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***In Vitro* Sensitivity Testing in the Assessment of Anti-CLL Drug Candidates**

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1. Introduction

Chronic lymphocytic leukemia (CLL) is characterized by the accumulation of morphologically mature, but immuno-incompetent B-lymphocytes in the bone marrow, peripheral blood, spleen and lymphoid organs. With an annual incidence of about 2-3/100000 in the general population (Hamblin, 2009), CLL represents a frequent leukemia type. Since CLL mostly affects persons of advanced age, the incidence among persons above 65 years reaches ten times this frequency (Eichhorst et al., 2009). Moreover CLL follows a remarkably heterogeneous course, emphasizing the need for personalized treatment approaches. Despite recent advances in CLL therapy, the disease still remains incurable and new treatment options need to be developed (Hallek et al., 2008). New insights into CLL biology have started to result in new targeted, sometimes patient group-specific treatment approaches (Pleyer et al., 2009; Zenz et al., 2010). Candidate substances for pre-clinical assays are mostly molecularly targeted drugs, i.e. either small molecules interfering with intracellular signaling (Wickremasinghe et al., 2011) or monoclonal antibodies (Jagłowski et al., 2010). As examples we will discuss in this chapter the pre-clinical assessment of protein and lipid kinase inhibitors and of monoclonal antibodies.

Since candidate substances with a potential for treating CLL become available at an increasing pace, there is a growing need for comprehensive laboratory assessment of these substances. For this purpose the effects of these drug candidates on fresh CLL lymphocytes are compared by means of viability and cytotoxicity assays with the aim of selecting suitable candidates for further development. In addition viability and cytotoxicity assays on CLL cells serve to prepare candidate substances for clinical trials and to determine, which subgroups of patients respond best, in the sense of personalized medicine. This endeavor constitutes the small excerpt of the drug discovery process immediately preceding clinical trials (Collins & Workman, 2006). Since it links laboratory investigation and clinical application it can be understood as translational research, which is further underscored by patient samples being subjected to cultivation and observation in the laboratory.

B lymphocytes freshly isolated from peripheral blood of CLL patients constitute a readily available source for the pre-clinical *in vitro* assessment of drugs and combinations with therapeutic potential for treating CLL. Due to the epidemiological features of CLL as a frequent chronic leukemia, with many patients living with the disease for extended periods, a continuous supply of blood samples can be made available and used in a meaningful manner by performing pre-clinical assays, which, according to the concept of translational medicine, in turn could lead to improved therapies. Drug assessment on fresh CLL samples can be performed rapidly and relatively conveniently by comparing untreated and treated *in vitro* cultures.

On the other hand, primary cultures of B cells freshly isolated from the peripheral blood of CLL patients often may represent an insufficient model for predicting clinical drug efficacy, since they are known to behave differently from CLL lymphocytes in their *in vivo* environment. This is evident from the obviously contrasting behavior of CLL cells *in vivo* and during *in vitro* culture. Whereas accumulation of CLL cells *in vivo* is thought to occur due to resistance towards apoptosis and a certain degree of cell proliferation, *in vitro* cultures spontaneously undergo apoptosis and show low viability and almost completely absent proliferation. Because this contrasting behaviour of CLL cells *in vitro* and *in vivo* can be attributed to a lack of the appropriate micro-environment during laboratory culture, the value of drug assessment on CLL lymphocytes *ex vivo* can be greatly enhanced by mimicking certain micro-environmental stimuli.

Commonly used cytotoxicity and viability assays are compiled in this chapter and will be discussed in the context of the assessment of potential anti-CLL drugs. On the level of individual susceptibility it is well established for chemotherapeutic agents that sensitivity of tumor cells *in vitro* corresponds to the probability of clinical response (Bosanquet et al., 2009). Therefore one would expect also for targeted drugs that *in vitro* assays enable to some degree the comparison of the efficacies of different agents and the prediction of the response of molecularly defined subgroups of CLL patients. As examples for correlations of molecularly defined patient subgroups with treatment susceptibility *in vitro* we name here the clearly higher dasatinib sensitivity of CLL samples with unmutated IgHV genes as compared to mutated ones (Veldurthy et al., 2008) or the correlation of the B cell depletion induced by CD20 antibodies with antigen expression on the surface of CLL cells (Patz et al., 2011).

In this chapter we review the pros and cons of pre-clinical drug assessment in comparatively simple *ex vivo* assays. The predictability of treatment out-come from *in vitro* cultures of CLL lymphocytes must be considered, since there are known limitations of the assay system, which can, however, be overcome to a certain degree by linking the results to investigations of target and cell type specificity.

2. Performing cytotoxicity assays on CLL samples

In the course of the pre-clinical assessment of anti-CLL drug candidates, *in vitro* cultures of CLL lymphocytes are treated with test substances. Subsequently dose-dependent treatment effects on the viability of CLL cells are recorded by means of established proliferation and cytotoxicity assays. These assays yield a first measure of drug potency for CLL lymphocytes, but certainly need to be rigidly controlled and standardized. Moreover they should be

accompanied by biochemical assays in order to assure cell type and target specificity. Such comprehensive approaches will allow the design of meaningful drug combinations, which then are to be subjected to another round of *in vitro* sensitivity assays.

