

Corrections

GENETICS. For the article “A non-ATP-competitive inhibitor of BCR-ABL overrides imatinib resistance,” by Kiranmai Gumireddy, Stacey J. Baker, Stephen C. Cosenza, Premila John, Anthony D. Kang, Kimberly A. Robell, M. V. Ramana Reddy, and E. Premkumar Reddy, which appeared in issue 6, February

8, 2005, of *Proc. Natl. Acad. Sci. USA* (**102**, 1992–1997; first published January 27, 2005; 10.1073/pnas.0408283102), the authors note that an incorrect image was published as Fig. 1*B*. The correct figure and its legend appear below. This correction does not affect the conclusions of the article.

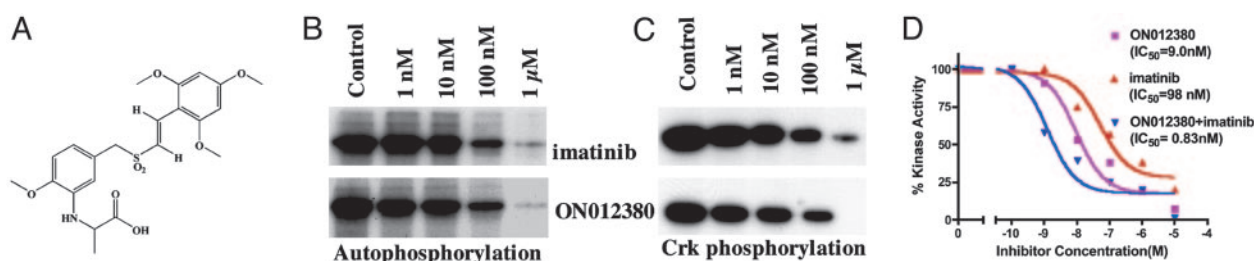


Fig. 1. BCR-ABL inhibitory activity of ON012380. (A) Structure of ON012380. (B and C) Ten nanograms of recombinant BCR-ABL protein was mixed with different concentrations of the indicated inhibitor, and kinase assays were performed by using Crk as a substrate to measure autophosphorylation and substrate (Crk) phosphorylation. (D) BCR-ABL kinase assays were performed as described in *Materials and Methods* by using c-Crk as a substrate. The reactions mixtures were spotted onto strips of P81 phosphocellulose paper, washed, and counted. In experiments for which a mixture of imatinib and ON012380 was used, the reaction mixtures contained a constant amount of imatinib (10 nM) and various amounts of ON012380. The values from individual samples were analyzed and plotted as a function of drug concentration. Data points represent an average of three independent experiments performed in duplicate.

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PLANT BIOLOGY. For the article “A plasma membrane H⁺-ATPase is required for the formation of proanthocyanidins in the seed coat endothelium of *Arabidopsis thaliana*,” by Ivan R. Baxter, Jeffery C. Young, Gordon Armstrong, Nathan Foster, Naomi Bogenschutz, Tatiana Cordova, Wendy Ann Peer, Samuel P. Hazen, Angus S. Murphy, and Jeffrey F. Harper, which appeared in issue 7, February 15, 2005, of *Proc. Natl. Acad. Sci. USA* (**102**, 2649–2654; first published February 4, 2005; 10.1073/pnas.0406377102), due to a printer’s error, Ivan R. Baxter should have been credited in all categories of author contributions, and Jeffery C. Young should have been credited for writing the paper. The corrected author contributions footnote, which appears online only, is below.

Author contributions: I.R.B., J.C.Y., W.A.P., A.S.M., and J.F.H. designed research; I.R.B., J.C.Y., G.A., N.F., N.B., T.C., W.A.P., S.P.H., and A.S.M. performed research; I.R.B., J.C.Y., W.A.P., S.P.H., A.S.M., and J.F.H. analyzed data; and I.R.B., J.C.Y., and J.F.H. wrote the paper.

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CELL BIOLOGY. For the article “Mitochondrial biogenesis by NO yields functionally active mitochondria in mammals,” by Enzo Nisoli, Sestina Falcone, Cristina Tonello, Valeria Cozzi, Letizia Palomba, Mara Fiorani, Addolorata Pisconti, Silvia Brunelli, Annalisa Cardile, Maura Francolini, Orazio Cantoni, Michele O. Carruba, Salvador Moncada, and Emilio Clementi, which appeared in issue 47, November 23, 2004, of *Proc. Natl. Acad. Sci. USA* (**101**, 16507–16512; first published November 15, 2004; 10.1073/pnas.0405432101), the affiliations and address for Maura Francolini appeared incorrectly. The correct address appears below.

