Dose and drug changes in chronic lymphocytic leukemia cell response in vitro: A comparison of standard therapy regimens with two novel cyclin-dependent kinase inhibitors

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Abstract. Chronic lymphocytic leukemia (CLL) treatment is improving; however, some patients do not respond to therapy. Due to the high heterogeneity in disease development, there is an urgent need for personalization of therapy. In the present study, the response of leukemic mononuclear cells to anticancer drugs used for CLL treatment (cladribine + mafosfamide; CM or CM combined with rituximab; RCM) was compared with the response to new cyclin-dependent kinase (CDK) inhibitors: BP14 and BP30. Viable apoptotic and necrotic cells were quantified by flow cytometry using propidium iodide and Yo-Pro stains. CDK inhibitors were studied in several doses to determine the reduction of necrosis and simultaneous increase of apoptosis in leukemic cell incubations with anticancer agents. The distinct cell response to applied doses/anticancer agents was observed. Results obtained in the current manuscript confirmed that modulation of doses is important. This was particularly indicated in results obtained at 24 h of cells incubation with anticancer agent. While an important time for analysis of anticancer response efficacy (monitoring of apoptosis induction potential) seems to be 48 h of cells exposition to anticancer agents. High variability in response to the drugs revealed that both the nature and the dose of the anticancer agents could be important in the final effect of the therapy. The present findings support the thesis that personalized medicine, before drug administration in the clinic, could be important to avoid the application of ineffective therapy.

Introduction

Although the growing range of treatment options for chronic lymphocytic leukemia and for other lymphoproliferative neoplasms has improved patient survival (1-4), these diseases remain incurable. In addition, there are patients who do not respond to the applied therapy. The serious problems associated with the diagnostic procedure and the design of suitable treatments seem to be linked to the coexistence in patient peripheral blood of quiescent and cycling cells population; this constitutes a special challenge in predicting an effective approach for treating CLL patients (5,6).

Differences in cell signaling trafficking, as well as in the expression of factors involved in apoptosis or microenvironmental factors, might contribute to differences (between patients) in the cell response to anticancer agents between patients. In addition, it is well accepted that diversity in the accumulation of genetic aberrations and epigenetic modifications could also account for heterogeneity in the clinical course of CLL (7-9) and the response to therapy (10,11). Moreover, another factor that could imply in the course of CLL as well as response to therapy is the expression of miR-155. This microRNA is associated with the progression of CLL and weak response to therapy (12). The presence of several factors important for disease development reveals the necessity for the use of personalized medicine, by testing the potential reaction of the patient’s cells to anticancer drugs before treatment, to avoid administration of an ineffective regimen (8,13-16). Therefore, it is very important to search for new anticancer agents with the potential to induce apoptosis in CLL cells (17-20).

Cyclin-dependent kinases (CDK) are fundamental factors involved in the regulation of the cell-cycle, transcription
and apoptosis. Their frequent deregulation in cancers provides novel targets for pharmacological intervention in oncology (21). Various small-molecule CDK inhibitors have been developed, including CDK4/CDK6-specific palbociclib and ribociclib, recently FDA-approved for multiple myeloma and breast cancer, respectively (22,23). Besides the cell cycle, CDKs play critical roles also in a non-proliferative CLL and in cell lines where the CDK inhibitor flavopiridol has been designed as an orphan drug for CLL (24). Flavopiridol however suffers several side effects, such as significant toxicity including high rates of major tumor lysis syndrome, cytokine release syndrome and secretory diarrhea (24). Other CDK inhibitors are therefore studied as new drugs for CLL, such as roscovitine, dinaciclib or SNS-032 (25). These compounds target multiple CDKs, including CDK1, CDK2, CDK5, CDK7 and CDK9, and trigger cytotoxic effects through interruption of the transcription of key antiapoptotic genes responsible for sustenance of the leukemia cell, such as MCL-1 (21,24).

