification of cancer initiating cells

A considerable number of markers, usually expressed on CICs and adult stem cells originating from specific tissues, was described in the last years. The detection of their expression is the most commonly used method for the identification of CICs [46]. To name only a few examples, the detection of CD133 [58, 59] respectively a combination of CD133 and nestin [60, 61] is considered indicative for brain tumor initiating cells (BTIC). The identification of breast cancer initiating cells (BCIC) is usually based on the differential expression of the cell surface proteins CD24 and CD44. The expression pattern of BCIC populations was reported as CD24⁻/CD44⁺ [62-64]. Nevertheless, there is no clear correlation between the expression of these proteins and stem cell properties (c.f. section 1.4.1) [65].

Besides the detection of cell surface proteins, the identification of CICs based on functional assays was proposed, e. g. by determining the activities of aldehyde dehydrogenase (ALDH) [66-68] or the proteasome [69].

Especially in the absence of other specific markers, the determination of so-called side populations (SP), expressing ATP-binding cassette (ABC) transporters such as ABCB1, ABCC1 and ABCG2 may be a valuable approach to the identification and enrichment of CICs [70]. However, the expression of such transporters alone is not strictly associated with stem cell properties, since CICs often represent only a fraction of the side population [71, 72]. In addition to the Hoechst 33342 efflux assay for the identification of SPs [73, 74], methods such as the ABCG2-mediated mitoxantrone efflux assay [75] or the immunodetection of transporter proteins are frequently performed. Apart from being considered as marker proteins, ABC-transporters such as ABCB1 (P-gp) [76] and ABCG2 [77-79] impart increased resistance against chemotherapy, as a considerable number of cytostatic drugs are substrates of these efflux pumps [80].

Commercially available antibodies against CIC markers in combination with flow cytometry led to the identification of CICs in a great variety of cancer types, including brain tumors [44, 58], breast cancer [62], colorectal cancer [81, 82], skin squamous cell carcinoma [83], head and neck cancer [84], lung cancer [85], pancreatic cancer [86], prostate cancer [87] and ovarian cancer [88, 89].

1.5 Cancer chemotherapy

Chemotherapeutic regimens for the treatment of the majority of cancer types comprise a combination of different drugs. It is conceivable that enhanced anti-cancer efficacy of drug combinations is accompanied by reduced systemic toxicity, since lower doses of the individual drugs are required, especially, if the mechanisms of action and the side effects of the individual drugs do not overlap. Profound knowledge of the phenotype (e.g. expression of HER2 or EGFR) or resistance mechanisms (e.g. increased expression of dihydrofolate reductase in case of methotrexate treatment) favors the identification of appropriate drug combinations. Most chemotherapeutic regimens are applied in repetitive cycles in order to achieve a reduction of tumor mass in each cycle [90].

1.5.1 Alkylating agents

Despite the development of targeted anti-tumor drugs, alkylating agents continue to play a central role amongst cytotoxic chemotherapeutics (e.g. antifolates, antimetabolites, antimitotics, platinum analogs, topoisomerase inhibitors). Such drugs are applied in conventional combination therapy as well as in high-dose protocols with subsequent hematopoietic cell transplantation [91]. Besides the classical alkylating agents (which will be discussed in the following), so-called nonclassical agents, which do not contain alkylating moieties such as the nitrosourea or the nitrogen mustard group are frequently used (e.g. procarbazine, temozolomide). These “monofunctional” drugs are cytotoxic due to covalent binding to the DNA, too, but they usually undergo complex metabolic transformation to active intermediates [91].

1.5.2 Classical alkylating agents

Alkyl alkanesulfonates (e.g. busulfan), nitrosureas (e.g. carmustine) and nitrogen mustard derivatives (e.g. bendamustine), the first non-hormonal anti-cancer drugs, are usually classified as classical alkylating agents. The primary mechanism of action is the covalent binding to DNA, whereby an active intermediate acts as alkylating agent rather than the parent compound. The alkylating reaction may proceed according to a S_N1 (e.g. chloroethylnitrosureas) or S_N2 (e.g. busulfan) mechanism, which particularly differ regarding the rate of reaction. Alkylation of DNA results in cytotoxic, mutagenic and carcinogenic effects. The effectiveness of alkylating agents can be enhanced, for example, by radiation, hyperthermia, inhibition of DNA repair mechanisms and glutathione depletion [91]. The examination of the sites of alkylation revealed that oxygen atoms in phosphates [92, 93] and bases as well as nitrogen atoms of purine bases [94] are preferentially

alkylated. However, DNA is not exclusively alkylated, but other cellular constituents such as proteins are alkylated to a minor extent, as shown for nitrogen mustard [95].

Bifunctional alkylators (e.g. nitrogen mustard derivatives) cause intra- and, in particular, interstrand crosslinks, responsible for effective inhibition of DNA replication and transcription [96, 97]. This is reflected by overall higher cytotoxic activities of bifunctional agents [98, 99].

Apart from different reactivity of the alkylating moiety, the antitumor activities of alkylating agents mainly depend on their (physico)chemical properties, influencing the pharmacokinetics including diffusion or transport across membranes or the activation of cellular resistance mechanisms [91]. For instance, the uptake of melphalan was reported to be enhanced by amino acid transporters [100] and the human organic cation transporter 3 (OCT3) [101], which was recently identified in some human cancer cell lines [101, 102]. Contrary, reduced drug uptake is a well-known resistance mechanism, for example, for melphalan [103] and mustine (nitrogen mustard) [104]. Further resistance mechanisms against alkylating agents include enhanced drug inactivation, changes in cell-cycle checkpoint function (e.g. loss of normal p53 function) and alterations in DNA repair mechanisms (e.g. enhanced base – and nucleotide excision repair) [91]. The induction of such resistance mechanisms is often accompanied by the development of cross-resistances against other alkylating agents [91].

1.5.2.1 Nitrogen mustard (N-Lost) derivatives

The term “Lost” for bis(2-chloroethyl) derivatives of nitrogen and sulfur goes back to the names of the inventors of this compound class, Lommel and Steinkopf. S-Lost (sulfur mustard, yellow cross) (Fig. 1.1) was used as a warfare gas in World War I. Massive lymphopenia, granulocytopenia, thrombocytopenia and moderate anemia in soldiers, who were accidentally exposed to S-Lost, suggested the therapeutic potential of Lost derivatives [105].

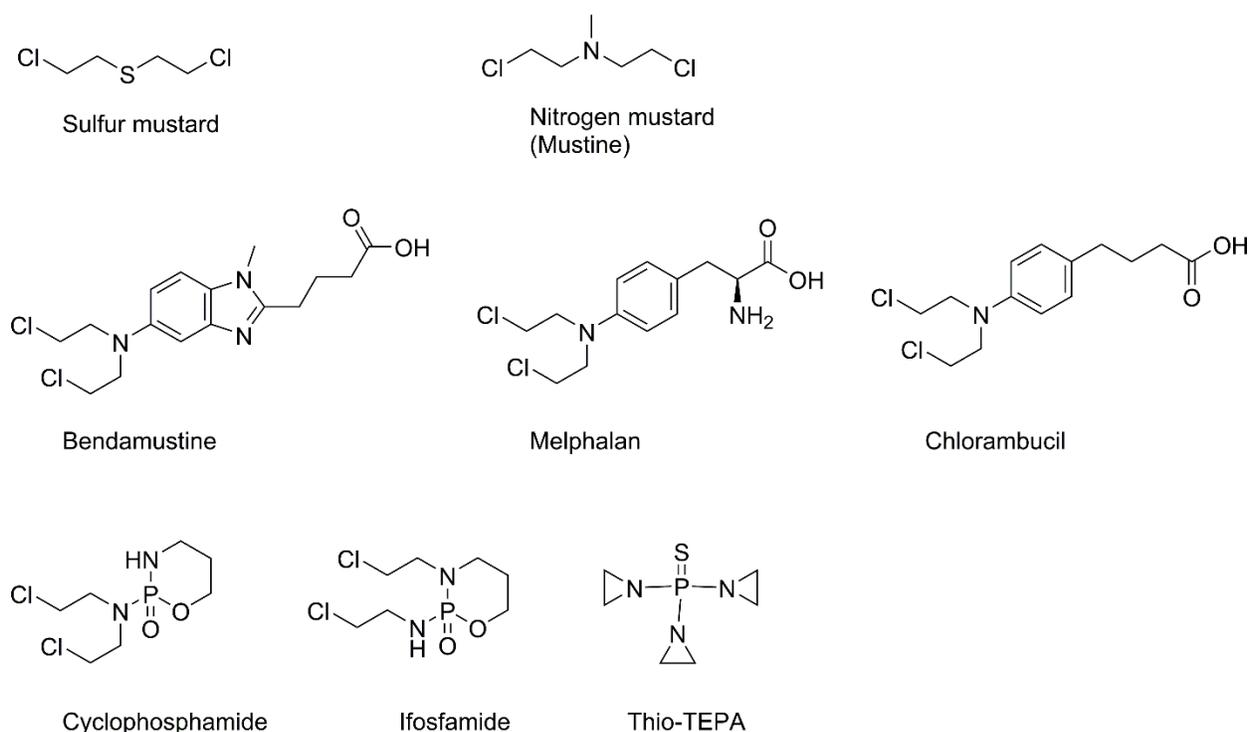


Figure 1.1: Structures of sulfur mustard (war gas), nitrogen mustard (N-Lost, mustine, chlormethine, mechlorethamine) and various derivatives.

Based on these findings, clinical trials on the effects of mustard derivatives against neoplasms of lymphoid tissue were performed in the US. Sulfur mustard derivatives proved to be inappropriate for parenteral administration due to severe adverse effects [105]. By contrast, the less reactive nitrogen mustard derivatives (e.g. mustine (mechlorethamine); c.f. Fig. 1.1) were better tolerated and led to an initial remission, especially in Hodgkin’s lymphoma [106, 107].

Subsequent investigations focused on structural modification of the N-Lost structure, aiming at higher chemical stability and less side effects. Therefore, the methyl group in mustine was replaced by electron-withdrawing substituents, rendering the nitrogen less nucleophilic (Fig. 1.1). Indeed, such N-Lost derivatives revealed lower systemic toxicities, though accompanied by

reduced antitumor activity. Melphalan, chlorambucil, bendamustine, cyclophosphamide and ifosfamide are examples of approved nitrogen mustard derivatives (c.f. Fig. 1.1). The aziridine derivative thio-TEPA is also related to the class of nitrogen mustards, although it does not comprise the bis(2-chloroethyl) group. Upon protonation, the aziridine rings correspond to aziridinium ions, reminiscent of the highly reactive alkylating intermediates formed from N-Lost derived alkylators.

Most nitrogen mustard derivatives are indicated to treat hematologic malignancies such as chronic lymphocytic leukemia (CLL) [108-110], non-Hodgkin's lymphoma (NHL) [111, 112] and multiple myeloma (MM) [113]. In contrast to classical mustard derivatives such as melphalan, spontaneous formation of aziridinium intermediates does not occur in case of cyclophosphamide and analogs such as ifosfamide. Instead, metabolic activation by CYP450 enzymes is required [114]. Cyclophosphamide and ifosfamide are also used in the treatment of solid tumors (e.g. breast cancer [115]; soft tissue sarcoma [116]).

The high alkylating potency of the N-Lost moiety is based on the presence of a nucleophilic center (S or N) in β -position relative to the halogen atom, enabling an intramolecular nucleophilic attack of a lone pair of electrons. This results in the displacement of the chloro substituent and the formation of a positively charged group, the highly reactive electrophilic aziridinium moiety, which subsequently reacts with nucleophiles in the DNA (Fig. 1.2). Due to the bifunctional character of the nitrogen mustard, this reaction can take place twice, resulting in DNA cross-links [105].

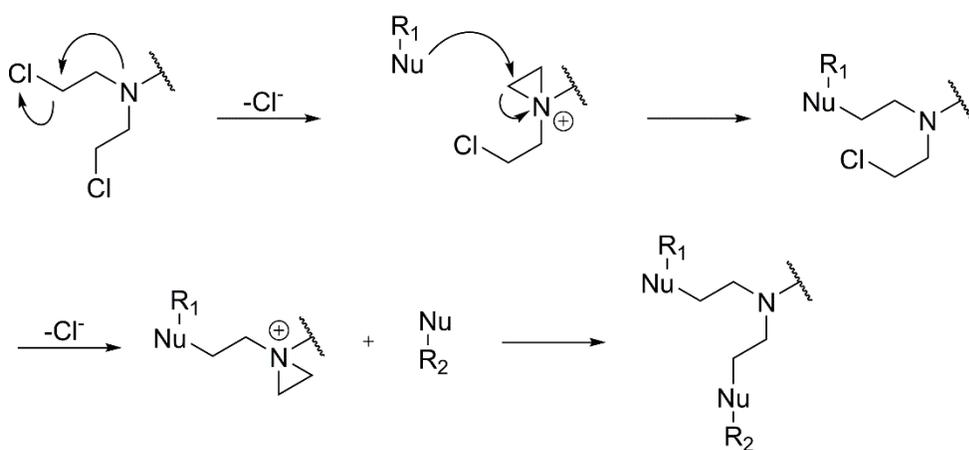


Figure 1.2: Mechanism of the alkylation of nucleophiles (Nu) by nitrogen mustard derivatives. The bifunctional character enables two consecutive alkylating steps.

1.6 Bendamustine

1.6.1 Development of bendamustine

Bendamustine (initially termed IMET3393) (c.f. Fig. 1.1) was synthesized in 1963 by Ozegowski et al. at the Institute for Microbiology and Experimental Therapy (IMET) in Jena [117, 118]. The inventors aimed at combining the alkylating activity of the nitrogen mustard group and potential antimetabolite properties of the benzimidazole scaffold. In addition, the electron-withdrawing benzimidazole structure should decrease the nucleophilicity of the nitrogen atom in the N-Lost group resulting in decreased reactivity and reduced toxicity [119]. To improve the solubility of the new compound, the butyric acid side chain was introduced, by analogy with the approach known from chlorambucil (c.f. Fig. 1.1).

First clinical trials with bendamustine in the treatment of several cancer types (e.g., hematologic malignancies, breast cancer, small cell lung cancer, ovarian cancer) were performed in 1965 [119]. However, bendamustine was barely used in Europe and the US before the end of the “cold war”. Subsequently, the drug was “re-discovered”, came into the focus of cancer research in Europe and the US at the end of the 1990s, and has been investigated in many clinical trials.

1.6.2 Bendamustine in cancer therapy

Bendamustine is approved for the treatment of CLL [110, 120], indolent NHL [121] and MM [122, 123] in Germany (Levact®) and for CLL and indolent NHL in the US (Treanda®). Most treatment regimens apply bendamustine in combination with other cytostatic drugs [124], often with rituximab [125-127]. Clinical research on bendamustine has been intensified in the last years. However, most trials with bendamustine as a single agent, for instance, for the treatment of bile duct cancer [128], soft tissue sarcoma [129], germ cell cancer [130], small cell lung cancer [131-133], pretreated metastatic [134, 135] or advanced [136] breast cancer revealed on the one hand good tolerability, but on the other hand limited benefit. Studies on bendamustine in combination therapy, for example with methotrexate and 5-fluorouracil for the treatment of metastatic breast cancer [137] or bendamustine with carboplatin for the treatment of small cell lung cancer [138], reported efficacies which were comparable to respective standard treatment regimens. Due to mild side effects and reduced cross resistances with other alkylating drugs, bendamustine was claimed to be an interesting drug for the treatment of patients in poor clinical condition [139] or as second line therapy [111, 140, 141].

1.6.3 Mechanism of action

Speculations about mechanisms of action of bendamustine different from the alkylation of DNA go back to the synthesis of the compound, based on the idea that the benzimidazole scaffold might confer antimetabolite activities. This was supported by the COMPARE analysis conducted at the national cancer institute [142]. In this study, bendamustine revealed a unique activity pattern against human cancer cell lines, which differed significantly from the activity pattern of other nitrogen mustard derivatives such as melphalan, chlorambucil and cyclophosphamide. Gene expression analyses indicated differential bendamustine-induced effects with respect to genes grouped into “response to DNA-damage stress” and “DNA-metabolism”. These findings led the authors to the assumption that bendamustine is unique in its ability to induce p53, regulate DNA-repair and inhibit mitotic checkpoints, leading to mitotic catastrophe [142]. However, detailed experiments verifying these mechanisms are lacking.

Several recent publications reported the induction of p53 [143, 144] and a concentration dependent induction of G2 phase arrest by bendamustine [143, 145, 146]. Additional evidence for effects on the cell cycle was provided by the analysis of proteins regulating cell cycle checkpoints, e. g. Chk1 [145] and Chk2 [143], and the induction of proteins involved in apoptosis (caspase-2 and caspase-8) [146]. Therefore, the induction of mitotic catastrophe, an alternate mechanism of cell death, which is particularly important in cells resistant to apoptosis, is discussed for bendamustine [146].

Recently, the putative activity as an antimetabolite was re-considered due to additive effects of various alkylating agents with bendamustine *in vitro* [147]. The fact that bendamustine was effective (e.g. induction of apoptosis) already after short exposure (3 hours), was interpreted as a hint to fast cellular uptake. Accordingly, the additive effect with other alkylating agents was explained by early effects of bendamustine combined with delayed cytotoxic effects of compounds entering the cells less rapidly. Moreover, co-incubation of bendamustine with cytosine arabinoside (CA) (pyrimidine antimetabolite) induced effects which are usually observed for combinations of pyrimidine with purine analogs (up-regulation of ENT1 expression and increased concentration of the active metabolite of CA, Ara-CTP). This was interpreted as further evidence for antimetabolite activity.

1.6.4 Stability and metabolism

The nitrogen mustard group is prone to chemical hydrolysis, in particular at neutral or basic pH values [148], resulting in the inactive bis(2-hydroxyethylamino)-substituted benzimidazole derivative upon twofold hydrolysis (Fig. 1.3). The hydrolytic decomposition is prevented or reduced at acidic pH, because the protonation of the nitrogen atom leads to decreased

nucleophilicity and, consequently, the tendency to form an aziridinium ion is considerably lowered. The hydrolysis rate is also retarded in the presence of high concentrations of chloride [148].

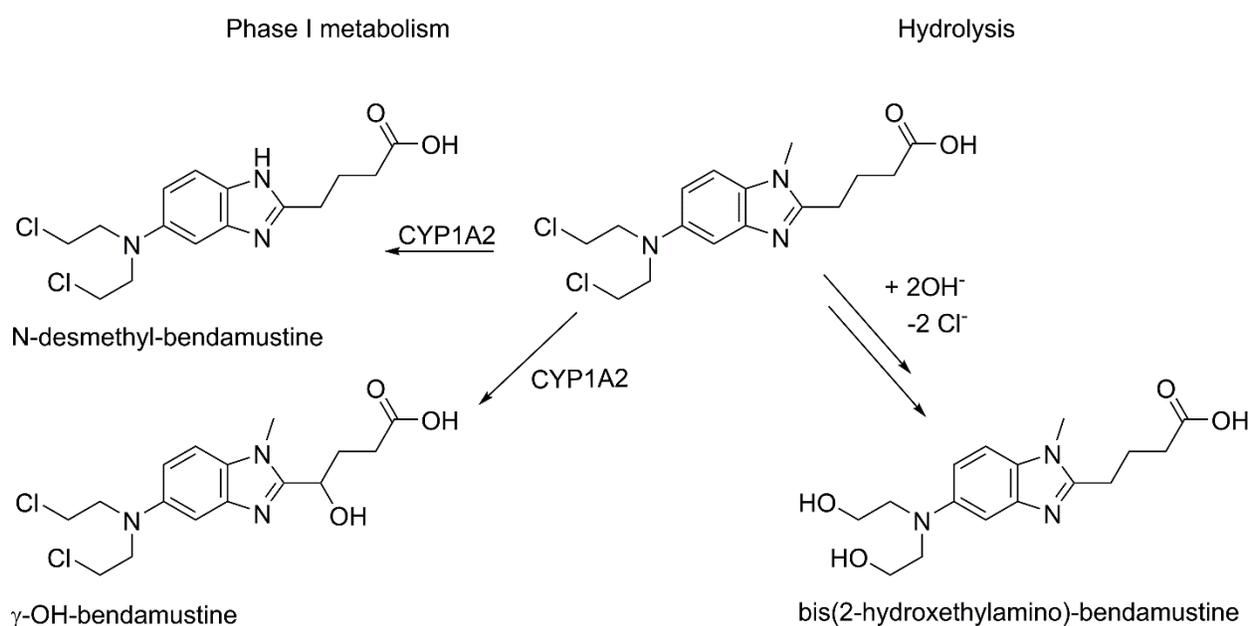


Figure 1.3: Phase I metabolism and chemical hydrolysis of bendamustine.

In addition to the chemical hydrolysis of the nitrogen mustard group in plasma, phase I and phase II metabolites of bendamustine are described in literature. Especially CYP1A2 was assumed to be relevant for the formation of the two known active phase I metabolites, N-desmethyl-bendamustine and γ -OH-bendamustine (c.f. Fig. 1.3) [149]. With regard to phase II metabolites, biliary excretion of N-acetyl-L-cysteine conjugates was previously described as a mechanism of bendamustine elimination [150, 151].

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2 Chapter II

Scope and objectives

2.1 Investigation of ABC transporter expression and cancer initiating cells in human cancer cell lines

The expression of ATP driven efflux pumps (ABC transporters) such as ABCB1 and ABCG2 is known to confer resistance to diverse cytostatic drugs [1]. The exposure of cancer cells to cytostatic drugs such as topotecan can cause elevated ABCG2-expression [2]. Moreover, side populations of cancer cells, expressing ABC-transporters (ABCB1, ABCG2) were recently reported to comprise cancer initiating cells [3]. However, knowledge whether the expression of ABC-transporters is actually linked to properties of cancer initiating cells such as *in vitro* clonogenicity and tumorigenicity is incomplete.

Aim of this project was the flow cytometric screening for cancer initiating- and side population cells in human brain tumor and breast cancer cells. Additionally, MCF-7 breast cancer cells were chosen for treatment with topotecan to examine the impact of this treatment on the expression of ABC-transporters and different marker proteins, which are commonly used for the identification of breast cancer initiating cells (CD24, CD44, EpCAM). Moreover, the effect of topotecan treatment on the *in vitro* clonogenicity and the tumorigenicity of MCF-7 cells was planned to be investigated.

2.2 Investigations on the stability and cytotoxic activity of bendamustine esters

Aiming at oral bioavailability, improved pharmacokinetic properties or increased stability, various bendamustine esters [4], oligomers [5] and complex pharmaceutical formulations [6] were claimed for patent in the last years. Among the recently reported bendamustine derivatives are a number of neutral and basic esters. Such structural modifications may not only influence the stability of the nitrogen mustard group, but also the cytotoxic activity. Moreover, in a biological environment the spectrum of possible degradation products for bendamustine esters is expected to become more complex compared to the parent compound, as, in addition to nucleophiles attacking the nitrogen mustard moiety, enzymatic process such as oxidation and, in particular, hydrolysis may contribute to degradation.

The second objective of this project was to establish and apply analytical procedures for the detailed investigation of the degradation of structurally diverse bendamustine esters in aqueous solution as well as in murine and human plasma. Moreover, as these esters might act not only as bendamustine prodrugs but also as cytotoxic agents of their own, investigations on the antiproliferative activity at a broad panel of human cancer cell lines, including hematologic

malignancies, osteosarcomas and different carcinomas, were envisaged. In addition, studies on the cellular uptake of representative esters and the parent compound as well as the effect on the expression of the p53 tumor suppressor were planned in case of eventually discovered significant differences in cytotoxicity.

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3 Chapter III

Investigation of ABC transporter expression and cancer initiating cells in human cancer cell lines

Note: As stated in the list of publications, parts of this chapter have already been submitted for publication, prior to the submission of this thesis. For detailed information on the contribution of other authors see also "Acknowledgements".

3.1 Introduction

The expression of ATP-driven efflux transporters such as the breast cancer resistance protein (BCRP, ABCG2) is a crucial mechanism that causes multidrug resistance in cancer cells [1]. Determining the efflux of different substrates, e.g. the fluorescent Hoechst dye [2] or mitoxantrone [3] allows to identify cancer cells expressing ABCG2, whereas Hoechst efflux defines so-called side populations [4]. Presupposed that resistance against cytostatics is also a property of cancer initiating cells (CICs), side populations are often considered to comprise an increased fraction of those cells compared to the bulk of cancer cells. Exposing cancer cells *in vitro* to cytostatic drugs like topotecan, mitoxantrone [5], and doxorubicin [6] can cause elevated ABCG2 expression. The effect of such cytotoxic treatments on the regulation and expression of other proteins related to poor clinical outcome has been less extensively examined.

Brain tumor initiating cells were previously described as a CD133⁺ subpopulation [7] and identified in different brain tumor types. Nowadays, CD133 is frequently used as a marker for cancer initiating cells originating from diverse tumor entities, for example, ovarian and liver cancer [8, 9]. However, the function of CD133, which was first described as a marker of primitive hematopoietic and neural stem cells [10], is not completely clarified.

Treatment efficiency and outcome of breast cancer patients is strongly affected by tumor heterogeneity, which therefore, constitutes a significant clinical problem. Nowadays the human epidermal growth factor receptor 2 (HER2) and the estrogen receptor (ER) represent well-established diagnostic markers and therapeutic drug targets whereas the importance of CD24, CD44 and EpCAM is less clear. Distinct expression profiles of these proteins are considered as markers of CICs and their expression has been related to mechanisms involved in metastasis and therefore with poor clinical outcome. Among other markers, the expression profile of a CIC (sub-)population in breast cancer has been described as CD24^{-/low}/CD44⁺ [11].

Considering the fact that increased CD24 expression is linked to poor prognosis in different solid tumor types including breast cancer [12, 13], low CD24 expression in CICs appears inexplicable at first glance. However, the association of high CD24 expression levels and metastasis [14], enhanced cell proliferation [15] and *in vitro* clonogenicity [16] is widely accepted. Nevertheless, regulation and function of CD24 are not fully elucidated yet.

The CD44 molecule, respectively variants thereof [17], is almost ubiquitously expressed on carcinoma cells [18]. It can interact with a variety of effectors [19, 20] and is involved in a multitude of regulatory mechanisms related to adhesion, cell proliferation, migration, invasiveness and chemoresistance [18, 21, 22].

Likewise, EpCAM, which was first described as a marker of human embryonic stem cells, is expressed on cancer cells of a variety of entities [23]. In many cancer types, including breast

cancer, high expression levels of EpCAM correlate with poor prognosis [24]. Although the functions of EpCAM as a cell adhesion molecule and promoter of cell proliferation are generally accepted and alterations in the expression level are closely related to epithelial-mesenchymal transition (EMT) during metastasis, the role of EpCAM in cancer cells is not yet fully elucidated [23, 25].

The HER2 receptor is overexpressed in up to 20 to 25 % of invasive breast cancers [26] and predicts poor clinical outcome. HER2 overexpression can result in constitutive kinase activity that causes increased cell proliferation and invasion [27].

In order to investigate correlations between side populations and cancer initiating cells, a panel of human brain tumor and breast cancer cell lines was screened for the expression of different CIC markers (CD133, CD24, CD44) as well as ABCB1 and ABCG2. Moreover, it was explored whether the topotecan induced ABCG2 expression manipulates the CD24, CD44, EpCAM, and HER2 related phenotype and if an enhanced ABCG2 expression affects *in vitro* clonogenicity or tumorigenicity in immunodeficient mice. To this end, ABCG2 expression was induced in MCF-7 breast cancer cells by long-term exposure to topotecan and the respective biomarker expression was assessed by flow cytometry. Moreover, the impact of topotecan treatment on cell propagation and the capacity of self-renewal was explored in mammosphere formation assays and the tumorigenicity of untreated vs. topotecan treated cells was evaluated in immunodeficient mice.

3.2 Materials and methods

3.2.1 Reagents and antibodies

Unless stated otherwise, all commercially available solvents and chemicals were of analytical grade and purchased from Merck (Darmstadt, Germany) or Sigma (Taufkirchen, Germany). For the purification of demineralized water, a Milli-Q system (Millipore, Eschborn, Germany) was used. A 1 mM stock solution of mitoxantrone was prepared by diluting Novatron® (Wyeth Pharma, Münster, Germany) in 70 % ethanol. Doxorubicin and topotecan were both from Sigma (Taufkirchen, Germany) and the stock solutions were also prepared in 70 % ethanol. Tariquidar was synthesized in our laboratory according to literature [28, 29] and dissolved in 70 % ethanol (100 µM). Calcein-AM (100 µM), 5-fluorouracil, methotrexate (all purchased from Sigma, Taufkirchen, Germany) and funitremorgin C (1 mM) (Merck, Darmstadt, Germany) were dissolved in DMSO. All stock solutions were stored at – 20 °C. A solution of 2 % glutardialdehyde (v/v) in PBS was prepared by diluting the commercially available 25 % (v/v) solution (Sigma, Taufkirchen, Germany). PBS containing 2 % fetal calf serum (FCS; Biochrom, Berlin, Germany) was used as buffer for all cytometric measurements. The following antibodies were used for the flow cytometric analyses: *APC anti-mouse/human CD44*, *Alexa Fluor 488 anti-mouse/human CD44*, *FITC anti-human CD24*, *Alexa Fluor 647 anti-human CD24*, *Brilliant Violet 421 anti-human CD326 (EpCAM)* and *pacific blue anti-human CD338 (ABCG2)*, as well as the isotype controls recommended by the manufacturer were purchased from BioLegend (London, UK). *FITC anti-human P-glycoprotein* and *IgG2b isotype control* as well as *Anti-Her-2/neu PE* respectively *PE Mouse IgG1 Isotype control* were from BD Biosciences (Heidelberg, Germany). *CD133/1-PE (clone AC133)* respectively *CD133/1-APC (clone AC133)* antibodies and the respective *mouse IgG1 isotype controls* were purchased from Miltenyi Biotec (Bergisch Gladbach, Germany).

3.2.2 Cell lines and standard cell culture

The human brain tumor cell lines LN-18 (ATCC® CRL-2610™; glioblastoma grade IV), U-87 MG (ATCC® HTB-14™; astrocytoma grad IV), SW-1783 (ATCC® HTB-13; astrocytoma grade III) and Daoy (ATCC® HTB-186™; desmoplastic cerebellar medulloblastoma) as well as the human breast cancer cell lines MCF-7 (ATCC® HTB-22™; breast adenocarcinoma), ZR-75-1 (ATCC® CRL-1500™; breast ductal carcinoma), T-47D (ATCC® HTB-133™; mammary gland ductal carcinoma), JIMT-1 (provided by Prof. Dr. Gero Brockhoff, Institute of Gynecology and Obstetrics, University of Regensburg), MDA-MB-231 (ATCC® HTB-26™; triple negative breast adenocarcinoma), HCC-1806 (triple negative primary acantholytic squamous cell carcinoma) and HCC-1937 (triple negative primary ductal carcinoma) (both provided by Dr. Jörg Engel, Frauenklinik, University of Würzburg) were maintained in RPMI-1640 (Sigma, Taufkirchen, Germany), supplemented with 2 g/L NaHCO₃

(Merck, Darmstadt, Germany), 110 mg/L sodium pyruvate, 10 mM HEPES (both from Serva, Heidelberg, Germany) and 10 % fetal calf serum (Biochrom, Berlin, Germany). For the incubation of MCF-7/Topo cells (ABCG2 overexpressing subclone of MCF-7) [29, 30] the culture medium was additionally supplemented with 500 nM topotecan. For experiments determining the expression stability of ABCG2 in MCF-7/Topo cells, the cells (hereafter referred to as MCF-7/Topo/wo) were maintained for several weeks in topotecan-free medium. All cell lines were maintained in cell culture flasks (Sarsted, Nuremberg, Germany) at 37 °C and 5 % CO₂ in a water-saturated atmosphere and serially passaged as required following the treatment with trypsin/EDTA (0.5 mg/mL / 0.22 mg/mL) (PAA, Pasching, Austria).

3.2.3 Neuro-/mammosphere culture and limiting dilution assay

Neuro- respectively mammosphere culture was basically performed as previously described [31-33]. Cells were maintained in suspension culture, using ultra-low attachment plates (Corning Inc., Corning, NY). MCF-7, MCF-7/Topo and MCF-7/Topo-CD24^{low} cells were maintained in DMEM/F12 supplemented with 20 ng/mL bFGF (both from Gibco®/Life Technologies, Carlsbad, CA), 20 ng/mL EGF and 5 µg/mL insulin (both from Sigma, Taufkirchen, Germany). For the incubation of Daoy and sorted CD133⁺-Daoy cells, the medium described above was additionally supplemented with B-27 (Gibco®/Life Technologies).

The growth capacity was determined by serial passaging of MCF-7, MCF-7/Topo, sorted MCF-7-CD24^{low}, Daoy and sorted Daoy-CD133⁺ cells. To this end, 5 x 10³ single cells were seeded per well in 100 µL medium. Every 3-4 days, fresh medium was added or, if the cells had reached confluency of approximately 80 %, spheres were enzymatically singularized and single cells were passaged. The cell growth was regularly monitored for at least 3 weeks. The expression of CD133 in sorted Daoy-CD133⁺ cells was again examined by flow cytometry (see 3.2.4.1) after 2 weeks of incubation.