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A non-ATP-competitive inhibitor of BCR-ABL overrides imatinib resistance

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Imatinib, which is an inhibitor of the BCR-ABL tyrosine kinase, has been a remarkable success for the treatment of Philadelphia chromosome-positive (Ph⁺) chronic myelogenous leukemias (CMLs). However, a significant proportion of patients chronically treated with imatinib develop resistance because of the acquisition of mutations in the kinase domain of BCR-ABL. Mutations occur at residues directly implicated in imatinib binding or, more commonly, at residues important for the ability of the kinase to adopt the specific closed (inactive) conformation to which imatinib binds. In our quest to develop new BCR-ABL inhibitors, we chose to target regions outside the ATP-binding site of this enzyme because these compounds offer the potential to be unaffected by mutations that make CML cells resistant to imatinib. Here we describe the activity of one compound, ON012380, that can specifically inhibit BCR-ABL and induce cell death of Ph⁺ CML cells at a concentration of <10 nM. Kinetic studies demonstrate that this compound is not ATP-competitive but is substrate-competitive and works synergistically with imatinib in wild-type BCR-ABL inhibition. More importantly, ON012380 was found to induce apoptosis of all of the known imatinib-resistant mutants at concentrations of <10 nM concentration *in vitro* and cause regression of leukemias induced by *i.v.* injection of 32Dcl3 cells expressing the imatinib-resistant BCR-ABL isoform T315I. Daily *i.v.* dosing for up to 3 weeks with a >100 mg/kg concentration of this agent is well tolerated in rodents, without any hematotoxicity.

ON012380 | substrate-competitive | Gleevec

The Philadelphia chromosome (Ph), which was discovered in 1960 by Nowell and Hungerford (1), results from a reciprocal translocation between chromosome 9 at band q34 and chromosome 22 at band q11 (2, 3). This translocation fuses the breakpoint cluster region (*Bcr*) and the *Abl* genes and creates the BCR-ABL oncogene (4). Because the BCR-ABL protein is active in >90% of CML cases, it has been possible to synthesize small molecules that inhibit BCR-ABL kinase activity in leukemic cells without adversely affecting the normal cell population. Imatinib (also called Gleevec or STI571) is a small-molecule inhibitor that binds to the kinase domain of BCR-ABL and stabilizes the protein in its closed, inactive conformation (5), thereby inhibiting its activity, and is now a first-line therapy for the majority of chronic myelogenous leukemia (CML) cases because of its high efficacy level and relatively mild side effects (6). Despite the fact that the majority of patients receiving imatinib respond to treatment at both the hematological and cytogenetic levels, relapse occurs in a large percentage of patients (reviewed in ref. 7). Although several studies have attempted to address the mechanism(s) by which CML cells acquire imatinib resistance (8–10), most studies indicate that mutation of the BCR-ABL gene itself accounts for the majority of imatinib-resistant leukemias *in vivo*. Mutation within the kinase domain is the most common phenomenon, and, to date, 17 different clinically relevant point mutations within this domain have been identified. It is believed that certain amino acid substitutions interfere with the ability of imatinib to interact directly with the BCR-ABL kinase domain, whereas others destroy or hinder the ability of the BCR-ABL kinase domain to adopt a conformation that is required for imatinib binding (reviewed in refs. 7 and 11).

Because of the frequency of mutations within the kinase domain, efforts are now focused on the identification of novel inhibitors that are active against imatinib-resistant mutants of BCR-ABL. Different approaches have recently been described to overcome this resistance in at least some CML cases. Farnesyltransferase inhibitors, such as SCH66336, and the proteasome inhibitor Bortezomib have been shown to have growth inhibitory effects on certain imatinib-resistant leukemias (12). Other compounds, such as PD180970 and CGP76030, both of which inhibit BCR-ABL by binding to the ATP-binding site, have been shown to induce apoptosis in a few select cases of imatinib-resistant CMLs (13–16). More recently, a broad-spectrum ATP-competitive Src kinase inhibitor, BMS-354825, was shown to inhibit the kinase activity of BCR-ABL (17). Although this compound was able to override imatinib resistance caused by the majority of documented mutations within the BCR-ABL kinase domain, it was ineffective against one of the most common mutations observed in imatinib-resistant patients, T315I. This particular amino acid substitution, thus far, has been resistant to all kinase inhibitors that are ATP mimetics, suggesting that other types of therapeutic strategies are required to overcome resistance caused by this mutation. In this report, we describe the profile of ON012380, a small-molecule inhibitor of BCR-ABL that is active against 100% of imatinib-resistant forms of BCR-ABL, including T315I. Because this compound does not compete with ATP to inhibit BCR-ABL but competes with its substrates, these data suggest that molecules that target sites outside the ATP-binding domain can function as effective therapeutic agents against imatinib-resistant leukemias.

