

Design, Synthesis, and Biological Activity of Prazosin-Related Antagonists. Role of the Piperazine and Furan Units of Prazosin on the Selectivity for α_1 -Adrenoreceptor Subtypes

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Prazosin-related quinazolines **4–20** were synthesized, and their biological profiles at α_1 -adrenoreceptor subtypes were assessed by functional experiments in isolated rat vas deferens (α_{1A}), spleen (α_{1B}), and aorta (α_{1D}) and by binding assays in CHO cells expressing human cloned α_1 -adrenoreceptor subtypes. The replacement of piperazine and furan units of prazosin (**1**) by 1,6-hexanediamine and phenyl moieties, respectively, affording **3–20**, markedly affected both affinity and selectivity for α_1 -adrenoreceptor subtypes in functional experiments. Cystazosin (**3**), bearing a cystamine moiety, was a selective α_{1D} -adrenoreceptor antagonist being 1 order of magnitude more potent at α_{1D} -adrenoreceptors (pA_2 , 8.54 ± 0.02) than at the α_{1A} - (pA_2 , 7.53 ± 0.01) and α_{1B} -subtypes (pA_2 , 7.49 ± 0.01). The insertion of substituents on the furan ring of **3**, as in compounds **4** and **5**, did not improve the selectivity profile. The simultaneous replacement of both piperazine and furan rings of **1** gave **8** which resulted in a potent, selective α_{1B} -adrenoreceptor antagonist (85- and 15-fold more potent than at α_{1A} - and α_{1D} -subtypes, respectively). The insertion of substituents on the benzene ring of **8** affected, according to the type and the position of the substituent, affinity and selectivity for α_1 -adrenoreceptors. Consequently, the insertion of appropriate substituents in the phenyl ring of **8** may represent the basis of designing new selective ligands for α_1 -adrenoreceptor subtypes. Interestingly, the finding that polyamines **11**, **16**, and **20**, bearing a 1,6-hexanediamine moiety, retained high affinity for α_1 -adrenoreceptor subtypes suggests that the substituent did not give rise to negative interactions with the receptor. Finally, binding assays performed with selected quinazolines (**2**, **3**, and **14**) produced affinity results, which were not in agreement with the selectivity profiles obtained from functional experiments. This rather surprising and unexpected finding may be explained by considering neutral and negative antagonism.

Introduction

It is clear now that α_1 -adrenoreceptors are comprised of multiple subtypes that can be classified by both pharmacological and binding studies into at least three subtypes, that is α_{1A} (α_{1a}), α_{1B} (α_{1b}), and α_{1D} (α_{1d}), with upper and lower case subscripts being used to designate native or recombinant receptors, respectively.^{1–3} However, the situation appears to be more complex as, in addition to α_{1A} -, α_{1B} -, and α_{1D} -adrenoreceptor subtypes, which share a high affinity for prazosin, the existence of additional α_1 -adrenoreceptors has been proposed. These are called α_{1L} -adrenoreceptors and are characterized by a low activity for prazosin. However, these receptors have not been cloned yet and their characterization is still difficult.⁴

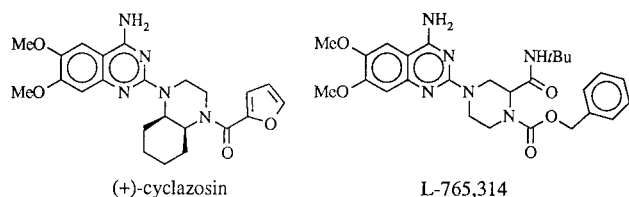
The existence of multiple α_1 -adrenoreceptor subtypes holds the promise of developing new molecules, which target only one receptor while not affecting others. Furthermore, the different localization of these receptor subtypes brought about the possibility of designing drugs that selectively interact with distinct subtypes, thus avoiding the occurrence of possible side effects.

Current evidence indicates that rat submaxillary gland,⁵ human liver,⁶ and various tissues such as prostatic rat vas deferens,⁷ rabbit prostate, and prostatic urethra⁸ contain predominantly the α_{1A} -adrenoreceptor, whereas rat liver and spleen⁹ are considered α_{1B} -adrenoreceptor preparations, and the α_{1D} -adrenoreceptor mediates the contraction in rat aorta.^{10,11} As a result, the effort to design agents selective for each of the three α_1 -adrenoreceptor subtypes has been an active area of research. Several relatively selective ligands for α_1 -adrenoreceptors are now available.¹² For example, SNAP 5089,¹² KMD-3213,¹³ Rec 15/2739,¹⁴ RS-17053,¹⁵ and (–)-mephendioxan¹⁶ are selective for α_{1A} -adrenoreceptors, (+)-cyclazosin¹⁷ and L-765,314¹⁸ are selective for the α_{1B} -subtype, and BMY-7378¹⁹ and cystazosin²⁰ are selective for the α_{1D} -subtype. Whereas it has been demonstrated that α_{1A} -adrenoreceptor antagonists can be useful in the treatment of benign prostatic hyperplasia, a potential therapeutic use for both α_{1B} - and α_{1D} -subtype antagonists has not been defined yet. Perhaps, the fact that only recently so-called selective α_1 -adrenoreceptor antagonists have become available has prevented the physiological roles of α_{1B} - and α_{1D} -adrenoreceptor subtypes in blood pressure control or other physiological functions from being revealed. It should be emphasized

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Chart 1



as well that the ideal selective ligands, which recognize only one among multiple receptor subtypes, are few and it remains a formidable challenge to medicinal chemists to realize useful drugs of this kind. For example, BMY-7378 is by far the most selective α_{1d} -adrenoreceptor antagonist reported to date.¹⁹ It displayed a 75- and 80-fold selectivity for the α_{1d} -adrenoreceptor relative to α_{1b} and α_{1a} subtypes, respectively. However, BMY-7378 is more potent (14-fold) at 5-HT_{1A} receptors and only 9-fold less potent at D₂ receptors with respect to α_{1d} -adrenoreceptors.¹²

Design Rationale

The starting point of this investigation was the observation that modifying the piperazine ring of prazosin (**1**) may afford antagonists, which are able to differentiate among α_1 -adrenoreceptor subtypes. It was shown that the piperazine ring of prazosin is not essential for activity and can be replaced with an α,ω -alkanediamine chain.²¹ Among a series of quinazolines bearing a polymethylene chain, compound **2** displayed the highest affinity for rat vas deferens α_1 -adrenoreceptors, being even more potent than prazosin. It was suggested that the hexane chain of **2** might contribute to the binding by interacting with a lipophilic site located between the sites where quinazoline and furan rings interact.²¹ More recently, it was demonstrated that the hexane chain could be constrained into a cyclohexyl moiety as in cyclazosin.²² The (+)-enantiomer of cyclazosin (Chart 1) can be considered the first α_{1b} -selective adrenoreceptor antagonist in binding assays, being about 100-fold selective versus the native α_{1A} and about 40-fold versus the recombinant α_{1a} - and α_{1d} -subtypes.¹⁶ Following a design rationale similar to that leading to cyclazosin, Patane et al.¹⁸ synthesized, by incorporating new structural elements into the prazosin piperazine moiety, L-765,314 (Chart 1) which turned out to be a selective α_{1b} -adrenoreceptor antagonist.

