Hybrid-Designed Inhibitors of p38 MAP Kinase Utilizing *N*-Arylpyridazinones

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Abstract: Imidazo[1,2-*a*]pyridyl *N*-arylpyridazinones were hybridized from the classic pyridinylimidazoles and the more recent dual hydrogen bond acceptors, resulting in a new structural class of selective p38 MAP kinase inhibitors.

Introduction. TNF α is one of several cytokines that plays a significant role in the development of an acute or chronic inflammatory response. The success of the soluble TNFa receptor fusion protein Enbrel (etaner- $(ept)^1$ and the monoclonal TNF α antibody Remicade (infliximab)² in the treatment of rheumatoid arthritis and Crohn's disease has provided a proof of concept for the treatment of these autoimmune diseases. These biologics are generally well tolerated to date but have drawbacks related to patient cost, efficiency of production, and administration by injection. Therefore, an orally active small-molecule drug that blocks or modulates circulating $TNF\alpha$ remains an attractive therapy. Of the different approaches toward this end, inhibition of p38 MAP kinase results in the suppression of not only TNF α but also IL-1 β , another significant proinflammatory cytokine.³

The biosynthesis of TNF α and IL-1 β occurs predominately in activated monocytes and macrophages via an intracellular signaling cascade that involves the dual phosphorylation (via MKK3 and MKK6) of Thr180 and Tyr182 within a TGY motif of p38 kinase. Phosphorylated p38 subsequently phosphorylates a variety of substrates, including kinases and transcription factors.⁴ Two of these substrates that are activated by p38 are MAPKAP kinases 2 and 3, which in turn phosphorylate heat shock protein HSP 27, ultimately leading to gene transcription and protein translation. This cascade of events can be initiated by a wide variety of extracellular stimuli or stresses such as endotoxin bacterial lipopolysaccharide and cytokines, or osmotic shock, heat shock, UV light, ionizing radiation, and oxidative stress. Four isozymes of p38 have been cloned and characterized to include the ubiquitously expressed $p38\alpha$ and p38 β ; p38 γ is primarily expressed in skeletal muscle, and p38 δ is highly distributed within lung, kidney, endocrine glandular, and small intestinal tissues.⁵

Historically, SmithKline Beecham demonstrated early on that an orally active small molecule, SB203580 (1), could reduce $TNF\alpha$ levels in vivo, validating the pyridinylimidazole structural class.⁶ Significant improvements to the kinase selectivity and whole blood potency of 1 were subsequently accomplished at Merck with the discovery of the (S)-sec-phenethylamine moiety represented in the second-generation pyridinylimidazole 2.7 Within a few years, Vertex demonstrated that even greater kinase selectivity could be obtained with VX-745 (3).8 Human clinical trials with 3 have been reported,⁹ reviewed,¹⁰ and claimed to have achieved proof of concept, although this press release lacks scientifically reviewed data.¹¹ In an effort to increase the whole blood potency of 3, chemists at Merck designed inhibitors such as 4 that retained the structural attributes of 3 in addition to the basic saturated heterocycle present in 2.12

Design. The modeling view in Figure 2 is based on the X-ray crystal structure of $p38\alpha$ in complex with the soaked inhibitor **4**.¹² Hybrid **6** was energy-minimized outside the protein and then docked into the enzyme active site using the binding orientation of **4** as a guide.¹³ The carbonyl oxygen of the urea moiety of **4** and that of the pyridazinone ring of **6** are able to accept two hydrogen bonds from the backbone amide NHs of Met109 and Gly110. The 2-chloro-4-fluorophenyl ring of **6** appears to superimpose quite well with the 2,4-difluorophenyl ring of **4** near Thr106.¹⁴ However, the tolyl moiety of **6** is situated in the binding pocket near Val30 differently from the 2,6-dichlorophenyl ring of **4**.

