Muscarinic Agonist SAR of Azaspirodioxolanes

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Abstract: The in vitro muscarinic activity and a model for binding to the m1 receptor are presented for a series of azaspirodioxolanes.

Central activity or blood brain barrier penetration is a prerequisite for any muscarinic agonist which is to have therapeutic potential for Alzheimer's Disease (AD). The potent agonists muscarine and cis-dioxolane (CD) are quaternary ammonium compounds which do not penetrate into the CNS. Furthermore the corresponding *nor*- tertiary amine analogs of these agonists are only weakly active. As part of a program to test muscarinic agonists in AD, we set out to prepare spirocyclic nonquaternary dioxolane analogs and determine if potent CNS active compounds of this substance class were in fact accessible.

Synthesis

Methods for synthesizing azaspirodioxolanes and their sulfur analogs have been published.^{2,3} There are three basic methods starting from a ketone. Method 1 involves preparation of a cyanohydrin then elaboration to a diol and finally acetal formation. Method 2 proceeds via one carbon homologation of the ketone to an oxirane or thiirane which is directly converted to an acetal by treatment with BF₃-Et₂O and a carbonyl component. Finally, with method 3, the oxirane or thiirane may be ring opened with acetate or thioacetate ion. In this case acetate hydrolysis is followed by standard acetalization with a carbonyl component.

SAR

At the outset of our studies there was already some evidence supporting the possibility of preparing potent nonquaternary analogs of CD. The quinuclidine dioxolane analog AF-30 was a known agonist of moderate potency.⁴ Furthermore, the spirocyclic piperidine succinimide RS86⁵, which had been clinically investigated, was a centrally active muscarinic agonist only tenfold less potent than muscarine. On this basis the first compound prepared was the N-methylpiperidine spirodioxolane 1. This compound proved to be equiactive to RS86, exhibiting high agonist potency with pD₂ values of 6.0 and 6.7 in the ileum and ganglion models and 100nM cortex affinity in displacing [³H]-CD. A very strict SAR relationship was observed for the series of subsequently prepared spirodioxolane analogs in which the activity of 1 was not surpassed. The *in vitro* pharmacological data for the spirodioxolane analogs 1-27 are summarized in Table 1. Compounds 1-15 are spiropiperidines and compounds 17-20 are spiropyrrolidines. A detailed description of

G. Shapiro et al.

the pharmacological tests employed may be found elsewhere.⁶ In brief, the ileum model serves as a functional

measure of peripheral muscarinic activity while the ganglion model is a functional model of central muscarinic activity. [³H]-CD-binding provides a measure of agonist affinity⁷ and [³H]-pirenzepine binding provides a measure of antagonist affinity.

Dioxolane ring variation

The C2 position of the dioxolane ring plays a key role in activity appearing to mimic the acyl carbon of Ach. A methyl group at C2 is optimal and necessary for the agonist activity of 1. The unsubstituted compound 2 retains only marginal activity. Bulkier substituents at C2 cause a drop in agonist activity and, from a certain size, a shift towards antagonism. The ethyl substituted analog 3 experiences a tenfold drop in potency relative to 1. Compound 4 having a methoxy group at C2 is equiactive to 3 while the acetylene analog 5 is slightly more active than 3 particularly regarding ganglion potency and efficacy. Compound 8 having dimethylsubstitution at C2 is inactive. A single phenyl substitution at C2 yields the weak antagonist 7 whereas diphenylsubstitution yields the potent antagonist 9. While the 2S absolute stereochemistry is optimal, 1-(R), the (+)-2R enantiomer of 1, is only 5 fold less potent than 1-(S), the (-)-2S antipode. Oxathiolane 12 in which the O1 acetal oxygen of 1 is exchanged for sulfur is equiactive to 1 in the ileum model and displays lowered ganglion potency but increased ganglion efficacy. The CD binding affinity of 12 is lowered by a factor of 3. Oxathiolane 13 in which the O3 acetal oxygen is exchanged for sulfur is about 10 fold less active than 1 in all tests. Dithiolane 14 displays activity comparable to 13. Exchange of the O3 dioxolane oxygen with sulfur for pyrrolidine 17, AF-30 and CD¹¹ leads to little or no loss of activity.

