Idelalisib induces G1 arrest and apoptosis in chronic myeloid leukemia K562 cells

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Abstract. Increasing resistance of imatinib, a BCR-ABL tyrosine kinase inhibitor, hinders its use in the therapy of chronic myeloid leukemia (CML). The PI3K pathway is known to be closely involved in BCR-ABL transformation and the tumorigenesis of CML, suggesting that PI3K may be a potential target for CML therapy. Idelalisib, a specific inhibitor of PI3K p1108, has been approved for the treatment of chronic lymphocytic leukemia (CLL). However, the antileukemia effect of idelalisib on CML remains unknown. In the present study, the antileukemia activity of idelalisib alone or in combination with imatinib was investigated by use of K562 cells. Idelalisib inhibited K562 proliferation in a dose-dependent manner. G1 arrest was induced, in which upregulation of p27 and p21, as well as downregulation of cyclin D1 and p-pRb, may be involved. Furthermore, idelalisib induced apoptosis in the K562 cells, with increased expression of pro-apoptotic molecules such as Bad and Bax, cleavage of caspase-9, -8 and -3, and PARP, in contrast to downregulation of anti-apoptotic protein Bcl-2. Combination of idelalisib with imatinib led to a synergistic antiproliferative effect on K562 cells, together with enhanced activity of G1 arrest and apoptosis induction. In conclusion, idelalisib exhibited in vitro antitumor activity on CML K562 cells alone or in combination with imatinib, suggesting potential application in CML therapy.

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Introduction

Chronic myeloid leukemia (CML), a clonal myeloproliferative disorder, is characterized by the Philadelphia (Ph) chromosome which originates from the t(9;22)(q34;q11) reciprocal translocation and leads to the BCR-ABL chimeric oncoprotein (1-3). This BCR-ABL oncoprotein bears constitutive tyrosine kinase activity and therefore promotes uncontrolled growth and proliferation of leukemia cells (4). Development of inhibitors targeting BCR-ABL has been generally successful in the past 15 years, with highly decreased mortality for CML patients. However, resistance to these drugs has been increasingly reported. CML remains an incurable disease (5,6). Therefore, novel drugs with targets other than BCR-ABL are expected.

Idelalisib, also named CAL101, was approved by the US Food and Drug Administration in July, 2014. It is a first-in-class oral PI3K inhibitor that has shown substantial and sustained antileukemia efficacy in patients with relapsed/refractory chronic lymphocytic leukemia (CLL) (7,8). Compared with other chemotherapy regimens, idelalisib showed advantages of long-term efficacy and reduced toxicity (9).

It has been reported that PI3K signaling contributes to BCR-ABL transformation and is essential for *in vivo* leukemogenesis of CML (10). Furthermore, the PI3K p110 δ isoform is preferentially expressed in hematopoietic cells (11), suggesting that idelalisib which targets p110 δ shows favorable antitumor efficacy against CML.

Therefore, in the present study, we investigated the antileukemia activities of idelalisib in CML K562 cells.

Materials and methods

Reagents. Idelalisib and imatinib were purchased from Selleck (London, ON, Canada). 3-(4,5-Dimethyl-2-thiazolyl)-2,5-diphenyl-2H-tetrazolium bromide (MTT) reagent was purchased from Amresco (Solon, OH, USA). Antibodies against Akt, phospho-Akt (Ser473), phospho-GSK-3 β (Ser9), caspase-3, -8 and -9, poly(ADP-ribose) polymerase (PARP), β -actin, and anti-mouse and anti-rabbit HRP-conjugated secondary antibodies were obtained from Cell Signaling Technology (Danvers, MA, USA). Antibodies against phospho-pRb (pS780), cyclin D1 and p27 were obtained from

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BD Biosciences Pharmingen (San Jose, CA, USA). Antibodies against lamin B, p21, Bad, Bcl-2 and Bax were purchased from Santa Cruz Biotechnology (Santa Cruz, CA, USA).

