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Benzimidazole Derivatives as Potent JAK1-Selective Inhibitors

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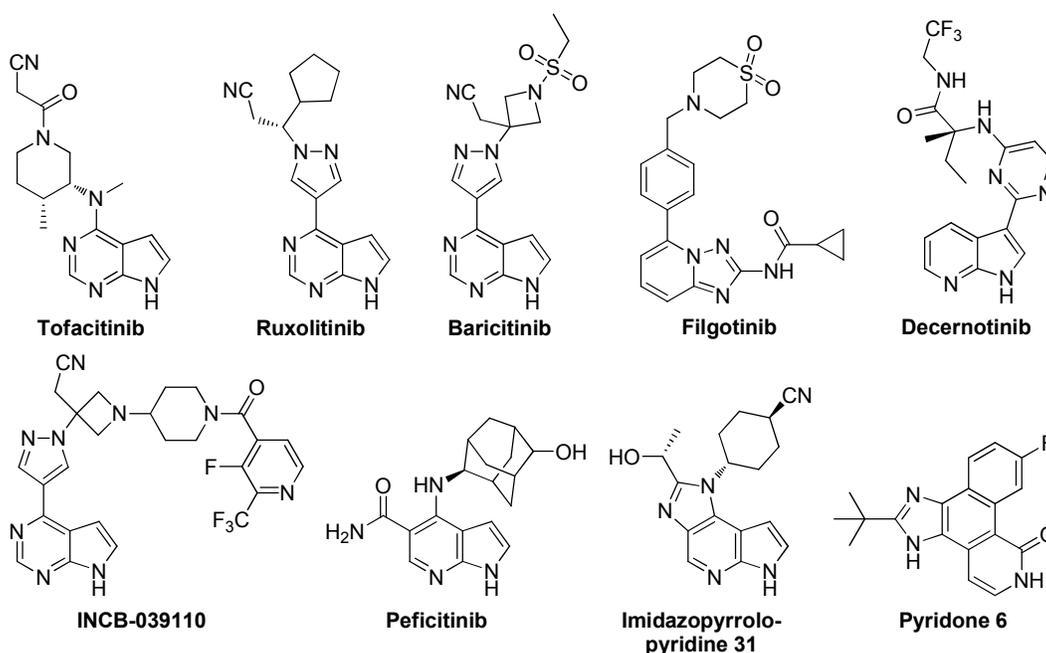
ABSTRACT: The Janus kinase (JAK) family comprises four members (JAK1, JAK2, JAK3, and Tyk2) that play a key role in mediating cytokine receptor signaling. JAK inhibition thus modulates cytokine-mediated effects. In particular, selective inhibition of JAK1 or JAK3 may provide an efficient therapeutic agent for the treatment of inflammatory diseases, with minimized side effects. In this study, as part of our continued efforts to develop a selective JAK1 inhibitor, a series of 1,2-disubstituted benzimidazole-5-carboxamide derivatives was prepared and their inhibitory activities against all four JAK isozymes were evaluated. A clear structure-activity relationship was observed with respect to JAK1 selectivity; this highlighted the importance of hydrogen bond donors at both *N*¹ and *R*₂ positions located within a specific distance from the benzimidazole core. One of the synthesized compounds, 1-(2-aminoethyl)-2-(piperidin-4-yl)-1*H*-benzo[*d*]imidazole-5-carboxamide (**5c**), showed remarkable JAK1 selectivity (63-fold vs JAK2, 25-fold vs JAK3, and 74-fold vs Tyk2). Molecular docking revealed that the 2-aminoethyl and piperidin-4-yl substituents of **5c** function as probes to differentiate the ATP-binding site of JAK1 from that of JAK2, resulting in preferential JAK1 binding. A kinase panel assay confirmed the JAK1 selectivity of **5c**, which showed no appreciable inhibitory activity against 26 other protein kinases at 10 μM.