Cytotoxicity assays play a pivotal role in pre-clinical drug testing (Kepp et al., 2011) and many of them are suitable for assessing treatment effects on fresh CLL cells (Table 1). Some of these assays are based on absolute cell counts or less laborious surrogate parameters determining total cellular activities e.g. in colorimetric or fluorimetric non-clonogenic microculture assays (Lindhagen et al., 2008). In contrast, flow cytometric assays usually yield percentages of cells with certain properties within the investigated cell population unless they are standardized for the examined volume, e.g. by absolute counting beads.

Method	Technological platform	parameters
cell counting	light microscope	trypan blue exclusion
metabolic activity	absorbance reader	tetrazolium salt reduction
ATP consumption	luminometer	luciferase activity
intracellular esterase activity	fluorimeter	conversion of non-fluorescent fluorescein diacetate
DNA replication	absorbance reader flow cytometry	bromo-deoxyuridine incorporation
phosphatidylserine exposure	flow cytometry	annexin V-binding
membrane disintegration	flow cytometry	staining with DNA intercalating dye
$\Delta\psi$ m dissipation	flow cytometry	fluorescent dye, e.g. DiOC6
morphology	flow cytometry	forward scatter / side scatter
production of reactive oxygen species	flow cytometry	fluorogenic substrate, e.g. CM-H2-DCF-DA
caspase activation	flow cytometry immunoblotting	fluorescent substrates detection of cleaved fragments

Table 1. Selected cytotoxicity and viability assays commonly used with CLL samples.

2.1 Flow cytometric cell death and viability assays

Flow cytometric assessment of phosphatidylserine exposure and membrane disintegration is among the viability and cytotoxicity assays most frequently applied for monitoring drug effects on CLL cells. Percentages of cell populations undergoing cell death can be determined by monitoring the loss of membrane asymmetry in early phases of apoptosis and subsequent membrane disruption (Fig. 1). This can be achieved by staining cells, e.g. B lymphocytes, with fluorescently labeled annexin V, which binds to phosphatidylserine with high affinity (Koopman et al., 1994). Phosphatidylserine exposure accompanies early phases of apoptosis, before membrane disintegration of the cytoplasmic membrane allows access of DNA intercalating dyes to the nucleus. Counter-staining with DNA intercalating dyes originally served the distinction of apoptotic from necrotic cells. Analysis of annexin V binding or DNA staining can be replaced for staining with dyes indicating mitochondrial membrane potential, e.g. 3,3'-dihexyloxacarbocyanine iodide (DiOC6) (Stanglmaier et al.,

2004; Veldurthy et al., 2008), or reactive oxygen species, e.g. the fluorogenic chloromethyl-2,7-dichlorodihydrofluorescein-diacetate (CM-H₂-DCF-DA) (Lilienthal et al., 2011). Usually populations of CLL cells with phosphatidylserine exposure coincide with those showing typical morphological signs of apoptosis, i.e. reduced size and increased granularity as indicated by forward-scatter (FSC) and side-scatter (SSC) in the flow cytometer. Concerns have been raised about a possible over-estimation of drug effects due to flow cytometry artefacts. For instance, the widely used determination of phosphatidylserine exposure was claimed to over-estimate apoptosis induction in the extraordinarily fragile CLL cells due to *in vitro* handling during sample preparation for flow cytometry (Groves et al., 2009). Similarly antibody effects on CLL samples determined by flow cytometry were suspected of being misinterpreted owing to cell aggregation (Golay et al., 2010). These concerns can be overcome by parallel biological effect monitoring in several different assay systems as controls.