We have recently modified roscovitine to increase its potency and the optimization yielded new 2-substituted-6-biarylmethylamino-9-cyclopentylpurine derivatives BP14 and BP30, which display selective and potent inhibition of CDKs 1, 2, 7 and 9 with low nanomolar IC50 values (26). Both BP14 and BP30 exhibit strong cytotoxicity in human cancer cell lines that correlate with robust CDK1 and CDK2 inhibition and caspase activation. BP14 has demonstrated efficacy against xenografted human liver carcinomas, effectively repressing tumor growth (27). In addition, BP14 potently inhibited transcriptional regulator CDK9 and downregulated anti-apoptotic protein MCL-1 (27,28), key mediator of CLL-cell survival.

The aim of the current work was to observe the importance of drug doses for anticancer response in leukemic cells. For this purpose we have compared the apoptosis induction potential of new CDK inhibitors as potential drugs for CLL and compare them with standard treatments. The present study compares the cytotoxicity (cell viability, apoptosis or necrosis level) of novel roscovitine derivatives BP14 and BP30 and anticancer drugs used in hematological clinics for treating CLL (CM, cladribine + mafosfamide; RCM, rituximab + cladribine + mafosfamide). The concentration of the drugs used for cell incubations were as follows: Cladribine 50 ng/ml (139 nM), mafosfamide 1 µg/ml (2.4 µM), rituximab 20 µg/ml (170 nM). Cells incubated only with medium or with appropriately diluted DMSO served as a control.

**Patients.** Blood samples were obtained from five CLL patients: One female patient and four male (cooperation with the Department of Hematology, Medical University of Lodz, Poland; Head: Prof. T. Robak). The median age was 65 years (54-81). Leukocytosis ranged from 60 to 700x10⁹/µl (Table I). Samples were collected on EDTA to sterile, disposable tubes. Cell marker studies were performed to confirm B-cell origin and monoclonal proliferation. All B lymphocytes were CD5, CD19 and CD23 positive.

**Ethics statement.** The study was approved by the Local Ethics Committee of the University of Lodz (20/KBBM-UL/2015 and 5/KBBMUL/2017).

**Isolation of mononuclear cells and cell cultures with anticancer agents.** Mononuclear cells were isolated from CLL samples by centrifugation on a Histopaque 1077 gradient (900 rpm, 19 min, 24°C). The obtained cells were resuspended in RPMI-1640 medium with 10% fetal bovine serum, 2 mM L-glutamine, 100 U/ml penicillin and 100 µg/ml streptomycin. The cell culture density was 3x10⁶/ml. In the next step, the cells were incubated with drugs/novel anticancer compounds. After 24 and 48 h of cell incubations with anticancer agents cell viability was tested.

**Flow cytometry analysis.** Cell viability, the rate of apoptosis and necrosis were estimated using a Vybrant Apoptosis Assay #4 kit after 0, 24 and 48 h, as in previous studies (14,15). The early apoptotic cells were marked with YO-PRO fluorescent dye, and late apoptotic and necrotic cells with propidium iodide. For flow cytometry analysis, 1x10⁶/ml cells were used, centrifuged (5,000 rpm, 5 min, 24°C), then resuspended in phosphate buffer saline (PBS) containing fluorescent dyes. Samples were incubated in the dark for 20 min and then analyzed.

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**Materials and methods**

**Drugs and treatments.** The studied compounds BP14 (N⁴-(trans-4-aminocyclohexyl)-9-cyclopentyl-N⁶-[6-(2-furanyl)-3-piridinyl]methyl)-9H-purine-2,6-diamine) and BP30 (N⁴-(trans-4-aminocyclohexyl)-9-cyclopentyl-N⁶-[6-(2-thienyl)-3-piridinyl]methyl)-9H-purine-2,6-diamine) were synthesized and characterized as described in our previous report (26). The identities of the compounds were confirmed by mass spectrometry, NMR and elemental analysis. The purity of the compounds was at least 99.5%. 10 mM stock solutions of the compounds were prepared in DMSO. The final DMSO concentrations did not exceed 0.1%, a concentration verified not to affect respective experimental parameters.

Cladribine (Biodyrin) and rituximab were obtained from the Institute of Biotechnology and Antibiotics (Warsaw, Poland) and from Roche (Basel, Switzerland), respectively.