Thus, the finding that the affinity profile of prazosin-related quinazolines can depend on the type of moiety linking the two nitrogen atoms of the piperazine ring of prazosin prompted us to further modify the structure of analogue **2**, in an attempt to improve the affinity and selectivity for different α_1 -adrenoreceptor subtypes. In particular, two types of structural modifications were performed on the structure of **2**, that is (a) replacement of the hexane spacer with a cystamine moiety and (b) insertion of substituents on the furan ring or its replacement by an (un)substituted phenyl unit. Thus, replacing the hexane chain of **2** by a cystamine moiety, which is a structural feature of benextramine,²³ an irreversible α -adrenoreceptor antagonist, afforded cystazosin (**3**) which displayed an interesting selectivity profile in comparison with both (+)-cyclazosin and the carbon analogue **2**, owing to a significantly lower affinity

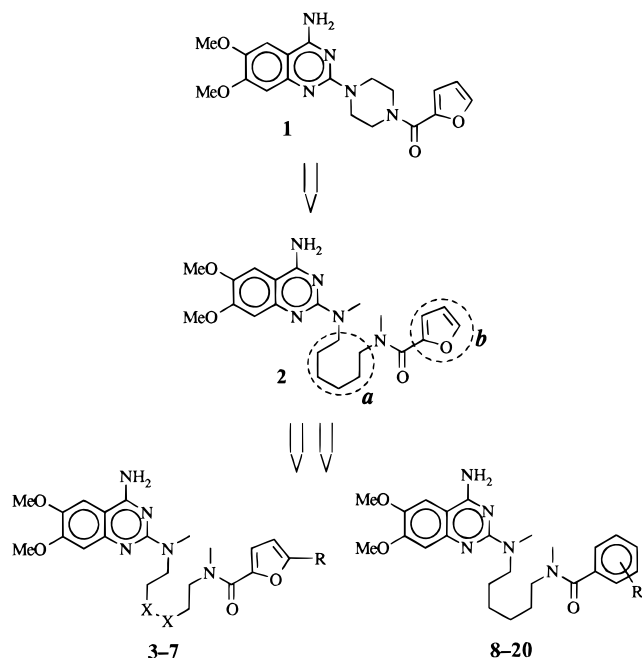


Figure 1. Design strategy for the synthesis of quinazolines **3–20** by replacing the hexane spacer and/or the furan ring of **2**, an open analogue of prazosin (**1**), by (a) a cystamine moiety and (b) an (un)substituted phenyl ring, respectively.

for α_{1A} - and α_{1B} -adrenoreceptor subtypes relative to the α_{1D} -subtype.²⁰ This observation formed the basis for further structural modifications. We thought that increasing the number of contacts between a ligand and its receptor would hopefully also increase receptor subtype selectivity. To this end we introduced a chloromethyl substituent at position 5 of furan ring of **3** and **2**, affording **4** and **6**, respectively, because it can be easily functionalized as in **5** and **7**. According to its properties, an amine function can be protonated at physiological pH giving rise to a possible additional interaction with a nucleophilic, complementary receptor group, which would increase the possibility to achieve receptor subtype selectivity. Subtle and unpredictable differences in the binding pockets may account for selectivity; thus incorporation of additional structural elements in the structure of a nonselective ligand may well lead to preferential recognition of a particular receptor subtype. Next, the furan ring of **2** was replaced by a phenyl ring affording **8**, which offered us the possibility to incorporate additional structural elements at different positions, yielding **9–20**. These 12 compounds allow us to investigate the effect of the substituents not only upon the affinity but also on the selectivity for α_1 -adrenoreceptor subtypes. The design strategy for our compounds is shown in Figure 1.

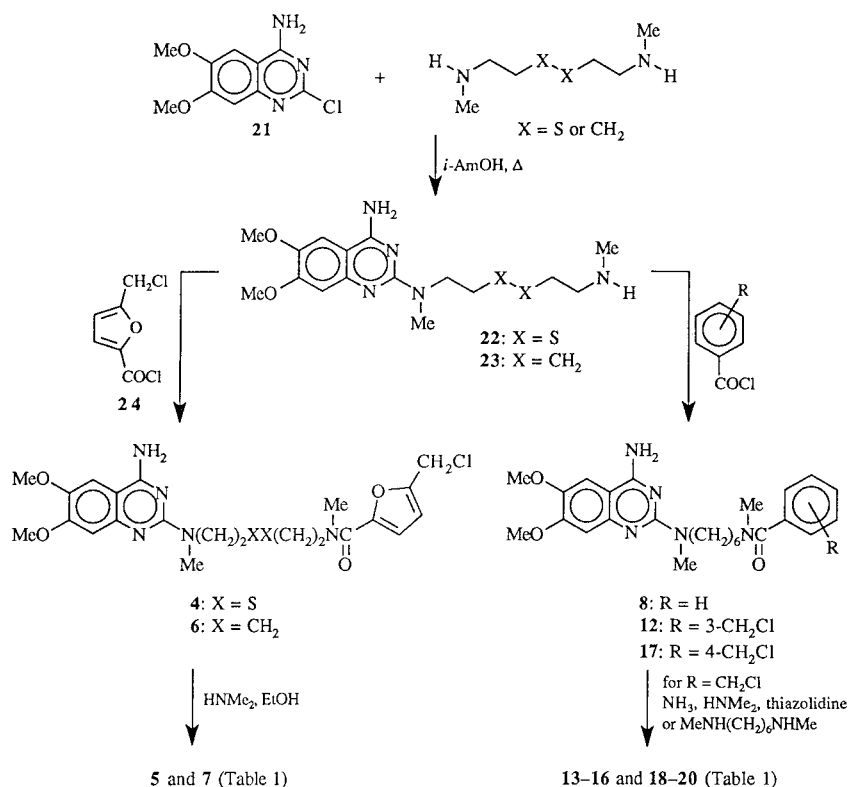
We describe here the synthesis and the pharmacological profile of quinazolines **4–20** in functional and binding experiments in comparison with prototypes prazosin (**1**) and its open analogues **2** and **3**.

A preliminary communication dealing with cystazosin (**3**) has been published recently.²⁰

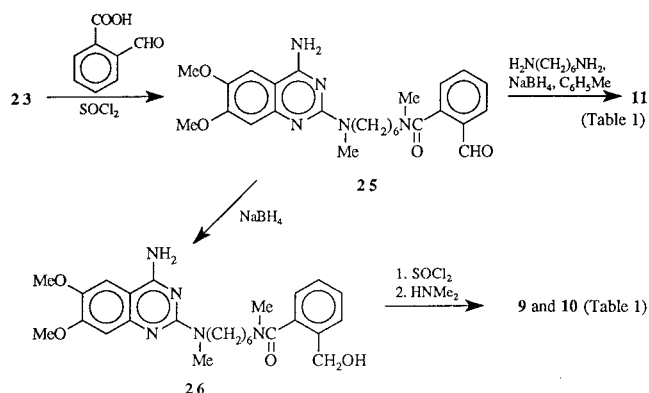
Chemistry

The new compounds were synthesized by standard methods as illustrated in Schemes 1 and 2 and characterized by ¹H NMR, IR, and elemental analysis.