INTRODUCTION

The cytoplasmic domains of cytokine receptors that mediate the intracellular signaling of type I/II cytokines are associated with specific Janus protein tyrosine kinases (JAK1, JAK2, JAK3, and Tyk2). For example, interleukins (IL) responsible for adaptive immune functions (IL-2, IL-4, IL-7, IL-9, IL-15, and IL-21) specifically bind to cytokine receptors that contain a common gamma chain (γ_c , CD132) and are associated with intracellular JAK3 and JAK1.¹ JAK3 is characterized by its exclusive association with γ_c , and loss-of-function mutation of JAK3 results in complete abrogation of signaling by γ_c -cytokines; this causes severe combined immunodeficiency.² In contrast, JAK1 is widely expressed and, by pairing with JAK2 and/or Tyk2, is involved in a broad range of cytokine signaling to affect several proinflammatory cytokines associated with the innate immune response (γ_c , gp130, type I/II interferon, IL-6, IL-10 subfamily).³ JAK2 is exclusively associated with the receptors for various cytokines and growth fac-

tors (IL-3, IL-5, granulocyte macrophage colony-stimulating factor, erythropoietin, and thrombopoietin) and functions as a homodimer to play an essential role in red blood cell formation.⁴ As a result, JAK2-knockout mice develop anemia⁵, while a unique gain-of-function JAK2 point mutation (V617F) results in myeloproliferative disorders (MPD) in humans.⁶ Taken together, studies of cytokine-mediated signaling have revealed pathologic roles of specific cytokines in inflammatory diseases and MPD, which can be abrogated by inhibition of the associated JAK isozymes.⁷ Consequently, enormous efforts have been devoted to the development of JAK inhibitors, which culminated in the recent approval of two JAK inhibitors, ruxolitinib⁸ and tofacitinib⁹, for the treatment of MPD and rheumatoid arthritis (RA), respectively (Chart 1); ruxolitinib is a JAK1/JAK2-selective inhibitor with reduced potency against JAK3⁸, while tofacitinib is a pan-JAK inhibitor that inhibits JAK1, JAK2, and JAK3 to an approximately equal extent.⁹ However, it is still unclear whether

Chart 1. Structures of the JAK inhibitors

Table 1. JAK₁-selectivity of known inhibitors in enzyme assay

Comps	IC ₅₀ (nM)				Selectivity ^a		
	JAK ₁	JAK ₂	JAK ₃	Tyk2	JAK ₁ /JAK ₂	JAK ₁ /JAK ₃	JAK ₁ /Tyk2
Baricitinib	6	6	>400	53	1	>67	9
Decernotinib	11	13	2.5 ^b	11	1	0.2	1
Filgotinib	10	25	810	116	3	81	12
INCB-039110 ^c	-	-	-	-	>20	>200	-
Peficitinib	3.9	5.0	0.71	4.8	1	0.2	1
ABT-494	-	-	-	-	74 ^d	-	-
Imidazopyrrolopyridine-31	1.9 ^e	-	280 ^e	12 ^e	36	147	6

^aSelectivity in enzyme assay = (IC₅₀ against JAK₂, JAK₃, or Tyk2) / (IC₅₀ against JAK₁). ^bK_i value. ^cUnder development for oncology indications. ^dSelectivity in cellular assays. ^eInhibition constants (K_i's)

inhibition of multiple JAK isozymes is a prerequisite for the clinical efficacy of JAK inhibitors, or if an isozyme-selective inhibitor would show improved efficacy and safety profiles. In this context, it is worth noting that, even though the clinical efficacy of tofacitinib in patients with RA has been attributed to the suppression of JAK₃ activity^{8,10}, a dominant role of JAK₁ over JAK₃ in RA has been strongly proposed¹¹. On the other hand, a recent report by Pfizer stated that inhibition of either JAK₁ or JAK₃ was enough to abrogate cytokine signaling¹². Accordingly, there is a compelling need for the development of selective JAK₁ or JAK₃ inhibitors. Moreover, from a therapeutic point of view, inhibition of JAK₂ should be avoided because the induction of anemia, a dose-limiting side effect of tofacitinib, has been ascribed to the concomitant inhibition of JAK₂¹³.

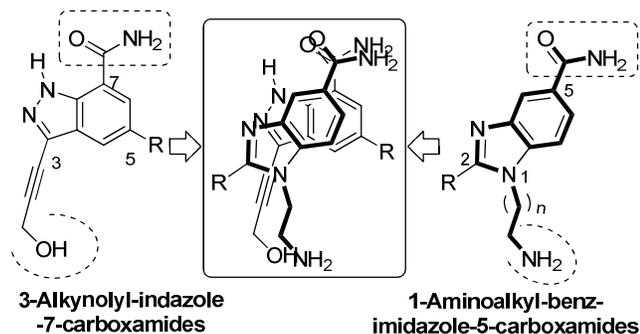
Currently, several JAK inhibitors with different isoform specificity profiles¹⁴⁻²⁰ are in the clinical or preclinical stage for the treatment of autoimmune diseases including