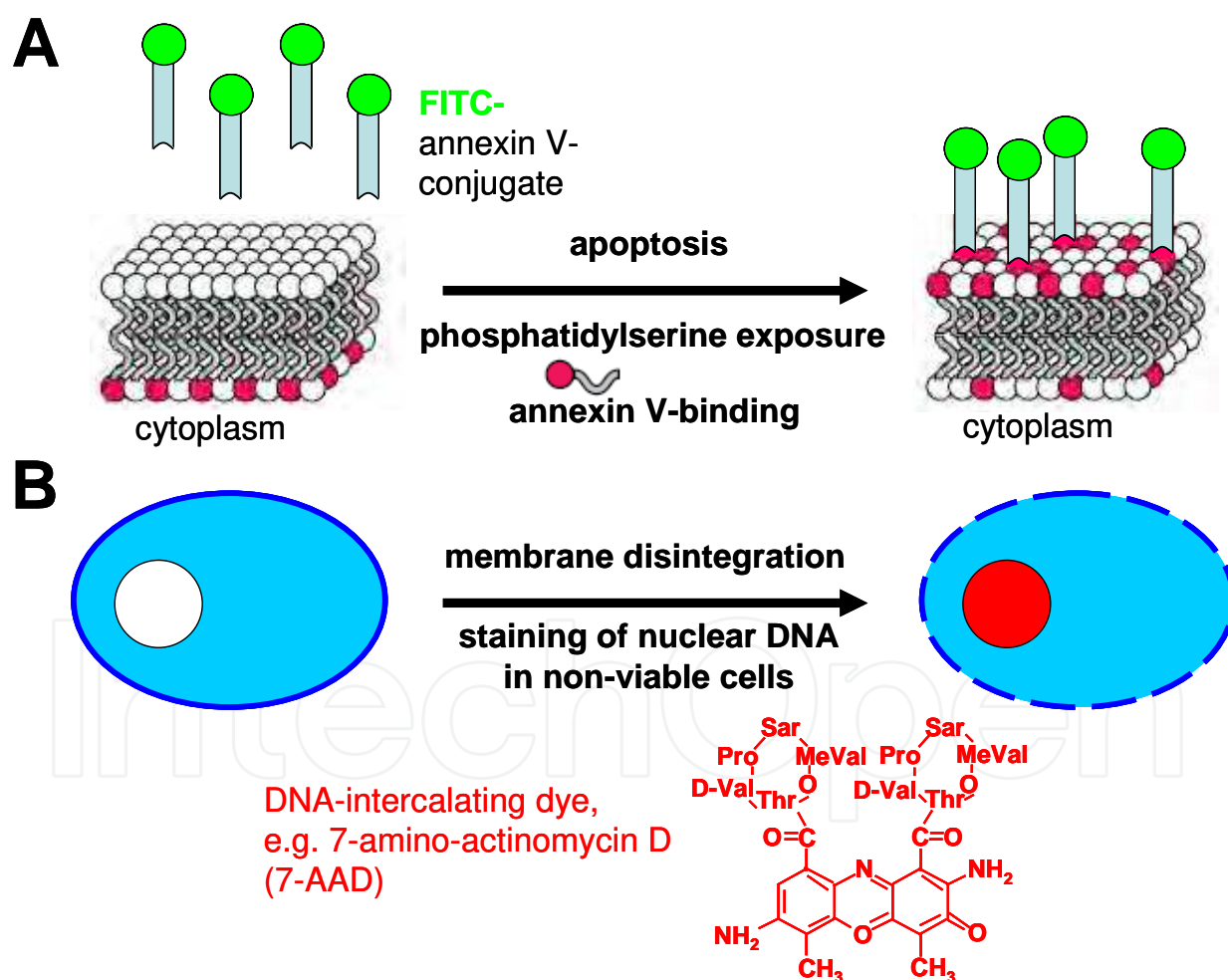


Fig. 1. Principle of a widely used assay for the determination of the percentages of apoptotic cells. The flow cytometric assessment of phosphatidylserine exposure (A) and membrane disintegration (B) can be performed simultaneously.

Part (A) adapted from Zhang et al., 1997 and Pharmingen, 1998.

2.2 Concentration dependence

Investigation of the dose-dependency of drug effects is an important confirmation of any observations that by far surpasses the importance of repeated measurements. In addition, the shape of dose response curves sometimes can provide mechanistic clues, e.g. in the case of saturation effects. When drugs are investigated for CLL that previously have been pre-clinically developed or admitted for the treatment of other cancers, it is possible to consider achievable plasma concentrations for the *in vitro* assessment. Biological effect measurements obtained at multiple concentrations can be conveniently summarized by concentrations inhibiting 50 % (IC_{50}). These IC_{50} concentrations can be a useful manner of comparing the effects of different drugs or diverging sensitivities of different samples. In the case of saturation effects, as in the dose response of CLL lymphocytes to dasatinib (Veldurthy et al., 2008), it might be more appropriate to indicate the individual saturation levels of response rather than extrapolating IC_{50} concentrations. Moreover, investigators must not be misled by the convenience of such tabulated values to regard them as sample-specific constants and therefore to apply them to different assay types, since IC_{50} concentrations strictly depend on the type of assay performed (Krause & Hallek, 2011).

3. Mimicking micro-environmental interactions

Considering the micro-environment of CLL cells may improve predictions of clinical drug efficacy from *in vitro* assays on fresh CLL samples. Regarding their resistance to apoptosis CLL cells in culture behave entirely differently from the situation *in vivo*, owing to dependence on their micro-environment. Therefore it is necessary to simulate certain micro-environmental stimuli for drug assessment *in vitro*, for instance following the approaches described in the following subchapters. A number of ligand/receptor pairs have been identified that activate CLL cells (Fig. 2) (Munk Pedersen & Reed, 2004), among them the chemokine stroma-derived factor 1, nowadays referred to as CXCL12 and its receptor CXCR4 on CLL cells, that belongs to the class of G-protein-coupled 7-transmembrane domain receptors (Burger & Kipps, 2006) and VCAM-1 (CD106) expressed on the surface of stroma cells that interacts with the integrin VAL-4 (CD49a) on CLL cells (Burger et al., 2009). The strict dependence of CLL cells on the interactions with their environment is also apparent from the absence of good cell line models for CLL.