The mammo- respectively neurosphere forming ability of freshly sorted MCF-7/Topo-CD24^{low}, and Daoy-CD133⁺ cells, was compared to unsorted cells using limiting dilution assays. For this purpose, different cell counts (1, 3, 10, 30, 100, 300, 1000 cells/well, 100µL) of singularized cells were plated in sextuplicate into 96-well ultra-low attachment plates (Corning). Wild type cells were trypsinized, washed 3 times with PBS in order to remove remaining FCS and trypsin and filtrated through a 40 µm cell strainer (BD Falcon, Heidelberg, Germany) to exclude agglomerates. Every 2-3 days, 20 µL of fresh medium were added and the number of spheres with diameters above 80 µm was determined in each well after 12 days [34, 35]. Pictures were taken with a DCM-510 ocular microscope camera (OCS.tec GmbH & Co. KG, Erding, Germany) controlled by ScopePhoto V3.0.12.498 software (Hangzhou Scopetek Opto-electric Co., Ltd., Hangzhou, China).

3.2.4 Determination of IC₅₀ values

The IC₅₀ values of 5-fluorouracil (5-FU), doxorubicin (Doxo) and methotrexate (MTX) on MCF-7 and MCF-7/Topo cells were determined after 72 hours of incubation according to the procedure described in section 5.2.4.

3.2.5 Flow cytometric screening of CIC - and side population markers

3.2.5.1 Cell surface markers

For the screening of cancer initiating cell-, respectively side population markers, cells in the log growth phase were detached by trypsinization. Prior to adjusting the cell number to 1×10^6 cells/mL in PBS/FCS (PBS + 2 % FCS), the cells were washed twice with PBS. Subsequently, the samples were probed with different antibodies, depending on the tumor entity of the particular cell line. Brain tumor cells were analyzed for the expression of CD133, ABCB1 and ABCG2. The expression of CD24, CD44, ABCB1 and ABCG2 was determined for all human breast cancer cell lines described in section 3.2.2. Additionally, MCF-7 and MCF-7/Topo cells were analyzed for the expression of EpCAM and HER2. All antibodies were used in the dilution recommended by the manufacturer. The staining procedure was conducted analogous for all used antibodies and the corresponding isotype controls. After incubating the cells with the respective antibody for 30 minutes on ice in the dark, they were washed with 800 μ L ice-cold PBS/FCS, resuspended in 300 μ L of the latter and stored on ice in the dark until just before the measurements at a FACSCalibur™ respectively a FACSCanto™ II cytometer (both BD Biosciences, Heidelberg, Germany). For the different fluorochromes, the appropriate excitation lasers (pacific blue, brilliant-violet 421: 405 nm (FACSCanto™ II); FITC, PE, alexa fluor 488: 488 nm; APC, alexa fluor 647: 635 nm) and emission filters (pacific blue,; 450/50 BP; brilliant-violet 421: 440/40 BP; alexa fluor 647: 670/20 BP (all FACSCanto™ II); FITC, alexa fluor 488: 530/30 BP; PE: 670 LP; APC: 661/16 BP) were selected, and photomultiplier voltages were adjusted on the autofluorescence of unstained cells. The discrimination of debris and cell agglomerates was performed by forward (FSC) and side scatter (SSC) gating. For reasons of significance, a suitable number of viable single cells (at least 2×10^5) was collected to distinguish rare events. Each experiment was performed at least twice. Data analysis was carried out with FlowJo V10 software (Treestar Inc., Ashland, OR). Viable single cells were gated according to forward and side scatter gates. The expression levels were reported as mean fluorescence intensity, respectively the percentage of cells, revealing a specific property, referred to living single cells.

3.2.5.2 Flow cytometric mitoxantrone efflux assay

The investigation of ABCG2-activity was based on the previously described mitoxantrone efflux assay with minor alterations [3, 36]. In principle, unstained control cells, cells incubated with the fluorescent substrate mitoxantrone, and cells incubated with mitoxantrone and the ABCG2 inhibitor fumitremorgin C (FTC) were compared for each tested cell line.

Cells (approx. 80 % confluence) were detached by trypsinization, washed two times with PBS and resuspended in PBS containing 2 % FCS (PBS/FCS). For each sample, the cell number was adjusted to 5×10^5 cells per 500 μL and 2.5 μL mitoxantrone (1 mM stock solution in 70 % ethanol) were added to reach a final concentration of 5 μM . Subsequently, 5 μL of the 1 mM FTC stock solution (final concentration 10 μM), respectively an equivalent amount of solvent (DMSO) was added. The samples were vortexed and incubated at 37 °C for 30 minutes to allow maximal mitoxantrone uptake. After adding 800 μL cold PBS/FCS (4 °C) and centrifugation (3 minutes, 300 g, 4 °C), the cell pellet was resuspended in 300 μL PBS/FCS and stored on ice in the dark until just before measurement. Mitoxantrone accumulation respectively efflux was determined at a FACSCalibur™ (BD Biosciences, Heidelberg, Germany), using an excitation wavelength of 635 nm while emission was detected at 661/16 nm bandpass filter (FL4). A minimum of 2×10^5 viable single cells, defined by side scatter (SSC) and forward scatter (FSC) gates, were collected and raw data was analyzed with FlowJo V10 software.

3.2.6 Fluorescence activated cell sorting (FACS)

Tumor cells displaying marker expression patterns of interest, such as CD133⁺ in the case of Daoy cells and CD24^{low} for MCF-7/Topo cells, were sorted using a FACSARIA™ IIu (BD Biosciences, Heidelberg, Germany) equipped with a 70 μm nozzle. Except for small alterations in the antibody dilution, the staining procedure was performed exactly as described under 3.2.5.1 except for the omission of isotype controls. In brief: for Daoy, approximately 5×10^7 cells were suspended in 800 μL ice-cold PBS/FCS and 100 μL of the antibody solution (*APC-anti-CD133/1 (AC133)*, Miltenyi) were added. About 9×10^7 MCF-7/Topo cells were suspended in 9 mL ice-cold PBS/FCS, followed by the addition of 125 μL *APC anti-mouse/human CD44* and 400 μL *FITC anti-human CD24* antibody (both BioLegend). After the incubation for 30 minutes on ice in the dark, the cells were washed and resuspended in ice-cold PBS/FCS to a final cell count of approximately 2×10^7 cells/mL. Prior to the sorting procedure, the cells were filtrated through a 35 μm cell strainer (BD Falcon, Heidelberg, Germany) and stored on ice in the dark. The flow- and sorting rate were regularly checked and adjusted to reach a sort efficacy of above 90 %. Cells were sorted using a three-way gating strategy. Initially, dead cells and debris were gated on a two physical parameter dot plot (FSC/SSC) followed by the exclusion of doublets by using pulse processing (FSC-height vs.

FSC-area). Finally, the CD24^{low} subpopulation was gated. Sorted cells were collected in RPMI-1640 medium, containing penicillin-streptomycin (100 U/mL / 100 µg/mL) (Sigma, Taufkirchen, Germany) and subsequently used for further experiments.

3.2.7 Tumorigenicity and growth kinetics of subcutaneous tumors in nude mice

Adult, female NMRI-*nu/nu* mice respectively new born (24-48 hours) NOD.Cg-Prkdc^{scid} Il2rg^{tm1Wjl}/SzJ (NSG) mice were employed to determine the tumorigenicity and growth kinetics of tumor cell lines. For cell implantation, adherently growing tumor cells were washed twice with serum-free RPMI-1640 to remove residual FCS. Depending on the adherence of the particular cell line, the cells were rinsed off with medium or detached by trypsinization. Subsequently, the cells were washed two times with serum-free medium and the cell number was adjusted in RPMI-1640. For sorted cell populations (see 3.2.6), the cell number was adjusted directly after the sorting procedure and the cells were injected immediately. For initial experiments regarding the growth kinetics, usually 1-3 x 10⁶ cells in 100 µL serum-free RPMI-1640 were injected under the thoracic dermis of each mouse. Differences of the *in vivo* tumorigenicity were determined by injecting 10³, 10⁴, 10⁵ respectively 10⁶ MCF-7, MCF-7/Topo, MCF-7/Topo-CD24^{low} (both mouse strains), Daoy wild type and Daoy-CD133⁺ cells (only NMRI-*nu/nu*) into each mouse. The body weight and tumor diameters of each mouse were monitored weekly.

3.2.8 Tumor processing

3.2.8.1 Flow cytometric characterization of tumors

Depending on the growth rate, the tumors were excised after 2-5 months when a sufficient size of at least 20 mm² was reached. Excised tumors were stored immediately in ice-cold FCS-free RPMI-1640. With respect to the following FACS-analysis of cell surface marker expression, tumors were dissected mechanically to prevent interference with proteolytic enzymes. The mechanical dissection was performed in ice-cold medium in 6 cm cell culture plates by cutting the tumor into small pieces. The tumor pieces were strained through a 40 µm mesh (BD Falcon, Heidelberg, Germany) which was subsequently rinsed with the medium in which the tumors were dissected and then washed two times with 5 mL of fresh, ice-cold RPMI-1640. For the exclusion of tissue and major cell agglomerates, the cell suspension was filtrated through a 40-µm cell strainer (BD Falcon) without mechanical pressure prior to the staining procedure (c.f. section 3.2.5.1)

3.2.8.2 Histology

For the determination of the morphology, proliferative activity and formation of metastasis of MCF-7-tumors, tissue samples of liver, lung and the formed tumors of NSG mice were fixed in 4 % formalin, embedded in paraffin and stained as previously described [37]. Briefly, samples were deparaffinized, microwave-heated for 30 min in 0.1 M citrate buffer (pH 7.3) and stained automatically on a Ventana Nexes autostainer (Ventana, Tuscon, USA) by using the streptavidin-biotin-peroxidase complex method and 3,3'-diaminobenzidine as chromogen. The following antibodies in the ultraView Universal DAB Detection Kit (Ventana, Roche) were used: anti-CK8 monoclonal mouse anti-human (clone: 35 β H11) anti-CK18 monoclonal mouse anti-human (clone: DC10) and anti-MIB-1/Ki-67 monoclonal mouse anti-human (clone: MIB-1) (all from DakoCytomation, Glostrup, Denmark). Samples were analyzed microscopically (Zeiss Axiovert 200).

3.3 Investigation of human brain tumor cells

3.3.1 Expression of CD133 and ABC-transporters

CD133 (AC133) is a 5-transmembrane cell surface protein which was originally described as a marker for hematopoietic stem and progenitor cells [38, 39]. CD133-expression was detected on several other cell types such as epithelial cells [40] or acute myeloid leukemia [41]. Moreover, CD133 is frequently used as a marker for brain tumor initiating cells (BTICs) as suggested by Singh et al. [7]. The glioblastoma cell lines U-87MG and LN-18, the astrocytoma cell line SW-1783 and the medulloblastoma cell line Daoy were analyzed for the expression of CD133. Only Daoy cells expressed CD133 (Fig. 3.1) and were therefore additionally analyzed for the expression of ABCB1 and ABCG2. However, both transporters were not expressed in Daoy cells (Fig. 3.1).

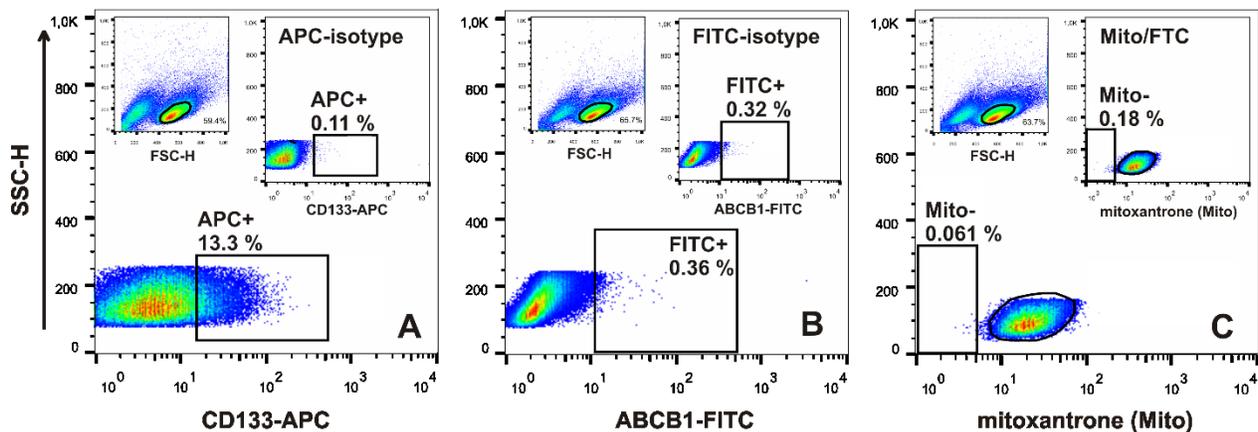


Figure 3.1: Expression of CD133 (A), ABCB1 (B) and ABCG2 (C) in Daoy cells.

A, B: Isotype controls are depicted in inserts. **C:** Mitoxantrone (Mito) efflux assay with Daoy cells. As a control experiment, Mito efflux was inhibited by the ABCG2 modulator fumitremorgin C (inset). The small insets (SSC/FSC) illustrate the applied physical gates (SSC/FSC).

According to the gating strategy illustrated in Fig. 3.1 A, the CD133⁺ subpopulation of Daoy cells was sorted to collect cells for the investigation of the stability of CD133 expression, the *in vitro* growth capacity and sphere formation ability (neurosphere culture) as well as the tumorigenicity in NMRI-*nu/nu* mice. However, Daoy cells proved to be inappropriate for high throughput sorting. The comparatively large cells caused unstable hydrodynamic focusing, resulting in low sorting efficiency at high flow rates. Additionally, the propensity to sediment and form agglomerates at high cell concentrations required an extensive dilution of the cell suspension and periodical resuspension of the cells. Consequently, it was impossible to sort high numbers of cells within an adequate period of time.

3.3.2 Neurosphere culture and stability of CD133 expression in Daoy-CD133⁺ cells

Critical properties of CICs, for example, the growth capacity and sphere formation ability, can be monitored under special *in vitro* culture conditions. For this purpose, a combination of a serum-free culture medium, only supplemented with specific growth factors, and the incubation in non-adherent cell culture conditions (ultra-low attachment plates) was used (sphere culture). Theoretically, these conditions allow for the maintenance of undifferentiated stem cells and the proliferation of multipotent, self-renewing stem cells [7]. By serial passaging under these conditions, it was possible to estimate the growth capacity of the respective cells.

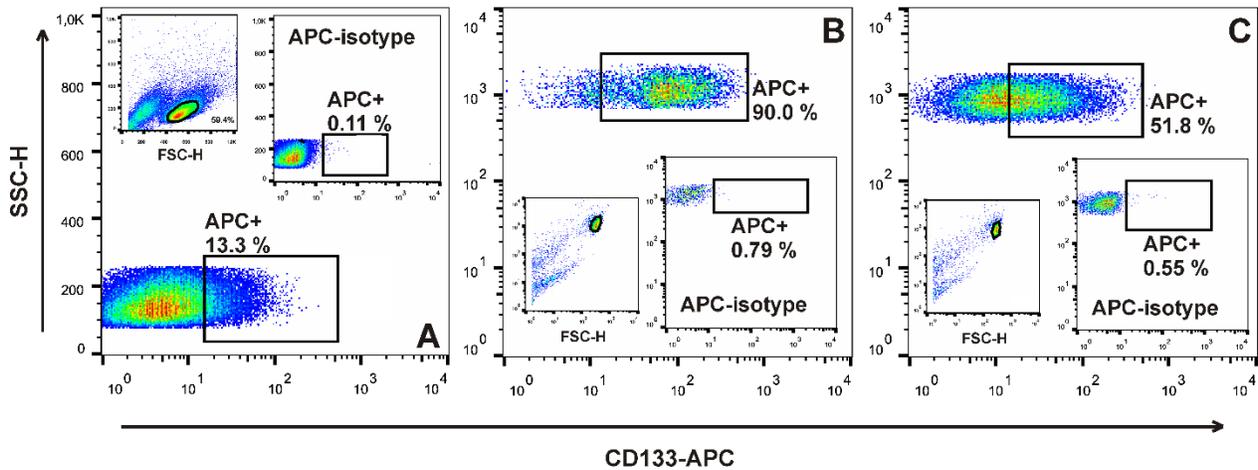


Figure 3.2: Expression of CD133 in Daoy wild type (A), freshly sorted Daoy-CD133⁺ (B) and Daoy-CD133⁺ cells after 14 days in neurosphere culture (C). Isotype controls are illustrated in the insets. Small inserts depict the applied gating strategy (SSC/FSC).

The sphere formation ability in limiting dilution assays was compared for Daoy wild type and Daoy-CD133⁺ cells. Within 24 hours after seeding of more than 300 cells per well, Daoy-CD133⁺ cells formed definite spheres, whereas Daoy wild type cells formed loosely adherent aggregates. However, both cell types could be cultured for several passages over a period of 3 weeks under these conditions without significant changes of growth. The CD133 expression by the sorted Daoy-CD133⁺ cell population, maintained in neurosphere culture for a period of 2 weeks, was re-analyzed by flow cytometry. Interestingly, the phenotype reverted with respect to CD133 expression, resembling the Daoy wild type cells (Fig. 3.2). In contrast to the comparable growth capacity of Daoy wild type and Daoy-CD133⁺ cells, the sphere formation ability in limiting dilution assays was significantly different. Daoy wild type cells formed irregularly shaped aggregates only at cell counts above 100 cells per well, whereas Daoy-CD133⁺ cells formed spheres with diameters of around 100 μm within 9 days (Fig. 3.3), even when only 1 cell/well was seeded (Table 3.1). Interestingly, the maximum number of spheres was comparable, regardless of the number of Daoy-CD133⁺ cells seeded: 300 or 1000 cells/well resulted in 3-5 spheres of different size.

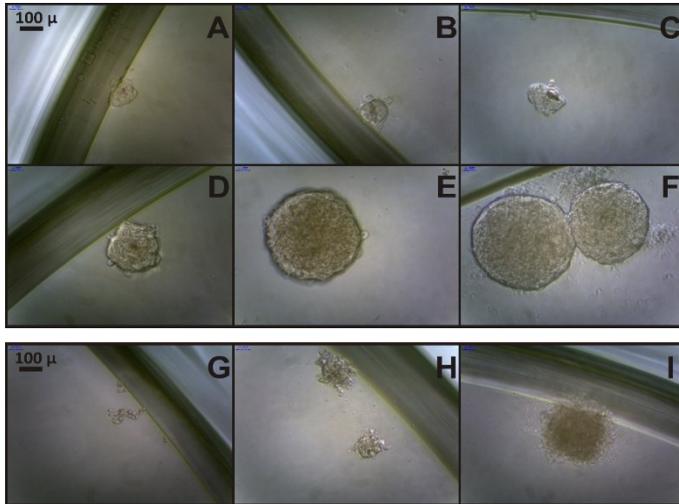


Figure 3.3: Limiting dilution assay with Daoy-CD133⁺ (A-F) and Daoy cells (G-I). Formed spheres, 9 days after seeding of 1 (A), 3 (B), 10 (C), 30 (D), 100 (G), 300 (E, H) and 10³ (F, I) cells/well.

Table 3.1: Number of spheres (\pm SEM) formed per well after seeding of different numbers of CD133⁺ cells (n=2).

Cells/well	Spheres/well
100	2.2 \pm 0.0
30	2.1 \pm 0.3
10	1.4 \pm 0.1
3	0.7 \pm 0.2
1	0.4 \pm 0.01

3.3.3 Tumorigenicity of Daoy and Daoy-CD133⁺ cells

Different numbers of Daoy and sorted Daoy-CD133⁺ cells were subcutaneously injected into nude mice (NMRI-*nu/nu*). The injection of 10⁶ Daoy wild type cells led to the formation of tumors in 7 out of 12 mice within 3 months. When 1.5 x 10⁵ Daoy cells were injected, 1 out of 3 mice developed a tumor. By contrast, tumor formation was not observed after injection of 1.5 x 10⁵ or 1.5 x 10⁴ sorted Daoy-CD133⁺ cells. Due to inefficient cell sorting, higher amounts of Daoy-CD133⁺ cells were not available to perform additional experiments. However, there is no hint to increased tumorigenicity of Daoy-CD133⁺ compared to wild type cells in this mouse model.

3.3.4 Flow cytometric characterization of Daoy tumors

The tumors formed after the injection of Daoy cells did not allow the preparation of single cell suspensions. Accordingly, flow cytometric analysis revealed mainly no live cells, precluding the determination of the expression of CD133, ABCB1 and ABCG2. With regard to possible impacts on flow cytometric measurements, enzymatic methods for tumor disaggregation were not applied.

3.3.5 Discussion

The flow cytometric examination of CD133 expression in human brain tumor cells (Daoy, LN-18, U-87MG, SW-1783) revealed a subpopulation of CD133⁺ cells only in Daoy medulloblastoma cells. A side population expressing ABCG2 or ABCB1 was not observed in Daoy cells. In contrast to the extent of sphere formation of sorted Daoy-CD133⁺ cells, compatible with the concept of CIC-like properties, the ability to form tumors in nude mice was not enhanced. On the contrary, slow restoration of a CD133⁻ subpopulation (50 % after 2 weeks) suggests a preferential growth of CD133⁻ cells. This slow process may result from asymmetric divisions, a stem cell property [42], or the preferential growth of a small number of residual CD133⁻ cells. Plasticity of CICs, as recently reported for breast cancer cells [43], should result in a faster re-adjustment of initial proportions of CD133⁻ and CD133⁺ cells. Additional investigations including markers of BTICs, such as nestin [44], could help to decide whether Daoy-CD133⁺ cells can be considered CICs. However, Daoy cells turned out to be inappropriate for high throughput cell sorting, compromising further experiments regarding cancer initiating cells, although a CD133⁺ subpopulation indicated at least increased clonogenicity *in vitro*.

Previously, CD133⁻ glioma cells were also described as highly tumorigenic [45]. According to a recent publication the validity of CD133 as a universal marker of cancer initiating cells is doubtful [46].

3.4 Investigation of human breast cancer cells

3.4.1 Expression of CD24, CD44, ABCB1 and ABCG2

Since Al-Hajj et al. [11] described a tumorigenic $CD24^{-low}/CD44^{+}$ subpopulation in breast cancer cells, this marker combination is most commonly used for the identification of breast cancer initiating cells (BCICs). All breast cancer cell lines examined in this study were $CD44^{+}$, whereas the expression of CD24 varied in a wide range (Fig. 3.4). The expression of ABCB1 and ABCG2 was additionally determined. Apart from a weak ABCG2-expression in JIMT-1 cells and the induced ABCG2 expression in MCF-7/Topo and MCF-7/Topo/wo cells, none of the cell lines comprised cells expressing ABCB1 or ABCG2. The comparison of results obtained by mitoxantrone efflux and the antibody-based determination of ABCG2 transporters in MCF-7/Topo cells revealed a better distinction between ABCG2⁻ and ABCG2⁺ cells using the specific antibody (Fig. 3.5).

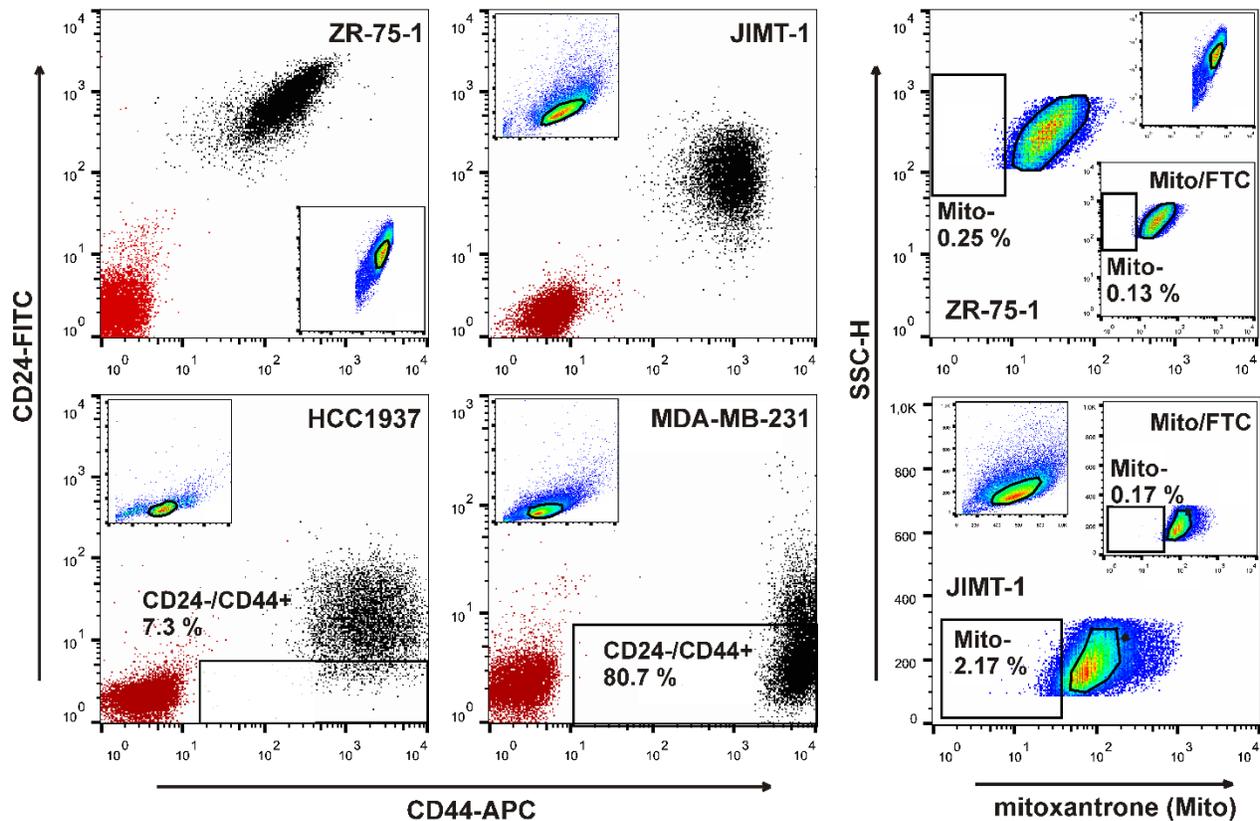


Figure 3.4: Expression of CD24/CD44 (left) and ABCG2 (right) in breast cancer cell lines.

Left panel: Expression of CD24 (y-axes) and CD44 (x-axes) in ZR-75-1, JIMT-1, HCC1937 and MDA-MB-231 cells. The respective isotype controls are depicted in red. The small insets illustrate the applied physical gating strategy (SSC/FSC). **Right panel:** Mitoxantrone efflux as indicator of ABCG2 activity in ZR-75-1 and JIMT-1 cells. Mitoxantrone efflux was inhibited by fumitremorgin C (FTC) (insets). The small insets illustrate the applied physical gates (SSC/FSC).

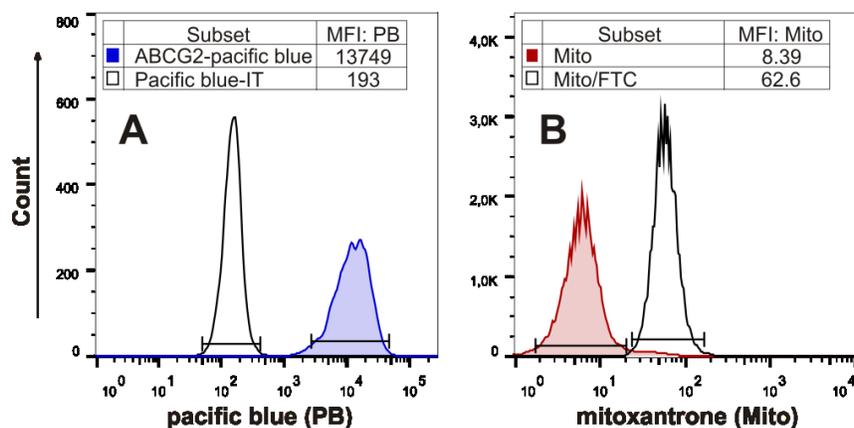


Figure 3.5: Flow cytometric determination of ABCG2 expression in MCF-7/Topo cells with anti-ABCG2-pacific blue antibody (A) and the mitoxantrone-efflux (Mito) assay (B). Isotype control respectively Mito efflux inhibited by fumitremorgin C (FTC) are illustrated as black line. Mean fluorescence intensities (MFI) are shown in the inserted tables.

3.4.2 Tumorigenicity of MDA-MB-231 cells

As stated in section 3.4.1, the triple negative breast adenocarcinoma cell line MDA-MB-231 exhibited the $CD24^{-/low}/CD44^{+}$ expression level which described for breast cancer initiating cells. The tumorigenicity of MDA-MB-231 cells was investigated in NMRI-*nu/nu* mice. The injection of 2×10^6 cells resulted in tumor formation in 4 out of 20 mice within 4 months.

3.4.3 Discussion

The $CD24^{-}/CD44^{+}$ expression pattern is the most frequently used marker combination to identify breast cancer initiating cells. However, the abundance of CD44 on carcinoma cells of diverse entities [18], but particularly on breast cancer cells appears surprising for a cancer initiating cell marker. The investigation of CD24/CD44 expression in diverse breast cancer cell lines revealed that the $CD24^{-/low}$ phenotype is the unique characteristic of the $CD24^{-}/CD44^{+}$ cell population since the vast majority of all cell lines was $CD24^{+}/CD44^{+}$. Only HCC1806 and HCC1937 exhibited a weak CD24 expression, comprising a small $CD24^{low}$ subpopulation. By contrast, more than 80 % of MDA-MB-231 cells were $CD24^{-}/CD44^{+}$. Besides, MDA-MB-231 cells revealed the highest CD44 expression of all examined cell lines. None of the examined cell lines revealed ABCB1 expression, and only JIMT-1 (and the ABCG2 induced MCF-7/Topo) cells comprised a side population expressing ABCG2. Contrary to the expected high tumorigenicity of the $CD24^{-}/CD44^{high}$ MDA-MB-231 cells, the take rates after the injection of 2×10^6 cells in NMRI-*nu/nu* mice were poor (4 tumors in 20 mice), suggesting that this phenotype is not necessarily linked to enhanced tumorigenicity. Notably, a lack of correlation between breast cancer stem cell markers ($CD24^{low}/CD44^{+}/EpCAM^{+}$) and tumorigenicity was recently reported [48].

3.5 Effects of topotecan treatment on MCF-7 cells

Note: This results have been submitted for publication under the title “Topotecan-induced ABCG2 expression in MCF-7 cells is associated with decreased CD24 and EpCAM expression and a loss of tumorigenicity”, prior to the submission of this thesis.

3.5.1 Expression profile of ABCG2 and ABCB1 in different MCF-7 variants

The topoisomerase inhibitor topotecan is widely used in cancer therapy, especially in the treatment of ovarian carcinoma and small cell bronchial carcinoma. It is a known substrate of the breast cancer resistance protein ABCG2 and long-term treatment of cancer cells *in vitro* leads to the induction of ABCG2. The function of the ABCG2 transporter was determined in the mitoxantrone efflux assay, and the expression was confirmed with a pacific blue coupled monoclonal antibody against an extracellular part of the protein (data not shown). While MCF-7 cells did not express the ABCG2 transporter, topotecan-treatment induced a high and homogenous expression of ABCG2 in MCF-7/Topo cells (Fig. 3.6). MCF-7/Topo cells cultivated in topotecan-free medium (MCF-7/Topo/wo cells) maintained the expression level of ABCG2 over a period of approximately 6 weeks. Later on, a small proportion of the cells (approx. 12 % after 10 weeks cultivation in the absence of topotecan) lost the ability of mitoxantrone efflux, whereas high expression of ABCG2 persisted in the majority of the cells. Comparing the ratio of the mean fluorescence intensities (R_{MFI}) between Mito⁺ (ABCG2 mediated mitoxantrone-efflux inhibited by FTC) and Mito⁻ (ABCG2 mediated mitoxantrone-efflux) cells revealed no significant difference for

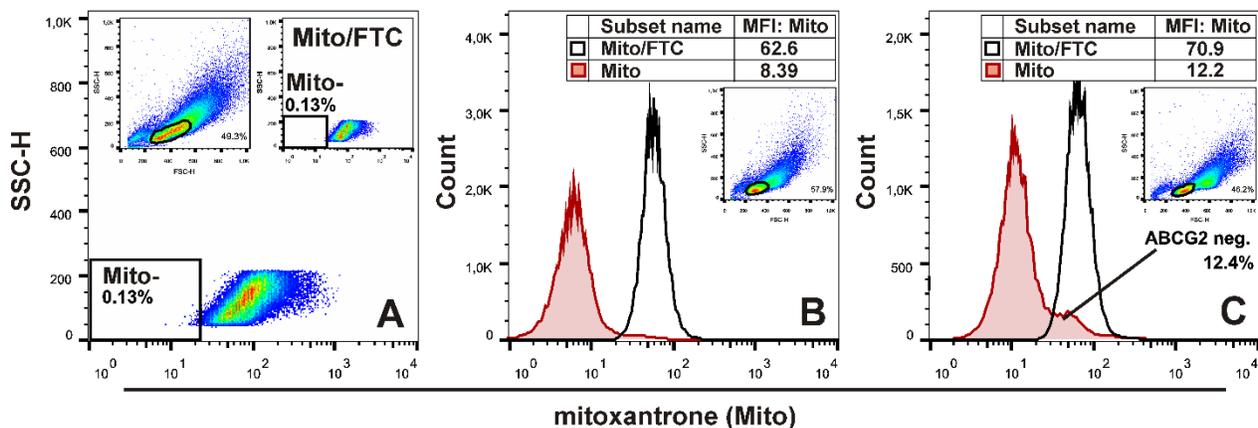


Figure 3.6: Expression of ABCG2 in MCF-7, MCF-7/Topo and MCF-7/Topo/wo cells.