Scheme 1



Scheme 2



4-Amino-2-chloro-6,7-dimethoxyquinazoline (**21**) was the common starting material. Adapting the procedure reported for **23**,²⁴ compound **22** was obtained by aromatic nucleophilic substitution of *N,N*-dimethylcystamine²⁰ on **21**. The reaction of **22** and **23** with 5-chloromethyl-2-furoyl chloride (**24**)²⁵ afforded **4** and **6**, respectively, which were transformed by reaction with dimethylamine into the corresponding *N,N*-dimethylamino analogues **5** and **7** (Scheme 1). Compound **8** was synthesized by reaction of **23** and benzoyl chloride. Similarly, **12** and **17** were obtained from **23** and 3- or 4-chloromethylbenzoyl chloride, respectively. Chlorides **12** and **17** were reacted with dimethylamine, thiazolidine, or *N,N*-dimethyl-1,6-hexanediamine²⁶ to give the corresponding amines **13–16** and **18–20**.

Finally, 2-substituted derivatives **9–11** were synthesized following the synthetic pathway shown in Scheme 2. Intermediate **25** was obtained through amidation of **23** with 2-formylbenzoyl chloride, which was generated in situ by treating 2-formylbenzoic acid with SOCl_2 . Reductive amination of **25** by 1,6-hexanediamine af-

forded **11** whereas reduction of **25** with NaBH_4 gave the corresponding alcohol **26**. The latter compound was transformed into the chloride **9** which, in turn, was alkylated to the amine **10**.

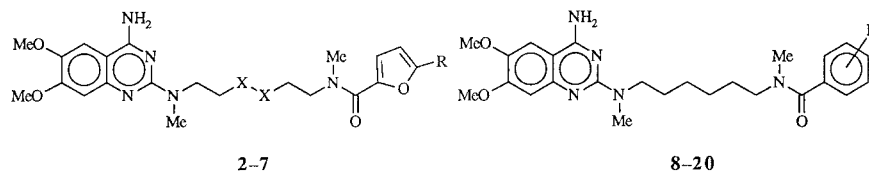
Biology

Functional Studies. The pharmacological profile of prazosin-related quinazolines **2–20** was evaluated at α_1 -adrenoreceptor subtypes and α_2 -adrenoreceptors on different isolated tissues using prazosin (**1**) and BMY-7378 as standard compounds. α_1 -Adrenoreceptor subtypes blocking activity was assessed by antagonism of (–)-noradrenaline-induced contraction of prostatic vas deferens (α_{1A})²⁷ or thoracic aorta (α_{1D})²⁸ and by antagonism of (–)-phenylephrine-induced contraction of spleen (α_{1B}),²⁸ while α_2 -adrenoreceptor blocking activity was determined by antagonism of the clonidine-induced depression of the twitch responses of the field-stimulated prostatic portion of rat vas deferens. The potencies of the drugs were expressed as pA_2 values.^{29–31}

Binding Experiments. Receptor subtype selectivity of few prazosin-related quinazolines was further determined by employing receptor binding assays. [^3H]-Prazosin was used to label cloned human α_1 -adrenoreceptors expressed in Chinese hamster ovary (CHO) cells.³² Furthermore, [^3H]rauwolscine and [^3H]spiperone were used to label α_2 -adrenoreceptors in rat cortex and D_2 receptors in rat striatum, respectively, whereas [^3H]8-hydroxy-2-(di-*n*-propylamino)tetralin ([^3H]8-OH-DPAT) was the radioligand to label cloned human 5-HT_{1A} receptors which were expressed in HeLa cells.^{33,34}

Results and Discussion

The biological activity expressed as pA_2 values, at α_1 -adrenoreceptor subtypes and α_2 -adrenoreceptors of com-

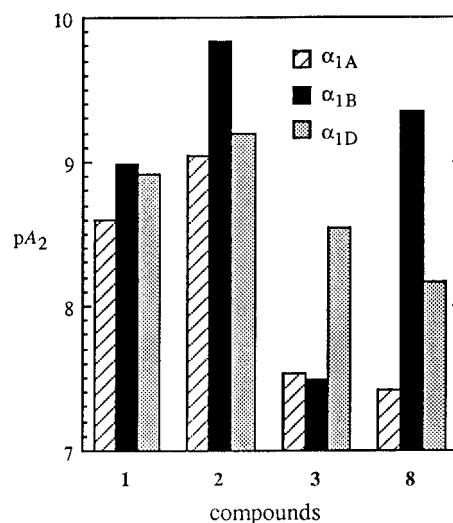
Table 1. Antagonist Affinities, Expressed as pA_2 , of **2–20** at α_1 - and α_2 -Adrenoreceptors on Isolated Tissue from the Rat, Namely, Prostatic Vas Deferens (α_{1A} and α_2), Spleen (α_{1B}), and Thoracic Aorta (α_{1D}), in Comparison to Reference Compounds Prazosin (**1**) and BMY-7378

no	X	R	pA_2^a			
			α_{1A}	α_{1B}	α_{1D}	α_2^b
1	(prazosin)		8.60 ± 0.07	8.99 ± 0.01	8.91 ± 0.04	5.43 ± 0.13
2	CH ₂	H	9.04 ± 0.02	9.84 ± 0.01	9.19 ± 0.03	6.66 ± 0.03
3	S	H	7.53 ± 0.01	7.49 ± 0.01	8.54 ± 0.02	<5
4	S	CH ₂ Cl	7.78 ± 0.07	7.37 ± 0.05	8.03 ± 0.09	<5
5	S	CH ₂ NMe ₂	6.70 ± 0.08	7.27 ± 0.03	8.22 ± 0.06	<5
6	CH ₂	CH ₂ Cl	8.17 ± 0.02	8.97 ± 0.06	9.39 ± 0.02	<5
7	CH ₂	CH ₂ NMe ₂	7.41 ± 0.04	8.62 ± 0.02	8.23 ± 0.01	<5
8		H	7.42 ± 0.03	9.35 ± 0.02	8.16 ± 0.02	6.52 ± 0.05
9		2-CH ₂ Cl	6.38 ± 0.05	8.73 ± 0.04	8.55 ± 0.02	5.31 ± 0.01
10		2-CH ₂ NMe ₂	7.04 ± 0.01	7.88 ± 0.04	8.26 ± 0.02	<5
11		2-CH ₂ NH(CH ₂) ₆ NH ₂	7.97 ± 0.01	9.17 ± 0.02	8.50 ± 0.04	5.47 ± 0.01
12		3-CH ₂ Cl	7.10 ± 0.01	7.18 ± 0.01	7.29 ± 0.05	<5
13		3-CH ₂ NH ₂	7.32 ± 0.01	8.61 ± 0.02	8.69 ± 0.01	5.44 ± 0.01
14		3-CH ₂ NMe ₂	7.26 ± 0.08	7.43 ± 0.02	8.26 ± 0.03	<5
15		3-CH ₂ N ₅	7.82 ± 0.09	8.03 ± 0.05	8.51 ± 0.06	5.94 ± 0.03
16		3-CH ₂ N(Me)(CH ₂) ₆ NHMe	8.05 ± 0.07	7.71 ± 0.05	7.84 ± 0.04	<5
17		4-CH ₂ Cl	7.11 ± 0.02	9.15 ± 0.04	7.71 ± 0.01	5.54 ± 0.02
18		4-CH ₂ NMe ₂	7.14 ± 0.01	8.53 ± 0.01	7.86 ± 0.08	5.12 ± 0.06
19		4-CH ₂ N ₅	7.23 ± 0.10	8.01 ± 0.03	8.17 ± 0.03	<5
20		4-CH ₂ N(Me)(CH ₂) ₆ NHMe	6.72 ± 0.04	9.21 ± 0.09	8.46 ± 0.07	5.60 ± 0.03
BMY-7378			6.94 ± 0.08	7.55 ± 0.07	8.34 ± 0.05	<5

