The ABL Switch Control Inhibitor DCC-2036 Is Active against the Chronic Myeloid Leukemia Mutant BCR-ABL$^{T315I}$ and Exhibits a Narrow Resistance Profile

Christopher A. Eide$^{1,2}$, Lauren T. Adrian$^{1,2}$, Jeffrey W. Tyner$^{1}$, Mary Mac Partlin$^{1}$, David J. Anderson$^{1}$, Scott C. Wise$^{3}$, Bryan D. Smith$^{3}$, Peter A. Petillo$^{3}$, Daniel L. Flynn$^{3}$, Michael W.N. Deininger$^{1}$, Thomas O’Hare$^{1,2}$, and Brian J. Druker$^{1,2}$

Abstract

Acquired point mutations within the BCR-ABL kinase domain represent a common mechanism of resistance to ABL inhibitor therapy in patients with chronic myeloid leukemia (CML). The BCR-ABL$^{T315I}$ mutant is highly resistant to imatinib, nilotinib, and dasatinib, and is frequently detected in relapsed patients. This critical gap in resistance coverage drove development of DCC-2036, an ABL inhibitor that binds the switch control pocket involved in conformational regulation of the kinase domain. We evaluated the efficacy of DCC-2036 against BCR-ABL$^{T315I}$ and other mutants in cellular and biochemical assays and conducted cell-based mutagenesis screens. DCC-2036 inhibited autophosphorylation of ABL and ABL$^{T315I}$ enzymes, and this activity was consistent with selective efficacy against Ba/F3 cells expressing BCR-ABL (IC$_{50}$ 19 nmol/L), BCR-ABLT315I (IC$_{50}$ 63 nmol/L), and most kinase domain mutants. Ex vivo exposure of CML cells from patients harboring BCR-ABL or BCR-ABLT315I to DCC-2036 revealed marked inhibition of colony formation and reduced phosphorylation of the direct BCR-ABL target CrkL. Cell-based mutagenesis screens identified a resistance profile for DCC-2036 centered around select P-loop mutations (G250E, Q252H, Y253H, E255K/V), although a concentration of 750 nmol/L DCC-2036 suppressed the emergence of all resistant clones. A decreased concentration of DCC-2036 (160 nmol/L) in dual combination with either nilotinib or dasatinib achieved the same zero outgrowth result. Further screens for resistance due to BCR-ABL compound mutations (two mutations in the same clone) identified BCR-ABLE255V / T315I as the most resistant mutant. Taken together, these findings support continued evaluation of DCC-2036 as an important new agent for treatment-refractory CML.

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Introduction

Use of ABL inhibitors to block the activity of the oncogenic BCR-ABL tyrosine kinase in the treatment of chronic myeloid leukemia (CML) serves as a model for molecularly targeted cancer therapies. The ABL inhibitor imatinib has an extensive and impressive clinical track record in CML, with newly diagnosed, chronic phase patients demonstrating 5-year overall and progression-free survival rates of 89% and 93%, respec-}

Authors’ Affiliations: $^{1}$Oregon Health & Science University Knight Cancer Institute and $^{2}$Howard Hughes Medical Institute, Portland, Oregon; and $^{3}$Deciphera Pharmaceuticals LLC, Lawrence, Kansas

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Thomas O’Hare and Brian J. Druker contributed equally to this work.

Corresponding Author: Thomas O’Hare, Howard Hughes Medical Institute/Oregon Health & Science University Knight Cancer Institute, Mailcode LS92, 3181 SW Sam Jackson Park Road, Portland, OR 97239. Phone: 503-494-1916; Fax: 503-494-3688. E-mail: oharet@ohsu.edu.

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off-rate for binding both ABL and ABL\textsubscript{T315I}, and demonstrates additional highly selective activity for FLT3, TIE2, and SRC-family kinases. DCC-2036 also showed significant efficacy and improved survival in a murine bone marrow transplantation model of BCR-ABL\textsubscript{T315I}-driven CML (5). Here, we evaluate the efficacy of DCC-2036 against BCR-ABL\textsubscript{T315I} and other mutants in CML cell lines and primary cells, and establish the anticipated resistance profile for DCC-2036 using cell-based screens.