Within the MAP kinase family, the JNK Ser/Thr protein kinases bear high homology to p38.¹⁵ Crystal structure data reveal that inhibitors with a dual hydrogen bond motif at Met109 and Gly110 of p38 α induce a peptide flip at Gly110 (Figure 2). Since JNK1–3 kinases contain an aspartic acid at a position equivalent to the position of Gly110 in p38 α , the Ramachandran space of this Asp residue is much more restricted. This rotational restriction could result in a hindered peptide flip, an inefficient dual hydrogen bond, and reduced affinity for these inhibitors.¹⁶ VX-745 (**3**) was reported to be more than 1000-fold selective for p38 α over JNK kinases.¹⁷ Since one of the closest homologues to p38 α kinase is JNK2 β 2, we counterscreened this JNK isozyme for inhibitor selectivity.

This study describes a new class of p38 inhibitors structurally hybridized from the prototypical pyridinylimidazoles and the dual hydrogen bond acceptors represented by **3** and **4**. The pyridazinone moiety¹⁸ was selected as a dual hydrogen bond acceptor because of its diversity based on available arylhydrazines. The imidazo[1,2-*a*]pyridine core¹⁹ was chosen for its basicity to suppress serum albumin binding.²⁰ On the basis of this design, compounds such as **5** and **6** were anticipated to be highly selective for p38 over JNK kinases and more potent in whole blood relative to VX-745 (**3**).

Chemistry. One highly convergent approach to the synthesis of these molecules is shown in Scheme 1. Intermediate **7** was prepared²¹ and then selectively cyclized to the energetically preferred six-membered



Figure 1. Selected p38 inhibitors.



Figure 2. Modeling view of 6 (yellow) overlaid on the crystal structure of $p38\alpha$ MAP kinase in complex with 4 (magenta).

Scheme 1^a



 a (a) LiHMDS, succinic anhydride, THF, $-78~^{\rm C}{\rm C}$ \rightarrow 23 °C, 2 h, 20%; (b) 2-chlorophenylhydrazine hydrochloride, NaOAc, HOAc, 110 °C, 2 h, 50%; (c) n-Bu₄NBr₃, 2:1 CH₂Cl₂/CCl₄, 0 °C \rightarrow 23 °C, 4 h, 99%.

ring dihydropyridazinone. After bromination to form **8**, a Chichibabin based imidazopyridine synthesis was envisioned using 2-aminopyridine. This reaction failed repeatedly apparently because of the absence of any microscopic tautomerization of **8** to the required α -bromoketone.

Subsequently a more linear synthesis was developed that provided the targeted imidazopyridyl *N*-arylpyridazinones (Scheme 2). Readily available benzoyl acetates were brominated under ionic conditions, and the α -monobromides were subjected to the Chichibabin based imidazopyridine synthesis using 2-aminopyridine to form haloaryl analogues of **9**.²² The methyl esters were converted to the Weinreb amides under basic conditions,²³ followed by methyl ketone generation to

Scheme 2^a



^{*a*} (a) *n*·Bu₄NBr₃, CH₂Cl₂, 0 °C → 23 °C, 3 h; (b) 2-aminopyridine, EtOH, 60 °C, 14 h, ~70% (two steps); (c) Me(MeO)NH/HCl, *i*·PrMgCl, THF, −10 °C, 1 h, ~74%; (d) MeMgBr, THF, 0 °C, 30 min, ~97%; (e) LiHMDS, BrCH₂CO₂*t*·Bu, THF, −78 °C → 23 °C, 2 h; (f) TFA, CH₂Cl₂, 0 °C → 23 °C, 4 h, then HCl_g, MeOH, 0 °C, 1 h, ~40% (two steps); (g) arylhydrazine hydrochloride, NaOAc, HOAc, H₂O, 130 °C, 20 h, ~30%; (h) CuCl₂, CH₃CN, 85 °C, 72 h, ~50%.

form **10** in good overall yield from the starting benzoyl acetates. The methyl ketones (**10**) were then homologated to the γ -ketoesters (**11**) in modest yield, followed by cyclization with several different phenylhydrazines to form the *N*-aryldihydropyridazinones. Oxidation proved to be extremely slow and often incomplete, yet absent of any byproducts to provide the desired pyridazinones (**12**).