Cell culture. The human CML K562 cell line was purchased from the Cell Resource Center, Peking Union Medical College (Beijing, China). The cells were routinely maintained in RPMI-1640 medium supplemented with 10% fetal bovine serum, 1% kanamycin (100 μ g/ml), and 1% glutamine (0.44 μ g/ml) at 37°C in a humidified atmosphere containing 5% CO₂.

MTT assay. The MTT assay was performed to assess cell viability as previously described (12,13). Briefly, cells ($2x10^4$ cells/ml) were cultured in a 96-well plate for 48 h in the presence of 0, 1, 5, 10, 20, 50, 100, 150 and 200 μ M of idelalisib. After addition of MTT (5 mg/ml) to each well, the cells were further incubated for 4 h. The produced formazan blue was dissolved with dimethyl sulfoxide (DMSO), and the absorbance was measured at 490 nm using the microplate reader iMark (Bio-Rad, Hercules, CA, USA).

Soft agar assay. The soft agar assay was carried out as previously described (14) with a small modification. K 562 cells were treated with 0, 20, 50 and 100 μ M of idelalisib for 48 h. Then, the treated cells were seeded on solidified agarose in 60-mm dishes (1.2x10⁴ cells/dish). After incubation for 10 days at 37°C, the cells were fixed with 4% paraformaldehyde and stained with 0.5% crystal violet for 30 min. Colonies were counted under a microscope. Each assay was performed 3 times.

Flow cytometric analysis of cell cycle distribution. Cell cycle analysis was carried out as previously described (15). The cell suspension (4x10⁵ cells/2 ml/well) of K562 cells was planted in a 6-well plate and exposed to various concentrations of idelalisib for 48 h. The cells were harvested, washed with ice-cold phosphate-buffered saline (PBS), and fixed with 75% ethanol. After centrifugation, the fixed cells were resuspended in propidium iodide (PI) solution (25 μ g/ml), and incubated in the dark for 30 min at 4°C to be available for analysis by BD Accuri C6 flow cytometer (BD Biosciences, San Jose, CA, USA). Data were analyzed using FlowJo 7.6 software.

Flow cytometric analysis of apoptosis with Annexin V-FITC/ PI staining. Analysis of apoptosis was carried out by Annexin V-FITC/PI double staining as previously described (16). K562 cells treated with or without idelalisib in a 6-well plate for 48 h were collected, washed with ice-cold PBS, and then stained with 2.5 μ l of Annexin V-FITC and 2.5 μ l of PI (5 μ g/ml) in binding buffer for 15 min at room temperature in the dark. Flow cytometric analysis was conducted using BD FACSVerse flow cytometer (BD Biosciences).

Synergism assay. Synergism was determined by the isobologram and Fa-CI plot based on Chou and Talalay method (17,18). K562 cells seeded in a 96-well plate were exposed to DMSO (as control), idelalisib, imatinib or their combination at a fixed ratio of IC_{50 idelalisib} to IC_{50 imatinib} (390:1) for 48 h. Cell growth inhibition was determined using the MTT assay, and the IC₅₀ values were calculated. The combination index (CI) was calculated using CalcuSyn software according to the method of Chou and Talalay: CI <1 is defined as synergism; CI =1 is defined as an additive effect; and CI >1 is defined as antagonism. All experiments were carried out in triplicate.

Western blot analysis. Cell lysate preparation and western blot analysis were performed as previously described (19,20). Briefly, total and nuclear proteins were prepared using RIPA lysis buffer (Roche Diagnostics, Basel, Switzerland) and NE-PER Nuclear and Cytoplasmic Extraction kit (Thermo Fisher Scientific, Waltham, MA, USA), respectively. Cell lysates with equal amount of protein were subjected to 10% SDS-polyacrylamide gel electrophoresis (PAGE), and the separated proteins were transferred onto a polyvinylidene fluoride (PVDF) membrane (Millipore, Billerica, MA, USA). The membrane was blocked in 5% non-fat dried milk, exposed to the specified primary antibodies overnight at 4°C, and then to the respective secondary antibodies. The blots were visualized using enhanced chemiluminescence (ECL) reagents and digitalized by scanning.