Materials and Methods

Cell Lines. 32Dcl3 cells (18) were maintained in Iscove's modified Dulbecco's medium supplemented with 10% FBS, 1 unit/ml penicillin–streptomycin, and 10% WEHI-3B conditioned medium as a source of IL-3 (19). K562 cells were maintained in RPMI medium 1640 supplemented with 10% FBS and 1 unit/ml penicillin–streptomycin.

Generation of Wild-Type and Imatinib-Resistant Mutants of BCR-ABL.

Oligonucleotides corresponding to published mutations (reviewed in ref. 7) that confer imatinib resistance were used to introduce these mutations into the full-length BCR-ABL cDNA by using PCR-based site-directed mutagenesis (20). All constructs were verified by sequence analysis. pCDNA3-based ex-

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Abbreviations: Ph, Philadelphia chromosome; CML, chronic myelogenous leukemia; STAT-5, signal transducer and activator of transcription 5.

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†This work is supported by a grant from Onconova Therapeutics. In addition, S.C.C., S.J.B., M.V.R.R., and E.P.R. act as consultants for Onconova Therapeutics, which is currently developing the compound that is described in this article for clinical trials. M.V.R.R. and E.P.R. are also shareholders in this company.

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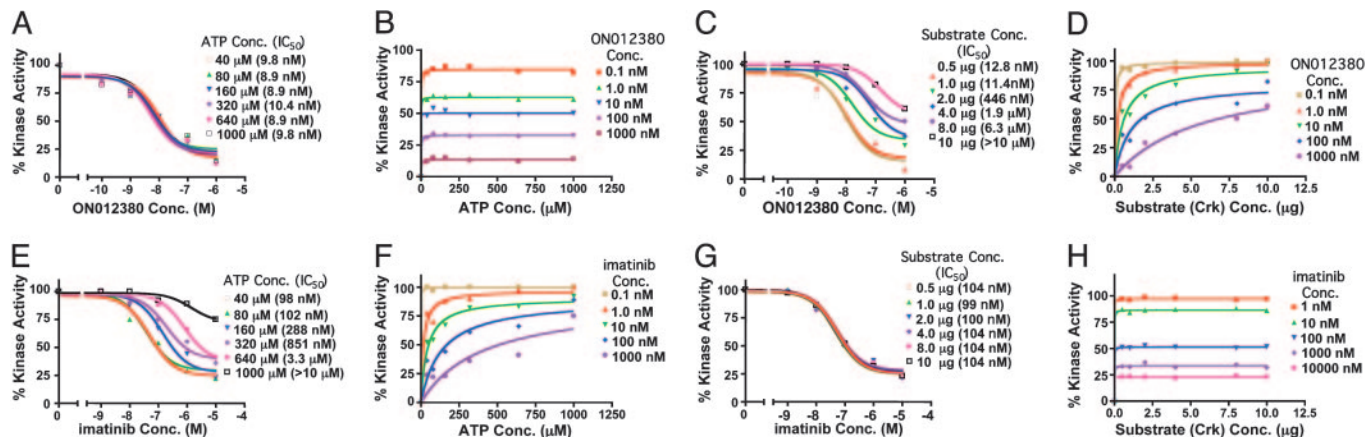