Flow cytometric measurement of Mito efflux as indicator for ABCG2 activity in MCF-7 (A), MCF-7/Topo (B) and MCF-7/Topo/wo (C) (10 weeks in topotecan-free medium) cells. Mito efflux was inhibited by fumitremogin C (FTC) (black line). A: MCF-7 cells do not express ABCG2. B, C: red line: uninhibited ABCG2-mediated efflux of Mito. black line: inhibition of Mito efflux by FTC. The inserts illustrate the applied physical gates (SSC/FSC). Mean fluorescence intensities are given in the inserted tables.

MCF-7/Topo ($R_{MFI} = 5.7 \pm 1.3$) and MCF-7/Topo/wo cells ($R_{MFI} = 5.4 \pm 0.3$) after cultivation over 10 weeks in the absence of topotecan (Fig. 3.6). ABCB1 (P-gp) expression was determined by the use of a monoclonal antibody. Topotecan treatment had no effect on the expression of ABCB1, since both cell variants, MCF-7 and MCF-7/Topo, were ABCB1 negative (data not shown).

3.5.2 Expression of CD24, CD44, EpCAM and HER2 in MCF-7 variants

The HER2 receptor and the cell surface proteins CD24, CD44, and EpCAM are important markers for breast cancer cell phenotyping. MCF-7 cells express all four marker proteins (CD24⁺, CD44⁺, EpCAM⁺, HER2⁺). However, long-term treatment of MCF-7 cells with topotecan caused a significantly altered CD24, EpCAM and HER2 expression profile with the most pronounced effect on CD24: The previously homogenous expression in MCF-7 cells became heterogenous, and the

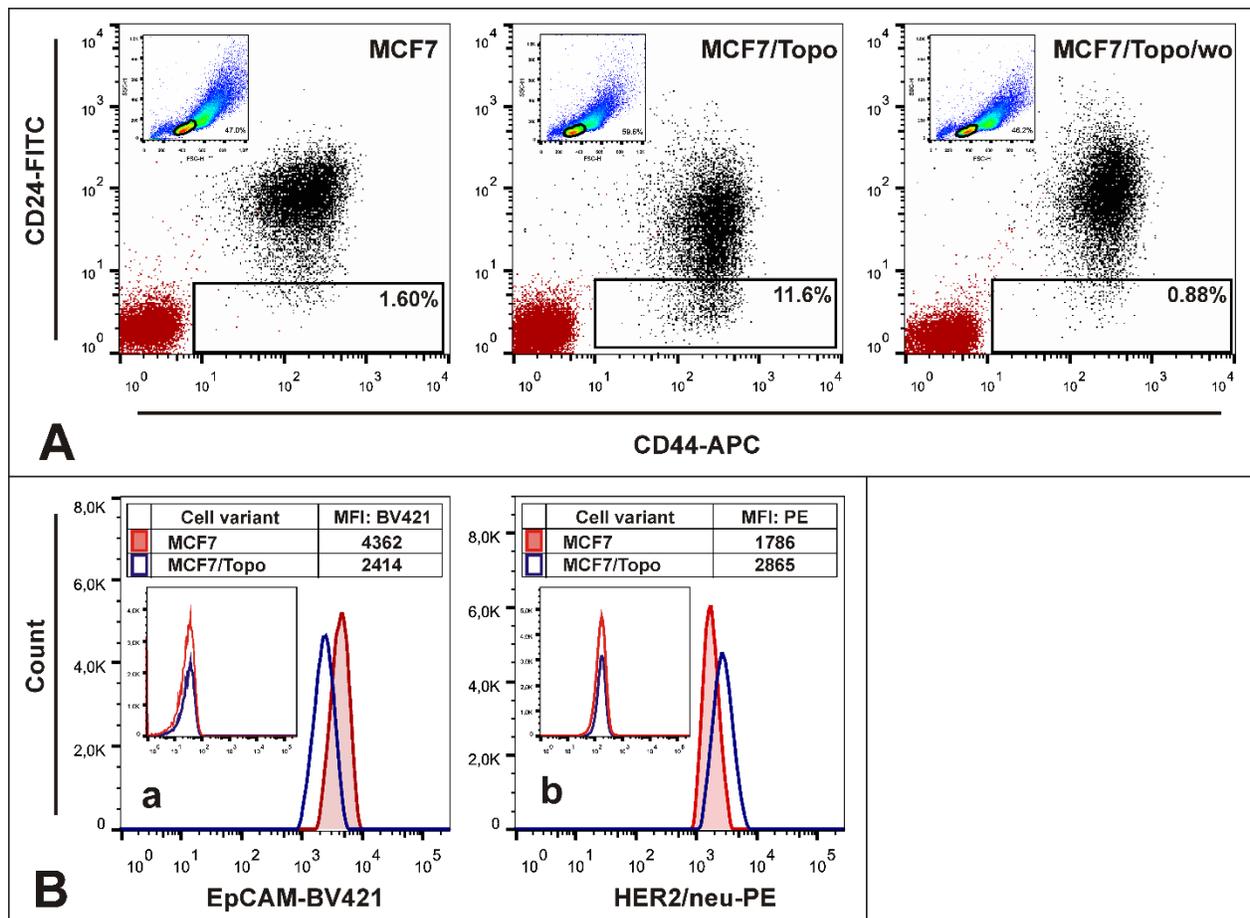


Figure 3.7: Expression of CD24, CD44, EpCAM and HER2 in different MCF-7 variants.

A: Expression of CD24 and CD44 in MCF-7, MCF-7/Topo and MCF-7/Topo/wo (6 weeks in topotecan-free medium) cells. Isotype controls in the dot plots are depicted in red. The applied gating strategy is illustrated in the small inserted two parameter (SSC/FSC) dot blots.

B: Expression of EpCAM (**a**) and HER2 (**b**) in MCF-7 (filled) and MCF-7/Topo (empty) cells. Mean fluorescence intensities (MFI) are shown in the inserted tables. Isotype controls are illustrated in the small inserts.

overall expression level decreased distinctly, resulting in 10 to 15 % CD24^{low} cells. (Fig. 3.7 A) While the expression of CD44 remained virtually unaffected by topotecan treatment, EpCAM was downregulated by 45 % and HER2 was up-regulated by 60 % (Fig. 3.7 B). Unlike the expression of ABCG2, which was quite stable in culture in the absence of topotecan, the expression level of CD24 in MCF-7/Topo/wo cells recovered within 6 weeks to the initially homogenous expression level as observed in MCF-7 cells (Fig. 3.7 A). The CD24^{low} subpopulation of MCF-7/Topo cells was sorted and tumorigenicity as well as mammosphere formation capacity *in vitro* was assessed. A high degree of purity in CD24^{low} sorted cells (referred to as MCF-7/Topo-CD24^{low}) was confirmed (>98 %, data not shown).

3.5.3 Chemosensitivity of MCF-7 and MCF-7/Topo cells

Doxo, MTX, and 5-FU are often used in different combinations with other drugs for chemotherapy of breast cancer [47]. The cytotoxic effect of these cytostatic drugs was determined as an indicator of the induction of ABCG2 transporter expression. The IC₅₀ values of Doxo and MTX on MCF-7 cells were in a low nanomolar range (Table 3.2). On the contrary, MCF-7/Topo cells revealed a 1.5 fold increase in the IC₅₀ value of MTX ($p < 0.05$) and 4.0 fold increased IC₅₀ value of Doxo ($p < 0.001$). However, MTX treatment did not induce a cytotoxic drug effect, neither in MCF-7 nor in MCF-7/Topo cells, but resulted in a decrease of the cell number by 70 % compared to the untreated control. 5-FU exhibited cytotoxic effects in considerably higher concentrations (Table 3.2) than Doxo and MTX, but revealed no significant difference regarding the IC₅₀ value in both cell lines.

Table 3.2: IC₅₀ values (\pm SEM, $n = 3$) in nM of Doxo, MTX, and 5-FU determined in MCF-7 and MCF-7/Topo cells after 72 hours of incubation.

Compd.	Cell variant	
	MCF-7	MCF-7/Topo
Doxo	19.7 \pm 4.7	77.4 \pm 6.9
MTX	19.2 \pm 0.3	32.1 \pm 3.1
5-FU	2440 \pm 235	2325 \pm 435

3.5.4 Mammosphere culture and limiting dilution assays

In adherent cell culture MCF-7 and MCF-7/Topo cells exhibited no significant difference in the growth characteristics: Both cell variants revealed epithelial, polygonal morphology and showed comparable doubling times (MCF-7: 22.8 \pm 1.7 hours; MCF-7/Topo: 23.4 \pm 0.9 hours; $n = 3$). Mammosphere formation under non adherent growth conditions was assessed to examine the tumor initiating capacity of MCF-7, MCF-7/Topo, and MCF-7/Top-CD24^{low} cells. Therefore, the

growth capacity (sphere size) and cell propagation after serial passaging and the ability to form spheres out of small cell numbers (limiting dilution) were determined. All cell variants were principally capable of forming spheres, but MCF-7 cells produced spheres with a larger diameter (200 – 300 μm) compared to MCF-7/Topo and MCF-7/Topo-CD24^{low} cells (diameter 100 μm , respectively), which formed an increased number of spheres instead. However, MCF-7 and MCF-7/Topo cells showed comparable long-term growth characteristics; serial passaging revealed no decline in growth for at least three weeks when cells were passaged 2-3 times a week. In contrast, the CD24^{low} cell population lost the capacity to proliferate after a few passages within 2-3 weeks which was due to the formation of vacuole-like structures and cell disintegration / lysis (Fig. 3.8 left panel). For limiting dilution assays cell numbers between 1 and 10³ (adjusted by serial dilution) MCF-7, MCF-7/Topo and sorted MCF-7/Topo-CD24^{low} cells were seeded per well and incubated for 12 days.

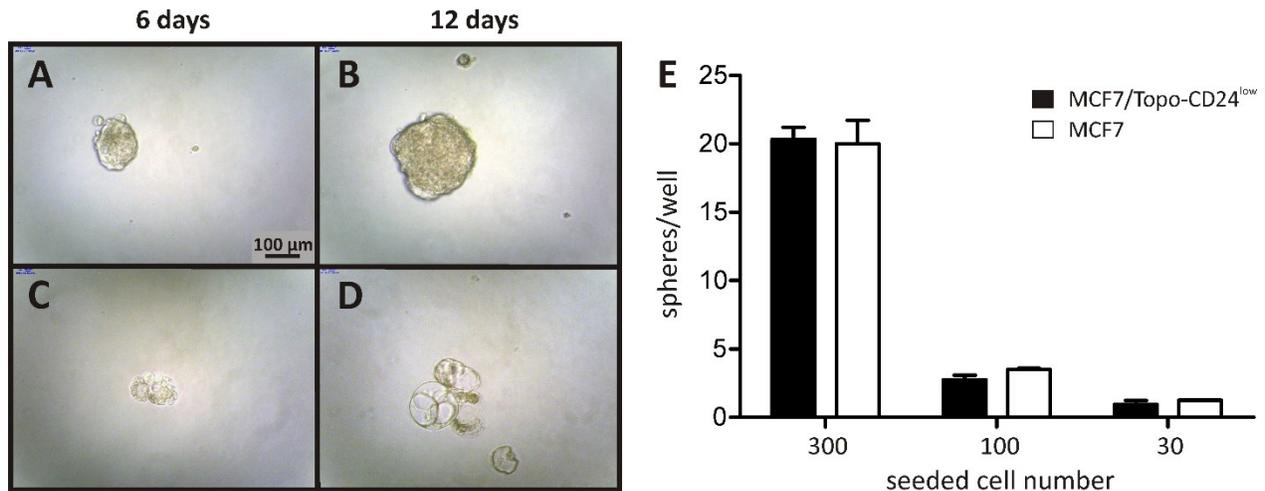


Figure 3.8: Mammosphere culture of MCF-7 and MCF-7/Topo-CD24^{low} cells.

Left panel: Formed spheres at day 6 (A, C) and day 12 (B, D) after seeding of MCF-7 (A, B) and MCF-7/Topo-CD24^{low} (C, D) cells, respectively. MCF-7 cells pursue to grow, while MCF-7/Topo-CD24^{low} underwent cell lysis and sphere disaggregation (D).

Right panel: Numbers of spheres formed per well after limiting dilution of MCF-7 and MCF-7/Topo-CD24^{low} cells. (Mean \pm SEM, n = 3)

The mammosphere forming ability was not significantly different for all cell variants. All cell types formed spheres when 30-1000 cells were seeded, while MCF-7 formed again larger spheres (Fig. 3.8 A and B) than MCF-7/Topo-CD24^{low} cells (Fig. 3.8 C and D). However, number of formed spheres after seeding 30 or 100 cells per well was not significantly different for all examined cell variants (Fig. 3.8 E). In none of the cell types sphere formation was observable when 10 cells or less were seeded.

3.5.5 *In vivo* investigations of MCF-7 variants

Beyond the assessment of mammosphere formation *in vitro* the tumorigenicity of MCF-7, MCF-7/Topo and MCF-7/TopoCD24^{low} cells was determined in newborn NSG respectively adult NMRI-*nu/nu* mice. For this purpose, different cell numbers of the respective cell variant were subcutaneously injected and the tumor growth as well as the body weight was monitored over a period of 3-5 months. The tumorigenicity was considerably higher in newborn NSG-mice than in NMRI-*nu/nu* mice (Table 3.3). The injection of only 1×10^5 MCF-7 cells led to the formation of tumors (3 of 3 mice; Fig. 3.9 A), while at least 1×10^6 cells were required to induce tumor growth in NMRI-*nu/nu* mice (2 of 6 mice). The *in vivo* tumorigenicity of MCF-7 and MCF-7/Topo cells was compared after injecting 1×10^6 cells into NMRI-*nu/nu* mice ($n = 6$). MCF-7 cells formed tumors in 2 of 6 mice, while the injection of MCF-7/Topo cells did not result in tumor growth. Additionally, the tumor-initiating efficacy of MCF-7 and sorted MCF-7/Topo-CD24^{low} cells in NMRI-*nu/nu* mice was compared as a function of 1×10^3 and 1×10^6 injected cells ($n = 3$ for each cell number). Unlike MCF-7 cells, which formed tumors after injection of 1×10^6 cells in 33 % of used mice, the injection of MCF-7/Topo and MCF-7/Topo-CD24^{low} cells did not result in tumor growth. The impaired tumorigenicity of MCF-7/Topo-CD24^{low} cells could be confirmed in the NSG-mouse model (Fig. 3.9 B). MCF-7 cells showed tumor growth even after the injection of only 10^5 cells while MCF-7/Topo-CD24^{low} cells only formed slowly growing tumors after the injection of 10^6 cells.

Table 3.3: Tumorigenicity of MCF-7, MCF-7/Topo and MCF-7/Topo-CD24^{low} cells in NMRI-*nu/nu* and NSG mice. Take rate, expressed as number of tumors per number of injected mice.

Cell number	NMRI- <i>nu/nu</i>			NSG	
	MCF-7	MCF-7/Topo	MCF-7/Topo-CD24 ^{low}	MCF-7	MCF-7/Topo-CD24 ^{low}
10^6	3/9	0/9	0/3	3/3	2/2
10^5	0/3	0/3	0/3	3/3	0/2
10^4	0/3	0/3	0/3	0/3	0/2
10^3	0/3	0/3	0/3	n. d.	n. d.

n. d.: not determined

Tumors were excised after 3-5 months and analyzed by histology and flow cytometry. In histological specimens CK8, CK18, and Ki67 stainings [48, 49] were performed to verify the presence (CK8, CK18) and proliferation capacity (Ki67) of injected, human tumor cells (Fig. 3.9 lower panel). All examined tumors arose at the site of injection and developed epithelial morphology. The histological examination of liver and lung revealed no evidence for metastases. Flow cytometric measurements of MCF-7 tumors revealed a CD24^{low}/CD44^{high} subpopulation ($2.8 \% \pm 1.0$ (SEM, $n = 3$)) while $46.9 \% \pm 2.9$ (SEM, $n = 3$) of solid cancer cells showed a CD24⁻, CD44⁻,

EpCAM⁻, and HER2⁻ phenotype (Fig. 3.9 C). In tumors which were formed after the injection of MCF-7/Topo-CD24^{low} cells, the majority of cells remained a CD24^{low}, CD44⁺, EpCAM⁺ and HER2⁺ expression. Only small subsets of approximately 5 % of CD24⁺/CD44⁺ and 3 % CD24^{low}/CD44^{high} cells were observed. In addition, the rate of EpCAM/HER2 negative cells was lower (27 %) than in MCF-7 tumors (47 %).

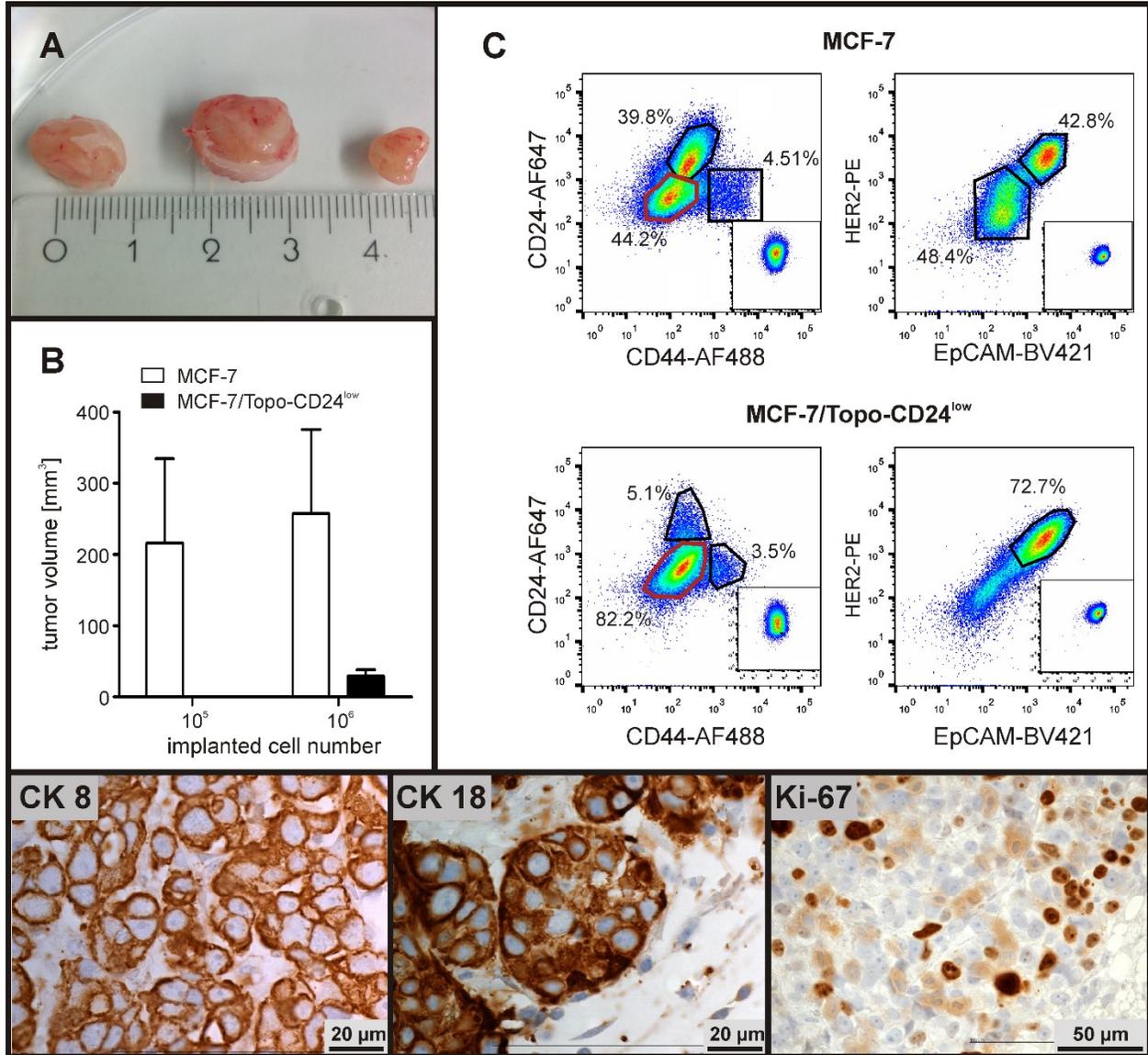


Figure 3.9: Flow cytometric and immunohistological analyses of excised tumors (NSG-mice).

A: Size and shape of MCF-7 tumors formed in different mice. **B:** Final size of formed tumors [mm³] after the injection of different cell numbers of MCF-7 and MCF-7/Topo-CD24^{low} cells. **C:** Flow cytometric analysis of tumor cell suspensions, prepared from MCF-7 or MCF-7/Topo-CD24^{low} tumor tissue. Respective flow cytometric analyses of MCF-7 respectively MCF-7/Topo cells are shown as inserts. **Lower panel:** Expression of CK8, CK18 and Ki67 in MCF-7 tumor cells visualized by immunohistochemistry.

3.5.6 Discussion

ABCG2 transporter expression in tumor cells is induced upon treatment with a variety of cytostatic drugs and enables the efflux of different chemotherapeutics (multidrug resistance). ABCG2 activity can be experimentally identified in so-called “side populations”. Since those “side populations” have been considered to comprise CICs, we examined the effect of cytotoxic treatment on the expression of CD24, CD44, EpCAM, and HER2, which all are supposed to enhance tumorigenicity, tumor progression, and metastasis. For example, a CD24⁻ phenotype has been linked to treatment-resistant breast cancer cells [50, 51], whereas another group reported the enrichment of a highly tumorigenic CD24⁻ subpopulation as a result of anti-Her2 treatment with Trastuzumab in the presence of natural killer cells [52]. Hence, we assessed the capacity of mammosphere formation in MCF-7 cells *in vitro* and tumor outgrowth in immunodeficient mice as a function of treatment with Topo.

We report that long-term exposure of MCF-7 breast cancer cells to Topo led to a homogeneously high expression level of ABCG2 (= MCF-7/Topo), which remained stable for several weeks after removal of Topo. As expected, MCF/Topo cells exhibited an increased chemoresistance against cytostatic drugs like Doxo and MTX (i.e. ABCG2-substrates) [1]. However, the induction of ABCG2 in MCF-7/Topo cells under chemotherapeutic pressure did not result in an increased capacity to form mammospheres. By contrast, MCF-7/Topo cells revealed a considerably decreased tumorigenicity compared to MCF-7 cells. These findings suggest that the induction of ABCG2 upon Topo treatment does not induce cells with enhanced cancer-initiating properties. It has been reported that ABCG2⁻ cells tended to form tumors more efficiently than ABCG2⁺ cells [53]. ABCG2 expression was indicative of fast-cycling tumor progenitors, whereas the ABCG2⁻ population expressed genes which are generally associated with “stemness”. The decreased tumorigenicity of the ABCG2⁺ cell variant MCF-7/Topo as observed in this study may be attributed to the concurrent Topo-treatment related downregulation of CD24 and EpCAM, which has not been analyzed in the context of ABCG2 expression so far. However, it remains to be examined whether Topo treatment causes both, increased ABCG2 and decreased CD24/EpCAM expression, or whether the reduced CD24/EpCAM expression is a consequence of ABCG2 upregulation. Apart from that, the increased CD24^{low}/EpCAM^{low} subpopulation could either originate from the selection of a pre-existing subpopulation or by a real downregulation of those markers. Nevertheless, the association of CD24 with tumor clonogenicity becomes obvious from the considerably reduced mammosphere formation *in vitro* and tumor outgrowth *in vivo* of sorted MCF-7/Topo-CD24^{low} cells. Notably, a positive correlation between high CD24 expression and enhanced cell proliferation *in vitro* and *in vivo* was reported previously [16]. For example, CD24⁺ cells from pleural effusions of different carcinomas revealed increased tumorigenicity in immunodeficient mice [54]. Further evidence for the impact of CD24 was previously provided by the comparison of CD24^{low} and CD24^{high} subpopulations of MCF-7 cells [55]. The CD24^{high}

subpopulation proliferated faster and exhibited increased adhesion and invasiveness *in vitro*. However, the expression of ABCG2 and the sensitivity against doxorubicin did not differ between both cell types. Moreover, CD24 has been suggested to be involved in the metastatic process. It has been described to be involved in tumor cell dissemination/seeding by mediating an interaction of tumor cells with platelets and endothelial cells [56].

The concurrent downregulation of EpCAM in MCF-7 cells upon treatment with Topo, as observed in this study, is likely to enhance the impact of reduced CD24 expression on clonogenicity and tumorigenicity. While high EpCAM expression levels obviously favor cell aggregation [57] lower expression of EpCAM impairs mammosphere formation of MCF-7/Topo cells. Alike CD24, the EpCAM molecule is apparently involved in cell migration and invasive growth [58, 59]. However, the regulation of EpCAM during metastasis is postulated to be more complex: EpCAM positive but also negative circulating and disseminated tumor cells have been reported, although cell migration would rather implicate a low EpCAM expression. Conversely, the subsequent tissue invasion as well as the formation of micrometastases would require distinct EpCAM expression [60].

In the past, a number of distinctive biomarker patterns have been associated to breast cancer stem cells. Nevertheless, a definite or unique stem cell phenotype could not be established, whereas even a pronounced plasticity of cell populations with distinct marker expression patterns (CD24, CD44, EpCAM) was recently reported [43]. Moreover, even a missing correlation of a typical profile of breast cancer stem cell markers (i.e., CD24^{low}, CD44^{high}, ALDH^{high}, EpCAM^{low}) and enhanced tumorigenicity has been reported [61]. The present study substantiates existing evidence for the importance of CD24 and EpCAM in tumor formation and outgrowth. Beyond that, the data indicate that induced ABCG2 expression is not necessarily linked to a more aggressive phenotype.

3.6 Conclusion

Distinct cancer initiating cell marker expression was found in Daoy (brain tumor), MDA-MB-231, HCC1806 and HCC1937 (triple negative breast cancer) cells. The sorted Daoy-CD133⁺ subpopulation revealed also significantly enhanced clonogenicity *in vitro*. However, Daoy cells showed inappropriate characteristics for cell sorting, impeding investigations with high numbers of sorted cells. Furthermore, overall poor take rates in NMRI-*nu/nu* mice hampered the *in vivo* investigations of brain (Daoy) and breast cancer (MCF-7) cell lines. Apart from that, evidence for lacking correlations between established markers of cancer initiating cells and tumorigenicity were provided for MDA-MB-231 cells, which underlines the importance of the combination of several markers with *in vivo* experiments.

The treatment of MCF-7 breast cancer cells with topotecan induces ABCG2-expression and thus leads to increased cellular resistance against cytostatic drugs which are substrates of the transporter molecule. Concurrently, topotecan treatment resulted in a persisting but reversible reduction of CD24 and EpCAM expression. This phenotype modification was not associated with an enhanced, but rather with a reduced clonogenicity and tumorigenicity. Nevertheless, further investigations are necessary to determine whether ABCG2 induction and CD24/EpCAM downregulation, respectively the selection of a CD224^{low}/EpCAM^{low} subpopulation are correlated or independent processes.

3.7 References

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4 Chapter IV

Stabilities of neutral and basic esters of bendamustine in plasma compared to the parent compound: Kinetic investigations by HPLC

Note: As stated in the list of publications, this chapter has already been published prior to the submission of this thesis [1]. For detailed information on the contribution of other authors see also "Acknowledgements".

4.1 Introduction

Bendamustine (**1**, Fig. 4.1), first synthesized in 1963 [2, 3], was brought on the market in East Germany under the name “Cytostasan” for the treatment of chronic lymphatic leukemia (CLL), non-Hodgkin lymphoma (NHL), Hodgkin disease, multiple myeloma (MM) and breast cancer [4-7]. After passing out of mind, it was rediscovered in the 1990s and marketed in North America. Today, bendamustine is approved for CLL [8, 9], indolent NHL [10] and MM [11, 12] in Germany and for CLL and indolent NHL in the US.

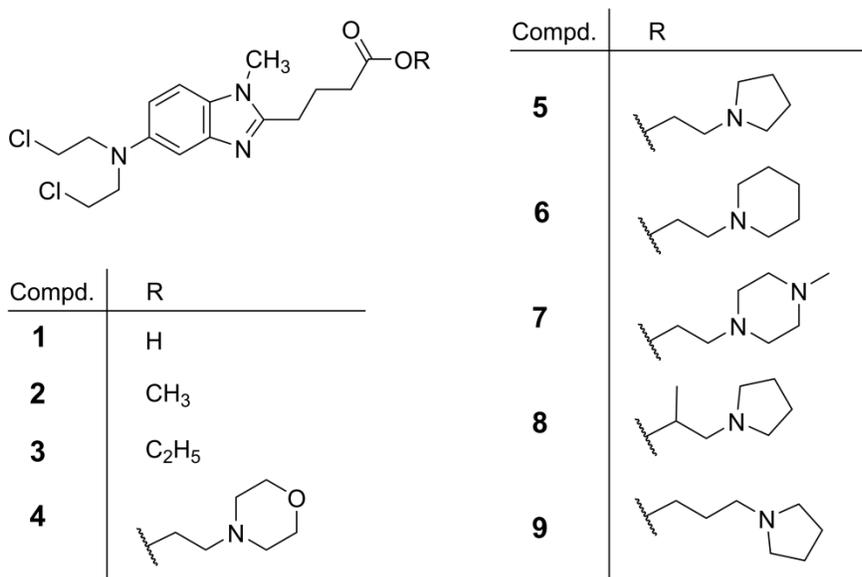


Figure 4.1: Structures of bendamustine (**1**) and the bendamustine esters **2-9**.

As a nitrogen mustard derivative, **1** is instable in aqueous solutions, in particular at neutral or basic pH values [13], resulting in the inactive bis(2-hydroxyethylamino)-substituted benzimidazole derivative upon twofold hydrolysis. Several HPLC methods for the determination of bendamustine and related compounds, degradation products and metabolites are reported in the literature, using UV [14-16], fluorescence [17] and mass spectrometric [18, 19] detection.

Bendamustine is provided as a lyophilized powder for preparation of solutions prior to intravenous administration [20]. Recent patent applications aimed at improved oral bioavailability of bendamustine by optimized formulations [21-23]. Simple alkyl esters (e. g. methyl and ethyl ester) of **1** are mainly known as synthetic intermediates [24, 25]. Dimeric and dendrimeric bendamustine derivatives were investigated with respect to cytotoxicity [26, 27], and C₁ – C₂₄ alkyl esters were reported as prodrugs to prepare formulations for intravenous injection [28]. In a different approach, exploring the potential of derivatives of **1**, very recently, esters of bendamustine (compounds **2-7** (Fig. 4.1) cf. ref. [29]) proved to possess considerably increased

antiproliferative activity compared to the parent compound against a panel of different cancer cell types [29]. The increase in potency strongly depended on the chemical properties of substituent R (cf. Fig 4.1), with highest activities residing in esters bearing a basic moiety such as pyrrolidinoethyl, piperidinoethyl, or morpholinoethyl (mofetil) ester, though ordinary alkyl (methyl, ethyl) esters were also more cytotoxic than the parent compound. In case of these derivatives of **1**, the spectrum of possible degradation products becomes more complex in a biological environment, where, in addition to nucleophiles attacking the nitrogen mustard moiety, hydrolytic enzymes may contribute to degradation. Here we report on the analytical investigation of the stability of esters of **1**, depending on the chemical nature of the substituent R in different media, taking into consideration the possible routes of chemical and enzymatic cleavage suggested in Figure 4.2. For this purpose, we established a RP-HPLC method with fluorescence detection for the determination of bendamustine, bendamustine esters and related hydrolysis products.

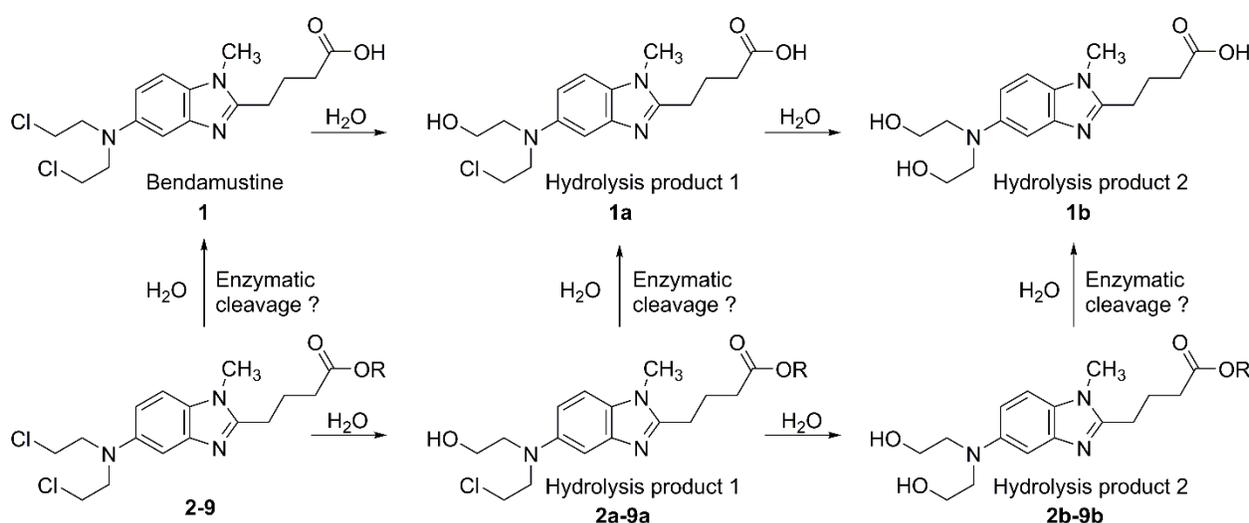


Figure 4.2: Possible routes of degradation of bendamustine (**1**) and esters **2-9** due to hydrolysis of the nitrogen mustard moiety and (enzymatic) ester cleavage, respectively.