3.1 B cell receptor stimulation

Like for normal lymphocytes, also the fate of CLL cells is to a high degree determined by B cell receptor (BCR) stimulation (Stevenson & Caligaris-Cappio, 2004). According to the degree of somatic hypermutation in rearranged antigen receptor genes, subgroups of CLL clones with immunoglobulin heavy chain variable region (IgHV) genes can be distinguished that reflect progressing B cell development stages corresponding to naïve or memory B cells. Usually the threshold separating these molecularly defined prognostic subgroups is set at 2 % sequence divergence of rearranged IgHV genes from the closest germline sequences. The CLL subgroups with unmutated or mutated IgHV genes have a widely different prognosis indicated by 24 versus 8 years median overall survival after diagnosis (Hamblin et al., 1999). High expression of zeta-associated protein 70 (ZAP-70) and of CD38 serve as surrogate markers of unmutated IgHV genes (Crespo et al., 2003; Hamblin et al., 2002).

Antigen contact for CLL cells can be mimicked *in vitro* by crosslinking surface IgM by means of anti-IgM antibodies. Long lasting stimulation of the BCR leads to prolonged survival of CLL cells (Deglesne et al., 2006). This can be achieved by using soluble anti-IgM or anti-IgM-coated surfaces.

3.2 CD40 stimulation

The CD40 molecule expressed on the surface of CLL cells belongs to the tumor necrosis factor family and participates in antigen recognition as a co-receptor. Its cognate ligand, CD40 ligand (CD40L), also known as CD154, is expressed on the surface of activated T lymphocytes. Engagement of CD40 on CLL lymphocytes mimics the micro-environment inside lymph nodes and leads to protection against DNA damaging substances, e.g. chemotherapeutic agents. CD40L stimulation of CLL cells can be provided by co-culture with fibroblasts expressing recombinant CD154. For instance co-culture with CD40L expressing fibroblasts protects CLL cells from DNA damaging agents, but this effect can be partly reversed by the kinase inhibitor dasatinib (Hallaert et al., 2008). The sensitivity of CLL cells for the Bcl-2 antagonist ABT737 is decreased by a factor of 1000, if the CLL cells are co-cultivated with fibroblasts expressing CD40L (Vogler et al., 2009).

3.3 Stroma cell-derived soluble factors and cell surface interactions

The soluble factors produced by bone marrow stromal cells include the chemokine CXCL12, which despite its original designation as stroma-cell derived factor is shown in Fig. 2 as a micro-environmental factor occurring in peripheral blood, owing to its alternate origin from nurse-like cells. Stimulation by purified recombinant CXCL12 induced the raf-dependent mitogen activated protein (MAP) kinase cascades in CLL cells, which augmented their survival and was targeted by the raf inhibitor sorafenib (Messmer et al., 2011).

In vivo, inhibition of apoptosis may occur preferentially in pseudofollicles containing CLL and accessory cells, due to cell contact and mutual paracrine and autocrine stimulation. *In vitro*, co-culture with bone-marrow-derived stromal cells, e.g. the cell line HS-5, may provide stimuli for long-term survival of CLL cells. In co-cultures of primary CLL cells with HS-5 cells, various chemokines attracting T lymphocytes, most prominently CCL4 and CCL3, were detected, which are not produced by HS-5 control cultures (Seiffert et al., 2010). The proteins found in the supernatant of HS-5 co-cultures included factors, which are commonly secreted by monocytes, e.g. soluble CD14. Among the soluble factors provided to CLL cells by co-culture with the bone marrow stromal cell line HS5, vascular endothelial growth factor (VEGF) is partly responsible for the increase in viability of co-cultivated CLL cells (Gehrke et al., 2011).

3.4 Oligonucleotides containing CpG dinucleotides

Toll like receptor 9 (TLR9) has been identified as a part of the innate immune response recognizing unmethylated foreign DNA that can be mimicked by phosphothioate oligodeoxynucleotides containing CpG dinucleotides (CpG-ODN) (Krieg, 2006). Survival and proliferation of CLL cells can be considerably enhanced by class B CpG-ODN, e.g. DSPN-30 (Decker et al., 2000). Like CD40 ligation, activation of CLL cells through TLR9

occupation by CpG-ODN was exploited as a mitogenic signal in order to obtain metaphase chromosomes for cytogenetic analysis by fluorescent in situ hybridization (Mayr et al., 2006). The importance of activated CLL cells in drug assessment is demonstrated by the example of mTOR inhibitors. In untreated CLL cells rapamycin showed an IC₅₀ of 10 µM for apoptosis induction (Aleskog et al., 2008). In contrast less than one thousandth of this concentration of RAD001 or 10 nM rapamycin was sufficient for complete inhibition of the cell proliferation induced by CpG-ODN (Decker et al., 2003).