RA (Table 1). Among those, filgotinib¹⁶, INCB-039110¹⁷, ABT-494¹⁹ and imidazopyrrolopyridine-31²⁰ are reported to show JAK₁-selectivity while decernotinib¹⁵ is the best characterized selective JAK₃ inhibitor. In particular, ABT-494, a highly selective JAK₁ inhibitor (74-fold vs. JAK₂), turned out to be effective in RA patients without unfavorable effects on erythropoietin signaling and peripheral NK cell counts.¹⁹ Thus, it is anticipated that the clinical studies with JAK₁ or JAK₃ selective inhibitors will confirm a relationship between the isoform selectivity and side-effect profiles. In this context, further investigation to identify novel scaffolds of JAK₁-selective inhibitors is required.

Recently, our research group reported that 3-alkynyl-indazole-7-carboxamide derivatives with a linear tether, 3-alkynol (dashed circle, Chart 2), which serves as an isozyme-specific probe group, showed selective inhibition of JAK₁ over other JAK isozymes²¹. Based on this observation, the present study developed compounds with a novel

benzimidazole-5-carboxamide scaffold and an aminoalkyl functionality (dashed circle, Chart 2). In addition, to optimize selectivity and potency for the target enzyme, the benzimidazole core structure was further substituted with an alkyl substituent at the 2-position. Here, we describe the synthesis and biological evaluation of a series of these 1-aminoalkyl-2-alkyl-1*H*-benzo[*d*]imidazole-5-carboxamide derivatives, resulting in the identification of a potent JAK1 inhibitor with remarkable JAK1 selectivity.

Chart 2. Design of the title compound



RESULTS AND DISCUSSION

The 1-aminoalkyl-2-alkyl-1*H*-benzo[*d*]imidazole-5-carboxamides (**4** and **5**), were prepared from commercially available 4-fluoro-3-nitrobenzoic acid (**1**) in 5 steps (Scheme 1). First, EDC [1-ethyl-3-(3-dimethylaminopropyl)carbodiimide] coupling of **1** with 1-hydroxy-1*H*-benzotriazole (HOBT)-ammonium salt ($\text{NH}_3 \cdot \text{HOBT}$) in a 4:1 mixture of CH_3CN and *N,N*-dimethylformamide (DMF) yielded the corresponding benzamide, which underwent nucleophilic aromatic substitution by 1,3-diaminopropane or ethylenediamine followed by *N*-Boc protection, to give the corresponding 4-alkylamino-3-nitrobenzamide (**2a** or **2b**), with 75% or 79% yield. The nitro functionality of **2a** or **2b** was then reduced to give the key intermediate, 4-alkylamino-3-aminobenzamide (**3a** or **3b**), with 67% or 91% yield. These diaminobenzamides were then cyclized to the corresponding benzimidazole-5-carboxamides by treatment with the appropriate aldehydes in the presence of sodium hydrogen sulfite (NaHSO_3)²² with 10–67% yields. Finally, deprotection of the Boc groups provided the desired compounds (**4** and **5**).

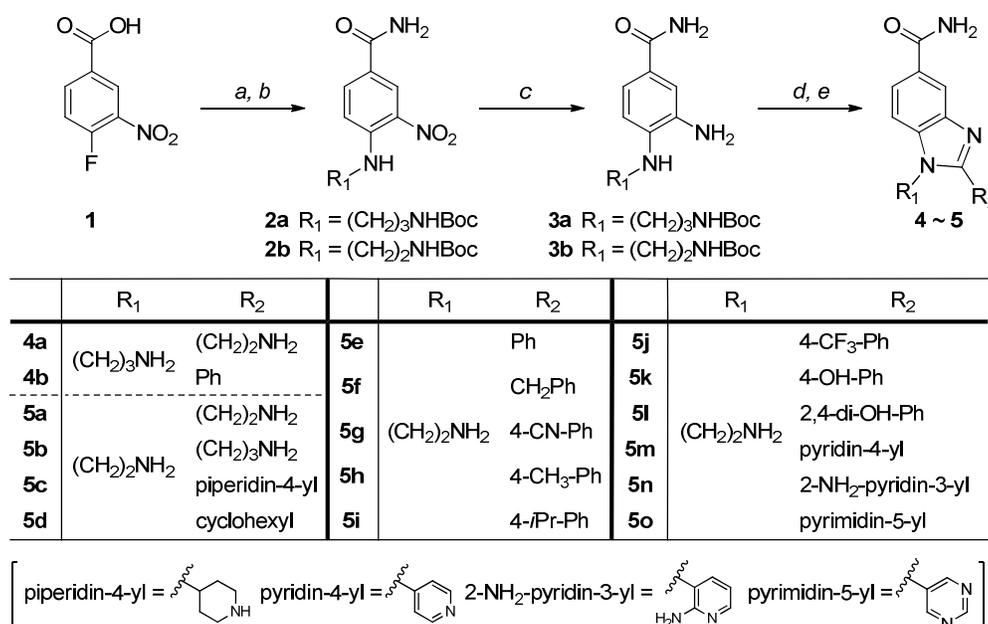
Z'-LYTE™ Kinase Assay Kits (Invitrogen) were used to determine the *in vitro* inhibitory activity of the benzimidazole derivatives against the JAK isozymes; the Tyr 6 peptide was used to analyze JAK1 and JAK3 effects and the Tyr 3 peptide was used for Tyk2. Pyridone-6 (Chart 1)²³, a pan-JAK inhibitor which is used for quality control of the assay kits, was used as a positive control. Assays were performed at the ATP K_m and the IC_{50} values of the synthesized compounds against each JAK isozyme, as well as their JAK1-selectivities, are summarized in Table 2. The benzimidazole derivatives with a 3-aminopropyl group substituted at the *N* position (**4a** and **4b**) showed only moderate inhibitory activity against JAK1 (IC_{50} = 6.1 and