^a pA_2 values ± SE were calculated from Schild plots,²⁹ constrained to a slope of -1.0 , unless otherwise specified.³⁰ pA_2 is the positive value of the intercept of the line derived by plotting $\log(DR - 1)$ vs \log [antagonist]. The $\log(DR - 1)$ was calculated at least at three different antagonist concentrations, and each concentration was tested from four to six times. Dose–ratio (DR) values represent the ratio of the potency of the agonist (EC_{50}) in the presence of the antagonist and in its absence. Parallelism of dose–response curves was checked by linear regression, and the slopes were tested for significance ($p < 0.05$). ^b pA_2 values were calculated at only one concentration (10 μ M) according to van Rossum.³¹

pounds used in the present study, is shown in Table 1. To make relevant considerations on structure–activity relationships, prototypes **1** and **2** and the reference compound BMY-7378 were included for comparison. All quinazolines behaved as competitive antagonists as revealed by the slopes of their Schild plots, which were not significantly different from unity ($p > 0.05$) (Table 1). By taking as a starting point the hexane analogue **2** of prazosin, it is possible to observe how affinity and selectivity for α_1 -adrenoreceptor subtypes can be markedly affected by replacing the hexane moiety and/or the furan ring by a cystamine unit and an (un)substituted phenyl unit, respectively. Interestingly, all prazosin-related compounds of the present investigation displayed a weak, if any, affinity for α_2 -adrenoreceptors in comparison with α_1 -adrenoreceptor subtypes.

An analysis of the results shown in Table 1 reveals that the replacement of the hexane spacer or the furan ring of **2** by a cystamine moiety or a phenyl group, affording **3** and **8**, respectively, caused a dramatic effect in the affinity profile for α_1 -adrenoreceptor subtypes as shown in Figure 2. Clearly, **2** is a very potent α_1 -adrenoreceptor antagonist but, at the same time, is not selective, displaying only a slight preference for the α_{1B} -subtype. Interestingly, the structural modifications performed on **2** did not improve affinity for α_1 -adrenoreceptor subtypes but, what is more important, gave rise

**Figure 2.** Affinity constants (pA_2) in rat prostatic vas deferens (α_{1A}), spleen (α_{1B}), and aorta (α_{1D}) α_1 -adrenoreceptor subtypes of cystazosin (**3**) and **8** in comparison with prazosin (**1**) and **2**.

to selectivity. Thus, cystazosin (**3**) resulted in a selective α_{1D} -adrenoreceptor antagonist, owing to a slight (4.5-fold) decrease in affinity for the α_{1D} -subtype and a large drop in affinity (32- and 224-fold, respectively) for α_{1A} - and α_{1B} -subtypes in comparison with **2**. On the other

Table 2. Affinity Constants (pK_i) of **2**, **3**, and **14** for Cloned α_1 -Adrenoreceptor Subtypes and 5-HT_{1A} Receptors, Native α_2 -Adrenoreceptors, and D₂ Receptors in Comparison to Reference Compounds^a

no.	pK_i , human cloned receptors			pK_i , native receptors (rat)		
	α_{1A}	α_{1B}	α_{1D}	5-HT _{1A}	α_2 (cerebral cortex)	D ₂ (striatum)
1	9.23 ± 0.07	9.39 ± 0.10	9.65 ± 0.08	<6	6.80 ± 0.03	<5
2	9.78 ± 0.04	9.96 ± 0.07	9.71 ± 0.01	<6	7.00 ± 0.16	5.63 ± 0.02
3	9.38 ± 0.05	8.97 ± 0.09	9.14 ± 0.07	<6	6.23 ± 0.07	≤5
14	9.49 ± 0.06	9.78 ± 0.06	9.61 ± 0.11	6.13 ± 0.04	7.72 ± 0.25	5.71 ± 0.15
BY ^b	6.36 ± 0.06	7.19 ± 0.04	8.89 ± 0.01	8.76 ± 0.28	5.98 ± 0.20	7.32 ± 0.04

^a Values are the mean ± SE of at least three separate experiments performed in triplicate. The pseudo-Hill coefficients (nH) were not significantly different from unity ($p > 0.05$). Equilibrium inhibition constants (K_i) were derived using the Cheng–Prusoff equation.⁴⁰ Scatchard plots were linear or almost linear in all preparations tested. The affinity estimates were derived from displacement of [³H]prazosin from α_1 -adrenoreceptors, [³H]rauwolscine from α_2 -adrenoreceptors, [³H]spiperone from D₂ receptors, and [³H]8-hydroxy-2-(di-*n*-propylamino)tetraline from 5-HT_{1A} receptors. ^b BY, BMY-7378.

hand, the phenyl analogue **8** displayed a significantly improved α_{1B} -selectivity (85- and 15-fold relative to α_{1A} - and α_{1D} -subtypes, respectively), owing to a much larger decrease in affinity for both α_{1A} - and α_{1D} -subtypes than for the α_{1B} -subtype in comparison with **2** (Figure 2).

The insertion of a 5-chloromethyl or a 5-*N,N*-dimethylaminomethyl substituent on the furan ring of **3**, affording **4** and **5**, respectively, did not improve the selectivity profile. The same structural modification performed on **2** to afford **6** and **7** caused a marked decrease in affinity which was more pronounced for 5-*N,N*-dimethylaminomethyl group. However, compound **6** was slightly more potent than **2** at α_{1D} -adrenoreceptors while displaying a significantly lower affinity at α_{1A} - and α_{1B} -subtypes as revealed by its pA_2 values (α_{1A} , 8.17 ± 0.02; α_{1B} , 8.97 ± 0.06; α_{1D} , 9.39 ± 0.02). Clearly, this finding suggests that appropriate substituents on the aromatic moiety may have a role in achieving receptor subtype selectivity.