Materials and Methods

**ABL autophosphorylation assays**

Kinase autophosphorylation assays with tyrosine-dephosphorylated ABL and ABL\textsubscript{T315I} (Invitrogen) were performed alone or with DCC-2036 (0.2–3.125 nmol/L) or imatinib (625–3,125 nmol/L) as described (6).

**Cell lines**

Certified Ba/F3, K562, KYO1, LAMA, HEL, CMK, and Marimo cells were obtained from the American Type Culture Collection and grown in the recommended culture medium. Ba/F3 transfectants expressing native BCR-ABL or BCR-ABL with a single kinase domain mutation were generated and maintained as described (7). The Ba/F3 BCR-ABL\textsubscript{T315A} cell line was a gift from N. Shah (UCSF). None of the cell lines used in this study was cultured for longer than 6 months from initial purchase or characterization. No further authentication of cell line characteristics was done.

**Cell proliferation assays**

Parental Ba/F3 cells and Ba/F3 cells expressing native or mutant BCR-ABL (4 × 10\textsuperscript{4} per well) were incubated alone or with DCC-2036 (0.2–15,625 nmol/L) for 72 hours. Proliferation measurements and IC\textsubscript{50} value determinations were done as described (7). Identical experiments were carried out for CML (K562, KYO1, LAMA) and non-CML cell lines (HEL, CMK, Marimo).

**Immunoblot analyses of CrkL phosphorylation**

For cell line experiments, Ba/F3 cells expressing CBR-ABL, BCR-ABL\textsubscript{L255V}, or BCR-ABL\textsubscript{T315I} (5 × 10\textsuperscript{6} cells/well) were cultured 4 hours in complete media alone or with DCC-2036 (0.2–3.125 nmol/L) or imatinib (0.2–3.125 nmol/L) as described (8). For primary cell experiments, following informed consent, peripheral blood mononuclear cells from a newly diagnosed CML patient and from an accelerated phase patient harboring BCR-ABL\textsubscript{T315I} were cultured (2 × 10\textsuperscript{6} per well) overnight in Iscove’s modified Dulbecco’s medium (Irvine) containing 20% bovine serum albumin, insulin, transferrin (BIT) serum substitute (Stemcell) alone or with DCC-2036 (0.2–625 nmol/L), imatinib (1,000 nmol/L), nilotinib (200 nmol/L), or dasatinib (50 nmol/L). For all experiments, cells were lysed in boiling SDS-PAGE loading buffer supplemented with 0.1 mmol/L AEBSF and 0.1 mmol/L Na\textsubscript{2}VO\textsubscript{4}, subjected to SDS-PAGE, and immunoblotted with antibodies against phosphorylated (anti-Y207; Cell Signaling) or total CrkL (C-20; Santa Cruz).

**Hematopoietic colony formation assays**

To assess granulocyte/macrophage colony formation (CFU-GM), mononuclear cells from bone marrow of a newly diagnosed CML patient, an accelerated phase patient harboring BCR-ABL\textsubscript{T315I}, or a healthy donor were obtained following informed consent and plated alone or with DCC-2036 (50 or 500 nmol/L) or imatinib (2,000 nmol/L) in triplicate (5 × 10\textsuperscript{4} cells/plate) in IMDM/methylcellulose as described (9). Results are reported as percentage of colonies relative to untreated.

**Cell-based resistance screens**

For DCC-2036 screens starting from Ba/F3 cells expressing native BCR-ABL, cells were treated overnight with N-ethyl-N-nitrosourea (ENU; 50 μg/mL) and resuspended in complete media (1 × 10\textsuperscript{5} cells/well) supplemented with DCC-2036 (50–1,250 nmol/L) as described (10). DCC-2036 (160 nmol/L) was also evaluated in dual combinations with imatinib (2,000 nmol/L), nilotinib (500 nmol/L), or dasatinib (25 nmol/L). Wells exhibiting outgrowth were expanded, sequenced, and analyzed for mutations (Mutation Surveyor; SoftGenetics) as described (8). Similar experiments were conducted starting from Ba/F3 BCR-ABL\textsubscript{T315I} cells treated with DCC-2036 (250–750 nmol/L), and from a pooled mix of equal cell numbers of all Ba/F3 BCR-ABL cell lines treated with a cocktail of ABL kinase inhibitors [DCC-2036 (250 nmol/L), nilotinib (500 nmol/L), and dasatinib (25 nmol/L)].