11.3 μM , respectively) and JAK3 (IC_{50} = 1.5 and 1.8 μM , respectively). Surprisingly, however, shortening the 3-aminopropyl chain of **4a** by one carbon potentiated the JAK1-inhibitory activity of the resulting benzimidazole derivative **5a** by more than 60-fold (IC_{50} = 0.1 μM) without altering inhibition of other JAK isozymes. As a result, the 2-aminoethyl-substituted benzimidazole derivative **5a** showed high JAK1 selectivity over other JAK isozymes (29-fold vs JAK2, 15-fold vs JAK3, and 31-fold vs Tyk2). Based on this observation, the *N* substituent was fixed as a 2-aminoethyl group and a series of benzimidazole derivatives with various R_2 substituents was prepared (**5b–5o**). Within the series, the R_2 -alkyl derivatives (**5a–5d**) showed a clear structure-activity relationship whereby an amino group located within a specific distance from the benzimidazole core played a key role in maintaining potent JAK1-inhibitory activity. Thus, compounds with 2-aminoethyl (**5a**) or piperidin-4-yl (**5c**) functionalities at R_2 showed potent inhibitory activity against JAK1, with IC_{50} values of 0.1 and 0.05 μM , respectively, while derivatives with a longer R_2 -alkylamino group [**5b**, R_2 = $(\text{CH}_2)_3\text{NH}_2$] or with an R_2 -hydrocarbon substituent [**5d**, R_2 = cyclohexyl] were significantly less potent. On the other hand, the R_2 -alkyl substituents did not affect the inhibitory activities of the corresponding benzimidazole derivatives (**5a–5d**) against other JAK isozymes (JAK2, JAK3, and Tyk2). Taken together, the R_2 -dependence of the inhibitory activities of these benzimidazole derivatives was specific to JAK1, and compound **5c**, with a piperidin-4-yl group at R_2 , showed remarkable JAK1 selectivity (63-fold vs JAK2, 25-fold vs JAK3, and 74-fold vs Tyk2). On the other hand, among the benzimidazole derivatives with an aromatic substituent at R_2 (**5e–5o**), only **5k** (R_2 = 4-OH-Ph) with a hydroxyl group at the 4-position of the aromatic group showed JAK1-selective inhibitory activity (11-fold vs JAK2, 20-fold vs JAK3, and 26-fold vs Tyk2). Based on these observations, common structural features within the set of JAK1-selective inhibitors (**5a**, **5c**, and **5k**) could be identified; an alkyl chain bearing a hydrogen bond donor ($-\text{NH}$ or $-\text{OH}$) is substituted at the 1-position of the benzimidazole core.

In a separate experiment, the inhibitory activities of **5c** against the JAK isozymes were measured in the presence of 1 mM ATP (Table 3), instead of at the K_m ^{12,24}, in order to mimic the physiological level of ATP. As expected^{12,24}, the observed IC_{50} values for **5c** were 114–205-fold higher in the presence of 1 mM ATP than those obtained at the K_m values (Table 2); these data shifts were smallest for JAK1, which has the highest K_m for ATP. As a result, the JAK1 selectivity of **5c** became more prominent under these physiologically relevant assay conditions.