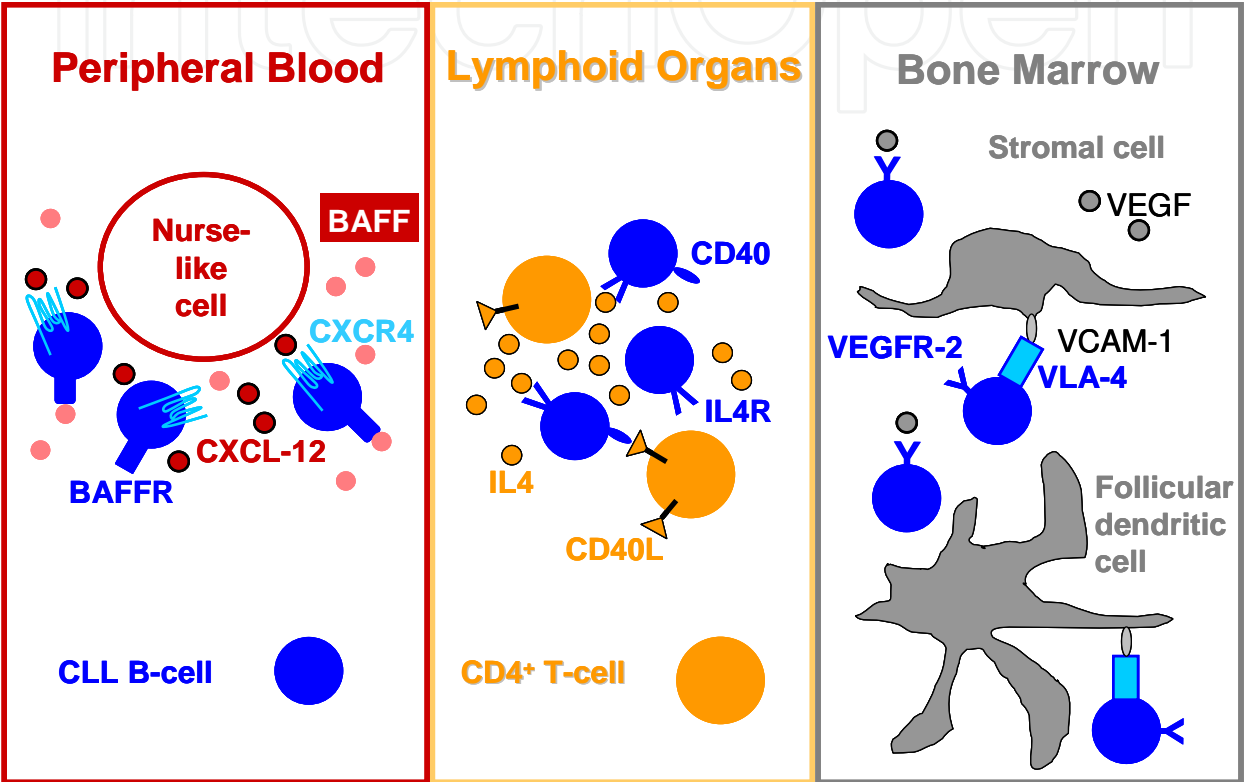


Fig. 2. Microenvironment interactions of CLL cells. The survival of CLL lymphocytes can be prolonged by contact with different accessory cells in the peripheral blood, lymphoid system and bone marrow. Some prominent interactions of CLL cells with soluble factors and cell surface molecules supplied by accessory cells are shown. On the surface of CLL cells receptors for various soluble factors are expressed, e.g. for the chemokine CXCL12, formerly designated as stroma-derived factor 1 (SDF-1), for the cytokine interleukin-4 (IL-4), and for vascular endothelial growth factor (VEGF). CD40 or the integrin VAL-4 on the surface of CLL cells interact with CD40 ligand (CD40L) on T cells or VCAM-1 on follicular dendritic cells and other stromal cells, respectively. Adapted from Munk Pedersen *et al.*, 2004.

In addition to activating CLL cells CpG-ODN were found to increase surface expression levels of co-stimulatory molecules including CD20 (Jahrsdorfer et al., 2001). Consequently, the same CpG-ODN DSPN-30 that is commonly used for activating CLL cells, increased CD20 expression on freshly isolated CLL cells, which in turn led to higher B cell depletion by the type II CD20 antibody GA101 (Patz et al., 2011).

3.5 B cell depletion from whole blood samples

Monoclonal antibodies (MABs) induce direct cell death (DCD) of tumor cells via signal transduction and additional Fc-mediated cytotoxic effects, namely antibody-dependent cell-mediated cytotoxicity (ADCC) and complement-dependent cytotoxicity (CDC) (Fig. 3). In order to include ADCC and CDC in measurements of overall MAB effects, the extent of tumor cell depletion by MABs from individual blood samples can be determined by multi-color flow cytometry comparing treated and untreated whole blood cultures.