Fig. 2. Steady-state kinetic analysis of BCR-ABL kinase inhibition by ON012380. (A) BCR-ABL kinase inhibition assays were performed as described for Fig. 1 in a reaction mixture containing [γ - 32 P]ATP and various concentrations (conc.) of ATP. The values from individual samples were analyzed and plotted as a function of inhibitor concentration. The IC_{50} of ON012380 for kinase activity was calculated. (B) The curves represent calculated best fits to the Michaelis–Menton equation with a constant amount of substrate and various amounts of ATP and ON012380. (C) BCR-ABL kinase inhibition assays with different concentrations of ON012380 and various concentrations of substrate (Crk) were performed, and the values from individual samples were analyzed and plotted as a function of inhibitor concentration. (D) Michaelis–Menton curves for BCR-ABL with a curve fit derived by using nonlinear regression analysis is shown for data obtained by using a constant amount of ATP and various amounts of substrate and ON012380. (E) Inhibition assays with recombinant BCR-ABL protein and different concentrations of imatinib were performed in the presence of various concentrations of ATP as described for A. (F) The curves represent calculated best fits to the Michaelis–Menton equation with a constant amount of substrate and various amounts of ATP and imatinib. (G) Inhibition assays with recombinant BCR-ABL protein and different concentrations of imatinib were performed in the presence of various concentrations of substrate (Crk) as described for E, and the values from individual samples were analyzed and plotted as a function of drug concentration. (H) The curves represent calculated best fits to the Michaelis–Menton equation with a constant amount of ATP and various amounts of substrate and imatinib.

phorylation as well as Crk phosphorylation, which was used as a substrate (Fig. 1 B and C). Filter binding assays performed with the recombinant protein showed the IC_{50} of this compound to be 9.0 nM. When these assays were performed by using a purified preparation of imatinib, the IC_{50} was found to be \approx 100 nM, which is in close agreement with published data (26). Thus, ON012380 appears to be \approx 10-fold more active than imatinib in BCR-ABL kinase inhibition assays (Fig. 1D). More interestingly, when a mixture of imatinib and ON012380 was used in these assays, these compounds were found to act synergistically to inhibit BCR-ABL, suggesting that they bind to the BCR-ABL kinase at different positions and that this dual binding may lead to potent inactivation of enzymatic activity. (Fig. 1D).

Biochemical Mechanism of Action of ON012380. To define the biochemical mechanism of action of ON012380, we examined the effects of increasing concentrations of ATP or a substrate protein (Crk) on the inhibitory activity of this compound by using steady-state analysis. Ten nanograms of recombinant BCR-ABL was mixed with different concentrations of ON012380, and kinase assays were performed as described in *Materials and Methods* by using [γ - 32 P]ATP and various concentrations of ATP. The values from individual samples were analyzed and plotted as a function of drug concentration (Fig. 2A), and values for K_m (62 μ M) and V_{max} (89 μ mol/min) for the reaction were obtained by fitting the data to the Michaelis–Menton equation (27) (Fig. 2B). These analyses showed that the velocity of substrate phosphorylation was unaltered in the presence of increasing ATP concentrations and that the K_m values remained unchanged, suggesting that ON012380 is not an ATP-competitive inhibitor (Fig. 2A and B). This is also demonstrated by the IC_{50} values which remained unchanged in the presence of increasing ATP concentrations. When the same experiments were performed with imatinib, an opposite result was obtained in which ATP effectively competed with the inhibitor as suggested by an increased IC_{50} with increasing concentration of ATP (Fig. 2E and F), which is accordance with data published in ref. 28. We next examined the effects of increasing substrate concentrations on the inhibitory activity of the compound in the

presence of a constant amount of ATP. Our results (Fig. 2C and D) showed that increasing the concentration of the substrate resulted in increased IC_{50} values for the drug. Data analysis with the Michaelis–Menton equation (27) demonstrated that the maximal velocity of BCR-ABL was not significantly affected by the inhibitor despite the fact that the K_m values were significantly increased. The increase in K_m combined with the unchanged V_{max} in the presence of an inhibitor is characteristic of competitive inhibition. The increased IC_{50} values in the presence of increased substrate concentration for the inhibition of BCR-ABL kinase activity (Fig. 2C) further demonstrate the substrate-dependent and ATP-independent nature of inhibition. When these assays were performed with imatinib, there was no change in the K_m values with increasing concentration of the substrate (Fig. 2G and H), confirming previously published data showing that imatinib acts as an ATP-dependent and substrate-independent inhibitor of BCR-ABL.

We next examined the inhibitory effects of ON012380 on a panel of serine/threonine and tyrosine kinases (Table 2, which is published as supporting information on the PNAS web site). These studies showed that ON012380 does not exhibit significant inhibitory activity against any of the other kinases tested. However, at higher concentrations, this compound was found to inhibit the platelet-derived growth factor receptor as well as the LynB and Fyn tyrosine kinases. These results suggest a common feature shared between these tyrosine kinases in their substrate binding sites. Of the kinases tested, ON012380 was most active against BCR-ABL, which is likely to be its primary target.