Statistical analysis. Data are presented as mean \pm standard deviation (SD) from 3 independent experiments. The Student's t-test was carried out for analysis of statistical significance using GraphPad Prism 5 software (GraphPad, San Diego, CA, USA). p<0.05 was regarded as indicative of a statistically significant difference.

Results

Idelalisib inhibits the proliferation of K562 cells. The inhibitory activity of idelalisib on the proliferation of K562 cells was assessed by MTT and soft agar assays, respectively. First, the MTT assay was utilized. As shown in Fig. 1A, after exposure to idelalisib at concentrations from 1 to 200 μ M for 48 h, K562 cells showed a dose-dependent decrease in viability, with an IC₅₀ value of 71.4 μ M.

We further examined the antiproliferative activity of idelalisib by use of soft agar assay, which is a suitable method for monitoring anchorage-independent cell growth (21). The cells treated with 0, 20, 50 and 100 μ M of idelalisib for 48 h were grown in soft agar for 10 days. As shown in Fig. 1B and C, treatment with idelalisib decreased both the number and size of the cell colonies, confirming that idelalisib dose-dependently inhibited K562 cell proliferation.

Idelalisib induces cell cycle arrest in the G1 phase in the K562 cells. To determine whether the suppression of K562 cell proliferation by idelalisib is attributed to cell cycle arrest, cell cycle distribution was examined by flow cytometry after idelalisib treatment for 48 h. As shown in Fig. 2A and B, idelalisib induced accumulation of the cell population in the G1 phase, with 60.8% for 50 μ M idelalisib-treated cells vs. 47.4% for control cells. Then, we investigated the effect of idelalisib on cell cycle-related proteins. Fig. 2C indicates that treatment with idelalisib decreased the expression of cyclin D1 and the phosphorylation of pRb, but increased the effect on the phosphorylation of Akt and GSK-3 β , which are known to regulate the cell cycle downstream of the

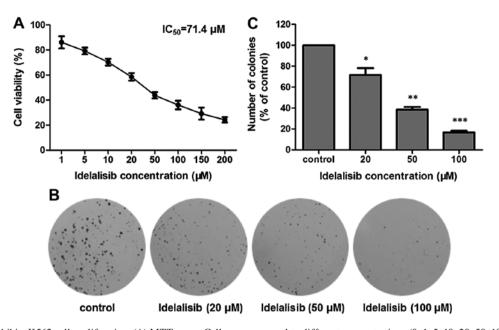


Figure 1. Idelalisib inhibits K562 cell proliferation. (A) MTT assay. Cells were exposed to different concentrations (0, 1, 5, 10, 20, 50, 100, 150 and 200 μ M) of idelalisib for 48 h. Cell viability was measured by determination of the absorbance at 490 nm after addition of MTT reagent. (B) Soft agar assay. K562 cells treated with different concentrations (0, 20, 50 and 100 μ M) of idelalisib for 48 h, were grown in soft agar for 10 days at 37°C, fixed with 4% paraformaldehyde and stained with 0.5% crystal violet for 30 min. The resulting colonies were counted under a microscope. (C) Bar graph showing the number (percentage of control) of colonies formed by K562 cells with or without idelalisib treatment. Data are mean ± SD, representative of 3 independent experiments; *p<0.05, **p<0.01, ***p<0.001 compared with the control.