4.2 Materials and Methods

4.2.1 Chemicals and reagents

Bendamustine-HCl as well as the esters **2-7** [29] were provided by Arevipharma GmbH (Radebeul, Germany). Compounds **8**⁽¹⁾ and **9**⁽¹⁾ (Fig. 4.1) were synthesized in our laboratory (for synthesis protocols and analytical data cf. master thesis of F. Antoni [30]). Physostigmine, umbelliferone, porcine liver esterase and o-nitrophenyl butyrate were from Sigma-Aldrich (Taufkirchen, Germany). HPLC-grade acetonitrile was obtained from Fisher Scientific (Schwerte, Germany). Demineralized water was purified with a Milli-Q system (Merck KGaA, Darmstadt, Germany). All other reagents, if not otherwise stated, were of analytical grade and from Sigma-Aldrich (Taufkirchen, Germany).

4.2.2 Analytical procedures

4.2.2.1 Instrumentation

HPLC analyses were performed with a system from Thermo Separation Products (Dreieich, Germany), consisting of a SN 4000 Controller, a P4000 pump, an AS3000 autosampler, a Spectra Focus 2000 UV-Vis detector a FL3000 fluorescence detector and a SCM 400 solvent degasser. A Synergi Hydro RP column (4 µm; 80Å; 250 x 4.0 mm), equipped with a guard cartridge (AQ C18, Phenomenex, Aschaffenburg, Germany) was used for separation of the analytes.

4.2.2.2 HPLC conditions

The mobile phase consisted of NaH₂PO₄ (10 mM, pH 3.0) (A), filtrated through a 0.22 µm Steritop-GP filter (Merck, Darmstadt, Germany) and acetonitrile (B). Both eluents were degassed with helium prior to use. For all analyses, the injection volume was 50 µL, and a linear gradient from 5 – 50 % B within 15 min at a flow rate of 0.75 mL/min and a column temperature of 30 °C was used. Fluorescence detection was performed at an excitation wavelength of 330 nm and emission at 420 nm. Void volume was determined with uracil.

4.2.2.3 Standards and calibration

Solutions of **1-9** and of the internal standard umbelliferone were prepared in dimethyl sulfoxide (DMSO). The identities of the reference compounds (“standards”), umbelliferone, bendamustine (**1**) and 4-[5-[bis(2-hydroxyethyl)amino]-1-methyl-1*H*-benzo[d]imidazol-2-yl]butanoic acid (**1b**)

¹ Synthesized and examined by F. Antoni [30].

were confirmed by LC-MS and ^1H NMR spectroscopy, respectively (data not shown). For calibration, solutions of **1-9** (at least 8 concentrations between 5 nM and 30 μM) were prepared by serial dilution of the stock solutions with an aqueous solution of NaH_2PO_4 (10 mM) adjusted to pH 2.0 with phosphoric acid. In case of the internal standard (umbelliferone), nine different concentrations (5 nM - 25 μM), obtained by dilution with phosphate solution pH 2.0, were used for calibration.

4.2.3 Determination of the stability of bendamustine and bendamustine esters

4.2.3.1 Stability in phosphate buffer

Stock solutions (3 mM) of **1-7** were diluted with phosphate buffer (10 mM Na_2HPO_4 , pH was adjusted to 7.4 with phosphoric acid) to a final concentration of 30 μM . After different periods of incubation (0, 7.5, 15 and 30 min) at 37 $^\circ\text{C}$, hydrolysis of the nitrogen mustard moiety was stopped by adding 20 μL of the sample to 180 μL ice cold acidic phosphate solution (10 mM, pH 2.0). After filtration (Phenex-RC 4-mm Syringe Filters 0.2 μm ; Phenomenex), 50 μL of this solution were injected into the HPLC system.

Half-lives were calculated from plots of peak areas versus time, using the “one phase exponential decomposition” option of GraphPad Prism 5 (GraphPad Software, La Jolla, USA).

With respect to the preferred deproteination procedure and the sample preparation and storage, the stability of the analytes was assessed in PBS containing 0.5 M perchloric acid at 4 $^\circ\text{C}$ over a period of 8 h.

4.2.3.2 Stability in human and murine plasma

Human CPD (citrate, phosphate, dextrose)-plasma was a gift from the Bavarian Red Cross. Murine blood was collected from nude mice (NMRI-*nu/nu*) by cardiac puncture using heparin-coated syringes. Blood samples were immediately stored on ice until the cellular components were removed by centrifugation at 4 $^\circ\text{C}$ and 5000 g for 10 min. The plasma samples were stored at -80 $^\circ\text{C}$.

Solutions of **1-9** (20 μL , 1 mM) were mixed with 180 μL of a 200 μM solution of the internal standard (umbelliferone) in phosphate buffered saline (PBS). Afterwards, this mixture was added to human or murine plasma, giving a concentration of 10 μM of **1-9**.

After incubation at 37 $^\circ\text{C}$ for different periods, depending on the individual compound, deproteination of the samples was performed by adding the same volume of ice-cold perchloric acid (1 M) and vortexing. After storage for 20 min at 4 $^\circ\text{C}$ and subsequent centrifugation of the

samples (3 min, 4 °C, 16,000 g; Eppendorf Centrifuge 5415 R, Hamburg, Germany), the supernatants were filtrated (Phenex-RC 4-mm Syringe Filters 0.2 µm; Phenomenex) and stored on ice until HPLC analysis. To inhibit unspecific esterases (butyrylcholinesterase) in human plasma, samples were incubated with physostigmine (100 µM) at 37 °C for 30 min prior to the addition of the esters **4** and **5**.

4.2.3.3 Stability depending on protein concentration

The influence of the protein concentration on the stability of different bendamustine derivatives was determined by analogy with the stability in plasma, except for the replacement of plasma by a solution of porcine liver esterase (5.48 U/mL) in PBS, containing different concentrations of bovine serum albumin (BSA) ranging from 20 mg/mL to 70 mg/mL. The most pronounced effect was observed at concentrations from 30 mg/mL to 50 mg/mL.

4.2.3.4 Determination of the activity of unspecific esterases

Esterase activity was determined photometrically using *o*-nitrophenyl butyrate as substrate [31, 32]. *o*-Nitrophenyl butyrate (40 µL of a 100 mM stock solution in DMSO) was mixed with 1 mL of phosphate buffer (0.1 M, pH 7.4). When the solution had reached 37 °C, 10 µL of murine or human plasma were added, and the increasing absorbance was recorded at 414 nm for 4 min. The volume activity V_A was determined from the initial rate according to the following equation:

$$V_A = (\Delta A / t) \cdot V / (\epsilon \cdot d \cdot v)$$

(Volume activity V_A ($\mu\text{mol} \cdot \text{min}^{-1} \cdot \text{mL}^{-1}$), ΔA = increase in absorbance at 414 nm, ϵ = 3190 L · mol⁻¹ · cm⁻¹ [33], d = path length (cm), V = total volume (mL), v = volume of enzyme/plasma containing solution (mL))

4.3 Results

4.3.1 Validation of the HPLC-method

The analytical procedure was validated according to the ICH guideline Q2 (R1) [34] with respect to selectivity, linearity, limits of quantification (LOQ), limit of detection (LOD), accuracy, precision (repeatability), and robustness (for details cf. supplementary material, Tables S4.1-S4.6). As shown in Figure 4.3, baseline separation of the parent compounds and the corresponding cleavage products was achieved.

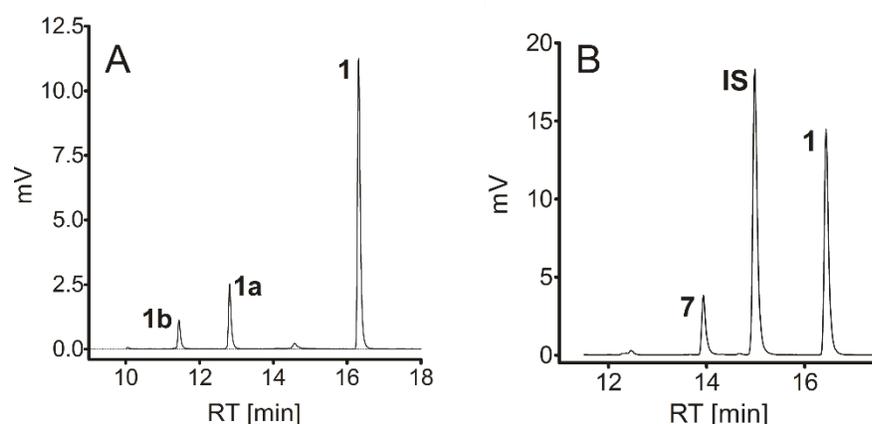


Figure 4.3: Representative chromatograms of **1** (A) and **7** (B) obtained after incubating the compounds in (A) phosphate buffer (pH 7.4) or (B) human plasma for 5 min. Fluorescence at an excitation wavelength of 330 nm and emission at 420 nm.

Table 4.1: Retention factors (k) of the analytes **1-9** and the internal standard (IS), and selectivities (α) of the HPLC method.

Compd. X	$k(X)$	Selectivity ^a			
		$\alpha(X; Xa)$	$\alpha(X; 1)$	$\alpha(X; IS)$	$\alpha(Xa; IS)$
1	2.50	1.54	-	1.14	1.36
2	2.88	1.47	1.15	1.31	1.12
3	3.07	1.48	1.23	1.40	1.06
4	1.89	1.58	1.32	1.16	1.83
5	1.91	1.54	1.31	1.15	1.77
6	1.98	1.61	1.26	1.11	1.79
7	1.86	1.54	1.34	1.18	1.82
8^b	2.01	1.50	1.24	1.09	1.64
9^b	1.81	1.39	1.38	1.22	1.69
IS	2.20	-	-	-	-

^a Selectivities α calculated from the retention factors (k) of the parent compounds X (**1-9**), and the k values of the corresponding products of the first hydrolysis of the nitrogen mustard moiety Xa (**1a – 9a**, structures cf. Fig. 4.2), bendamustine (**1**) and the internal standard (IS) as indicated in parentheses. ^b Data provided by F. Antoni.

The retention factors (k) of **1-9** and of the internal standard, as well as the separation factors (α) are summarized in Table 4.1. Calibration curves (examples, cf. Figure 4.4) of the test compounds **1-9** and the internal standard in the concentration range between 50 nM and 30 μ M were linear with correlation coefficients $r^2 \geq 0.996$ for all test compounds (Table S4.1). The LOD and LOQ of compounds **1-9** and the internal standard were calculated based on peak heights for a signal-to-noise ratio of 3:1 and 10:1, respectively. The LOD was in the range between 1.6 nM and 9.3 nM, and LOQ was between 5.4 nM and 31.2 nM (Table S4.2). Repeatability and accuracy were determined by analyzing three different concentrations within the specified range (250 nM – 10 μ M) (Table S4.3). The relative standard deviation (RSD) was < 5% for all analytes and the accuracy was between 96.3% and 107.1%. Inter-day precision (coefficients of variation were $\leq 6.3\%$) was determined over a period of six months (Table S4.4).

The method was demonstrated to be robust with respect to changes of flow rate (0.7 mL/min and 0.8 mL/min), gradient (0 – 45% and 10 - 55% acetonitrile), pH of the mobile phase (pH 3 and 4), column temperature (25 °C and 35 °C), excitation and emission wavelengths (λ_{ex} : 320 nm, λ_{em} : 410 nm and λ_{ex} : 340 nm, λ_{em} : 430 nm) and duration of treatment with perchloric acid (1 M) for deproteination of BSA (50 mg/mL) containing solutions (Table S4.5, S4.6). The specificity and the baseline separation remained unaffected by varying mobile phase, flow rate and temperature. The peak areas were increased (up to approximately 12%) when the samples were treated with perchloric acid for 25 minutes compared to 15 minutes.

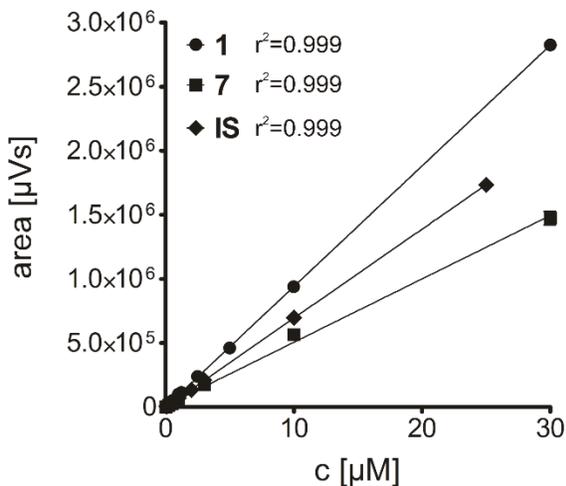


Figure 4.4. Concentration-dependent intensity of the fluorescence signal of **1**, **7** and the internal standard (**IS**) umbelliferon. Analytes were dissolved in acidic phosphate solution (pH 2), prior to HPLC analysis. Excitation at 330 nm, emission at 420 nm.

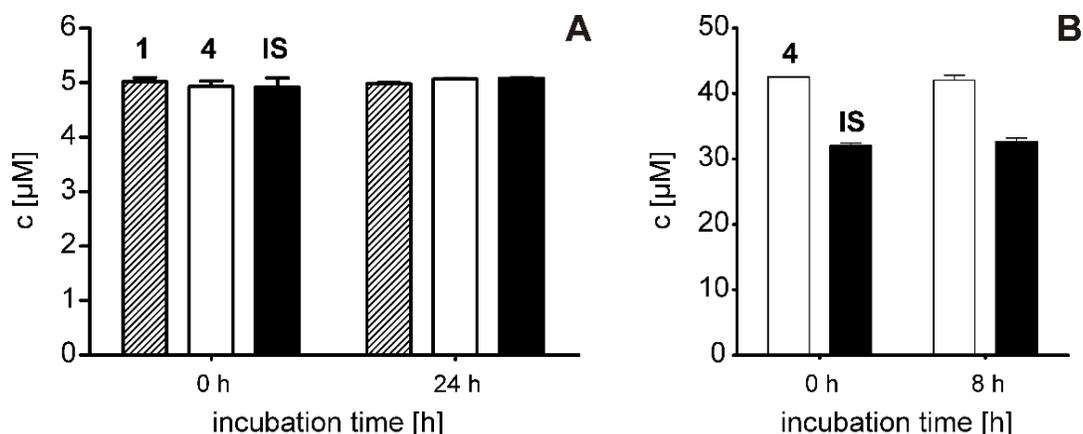


Figure 4.5: Stability of **1**, **4** and the internal standard (**IS**) umbelliferone. Conditions: Storage at 4 °C in phosphate solution pH 2 for 24 hours (**A**) or in phosphate solution containing 0.5 M perchloric acid for 8 hours (**B**) to stop the hydrolysis of the nitrogen mustard group. Mean values \pm SEM (N = 3).

The analytes were stable in 10 mM phosphate buffer (pH 2.0) over a period of 24 h, as shown in Figure 4.5 A for **1**, **4** and umbelliferone as examples. We compared different deproteination procedures (acetonitrile, 2 M trichloroacetic acid, 2 M trichloroacetic acid in ethanol, 1 M perchloric acid) with regard to stability and recovery (Table 4.2) of the analytes. Treatment with 1 M ice-cold perchloric acid gave the best results. With respect to sample processing, stability of the analytes was guaranteed in the presence of 0.5 M perchloric acid over a period of 8 hours (examples cf. Figure 4.5 B). By contrast, in human and, especially, in murine plasma, the ester cleavage was too fast in case of **5-7** and **9** to determine the recovery of the intact compounds; considerable amounts degraded within seconds.

Table 4.2: Recovery from human plasma (mean values \pm SEM, n = 4)

Compd.	% Recovery from human plasma
1	89.2 \pm 0.7
2	74.5 \pm 2.9
3	66.1 \pm 1.6
4	78.7 \pm 4.0
5	n. d. ^a
6	n. d. ^a
7	n. d. ^a
8^b	55.4 \pm 5.0
9^b	n. d. ^a
IS	83.9 \pm 1.6

^a Recovery not determined due to the short half-life.

^b Data provided by F. Antoni.

4.3.2 Stability in phosphate buffer (pH 7.4)

The hydrolysis rate of the nitrogen mustard group of bendamustine, yielding hydrolysis product **1a** (Fig. 4.2) (*4-{5-[(2-chloroethyl)(2-hydroxyethyl)amino]-1-methyl-1H-benzo[d]imidazol-2-yl}butanoic acid*) in the first and hydrolysis product **1b** (Fig. 4.2) (*4-{5-[bis(2-hydroxyethyl)amino]-1-methyl-1H-benzo[d]imidazol-2-yl}butanoic acid*) in the second step [24], is significantly reduced at low temperatures, at acidic pH values and high concentrations of chloride [13]. After incubation

Table 4.3: Half-lives of compounds **1-7** (hydrolysis of the nitrogen mustard group) in phosphate buffer (10 mM, pH 7.4) at 37 °C (mean values \pm SEM, n=3).

Compd.	Half-life [min]
1	10.8 \pm 2.9
2	8.7 \pm 1.4
3	8.9 \pm 1.4
4	10 \pm 1.0
5	8.8 \pm 1.0
6	10.3 \pm 1.3
7	9.0 \pm 1.7

at physiological pH (phosphate buffer 10 mM, pH 7.4, 37 °C), the samples were stored at 4 °C and at pH 2.0 until HPLC-analysis to prevent further hydrolysis.

As expected, the reactivity of the nitrogen mustard against nucleophiles, for instance water, was not affected by esterification of the carboxylic group in bendamustine. The hydrolysis of the parent compounds **1-7**, resulting in the corresponding monohydroxy compounds **1a-7a** occurred at comparable rates with half-lives of about 10 min (Table 4.3).

4.3.3 Stability in murine and human plasma

Interestingly, in murine plasma the hydrolysis rate of the nitrogen mustard moiety in bendamustine was significantly lower ($t_{1/2} = 35 \pm 2.2$ min) than in aqueous solution at physiological pH ($t_{1/2} = 10.8 \pm 2.9$ min). By contrast, the ester moieties of **2-7**, irrespective of the chemical nature of the substituents R, were prone to rapid cleavage ($t_{1/2} < 2$ min; cf. Table 4.4). Thus, in murine plasma the cleavage of the ester group in **2-7**, resulting in bendamustine (**1**), was considerably faster than the hydrolysis of the nitrogen mustard moiety.

Table 4.4: Half-lives of the esters **2-9**, determined in murine and human plasma at 37 °C (mean values \pm SEM, n=3-4)

Compd.	Half-life [min]	
	Murine plasma	Human plasma
2	< 2	63.9 \pm 8
3	< 2	116.6 \pm 9
4	< 2	42.2 \pm 2.7
5	< 2	< 5
6	< 2	< 5
7	< 2	< 5
8^b	n. d.	39.9 \pm 1.9
9^b	n. d.	< 5

n. d.: not determined

^b Data provided by F. Antoni.

Surprisingly, compounds **1-9** behaved differently in human compared to murine plasma. Firstly, the rate of the hydrolysis of the nitrogen mustard group was significantly slower in human plasma (**1**: $t_{1/2} = 131 \pm 10.2$ min). Secondly, the chemical nature of the substituent R substantially determined the rates of ester cleavage (Table 4.4). Representative chromatograms of the decomposition in human plasma of compounds **2**, **4**, **7** and **8** are shown in Fig. 4.6.

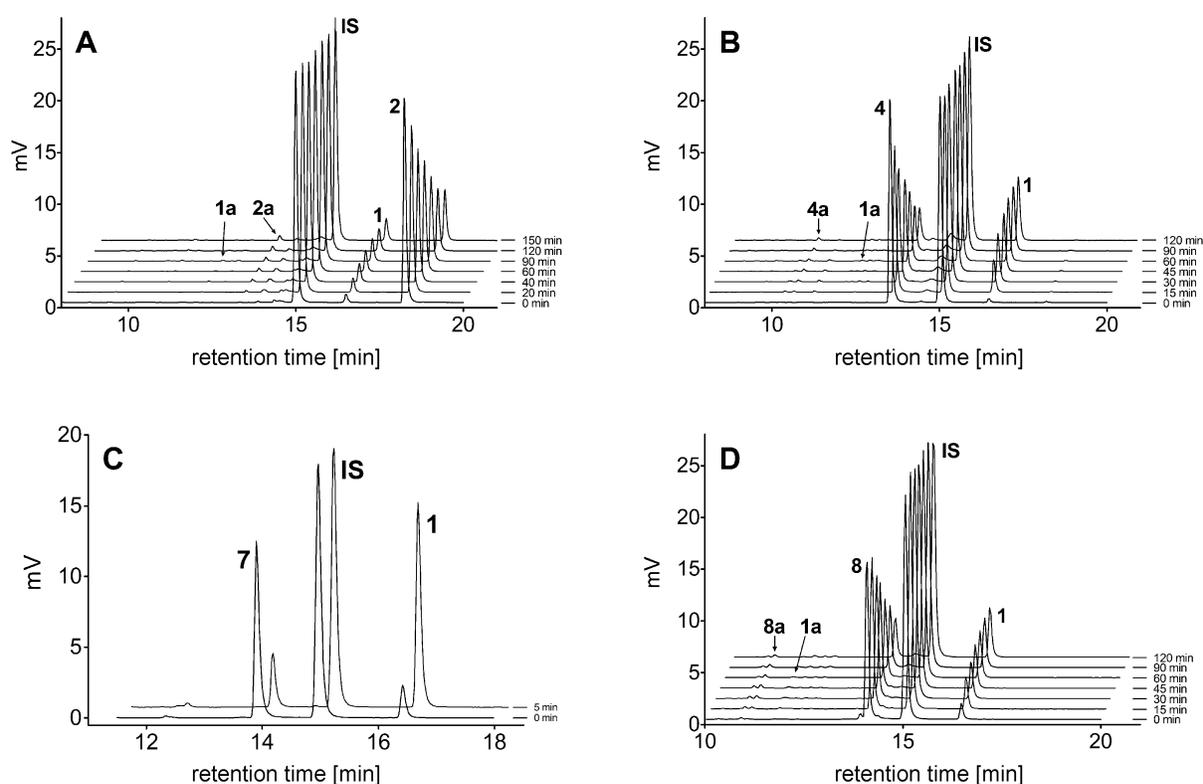


Figure 4.6. Representative chromatograms illustrating the different rates of decomposition of compounds **2** (A), **4** (B), **8** (C) and **7** (D; data provided by F. Antoni) after the incubation in human plasma at 37 °C for different periods of time.

The sequence of degradation of the aminoalkyl esters **5**, **6**, **7** and **9**, comprising unsubstituted ethylene (**5-7**) or trimethylene (**9**) chains, was the same as in murine plasma, yielding bendamustine as the first degradation product, due to rapid ester cleavage ($t_{1/2} < 5$ min, Table 4.4), followed by hydrolysis of the nitrogen mustard group.

The neutral alkyl esters **2** ($t_{1/2} = 63.9 \pm 8$ min) and **3** ($t_{1/2} = 116.6 \pm 9$ min), the weakly basic morpholinoethyl (mofetil) ester **4** ($t_{1/2} = 42.2 \pm 2.7$ min) and the branched, more basic pyrrolidino-substituted ester **8** ($t_{1/2} = 39.9 \pm 1.9$ min) showed considerably increased half-lives in human plasma compared to murine plasma (Table 4.4). The fast cleavage of **5**, **6**, **7** and **9**, reminiscent of substrates of unspecific cholinesterases, in comparison to the retarded degradation of neutral,

less basic or branched esters (**2**, **3**, **4** and **8**) suggested an enzyme-mediated hydrolysis of the ester bond. Indeed, the inhibition of unspecific plasma esterases by physostigmine resulted in prolonged half-lives of compounds **4** and **5**, which were investigated as examples. The effect of physostigmine was much more pronounced in case of the heterocyclic basic ester **5** (48.3 compared to <5 min) than for the mofetil ester **4** (52.8 compared to 42.2 min) (Fig. 4.7).

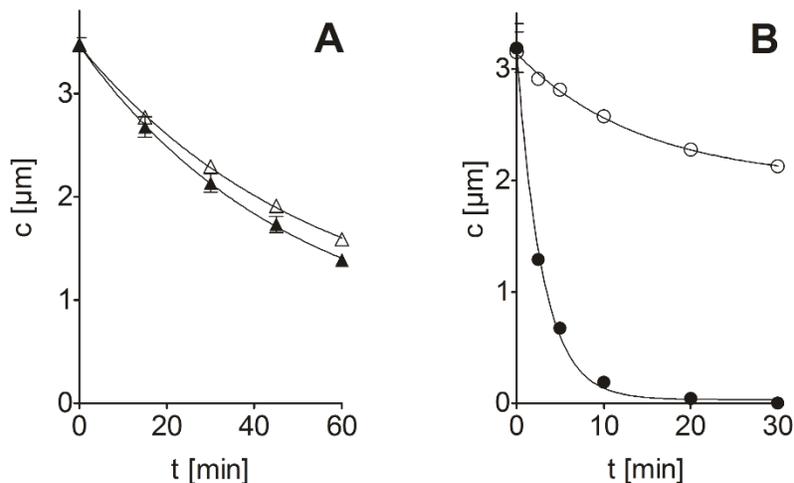


Figure 4.7. Decomposition of bendamustine mofetil ester **4** (A) and bendamustine pyrrolidinoethyl ester **5** (B) in human plasma at 37 °C in the presence (empty symbols) and the absence (filled symbols) of the esterase inhibitor physostigmine (100 μM).

4.3.4 Influence of protein concentration and esterase activity on the degradation of bendamustine esters

The discrepancies between the decomposition kinetics of **1-7** in human and murine plasma supported the hypothesis that species-specific differences in protein content and esterase activity were associated with this phenomenon. Therefore, we determined the decomposition kinetics of **1** and **4** as examples in the presence of both porcine liver esterase and different amounts of bovine serum albumin. The results revealed a significant correlation between the half-lives and the albumin concentration. The increase in the albumin concentration from 30 mg/mL to 50 mg/mL led to the prolongation of the half-life by 42 % for compound **1** ($t_{1/2} = 120.4 \pm 7.0$ min vs. $t_{1/2} = 170.7 \pm 14.2$ min) and by 70% for compound **4** ($t_{1/2} = 7.3 \pm 1.3$ vs. $t_{1/2} = 12.4 \pm 0.5$) (Fig. 4.8).

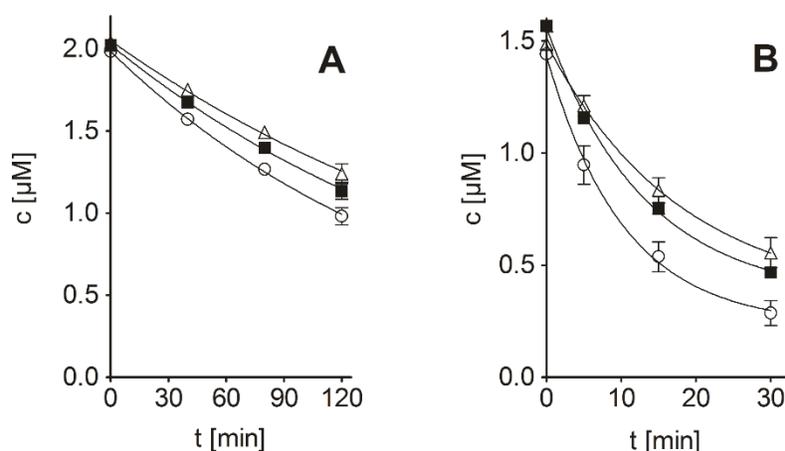


Figure 4.8: Decomposition of bendamustine **1** (A) and bendamustine mofetil ester **4** (B) in a solution of porcine liver esterase in phosphate buffer (pH 7.4) containing different concentrations (circles: 30 g/L; squares: 40 g/L; triangles: 50 g/L) of BSA. Compound **1** decomposed by hydrolysis of the nitrogen mustard moiety, whereas compound **4** primarily underwent ester cleavage to give **1** under the conditions of this experiment (cf. Fig. 4.2).

Additionally, the determination of the amounts of unspecific esterases revealed nearly 1.5 fold higher activity in murine plasma ($6.04 \pm 0.2 \mu\text{mol}\cdot\text{min}^{-1}\cdot\text{mL}^{-1}$) compared to human plasma ($4.26 \pm 0.23 \mu\text{mol}\cdot\text{min}^{-1}\cdot\text{mL}^{-1}$). In this context it should be noted that the protein content of murine plasma (total protein: 40-60 g/L, albumin: 20-35 g/L [35]) is lower than that of human plasma (total protein: 64-83 g/L, albumin: 35-55 g/L [36]). The combination of both, the lower protein content and the higher esterase activity appear to be responsible for the considerably lower stability of bendamustine and esters of **1** in murine plasma.

4.4 Discussion

Previously, solid phase extraction and mass spectrometric detection, requiring special equipment, were applied for the analysis of bendamustine in biological material [18, 37]. We aimed at establishing a simple, specific, sensitive and fast method to analyze esters of bendamustine (**2-7**) and the corresponding decomposition products, including **1**, in aqueous solution and in biological material. Pencheva et al. (**1**: LOD 9.6 µg/mL, LOQ 34 µg/mL) [15] and Mathrusri Annapurna et al. (**1**: LOD 0.24 µg/mL, LOQ 0.73 µg/mL) [15] described HPLC methods using UV detection. Xie et al. [17] reported a more sensitive, convenient method for the determination of **1** and γ -hydroxy-bendamustine in plasma, using perchloric acid for deproteination and fluorescence detection (**1**: LOD 5 ng/mL in plasma and urine). As the fluorescence-based method was most promising, we adapted this approach especially to the quantification of the new bendamustine esters and the corresponding hydrolysis products. Regarding the sensitivity and recovery of bendamustine, the modified HPLC procedure described here was comparable to the data reported by Xie et al. [17]. It should be stressed that, according to the optimized protocol, fluorescence detection allowed for the specific quantification of bendamustine derivatives with high signal-to-noise ratios despite protein precipitation with perchloric acid, which was reported unfavorable by Dubbelmann et al. [18].

Recently, several esters (**2-7**) of the cytostatic bendamustine (**1**) were demonstrated to be significantly more potent than the parent compound as cytostatic agents against a variety of human cancer cell types in vitro [29]. These results suggest that the antiproliferative activity has to be attributed to the derivatives on their own rather than to the conversion of the esters to bendamustine as the active principle. With respect to exploring the potential value as anticancer drugs in vivo, information on the stability of the esters in biological matrices are of utmost importance.

The alkylating activity of nitrogen mustard derivatives such as **1-9** originates from the bis(2-chloroethylamino) group, which is, apart from the reaction with nucleophiles, a labile group, undergoing hydrolysis in solutions, especially at neutral or basic pH [13]. This reaction can be stopped by addition of acid, as demonstrated under the conditions of the HPLC analysis, when acidic phosphate solution was used as a component of the mobile phase, and perchloric acid and low temperature to precipitate plasma proteins. The established HPLC method enabled the analysis of the stability of **1** and corresponding neutral and basic esters in buffer, plasma, and against porcine liver esterase in the presence of different amounts of albumin.

In buffer at pH 7.4 hydrolysis of the nitrogen mustard group was the prevailing reaction, occurring with comparable half-lives of around 10 min (**1-7**, Table 4.3). Interestingly, this decomposition reaction slowed down by a factor of up to 13 in the presence of plasma protein. This can be

explained by protection of the parent compounds in a hydrophobic environment when bound to albumin [38].

Due to different rates of hydrolysis of the ester and the nitrogen mustard moiety, the cleavage of the derivatives proved to be much more complex than that of **1**. Cleavage of the ester group was slowest in case of the ethyl ester (**3**, $t_{1/2}$ 116 min) followed by compounds **2** ($t_{1/2}$ 64 min), **4** ($t_{1/2}$ 42 min) and **8** ($t_{1/2}$ 40 min), i.e., in these cases the degradation kinetics was dominated by the hydrolysis of the nitrogen mustard moiety. These compounds having a more stable ester group (**2**, **3**, **4** and **8**) were converted to **1a** (**1b** not determinable with this method) via bendamustine (**1**) and underwent hydrolysis at the nitrogen mustard moiety to give **2a**, **3a**, **4a**, **8a** as intermediates (examples cf. Fig. 4.6). By contrast, the cleavage of the ester bond was rapid ($t_{1/2}$ <5 min) in case of the 2-pyrrolidinoethyl (**5**), 2-piperidinoethyl (**6**), 2-(4-methylpiperazino)ethyl (**7**), and 3-pyrrolidinopropyl ester (**9**). For compounds such as **7**, the corresponding hydrolysis product **7a** was not detectable due to very fast conversion to **1** (Fig. 4.2). It should be mentioned that unidentified reaction products were observed to a minor extent in the chromatograms (Fig. 4.6). Chemical reactions of the alkylating molecular species with bionucleophiles in the incubation mixture could have taken place, lowering the portion of residual analytes detectable with the applied method.

The enzymatic hydrolysis in human plasma strongly depended on the presence or absence of a strongly basic group (cf. compds. **5-9**) and on sterical hindrance (cf. compd. **8**). These structural characteristics are reminiscent of substrates of unspecific cholinesterases [39]. Accordingly, the enzymatic cleavage was strongly inhibited by physostigmine in case of **5**, but only slightly affected for the less basic ester **4**. Interestingly, in contrast to human plasma, the same analytes were much less stable in murine plasma ($t_{1/2}$ <2 min). These differences can be explained by the lower protein content and the higher enzymatic activity in murine compared to human plasma. The determination of the decomposition kinetics of **1** and **4** in presence of porcine liver esterase and different concentrations of bovine serum albumin (Fig. 4.8) revealed that the concentration of albumin had a significant effect on the hydrolytic stability of both, the nitrogen mustard and the ester bond.