Activity of ON012380 Against BCR-ABL-Expressing Cell Lines. To determine the activity ON012380 against BCR-ABL-expressing cell lines, we incubated K562 cells as well as 32Dcl3 cells that ectopically expressed the wild-type p210^{BCR-ABL} oncoprotein with various concentrations of ON012380 for a period of 72 h and determined their viability by trypan blue exclusion. Incubation of both cell lines with ON012380 (Fig. 3A and B) resulted in a rapid loss of viability, with a LD_{50} of 10–15 nM. When these assays were performed by using imatinib, the IC_{50} of this compound was found to be $>$ 100 nM (Fig. 3A and B), which is

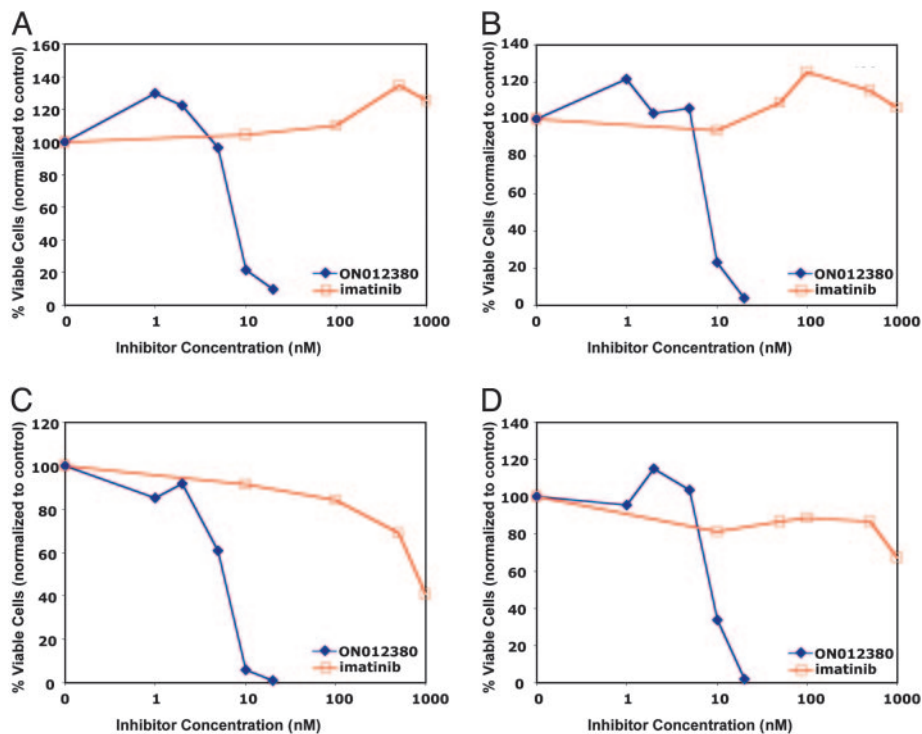


Fig. 5. *In vitro* tumor-cell-killing activity of cells expressing an imatinib-resistant mutant of BCR-ABL by ON012380. The four representative imatinib-resistant cell lines T315I (A), E255K (B), Y253H (C), and G250E (D) were incubated with increasing concentrations of the indicated compounds, and the total number of viable cells was determined 72 h after treatment by trypan blue exclusion.

in the presence of 1, 10, 100, and 1,000 nM ON012380 or imatinib were subjected to immunoprecipitation followed by kinase assays (Fig. 4B and C), we observed a dose-dependent reduction in the level of BCR-ABL kinase activity in the ON012380-treated cells, whereas no such reduction was observed in the imatinib-treated samples. When the status of STAT-5 phosphorylation in ON012380 and imatinib-treated cells was examined (Fig. 4D), there was a substantial inhibition of STAT-5 phosphorylation in the ON012380-treated cells, whereas the phospho-STAT-5 levels were unaffected in the imatinib-treated cells. Similar results were obtained by using 32Dcl3 cells that express the other imatinib-resistant mutants listed in Table 1 (data not shown). Taken together, these results suggest that ON012380 effectively inhibits the activity of the imatinib-resistant BCR-ABL proteins, including BCR-ABL^{T315I}.