Aza ring variation

The geometrical disposition for the amino nitrogen of 1, two methylene units removed from C4 of the dioxolane ring, is well defined with two alternative chair conformations of the piperidine ring being highly favored. Compound 15 where only one methylene unit separates the spatially well defined piperidine ring nitrogen is virtually inactive. When the amine function is incorporated in the conformationally distinct and more flexible pyrrolidine ring of 17, activity is restored. As with CD¹² and AF-30⁴ the *cis* configuration of the methyl group and aminomethylene element in 17 is preferred and *trans* compound 18 is less active. Compound 17 is roughly equiactive to AF-30 but tenfold less potent than 1 in functional tests. The azabicycloheptane-spirodioxolane 21 was prepared¹³ as a 70:30 mixture of C2 methyl epimers where the O3 of the dioxolane ring system is *exo* to the azanorbornane ring. In functional tests this mixture is only slightly less potent than 1. Compound 22¹⁴ where O3 is *endo* and the methyl group and aminomethylene unit are *cis* is less active than 21 and *endo-trans* isomer 23 is only weakly active. Tropane analogs 26,27¹⁵ and *cis*-2,6-dimethylpiperidine analogs 24,25¹⁶ which incorporate the structure of 1 but do not allow piperidine ring inversion were all inactive. Quaternization of 1 to the dimethylammonium analog 10 results in a ca. tenfold drop in potency but an increase in ganglion efficacy. The secondary amine analog 11 lacking the N-methyl

Piperidines

Pyrrolidines

AF30 analogs

Compound#	X	Y	R1	R2	A	В	pD2 ileum	eff.%	pD2 ganglion	eff.%	CD nM	Pir nM	
1-racemic	0	0	Me	н	NMe	CH2	6.0	100	6.7	80	100	5825	
(+)-1-(R)	0	0	Me(R)	Н	NMe	CH2	5.5	100	64	80	240	7690	
(·)-1-(S)	0	0	Me(S)	Н	NMe	CH2	6.3	100	7.0	80	85	5325	
2	0	0	Н	Ŧ	NMe	CH2	4.6	70	46	90	1200	>10,000	
3	0	0	Et	Н	NMe	CH2	5.0	70	5.5	85	745	1325	
4	0	0	OMe	Н	NMe	CH2	4.7	80	5.5	100	870	>10,000	
5	0	0	C≅CH	Н	NMe	CH2	52	100	5.9	120	295	3000	
6	0	0	CCI3	Н	NMe	CH2	4.1	60	<40		960	200	
7	0	0	Ph	Н	NMe	CH2	pA2 5.3		n.d.		2800	620	
8	0	0	Me	Me	NMe	CH2	<4.0		<4.0		>10,000	>10,000	
9	0	0	Ph	Ph	NMe	CH2	pA2 8.0		pA2 8.2		8.1	0.60	
10	0	0	Me	H	NMe2	Œ.	5.2	105	6.1	110	210	5000	
11	0	0	Me	Н	NH	CH2	5.1	100	5.9		145	>10,000	
12	0	s	Me	н	NMe	CH2	60	100	61	100	185	2650	
13	S	0	Me	Н	NMe	CH2	5.1	100	5.6	70	720	3160	
14	S	S	Me	Н	NMe	CH2	4.9	100	5.0	100	530	2100	
15	0	0	Me	Н	CH2	NMe	<4.0		<4.0				
RS 86	co	8	Et	Н	NMe	CH2	6.1	100	67	80	87	765	
16	CO	co	Me	Н	NMe	5	5.5	100	6.2		175	4050	
									-				
pyrrolidines													
17	0	0	Me cis				5.2	100	56	90	550	>10,000	
18	0	0	Me trans				4.5	100	n d.		1750	>10,000	
19	0	s	Me cis				5.2	100	5.7	100	585	6700	
20	S	0	Me cis								9150	>10,000	
AF-30	0	0	Me as				48	100	6.0	60	90	1575	
AF-102b	0	S	Me cis				4.9	50	6.1	80	105	430	
AF-102a	0	S	Me trans			-	5.4	30	5.7	70	515	1250	
21							5.9	100	6.5	75	30_	1950	
22							5.5	100	5.7	85	180	6400	
23							4.1	100	44		1850	>10,000	
muscarine	L						68	100	7.4	100	12.0	5680	
CD	L						7.5	100	7.8	100	6.4	2100	

CH₃

21

CH₃

CH4

N CH₃

25

26

27

28

G. Shapiro et al.

group suffers a roughly tenfold drop in potency with a somewhat less marked loss of CD-affinity.