Next, the insertion of a substituent on the benzene ring of **8**, affording **9–20**, affected differently, according to substituent type and position, the affinity and, as a consequence, the selectivity for α_1 -adrenoreceptor subtypes. The affinity profiles of **9**, **12**, and **17** reveal that a chloromethyl substituent at any position did not improve potency relative to **8**, the exception being **9** with a slightly higher affinity at α_{1D} -adrenoreceptors. However, it should be noted that when the substituent is at position 4, as in **17**, both affinity and selectivity profiles were not markedly influenced relative to **8**, whereas at position 2 and 3 the affinity for α_1 -adrenoreceptor subtypes was affected in such a way that **9**, bearing a 2-substituent, was highly selective (224–148-fold) for α_{1B} - and α_{1D} -adrenoreceptors versus the α_{1A} -subtype, and **12**, having a 3-substituent, resulted a weak and nonselective α_1 -adrenoreceptor antagonist.

Replacing chlorine atom of **9**, **12**, and **17** with an *N,N*-dimethylamino group afforded **10**, **14**, and **18**, respectively, which were almost as active as **8** at α_{1A} - and α_{1D} -adrenoreceptors while being significantly less potent at α_{1B} -adrenoreceptors. Consequently, the selectivity profile of these analogues was different from that of **8** and parent chlorides. It turned out that amines **10** and **14** had a reversed selectivity profile relative to **8** and **9** or **12**, respectively, being more potent at α_{1D} -adrenoreceptors, whereas amine **18** retained a selectivity profile similar to that of **8** and **17** owing to its higher affinity for α_{1B} -adrenoreceptors.

The finding that inserting an *N,N*-dimethylaminomethyl moiety onto the phenyl group of **8** resulted into

a marked effect on the selectivity profile prompted us to further modify the amine function, as in **13**, **15** and **19**. Thiazolidine analogues **15** and **19** did not display a better affinity profile than prototypes **14** and **18**, whereas **13** was as active as **14** at α_{1A} - and α_{1D} -adrenoreceptors, while showing a marked increase in affinity for the α_{1B} -subtype.

Finally, replacing the chlorine atom of **9**, **12**, and **17** with a substituted 1,6-diaminohexane moiety, affording **11**, **16**, and **20**, respectively, resulted in a marked effect on the affinity for α_1 -adrenoreceptors. Polyamine **11** was as active as, if not more active than, prototype **8**, whereas analogue **16**, bearing the *N,N*-dimethyl-1,6-diaminohexane moiety in position 3, turned out to be slightly or markedly less potent at α_{1D} - and α_{1B} -adrenoreceptors, respectively, while being more potent at the α_{1A} -subtype than **8**, and **20** retained a comparable affinity at α_{1B} -adrenoreceptors while being markedly less potent at the α_{1A} -subtype and more potent at the α_{1D} -subtype in comparison with **8**.

A most intriguing finding of the present investigation is the observation that polyamines **11**, **16**, and **20** retained high affinity for α_1 -adrenoreceptor subtypes, which suggests clearly that a 1,6-diaminohexane chain on the benzene ring did not give rise to negative interactions with the receptor. This observation may have relevance for the development of new quinazolines bearing a polyamine backbone on which additional substituents can be mounted to improve selectivity for α_1 -adrenoreceptor subtypes. Clearly, the site where the terminal aromatic ring of **8** interacts does not seem to present steric hindrance and particularly stringent requirements.

The binding affinities, expressed as pK_i values, in CHO cells expressing human cloned α_1 -adrenoreceptor subtypes and HeLa cells expressing human 5-HT_{1A} receptors and in membranes of rat cerebral cortex (α_2 -adrenoreceptors) and striatum (D₂ receptors) of quinazolines **2**, **3**, and **14** are shown in Table 2 in comparison with those of prazosin (**1**) and BMY-7378. As anticipated for cystazosin (**3**),²⁰ the results obtained in binding experiments did not show the same selectivity profile observed in functional assays. It can be seen that while binding affinities of reference compounds prazosin (**1**), **2**, and BMY-7378 are qualitatively and quantitatively comparable with pA_2 values derived from functional experiments, those observed for **3** and **14** are not in agreement at all from both a qualitative and a quantitative point of view with functional affinities. Both compounds were devoid of selectivity for α_1 -adrenore-

ceptor subtypes in binding assays, owing to a marked increase in affinity of about 2 orders of magnitude for α_{1a} - and α_{1b} -adrenoreceptors and of about 1 order of magnitude for the α_{1d} -subtype. As a matter of fact, the theory states that the affinity of an antagonist assessed in functional assays should not differ from that determined in binding experiments using both native and recombinant receptors. For this reason competitive antagonists are considered better tools for receptor characterization and classification than agonists because for the latter ones, in addition to affinity, other pharmacological parameters must be taken into account.^{35,36} Consequently, there is no apparent explanation for the discrepancy observed between our functional and binding results. However, very recently we discussed the possibility that if an antagonist does not adhere perfectly to the concept of neutral antagonism in the interaction with the receptor but behaves as a negative antagonist (inverse agonist), then its affinity may not be, as assumed by theory, system-independent, giving rise to affinity values which might be different according to the system employed for the determination.³⁵ In other words, as pointed out by Leff,³⁶ the use of inverse agonists as neutral antagonists may have, like agonists, problems since their estimated affinities are system-dependent. Thus, for inverse agonists the affinity values estimated in functional assays may not necessarily be comparable with those obtained in binding experiments. Interestingly, a survey of literature has revealed that some of the so-called competitive antagonists behave as inverse agonists when tested in the appropriate model. In the field of α_1 -adrenoreceptor antagonists, prazosin (**1**), WB 4101, and benoxathian were shown to be inverse agonists in a vascular model.³⁷ Thus, the difference, which is often observed for functional and binding affinities of antagonists, might be explained by the fact that these compounds are inverse agonists, and hence their affinity is system-dependent. Work is in progress to determine the nature of antagonism displayed by **3** and **14** and the other quinazolines of the present investigation.

Experimental Section

Chemistry. Melting points were taken in glass capillary tubes on a Büchi SMP-20 apparatus and are uncorrected. IR, electron impact (EI) mass, and ¹H NMR spectra were recorded on Perkin-Elmer 297, VG 7070E, and Varian VXR 300 instruments, respectively. Chemical shifts are reported in parts per million (ppm) relative to tetramethylsilane (TMS), and spin multiplicities are given as s (singlet), br s (broad singlet), d (doublet), t (triplet), or m (multiplet). Although the IR spectra data are not included (because of the lack of unusual features), they were obtained for all compounds reported and were consistent with the assigned structures. The elemental compositions of the compounds agreed to within $\pm 0.4\%$ of the calculated values. When the elemental analysis is not included, crude compounds were used in the next step without further purification. Chromatographic separations were performed on silica gel columns (Kieselgel 40, 0.040–0.063 mm, Merck) by flash chromatography. Compounds were named following IUPAC rules as applied by AUTONOM, a PC software for systematic naming in organic chemistry (Beilstein-Institut and Springer-Verlag).