**Results and Discussion**

We established that DCC-2036 (Fig. 1A) directly inhibits the catalytic activity of ABL and ABL\textsubscript{T315I} by evaluating kinase autophosphorylation activity. Although both imatinib and DCC-2036 attenuated activity of ABL, only DCC-2036 blocked ABL\textsubscript{T315I} tyrosine autophosphorylation (Fig. 1B). Unlike imatinib, nilotinib, and dasatinib, the binding mode of DCC-2036 to ABL or ABL\textsubscript{T315I} does not require a hydrogen bond to the side chain hydroxyl of native T315 and avoids a steric clash with mutated T315. On binding, DCC-2036 induces and stabilizes an aspartate-phenylalanine-glycine motif (DFG)-out, catalytically inactive conformation of the kinase domain, precluding phosphorylation of activation loop residue Y393, a critical event preceding full catalytic activation of ABL kinase (5).

Cellular assays demonstrated that DCC-2036 selectively inhibits most clinically relevant imatinib-resistant mutants (Fig. 1C). DCC-2036 inhibited growth of Ba/F3 cells expressing BCR-ABL (IC\textsubscript{50} 19 nmol/L) with approximately 16-fold higher potency than imatinib and, of key importance, cells expressing the imatinib-, nilotinib-, and dasatinib-resistant BCR-ABL\textsubscript{T315I} mutant (IC\textsubscript{50} 63 nmol/L). The selectivity of DCC-2036 for BCR-ABL–positive cells was evidenced by its marked inhibition of CML cell lines compared with non-CML leukemia lines (Fig. 1C). Sensitivity of BCR-ABL mutants to DCC-2036 segregated into 3 categories: (a) IC\textsubscript{50} ≤ 50 nmol/L: 1/14; M351T, (b) IC\textsubscript{50} < 100 nmol/L: 8/14; M244V, G250E, Q252H, Y253F, T315A, T315I, F317L, H396P, and (c) IC\textsubscript{50} > 100 nmol/L: 5/14; Y253H, E255K, E255V, F317V, F359V. Among these, BCR-ABL\textsubscript{L255V} was least sensitive to DCC-2036 (IC\textsubscript{50} 410 nmol/L).
Immunoblot analyses examining the ability of DCC-2036 to block tyrosine phosphorylation of the direct BCR-ABL substrate CrkL revealed greater inhibition in cells expressing BCR-ABL or BCR-ABL<sup>T315I</sup> than BCR-ABL<sup>E255V</sup> (Fig. 1D). These findings suggest that, although DCC-2036 exhibits activity against the T315I mutant, select mutations of the P-loop such as E255V may be more problematic. Notably, BCR-ABL<sup>E255V</sup> is highly resistant to imatinib and confers moderate resistance to both nilotinib and dasatinib in vitro (7), and has been reported in clinical failures of each of these therapies (4, 11, 12).

As follow-up to the efficacy of DCC-2036 observed in BCR-ABL–positive cells, particularly the BCR-ABL<sup>T315I</sup> mutant, we evaluated DCC-2036 against mononuclear cells from a newly diagnosed CML chronic phase patient and an accelerated phase patient harboring BCR-ABL<sup>T315I</sup>. <i>In vivo</i> exposure of primary BCR-ABL<sup>T315I</sup> cells to DCC-2036 sharply reduced CrkL phosphorylation, whereas imatinib, nilotinib, and dasatinib were ineffective (Fig. 2A). All inhibitors reduced CrkL phosphorylation in primary cells from the newly diagnosed CML patient (Fig. 2B), although imatinib (1 μmol/L) showed limited effect. CrkL phosphorylation is a clinical biomarker of BCR-ABL activity, and its inhibition in primary CML cells has been correlated with degree of response achieved on therapy (13). Although complete pharmacodynamic data for DCC-2036 have not yet been reported, our results demonstrate that DCC-2036 is active in clinical isolates from CML patients harboring BCR-ABL or BCR-ABL<sup>E255V</sup>. This is corroborated by colony formation data for primary CML cells, wherein exposure of cells from the same BCR-ABL<sup>T315I</sup> CML patient (Fig. 2A) and newly diagnosed CML patient (Fig. 2B) to DCC-2036 substantially reduced outgrowth of CML cells, with no toxicity toward mononuclear cells from a healthy individual (Fig. 2C).