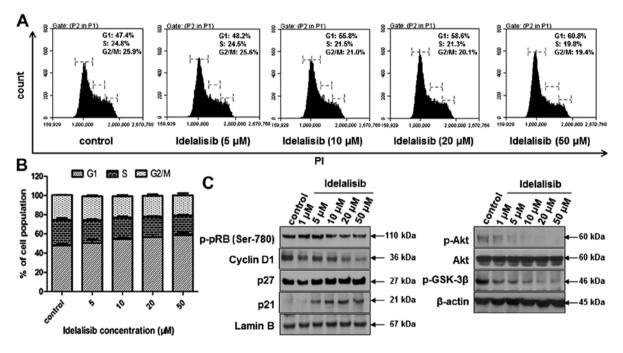


Figure 2. Idelalisib induces K562 cell cycle arrest in the G1 phase. (A) Cell cycle distribution analyzed by flow cytometry. Cells were treated with idelalisib at 0, 5, 10, 20 and 50 μ M for 48 h, stained with PI and subjected to flow cytometry. (B) Bar graph showing the percentage of K562 cells in the G1, S and G2/M phase, respectively. Data represent mean ± SD of 3 independent experiments. (C) Effect of idelalisib on cell cycle-related proteins (left panel) and downstream effectors of the PI3K/Akt pathway (right panel). K562 cells were treated with the indicated concentrations of idelalisib (0, 1, 5, 10, 20 and 50 μ M) for 48 h. Cell lysates were prepared respectively to be available for analysis of protein levels in the nucleus (p-pRb, cyclin D1, p21 and p27) or whole cell (p-Akt, Akt and p-GSK-3 β) by western blotting.

PI3K/Akt pathway (22). Phosphorylation of Akt and GSK- 3β was dose-dependently inhibited by idelalisib, suggesting that the cell cycle arrest effect of idelalisib may be attributed to its blockade of the PI3K/Akt pathway.

Idelalisib induces apoptosis in the K562 cells. Since apoptosis may contribute to a decrease in cell viability, we also examined whether idelalisib induces apoptosis in the K562 cells. Flow cytometric analysis was carried out after Annexin V-FITC/

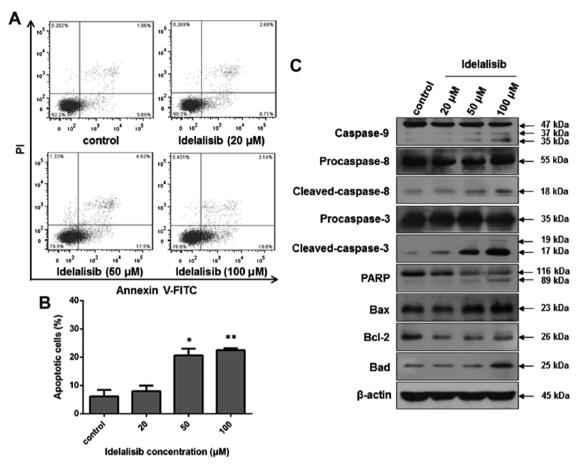


Figure 3. Idelalisib induces apoptosis in the K562 cells. (A) Cell apoptosis analyzed by flow cytometry. Cells were treated with indicated concentrations of idelalisib (0, 20, 50 and 100 μ M) for 48 h, double-stained with Annexin V-FITC/PI and subjected to flow cytometric analysis. (B) Bar graph showing the percentage of apoptotic K562 cells. Data represent mean ± SD of 3 independent experiments; *p<0.05, **p<0.01, compared with the control. (C) Effect of idelalisib on apoptosis-related proteins. K562 cells were treated with idelalisib (0, 20, 50 and 100 μ M) for 48 h. The levels of cleaved caspase-3, -8 and -9, and PARP, as well as the expression of Bcl-2, Bax and Bad were examined by western blotting.

PI staining of K562 cells with or without idelalisib treatment. As indicated in Fig. 3A and B, the cell population in the upper- and lower-right quadrants was increased in a dosedependent manner after idelalisib treatment, with 22.94% for cells treated with 100 μ M idelalisib vs. 7.55% for control cells, suggesting that idelalisib induced apoptosis in the K562 cells. Notably, the increased apoptotic cells were mainly in the early-stage (lower-right quadrant; 19.8% for cells treated with 100 μ M idelalisib vs. 5.69% for control cells).

Then, we investigated the effect on apoptosis-related proteins by western blot analysis. As shown in Fig. 3C, idelalisib treatment increased the level of cleaved caspase-9, -8 and -3, and PARP, as well as the expression of Bad and Bax. In contrast, the expression of anti-apoptotic protein Bcl-2 was reduced. These results suggest that the apoptosis induction by idelalisib in K562 cells may be related to the Bad/Bcl-2/Bax family and the cleavage of caspases and PARP.