Mafosfamide was purchased from Niomech IIT GmbH (Bielefeld, Germany). These three drugs were used in the following combinations: CM (cladribine + mafosfamide), RCM (rituximab + cladribine + mafosfamide). The concentration of the drugs used for cell incubations were as follows: Cladribine 50 ng/ml (139 nM), mafosfamide 1 µg/ml (2.4 µM), rituximab 20 µg/ml (170 nM). Cells incubated only with medium or with appropriately diluted DMSO served as a control.

**Table I. Characteristics of patients.**

| Patient number | Sex | Age, years | Leukocytosis (x10⁹/µl) | Rai stage |
|----------------|-----|------------|------------------------|-----------|
| 12             | M   | 81         | 470                    | 4         |
| 14             | M   | 57         | 700                    | 3         |
| 15             | M   | 65         | 60                     | 4         |
| 16             | F   | 66         | 153                    | 4         |
| 17             | M   | 54         | 262.3                  | 2         |

M, male; F, female.
Figure 1. Dose-response effect of BP14 and BP30 on the viability of CLL cells following 48-h incubations. Data are presented as the mean ± standard error of the mean (n=6).

Figure 2. (A) Determination of (a) cell viability, (b) level of apoptosis or (c) necrosis. CLL cells (Patient no. 15) were incubated for 48 h without drugs (Co. or Co. DMSO) or with CM, RCM or CDK inhibitors (A) BP14 and (B) BP 30. Co., untreated control; Co. DMSO, control with dimethyl sulfoxide; CM, cladribine combined with mafosfamide; RCM, CM combined with rituximab; CDK, cyclin-dependent kinase.
**Statistical analysis.** Statistical analysis were performed by Kruskal Wallis test with Dunn's multiple comparison test. The statistical differences between the control sample (Co.) and samples with the addition of the anticancer agent was analyzed, respectively. Statistical analysis were obtained for N=5; *P<0.05, **P<0.001, ***P<0.0001.

**Results**

Peripheral mononuclear cells from the group of 6 patients with CLL were exposed to varying concentrations of novel CDK inhibitors (BP14 and BP30). The representative results for patients (3 presented as single cases and 2 combined figures with statistics) were included in current manuscript, showing distinct patients response to anticancer agents. The effect of CDK inhibitors on CLL cells were analysed for 6 CLL patients and compared with standard chemotherapeutic regimens (CM, cladribine + mafosfamide; RCM, rituximab + cladribine + mafosfamide), as described in our previous studies with more than 50 patients (14,15). The CLL cells isolated from peripheral blood of patients were incubated with anticancer agents for 24 and 48 h, harvested and then analyzed for viability, apoptosis and necrosis by flow cytometry. Both studied CDK inhibitors decreased the viability of the CLL cells in a dose-dependent manner (Fig. 1). At the concentration of 25 nM, the median reduction (IC_{50}) for all patients in cell viability for both CDK inhibitors was similar, 39.1 and 31.5% for BP14 and BP30, respectively (Fig. 1).

Next we have analysed CLL viability and the level of necrosis upon treatment with anticancer agents. The cells of patient no. 15 showed increasing level of necrosis after 24 h exposition to all used anticancer agents but with distinct extend. The cells obtained from patient no. 15 were very sensitive toward all anticancer agents (Figs. 2 and 3). The strong reduction in cell viability was observed for all used types of treatment, but the most spectacular effect was noticed when cell were incubated for 48 h with CM and RCM. Moreover, the increase in apoptosis level (24 h incubation) and the high value of necrosis (48 h) was a consequence of dynamics in apoptosis realization. The high level of necrosis after 48 h of cell exposition to anticancer agents could be a compilation of necrosis as well as advanced apoptosis. Therefore, for a
proper modulation of doses value a special importance should be given to results obtained at 24 h of cells incubation with anticancer agents. While for monitoring of apoptosis induction potential an important time for analysis of anticancer response efficacy seems to be 48 h of cells exposition to anticancer agents (Fig. 2).

The results obtained for representative patient no. 16 show the potential resistance of patient's cells to drug combinations used in hematological clinics for CLL treatment. For this point only BP14/BP30 reflect to be active in apoptosis induction. Moreover, BP30 and BP14 in all used doses induced very fast apoptosis. The cells of patient no. 16 (Figs. 4 and 5) demonstrated resistance to CM and RCM, as well as sensitivity to each analyzed dose of CDK inhibitors (BP14 and BP30). Interestingly, the level of apoptosis in patients no. 16's cells for each dose of BP14 and BP30 was relatively constant, but increasing levels of necrosis/late apoptosis were observed.