Given the unique binding characteristics of DCC-2036, we screened for resistance-conferring mutations specific to DCC-2036 but susceptible to other ABL inhibitors. Results from a cell-based resistance screen for BCR-ABL mutants persisting in the presence of DCC-2036 revealed a concentration-dependent
reduction in outgrowth and the spectrum of resistant subclones recovered (Fig. 3A; Supplementary Table S1). The resistance profile of DCC-2036 narrowed to a subset of mutations described for imatinib (G250E, Q252H, Y253H, E255K, E255V; refs. 4, 10), as has been largely the case for other ABL inhibitors [dasatinib (10), nilotinib (10), SGX393 (9), AP24534 (8)], suggesting a limited set of resistance-conferring mutations can be tolerated without crippling kinase function. Structurally, the vulnerability of DCC-2036 to P-loop mutations (e.g., E255V) suggests subtle local alterations in the ATP binding site may effectively destabilize the inactive conformation, as for imatinib (3). A full explanation pertaining to DCC-2036 will require further crystallographic, dynamic (e.g., NMR), and in silico analysis (5).

Impressively, resistant outgrowth was completely suppressed by DCC-2036 at 750 nmol/L (Fig. 3A; Supplementary Table S1). As clinically achievable plasma levels of DCC-2036 have not yet been reported and select P-loop mutants confer partial resistance to DCC-2036, nilotinib (7), and dasatinib (7), we evaluated dual combinations of DCC-2036 (160 nmol/L) with each clinical ABL inhibitor in resistance screens. Although the combination of DCC-2036 with imatinib (2 μmol/L) reduced the fraction of wells with outgrowth, P-loop vulnerabilities at residues G250, Y253, and E255 were detected (Fig. 3B; Supplementary Table S2). No resistant subclones were recovered with dual combinations of DCC-2036 and clinically achievable concentrations of nilotinib (500 nmol/L) or dasatinib (25 nmol/L). These findings are similar to those from studies with another ABLT315I inhibitor, SGX393 (9), and suggest that ABL inhibitor cocktails that include an ABLT315I inhibitor such as DCC-2036 may represent a rational therapeutic approach to mitigating resistance.

As the immediate clinical application of an ABLT315I inhibitor is in refractory CML patients harboring this mutation, we performed resistance screens starting from Ba/F3 cells expressing BCR-ABL T315I to identify BCR-ABL compound mutations (2 mutations in the same clone) conferring increased resistance to DCC-2036. Such mutations have been reported in clinical failures to dasatinib or nilotinib salvage therapy, suggesting potential for selection on sequential treatment with ABL inhibitors (14–17). The compound mutation-based resistance profile for DCC-2036 narrowed predominantly to BCR-ABL E255V / T315I (83.3% of recovered mutants at 750 nmol/L; Fig. 3C; Supplementary Table S3). An additional mutant featuring substitution of the baseline isoleucine at residue 315 for methionine (I315M) was also recovered. To our knowledge, mutation of the gatekeeper residue (either the native threonine or mutant isoleucine) to methionine has not been observed in resistance to other ABL tyrosine kinase inhibitors. DCC-2036 forms hydrogen bonds with the nearby ATP hinge

Figure 2. BCR-ABL–mediated signaling and colony formation of primary CML cells is selectively inhibited by DCC-2036. Effect of DCC-2036 on CrkL phosphorylation and colony formation in primary CML cells harboring BCR-ABL T315I (patient 07/316; A) or native BCR-ABL (patient 07/207; B). Immunoblot analysis for CrkL phosphorylation (p-CrkL) was performed following ex vivo exposure of mononuclear cells to DCC-2036, imatinib, nilotinib, or dasatinib using an anti-Y207 CrkL antibody. Colony assays (CFU-GM) were conducted by plating mononuclear cells in methylcellulose containing DCC-2036 or imatinib. Similar assays were done using mononuclear cells from a healthy individual (C). Results are expressed as percent of the number of no treatment colonies; error bars represent SEM.
DCC-2036 Inhibits BCR-ABL<sup>T315I</sup>