The JAK1 selectivity of this benzimidazole derivative was then investigated using a molecular docking study. The most potent JAK1-selective inhibitor, **5c**, was docked into the ATP-binding site of JAK1 (PDB ID = 4EHZ)²⁰ or JAK2 (PDB ID = 4F09)²⁰ using the flexible ligand docking software, Glide, incorporated into the Schrödinger molecular modeling software suite; the best docking poses are shown in Figure 1. The ATP-binding sites of JAK1 and JAK2 are differentiated by two amino acids, one in the

Scheme 1. Synthesis of the benzimidazole derivatives (4 and 5).



Reagent & Conditions : (a) EDC, NH₃-HOBT, DMF/CH₃CN; (b) R₁NH₂, DIPEA, *i*-PrOH, 80 °C; (c) H₂, Pd/C, MeOH; (d) R₂CHO, NaHSO₃, MeOH, reflux (e) HCl, MeOH

Table 2. Inhibitory activities of the benzimidazole derivatives for the indicated enzymes

Compds	IC ₅₀ (μM) ^a				Selectivity ^b		
	JAK ₁	JAK ₂	JAK ₃	Tyk ₂	JAK ₁ /JAK ₂	JAK ₁ /JAK ₃	JAK ₁ /Tyk ₂
4a	6.1±0.8	2.2±0.3	1.5±0.1	6.4±0.9	0.4	0.2	1
4b	11.3±0.8	20.5±1.2	1.8±0.3	2.7±0.1	2	0.2	0.2
5a	0.1±0.01	2.9±1.3	1.5±0.3	3.1±0.2	29	15	31
5b	2.2±0.3	1.6±0.2	1.1±0.1	2.5±0.2	0.7	0.5	1
5c	0.05±0.01	3.15±0.2	1.26±0.1	3.7±0.3	63	25	74
5d	4.3±0.3	4.1±0.2	2.5±0.1	3.3±0.2	1	0.6	0.8
5e	11.6±1.2	2.8±0.3	14.1±0.6	4.2±0.2	0.2	1	0.4
5f	18.2±0.7	2.7±0.3	2.1±0.3	7.5±0.2	0.1	0.1	0.4
5g	2.9±0.2	2.6±0.3	1.9±0.3	2.5±0.1	0.9	0.7	0.9
5h	3.7±0.2	2.6±0.1	2.3±0.1	3.5±0.3	0.7	0.6	0.9
5i	3.1±0.1	5.9±0.5	1.9±0.2	2.7±0.2	2	0.6	0.9
5j	9.3±0.5	4.8±0.2	2.1±0.2	3.2±0.3	0.5	0.2	0.3
5k	0.1±0.0	1.3±0.2	2.4±0.1	3.2±0.1	11	20	26
5l	3.1±0.3	2.5±0.2	2.9±0.3	3.7±0.1	0.8	0.9	1
5m	32.4±1.4	13.5±0.9	10.5±0.4	5.8±0.2	0.4	0.3	0.2
5n	8.1±0.4	4.7±0.6	1.9±0.2	4.3±0.5	0.6	0.2	0.5
5o	3.4±0.1	1.6±0.2	2.5±0.1	2.8±0.3	0.5	0.7	0.8
Pyridone-6^c	0.004	0.001	0.01	0.005	2	0.2	0.2

The ATP concentration (K_m) was 87 μM for JAK₁, 35 μM for JAK₂, 16 μM for JAK₃, and 25 μM for Tyk₂. ^aThe means ± SD of three independent experiments are shown. ^bSelectivity = (IC₅₀ against JAK₂, JAK₃, or Tyk₂) / (IC₅₀ against JAK₁). ^cPan-JAK inhibitor²³ used as a positive control in this study.

αD-helix (Glu966 in JAK₁, Asp939 in JAK₂) and the other in the glycine-rich loop (His885 in JAK₁, Asn859 in JAK₂)

(Figure 1). Comparison of these two pairs of amino acid residues (Glu966 vs Asp939, His885 vs Asn859) revealed

Table 3. Inhibitory activity of **5c** against JAK isozymes in the presence of 1 mM ATP.