The comparative analysis of patient no. 17 reflects the opposite reaction to patient no. 16. The cells of patient no. 17 (Fig. 6) were sensitive to both standard CLL treatments (CM and RCM), but displayed resistance to CDK inhibitors, which was clear in all assays, including cell viability, apoptosis and necrosis.

Finally, the comparative analysis of the results of the 24 h (A) and 48 h (B) CLL cell incubations with anticancer agents for five patients (Table I) are presented on Fig. 7 (for BP14) and Fig. 8 (for BP30), with a statistical analysis performed by the Kruskal-Wallis test. It must be stated that these combined results are difficult to compare with those of each single patient. It is important to note that the results for patients 15,
16 or 17 presented here indicate that the median results presented in scientific papers usually provide incorrect recommendations (13-15) concerning patient resistance/sensitivity to anticancer agents, and these would not be effective in treating patient no. 16.

**Discussion**

During CLL development, long-lived B lymphocytes accumulate in peripheral blood and in other lymphatic organs of patients with the advanced form of the disease (29). This accumulation occurs due to suppression of apoptosis and usually happens in the resting form of the disease (9,29). Since the disease could transform into an active form, the coexistence of quiescent and cycling cell populations complicate the diagnostics and prediction of therapy efficacy (5,6). Therefore, an analysis of the chemosensitivity profile of patient cells prior to therapy followed by the use of personalized medicine based on apoptosis induction would markedly reduce the ineffective treatment (8,13-15).

The current study compares cytometric analyses of cell viability, the levels of apoptosis and necrosis in *ex vivo* CLL cells isolated from five patients to determine whether the response has some connection with the type of used anticancer agents. The special attention was given for observation of the role of anticancer agent doses on cell response.

Our findings reveal significant differences in the response to anticancer agents among leukemic patients, as noticed in previous studies (8,14,15). Moreover, our results show that the comparison of cell viability, apoptosis rate, and the level of necrosis underline the importance of proper anticancer agent dose selection to avoid the occurrence of a high necrosis level that could transform *in vivo* into inflammation. To eliminate potential resistance to therapy, the therapeutic approach should be chosen before drug administration in the clinic using personalized therapy tests (14,15). Such techniques have good value for predicting CLL cell sensitivity to anticancer drugs. This prediction is of importance as it allows for the optimal course of therapy to be chosen for the patient, thus avoiding ineffective anticancer administration and minimizing the level

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Figure 5. Graph plots of cell viability (Q3), apoptotic (Q4), necrotic and late apoptotic cells (Q2), debris (Q1). CLL cells (Patient no. 16) were incubated for 48 h (Co. or Co. DMSO) or with anticancer drugs (CM or RCM) or CDK inhibitors in several doses BP14.25, BP14.35, BP14.50, and BP 30.15, BP 30.15, BP 30.25. Co., untreated control; Co. DMSO, control with dimethyl sulfoxide; CM, cladribine combined with mafosfamide; RCM, CM combined with rituximab; CDK, cyclin-dependent kinase.
of necrosis that could influence the activity of the drug in vivo. In order to expand the range of possible therapeutic options for leukemia treatment, there is an urgent need to identify new compounds with anticancer properties.

Recent developments in the field of drug discovery include the identification of potent CDK inhibitors. Currently, it is not clear whether highly CDK specific inhibitors, such as palbociclib, or pan-CDK specific inhibitors, such as flavopiridol or dinaciclib, are more suitable for cancer therapy. A number of studies have found them to have promising efficacy when used in combination with standard anticancer drugs or with other molecularly targeted agents (21). The present study examines the anti-CLL potential of two novel CDK inhibitors that have already displayed high