4.5 Conclusion

The properties of the introduced ester moieties define whether bendamustine esters are preferred substrates of unspecific plasma esterases and thus strongly determine the stability of the particular ester bond. The stabilization of bendamustine and esters by plasma proteins became obvious from the studies in the presence of different amounts of bovine serum albumin and from the investigations in murine and human plasma, which considerably differ regarding protein content. The activity of unspecific esterases was much higher in murine compared to human plasma. In concert, these species-dependent differences proved to cause tremendous discrepancies regarding the stability of the title compounds. This phenomenon has to be taken into account with respect to translational animal studies, and may also be relevant to pharmacological investigations of structurally diverse bioactive compounds other than cytostatics.

4.6 Supplementary material: Method validation

Table S4.1: Slopes and intercepts of the equation $y = a \cdot x + b$. Coefficients of determination (r^2) calculated by simple linear regression from the data of three independent experiments.

Compound	a [$\mu\text{V} \cdot \text{s} \cdot \mu\text{M}^{-1}$]	b [$\mu\text{V} \cdot \text{s}$]	r^2
1	93570 ± 281.1	275.5 ± 941.2	0.9999
2	103700 ± 572.3	-755.7 ± 1916	0.9997
3	65030 ± 1107	-5786 ± 4702	0.9998
4	23440 ± 1143	8122 ± 4899	0.9926
5	45260 ± 782.8	2077 ± 3357	0.9983
6	66850 ± 1286	7719 ± 5516	0.9963
7	56700 ± 344.2	-2240 ± 1476	0.9994
8 ^a	32990 ± 184.2	-2073 ± 794.5	0.9998
9 ^a	35970 ± 146.1	-3619 ± 630.1	0.9999
IS	69430 ± 218	-715.4 ± 1976	0.9994

^a Data provided by F. Antoni.

Table S4.2: Limit of detection (LOD) and quantification (LOQ) calculated with a signal to noise ratio of the peak heights of 3:1 and 10:1, respectively.

Compound	LOD [nM]	LOQ [nM]
1	1.6	5.4
2	2.4	7.9
3	8.0	26.6
4	9.3	31.2
5	5.0	16.7
6	4.8	16.1
7	4.5	15.1
8 ^a	5.0	16.5
9 ^a	4.9	16.3
IS	3.6	12.1

^a Data provided by F. Antoni.

Table S4.4: Inter-day precision of the HPLC-method over a period of 6 months. (Coefficient of variation (CV) in %, n = 6)

Compound	CV (%)
1	2.37
2	5.26
3	3.14
4	7.00
5	3.58
6	3.32
7	4.46
8 ^a	6.30
9 ^a	4.73
IS	4.91

^a Data provided by F. Antoni.

Table S4.3: Repeatability and accuracy of the HPLC-method. (n = 3-4)

Compound	Repeatability			95 % confidence interval	Accuracy [%]
	nominal concentration [μM]	determined concentration ± SD [μM]	RSD [%]		
1	10	10.03 ± 0.146	1.45	9.670 - 10.39	99.7
	1	1.079 ± 0.018	1.63	1.035 - 1.122	92.7
	0.25	0.273 ± 0.016	5.77	0.234 - 0.312	91.7
2	10	9.948 ± 0.214	2.15	9.417 - 10.48	100.5
	1	1.134 ± 0.065	5.71	0.973 - 1.294	88.2
	0.25	0.243 ± 0.008	3.32	0.223 - 0.263	102.7
3	10	10.05 ± 0.071	0.71	8.195 - 11.31	99.5
	1	1.017 ± 0.031	3.06	0.861 - 1.097	98.3
	0.25	0.238 ± 0.007	2.76	0.215 - 0.252	105.0
4	10	9.750 ± 0.626	6.42	8.195 - 11.31	102.6
	1	0.979 ± 0.047	4.84	0.861 - 1.097	102.1
	0.25	0.234 ± 0.001	3.25	0.215 - 0.252	107.1
5	10	9.910 ± 0.202	2.04	9.410 - 10.41	100.9
	1	0.983 ± 0.055	5.61	0.846 - 1.120	101.7
	0.25	0.254 ± 0.011	4.18	0.228 - 0.280	98.4
6	10	9.885 ± 0.177	1.79	9.440 - 10.33	101.2
	1	1.024 ± 0.045	4.37	0.913 - 1.135	97.7
	0.25	0.243 ± 0.012	4.97	0.213 - 0.273	102.8
7	10	9.893 ± 0.142	1.43	9.540 - 10.24	101.1
	1	0.916 ± 0.027	2.93	0.849 - 0.983	109.2
	0.25	0.256 ± 0.011	4.17	0.229 - 0.282	97.7
8^a	10	9.918 ± 0.450	4.54	8.800 - 11.04	100.8
	1	0.974 ± 0.012	1.23	0.944 - 1.004	102.7
	0.25	0.244 ± 0.001	0.21	0.243 - 0.245	102.5
9^a	10	10.38 ± 0.283	2.73	9.674 - 11.08	96.3
	1	0.995 ± 0.01	0.98	0.972 - 1.020	100.5
	0.25	0.256 ± 0.01	2.84	0.238 - 0.273	97.7
IS	10	10.05 ± 0.143	1.42	9.700 - 10.41	99.5
	1	0.987 ± 0.015	1.52	0.950 - 1.025	101.3
	0.25	0.256 ± 0.012	4.84	0.225 - 0.286	97.7

^a Data provided by F. Antoni.

Table S4.5: Effect of the flow rate, the composition and the pH of the mobile phase and the column temperature on the retention factors relative to the respective retention factors of the optimized method (100%).

Compound	Flow rate		Gradient ^{a)}		pH = 4	Temperature	
	0.7 mL/min	0.8 mL/min	0 - 45% MeCN	10 - 55% MeCN		25 °C	35 °C
1b	96.64	104.80	80.85	81.35	100.44	102.79	98.38
1	94.72	105.39	76.85	100.40	101.64	101.36	100.16
2	94.77	105.54	73.12	102.00	105.24	101.16	100.45
4	94.72	105.43	77.61	96.59	112.31	101.41	99.97
IS	95.66	104.31	78.07	99.12	100.72	102.53	98.74

^{a)} MeCN = acetonitrile

Table S4.6: Effects of the flow rate, the composition and the pH of the mobile phase, the column temperature, the wavelengths and the duration of treatment with perchloric acid on the peak areas relative to the respective peak areas of the optimized method (100%).

Compound	Flow rate		Gradient ^{a)}		pH = 4	Temperature		Wavelengths ^{b)}		Incubation period ^{c)}	
	0.7	0.8	0-45 %	10-55 %		25 °C	35 °C	λ_{ex} : 320	λ_{ex} : 340	15 min	25 min
	mL/min	mL/min	MeCN	MeCN				λ_{em} : 410	λ_{em} : 430		
1b	100.24	97.29	109.00	99.35	106.52	100.25	98.15	69.47	113.42	102.30	107.62
1	107.07	94.43	96.78	101.97	103.10	98.82	99.06	90.40	79.87	99.57	114.93
2	108.44	94.12	102.05	101.11	100.23	98.92	99.03	89.96	80.33	102.21	115.63
4	107.22	92.00	97.32	97.57	102.67	98.58	97.87	88.09	82.48	101.16	114.86
IS	106.91	96.08	101.15	99.39	101.53	97.01	99.17	46.59	156.49	100.50	113.30

^{a)} MeCN = acetonitrile; ^{b)} λ_{ex} : excitation wavelength, λ_{em} : emission wavelength; ^{c)} Incubation period for deproteination of BSA (50 mg/mL) containing solutions with perchloric acid (1 M).

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5 Chapter V

Esters of bendamustine are by far more potent cytotoxic agents than the parent compound against human sarcoma and carcinoma cells

Note: As stated in the list of publications, this chapter has been submitted for publication prior to the submission of this thesis. For detailed information on the contribution of other authors see also "Acknowledgements".

5.1 Introduction

The alkylating agent bendamustine (**1**, Fig. 5.1) was synthesized in 1963 [1, 2] and developed as an anticancer drug in the German Democratic Republic [3-5]. In the 1990s bendamustine got into the focus of research again [6]. It is approved for the treatment of chronic lymphocytic leukemia (CLL) [7, 8], indolent non-Hodgkin lymphoma (NHL) [9] and multiple myeloma (MM) [10, 11] or as second line therapy of refractory diseases [12-14] in various countries. Current clinical trials suggest beneficial effects in the treatment of solid cancer types such as breast cancer [15] or small-cell lung cancer [16]. Most treatment regimens include bendamustine in combination with other anticancer drugs including biologicals [17-20], for instance rituximab [17, 19, 20]. It has been hypothesized that, apart from the alkylating nitrogen mustard group (N-Lost), the benzimidazole scaffold may contribute to the antitumor activity for instance, due to antimetabolite properties, by facilitating nuclear transport or inhibiting DNA repair [6]. Besides the alkylation of DNA, causing strand breaks, bendamustine induces the expression of p53 [21], triggers apoptosis [22] and down-regulates mitotic checkpoints, leading to mitotic catastrophe [23, 24]. Recent patent applications aimed at formulations for oral administration [25-27]. Alkyl ($C_1 - C_{24}$) esters of bendamustine were reported as potential prodrugs for intravenous application [28], and biodegradable polyphosphoesters were described as an approach to stabilize bendamustine in solution [29]. Another approach aimed at increasing the cytotoxicity by constructing dimeric and dendrimeric bendamustine derivatives [30].

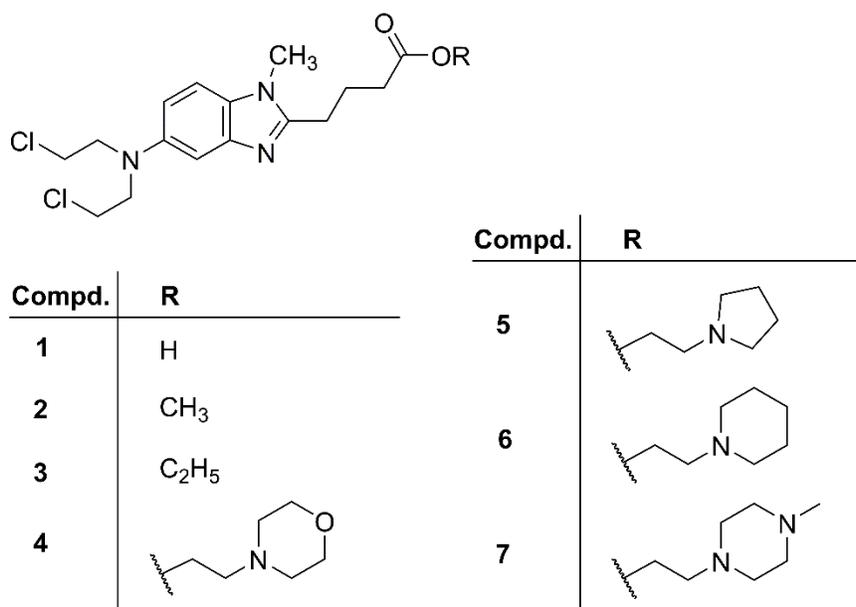


Figure 5.1: Structures of bendamustine (**1**) and the bendamustine esters **2-7**.

Exploring the properties of derivatives of bendamustine (**1**), esters of bendamustine (**2-7**, Fig. 5.1) [31] comprising basic moieties (**4-7**) were prepared as potential prodrugs with higher solubility compared to simple esters such as compounds **2** and **3**, which were mainly prepared as synthetic intermediates [32, 33]. Very recently, we reported on the stability of the nitrogen mustard and the ester moieties in compounds 2-7 against hydrolysis and enzymatic cleavage in buffer, in the presence of porcine butyrylcholine esterase as well as in human and murine plasma [34]. The moderately basic morpholinoethyl ester (**4**) proved to be of particular interest with respect to both solubility and stability [34]. Preliminary data suggested considerably increased cytotoxicity of the esters compared to the parent compound **1**, the basic compounds being of particular interest [31]. Based on the assumption that higher antiproliferative activity may result from increased cellular accumulation, additional mechanisms of action or both, we compared compounds 2-7 with bendamustine (**1**) regarding cytotoxicity against a panel of human sarcoma and carcinoma cell types, representing hematologic and solid malignancies. Additionally, the induction of p53 expression and apoptosis, cellular enrichment and the involvement of the organic cation transporters OCT1 and OCT3 were investigated.

5.2 Material and Methods

5.2.1 Ethics Statement

Human embryonal kidney (HEK293) cells were purchased from the German Collection of Microorganism and Cell Cultures (DSMZ, Braunschweig, Germany). Cancer cell lines were obtained from the American Type Culture Collection (ATCC; Rockville, MD, USA): SK-MEL-3 (malignant melanoma; HTB-69), Capan-1 (pancreatic adenocarcinoma; HTB-79), HT-29 (colorectal adenocarcinoma; HTB-38), SK-ES-1 (Ewing's sarcoma; HTB-86), HEL92.1.7 (erythroleukemia; TIB-180), Jurkat (acute T-cell leukemia; TIB-152), U-937 (histiocytic lymphoma; CRL-1593.2), LNCaP (prostate cancer; CRL-1740) and NCI-H460 (large cell lung cancer; HTB-177), and MG-63 (osteosarcoma; CRL-1427) cells.

5.2.2 Chemicals and reagents

Bendamustine hydrochloride (**1**) and the esters **2-7** were kindly provided by Arevipharma (Radebeul, Germany). Stock and working solutions of **1-7** were prepared in dimethylsulfoxide (DMSO). Thiazolyl blue tetrazolium bromide (MTT), umbelliferone, 4-[4-(dimethylamino)styryl]-N-methylpyridinium iodide (ASP⁺), penicillin-streptomycin and tetrapentyl ammonium (TPA) were from Sigma (Taufkirchen, Germany). Hygromycin B was from A.G. Scientific, (San Diego, CA, USA). Fetal calf serum (FCS) was from Biochrom (Berlin, Germany). A 1 mM stock solution of TPA was prepared in phosphate buffered saline (PBS). The 100 μ M stock solution of ASP⁺ was prepared in PBS containing 10 % DMSO. HPLC-grade acetonitrile was from Fisher Scientific (Schwerte, Germany). Demineralized water was prepared with a Milli-Q system (Merck; Schwalbach, Germany). If not otherwise stated, all other reagents were of analytical grade and purchased from Sigma (Taufkirchen, Germany).

5.2.3 Cell culture

Cancer cells: All media were from Sigma (Taufkirchen, Germany) and supplemented with 10 % FCS. SK-MEL-3, Capan-1, HT-29 and SK-ES-1 cells were cultured in McCoy's 5A medium, supplemented with 2.2 g/L NaHCO₃ (Merck, Darmstadt, Germany). HEL92.1.7, Jurkat, U-937, LNCaP and NCI-H460 were cultured in RPMI-1640 containing 110 mg/L sodium pyruvate, 10 mM HEPES (both from Serva, Heidelberg, Germany) and 2.0 g/L NaHCO₃. MG-63 cells were cultured in EMEM supplemented with 2.2 g/L NaHCO₃. Genetically modified HEK293 cells: Transfected human embryonic kidney cells (HEK293) were cultured in EMEM, supplemented with penicillin/streptomycin (100 U/mL / 100 μ g/mL) and 250 μ g/mL hygromycin B. The cells were cultured in a water-saturated atmosphere with 5 % CO₂ at 37 °C and passaged once to twice a

week. Adherently growing cells were treated with trypsin/EDTA (0.5 mg/mL / 0.22 mg/mL) (PAA, Pasching, Austria) and washed with medium prior to transfer into new culture flasks. Cells growing in suspension were passaged after mechanical separation of cell agglomerates.

5.2.3.1 HEK293 cells expressing the human organic cation transporters OCT1 or OCT3

Parental HEK293 cells were transfected with the plasmid pcDNA3.1/Hygro(-)-OCT1, containing the full length *SLC22A1* cDNA (NM_003057) encoding the human OCT1 protein, as described [35]. After hygromycin selection, single colonies were characterized for *SLC22A1* mRNA expression by real-time qRT-PCR, and the cell clone with the highest expression was further analyzed with respect to protein expression and [³H]-1-methyl-4-phenylpyridinium ([³H]MPP⁺) uptake (cf. Supporting Information). HEK-Co cells (control) were established by the same method using the empty plasmid pcDNA3.1/Hy(-) for transfection [36] (characterization of HEK-OCT1 and HEK-Co cells cf. Supporting Information Fig. S5.1). HEK-OCT3 cells were recently described [37]. HEK-Co, HEK-OCT1 and HEK-OCT3 cells were provided by Prof. Dr. Jörg König.

5.2.4 Chemosensitivity assays

Depending on the characteristics of the cells, two different assays were performed. For adherently growing cells (HT-29, NCI-H460, SK-MEL-3, MG-63, Capan-1), cytotoxic and cytocidal effects were determined in the crystal violet assay [38], whereas for loosely adherent cells (LNCaP, SK-ES-1) and cells growing in suspension (HEL 92.1.7, Jurkat, U-937) the MTT-assay [39] was used with minor modifications [38]. In brief, $1.5 \cdot 10^3$ (HT-29, NCI-H460, SK-MEL-3, MG-63, Capan-1) or $5 \cdot 10^3$ (Jurkat, U-937, HEL92.1.7, LNCaP, SK-ES-1) cells per well were seeded into flat-bottom 96-well plates (Sarstedt, Nümbrecht, Germany). All compounds were used as solutions prepared in DMSO (final concentration of DMSO in the assays: 0.1 %) and tested at final concentrations (8 replicates per concentration) of 1, 3, 10, 30 and 50 μ M. Absorbance was measured at 580 nm in a 96-well plate reader (GENios pro microplate reader, Tecan, Salzburg, Austria). Effects were quantified as previously described [40] for both assays.

In addition to long-term exposure, IC₅₀ values were determined after an incubation period of 96 hours, following the same procedure as described above (4 instead of 8 replicates). As suggested by the National Cancer Institute [41] the corrected T/C values (T/C_{corr}) were plotted against the logarithm of the concentrations, and the IC₅₀ values were calculated using Prism 5.01 (GraphPad Software, La Jolla, CA, USA) according to the “*log (inhibitor) vs. normalized response-variable slope*” equation.

5.2.5 Detection of apoptosis (annexin V/propidium iodide assay)

Apoptosis was determined by incubating progressively growing Jurkat cells with medium containing **1**, **2**, **4** or **5** at a concentration of 10 μ M or 0.1 % of DMSO (untreated control). After different periods of incubation (6, 24, 48 hours), samples were analyzed using the Annexin V-FITC apoptosis detection kit I (BD Biosciences, Heidelberg, Germany) according to the manufacturer's protocol using 10⁶ cells/mL. Cells were analyzed using a FACSCalibur flow cytometer (BD Biosciences, Heidelberg, Germany). The compensation was performed for each experiment with annexin V-FITC (530/30 BP filter) and propidium iodide (585/42 BP filter), respectively. At least 1 · 10⁴ events were registered per sample and debris as well as cell aggregates were excluded by forward (FSC) versus side scatter (SSC) gates. Raw data were analyzed using FlowJo V10 software (Treestar Inc., Ashland, OR, USA).

5.2.6 Detection of p53 expression by immunoblotting

The expression of the tumor suppressor p53 by NCI-H460 and HT-29 cells was determined after incubating the cells with compounds **1**, **2**, **4** and **5** at different concentrations for 24 hours. The cells (from a 10-cm culture dish, 70 % confluency) were washed twice with PBS and harvested by scraping after addition of ice-cold buffer A (10 mM HEPES pH = 7.9, 10 mM KCl, 0.1 mM EDTA, 0.1 mM EGTA, protease inhibitor mix (Sigma-Aldrich)). After adding Nonidet P-40 (NP-40) (Sigma, Taufkirchen, Germany) to a final concentration of 1 %, the cell suspensions were vortexed and subsequently centrifuged at 13000 g and 4 °C for 30 seconds. The pellets were re-suspended in buffer B (buffer A + 400 mM NaCl, 1% NP-40) and gently agitated using a Sarmix M2000 (Sarstedt, Nümbrecht, Germany) at 4 °C for 15 min, followed by centrifugation (13000 g, 4 °C, 5 min; Microfuge; Eppendorf, Hamburg, Germany). The concentration of soluble protein was determined according to Bradford using the Bio-Rad Protein Assay (Bio-Rad Laboratories, Munich, Germany). An amount of 30 μ g total protein of each sample and 8 μ L of a biotinylated molecular weight standard (1:6 dilution) (Cell Signaling, Danvers, MA, USA) was separated by SDS-PAGE (12 % gel) and afterwards electroblotted (150 V, 30 min) to nitrocellulose membrane (PeqLab, Erlangen, Germany). The blot was incubated with anti-p53 rabbit mAb (dilution 1:1000) (Cell Signaling) and anti-histone H2B Ab (dilution 1:1000) (Cell Signaling) as loading control. The primary antibodies and the "protein ladder" were simultaneously detected with a horseradish peroxidase (HRP)-coupled donkey anti rabbit mAb (dilution 1:1000) (Santa Cruz biotechnology, Heidelberg, Germany) or the anti-biotin HRP-coupled antibody (dilution 1:5000) (Cell Signaling), respectively. The bands were detected by bioluminescence using the Pierce ECL Western Blotting Substrate (Thermo Scientific, Dreieich, Germany). After exposure, the x-ray film (Amersham Hyperfilm™ ECL, GE Healthcare, München, Germany) was developed (CAWOMAT 2000 IR, CAWO,

Schrobenhausen, Germany) and analyzed using a GS-710 imaging densitometer and Quantity One V. 4 software (Bio-Rad Laboratories, Munich, Germany).

5.2.7 Quantification of cell-associated bendamustine and derivatives

Accumulation of bendamustine and selected bendamustine esters by HT-29 and NCI-H460 cells was determined by HPLC. For this purpose, $1.5 \cdot 10^6$ cells/well were seeded into 6-well plates (Sarstedt, Nümbrecht, Germany). After two days of cultivation, the cells were incubated in PBS containing 30 μ M of **1**, **2**, **4** or **5** and 200 μ M of umbelliferone as internal standard at 37 °C for 10 minutes. After removing the medium, the cells were washed three times with PBS. Untreated cells were detached with trypsin/EDTA and counted. The treated cells were harvested by adding 200 μ L of ice-cold perchloric acid (1 M) and scraping. Subsequently, the samples were vortexed and sonicated (Branson 3200 ultrasonic cleaner; Branson, Danbury, USA) for 10 minutes. After 5 minutes of centrifugation (13000 g, 4 °C), the supernatants were filtered (0.2 μ m Phenex; Phenomenex, Aschaffenburg, Germany) and directly analyzed by HPLC with fluorescence detection according to a recently reported validated procedure [34]. The normalization of the measured concentration to the cell count allowed for a calculation of the cell-associated amount of the respective test compound. Additionally, the ratio of cell-associated substance compared to the applied concentration (30 μ M) was calculated based on an average cell volume of 3 pL.

5.2.8 Flow cytometric determination of OCT1 and OCT3 activity

The previously described fluorescent substrate of organic cation transporters, 4-(4-dimethylaminostyryl)-N-methylpyridinium (ASP⁺) [42-45], was used to determine the function of OCT1 and OCT3 by flow cytometry in the absence and the presence of bendamustine and selected bendamustine esters. HEK293 cells expressing the transporter of interest were trypsinized and washed twice with PBS, prior to re-suspension of $0.5 \cdot 10^6$ cells in 500 μ L of PBS containing 2 % (v/v) FCS. The cells were incubated with ASP⁺ (100 μ M stock solution in PBS containing 10 % DMSO) for 5 minutes, allowing for cellular accumulation, and subsequently analyzed with a FACSCalibur flow cytometer. ASP⁺ was excited at 488 nm and the fluorescence was measured using a 530/30 nm and a 585/42 band-pass filter. At least $1 \cdot 10^4$ single cells were analyzed by appropriate FSC/SSC gating.

For the determination of K_m , cell suspensions of HEK-OCT1 and HEK-OCT3 cells as well as HEK-Co cells as control for unspecific uptake of ASP⁺ were incubated with ASP⁺ at increasing concentrations at 37 °C for 5 minutes. Subsequently, two volumes of an ice-cold 200 μ M solution of the standard inhibitor tetrapentyl ammonium (TPA) [46-48] in PBS/FCS were added to one volume of cell suspension. The mixture was immediately analyzed. The mean fluorescence

intensities (MFI) were calculated using FlowJo V10, and the difference between total (HEK-OCT1 or HEK-OCT3 cells) and unspecific (HEK-Co cells) uptake was plotted as specific uptake against the concentration of ASP⁺. The linearity of ASP⁺ uptake (1 μM) at 37 °C was assessed between 30 seconds and 7 (HEK-OCT1) respectively 9 (HEK-OCT3) minutes. K_m was calculated (GraphPad Prism 5.01) using the Michaelis-Menten equation.

The inhibition of ASP⁺ uptake was measured to determine the IC₅₀ values of TPA and compounds **1**, **2**, **4** and **5**. The cells were pre-incubated with the respective test compound at different concentrations at room temperature for 10 minutes. Afterwards, ASP⁺ was added to a final concentration of 1 μM, and the samples were incubated at 37 °C in the dark for 5 minutes. Subsequently, the cells were washed, re-suspended in ice-cold PBS/FCS and stored on ice in the dark until measurement. The normalized mean fluorescence intensities of the OCT expressing cells, set to 100 % in the absence of an inhibitor, were calculated by subtracting the mean fluorescence intensity of unspecific ASP⁺ uptake into HEK-Co control cells. The normalized fluorescence intensities in the presence of an inhibitor were plotted against the logarithm of the concentrations of the test compounds to calculate IC₅₀ values. The “*log (inhibitor) vs. normalized response-variable slope*” equation of GraphPad Prism 5.01 was used for this purpose.

5.2.9 Imaging of cellular ASP⁺ uptake by confocal laser scanning microscopy

HEK-Co, HEK-OCT1 and HEK-OCT3 cells were seeded into 8-well μ-Slides (Ibidi, Munich, Germany) (2 · 10⁴ cells/well) and allowed to attach for 48 hours. Prior to the staining procedure, the medium was replaced by PBS and the DNA probe Draq5 (5 μM) (Biostatus, Shephed, UK) was added to each well. To monitor the inhibition of ASP⁺ uptake, 200 μM of TPA or 15 μM of compound **5**, respectively, were added and incubated at 37 °C for 5 minutes. Subsequently, ASP⁺ was added to reach a concentration of 1 μM, and the cells were incubated for 5 minutes in the dark. Imaging was performed with an Axiovert 200 M confocal microscope coupled to a Zeiss LSM 510 scanning device (Carl Zeiss, Oberkochen, Germany) using a Plan-Apochromat 63x/1.40 oil immersion objective. Draq5 was excited at 633 nm and fluorescence was detected using a 650 nm long-pass filter, ASP⁺ was excited at 488 nm and detected using a 565/35 nm band-pass filter.

5.2.10 Determination of OCT1 and OCT3 expression by various cancer cells

The expression of *SLC22A1* and *SLC22A3* mRNA in different cancer cell types was analyzed by quantitative RT-PCR using the Roche LightCycler system. Total RNA was isolated using the NucleoSpin RNA Purification Kit (Macherey-Nagel, Dueren, Germany) according to the manufacturer's instructions. One μg of total RNA was reversely transcribed using the iScript Kit (Biorad, Munich, Germany) according to the manufacturer's instructions. *SLC22A1*, *SLC22A3* and

β-actin mRNA levels were determined using the LightCycler System and the FastStart DNA Master SYBR Green I Kit (both from Roche, Mannheim, Germany). The primer pair for the amplification of the *SLC22A1* cDNA fragment was oOCT1-RT.for (CTGCCTGGTGAATGCTGAGC) and oOCT1-RT.rev (ACATCTCTCTCAGGTGCCCG), for the *SLC22A3* cDNA fragment oOCT3-RT.for (CAAGCAATATAGTGGCAGGGG) and oOCT3-RT.rev (CCTCAAAGGTGAGAGCGGGA) and for the *β-actin* fragment oActin.for (TGACGGGGTCACCACACACTGTGTGCCCATCTA) and oActin.rev (CTAGAAGCATTGCGGTGGACGATGGAGGG). PCR was performed according to the manufacturer's instructions with 0.5 μM of the respective sense and antisense primers, 4 mM MgCl₂ and 1-fold FastStart DNA Master SYBR green I mix in a total volume of 20 μL including 1 μL of the synthesized sscDNA. Cycling conditions were as follows: 10 min denaturation at 94 °C, followed by 45 cycles of 10 s denaturation at 94 °C, 15 s primer annealing at 64 °C and 30 s of elongation at 72 °C. The amount of *β-actin*, *SLC22A1* and *SLC22A3* cDNAs were determined using a serial plasmid dilution (pOCT1.31; from 10⁶ to 10⁴ fg) as amplification standard. The *β-actin* concentration, calculated in relation to the standard curve, was set to 100 % and the respective *SLC22A1* and *SLC22A3* mRNA values are given as a percentage of *β-actin* amplification.

5.2.11 Western blot analysis of hOCT1 expression

HEK-Co and HEK-OCT1 cells were lysed with 0.2% ice-cold SDS containing protease inhibitor mixture (Mini-complete protease inhibitor cocktail tablets; Roche Diagnostics-Applied Science, Mannheim, Germany). Protein concentration of each lysate was measured with a bicinchoninic acid assay (BCA protein assay kit, Thermo Fisher Scientific, Waltham, MA, USA). 20 μg of cell lysates were subjected to a 10% SDS-polyacrylamide gel electrophoresis under reducing conditions. Separated proteins were transferred onto nitrocellulose membranes (Protran nitrocellulose transfer membrane; Whatman, Dassel, Germany). Blots were probed with the KEN antiserum (1:10000) directed against human OCT1 [49] overnight at 4 °C. Membranes were then incubated with a horseradish peroxidase-conjugated goat-anti-rabbit secondary antibody (1:10 000; Sigma Aldrich, Munich, Germany). Immunoreactive bands were visualized using an ECL western blotting detection reagent (GE Healthcare, Buckinghamshire, UK) and a Chemidoc XRS imaging system (Bio-Rad, Munich, Germany). To control sample loading, membranes were incubated for 30 min with restore western blot stripping buffer (Pierce, Rockford, USA) at 37 °C and reprobed with a mouse monoclonal anti-human *β-actin* antibody (1:10000; Sigma Aldrich, Munich, Germany). A horseradish peroxidase-conjugated goat-anti-mouse antibody (GE Healthcare Europe, Freiburg, Germany) was used as secondary antibody (dilution 1:4000).

5.2.12 Immunofluorescence detection of hOCT1 expression

The cellular localization of the OCT1 protein in the stably transfected HEK-OCT1 cells was analyzed by immunofluorescent staining. The OCT1-specific KEN antiserum was used at a dilution of 1:1000. Subsequently, cells were incubated with an Alexa Fluor 488 conjugated goat-anti-rabbit secondary antibody (1:10,000; Invitrogen, Carlsbad, CA).

5.2.13 [³H]MPP⁺ transport assay

HEK-OCT1 and HEK-Co cells were seeded in poly-D-lysine (Sigma Aldrich, Taufkirchen, Germany) coated 48-well plates at a density of $1.2 \cdot 10^5$ cells/well. After incubation at 37 °C and 5% CO₂ for 48 h medium was replaced by warm (37 °C) uptake buffer (142 mM NaCl, 5 mM KCl, 1 mM K₂HPO₄, 1.2 mM MgSO₄, 1.5 mM CaCl₂, 5 mM glucose and 12.5 mM HEPES, pH 7.3). The uptake assay was started by addition of [³H]MPP⁺ (80 Ci/mmol, American Radiolabeled Chemicals, St Louis, MO) at indicated concentrations (concentration dependency) or at a concentration of 50 μM (time dependency). Cells were incubated at 37 °C for 3 min (concentration dependency) or at indicated time points (time dependency). Uptake was stopped by washing the cells three times with ice-cold uptake buffer. Afterwards, cells were lysed with 5 mM Tris buffer (pH 7.3) containing 0.1% Triton X-100. The intracellular accumulation of radioactivity was determined by liquid scintillation counting (PerkinElmer, Rodgau-Jügesheim, Germany) and protein concentration of each lysate was measured with the BCA assay. We performed two experiments each on two different days ($n = 4$). The OCT1-mediated net uptake of [³H]MPP⁺ was determined as the difference in substrate uptake between HEK-OCT1 and HEK-Co cells. Data were presented as means ± standard error of the mean. A value of $p < 0.05$ was considered statistically significant.

5.3 Results and discussion

5.3.1 Cytotoxicity of bendamustine and derivatives

The cytotoxicity of compounds **1-7** against tumor cells was determined both as an end point and kinetically. IC₅₀ values of compounds **1-7** (Table 5.1 and Fig. S5.2) were calculated after 96 hours of incubation and the cytotoxic drug effect was measured over a period of 5 days (Fig. 5.2 and Supplementary Material, Fig. S5.3 – S5.12). The kinetic approach allows the distinction between cytotoxic, cytostatic and cytocidal drug effects [40].

Table 5.1: Cytotoxicity of compounds **1-7** against different human tumor cell lines, expressed as IC₅₀ values (μM) after 96 hours of incubation.