***In Vivo* Efficacy of ON012380.** To assess the *in vivo* efficacy of ON012380, athymic nude mice were injected i.v. with 32Dcl3 cells

expressing the T315I mutant form of BCR-ABL. For these studies, the mice were injected i.v. through the tail vein with 1×10^6 32Dcl3 cells expressing the T315I mutant; 24 h after injection, the mice were divided into three groups (10 mice per group), with each group receiving either saline (vehicle), ON012380 (100 mg/kg), or imatinib (100 mg/kg) i.p. on a daily basis. The dose and treatment schedule was chosen to mimic the maximum tolerated dose of imatinib (unpublished results) in this particular strain of mice.

On days 7 and 14 after the beginning of treatment, the number of T315I cells in the blood of mice treated with ON012380 was compared with the number of T315I cells found in mice treated with either imatinib or saline. The results of this study (Fig. 6A) showed that the number of T315I cells in the blood of mice treated with ON012380 was significantly reduced on days 7 and 14 as compared with the number of cells found in both the vehicle- and imatinib-treated groups. The ON012380-treated mice showed no signs of toxicity, body weight loss, ruffled coats,

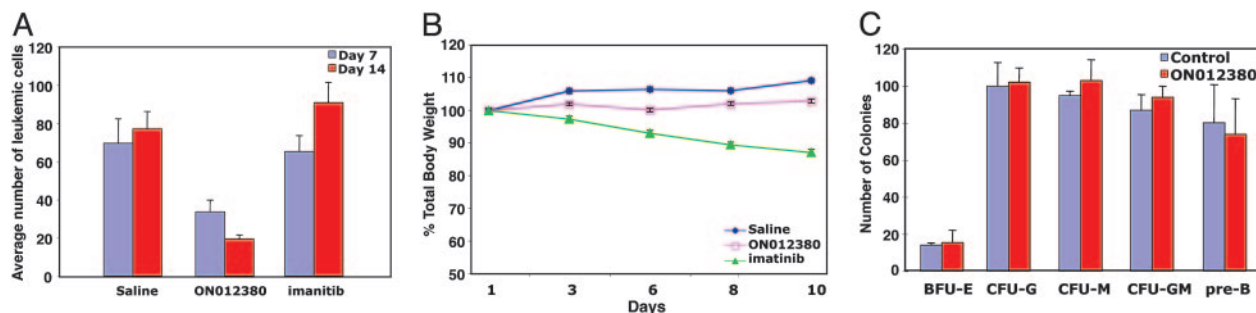


Fig. 6. Effect of ON012380 on the *in vivo* growth of T315I cells. (A) Nude mice (10 mice per group) were injected i.v. through the tail vein with 1×10^6 32D/BCR-ABL^{T315I} cells. Treatment with daily i.p. injections of 100 mg/kg ON012380, 100 mg/kg imatinib, or an equal volume of saline was initiated 24 h later. Blood smears from each mouse were performed on days 7 and 14, and the number of T315I-expressing cells per 10 fields was determined. The data were plotted as the average number of 32D/BCR-ABL^{T315I} cells per 10 fields \pm SEM ($n = 10$). (B) The total body weight of individual mice in the three groups was determined daily, and the average body weights were plotted as the percent of starting body weight. (C) CD-1 mice were injected i.v. (tail vein injection) with saline or ON012380 (200 mg/kg) dissolved in saline. Bone marrow cells were extracted from the mice after 24 h, and 2×10^5 cells were plated on methycellulose containing appropriate cytokines for lineage-specific colony formation. Colonies were counted after 5–14 days of incubation. BFU-E, erythroid burst-forming unit; CFU-G, granulocyte colony-forming unit; CFU-M, macrophage colony-forming unit; CFU-GM, granulocyte/macrophage colony-forming unit; pre-B, pre-B lymphocyte.

lethargy, or abnormal feces. In contrast, administration of imatinib for 10 days produced severe toxicity as judged by a >20% loss of bodyweight, which resulted in the termination of drug administration (Fig. 6B). ON012380 has been shown to be well tolerated over a 21-day period of daily i.p. injections (data not shown) and is therefore well suited for long-term treatments. We have also carried out single dose and repeated dose (28 daily injections) toxicology studies. In mice, a single dose of 300 mg/kg produced no toxicity. During 7 days of repeated dosing, 200 mg/kg per dose was well tolerated. During 28 days of repeated dosing, fixed daily doses of 100 mg/kg were well tolerated. To examine the effects of ON012380 on normal hematopoietic cell population *in vivo*, we injected an i.v. dose of this compound at a concentration of 200 mg/kg and examined its effect on the *in vitro* hematopoietic colony formation of normal bone marrow cells derived from these mice 24 h after administration. These studies (Fig. 6C) show that there was no reduction in colony formation of the erythroid, myeloid, or lymphoid lineages. Taken together, this study shows that ON012380 exhibits a very desirable safety profile with a high therapeutic index and can reduce the *in vivo* growth of imatinib-resistant cells in an efficacious manner.