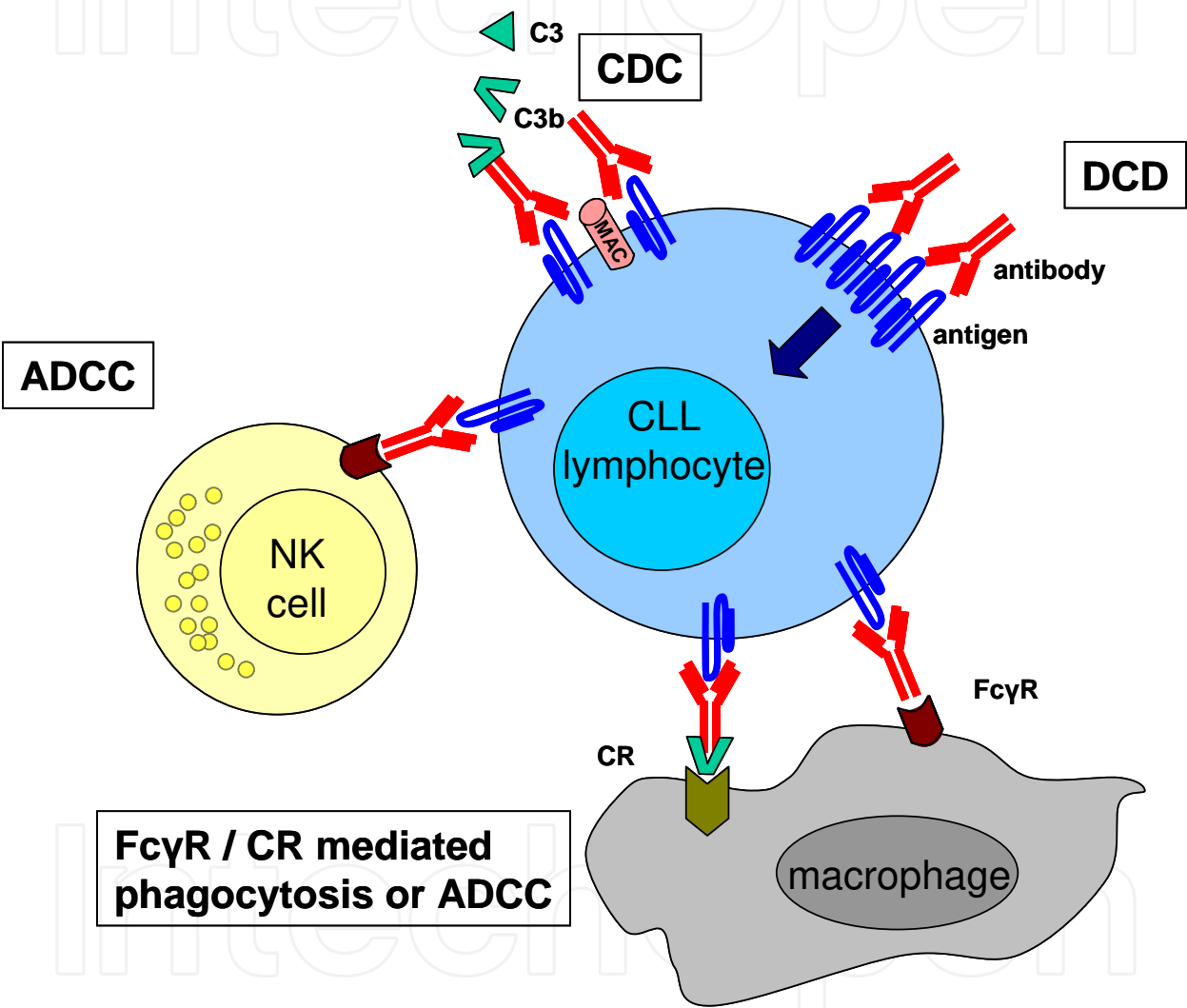


Fig. 3. Killing mechanisms of monoclonal antibodies and their assessment. Apart from direct cell death (DCD) induction in tumor cells, monoclonal antibodies exert their action via Fc-mediated functions, namely complement-mediated cytotoxicity (CDC) and antibody-dependent cell-mediated cytotoxicity (ADCC) and phagocytosis. Due to the importance of CD20 antibodies in CLL therapy, a structure crossing the cell membrane four times is shown as the surface antigen on CLL cells. This overall structural organization is shared by another emerging target for immunotherapy, the tetraspanin CD37. Ways to examine the above mechanisms on isolated CLL cells and whole blood samples are compiled in Table 2 and described in the text. Adapted from Olszewski & Grossbard, 2004 and Jaglowski et al., 2010.

4. Pre-clinical assessment of kinase inhibitors

Prototypic targeted therapy by the tyrosine kinase inhibitor imatinib was developed for Bcr-Abl positive leukemias, in which deregulated Abl activity is a predominant driving force (Druker et al., 2001). In contrast, the pathogenesis of CLL appears to be multi-factorial. The second generation of Abl inhibitors for treatment of imatinib-resistant Bcr-Abl positive leukemias achieves greater efficacy against mutant forms of the Abl kinase (Weisberg et al., 2007). Some of these inhibitors, e.g. dasatinib (Shah et al., 2004) and bosutinib (Puttini et al., 2006) are dual-specific and target Abl and additionally Src kinases. Since members of the latter tyrosine kinase family, e.g. Lyn (Contri et al., 2005) and Lck (Majolini et al., 1999) have been suggested to be involved in CLL pathogenesis, we conducted an assessment of dasatinib on CLL cells (Veldurthy et al., 2008). This pre-clinical investigation indicated an influence of Src kinase inhibition on the cellular survival of CLL cells with preference for the subgroups with unmutated immunoglobulin heavy chain genes or with high ZAP70 expression and thus indicated patient groups that might profit most from Src kinase inhibition. Since the fate of CLL B lymphocytes critically depends on BCR signaling (Stevenson & Caligaris-Cappio, 2004), inhibition by kinase inhibitors of survival pathways emanating from the BCR or from micro-environmental stimuli (Burger et al., 2009) represents a promising strategy for treating CLL (Gandhi, 2010).

