



BIOORGANIC & MEDICINAL CHEMISTRY LETTERS

Bioorganic & Medicinal Chemistry Letters 13 (2003) 3401-3404

Synthesis of *Para*-Alkyl Aryl Amide Analogues of Sphingosine-1-phosphate: Discovery of Potent S1P Receptor Agonists

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Received 29 April 2003; revised 22 July 2003; accepted 29 July 2003

Abstract—Sphingosine-1-phosphate (S1P) is a biologically active lysophospholipid with the capacity to induce a broad range of cellular responses via its interaction with the S1P family of G-protein coupled receptors. This report describes the synthesis of several potent S1P receptor agonists. For instance, compound **9c** displayed an $EC_{50} = 8.6$ nM at the S1P₁ receptor using a $[\gamma^{-35}S]$ GTP binding assay as compared to an $EC_{50} = 4.5$ nM for the endogenous ligand. We also report the effects associated with introduction of a phenyl ring between the 'linker' and 'lipophilic tail' regions of the analogues, for example total loss of activity at S1P₂ and increased agonism at S1P₅.

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Sphingosine-1-phosphate (S1P) has been demonstrated to act as both an intercellular signaling molecule and an intracellular second messenger (Fig. 1). Extracellular S1P binds a specific subset of seven trans-membrane G-protein coupled receptors, namely the S1P₁ (formerly Edg-1), S1P₂ (Edg-5), S1P₃ (Edg-3), S1P₄ (Edg-6), and S1P₅ (Edg-8) receptors. Through its interaction with these receptors, S1P has been shown to induce a wide variety of cellular effects including differentiation, motility, and escape from apoptosis.

The ability of S1P to mediate biological processes such as immune modulation make S1P receptor agonists and antagonists attractive candidates for use as therapeutic agents. For example, the sphingosine analogue, FTY720, is a novel immunosuppressive agent that acts by reversibly sequestering lymphocytes to secondary lymphoid tissues. FTY720 is a pro-drug that, after *O*-phosphorylation by a yet to be defined kinase, elicits lymphopenia via its interaction with one or more of the S1P receptors. Phosphorylated FTY720 (phospho-FTY720) shares several structural similarities with S1P including a lipophilic tail region, a 2-amino group, and a phosphate head group.

To understand better the processes associated with S1P binding to its receptors, it is vital to establish a structure–activity relationship (SAR) for S1P with respect to each individual receptor. The generation of S1P analogues with different binding profiles across the S1P receptors proves useful in elucidating this SAR. More specifically, the synthesis of S1P receptor specific agonists and antagonists could prove useful in the discovery of novel drugs. To supplement our previously reported S1P₁/S1P₃ receptor agonist,⁷ we now wish to report the synthesis of potent S1P₁, S1P₃, S1P₄ and S1P₅ receptor agonists along with their activity profiles across all five recombinant human S1P receptors. We also report several interesting effects in activity with the incorporation

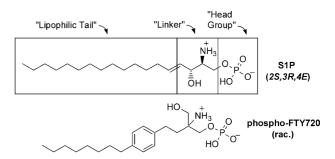


Figure 1. Structures and regions of S1P and phospho-FTY720.

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of a phenyl ring between the 'linker' and 'lipophilic tail' regions of the S1P analogues, as is the case in phospho-FTY720. These effects include a total loss of activity at S1P₂, increase in agonism at S1P₅, and a loss of C2-stereospecificity at the S1P₁ and S1P₃ receptors.