Compds	IC ₅₀ (μM) ^a				Selectivity ^b		
	JAK1	JAK2	JAK3	Tyk2	JAK1/JAK2	JAK1/JAK3	JAK1/Tyk2
5c	5.7±0.8	437.3±3.1	232.5±7.5	759.5±5.6	76	41	133
Pyridone-6	0.6±0.1	1.4±0.3	0.9±0.2	1.7±0.3	2	1	3

^aThe means ± SD of three independent experiments. ^bSelectivity = (IC₅₀ against JAK2, JAK3, or Tyk2) / (IC₅₀ against JAK1).

that they had different side chain lengths. Thus, amino acids with longer side chains, Glu966 (JAK1) and Asn859 (JAK2), protrude into the ATP-binding site, which results in constriction of ligand binding in the “upper” portion of JAK1 (Glu966–Leu881, 5.2 Å, Figure 1a) and the “lower” portion of JAK2 (Arg980–Asn859, 4.1 Å, Figure 1b).

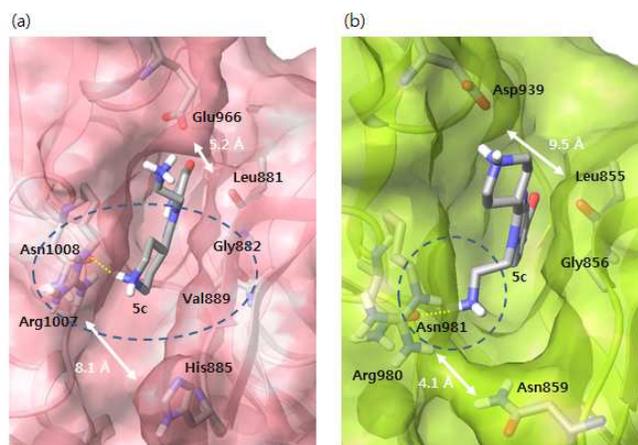


Figure 1. Docked structures of **5c** to (a) JAK1 (PDB ID: 4EHZ) and (b) JAK2 (PDB ID: 4F09). Dotted lines show hydrogen bonds. Hydrophobic contacts between **5c** and the ATP-binding sites are shown as dashed circles. Block arrows indicate the widths of the ATP-binding sites.

Interestingly, the best docking poses of **5c** to JAK1 and JAK2 showed that the more sterically demanding piperidin-4-yl moiety of **5c** was positioned away from the constricted area. The binding mode of **5c** to JAK1 is characterized by positioning the piperidin-4-yl group at the “lower” open side of the ATP-binding pocket (Figure 1a), forming a hydrogen bond with Asn1008 and multiple favorable contacts with Arg1007, Gly882, Val889, and Asp1021. In addition, the smaller and charged 2-aminoethyl group is placed near the polar side chain of Glu966. On the other hand, upon binding to JAK2, the benzimidazole moiety of **5c** rotates about the aryl-carbonyl bond (~190.4°) to locate the 2-aminoethyl and piperidin-4-yl substituents at the opposite side of the ATP-binding pocket (Figure 1b); the piperidin-4-yl group, at the upper side of the ATP-binding pocket, is located between Asp939 and Leu855, while the 2-aminoethyl functionality forms a hydrogen bond to Asn981 at the bottom of the ATP-binding site. Overall, the binding mode of **5c** to JAK2 is less favorable because of the lack of hydrophobic contacts around the 2-aminoethyl substituent, as well as the positioning of the nonpolar hydrocarbons in the piperidin-4-yl ring between the polar side chain of Asp939 and the backbone carbonyl group of Leu855. The preference of **5c** for JAK1 was re-

flected by the docking scores (G-score; -8.95 for JAK1 and -7.75 for JAK2). Taken together, this molecular docking information clearly showed that the 2-aminoethyl and piperidin-4-yl moieties (attached to the benzimidazole core of **5c**) differentiate the ATP-binding sites of JAK1 and JAK2; **5c** preferentially binds to the ATP-binding site of JAK1 by positioning the bulkier piperidin-4-yl substituent in the open space and forming multiple favorable contacts with the enzyme. The commercial KinaseProfiler™ Service (Eurofins Scientific, Inc.) was utilized to evaluate the selectivity of compound **5c** against a panel of 27 kinases (Figure 2). KinaseProfiler™ assay protocols measure the percent inhibition of phosphorylation of a peptide substrate in the presence of fixed concentrations of ATP that were close to the *K_m* for each kinase (Figure 2). Except JAK1, no appreciable inhibition of these protein kinases was observed in the presence of 10 μM **5c**, with most maintaining >65% of their control activity.