Drug assessment on primary CLL cells serves as preparation for clinical trials and to some degree enables comparison of the efficacies of different agents and the prediction of the response of molecularly defined subgroups of CLL patients. The clearly higher dasatinib sensitivity of CLL samples with unmutated IgV_H genes as compared to mutated ones is an example for this type of correlation (Veldurthy et al., 2008). Signaling analysis revealed that treatment of primary CLL cells with dasatinib drastically reduces the level of activated SFK in CLL cells, but inhibits downstream BCR signaling pathways and induces apoptosis more strongly in the patient subgroup with aggressive disease. The extent of dasatinib-induced apoptosis in CLL cells corresponds to the concomitant decrease in the phosphorylation of the direct SFK substrates Syk and phospholipase C- γ (Song et al., 2010). Signaling analysis during SFK inhibition thus contributed to the rationale for pre-clinical assessment of Syk inhibitors on CLL cells (Baudot et al., 2009; Buchner et al., 2009). For another second generation dual Abl/Src inhibitor, bosutinib, inhibition of the receptor tyrosine kinase Axl was found to be partially responsible for its apoptosis induction in CLL cells (Ghosh et al., 2010).

Inhibition of Abl does not reduce viability of CLL cells on its own, but can sensitize CLL cells for chemotherapeutic agents, e.g. chlorambucil, by interfering with DNA repair (Aloyz et al., 2004). Inhibitors of the delta isoform of the catalytic p110 subunit of phosphatidylinositol-3-kinase (PI3K) show moderate efficiencies on primary CLL lymphocytes without activation that contrast the promising effects in clinical trials. The observed pre-clinical efficiency of the PI3K-delta inhibitor CAL-101 is not abrogated by micro-environmental stimulation and other cell types, while other cell types, e.g. natural killer cells are not influenced by CAL-101 (Herman et al., 2010).

5. Pre-clinical assessment of monoclonal antibodies

Apart from small molecules, monoclonal antibodies constitute another group of targeted therapeutics for the treatment of CLL. This group includes the first biological anti-tumor

agent, namely the CD20 antibody rituximab. As a common cell surface antigen of all B cells except stem or plasma cells, CD20 has become a very effective antibody target for the treatment of B cell malignancies (Molina, 2008) including CLL despite variable surface expression on CLL cells. Together with the monoclonal anti-CD52 antibody alemtuzumab, rituximab thus may be counted among the most efficient targeted treatment options for CLL achieved so far. In a recent phase III trial inclusion of rituximab was shown to substantially improve the established fludarabine / cyclophosphamide chemotherapy regimen (Hallek et al., 2010).

Apart from DCD induction in tumor cells, monoclonal antibodies exert their action via Fc-mediated functions, namely CDC, ADCC and phagocytosis (Jagłowski et al., 2010; Olszewski & Grossbard, 2004) (Fig. 3). Therefore an assessment of antibody effects on CLL cells *ex vivo* can either be performed on freshly isolated CLL cells in separate dedicated assays for each mentioned mechanisms, or in a comprehensive assay from whole (Table 2). For assessing antibody effects on isolated CLL cells, the same procedures as for other anti-CLL agents can only be applied for the determination of DCD. For instance DCD induction by rituximab in freshly isolated CLL cells was assessed according to viable cell counts, metabolic activity and phosphatidylserine exposure and was found variable among individual samples and considerably smaller than in lymphoma cell lines (Patz et al., 2011; Stanglmaier et al., 2004). Since DCD induction in primary CLL cells may constitute only a minor fraction of overall B cell depletion as in the case of rituximab (Voso et al., 2002), it is indispensable to additionally assess Fc-mediated mechanisms. For performing ADCC assays on isolated CLL lymphocytes, effector cells need to be externally added, e.g. autologous or heterologous peripheral blood mononuclear cells or a natural killer cell line (Weitzman et al., 2009). Similarly, CDC can be assessed by monitoring changes in the membrane integrity of CLL cells after incubation in medium containing high concentrations of complete as compared to heat-inactivated serum (Golay et al., 2000; Patz et al., 2011).

Mechanism	Isolated CLL cells	Whole blood
DCD	Cytotoxicity or viability assays	Requires distinction of CLL lymphocytes from other cell populations
CDC	Comparison of effects with complete or heat-inactivated serum	Complement inhibition by cobra venom factor
ADCC	Externally added effector cells	Blocking antibodies for Fc receptors

Table 2. Determination and dissection of different mechanisms of antibody-induced cytotoxicity.