Cell line	Compound						
	1	2	3	4	5	6	7
HEL 92.1.7 ^b	86.2 ± 2.6	2.52 ± 0.09	2.85 ± 0.02	2.35 ± 1.02	1.04 ± 0.01	1.29 ± 0.01	1.72 ± 0.04
Jurkat ^b	43.4 ± 4.8	3.14 ± 0.11	3.84 ± 0.04	3.19 ± 1.01	0.61 ± 0.02	0.83 ± 0.01	0.90 ± 0.03
U-937 ^b	83.4 ± 8.5	1.83 ± 0.01	1.94 ± 0.01	3.2 ± 0.96	0.83 ± 0.19	1.08 ± 0.20	1.28 ± 0.28
MG-63 ^a	55.8 ± 1.90	1.85 ± 0.05	2.14 ± 0.06	3.45 ± 0.06	0.38 ± 0.06	0.53 ± 0.01	0.55 ± 0.01
SK-ES1 ^b	9.6 ± 0.3	0.18 ± 0.02	0.18 ± 0.01	0.91 ± 0.08	0.09 ± 0.01	0.19 ± 0.01	0.19 ± 0.03
Capan-1 ^a	18.9 ± 4.5	0.60 ± 0.04	1.00 ± 0.01	1.38 ± 0.01	0.12 ± 0.02	0.17 ± 0.01	0.25 ± 0.02
LNCaP ^b	77.9 ± 5.6	0.95 ± 0.15	1.52 ± 0.05	2.2 ± 0.1	0.50 ± 0.08	0.75 ± 0.03	0.93 ± 0.12
NCI-H460 ^a	90.0 ± 4.3	4.82 ± 0.03	4.80 ± 0.01	9.65 ± 1.27	0.90 ± 0.01	1.17 ± 0.02	1.24 ± 0.06
HT-29 ^a	> 100 ^c	9.35 ± 0.04	10.0 ± 0.10	8.99 ± 0.47	1.17 ± 0.06	1.71 ± 0.01	2.35 ± 0.01
SK-Mel3 ^a	> 100 ^c	13.8 ± 0.4	21.7 ± 4.10	20.9 ± 0.9	0.83 ± 0.18	1.24 ± 0.07	1.0 ± 0.27

Data are mean values ± SEM of 3 independent assays with 4 replicates per compound concentration.

^a) Crystal violet assay

^b) MTT-assay

^c) Mean value ± SEM of 2 independent assays

Although **1** is approved for hematologic malignancies (sarcoma), the effects on representative tumor entities were rather weak (Table 5.1, Fig. 5.2 and Supporting Information). The IC₅₀ values were > 80 μM against HEL 92.1.7 (erythroleukemia) and U-937 (histiocytic lymphoma), and 43 μM against Jurkat cells (acute T-cell leukemia, Fig. 5.2A). By contrast, the Ewing's sarcoma cells SK-ES-1 showed a distinct response upon treatment with bendamustine (IC₅₀ ~ 10 μM; Table 5.1, Fig. 5.2B), whereas the chemosensitivity of MG-63 osteosarcoma cells was markedly lower (IC₅₀ ~ 56 μM) with cytotoxic drug effects at concentrations above 30 μM (Fig. S5.6). Surprisingly, with an IC₅₀ value < 20 μM Capan-1 pancreatic cancer cells showed moderate chemosensitivity. On the contrary, **1** was ineffective (SK-MEL-3 melanoma, HT-29 colorectal carcinoma) or only very weakly active (LNCaP prostate cancer; NCI-H460 large cell lung cancer) against the other carcinoma cell types (IC₅₀ values 78 to > 100 μM).

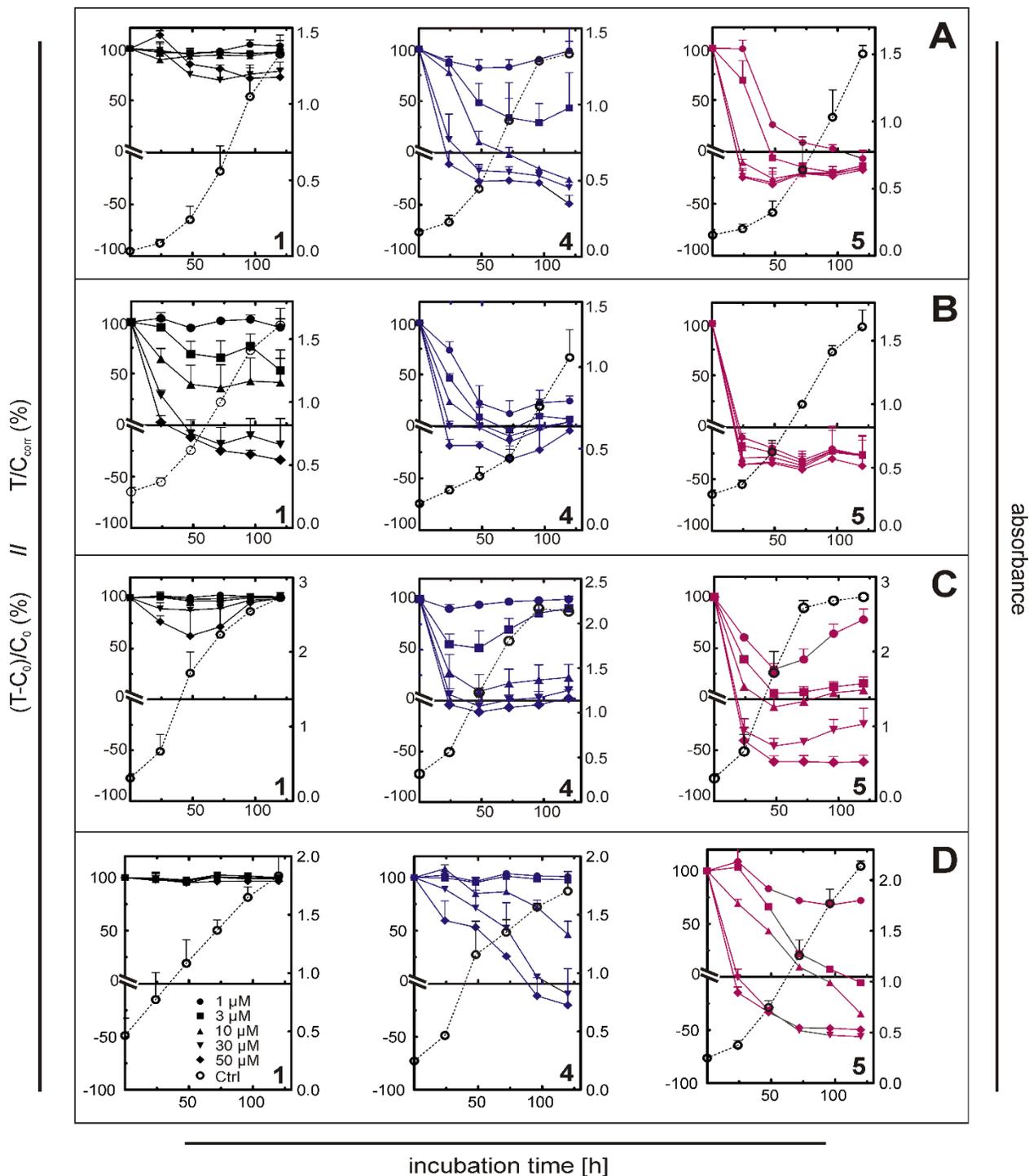


Figure 5.2: Antiproliferative activity of compounds 1, 4 and 5 against selected sarcoma and carcinoma cells upon long-term incubation. Jurkat (A; acute T cell leukemia), SK-ES-1 (B; Ewing's sarcoma), NCI-H460 (C; large cell lung cancer) and HT-29 (D; colorectal cancer) cells were treated with compounds 1, 4 and 5 at concentrations between 1 μM and 50 μM over an incubation period of 5 days. Antiproliferative and cytotoxic effects correspond to the left y-axes. The growth curves of untreated control cells (open circles) correspond to the right y-axes. Data are mean values \pm SEM of at least 2 independent assays with 8 replicates per compound concentration.

In contrast to the parent compound bendamustine, the derivatives **2-7** exhibited considerably higher potencies up to factors > 100 both, against the sarcoma and the carcinoma cells investigated. With respect to the cytotoxic effects, compounds **2-7** fall into two groups: the alkyl esters (**2, 3**) and the mofetil ester **4** on one hand, and the basic heterocyclic esters **5-7** on the other hand. Compared to bendamustine, compounds **2-4** showed a 10- to 30-fold increase in potency, and **5-7** were 60- to 120-fold more potent than **1**.

Compounds 2-4: Among compounds **2-4**, minor differences were observed at SK-ES-1 and NCI-H460 cells (Table 5.1) with a more pronounced effect of the alkyl esters **2** and **3** against SK-ES-1 cells (IC₅₀ values: **2** and **3**: ~ 0.2 μM, **4**: 0.9 μM). The increase in toxicity compared to **1** was most striking at HEL92.1.7, LNCaP, MG-63 and, in case of the alkyl esters **2** and **3**, at SK-ES-1 cells. Comparable IC₅₀ values of **2-4** were confirmed by the cytotoxic drug effects over a period of 5 days (Fig. S5.3-S5.12). Compounds **2-4** revealed cytotoxic effects at all investigated cancer cell types at concentrations of 30 and 50 μM, except for SK-MEL-3, HT-29 and NCI-H460.

Compounds 5-7: Except for the hematologic malignancies (HEL92.1.7, Jurkat, U-937), SK-ES-1 and LNCaP cells, compounds **5-7** exhibited a distinct increase in cytotoxicity compared to **2-4**, showing IC₅₀ values in low micro- to nanomolar range and cytotoxic effects at concentrations as low as 1-10 μM at all treated cancer cells. The antiproliferative activity of compounds **5-7** was comparable both, in the end-point (cf. Table 5.1) and in the kinetic assays over a period of 5 days (Supporting Information, Fig. S5.3 – S5.12). Remarkably, compared to **1**, compounds **5-7** were up to > 100 times more potent against tumor cells such as SK-MEL3, NCI-H460, HT-29 and MG-63, which were refractory against treatment with bendamustine.

Data on the in vitro cytotoxicity of bendamustine are scarce. In the literature, for myeloma cells IC₅₀ values around 100 μM or even higher are reported [22, 23]. A very recent study on several hematologic malignancies revealed IC₅₀ values between approximately 10 μM and 250 μM. Especially mantle cell lymphoma, Burkitt's lymphoma and T-cell acute lymphoblastic leukemia derived cell lines were relatively sensitive to bendamustine treatment [50]. The IC₅₀ value (approximately 50 μM) reported by Hiraoka et al. [50] for Jurkat cells is in good agreement with our data. Published plasma levels of **1** after intravenous administration (C_{max} = 6 μg/mL (≈ 17 μM) [51]; C_{max} = 11 μg/mL (≈ 31 μM) [52]) suggest that the chemotherapy with bendamustine must be considered ineffective in case of tumor entities showing IC₅₀ values in the two- to three-digit micromolar range in vitro. In this context the up to 100-fold antiproliferative activity of the bendamustine esters, in particular **5-7**, suggest both, higher efficacy in case of malignancies for which the parent compound is approved and a possible extension of the scope of indications.

5.3.2 Induction of apoptosis and p53 expression by compounds 1, 2, 4 and 5

Bendamustine was reported to trigger apoptosis [22]. In search for an explanation of the higher antiproliferative activity of the bendamustine esters compared to the parent compound, we determined early and late stage of apoptosis by flow cytometry (annexin V/propidium iodide staining) after treatment of Jurkat cells with 10 μM of compounds **1**, **2**, **4** or **5** for 6, 24 and 48 hours (Fig. 5.3; Fig. S5.13). As becomes obvious from Fig. 5.3, annexin V⁺/PI⁺ cells, defined as late apoptotic/necrotic cells, amounted to approximately 10 % of the total cell population, most probably resulting from sample preparation. As expected from the results of the MTT assay, at a concentration of 10 μM , bendamustine had no effect compared to the control cells, regardless of the period of incubation. After 6 hours, the fraction of early apoptotic cells was less than 1 % except for compound **5** (~2.5 %; cf. Supporting Information, Fig. S5.13). Compounds **2** and **4** showed induction of apoptosis in around 10 % of the cells after 24 hours of incubation. After 48 hours of incubation, 15 % of the cells were in an early apoptotic state and a considerable fraction (20-25 %) was late-apoptotic/necrotic cells. In agreement with the results from the chemosensitivity assays, compound **5** was more potent, exhibiting a more rapid onset of action and a higher maximal response compared to **1**, **2** and **4** regarding induction of apoptosis. Approximately 50 % of the cells were either early apoptotic (20 %) or late apoptotic/necrotic (30 %) after 24 hours. Two days after treatment, only around 15 % of the cells were viable, whereas the majority of the cells was late apoptotic (30 %) or necrotic (40 %).

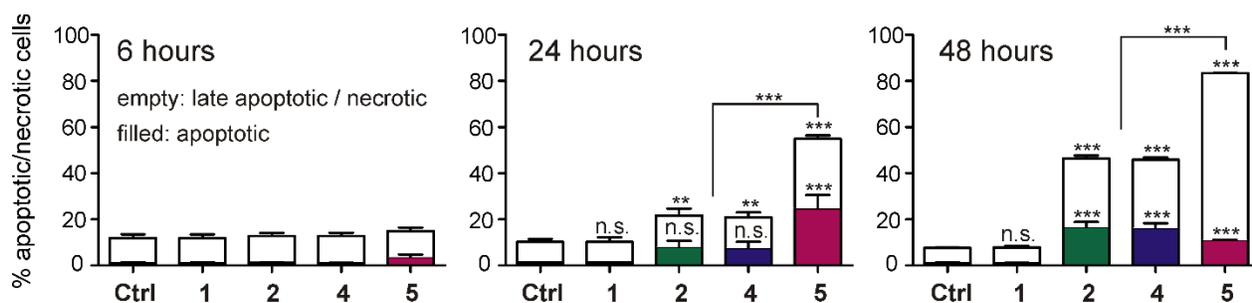


Figure 5.3: Induction of apoptosis in Jurkat cells.

Results of annexin V/propidium iodide staining, performed after incubating Jurkat cells with compounds **1**, **2**, **4** and **5** at a concentration of 10 μM for different periods. Annexin V⁺/PI⁻ cells were defined as early apoptotic, whereas late apoptotic/necrotic cells were annexin V⁺/PI⁺. Significances illustrated on top of the columns refer to the overall effect (apoptotic + necrotic cells), those depicted in the columns refer to apoptotic cells. After 48 hours, apoptotic and necrotic cells were significantly different ($p < 0.001$) (mean \pm SEM, $N = 3$). One-way ANOVA and Bonferroni's post-test were applied to calculate the significance of the apoptotic fraction and the sum of apoptotic and necrotic cells; n.s.: not significant; *: $p < 0.05$; **: $p < 0.01$, ***: $p < 0.001$.

Bendamustine was reported to induce the expression of p53 [21, 23, 50]. As the derivatives of **1** were much more potent against large cell lung and colorectal cancer cells, we investigated the induction of p53 expression in NCI-H460 and HT-29 cells after 24 hours of incubation with compounds **1**, **2**, **4** and **5** (Fig. 5.4).

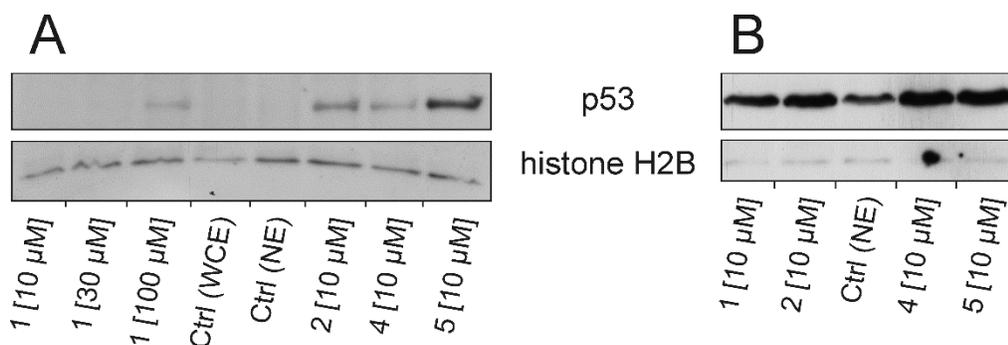


Figure 5.4: Effect of **1**, **2**, **4** and **5** on the expression of p53 in NCI-H460 and HT-29 cells. Western blot of p53 and histone H2B (loading control) of NCI-H460 (A) and HT-29 (B) cells. **A:** Nuclear extracts of NCI-H460 cells, incubated with different concentrations of **1** (10, 30 and 100 μM) or 10 μM of **2**, **4**, or **5** for 24 hours. Nuclear extracts (NE) and whole cell lysates (WCE) of untreated cells were used as control (Ctrl). **B:** Nuclear extracts of HT-29 cells, incubated with 10 μM of **1**, **2**, **4**, **5** for 24 hours. Untreated cells served as control (Ctrl).

Especially in NCI-H460 cells, p53 expression was significantly induced by the treatment with 10 μM of compounds **2**, **4** and **5**, while **5** led to the most pronounced effect. Bendamustine also weakly induced the expression of p53, although at a tenfold higher concentration (100 μM). Interestingly, p53 was only detectable in nuclear extracts of NCI-H460 cells after treatment, whereas HT-29 cells exhibited a high constitutive expression of the tumor suppressor protein. Nevertheless, the treatment with compounds **1**, **2**, **4** and **5** increased the expression of p53 in HT-29 cells but, similar to NCI-H460 cells, compounds **2**, **4** and **5** caused a higher expression level than **1**. Another notable observation was a slight induction of p53 expression in HT-29 cells after treatment with 10 μM of **1**, since proliferation assays revealed no toxic effect at this concentration. It is known from the literature that concentrations of ≥ 80 μM of bendamustine are required to produce a significant induction of p53 expression in various tumor cell types [22]. Both, the induction of p53 and apoptosis, correlated with the cytotoxic potency of bendamustine and derivatives.

5.3.3 Cellular accumulation of bendamustine and derivatives

Higher cellular uptake of the neutral (**2**, **3**) and basic (**4-7**) bendamustine esters compared to the parent compound could account for the increased potency in terms of antiproliferative activity,

induction of apoptosis and p53 expression. Therefore, we performed HPLC analyses to determine the amount of cell-associated test compounds **1**, **2**, **4** and **5** at a concentration of 30 μM after 10 minutes of incubation. NCI-H460 and HT-29 cells were selected as examples of solid tumors, which surprisingly proved to be sensitive against bendamustine esters (Fig. 5.5). The recently reported validated HPLC method was applied [34]. To prevent hydrolysis of the nitrogen mustard group during sample preparation and analysis, cell lysis and deproteination were performed under acidic conditions (1 M perchloric acid, sonication). The esters were proven to be stable over the incubation period of 10 minutes in the presence of cells (Supporting Information; Fig. S5.14). The cellular association of the internal standard umbelliferone did not significantly differ between both cell lines ($p > 0.05$). The amounts of cell-associated bendamustine (**1**) were extremely low (HT-29: 0.07 ± 0.011 nmol/ 10^6 cells; NCI-H460: 0.03 ± 0.001 nmol/ 10^6 cells). The ratio of cell-associated to the applied concentration of the test compound (accumulation factor) was 1:3 for NCI-H460 cells and 2:3 for HT-29 cells, indicating an incomplete uptake. In contrast, compounds **2**, **4** and **5** revealed considerably higher cell-associated amounts, with **5** reaching the highest cellular concentrations. A pronounced cellular enrichment of **2**, **4** and **5** was particularly observed in HT-29 cells. Compounds **2** (2.05 ± 0.39 nmol/ 10^6 cells) and **4** (1.59 ± 0.23 nmol/ 10^6 cells) revealed a 20- to 30-fold, and **5** (5.01 ± 0.34 nmol/ 10^6 cells) an approximately 70-fold higher cellular enrichment than bendamustine. Qualitatively, the cellular accumulation of the bendamustine derivatives was comparable in HT-29 and NCI-H460 cells, though at a lower level (factor of approximately three) in case of the latter. The amounts of cell-associated **2**, **4** and **5** were 0.28 ± 0.06 nmol/ 10^6 cells, 0.59 ± 0.04 nmol/ 10^6 cells and 1.89 ± 0.14 nmol/ 10^6 cells, respectively.

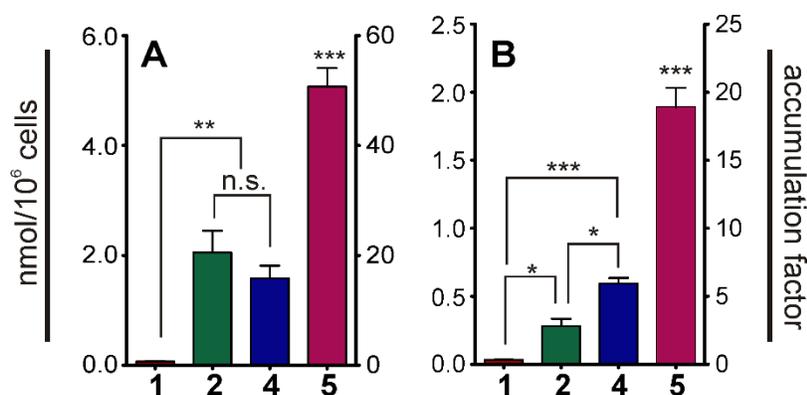


Figure 5.5: Accumulation of **1** and compounds **2**, **4** and **5** by HT-29 and NCI-H460 cells. Amount (mean \pm SEM, $N = 3-4$) of cell-associated **1**, **2**, **4** and **5**, expressed as nmol/ 10^6 cells (left y-axis) and cellular enrichment (right y-axis) in HT-29 (A) and NCI-H460 (B) cells. A mean cell volume of 3 pL was used to calculate accumulation factors. Significance was calculated using one-way Anova: n.s.: not significant; *: $p < 0.05$; **: $p < 0.01$, ***: $p < 0.001$.

The levels of cellular enrichment correlate very well with the antiproliferative activities, underlining a crucial role of the ester moiety depending on the chemical nature, covering a neutral group (**2**) or substructures with different degree of basicity (**4** and **5**). In particular, the contribution of the pyrrolidino group in **5**, which is positively charged under assay conditions, becomes obvious from the increased cellular accumulation which is paralleled by the antiproliferative activity, the induction of apoptosis and p53 expression. The high cellular accumulation of **5** and the comparable effects of **5-7** concerning the toxicity suggest an important role of the basic substituent and the positive charge for the cellular association and thus for the toxicity. Apart from that, the significant difference between HT-29 and NCI-H460 cells might result from a different extent of diffusion and transporter-mediated uptake. Therefore, a possible contribution of organic cation transporters (OCT) was taken into account [53]. To test this hypothesis, functional studies on recombinant OCT1 and OCT3, expressed in HEK293 cells, were performed. Additionally, the expression of the respective transporters by the tumor cell types selected for cytotoxicity studies was investigated.

5.3.4 Effect of bendamustine derivatives on the activities of OCT1 and OCT3

5.3.4.1 ASP⁺ uptake by OCT1 and OCT3 expressing HEK293 cells

Confocal laser scanning microscopy revealed specific uptake of the fluorescent substrate ASP⁺ by organic cation transporter expressing HEK293 cells (Fig. 5.6). OCT1 activity was lower than that of OCT3. The activities of both transporters were inhibited by compound **5** and the reference inhibitor TPA (Fig. 5.6).

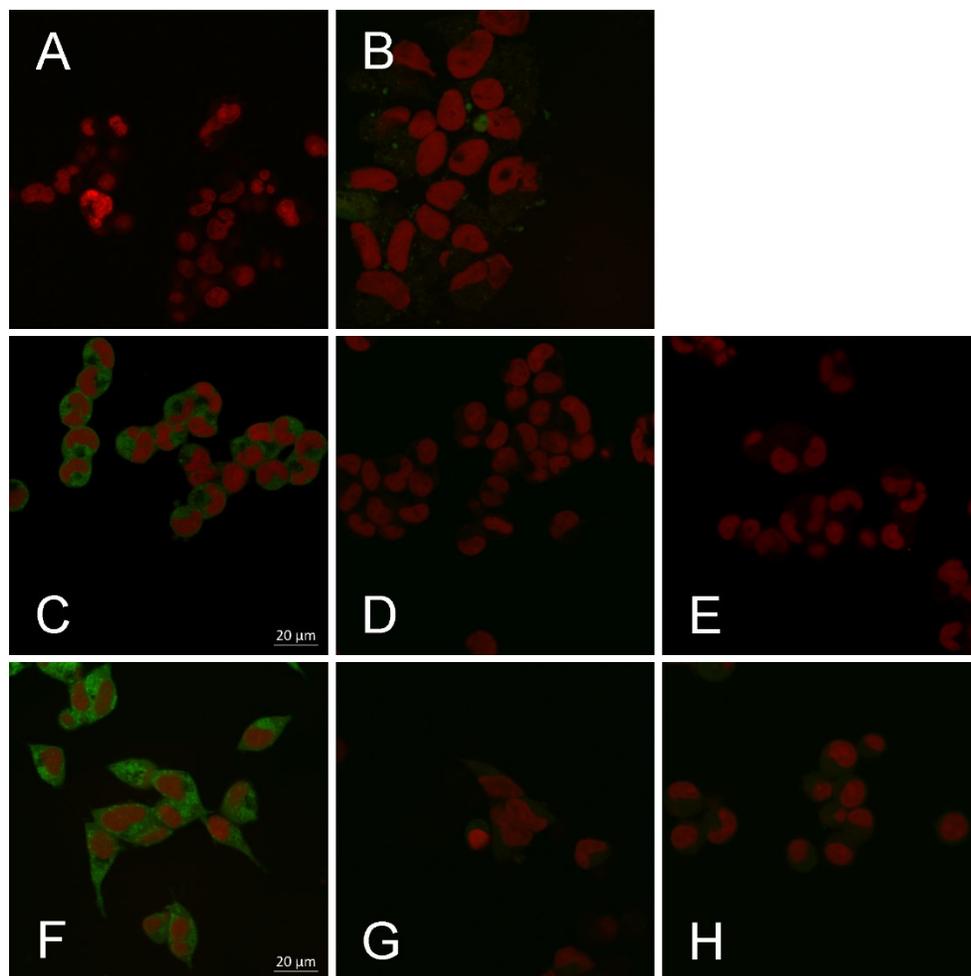


Figure 5.6: Confocal laser scanning microscopy of Cellular ASP⁺ uptake via OCT1 and OCT3.

Uptake of fluorescent ASP⁺ [1 μM] (green) in the absence and presence of compound **5** or TPA after 5 minutes of pre-incubation. Nuclei were stained with Draq5 [5 μM] (red). A, B: HEK-Co (control) cells, treated with Draq5 (A) or ASP⁺ plus Draq5 (B). C-E: HEK-OCT1 cells, treated with ASP⁺ plus Draq5 (C), ASP plus Draq5 plus compound **5** [15 μM] (D), or ASP⁺ plus Draq5 plus TPA [200 μM] (E). F-H: HEK-OCT3 cells, treated with ASP⁺ plus Draq5 (F), ASP⁺ plus Draq5 plus compound **5** [15 μM] (G), or ASP⁺ plus Draq5 plus TPA [200 μM] (H).

5.3.4.2 Determination of the affinities of OCT1 and OCT3 to ASP⁺ as substrate.

Flow cytometry was applied to determine K_m values (Fig. 5.7). The kinetics of ASP⁺-uptake and thus the increase in fluorescence were linear for both transporters (Fig. 5.7A), allowing the determination of initial velocities (v_0) at different concentrations (Fig. 5.7B). The affinity to ASP⁺ was not significantly different for OCT1 ($K_m = 3.1 \pm 0.1 \mu\text{M}$) and OCT3 ($K_m = 2.7 \pm 0.1 \mu\text{M}$). By contrast, the maximal uptake rate for ASP⁺ was significantly higher for OCT3 ($v_{\text{max}} = 5321 \pm 89 \text{ MFI}/(10^4 \text{ cells} \cdot \text{min})$) than for OCT1 ($v_{\text{max}} = 2808 \pm 78 \text{ MFI}/(10^4 \text{ cells} \cdot \text{min})$); $p < 0.001$).

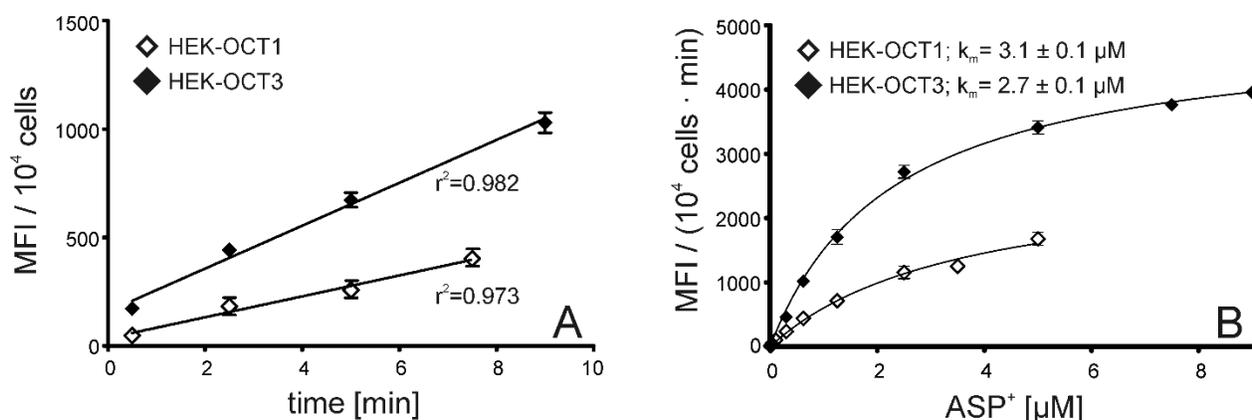


Figure 5.7: Specific uptake of ASP⁺ and Michaelis-Menten kinetics determined by flow cytometry. A: Time dependency of the specific mean fluorescence intensity (MFI) after incubating HEK-OCT1 and HEK-OCT3 cells with 1 μM ASP⁺ (means \pm SEM, $N = 3$).

B: Concentration dependency of the initial velocity of the specific increase in the mean fluorescence intensity (MFI) caused by ASP⁺-uptake into HEK-OCT1 and HEK-OCT3 cells. HEK cells transfected with an empty vector served as control for unspecific uptake (means \pm SEM, $N = 3$).

5.3.4.3 Inhibition of OCT mediated ASP⁺ uptake by bendamustine derivatives.

Bendamustine (**1**), the derivatives **2**, **4** and **5** and the standard inhibitor tetrapentyl ammonium (TPA) were investigated in the flow cytometric ASP⁺-uptake assay to determine IC_{50} values. The IC_{50} values of TPA in HEK-OCT1 ($\text{IC}_{50} = 6.04 \pm 0.83 \mu\text{M}$) and HEK-OCT3 ($\text{IC}_{50} = 19.9 \pm 2.23$) cells were consistent with previously published data, obtained in a microtiter assay (OCT1: $\text{IC}_{50} = 7.9 \mu\text{M}$ [46], OCT3: $\text{IC}_{50} = 28 \mu\text{M}$ [47]). Whereas compounds **2**, **4** and **5** (Fig. 5.8) revealed concentration-dependent inhibition of both transporters, incomplete inhibition of OCT1 (35 %) and OCT3 (37 %) was observed at the highest examined concentration of bendamustine (200 μM). Compounds **2** and **4** were comparable to TPA at OCT1, whereas **5** was more potent by a factor of approximately 20 (Table 5.2). By contrast, compounds **2**, **4** and **5** were almost equipotent at OCT3 and approximately four times more potent than the reference compound TPA ($p \leq 0.05$). Several

Table 5.2: IC₅₀ (μM) values (± SEM, N=3) of the ASP⁺-uptake inhibition into HEK-OCT1 and HEK-OCT3 cells.

Compound	OCT1	OCT3
TPA	6.04 ± 0.83	19.9 ± 2.23
1	> 200	> 200
2	11.32 ± 1.93	5.07 ± 1.41
4	10.05 ± 2.49	5.11 ± 0.02
5	0.35 ± 0.03	3.41 ± 0.16

studies conducted in the last years focused on the uptake of cytostatic drugs by organic cation transporters. Oxaliplatin [54], picoplatin [55], imatinib [56], paclitaxel and irinotecan [57] were identified as OCT1 substrates and oxaliplatin [58], melphalan, irinotecan and vincristine [59] as substrates for OCT3. Compared to other cytostatic drugs such as cisplatin (IC₅₀ value > 100 μM at OCT1 [60]) and mitoxantrone (IC₅₀ 16 μM at OCT1; 440 μM at

OCT3 [61]), especially **5** revealed remarkably higher potency at OCT1 (IC₅₀ 0.35 ± 0.03 μM). Whether compounds **2**, **4** and **5** are inhibitors or substrates of OCT1 and OCT3 is a matter of question. Supposed that these bendamustine derivatives, in particular the basic ester **5**, are substrates, accumulation and cytotoxicity should be related to the OCT expression.