Synergistic effect of idelalisib and imatinib in K562 cells. A well-designed drug combination may enhance efficacy while reducing toxicity. To investigate whether idelalisib can enhance the antileukemia activity of the first-line drug imatinib, we carried out a combination study using Chou and Talalay method. K562 cells were treated with idelalisib and imatinib

alone or in combination for 48 h, respectively. MTT assay was conducted to determine the inhibitory activities of each drug and the combination. The IC₅₀ of imatinib was calculated to be 0.183 μ M (Fig. 4A). A synergism study was performed using a series of drug combinations (20, 40, 60, 80 and 100% of IC₅₀ value of each drug) with a fixed ratio of IC₅₀ idelalisib to IC₅₀ imatinib (390:1). As shown in Fig. 4B, co-treatment with the two drugs led to an enhanced cell growth inhibition compared to either treatment alone. Analysis of the data by CalcuSyn software indicated a synergistic effect for the combination, since CI values were <1 at all fraction affected (Fa) levels (Fig. 4C). All of the 3 data points (ED₅₀, ED₇₅ and ED₉₀) were far below the additivity line (Fig. 4D), with the respective CI values as 0.62, 0.47 and 0.35 (Table I), suggesting strong synergy of the combination in regards to the growth inhibition of K562 cells.

Then, we further confirmed the combinational effect of idelalisib and imatinib on K562 cells using various assay methods. Soft agar assay showed that co-treatment with idelalisib (50 μ M) and imatinib (0.128 μ M) decreased cell proliferation more potently than either drug alone (Fig. 4E and F). Cell cycle distribution analysis indicated an increased G1 arrest (Fig. 5A and B), accompanied by further reduction in the level of p-pRb and cyclin D1, and further enhancement in p21 and p27 expression (Fig. 5C). In addition,

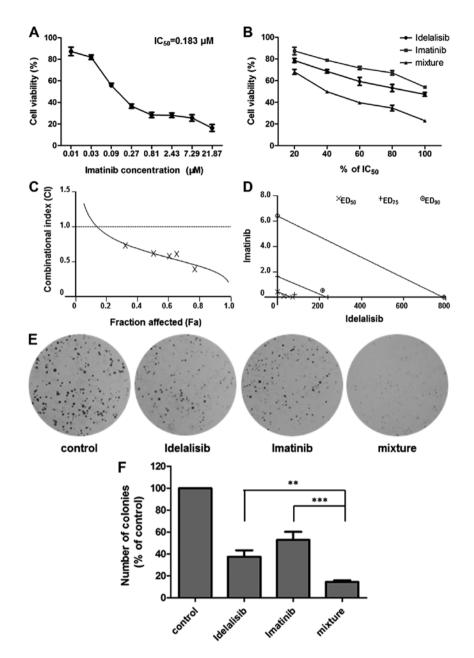


Figure 4. Combination effect of idelalisib and imatinib on K562 cell proliferation. (A) Imatinib inhibited the proliferation of K562 cells. Cells were treated with imatinib (0, 0.01, 0.03, 0.09, 0.27, 0.81, 2.43, 7.29 and 21.87 μ M). Cell viability was determined by MTT assay. IC₅₀ value of imatinib was calculated to be 0.183 μ M. Data are mean ± SD, representative of 3 independent experiments. (B) Combination with idelalisib enhanced the inhibitory activity of imatinib. K562 cells were exposed to a series of concentrations of idelalisib and imatinib (20, 40, 60, 80 and 100% IC₅₀ of each drug), alone or in combination. Cell viability was determined by MTT assay. Data are mean ± SD, representative of 3 independent experiments. (C) Analysis of the combinational effect using CalcuSyn software. Combinational index (CI) values of drug combinations below the horizontal line (CI=1) represent synergy. Fa, fraction affected. (D) Isobologram of idelalisib and imatinib combination. Data points of growth inhibition at 50% (ED₅₀), 75% (ED₇₅) and 90% (ED₉₀) are on the left side of the respective lines, indicating a synergistic effect. (E and F) Combination of idelalisib and imatinib (0.128 μ M) alone or in combination for 48 h, were grown in soft agar for 10 days at 37°C. Colonies were counted under a microscope. Data are mean ± SD, representative of 3 independent experiments; **p<0.01, ***p<0.001, compared with mixture (combination).