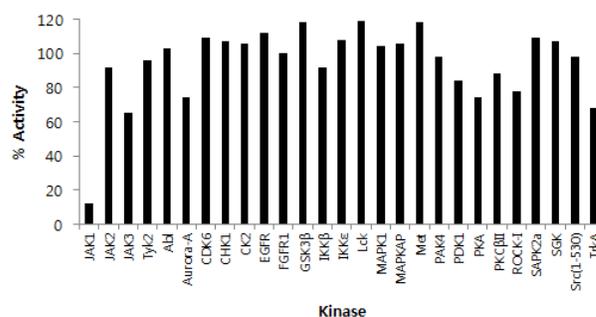


Figure 2. Kinase panel assays were performed in the presence of **5c** (10 μM) and ATP (*K_m* concentration) using Eurofins KinaseProfiler™ service. The y-axis represents percentage of remaining kinase activity (% Activity).

However, due to high polarity, the JAK1-selective inhibitor **5c** showed low PAMPA permeability (*P_e* = 0.1 × 10⁻⁶ cm/s), lack of activity in cell-based assay²⁵ (Supporting Information) and poor oral bioavailability (0.7%) in rat. On the other hand, **5c** demonstrated generally favorable pharmacokinetic properties: following intravenous (i.v.) administration to rats, **5c** exhibited moderate volumes of distribution (*V_{ss}*, 0.5 l/kg) and low systemic plasma clearance (CL, 0.4 l/hr/kg) resulting in plasma half-life (*t_{1/2}*) of 9.1 h (Table S1, Supporting Information). In light of the unsatisfactory pharmacokinetic properties, the JAK1-selective 1,2-disubstituted benzimidazole-5-carboxamide scaffold needs further optimization.

CONCLUSIONS

Since each JAK isozyme is employed in various cytokine pathways regulating immune function, specific inhibition of a single JAK isozyme can provide a beneficial therapeutic option in patients with immunologic disorders such as RA. Pfizer's pan-JAK inhibitor, tofacitinib, was recently approved for the treatment of RA, and its clinical efficacy has been attributed to the suppression of JAK3 activity. However, since both JAK1 and JAK3 are associated with γ_c cytokine receptors, it is unclear whether the clinical efficacy of tofacitinib results from its concomitant inhibition of JAK3 and JAK1, or selective inhibition of one JAK isozyme. The development of selective JAK1 or JAK3 inhibitors, and investigation of their effects, is therefore required in order to address this issue. For this purpose, we set out to develop JAK1-selective inhibitors, based on our previous observations that a linear alkyl tether can function as a probe to differentiate the ATP-binding site of JAK1 from those of the other JAK isozymes. We synthesized a series of 1,2-disubstituted benzimidazole derivatives and evaluated their inhibitory activities against all four JAK isozymes. These benzimidazole derivatives showed clear structure-activity relationships that confirmed the importance of hydrogen bond donors at both N^1 and R_2 positions for JAK1 selectivity; a 2-aminoethyl group and an alkyl chain bearing a hydrogen bond donor were favored at N^1 and R_2 , respectively. Compound **5c**, with a piperidin-4-yl group at R_2 , showed unprecedented JAK1 selectivity (63-fold vs JAK2, 25-fold vs JAK3, and 74-fold vs Tyk2). A molecular docking study revealed that **5c** preferentially bound to the ATP-binding site of JAK1 by positioning the bulkier piperidin-4-yl substituent at the open space and by forming multiple favorable contacts with the enzyme. A kinase panel assay also showed that compound **5c** (10 μ M) did not inhibit a panel of 26 other protein kinases. The JAK1-selective benzimidazole derivatives identified in this study thus have the potential to elucidate the biological consequences of specific JAK1 inhibition, specifically in terms of therapeutic intervention for RA.

EXPERIMENTAL SECTION

General procedure for the preparation of 3-amino-4-alkylaminobenzamide (3a and 3b). Palladium on charcoal (10% w/w) was added to a solution of **2** (1.4 mmol) in MeOH (15 ml) and charged with hydrogen (balloon, 1 atm). After stirring for 8 h at room temperature, solvent was removed under reduced pressure to give a crude product which was purified by column chromatography (SiO_2 , CH_2Cl_2 :MeOH: NH_4OH :water = 60:35:5:5) to give **3**.