In Vitro SAR. The biological data for the *p*-fluoro derivative **13** represent a baseline for SAR in the *N*-(2,6-dichlorophenyl)pyridazinone series (**A**, Table 1). Unlike pyridinyl imidazole **2**, the *m*-trifluoromethyl group was not tolerated in **14** at the level of the enzyme, indicating a different binding orientation from **2**. This result is consistent with the activity of the dual hydrogen bond acceptor inhibitors relative to the pyridinylimidazoles. Likewise the *o*,*m*-dichloro analogues **15** and **16** displayed poor enzyme inhibition. As with **4**,¹² combined substitution at the ortho and para positions was ideal for enzyme affinity and is exemplified by the 2-chloro-4-fluoroaryl analogues **17** and **18**.

To reduce the lipophilicity of **17** and **18**, the 2,4difluoroaryl analogues **5** and **19** were prepared.²⁴ As anticipated, the IC_{50} shift from enzyme inhibition to THP-1 cellular activity was reduced with **5** relative to **17**. However, the oxidation of **5** to **19** resulted in an overall decrease in activity observed not only in whole blood but also in the p38 inhibition assay.

Another approach to reduce the lipophilicity of **17** and **18** was to remove one chlorine atom and prepare the N-(2-chlorophenyl)pyridazinone series (**B**, Table 1). Although the IC₅₀ shift between enzyme affinity and THP-1 activity was attenuated for compounds **20** and **21** relative to **17** and **18**, a corresponding reduction in p38 affinity was also observed, resulting in THP-1 activity similar to that of **17** and **18**. The *m*-trifluoromethylaryl analogues **22** and **23** displayed weak activity similar to that of **14**.

On the basis of the patent literature²⁵ and the SAR of **4**, the *N*-(2-chlorophenyl)pyridazinone (**B**) binding subunit was replaced with an *N*-(2-tolyl)pyridazinone moiety (**C**, Table 1). Although a reduction in lipophilicity was anticipated, a potential reduction in p38 affinity relative to the *N*-(2,6-dichlorophenyl)pyridazinone series







			IC ₅₀ (nM)						
compd	R	bond order	p38 α ^b	p38β ^b	$JNK2\beta 2^{c}$	THP-1 ^{d}	hWB ^e		
(A) N-(2,6-Dichlorophenyl)pyridazinone Analogues									
13	4-F	2	49	153	40% @ 10000	70	60% @ 4000		
14	$3-CF_3$	2	1000	50% @ 5000	0% @ 10000	41% @ 4000			
15	2,3-diCl	1	2000	57% @ 5000	26% @ 5000	59% @ 4000			
16	2,3-diCl	2	270	890	14% @ 5000	65% @ 4000			
17	2-Cl-4-F	1	42	670	0% @ 10000	180			
18	2-Cl-4-F	2	14	64	23% @ 10000	161			
5	2,4-diF	1	8	63	22% @ 10000	10	205		
19	2,4-diF	2	22	112	11% @ 5000	89% @ 4000	1000		
(B) N-(2-Chlorophenyl)pyridazinone Analogues									
20	2-Cl-4-F	1	208	2600	0% @ 10000	400			
21	2-Cl-4-F	2	76	171	0% @ 10000	161			
22	$3-CF_3$	1	171	355	0% @ 10000	72% @ 4000			
23	3-CF ₃	2	344			48% @ 4000			
(C) N-(2-Tolyl)pyridazinone Analogues									
24	2,3-diCl	1	36% @ 100	33% @ 100	0% @ 1000	170			
25	2,3-diCl	2	22	44	15% @ 10000	6	500		
26	2-Cl-4-F	1	14	94	4% @ 5000	22	220		
6	2-Cl-4-F	2	9	28	16% @ 5000	20	100		
27	2,4-diF	1	8	78	4% @ 10000	14	440		
28	2,4-diF	2	26	140	15% @ 5000	68	870		

^{*a*} Values are an average of two or greater individual assays. On average, repeat determinations differed by ±20%. ^{*b*} Filter plate binding of compound versus [γ -³³P]ATP in active recombinant murine FLAG-p38 fusion protein (GST-ATF2).⁷ ^{*c*} HTRF detection of GST-ATF2 phosphorylated by recombinant human flag-tagged JNK2 β 2 fusion protein. ^{*d*} LPS-induced human TNF α inhibition detected by ELISA in THP-1 cell supernatants. ^{*e*} LPS-induced human TNF α inhibition in human whole blood quantified by immunoassay.⁷

(A) was a concern. In practice, the 2-tolyl analogues **24**–**26** and **6** were more active in functional assays and enzyme inhibition relative to the corresponding 2,6-dichlorophenyl analogues **15–18**. Analogues **26** and **6** were also superior to the related 2-chlorophenyl derivatives **20** and **21**.