Our preparation of S1P analogues began with the synthesis of the (2S)-N-alkyl amide, (2S)-O-alkyl ester, and (2S)-N-aryl amide compounds, 5a-c, 5d, and 9e-f, respectively (Scheme 1).7 N-Boc-(L)-Serine was protected as the benzyl ester using benzyl bromide under standard conditions. The protected amino acid 1 was then phosphorylated using di-tert-butyl diisopropylphosphoramidite with subsequent hydrogen peroxide oxidation of the phosphite to afford 2 which underwent hydrogenolysis to give acid 3. Coupling of 3 to the appropriate alkyl amine using DCC afforded the protected N-alkyl amides 4a-c. Coupling of 3 to 1-tetradecanol using the PyBOP reagent gave the protected Oalkyl ester 4d. Coupling of 3 to the appropriate p-alkyl aniline using the PyBOP reagent afforded the protected N-aryl amides 4e-f. Deprotection of the amides 4a-c and 4e-f and ester 4d using 1:1 TFA/CH₂Cl₂ gave the final products 5a-d and 9e-f as the TFA salts.

The synthetic route to the (2R)-N-alkyl amide and (2R)-N-aryl amide compounds, **5e** and **9a**-**d**, respectively,

commenced with PyBOP mediated coupling of *N*-Boc-(D)-Serine(Bzl)-OH with the appropriate alkyl amine or *p*-alkyl aniline (Scheme 2). The resulting amides **6a**—**e** were then subjected to hydrogenolysis to afford the deprotected alcohols **7a**—**e**. These alcohols were then phosphorylated using di-*tert*-butyl diisopropylphosphoramidite with subsequent oxidation of the phosphite by hydrogen peroxide to give the protected phosphates **8a**—**e**. Deprotection of **8a**—**e** using 1:1 TFA/CH₂Cl₂ gave the final products **5e** and **9a**—**d** as the TFA salts.

Receptor activation by S1P and the synthetic analogues was determined in vitro by measuring the ligand dependant binding of $[\gamma^{-35}S]$ GTP to membranes containing each of the five human S1P receptors expressed in HEK293T cells.⁸ None of the compounds in these series showed any activity at the related lysophosphatidic acid receptors (LPA₁₋₃) at concentrations up to 10 μ M in this assay (data not shown). As was previously reported, *R*-configuration at the C2 position of the *N*-alkyl amide analogues appears to result in increased binding at S1P₁ and S1P₃, but not at the other S1P receptors (Table 1).⁷

Although the *N*-alkyl amide and *O*-alkyl ester compounds were poor agonists at S1P₂, introduction of a phenyl ring between the 'linker' and 'lipophilic tail'

Scheme 1. Reagents and conditions: (i) Cs_2CO_3 , BnBr, DMF, rt, 12 h, quant.; (ii) tetrazole, di-*tert*-butyl diisopropylphosphoramidite, 1:1 CH_2Cl_2/THF , rt, 12 h; (iii) H_2O_2 , rt, 4 h, quant. (2 steps); (iv) H_2 , 10% Pd/C, EtOH, rt, 12 h, 91%; (v) DCC, DMAP, HOBT, CH_2Cl_2 , rt, 12 h, 26–41%; (vi) PyBOP, DIEA, CH_2Cl_2 , rt, 6 h, 43–60% (15% for 4d); (vii) 1:1 TFA/CH_2Cl_2 , rt, 4 h, 96–100%.

Scheme 2. Reagents and conditions: (i) PyBOP, DIEA, CH_2Cl_2 , rt, 6 h, 65–77%; (ii) H_2 , 10% Pd/C, EtOH, rt, 12 h, 84–96%; (iii) tetrazole, di-*tett*-butyl diisopropylphosphoramidite, 1:1 CH_2Cl_2/THF , rt, 12 h; (iv) H_2O_2 , rt, 4 h, 60–100% (2 steps, 9% for **9d**); (v) 1:1 TFA/CH_2Cl_2 , rt, 4 h, 56–100%.