General procedure for the preparation of benzimidazole derivatives 4 and 5. Preparation of 2-(2-aminoethyl)-1-(3-aminopropyl)-1*H*-benzo[d]imidazole-5-carboxamide (**4a**) is provided as a representative example. A solution of **3a** (22 mg, 0.07 mmol) in MeOH (3 ml) was added to *tert*-butyl (3-oxopropyl)carbamate (14 mg, 0.08 mmol) and sodium bisulfate (13 mg, 0.1 mmol). After stirring for 1 h at 60°C, inorganic material was collected by filtration and rinsing with MeOH. The filtrate was concentrated under reduced pressure to give a crude product,

which was purified by column chromatography (SiO_2 , CH_2Cl_2 :MeOH: NH_4OH :water = 80:20:1:1) to give the benzoimidazole-derivative as a dark yellow powder. This benzoimidazole-derivative was then dissolved in MeOH (3 ml) and treated with 6 N HCl (0.1 ml). The reaction mixture was stirred at room temperature for 8 h and then concentrated under reduced pressure. The crude product was purified by column chromatography (SiO_2 , CH_2Cl_2 :MeOH: NH_4OH :water = 60:35:5:5) to give **4a** (11 mg, 60% yield) as a yellow powder.

ASSOCIATED CONTENT

Supporting Information

General experimental procedures, ^1H and ^{13}C nuclear magnetic resonance (NMR) spectra for new compounds, and methods for the molecular docking study. This material is available free of charge via the Internet at <http://pubs.acs.org>.

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Author Contributions

All authors have given approval to the final version of the manuscript.

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ABBREVIATIONS

JAK, Janus kinase; IL, interleukin; MPD, myeloproliferative disorders; RA, rheumatoid arthritis; EDC, 1-ethyl-3-(3-dimethylaminopropyl)carbodiimide; HOBT, 1-hydroxy-1*H*-benzotriazole; DMF, *N,N*-dimethylformamide; Boc, *tert*-butoxycarbonyl; EtOAc, ethyl acetate; MeOH, methanol; PAMPA, parallel artificial membrane permeability assay.

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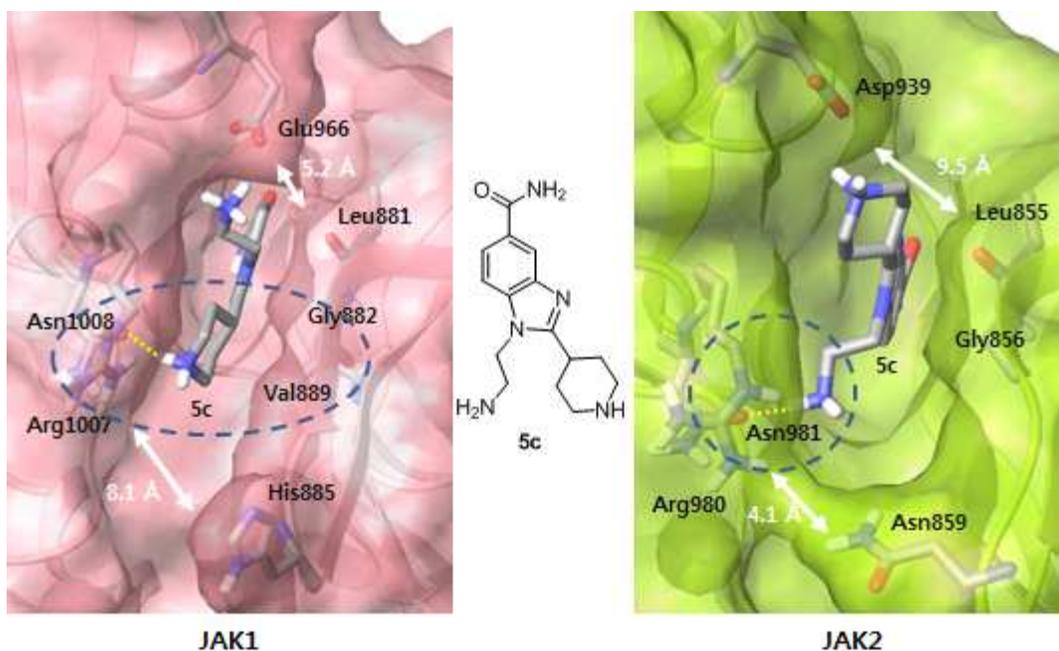
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JAK1

JAK2

[ATP]	IC ₅₀ (μM)				Selectivity		
	JAK1	JAK2	JAK3	Tyk2	JAK2	JAK3	Tyk2
<i>K_m</i>	0.05±0.01	3.2±0.2	1.3±0.1	3.7±0.3	63	25	74
1 mM	5.7±0.8	437.3±3.1	232.5±7.5	759.5±5.6	76	41	133