Receptor Binding Model

Recently, we have proposed a hypothetical model for the binding of muscarinic agonists to the m1 receptor.⁶ In this dynamic model the triad of residues Asp-105, Ser-109 and Ser-112 aligned on one side of the third putative transmembrane idealized alpha helix binds agonists via a salt bridge and one or two hydrogen bonds. For different agonists distinct receptor conformations were modeled demonstrating the dynamic nature of the receptor and the limitations intrinsic to a precise pharmacophore where receptor mobility is neglected. The extraordinary potency of the azabicycloheptane oxadiazole 2817 shown in Figure 1 (crossed stereoscopic view) was ascribed to an ideal receptor fit with a strong salt bridge and two equally strong hydrogen bonding interactions.⁶ The most active (2'S,3R) isomer of the quinuclidine spirodioxolane AF-30 can be accommodated in the same dynamic model (Fig. 2). The protonated quinuclidine nitrogen forms an ideal linear salt bridge (2.7Å) to Asp-105, Ser-109 forms a good hydrogen bond to the O3 dioxolane (1.9Å) and Ser-112 forms a weaker hydrogen bond to O1 (2.3Å). The azabicycloheptane analogs 21 and 22 fit in an analogous manner. In the case of the exo-isomer 21, the methylene bridge occupies the same region as the ethylene unit of AF-30 which is seen extending out of the plane (Fig. 2). One may postulate this orientation of the azabicyloheptane ring, which was also found for 28, to be preferrable. Both chair conformations of spiropiperidine 1-(S) gave equally good fits. To obtain a good fit for the conformation where O3 is equatorial, the N-methyl group must also be equatorial, and the torsion angle for Asp105 differs slightly (Fig. 3). An optimal salt bridge is formed and there is a good hydrogen bond (1.9Å) between Ser109 and O3 of the dioxolane ring. In the conformation where O3 is axial the N-methyl group must also be axial (fit not shown). In this fit there is also only a single hydrogen bonding interaction (Ser109 to O3). A binding mode for 1-(S) in which there are two hydrogen bonds as for AF-30 requires a large deviation of the salt bridge from optimal linearity. Pyrrolidine dioxolane 17 could be fit analogously to AF-30, or more exactly 21, with the N-methyl group occupying a pseudoaxial position (not shown). Alternatively a fit similar to that shown for 1-(S) was possible (Fig. 4). The fits shown for AF-30, 1-(S) and 17 are consistent with the effects of sulfur oxygen exchange on activity. Substitution of O1 of AF-30 with sulfur yields AF-102b which does not lose potency but does lose efficacy and is therefore a weaker agonist. This is consistent with the moderate hydrogen bonding role for O1 in AF-30. Substitution of sulfur for O1 in 1 and 17 results in slight and no loss of agonist activity, respectively. This is consistent with the depicted binding modes for these compounds where O1 is not involved in hydrogen bonding. It should be noted that the important hydrophobic binding site occupied by the dioxolane methyl group is not identified in this model. For 1, the 1-(R) enantiomer must have an active conformation and/or a binding mode distinct from the 1-(S) since the dimethylsubstituted 8 is completely inactive. When the dioxolane ring of 1 adopts an envelope conformation with C2 at the flap position, a good superposition of the C2-methyl groups of the 1-(S) and 1-(R) enantiomers may be achieved by simple inversion of the flap position. Interestingly, the same conformational relationship exists between the cis and trans isomers of the pyrrolidine, quinuclidine and azabicycloheptane analogs for which, pairwise. a similar 5-10fold difference in agonist activity is observed. 18 The SAR clearly shows that the agonist binding site is a very tight pocket with little tolerance for excess volume. Greater steric bulk generally leads to loss of agonist potency and/or efficacy (shift toward antagonism). Finally, this study emphasizes the potential importance of multiple binding modes for a given agonist.

Conclusion

The (-)-S isomer of 1 was developed under the classification of SDZ 210-086 for the indication Alzheimer's Disease. Clinical studies and further characterization with regard to the in vivo pharmacological actions of 210-086 have been performed.¹⁹

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- 14. This material was prepared via method 2 on the endo-epoxide followed by chromatographic separation of the two diastereomers 22,23 formed.
- 15. From a homogeneous tropane (N-benzyl) endo-epoxide, 26 was prepared via method 3 and 27 was
- prepared by method 2 (stereochemistry assigned by NOE-NMR). From a pure epoxide, a pure diol (N-carboethoxy) precursor was obtained by method 3. Acetalization of this diol could not be performed without facile equilibration at the spirocenter (60:40). Purification of the individual isomers to ca. 85% was achieved by simple chromatography and recrystallization. After carbamate reduction and fumarate salt formation 24 was isolated ca. 95% pure while 25 contained 30% 24 even after recrystallization (stereochemical assignment by NOE-NMR).
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