Figure 6. (A) Determination of (a) cell viability, (b) level of apoptosis or (c) necrosis. CLL cells (Patient no. 17) were incubated for 48 h without drugs (Co. or Co. DMSO) or with CM, RCM or CDK inhibitors (A) BP14 and (B) BP 30. Co., untreated control; Co. DMSO, control with dimethyl sulfoxide; CM, cladribine combined with mafosfamide; RCM, CM combined with rituximab; CDK, cyclin-dependent kinase.
Our results further support previous findings that CDK inhibitors induce apoptosis usually much faster than standard treatments used for CLL patients (24). However, a high variability noticed in disease development (29) translates on individual patient sensitivity to anticancer agents. Our recent findings reveal that the dose of the anticancer drug strongly influences the response to treatment. While the correct dose of the active drug will induce a high level of apoptosis and should lead to natural cell elimination by programmed cell death. The high dose usually induces necrosis, which could induce inflammation. In addition, our findings demonstrate that drug dose could play an important role in the modulation of leukemic cell response to the anticancer agent(s). It has also been found that 5-HT7 receptors play a role in the induction of inflammation by release of sirtuin, a nicotinamide adenine dinucleotide-dependent deacetylase that could influence gene expression (35), as well as in the level of serotonin regulation (36). Therefore, it is possible that an excessive dose of a drug induces a high level of necrosis, and that could be transferred into inflammation. It might also have an impact on the diversities in cell signaling. Interestingly, it has been suggested that inflammation could cause conditions directed

Figure 7. (A) Determination of (a) cell viability, (b) level of apoptosis or (c) necrosis. CLL cells were incubated for 48 h without anticancer agents (Co. or Co. DMSO) or with anticancer drugs (CM or RCM) or purine analog BP14. N=5; data are presented as the mean ± standard deviation (n=5). *P<0.05, **P<0.001 and ***P<0.0001. Statistical analysis were performed using the Kruskal Wallis test. Results were obtained after (A) 24 and (B) 48 h of leukemic cell exposure to anticancer agents. Co., untreated control; Co. DMSO, control with dimethyl sulfoxide; CM, cladribine combined with mafosfamide; RCM, CM combined with rituximab.

anticancer activity (26). Our results further support previous findings that CDK inhibitors induce apoptosis usually much faster than standard treatments used for CLL patients (24). However, a high variability noticed in disease development (29) translates on individual patient sensitivity to anticancer agents. Our recent findings reveal that the dose of the anticancer drug strongly influences the response to treatment. While the correct dose of the active drug will induce a high level of apoptosis and should lead to natural cell elimination by programmed cell death. The high dose usually induces necrosis, which could induce inflammation. In addition, our findings demonstrate that drug dose could play an important role in the modulation of leukemic cell response to the anticancer agent(s). It has also been found that 5-HT7 receptors play a role in the induction of inflammation by release of sirtuin, a nicotinamide adenine dinucleotide-dependent deacetylase that could influence gene expression (35), as well as in the level of serotonin regulation (36). Therefore, it is possible that an excessive dose of a drug induces a high level of necrosis, and that could be transferred into inflammation. It might also have an impact on the diversities in cell signaling. Interestingly, it has been suggested that inflammation could cause conditions directed
to osteoporosis (37), which can suggest that factors involved in the development of inflammation could also disturb other metabolic pathways, leading to the possible occurrence of a range of other metabolic alterations that could lead to osteoporosis. Interestingly, the aberrant products observed in cancer cells involved in the Krebs cycle could promote cancer progression (38).

The results of our studies confirm the need for personalized therapy for CLL, as such an individual approach would greatly avoid the chance of ineffective treatment being used. Our findings also highlight the importance of the application of the correct dose for treatment, and demonstrate that the drugs could induce apoptosis at different rates, which can be also monitored in vitro by incubating cells with anticancer agents.

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The authors declare that they have no competing interests.

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Availability of data and materials

The datasets used and/or analyzed during the current study are available from the corresponding author on reasonable request.

Authors’ contributions

MK and AS performed the experiments. JZB and PR determined the clinical material characteristics. MM performed and organised the blood transfers in the clinic. TG and VK synthesised the new purine analogs. MR wrote the manuscript, and conceived and designed the experiments. All authors read and approved the final manuscript.

Ethics approval and consent to participate

The present study was approved by the Local Ethics Committee from the University of Lodz (nos. 20/KBBM-UŁ/2015 and 5/KBBMUL/2017). The requirement for written, informed consent was waived.

Patient consent for publication

Not applicable.

Competing interests

The authors declare that they have no competing interests.

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