Discussion

Recent advances in our understanding of the molecular changes that accompany cell transformation has provided new approaches for the development of cancer therapeutic agents that can potentially target specific oncoproteins that are activated in a cancer cell. Imatinib is one of the first in this new generation of drugs that specifically target the BCR-ABL tyrosine kinase (5), and it has been a great success. However, it is becoming clear that a significant portion of patients chronically treated with imatinib develop resistance because of the acquisition of mutations in the kinase domain of BCR-ABL, which are thought to interfere with the ability of the enzyme to adopt the inactive conformation required for imatinib binding. Because selection of highly conserved mutable residues in the ATP binding site appears to be relatively common for many kinases, it has been argued that substrate-competitive inhibitors might constitute better drug candidates (30). Therefore, in our quest to develop other BCR-ABL inhibitors, we chose to develop compounds that do not compete for the ATP binding site of this enzyme, because such compounds offer the potential to be unaffected by mutations that make Ph⁺ CML cells resistant to imatinib. The results presented in this study describe the identification of such a pharmacophore, ON012380, which targets the BCR-ABL kinase and induces cell death of Ph⁺ CML cells at a concentration of 10 nM, which is >10-fold more potent than imatinib. Results presented in our study also demonstrate that this compound targets a

site that is different from the binding site of imatinib because ATP fails to compete with ON012380. Our results also suggest that ON012380 probably binds to the substrate binding site of the enzyme because a natural substrate, such as Crk, readily competes with ON012380 and interferes with its ability to inhibit BCR-ABL kinase. Furthermore, imatinib and ON012380 were found to synergistically inhibit wild-type BCR-ABL, suggesting that these two compounds bind to different sites on the enzyme.

Emerging data suggests that the majority of CML patients who achieve complete cytogenetic remission rarely achieve complete molecular remission (31, 32). This phenomenon appears to be due to the existence of rare resistant mutations within the pool of BCR-ABL⁺ cells, which, in the absence of imatinib, lack any growth advantage. It is now believed that imatinib treatment results in the elimination of wild-type BCR-ABL⁺ cells, leading to a cytogenetic remission. However, given adequate time, populations of CML cells harboring imatinib-resistant mutations expand, leading to a relapse of the disease (17). Our results show that ON012380 is very effective at inhibiting all of the imatinib-resistant mutants of BCR-ABL tested, including T315I. In addition to mutation within BCR-ABL, imatinib-resistance has also been recently attributed to overexpression and/or activation of Lyn (33, 34), which functions downstream of BCR-ABL. Studies have shown that inhibition of Lyn expression or activity in imatinib-resistant cells with short interfering RNA (35) or Src inhibitors reduced their proliferative capacity and ultimately induced apoptosis (33, 34). Because ON012380 is highly selective against BCR-ABL, it remains to be determined whether it will be effective against those samples in which Lyn function (as opposed to BCR-ABL mutation) mediates the resistance to imatinib. However, it is of interest to note that this compound also inhibits Lyn kinase activity in the nanomolar range (85 nM) (Table 2). It is therefore likely that ON012380 will also be effective against cells in which imatinib resistance is conferred by this pathway.

Another important feature of ON012380 is the very desirable safety profile that is not often seen in conventional chemotherapeutic agents. In mice, single doses of 300 mg/kg of ON012380 produced no toxicity. The low *in vivo* toxicity and the potent BCR-ABL inhibitory activity seen in *in vitro* and nude mouse assays bodes well for further development of this compound for the treatment of imatinib-resistant myeloid leukemias. Because this compound is highly effective in combination with imatinib, the lack of bone marrow toxicity may be beneficial for testing novel combinations for Ph⁺ CML, which may lead to complete molecular remission of the disease and prevent the appearance of imatinib-resistant forms of this disease over the course of time.

This work was supported by grants from Onconova Therapeutics and the Fels Foundation.

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