6,7-Dimethoxy-*N*²-methyl-*N*²-[2-(2-methylaminoethyl)disulfanyl]ethylquinazoline-2,4-diamine (22**).** A mixture of **21** (1.0 g, 4.2 mmol) and *N,N*-dimethylcystamine²⁰ (3.5 g, 19.4 mmol) in *i*-AmOH (15 mL) was refluxed for 30 h. Removal of the solvent under reduced pressure gave a residue that was

purified by chromatography. Eluting with methylene chloride–methanol–aqueous 30% ammonia (9:1:0.05) afforded 0.4 g (25% yield) of **22** as the free base: ¹H NMR (CDCl₃) δ 2.03–2.39 (m, 2 + 1 exchangeable with D₂O), 2.43 (s, 3), 2.79–3.08 (m, 6), 3.21 (s, 3), 3.93 (s, 3), 3.97 (s, 3), 5.59 (br s, 2, exchangeable with D₂O), 6.81 (s, 1), 6.94 (s, 1).

5-Chloromethyl-2-furoyl Chloride (24**).** A solution of 5-hydroxymethylfuran-2-carboxylic acid²⁵ (0.3 g, 2.11 mmol) and SOCl₂ (1.5 mL, 10 mmol) in benzene (10 mL) was refluxed for 1 h. Removal of the solvent under reduced pressure afforded crude **24** in a quantitative yield.

5-Chloromethylfuran-2-carboxylic Acid (2-{2-[(4-Amino-6,7-dimethoxyquinazolin-2-yl)methylamino]ethyl}disulfanyl)ethylmethylamide Hydrochloride (4**).** A solution of **24** (0.19 g, 1.1 mmol) in THF (20 mL) was added dropwise to a solution of **22** (0.4 g, 1.1 mmol) in THF (15 mL). After the mixture was stirred at room temperature for 3 h, the solvent was removed under reduced pressure to give a residue that was purified by chromatography. Eluting with methylene chloride–methanol–aqueous 30% ammonia (9.5:0.5:0.05) afforded 0.34 g (65% yield) of **4** as the free base that was transformed into the hydrochloride salt: mp 170 °C (from EtOH/ether); ¹H NMR (free base; CDCl₃) δ 2.96–3.42 (m, 10), 3.82–4.05 (m, 4), 3.93 (s, 3), 3.97 (s, 3), 4.61 (s, 2) 5.32 (br s, 2, exchangeable with D₂O), 6.48 (d, 1), 6.85 (m, 1), 6.92–7.07 (m, 2). Anal. (C₂₂H₂₉Cl₂N₅O₄S) C, H, N.

5-Dimethylaminomethylfuran-2-carboxylic Acid (2-{2-[(4-Amino-6,7-dimethoxyquinazolin-2-yl)methylamino]ethyl}disulfanyl)ethylmethylamide Dioxalate (5**).** A 5.6 M ethanolic solution of dimethylamine (4 mL) was added to a solution of **4** (0.08 g, 0.14 mmol) in absolute ethanol (10 mL). After the mixture was stirred at room temperature for 2 h, the solvent was removed under reduced pressure to give a residue that was purified by chromatography. Eluting with methylene chloride–methanol–aqueous 30% ammonia (9.5:0.5:0.04) afforded 0.03 g (40% yield) of **5** as the free base that was transformed into the dioxalate salt and crystallized: mp 106 °C (from EtOH/ether); ¹H NMR (free base; CDCl₃) δ 2.25 (s, 6), 2.93–3.01 (m, 4), 3.20 (s, 6), 3.50 (s, 2), 3.73–3.93 (m, 10), 5.49 (br s, 2, exchangeable with D₂O), 6.28 (d, 1), 6.85–7.01 (m, 3). Anal. (C₂₆H₃₈N₆O₁₂S₂) C, H, N.

5-Chloromethylfuran-2-carboxylic Acid {6-[(4-Amino-6,7-dimethoxyquinazolin-2-yl)methylamino]hexyl}methylamide Hydrochloride (6**).** A solution of **24** (0.3 g, 1.68 mmol) in THF (20 mL) was added dropwise to a solution of **23**²⁴ (0.58 g, 1.7 mmol) in THF (15 mL). After the mixture was stirred at room temperature for 2 h, the solid was collected by filtration, washed with ether, and purified by chromatography. Eluting with methylene chloride–methanol–aqueous 30% ammonia (9.5:0.5:0.03) gave 0.16 g (20% yield) of **6** as the free base that was transformed into the hydrochloride salt and crystallized: mp 130 °C (from EtOH/ether); ¹H NMR (free base; CDCl₃) δ 1.18–1.69 (m, 8), 2.95–3.29 (m, 7), 3.42–3.71 (m, 5), 3.87 (s, 3), 3.92 (s, 3), 4.56 (s, 2), 5.49 (br s, 2, exchangeable with D₂O), 6.33–6.43 (m, 1), 6.81–7.03 (m, 3). Anal. (C₂₄H₃₃Cl₂N₅O₃) C, H, N.

5-Dimethylaminomethylfuran-2-carboxylic Acid {6-[(4-Amino-6,7-dimethoxyquinazolin-2-yl)methylamino]hexyl}methylamide Dihydrochloride (7**).** This was synthesized from **6** and dimethylamine following the procedure described for **5** and purified by chromatography. Eluting with methylene chloride–methanol–aqueous 30% ammonia (9.5:0.5:0.03) afforded 0.08 g (80% yield) of **7** as the free base that was transformed into the dihydrochloride salt and crystallized: mp 145 °C (from EtOH/ether); ¹H NMR (free base; CDCl₃) δ 1.18–1.69 (m, 8), 2.25 (s, 6), 3.01–3.22 (m, 7), 3.42–3.54 (m, 5), 3.66 (t, 2), 3.89 (s, 3), 3.94 (s, 3), 5.51 (br s, 2, exchangeable with D₂O), 6.27 (d, 1), 6.82–7.01 (m, 3). Anal. (C₂₆H₄₀Cl₂N₆O₄) C, H, N.

***N*-{6-[(4-Amino-6,7-dimethoxyquinazolin-2-yl)methylamino]hexyl}-*N*-methylbenzamide Hydrochloride (**8**).** A solution of benzoyl chloride (0.06 mL, 0.5 mmol) in dioxane (10 mL) was added dropwise to a solution of **23**²⁴ (0.17 g, 0.5 mmol) in dioxane (15 mL). After the mixture was stirred at

room temperature for 4 h, the solid was filtered and washed with ether affording **8**: 90% yield; mp 133–135 °C (from EtOH/ether); ¹H NMR (DMSO-*d*₆) δ 1.03–1.68 (m, 8), 2.86–2.94 (m, 3), 3.17–3.21 (m, 4), 3.32–3.44 (m, 1), 3.53–3.80 (m, 2), 3.84 (s, 3), 3.89 (s, 3), 7.31–7.43 (m, 5), 7.48 (s, 1), 7.72 (s, 1), 8.42 (br s, 1, exchangeable with D₂O), 11.85 (br s, 1, exchangeable with D₂O). Anal. (C₂₅H₃₄ClN₅O₃) C, H, N.