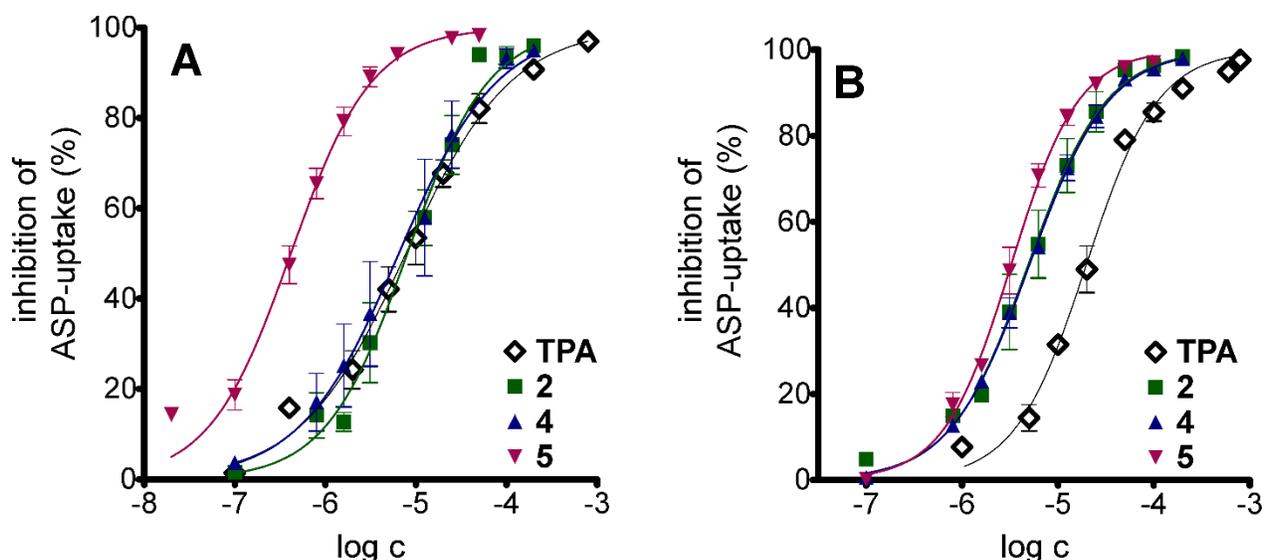


Figure 5.8: Concentration-dependent inhibition of ASP⁺-uptake by OCT expressing HEK-cells. Inhibition of ASP⁺-uptake (1 μM) into HEK-OCT1 (A) and HEK-OCT3 (B) cells by TPA, **2**, **4** or **5**. The mean fluorescence intensities were normalized to uninhibited ASP⁺-uptake (N=3).

5.3.4.4 Expression of OCT1 (*SLC22A1*) and OCT3 (*SLC22A3*) by cancer cells.

Previously, OCT1-mediated uptake of cytostatics in tumor cells was reported for chronic myeloid leukemia [56], chronic lymphocytic leukemia [57], and colon carcinoma [54], whereas OCT3 expression was associated with colorectal [58] and renal cancer [59]. We applied quantitative RT-PCR for the examination of OCT1 and OCT3 expression in all used cancer cell lines (Fig. 5.9).

Whereas OCT1 was weakly expressed in SK-ES1 and SK-MEL3 cells, OCT3 expression was high in HT-29 cells, and clear bands were detectable upon analysis of Capan-1 and LNCaP cells. The differential expression (Fig. 5.9) does not support the hypothesis that the chemosensitivities of the investigated cancer cell types (Table 5.1) are primarily determined by OCT-mediated uptake of the bendamustine esters.

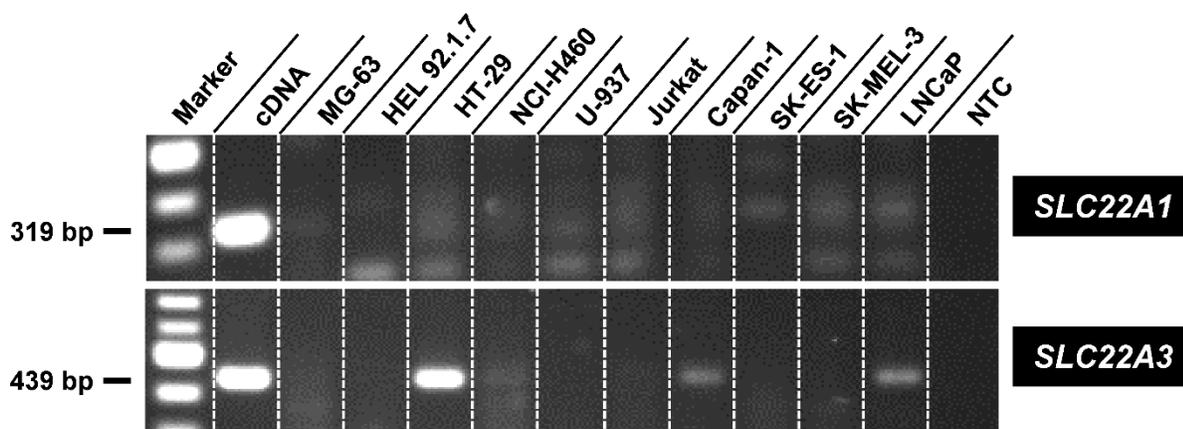


Figure 5.9: Gel electrophoretic analysis of *SLC22A1* (encoding human OCT1) and *SLC22A3* (encoding human OCT3) mRNA expression by different human cancer cell types after quantitative RT-PCR. Human *SLC22A1* mRNA was detected using the primer pair oOCT1-RT.for – oOCT1-RT.rev, resulting in a specific band of 319 bp, human *SLC22A3* mRNA by the primer pair oOCT3-RT.for – oOCT3-RT.rev, resulting in a band of 439 bp. As a positive control, the plasmids pOCT1.31 and pOCT3.31 (cDNA) were used as templates of *SLC22A1* and *SLC22A3*, respectively. NTC = non-template control. Data provided by Prof. Dr. Jörg König.

5.4 Conclusion

Although the investigated compounds are sufficiently stable to act as antitumor agents on their own, it cannot be precluded that the esters are only prodrugs, allowing for increased intracellular accumulation of bendamustine. Having the N-Lost moiety in common, alkylating property is a characteristic feature of both, the parent compound and the derivatives. In concert with the increase in apoptotic processes and elevated p53 expression, the data may be interpreted as a hint to a dual mechanism of action, in particular in case of the basic bendamustine esters.

5.5 Supporting Information

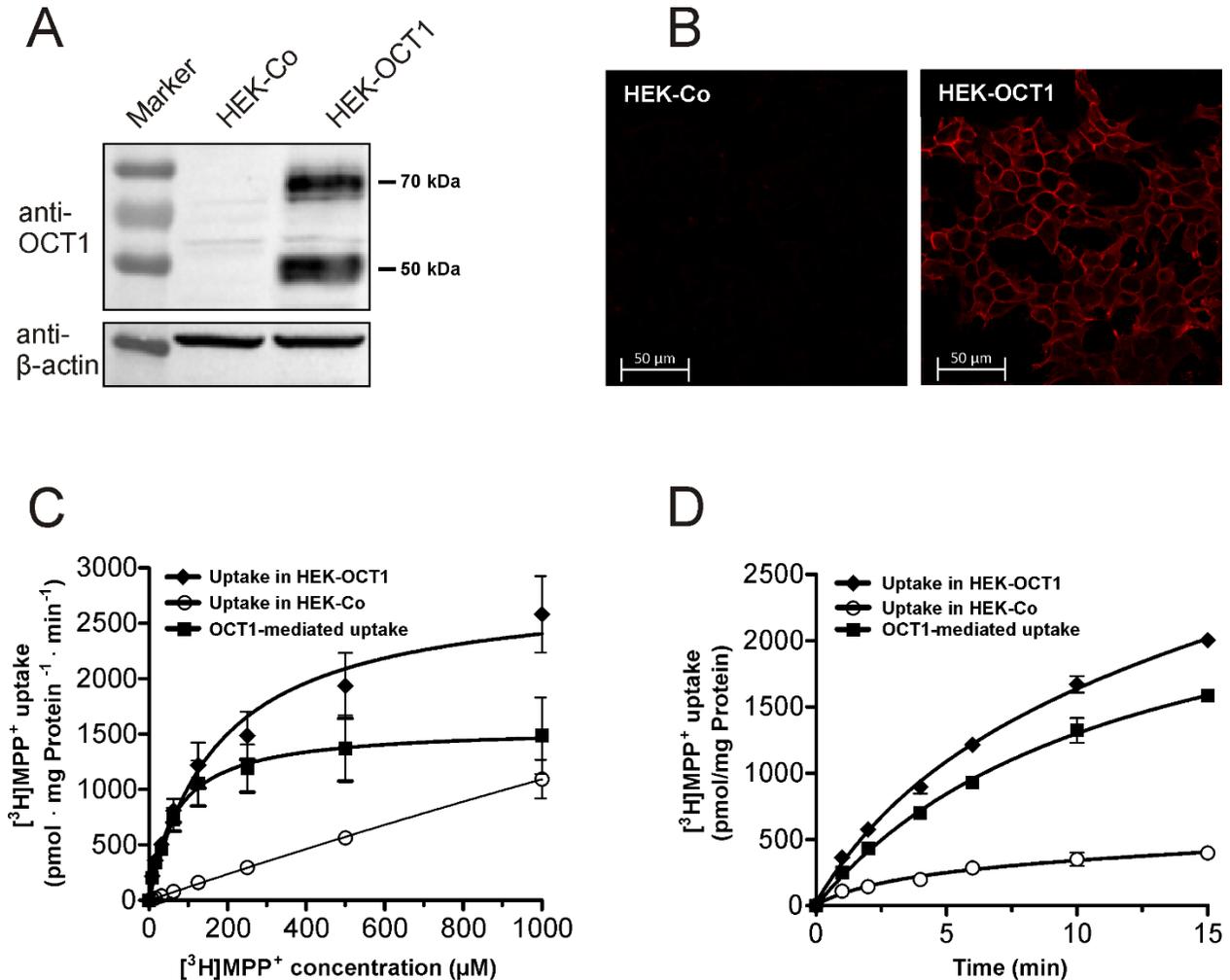


Figure S5.1: Characterization of HEK293 cells stably expressing human OCT1. A: Immunoblot analysis of lysates (20 μg each) of HEK-Co and HEK-OCT1 cells. OCT1 was detected with the antiserum KEN (see: Material and Methods). B: Immunolocalization of human OCT1 in HEK-Co and HEK-OCT1 cells by confocal microscopy. The OCT1 protein was detected in the plasma membrane of transfected HEK-OCT1 cells and no OCT1-specific staining was detectable in HEK-Co cells. C: Concentration dependent $^3\text{H}]\text{MPP}^+$ uptake in HEK-Co and HEK-OCT1 cells (incubation time: 3 min). D: Time-dependent uptake of $^3\text{H}]\text{MPP}^+$ (50 μM) by HEK-Co and HEK-OCT1 cells. In both cases OCT1-mediated uptake (squares) of $^3\text{H}]\text{MPP}^+$ was determined by subtracting uptake in HEK-Co cells from the uptake into HEK-OCT1 cells. Data provided by Prof. Dr. Jörg König.

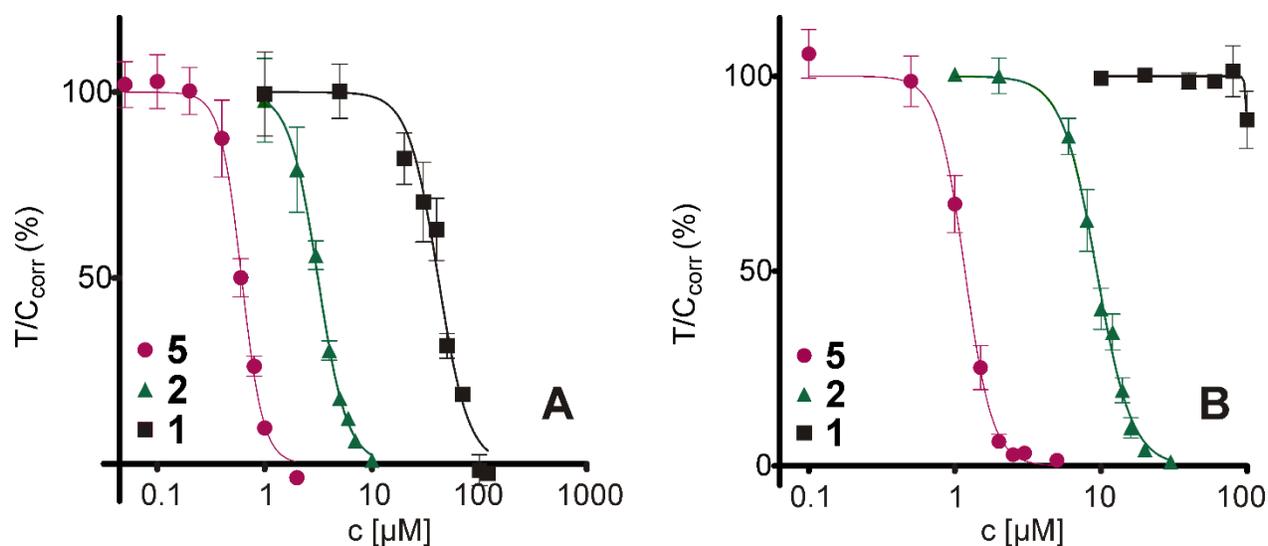


Figure S5.2: Concentration-response curves of 1, 2 and 5 against Jurkat (A) and HT-29 (B) cells after 96 hours of incubation. T/C_{corr} values represent the net proliferation of the treated cells referred to the vehicle treated control cells (set to 100%). The obtained data were used for the calculation of IC_{50} values (mean values \pm SEM of 2-3 independent experiments with 4 replicates per concentration).

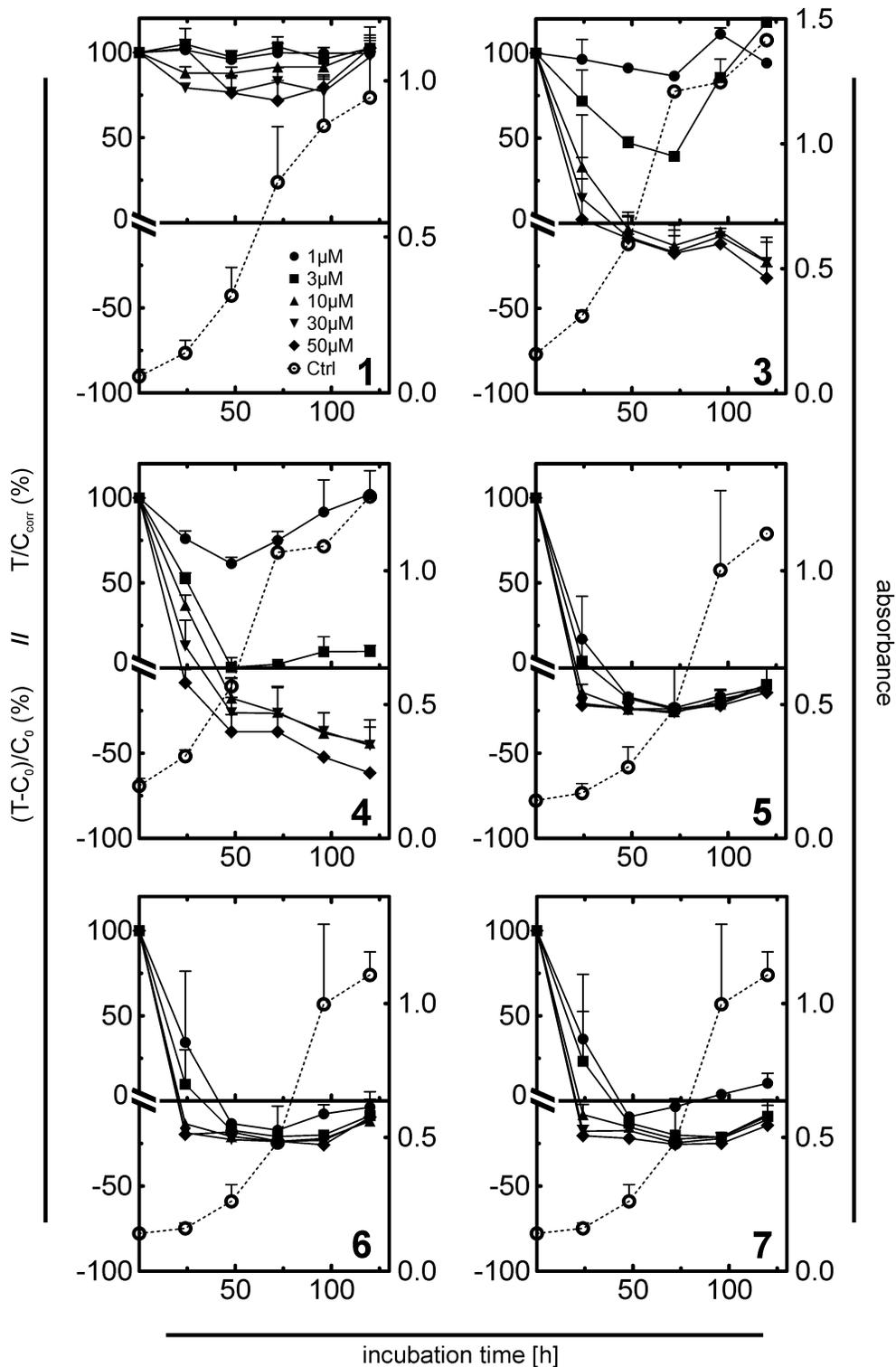


Figure S5.3: Chemosensitivity of HEL92.1.7 cells against compounds 1 and 3-7. Antiproliferative and cytotoxic effects correspond to the left y-axes. The growth curves of untreated cells (open circles) correspond to the right y-axes. Mean values \pm SEM of 2-3 independent assays with 8 replicates per concentration and time point.

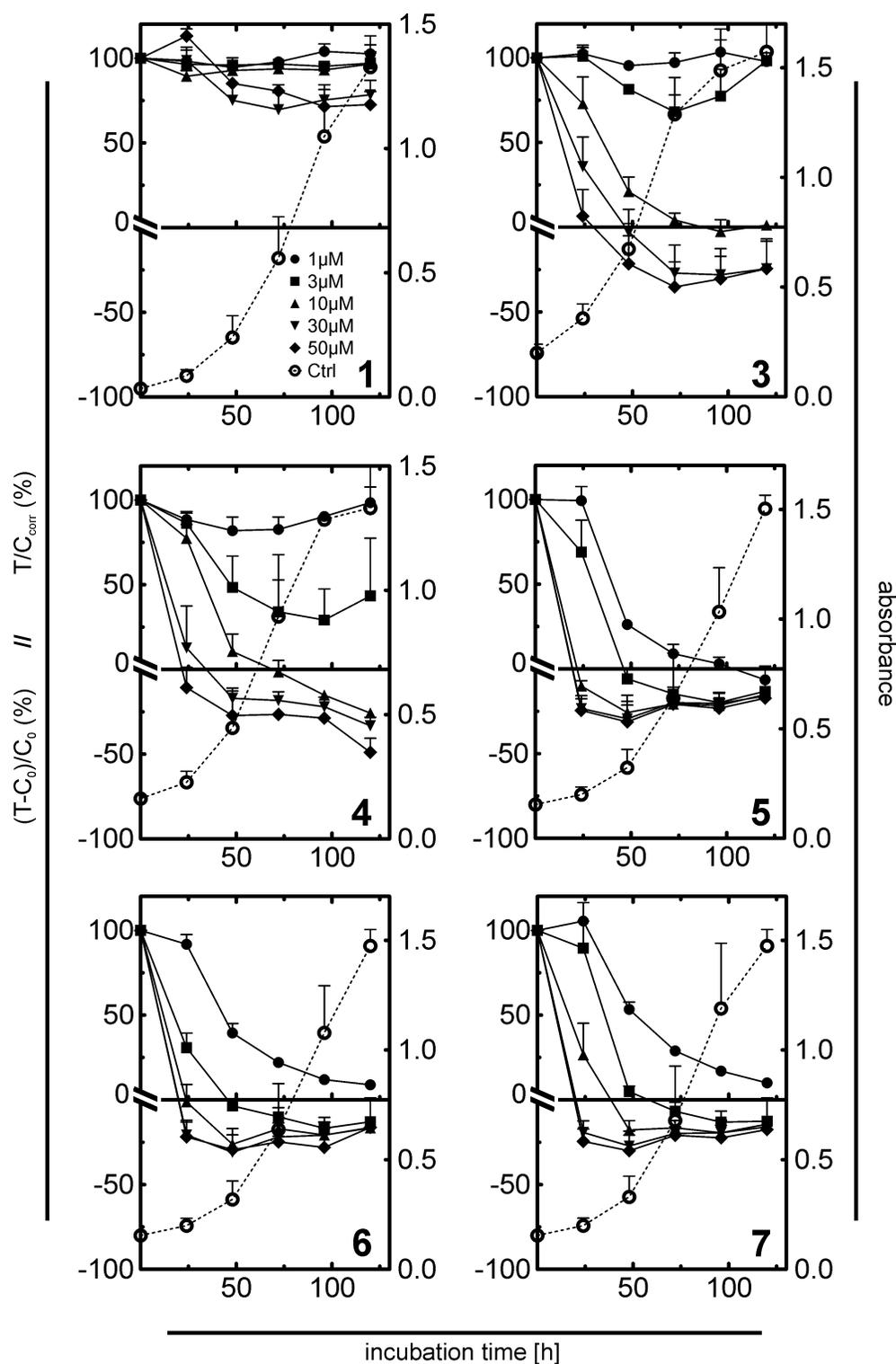


Figure S5.4: Chemosensitivity of Jurkat cells against compounds 1 and 3-7. Antiproliferative and cytotoxic effects correspond to the left y-axes. The growth curves of untreated cells (open circles) correspond to the right y-axes. Mean values \pm SEM of 2-3 independent assays with 8 replicates per concentration and time point.

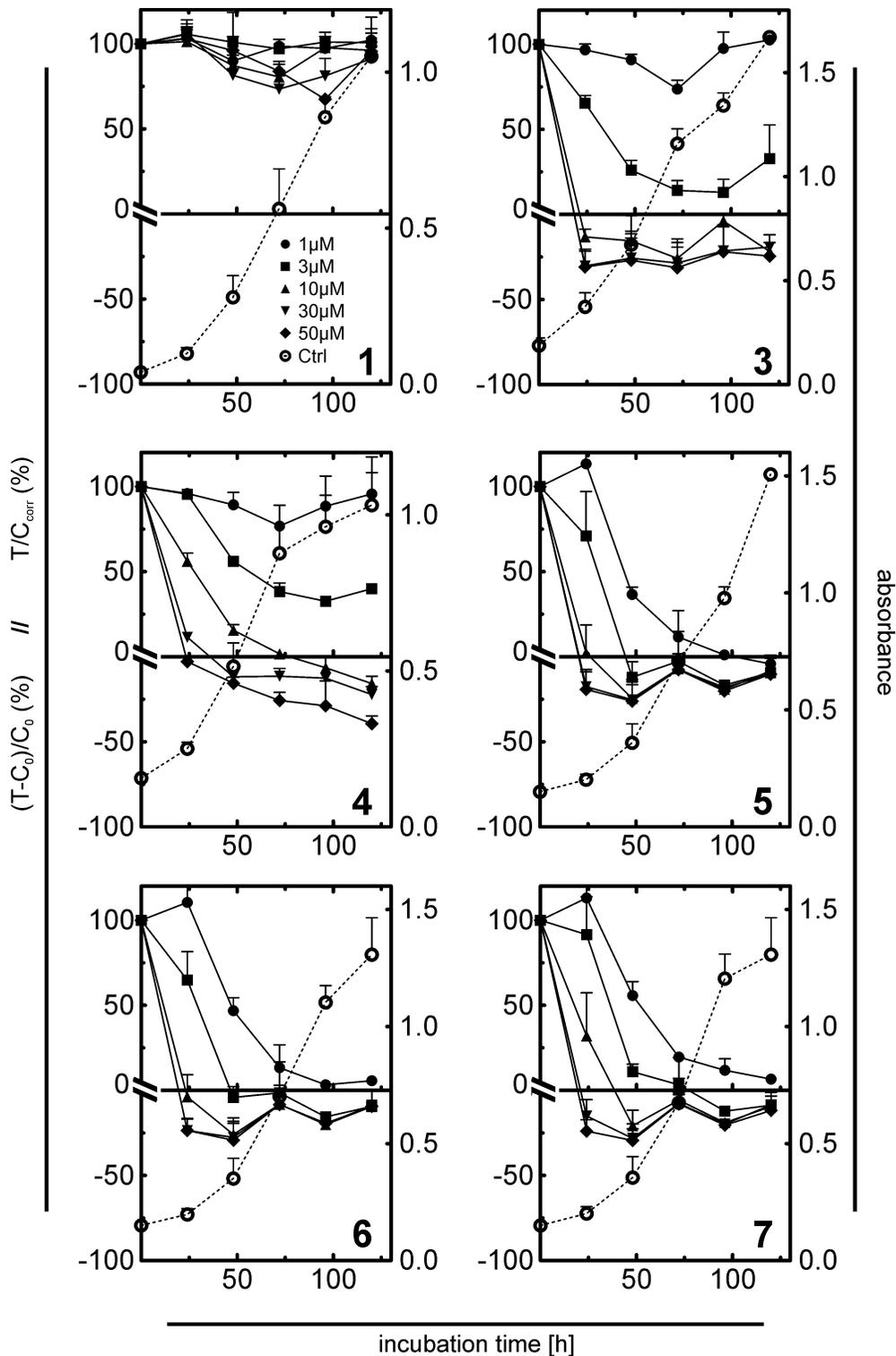


Figure S5.5: Chemosensitivity of U-937 cells against compounds 1 and 3-7. Antiproliferative and cytotoxic effects correspond to the left y-axes. The growth curves of untreated cells (open circles) correspond to the right y-axes. Mean values \pm SEM of 2-3 independent assays with 8 replicates per concentration and time point.

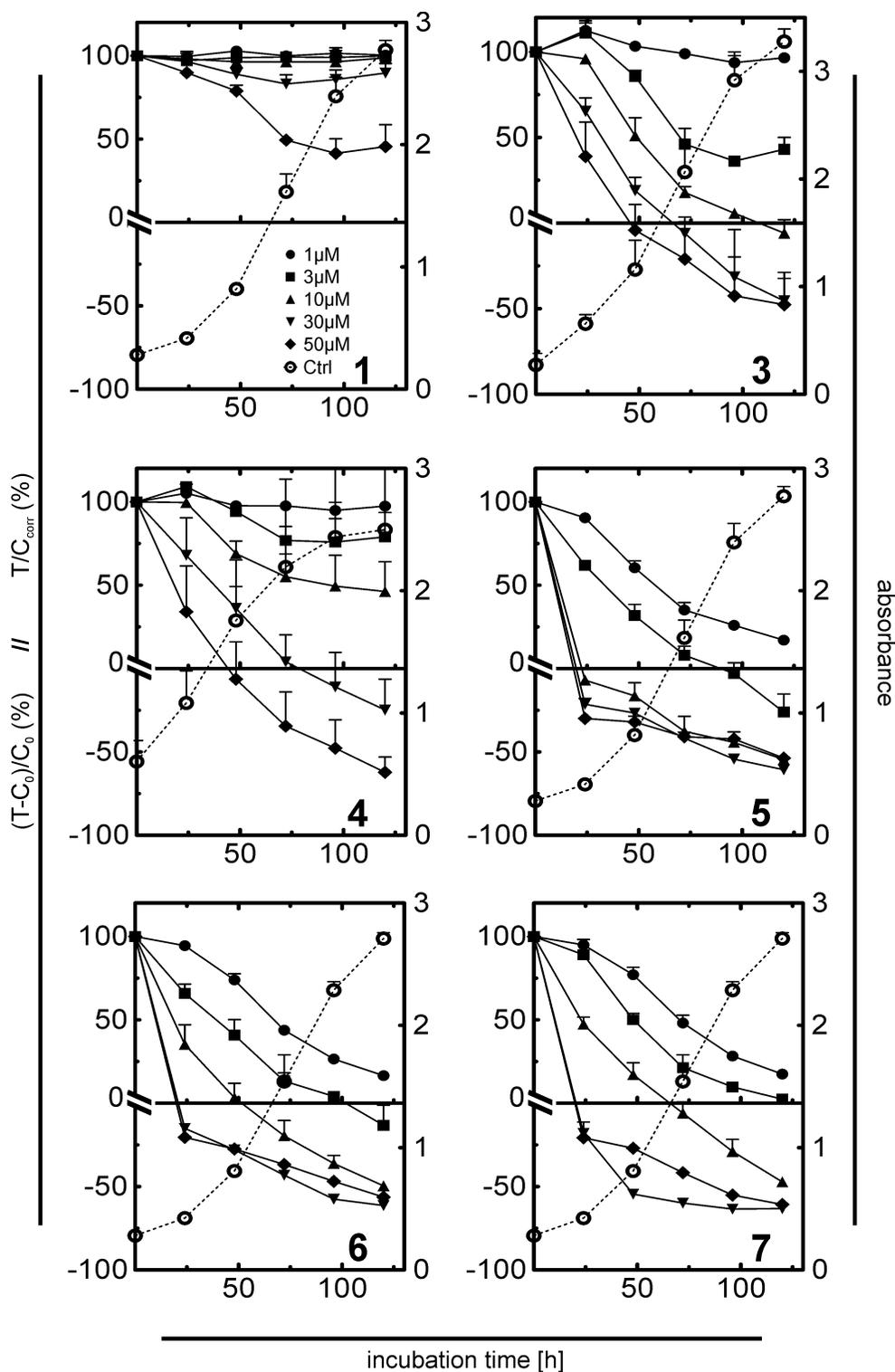


Figure S5.6: Chemosensitivity of MG-63 cells against compounds 1 and 3-7. Antiproliferative and cytotoxic effects correspond to the left y-axes. The growth curves of untreated cells (open circles) correspond to the right y-axes. Mean values \pm SEM of 2-3 independent assays with 8 replicates per concentration and time point.

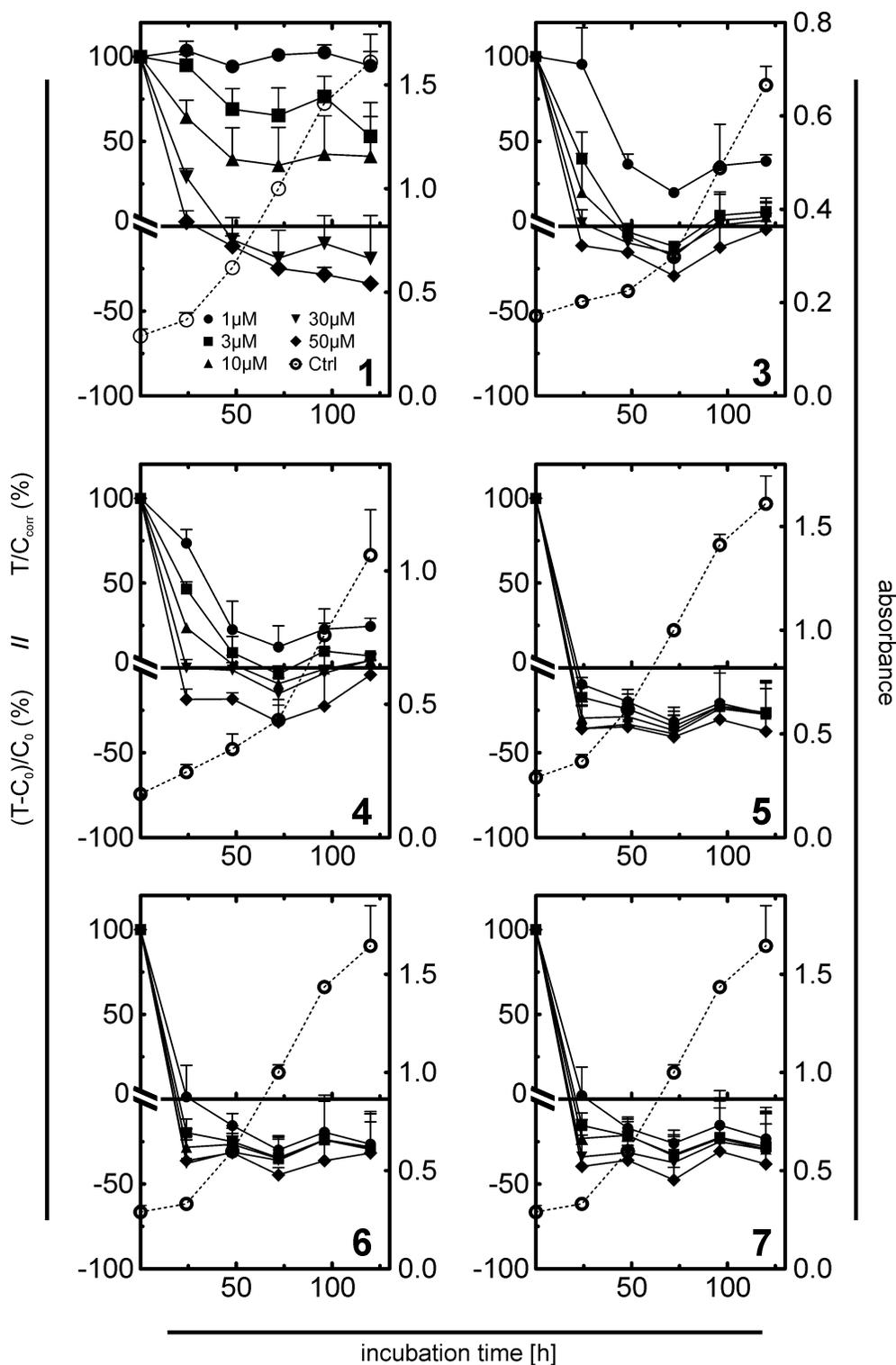


Figure S5.7: Chemosensitivity of SK-ES-1 cells against compounds 1 and 3-7. Antiproliferative and cytotoxic effects correspond to the left y-axes. The growth curves of untreated cells (open circles) correspond to the right y-axes. Mean values \pm SEM of 2-3 independent assays with 8 replicates per concentration and time point.

