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2-SUBSTITUTED-4-METHOXYBENZIMIDAZOLE-BASED PDE4 INHIBITORS

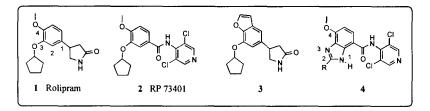
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Abstract: A new family of PDE4 inhibitors based on a benzimidazole framework is described. Several of these compounds are orally bioavailable and show efficacy in in vivo models of inflammatory disease. © 1998 Elsevier Science Ltd. All rights reserved.

Cyclic nucleotide phosphodiesterases (PDE1-7) are a family of enzymes that degrade the intracellular secondary messengers cAMP and cGMP¹ thereby limiting a host cell's response to extracellular signals mediated by these carriers. In a number of immunological cell types, elevation of cAMP levels by PDE4 inhibition² causes activation of negative feedback mechanisms resulting in suppression of the cell's inflammatory activities.³ In particular, this type of manipulation brings about a marked reduction in the release of the proinflammatory cytokine TNF- α into the blood.⁴ These observations have generated considerable interest in the use of PDE4 inhibitors for the treatment of various immunological disorders, including asthma and rheumatoid arthritis.^{5,6}

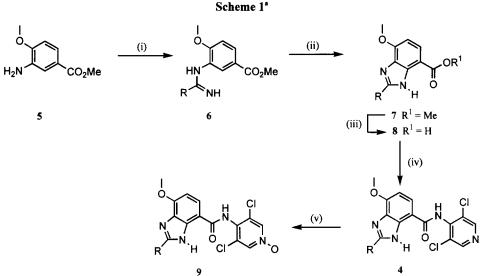


The discovery of the first PDE4 selective inhibitor, rolipram (1), more than two decades ago⁷ provided a benchmark for the development of more potent and selective analogues. Initially, the majority of research in this area focused on replacement of the pyrrolidinone of **1** with a wide range of other functionality.⁸ In our laboratories, this approach led to the discovery of clinical candidate RP 73401 (**2**).⁹ By contrast, modification of the 3,4-dialkoxyphenyl moiety of **1** (or its analogues) has been largely restricted to changes in the nature of the alkoxy substituents. The first example of a more radical alteration of the 4-methoxy-phenyl subunit of **1** was benzofuran **3**.^{10a} In this paper, we report the synthesis and biological properties of a series of PDE4 inhibitors

represented by 4, in which the 3,4-dialkoxyphenyl group of 2 has been replaced with a 2-substituted-7-methoxybenzimidazole.^{10b,c} In implementing this modification, we hoped to retain the necessary recognition elements for enzyme inhibition while (a) reducing the susceptibility of the system to oxidative metabolism¹¹ (b) increasing water solubility by introduction of an additional heteroatom and (c) providing convenient attachment points (imidazole nitrogens) for the introduction of additional functionality to further probe and define the underlying pharmacophore.

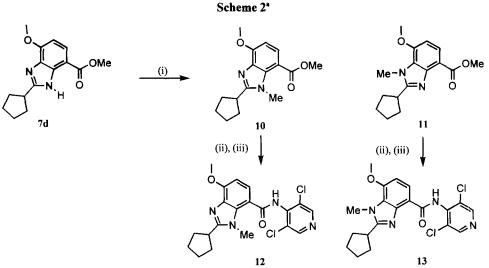
Chemistry

The synthesis of the requisite benzimidazoles is illustrated in Schemes 1 and 2. Fusion of the appropriate nitrile (Scheme 1) with 3-amino-4-methoxy benzoic acid methyl ester 5, followed by oxidative cyclization of the intermediate amidine 6 furnished the benzimidazole framework 7.¹² After saponification of the ester group in 7, the resulting acid 8 was activated (TBTU/DIEA) then coupled to 4-amino-3,5-dichloropyridine, using NaEt₂AlH₂,¹³ to give the target amide 4. Where desired, the pyridine nitrogen of 4 was oxidized to the corresponding *N*-oxide 9 using *m*-CPBA.



^aReagents and Conditions: (i) R-CN/p-TSA/Heat (60–90%). (ii) aq NaOCl then NaHCO₃/reflux (62-95%). (iii) aq NaOH, 50 °C (90–98%). (iv) TBTU/DIEA, rt then NaEt₂AlH₂/4-amino-3,5-dichloropyridine/toluene/THF (20–42% based on 8). (v) *m*-CPBA, rt.

The N-methylated benzimidazoles 12 and 13 (Scheme 2) were prepared by alkylation of 7d to give 10 and 11 as a 1/7 mixture which was separated by chromatography. Subsequent processing of 10 and 11 individually, as described above, provided the target compounds 12 and 13. Structural assignment of the isomers was made on the basis of proton-carbon HETCOR analysis.