For in the 2-tolyl series, attention was turned to the 5- to 10-fold IC₅₀ shift observed between the THP-1 and whole blood assays of **6** and **26**. The 2,4-difluoroaryl analogues were targeted again, this time toward the preparation of **27** and **28**. Once again, a reduction in affinity and functional activity was observed between the 2,4-difluoroaryl analogues **27** and **28**, similar to that between the 2,4-difluoroaryl analogues **5** and **19**.²⁶ Last, the p38 inhibitors described in Table 1 show consistent p38 α selectivity relative to JNK2 β 2. The most active hybrids, **5** and **6**, demonstrated >1000-fold and >500-fold kinase selectivity, respectively.

PD Profile and in Vitro Metabolism of Compound 6. After the in vitro data shown in Table 1 were obtained, analogue **6** was further evaluated against VX-745 (**3**) in a murine MAPKAP pharmacodynamic (PD) assay (Table 2). In this study, both **3** and **6** were each dosed ip at 3 mg/kg in five mice, resulting in mMAPKAP inhibition of 29% and 42%, respectively. The mMAPKAP inhibition with **6** was statistically significant (p < 0.05) relative to vehicle. Importantly, the compound exposure in mice was different for **3** relative to **6**. Plasma concentrations of 1743 and 810 nM for **3** and **6**,

Table 2. Biological Profiles for 3	and
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	in vi	itro IC ₅₀	(nM)	ex vivo mMAPKAP ^a		
compd	p38α	THP-1	hWB	% inhibition	plasma concn (nM)	
3	22	150	700	29	1734	
6	9	20	100	42	810	

^{*a*} BALB/c mice **3** (n = 5) and **6** (n = 5) were dosed ip at 3 mg/kg in a vehicle of DMSO/EtOH/PEG400/saline (10:10:60:20). Blood was collected for MAPKAP inhibition and plasma concentration at 1 h after dosing.

respectively, were measured from the same experiment by an ex vivo p38 kinase inhibition assay.²⁷

An additional comparison of **3** with **6** regarding in vitro metabolism served to correlate these plasma concentrations with microsomal stability. Both **3** and **6** were incubated with mouse liver microsomes, resulting in a respective 46% and 9% compound presence after 10 min (Supporting Information). From both this microsomal data and the in vivo plasma concentrations, it is apparent that **6** is more readily metabolized in the mouse relative to **3**, yet **6** appears to be equally, if not slightly more, efficacious relative to **3** in the mMAPKAP PD assay. This result is consistent with the greater in vitro activity of **6** relative to **3** (Table 2).

Conclusion. In summary, a new structural class of p38 kinase inhibitors was designed based ultimately on SB203580 and VX-745. Representative hybrids include imidazopyridyl *N*-arylpyridazinones **5** and **6**, both of which demonstrated low nanomolar enzyme inhibition, ≥ 1000 -fold p38 α /JNK2 β 2 kinase selectivity, and efficacy

in human whole blood. Analogue **6** was evaluated ex vivo in mice, providing 42% inhibition of mMAPKAP activity and further validating this class of p38 inhibitors. Relative to the clinically used VX-745 (**3**), analogue **6** displayed similar efficacy in this pharmacodynamic mouse model.

Supporting Information Available: Methods for molecular modeling, experimental procedures for compound preparation and characterization data, biological assay protocols, and liver microsome stability plots. This material is available free of charge via the Internet at http://pubs.acs.org.

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- (27) Blood plasma concentrations of **3** and **6** were calculated by bioassay. The free compound p38 inhibition IC_{50} is multiplied by 1 over the dilution factor of the plasma required to reach 50% inhibition in the p38 assay (EC₅₀). This method does not address active metabolites.

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