N-{6-[(4-Amino-6,7-dimethoxyquinazolin-2-yl)methylamino]hexyl}-2-formyl-*N*-methylbenzamide (**25**). 2-Formylbenzoyl chloride was synthesized in a quantitative yield from 2-formylbenzoic acid following the procedure described for **24**. A solution of this chloride (0.12 g, 0.72 mmol) in dioxane (5 mL) was added dropwise to a solution of **23** (0.25 g, 0.72 mmol) and triethylamine (0.1 mL, 0.72 mmol) in dioxane (10 mL). After the mixture was stirred at room temperature for 24 h, the solvent was removed under reduced pressure to give a residue that was purified by chromatography. Eluting with methylene chloride–methanol–aqueous 30% ammonia (9.5:0.5:0.04) afforded 0.3 g (87% yield) of **25** as the free base: ¹H NMR (CDCl₃) δ 1.20–1.82 (m, 8), 3.15–3.22 (m, 6), 3.86–3.93 (m, 4), 4.03 (s, 3), 4.10 (s, 3), 5.82 (br s, 2, exchangeable with D₂O), 7.19–7.15 (m, 2), 7.30–7.38 (m, 1), 7.89–7.95 (m, 2), 8.07 (d, 1), 10.11 (s, 1).

N-{6-[(4-Amino-6,7-dimethoxyquinazolin-2-yl)methylamino]hexyl}-2-hydroxymethyl-*N*-methylbenzamide Hydrochloride (**26**). Sodium borohydride (0.04 g, 1.0 mmol) was added portionwise to a cooled (0 °C) and stirred solution of **25** (0.5 g, 1.0 mmol) in absolute ethanol (18 mL). After the mixture was stirred and cooled for 3 h, water was added to the solution, and the solvent was removed under reduced pressure to give a residue that was purified by chromatography. Eluting with methylene chloride–methanol–aqueous 30% ammonia (9.5:0.5:0.04) afforded 0.37 g (79% yield) of **26** as the free base that was transformed into the hydrochloride salt: mp 172 °C (from EtOH/ether); ¹H NMR (free base; CDCl₃) δ 1.20–1.79 (m, 8), 3.07–3.21 (m, 6), 3.52–3.74 (m, 4 + 1 exchangeable with D₂O), 3.84 (s, 3), 3.92 (s, 3), 4.52 (s, 2), 5.38 (br s, 2, exchangeable with D₂O), 6.77–6.92 (m, 2), 7.31–7.44 (m, 4).

N-{6-[(4-Amino-6,7-dimethoxyquinazolin-2-yl)methylamino]hexyl}-2-chloromethyl-*N*-methylbenzamide Hydrochloride (**9**). A mixture of **26** (0.25 g, 0.52 mmol) and SOCl₂ (0.1 mL, 1.5 mmol) in anhydrous chloroform (20 mL) was refluxed for 2 h. Removal of the solvent gave 0.26 g of **9**: mp 155 °C (from EtOH/ether); ¹H NMR (DMSO-*d*₆) δ 1.10–1.97 (m, 8), 2.86–2.97 (m, 3), 3.10–3.24 (m, 3), 3.35–3.48 (m, 2), 3.61–3.81 (m, 2), 3.91 (s, 3), 3.97 (s, 3), 4.85 (s, 2), 7.28–7.45 (m, 5), 7.81 (s, 1), 8.58 (br s, 1, exchangeable with D₂O), 8.85 (br s, 1, exchangeable with D₂O), 11.85 (br s, 1, exchangeable with D₂O). Anal. (C₂₆H₃₅Cl₂N₅O₃) C, H, N.

N-{6-[(4-Amino-6,7-dimethoxyquinazolin-2-yl)methylamino]hexyl}-2-dimethylaminomethyl-*N*-methylbenzamide Dihydrochloride (**10**). This was synthesized from **9** (0.20 g, 0.37 mmol) following the procedure described for **5** and purified by chromatography. Eluting with methylene chloride–ethanol–aqueous 30% ammonia (9.2:0.8:0.04) gave 0.14 g (75% yield) of **10** as the free base that was transformed into the dihydrochloride: mp 158 °C (from EtOH/ether); ¹H NMR (CD₃OD) δ 1.08–1.80 (m, 8), 2.86 (s, 6), 3.01–3.15 (m, 3), 3.22–3.27 (m, 3), 3.60–3.81 (m, 4), 3.92 (s, 3), 3.98 (s, 3), 4.27 (s, 2), 7.29 (s, 1), 7.50–7.68 (m, 5), 8.58 (br s, 2, exchangeable with D₂O). Anal. (C₂₈H₄₂Cl₂N₆O₃) C, H, N.

N-{6-[(4-Amino-6,7-dimethoxyquinazolin-2-yl)methylamino]hexyl}-2-[(6-aminohexylamino)methyl]-*N*-methylbenzamide Trihydrochloride (**11**). A solution of **24** (0.1 g, 0.2 mmol) and 1,6-hexanediamine (0.23 g, 2.0 mmol) in toluene (50 mL) was refluxed, and the water formed was continuously removed for 24 h. The cooled mixture was filtered and the filtrate evaporated to give the corresponding Schiff base that was dissolved in ethanol (10 mL) and treated with NaBH₄ (0.07 g). The mixture was stirred at room temperature for 3 h, acidified with 2 N HCl (6 mL), made basic with 2 N NaOH, and finally extracted with chloroform. Removal of dried (Na₂SO₄) solvents gave a residue that was purified by chromatog-

raphy. Eluting with methylene chloride–ethanol–aqueous 30% ammonia (8:2:0.2) afforded 0.08 g (70% yield) of **11** as the free base that was transformed into the trihydrochloride: mp 148 °C (from EtOH/ether); ¹H NMR (CD₃OD) δ 1.10–1.89 (m, 16), 3.14–3.31 (m, 7), 3.38–3.47 (m, 3), 3.80–4.01 (m, 4), 4.10 (s, 3), 4.19 (s, 3), 4.38–4.42 (m, 2), 5.33 (br s, 2, exchangeable with D₂O), 7.50–7.57 (m, 1), 7.67–7.89 (m, 5). Anal. (C₃₂H₅₂Cl₃N₇O₃) C, H, N.

N-{6-[(4-Amino-6,7-dimethoxyquinazolin-2-yl)methylamino]hexyl}-3-chloromethyl-*N*-methylbenzamide Hydrochloride (**12**). This was synthesized from **23** (0.80 g, 2.3 mmol) and 3-chloromethylbenzoyl chloride (0.32 mL, 2.3 mmol) following the procedure described for **8**: 1.1 g (90% yield); mp 145 °C (from 2-PrOH/dioxane); ¹H NMR (DMSO-*d*₆) δ 1.03–1.94 (m, 8), 2.83–2.95 (m, 3), 3.17–3.22 (m, 4), 3.33–3.45 (m, 1), 3.58–3.80 (m, 2), 3.84 (s, 3), 3.90 (s, 3), 4.80 (s, 2), 7.30–7.45 (m, 5), 7.71 (s, 1), 8.56 (br s, 1, exchangeable with D₂O), 8.81 (br s, 1, exchangeable with D₂O), 11.57 (br s, 1, exchangeable with D₂O). Anal. (C₂₆H₃₅Cl₂N₅O₃) C, H, N.