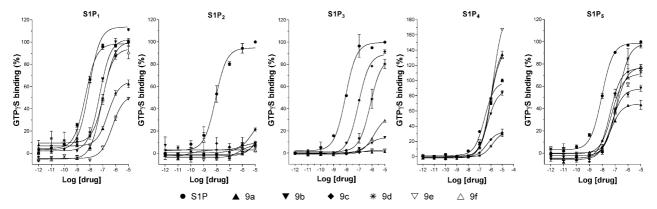


Figure 2. $[\gamma^{-35}S]GTP$ binding to HEK293T cell membranes in response to S1P and N-aryl amide S1P analogues. Each data point represents the mean of three determinations.

regions of the analogues, as in compounds 9a–f, resulted in a complete loss of activity at that receptor type (Table 1, Fig. 2). This structural modification also resulted in increased binding affinity towards $S1P_5$. In contrast to the N-alkyl amide compounds, stereochemical configuration at the C2 position of the N-aryl amide analogues does not appear to have an effect on activity at any of the S1P receptors. This finding also differs from data reported by Lynch, et al. for similar compounds containing 2-methyl groups, (S)-AFD and (R)-AFD, in which only the 2R enantiomer acted as a potent agonist at the S1P receptors (Fig. 3).

Table 1. EC₅₀ Values (nM) for S1P and N/O-alkyl amide/ester and N-aryl amide analogues at S1P receptors determined by a $[\gamma^{-35}S]GTP$ binding assay^{a,b}

N/O-Alkyl Compd	S1P ₁	S1P ₂	S1P ₃	S1P ₄	S1P ₅
S1P	0.9	2.9	1.1	nd	43.9
5a	598.4	973.2	845.4	nd	645.5
5b	397.0	2685.0	862.4	nd	1606.0
5c	1805.0	> 5000	878.6	nd	1220.0
5d	322.1	2760.0	601.9	nd	> 5000
5e	12.7	2107.0	50.8	nd	> 5000
N-Aryl Compd	S1P ₁	S1P ₂	S1P ₃	S1P ₄	S1P ₅
S1P	4.5	8.3	8.7	270	9.2
9a	260	na	na	880	43
9b	58	na	450	500	52
9c	8.6	na	89	450	41
9d	130	na	740	1400	160
				4 = 0 0	
9e	550	na	na	1500	55
9e 9f	550 82	na na	na 1500	1500 910	110

^aValues are means of three experiments (na = not active, nd = no data). ^bData for N/O-alkyl compounds are taken from ref 7.

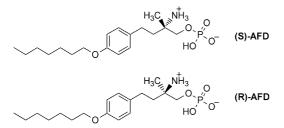


Figure 3. Structures of (S)-AFD and (R)-AFD (see ref 6).

Regarding the $S1P_1$ and $S1P_3$ receptors, the length of the alkyl chain of the *N*-aryl amide compounds was optimized at 10 carbons, compound **9c** displayed $EC_{50}s = 8.6$ nM and 89 nM, respectively, on those receptors. With respect to the $S1P_4$ receptor, shorter chain lengths of six and eight carbons on the *N*-aryl amide compounds exhibited higher potency and efficacy than the longer chain lengths of 10 and 12 carbons. At the $S1P_5$ receptor, the greatest efficacy of the *N*-aryl amide compounds was achieved with the longest chain length, 12 carbons. The potency, however, of the *N*-aryl amide compounds on $S1P_5$ was approximately equal, about one log order less than that of S1P itself.

To summarize, we have synthesized a second generation of S1P analogues that incorporate a phenyl ring between the 'linker' and 'lipophilic tail' regions of the pharmacophore, as is the case in phospho-FTY720. This structural modification has resulted in the generation of highly potent, fully efficacious agonists on four of the S1P receptors; S1P₁, S1P₃, S1P₄ and S1P₅. In particular, introduction of the phenyl ring between the 'linker' and 'lipophilic tail' regions of the S1P analogues described here results in complete loss of activity at S1P₂, increased activity at S1P₅, and a loss of C2-stereospecificity at the S1P₁ and S1P₃ receptors. These findings have helped to decipher further the SAR of S1P on the five S1P receptors and will serve as the basis for additional studies along this route.

Acknowledgements

This work was supported by grants from the NIH [NIGMS R01 GM052722 (to K.R.L.) and NIGMS F31 GM064101 (to M.D.D.)].

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