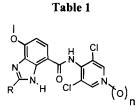
*Reagents and Conditions: (i) NaH/MeI/DMF/4 °C, (89% combined). (ii) aq NaOH, 50 °C (95%). (iii) TBTU/DIEA, rt then NaEt₂AlH₂/4-amino-3,5-dichloropyridine/toluene/THF (12: 56%; 13: 43%).

Biological Results

Table 1 summarizes the in vitro activity of a selection of benzimidazoles with respect to both inhibition of isolated guinea-pig macrophage PDE4 $(IC_{50})^{14}$ and association with the high affinity rolipram binding site (K_i) .¹⁵ In general, inhibition of PDE4 catalytic activity proved to be quite sensitive to side chain modifications at *C*-2 of the benzimidazole ring. The activities (IC_{50}) of **4a**-**f** clearly establish that substituents at this position play a role in the binding interaction but that a small group, such as a cyclopropyl, provides a sufficient presence to attain high activity. Indeed, increasing the size of the *C*-2 substituent (**4c**-**4f**) resulted in a reduction in potency. Incorporation of an oxygen atom into the side chain (**4g**) was tolerated providing a useful opportunity for increasing the polarity of these lipophilic compounds (e.g., **4d**, Clog P = 3.5 vs **4g**, Clog P = 1.5). Interestingly, the *N*-oxides **9c**,**g** were found to be equipotent with the corresponding pyridines **4c**,**g**. This observation would seem to argue against a direct H-bond between the pyridine ring of **4** and the enzyme and it may be that this interaction is, in fact, mediated by a water bridge. Alkylation of either nitrogen of the benzimidazole ring, as in **12** and **13**, resulted in a significant loss of potency (relative to **4d**). The loss in activity was most severe for **13**, which, although steric effects cannot be ruled out, may well reflect the importance of having an H-bond acceptor at *N*-3 of the benzimidazole.

Kinetic studies have shown that 1 binds to PDE4 in two distinct modes;¹⁶ the so called low affinity site, which is associated with inhibition of catalytic activity and the high affinity site, the physiological significance of which remains unclear.¹⁷ With respect to this discussion, the observation that the benzimidazoles in Table 1

inhibit catalytic activity and bind to the high affinity site, strongly supports the view that these compounds belong to the same pharmacophore family as 1.



Compound	R	n	IC ₅₀ PDE 4 (nM)	K, High Affinity Binding Site (nM)
1 (rolipr a m)			300	5
2 (RP73401)			1	1
4a	CH3	0	450	712
4b	Cyclopropyl	0	2	1
4c	<i>Iso</i> -propyl	0	27	24
4d	Cyclopentyl	0	100	540
4e	PhCH ₂	0	25	75
4f	PhCH ₂ CH ₂	0	23	56
4g	CH ₃ OCH ₂	0	42	25
9c	<i>Iso</i> -propyl	1	28	24
9g	CH ₃ OCH ₂	1	45	45
12ª			540	340
13ª			4400	10000

^aSee Scheme 2 for structure

Several of the compounds from the benzimidazole series displayed good oral activity. Table 2 shows a selection of animal data for these compounds as measured by (a) inhibition of TNF- α release in mice upon lipopolysaccaride (LPS) challenge¹⁸ (b) inhibition of streptococcal cell wall (SCW) induced arthritis in rats¹⁹ and (c) oral bioavailability in mice.

Table 2

Compound	Rª	nª	TNF-α Release ED ₅₀ (mg/kg, PO)	SCW-induced Arthritis ED ₅₀ (mg/kg, PO)	Bioavailability Mouse ^b (F%)			
4c	(CH ₃) ₂ CH	0	10	-	16			
4g	CH ₃ OCH ₂	0	2	10, tid	35			
9g	CH ₃ OCH ₂	1	7	5, bid	87			

^aRefers to the structure at the top of Table 1.

^bStudies were conducted on female Balb/c mice at a dose of 1mg/Kg.

All three compounds were effective in the TNF- α release assay when dosed orally. Two of the compounds (4g and 9g) were sufficiently bioavailable to warrant further evaluation in the SCW arthritis model. Here again, both compounds were effective, although for optimum results, 4g required dosing three times a day, presumably due to its rapid clearance from plasma (4g T_{1/2} = 1.75 h cf 9g T_{1/2} = 2.5 h).

In conclusion, replacement of the 3,4-dialkoxyphenyl substructure of 2 with a 2-substituted-4methoxybenzimidazole framework conserves the recognition elements that are required for inhibitory activity against PDE4. Furthermore, appropriate substitution at C-2 of these benzimdazoles results in compounds with good oral bioavailability suggesting that the benzimidazole system may be a useful alternative scaffold to consider when designing new classes of PDE4 inhibitor with improved pharmacokinetic profiles.

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