Figure 3. DCC-2036 suppresses resistance alone or in ABL inhibitor combinations and exhibits few BCR-ABL compound mutant vulnerabilities. A, BCR-ABL mutants recovered from cell-based resistance screens for DCC-2036, starting from Ba/F3 BCR-ABL cells. ENU-mutagenized cells were plated with inhibitor, monitored for outgrowth, expanded, and sequenced for mutations. Bars represent frequency of a given mutant among all recovered clones at a given inhibitor concentration; percent of wells demonstrating outgrowth and number of clones sequenced is indicated. B, BCR-ABL mutants recovered in the presence of dual combinations of DCC-2036 and imatinib, nilotinib, or dasatinib, starting from Ba/F3 BCR-ABL cells. C, BCR-ABL compound mutants recovered in the presence of DCC-2036, starting from Ba/F3 BCR-ABL<sup>T315I</sup> cells. See also Supplementary Tables S1, S2, and S3.
site residue M318 and the K271-E286 salt bridge, allowing for accommodation of the bulky isoleucine substitution in BCR-ABL T315I. Electrostatic interaction with E282 aids in stabilizing the E282-R386 switch control pair interaction and, consequently, the inactive kinase conformation (5). One explanation for the resistance of the mutant featuring methionine at residue 315 may be that the methionine sidechain impinges on DCC-2036 binding. Alternatively, introducing methionine at the gatekeeper position may induce the ABL kinase domain to adopt an active conformation.

To both broaden the screen for BCR-ABL compound-mutant-mediated resistance and evaluate efficacy of ABL inhibitor cocktails in this setting, we carried out a similar screen starting from a pooled mix of Ba/F3 BCR-ABL mutant cell lines (representing >70% of clinically observed imatinib-resistant mutations; ref. 4) using a combination of DCC-2036 (250 nmol/L), nilotinib (500 nmol/L), and dasatinib (25 nmol/L; Supplementary Fig. 1A). Strikingly, only 3 compound mutations were recovered: G250E/T315I, E255V/T315A, and E255V/T315I. Among these, the BCR-ABL T315K/T315I mutant has been observed clinically and reported to confer high level resistance to multiple other ABL T315I inhibitors (Supplementary Table S4; refs. 8, 15). Thus, although ABL inhibitor cocktails that include an ABL T315I inhibitor may prove an effective strategy in minimizing resistance, certain BCR-ABL compound mutations are predicted to be recalcitrant to such an approach.

Our investigation of the switch control inhibitor DCC-2036 reveals substantial activity in CML cells, including cells expressing BCR-ABL T315I. DCC-2036 is undergoing phase I evaluation for use in imatinib-refractory CML (NCT00827138; www.clinicaltrials.gov), and our results suggest that it may provide a treatment option for relapsed patients with a T315I mutation. DCC-2036 adds to a small set of ABL T315I inhibitors currently in development, each of which targets the BCR-ABL T315I mutant differently. Recent approaches include: dodging I315 via a carbon–carbon triple bond (ponatinib, formerly AP24534; ref. 8; pivotal phase 2 trial; NCT01207440; www.clinicaltrials.gov), utilizing a modified nilotinib–dasatinib hybrid structure to avoid gatekeeper mutations (HG-7–85-01; ref. 18; preclinical), and combining ATP-competitive and allosteric ABL inhibitors (GN-2; refs. 19, 20 preclinical). Although disease eradication remains on the horizon, the much anticipated, imminent clinical availability of ABL T315I inhibitors represents an important step toward maximal disease control.

Disclosure of Potential Conflicts of Interest

OSU and B.J. Druker have a financial interest in MolecularMD. Technology used in this research has been licensed to MolecularMD. This potential conflict of interest has been reviewed and managed by the OSU Conflict of Interest in Research Committee and the Integrity Program Oversight Council.

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