As an alternative to these separate assays, we applied a B cell depletion assay from whole blood encompassing Fc-mediated antibody-induced cytotoxicity. This assay is based on the enumeration of B lymphocytes in differentially treated whole blood samples after staining the general, B- and T- lymphocyte antigens CD45, CD19 and CD3 using three color flow cytometry and commercially available fluorescently labeled immunoreagents. B cell depletion can be calculated from the B cell counts in antibody-treated versus untreated control samples. B/T cell ratios with the T cell population as an internal standard can be

used for this calculation, if T cell counts are proven to be unaffected by the antibody treatment. Alternatively absolute B cell counts can be determined using externally added fluorescent counting beads. In part the contributions of DCD, CDC and ADCC to the observed B cell depletion from whole blood samples can be dissected. Thus, we were able to show a substantial contribution of ADCC to the B cell depletion by the novel type II CD20 antibody GA101 (Moessner et al., 2010) by blocking the interaction of FcγIIIa on NK cells and macrophages and the Fc exposed on antibody-coated target cells by incubation with anti-CD16 antibody in whole blood from healthy donors but not CLL patients (Patz et al., 2011). In summary, flow cytometric determination of B cell depletion from whole blood has the potential of comparing antibody effects on individual CLL samples and of predicting clinical responses.

Resistance mechanisms to anti-CLL antibodies have been unraveled by pre-clinical models and some of the influences interfering with antibody efficacy may be recapitulated in the the present B cell depletion from whole blood samples (*Table 3*) (Reslan et al., 2009). Thus the shape of the dose response curves of GA101 or mAb37.1 observed in the B cell depletion assay (Krause et al., 2011; Patz et al., 2011) is of saturation type and suggests an influence on antibody effects of individually different levels of endogenous human IgG in the assay matrix (Preithner et al., 2006). Similarly, varying ratios of effector to target cells, the Phe158Val polymorphism of FcγIIIa (Cartron et al., 2002) as well as complement depletion (Kennedy et al., 2004) will influence B cell depletion from whole blood samples. Thus, the assay described here has the advantage of reflecting both, the efficiency of antibody-induced B cell depletion and the potential to supply host-dependent immune functions and thus should be able to predict at the individual level the clinical efficiency of therapeutics assayed *in vitro*.

Influence	References
Antigen density on target cells	Golay et al., 2001, Patz et al., 2011
Complement depletion	Kennedy et al., 2004
Complement inhibitors CD59 and CD55	Golay et al., 2001
Ratio of effector to target cells	
Plasma levels of IgG	Preithner et al., 2006
Fc receptor polymorphisms	Cartron et al., 2002

Table 3. Influences on antibody efficacy in a whole blood matrix.

6. Conclusions

Due to the dependence of CLL lymphocytes on their micro-environment, the predictive value of simple cytotoxicity assays on freshly isolated CLL cells can be enhanced by activating CLL cells using procedures that mimic certain micro-environmental stimuli. In the case of monoclonal antibodies, effector cells and complement system need to be included in order to comprise indirect antibody-mediated mechanisms. For arriving at valid predictions, results of the individualized *in vitro* sensitivity testing should be linked to mechanistic and biochemical target validation studies, ideally involving genetically defined systems. Remarkably three major topics addressed in this chapter, namely the importance of accompanying signaling analysis, consideration of the micro-environment of CLL cells and

combination with chemotherapeutic agents conceptually strongly overlap with general targeting strategies in contemporary pre-clinical anticancer drug discovery (Caponigro & Sellers, 2011).

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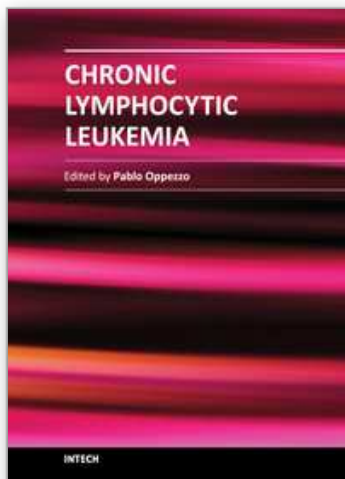
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B-cell chronic lymphocytic leukemia (CLL) is considered a single disease with extremely variable course, and survival rates ranging from months to decades. It is clear that clinical heterogeneity reflects biologic diversity with at least two major subtypes in terms of cellular proliferation, clinical aggressiveness and prognosis. As CLL progresses, abnormal hematopoiesis results in pancytopenia and decreased immunoglobulin production, followed by nonspecific symptoms such as fatigue or malaise. A cure is usually not possible, and delayed treatment (until symptoms develop) is aimed at lengthening life and decreasing symptoms. Researchers are playing a lead role in investigating CLL's cause and the role of genetics in the pathogenesis of this disorder. Research programs are dedicated towards understanding the basic mechanisms underlying CLL with the hope of improving treatment options.

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