	Drug(s)	IC ₅₀ (µM)	r	CI value (mean ± SD)		
Cell line				ED ₅₀	ED ₇₅	ED ₉₀
K562	Idelalisib Imatinib Idelalisib + imatinib	71.4 0.183	0.982 0.998 0.987	0.62±0.09	0.47±0.05	0.35±0.09

Table I. Combination indices (CI) for idelalisib and imatinib.

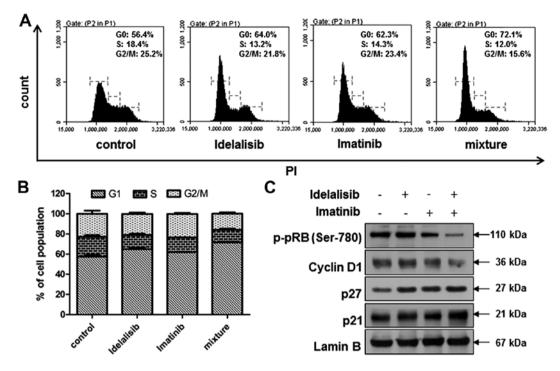


Figure 5. Combination effect of idelalisib and imatinib on K562 cell cycle distribution. (A) Cells treated with idelalisib (50 μ M), imatinib (0.128 μ M) or their combination, were analyzed by flow cytometry after PI staining. (B) Bar graph showing the percentage of K562 cells in the G1, S and G2/M phases, respectively. Data represent mean ± SD of 3 independent experiments. (C) Effect of idelalisib (50 μ M), imatinib (0.128 μ M) or their combination, on cell cycle-related proteins. The levels of cyclin D1, p21, p27 and p-pRb in the nucleus of K562 cells after treatment were determined by western blotting.

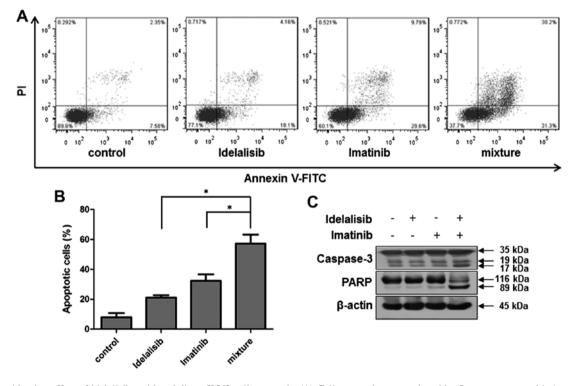


Figure 6. Combination effect of idelalisib and imatinib on K562 cell apoptosis. (A) Cell apoptosis was analyzed by flow cytometry with Annexin V-FITC/ PI staining after treatment with idelalisib (50 μ M), imatinib (0.128 μ M) or their combination. (B) Bar graph showing the percentage of apoptotic cells. Data represent mean ± SD of 3 independent experiments; *p<0.05, compared with mixture. (C) Effect of idelalisib (50 μ M), imatinib (0.128 μ M) or their combination on apoptosis-related proteins. The levels of cleaved caspase-3 and PARP in K562 cells after treatment were examined by western blotting.

combined treatment with idelalisib (50 μ M) and imatinib (0.128 μ M) led to increased cell apoptosis than that induced

by either drug alone (Fig. 6A and B). Notably, co-treatment with the two drugs induced highly increased cell population

in both the upper- and lower-right quadrants, suggesting that the combination of idelalisib and imatinib treatment induced apoptosis in both late and early stages. Consistently, the levels of cleaved caspase-3 and PARP were significantly increased following the combination treatment, as compared with each drug alone (Fig. 6C).