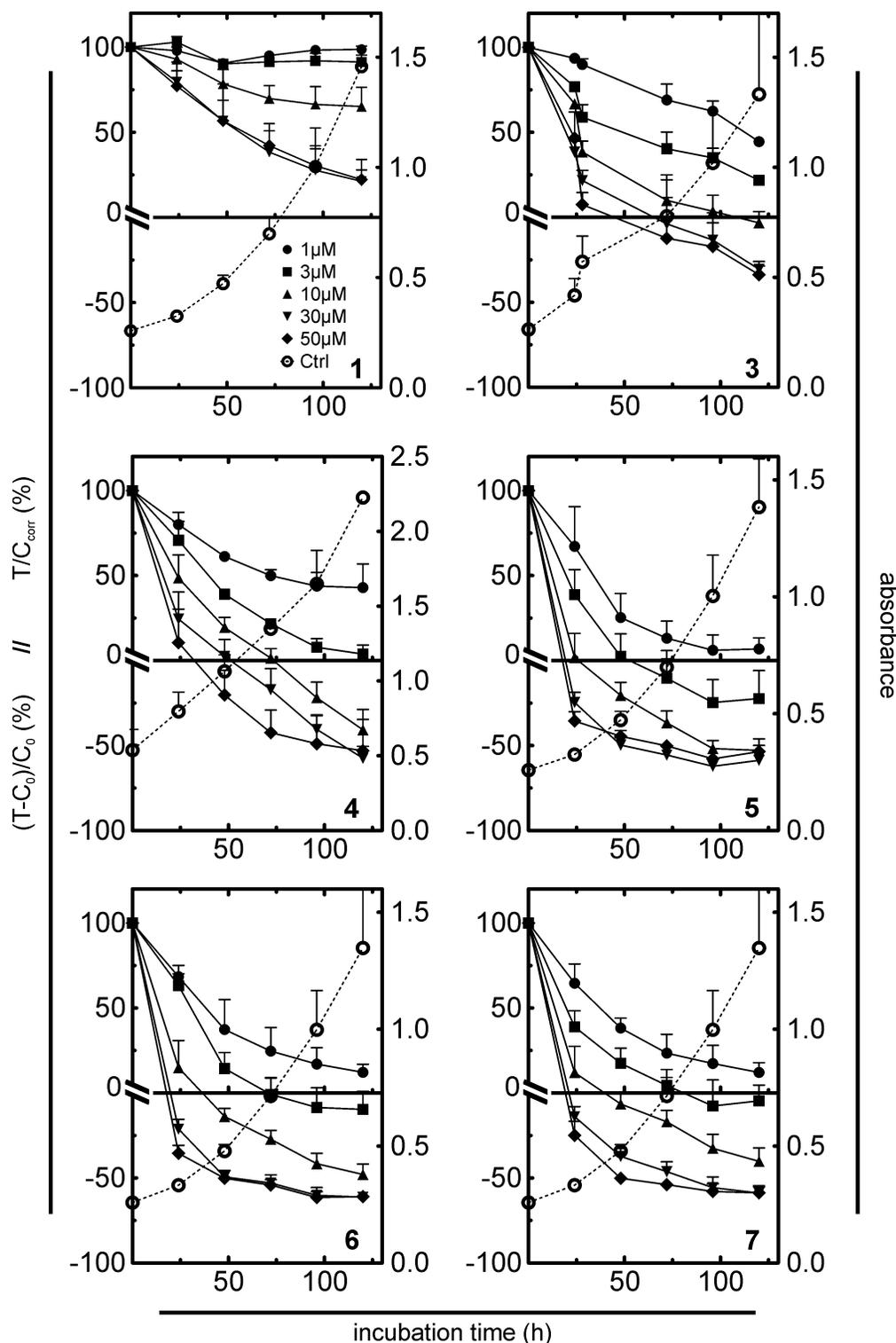


Figure S5.8: Chemosensitivity of Capan-1 cells against compounds 1 and 3-7. Antiproliferative and cytotoxic effects correspond to the left y-axes. The growth curves of untreated cells (open circles) correspond to the right y-axes. Mean values \pm SEM of 2-3 independent assays with 8 replicates per concentration and time point.

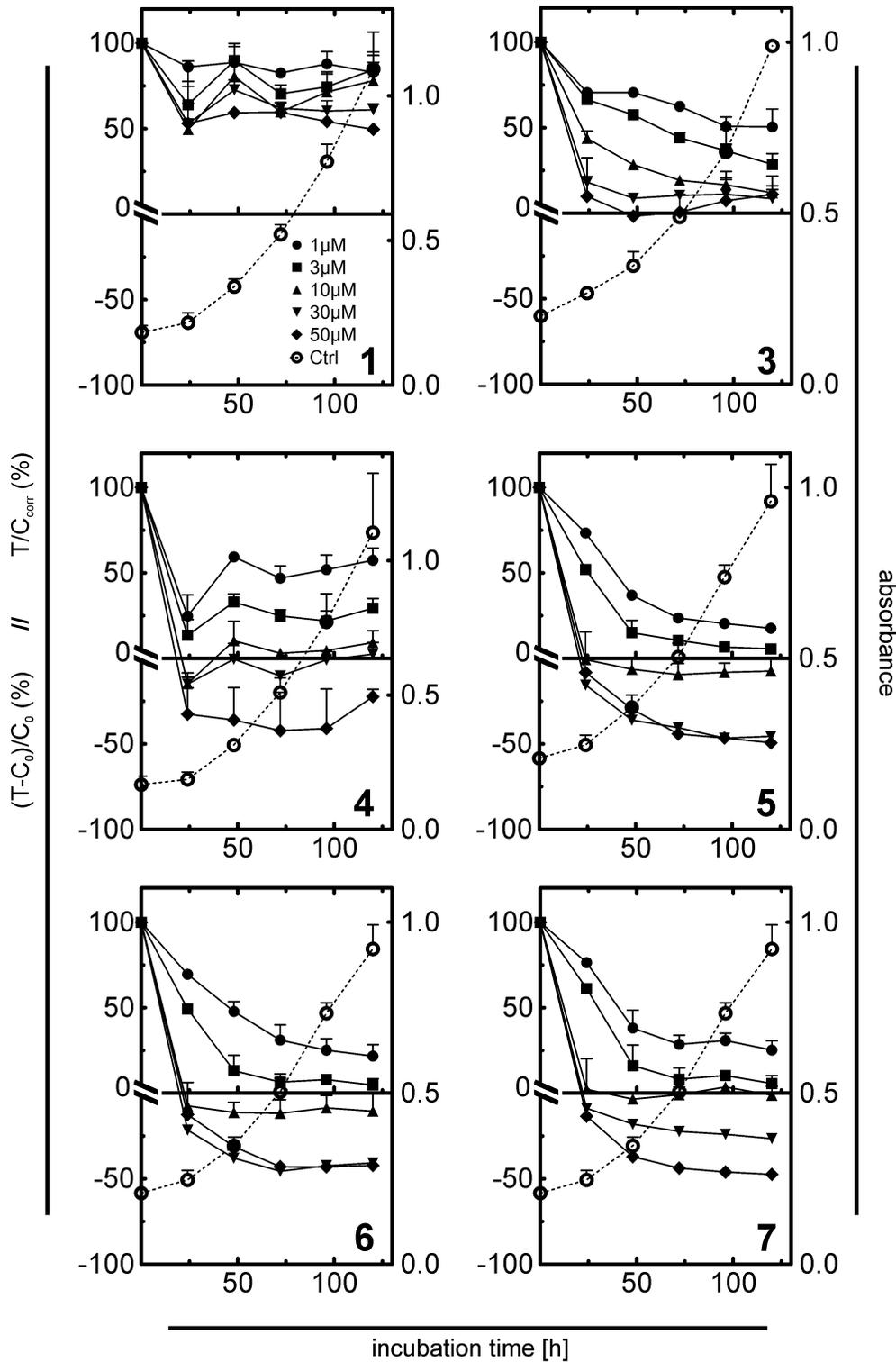


Figure S5.9: Chemosensitivity of LNCaP cells against compounds 1 and 3-7. Antiproliferative and cytotoxic effects correspond to the left y-axes. The growth curves of untreated cells (open circles) correspond to the right y-axes. Mean values \pm SEM of 2-3 independent assays with 8 replicates per concentration and time point.

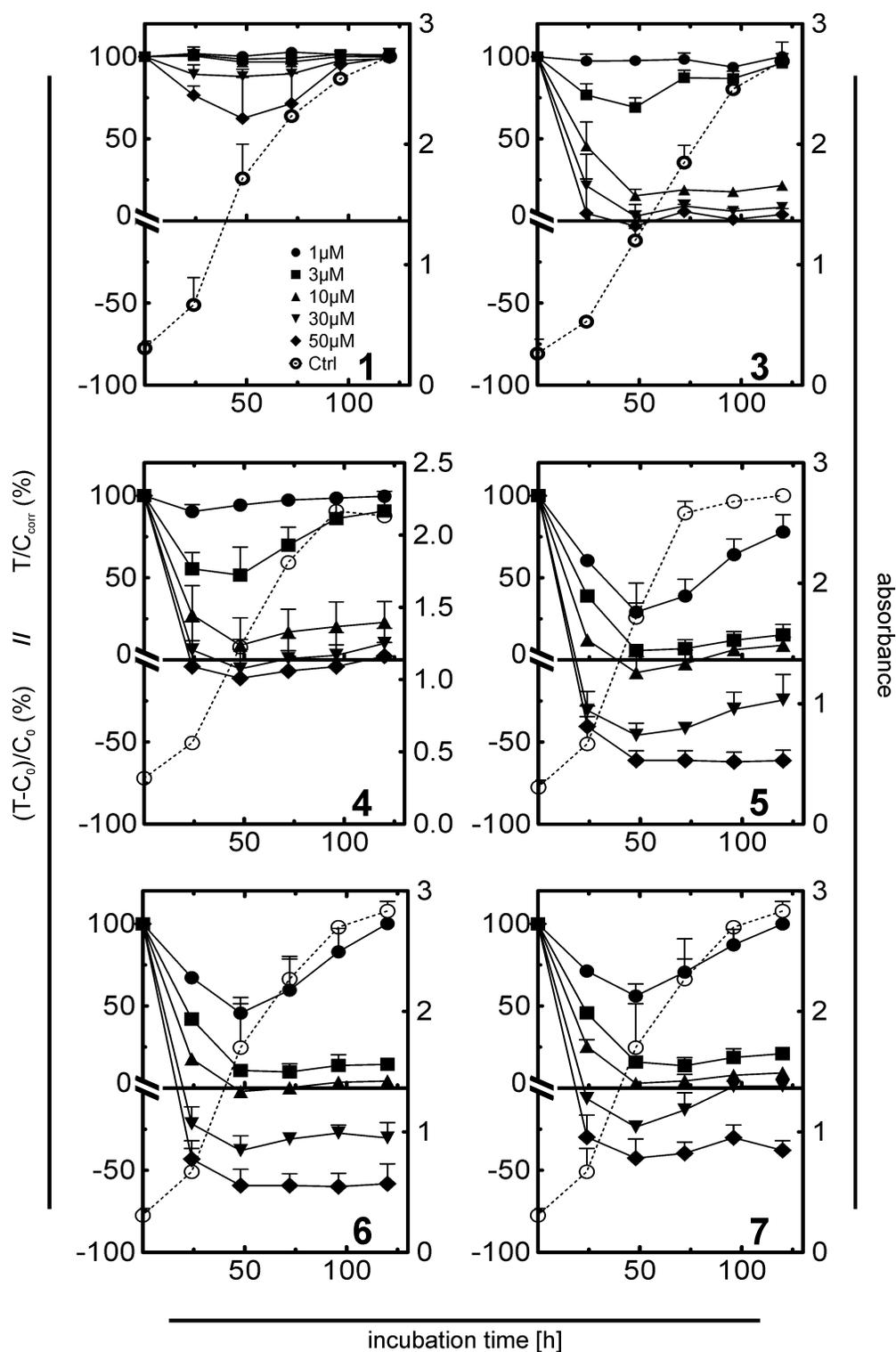


Figure S5.10: Chemosensitivity of NCI-H460 cells against compounds 1 and 3-7. Antiproliferative and cytotoxic effects correspond to the left y-axes. The growth curves of untreated cells (open circles) correspond to the right y-axes. Mean values \pm SEM of 2-3 independent assays with 8 replicates per concentration and time point.

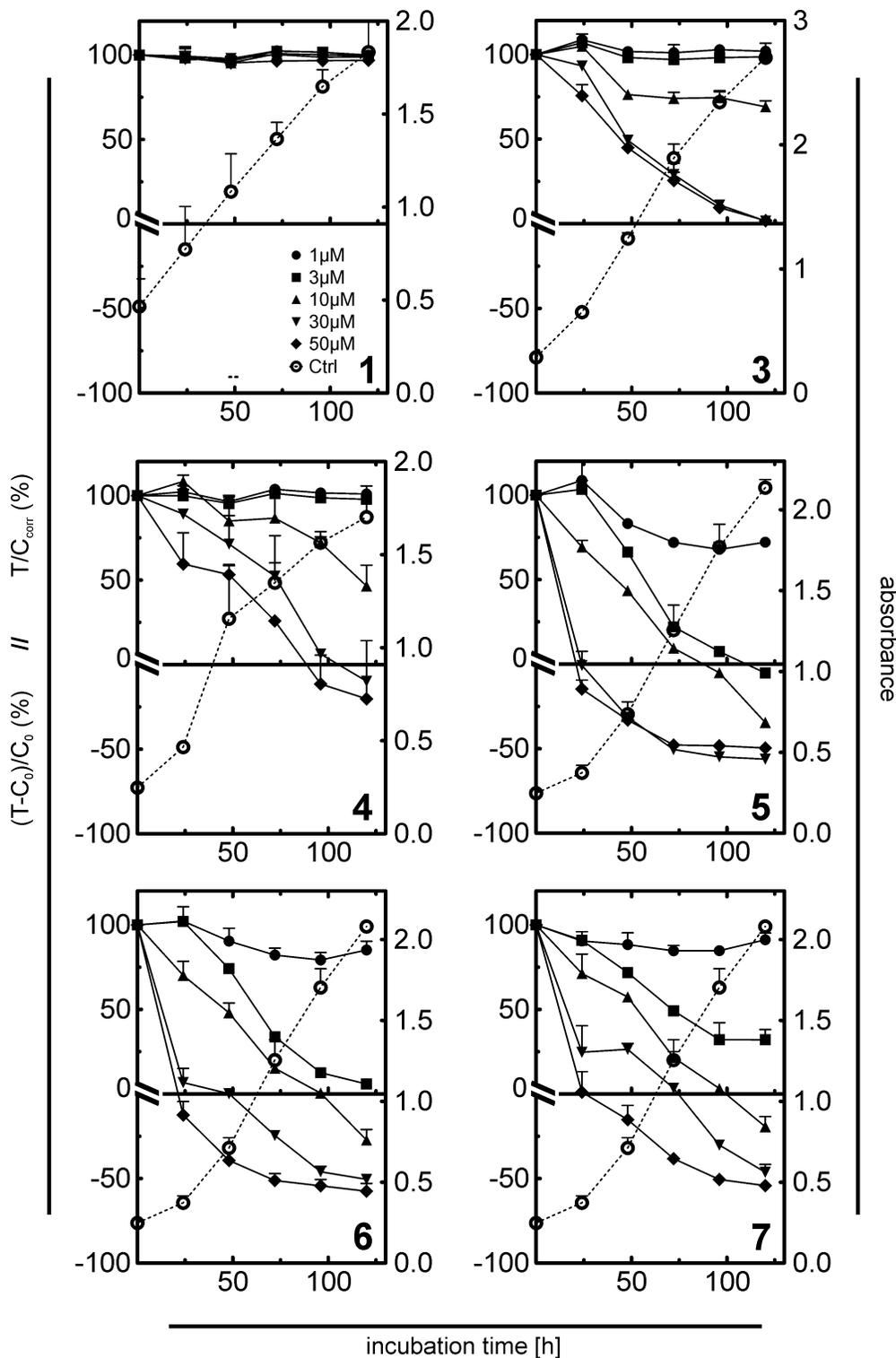


Figure S5.11: Chemosensitivity of HT-29 cells against compounds 1 and 3-7. Antiproliferative and cytotoxic effects correspond to the left y-axes. The growth curves of untreated cells (open circles) correspond to the right y-axes. Mean values \pm SEM of 2-3 independent assays with 8 replicates per concentration and time point.

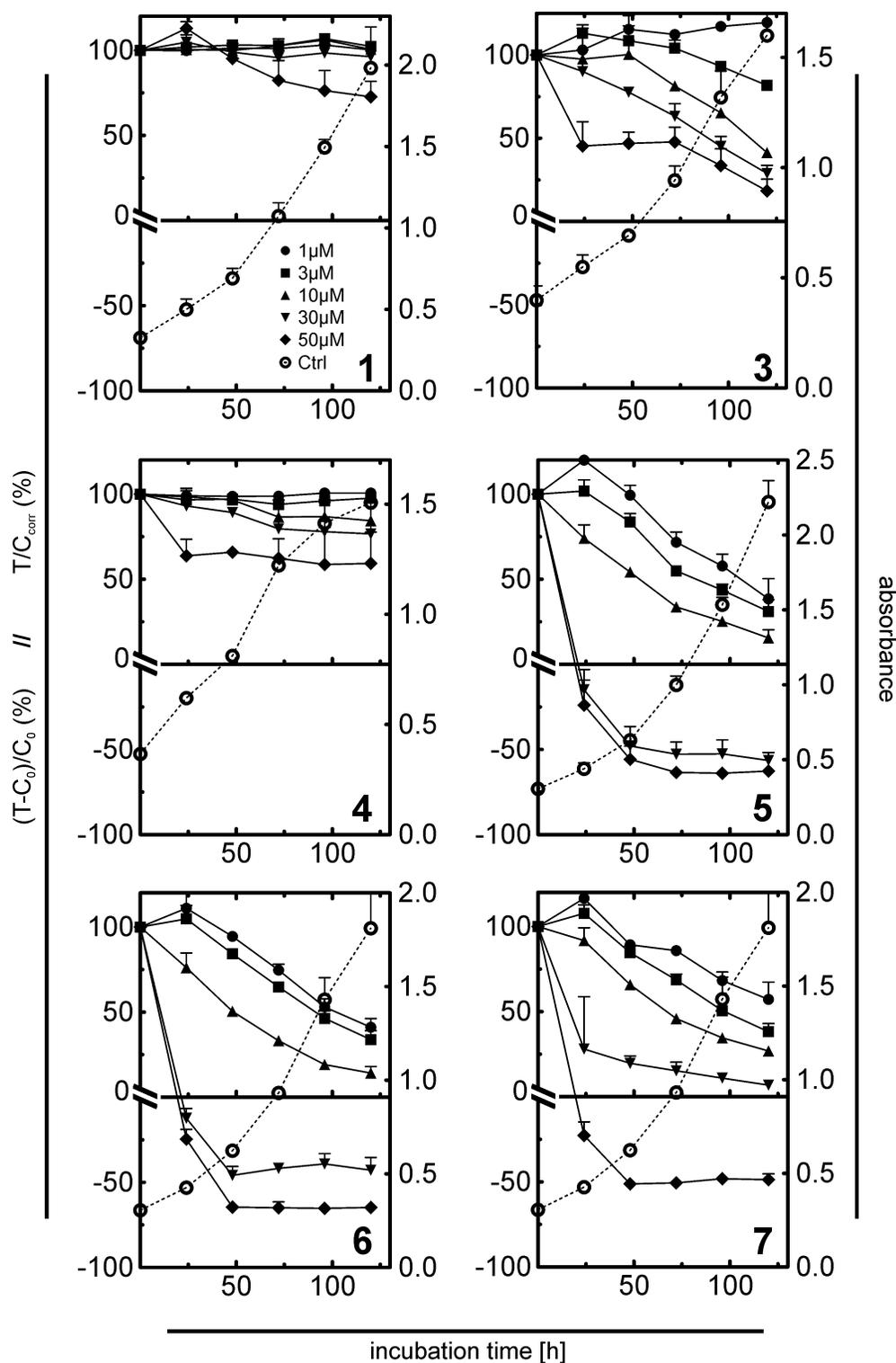


Figure S5.12: Chemosensitivity of SK-MEL-3 cells against compounds 1 and 3-7. Antiproliferative and cytotoxic effects correspond to the left y-axes. The growth curves of untreated cells (open circles) correspond to the right y-axes. Mean values \pm SEM of 2-3 independent assays with 8 replicates per concentration and time point.

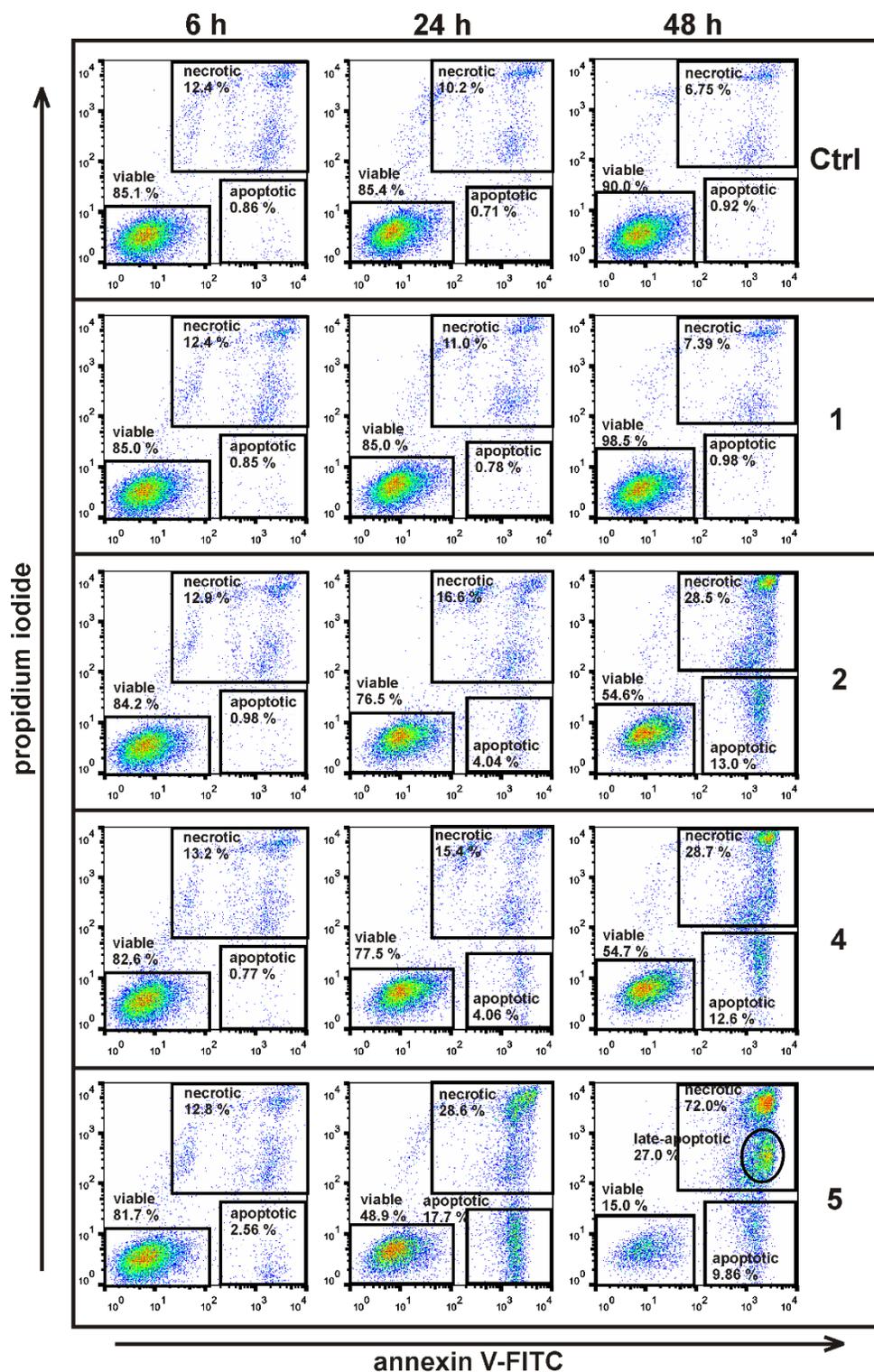


Figure S5.13: Flow cytometric analysis of Jurkat cells. Cells were treated with 10 μ M **1**, **2**, **4** and **5** after different periods of incubation (6, 24, 48 hours) by annexin V-FITC (x-axis) and propidium iodide (y-axis) staining. Samples of untreated cells were collected at the same time points and served as control (**Ctrl**). The indicated percentages are related to the total number of collected single cells. The different test compounds are arranged horizontally.

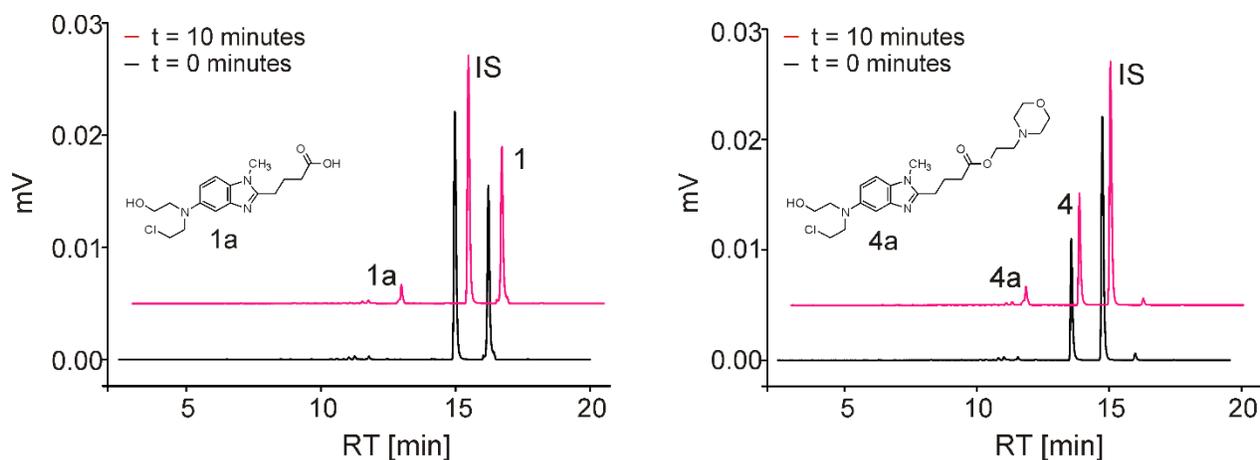


Figure S5.14: Stability of compounds **1 and **4** in the presence of NCI-H460 cells.** Representative chromatograms, indicating the stability of compounds **1** and **4** in the presence of NCI-H460 cells (IS = internal standard, umbelliferone). Samples were measured immediately before the incubation ($t = 0$ minutes) and after 10 minutes of incubation at 25 °C ($t = 10$ minutes). The kinetics of the hydrolysis of the N-Lost group, yielding **1a** or **4a**, is the same in case of bendamustine (**1**) and the morpholinoethyl ester **4**, respectively. Other decomposition products were not detected. The same holds for compounds **2** and **5** upon incubation with cells under the same conditions (data not shown).

5.6 References

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6 Chapter VI

Summary

ABC-transporter expression is a crucial mechanism of tumor resistance and was recently also associated with cancer initiating cells (CIC). Flow cytometry was applied to detect the expression of CIC-markers, ABCB1 and ABCG2 in human brain- and breast cancer cell lines and to examine possible correlations between ABC-transporter expression and CICs. Only one of the investigated brain tumor cell lines (Daoy) comprised cells expressing CIC-markers which revealed increased clonogenicity *in vitro*.

For investigations on the correlation between CICs and ABCG2, MCF-7 breast cancer cells were treated with topotecan (Topo) to induce ABCG2 expression. Topo treatment led to increased resistance against cytostatic drugs, which are substrates of ABCG2. Concurrently, Topo treatment induced a reversible decrease in the expression of the CIC-markers CD24 and EpCAM, but did not result in increased clonogenicity *in vivo*. In contrast, Topo treated cells revealed decreased tumorigenicity, most probably due to the down-regulation of CD24 and EpCAM. Nevertheless, further investigations are necessary to determine whether ABCG2 induction and CD24/EpCAM down-regulation are interrelated or independent processes.

The nitrogen mustard (N-Lost) derivative bendamustine (BM) has been approved as an anticancer drug for the treatment of hematopoietic malignancies for decades. Nevertheless, as becomes obvious from a series of recent publications and patent applications, there is increasing interest in this cytostatic with respect to, for example, improved oral bioavailability of BM by optimized formulations. Following a different approach, in this thesis the potential of esters of BM was explored, with a focus on representative aminoalkyl esters. Alkyl- and various aminoalkyl (e.g. 2-morpholinoethyl or 2-pyrrolidinoethyl) esters of BM were investigated for stability in various media and cytotoxicity *in vitro* against a broad panel of human malignancies.

For this purpose, a fast, selective RP-HPLC method using fluorescence detection was established and validated. This method enabled the separation of complex mixtures of analytes and degradation products with low limits of quantification and high accuracy and precision. Liquid phase extraction with perchloric acid yielded high recovery rates and proved to be applicable to the determination of the stability of BM and BM esters in biological material.

In aqueous solution, the half-life of the N-Lost moiety remained essentially unchanged upon esterification of BM. The N-Lost group was significantly more stable in murine plasma, whereas the ester bonds were prone to enzymatic cleavage. Interestingly, the nitrogen mustard group was even more stable in human than in murine plasma. The stability of the ester bond substantially depended on the chemical nature of the substituents at the ester group. Kinetic studies in the presence of plasma, butyrylcholine esterase and physostigmine revealed that the basic esters are substrates of unspecific choline esterases. While alkyl esters and the mofetil ester had significantly longer half-lives in human compared to murine plasma, basic esters, except for the branched 1-methyl-2-pyrrolidinoethyl ester, were cleaved rapidly. The discrepancies between the

species-dependent stabilities of the BM derivatives can be attributed to lower albumin content and higher enzymatic activity in mouse compared to human plasma.

Surprisingly, cytotoxicity studies revealed that the BM derivatives possess considerably increased antiproliferative activity against a panel of human cancer cells, representing not only hematologic malignancies but also solid tumors (e.g. malignant melanoma, colorectal carcinoma and lung cancer), which are resistant to the parent compound BM. Although the investigated compounds are sufficiently stable to act as antitumor agents on their own, it cannot be precluded that the esters are prodrugs, allowing for increased intracellular accumulation of BM. Especially, basic esters, positively charged under physiological conditions, were up to approximately 100 times more effective than BM. This was paralleled by a higher fraction of early apoptotic cancer cells and increased expression of p53.

As HPLC analyses revealed cellular enrichment of the esters with the highest accumulation factor in case of the pyrrolidinoethyl ester, transport by organic cation transporters OCT1 and OCT3 was taken into account. Although differential interactions of the BM derivatives with these transporters were confirmed *in vitro*, the OCT expression (mRNA level) of the investigated human sarcoma and carcinoma cells did not support the hypothesis of an OCT mediated uptake contributing to the drastic increase in chemosensitivity.

The presented approach to increasing cellular access, enhancing antiproliferative activity – also against cancer cells refractory to the parent compound - by chemical derivatization might also add value to alkylating agents other than bendamustine.

Abbreviations

5-FU	5-fluorouracil
ABC-transporter	ATP-binding cassette transporter
AF	alexa fluor
ALDH	aldehyde dehydrogenase
APC	allophycocyanin
ASP ⁺	4-(4-(dimethylamino)styryl)-N-methylpyridinium iodide
BCIC	breast cancer initiating cell
BCRP	breast cancer resistance protein
bFGF	basic fibroblast growth factor
BM	bendamustine
BP	band pass
BTIC	brain tumor initiating cell
BV-421	brilliant violet 421
CIC	cancer initiating cell
CK	cytokeratin
CLL	chronic lymphocytic leukemia
CSC	cancer stem cell
Ctrl	control
DMSO	dimethyl sulfoxid
Doxo	doxorubicin
EDTA	ethylenediaminetetraacetic acid
EGF	epidermal growth factor
EGFR	epithelial growth factor receptor
EGTA	ethyleneglycol-bis(aminoethylether)-N,N,N',N'-tetraacetic acid
EMT	epithelial mesenchymal transition
FACS	fluorescence activated cell sorting
FCS	fetal calf serum
FITC	Fluorescein isothiocyanate
FSC	forward scatter
FSC-H	forward scatter height
FSC-W	forward scatter width
FTC	fumitremorgin C
HE	hematoxylin-eosin
HER	human estrogen receptor
HPLC	high performance liquid chromatography
HRP	horseradish peroxidase
IS	internal standard
LOD	limit of detection
LOQ	limit of quantification
LP	long pass
mAb	monoclonal antibody
MeCN	acetonitrile

MFI	mean fluorescence intensity
MG	Masson-Goldner
Mito	mitoxantrone
MM	multiple myeloma
MOA	mechanism of action
MTT	thiazolyl blue tetrazolium bromide
MTX	methotrexate
NHL	non-Hodgkin lymphoma
NP-40	nonidet P-40
OCT	organic cation transporter
PB	pacific blue
PBS	phosphate buffered saline
PBS/FCS	PBS + 2 % FCS
PCR	polymerase chain reaction
PE	phycoerythrin
P-gp	P-glycoprotein
PI	propidium iodide
RP	reverse phase
RSD	relative standard deviation
RT	room temperature
SDS	sodium dodecyl sulfate
SP	side population
SSC	side scatter
$t_{1/2}$	half life
Topo	topotecan
TPA	tetrapentyl ammonium
TPC	tumor propagating cell
WT	wild type