Discussion

Phosphatidylinositol 3-kinases (PI3Ks), consisting of 3 classes (class I, II and III), are closely involved in cell growth and survival (23,24). Among these 3 classes, class I PI3K is the most studied and is closely related to signaling in hematopoietic cells (25). Class I PI3K is further divided into 2 subtypes as class IA and class IB. Class IA PI3K comprises a regulatory subunit and a catalytic subunit ($p110\alpha$, $p110\beta$ or $p110\delta$), whereas class IB PI3K comprises a p101 regulatory subunit and a p110y catalytic subunit. Catalytic isoform p1108 is preferentially expressed in hematopoietic cells (26), suggesting that targeting p110 δ may be a promising strategy for leukemia therapy. Furthermore, it has been reported that PI3K signaling contributes to BCR-ABL transformation and is essential for leukemogenesis of chronic myeloid leukemia (CML) (10). In addition, sustained activation of the PI3K/Akt signaling pathway may contribute to drug resistance due to enhanced drug efflux by ATP-binding cassette transporters such as P-gp (27,28). Occurrence of resistance has become a big challenge in the chemotherapy of CML in recent years. Therefore, targeting PI3K p1108 may be an alternative approach for CML treatment.

In the present study, the antileukemia activity of idelalisib, a specific inhibitor of PI3K p1108, on CML K562 cells was investigated. Our results demonstrated that idelalisib dose-dependently inhibited K562 cell proliferation, which were supported by MTT and soft agar assays. Both MTT and soft agar assays are well known assays which are used to evaluate cell proliferation in vitro, whereas the latter one can additionally predict tumorigenic ability which is correlated with anchorage-independent growth in vivo (14). G1 cell cycle arrest was induced by idelalisib treatment, accompanied by the decreased expression of cyclin D1 and phosphorylation of pRb in contrast to the increased expression of cyclin-dependent kinase (CDK)-inhibitors p27 and p21. Meanwhile, idelalisib treatment blocked the phosphorylation of Akt and GSK-3ß in a dose-dependent manner. Since Akt is known to promote cell cycle progression by upregulating GSK-3 β and cyclin D1 (29), and downregulating CDK inhibitors p27 and p21 (30), the cell cycle arrest effect of idelalisib may be attributed to the blockade of the PI3K/Akt pathway.

Apoptotic cell death is triggered either by the mitochondrial pathway or the death receptor pathway. The former is mainly regulated by the Bcl-2 family which comprises pro-apoptotic proteins such as Bax and Bak, anti-apoptotic proteins such as Bcl-2, and the BH3-only proteins such as Bad, while the latter is controlled by cell surface death receptors such as Fas (31). In the mitochondrial pathway, Bax and Bak are known to disrupt mitochondrial outer membrane integrity through multimerization and therefore release cytochrome c into the cytosol, which in turn activates caspases including caspase-9 and -3, finally leading to apoptosis (32). Bcl-2 exhibits an anti-apoptotic effect by inhibiting formation of the Bax/Bak complex, while Bad promotes apoptosis via neutralizing the inhibitory activity of Bcl-2 on Bax/Bak (32). For the death receptor pathway, Fas induces activation of caspase-8, which cleaves downstream caspases such as caspase-3, promoting apoptosis (31). In the present study, after treatment with idelalisib, expression of Bad and Bax was increased while that of Bcl-2 was reduced; the levels of cleaved caspase-9, -8 and -3 and PARP were enhanced. Akt is known to upregulate Bcl-2 via inhibition of the antagonist Bad in the mitochondrial pathway (33), and to mediate Fas via regulation of FoxO in the cell death receptor pathway (34). Therefore, idelalisib promoted cell apoptosis by activating both the mitochondrial and death receptor pathways, in which targeting PI3K p110& and the downstream effectors may be involved.

Development of imatinib for CML treatment has met with great success. However, the efficacy has been challenged by the increasing occurrence of acquired resistance, and the original insensitivity of a population of patients. The present study indicates that idelalisib could highly enhance the anti-leukemia activity of imatinib on K562 cells, suggesting the possibility for the combinational use of the two drugs in the future.

In conclusion, idelalisib, a novel PI3Kδ-specific inhibitor, alone or in combination with imatinib, exhibited potential antileukemia activity against CML K562 cells, suggesting the possible future application in the treatment of CML patients.

Acknowledgements

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