N-{6-[(4-Amino-6,7-dimethoxyquinazolin-2-yl)methylamino]hexyl}-3-aminomethyl-*N*-methylbenzamide Dihydrochloride (**13**). A solution of **12** (0.05 g, 0.09 mmol) in ethanol (10 mL) was saturated with gaseous NH₃ at –10 °C. After the mixture was stirred at room temperature for 48 h, the solvent was removed under reduced pressure affording a residue that was purified by chromatography. Eluting with methylene chloride–methanol–aqueous 30% ammonia (9:1:0.1) gave **13** as the free base that was transformed into the dihydrochloride salt and crystallized: 0.04 g (80% yield); mp 150 °C (from EtOH/ether); ¹H NMR (CD₃OD) δ 1.10–1.83 (m, 8), 2.91–3.10 (m, 3), 3.17–3.36 (m, 4), 3.44–3.81 (m, 3), 3.92 (s, 3), 3.98 (s, 3), 4.18 (s, 2), 7.19–7.31 (m, 1), 7.39–7.63 (m, 5). Anal. (C₂₆H₃₈Cl₂N₆O₃) C, H, N.

N-{6-[(4-Amino-6,7-dimethoxyquinazolin-2-yl)methylamino]hexyl}-3-dimethylaminomethyl-*N*-methylbenzamide Dihydrochloride (**14**). This was synthesized from **12** (0.30 g, 0.56 mmol) following the procedure described for **5** and purified by chromatography. Eluting with methylene chloride–ethanol–aqueous 30% ammonia (9.2:0.8:0.04) afforded 0.15 g (87% yield) of **14** as the free base that was transformed into the dihydrochloride salt and crystallized: mp 165 °C (from EtOH/ether); ¹H NMR (DMSO-*d*₆) δ 1.03–1.73 (m, 8), 2.68 (s, 3), 2.70 (s, 3), 2.90–2.97 (m, 3), 3.21–3.25 (m, 4), 3.41–3.53 (m, 1), 3.62–3.80 (m, 2), 3.86 (s, 3), 3.90 (s, 3), 4.31–4.35 (m, 2), 7.38–7.63 (m, 5), 7.79 (s, 1), 8.56 (br s, 1, exchangeable with D₂O), 8.80 (br s, 1, exchangeable with D₂O), 11.01 (br s, 1, exchangeable with D₂O), 11.93 (br s, 1, exchangeable with D₂O); EI MS *m/z* 508 (M⁺). Anal. (C₂₈H₄₂Cl₂N₆O₃) C, H, N.

N-{6-[(4-Amino-6,7-dimethoxyquinazolin-2-yl)methylamino]hexyl}-*N*-methyl-3-thiazolidin-3-ylmethylbenzamide Dioxalate (**15**). A mixture of **12** (0.3 g, 0.56 mmol), thiazolidine (0.05 mL, 0.62 mmol), triethylamine (0.09 mL, 0.62 mmol), and KI (few crystals) in absolute ethanol (15 mL) was refluxed for 27 h. Removal of the solvent gave a residue that was purified by chromatography. Eluting with methylene chloride–methanol–aqueous 30% ammonia (9.5:0.5:0.04) afforded 0.16 g (52% yield) of **15** as the free base that was transformed into the dioxalate salt and crystallized: mp 101–103 °C (from EtOH/ether); ¹H NMR (free base; CDCl₃) δ 1.19–1.70 (m, 8), 2.95 (s, 3), 3.06 (s, 3), 3.16–3.25 (m, 4), 3.56–3.72 (m, 6), 3.89 (s, 3), 3.95 (s, 3), 4.03 (s, 2), 6.24 (br s, 1, exchangeable with D₂O), 6.60 (br s, 1, exchangeable with D₂O), 7.10–7.21 (m, 2), 7.25–7.46 (m, 4). Anal. (C₃₃H₄₄N₆SO₄) C, H, N.

N-{6-[(4-Amino-6,7-dimethoxyquinazolin-2-yl)methylamino]hexyl}-*N*-methyl-3-[(methyl(6-methylamino)hexyl)amino]methylbenzamide Trioxalate (**16**). A solution of **12** (0.13 g, 0.26 mmol) and *N,N*-dimethyl-1,6-hexanediamine²⁶ (0.38 g, 2.6 mmol) in ethanol (15 mL) was stirred at 60 °C for 1 h and at room temperature for 24 h. Removal of the solvent under reduced pressure gave a residue that was purified by chromatography. Eluting with methylene chloride–methanol–aqueous 30% ammonia (9:1:0.1) afforded 0.11 g (70% yield) of

16 as the free base that was transformed into the trioxalate salt: mp < 50 °C (from EtOH/ether); ¹H NMR (free base; CDCl₃) δ 1.15–1.83 (m, 16), 2.16–2.30 (m, 5), 2.46 (s, 3), 2.54–2.66 (m, 2), 2.85–3.04 (m, 3), 3.10–3.27 (m, 4), 3.38–3.77 (m, 5), 3.90 (s, 3), 3.94 (s, 3), 5.55 (br s, 2, exchangeable with D₂O), 6.87–7.01 (m, 2), 7.19–7.38 (m, 5). Anal. (C₄₀H₅₉N₇O₁₅) C, H, N.

N-{6-[(4-Amino-6,7-dimethoxyquinazolin-2-yl)methylamino]hexyl}-4-chloromethyl-N-methylbenzamide Hydrochloride (17). This was synthesized from **23** (0.8 g, 2.3 mmol) and 4-chloromethylbenzoyl chloride (0.43 g, 2.3 mmol) following the procedure described for **8** and purified by chromatography. Eluting with methylene chloride–ethanol–aqueous 30% ammonia (9.5:0.5:0.03) afforded 0.45 g (40% yield) of **17** as the free base that was transformed into the hydrochloride salt and crystallized: mp 162–164 °C (from EtOH/ether); ¹H NMR (DMSO-*d*₆) δ 1.03–1.94 (m, 8), 2.85–2.93 (m, 3), 3.17–3.22 (m, 4), 3.37–3.44 (m, 1), 3.58–3.78 (m, 2), 3.83 (s, 3), 3.87 (s, 3), 4.79 (s, 2), 7.31–7.42 (m, 2), 7.43–7.50 (m, 2), 7.61–7.66 (m, 1), 7.75 (s, 1), 8.56 (br s, 1, exchangeable with D₂O), 8.50–8.62 (br s, 1, exchangeable with D₂O), 8.86 (br s, 1, exchangeable with D₂O), 11.82 (br s, 1, exchangeable with D₂O). Anal. (C₂₆H₃₅Cl₂N₅O₃) C, H, N.

N-{6-[(4-Amino-6,7-dimethoxyquinazolin-2-yl)methylamino]hexyl}-4-dimethylaminomethyl-N-methylbenzamide Dihydrochloride (18). This was synthesized from **17** (0.11 g, 0.2 mmol) following the procedure described for **5** and purified by chromatography. Eluting with methylene chloride–ethanol–aqueous 30% ammonia (9.2:0.8:0.04) afforded 0.09 g (88% yield) of **18** as the free base that was transformed into the dihydrochloride salt and crystallized: mp 162–164 °C (from EtOH/ether); ¹H NMR (free base; CDCl₃) δ 1.03–1.71 (m, 8), 2.21 (s, 6), 2.90–3.27 (m, 7), 3.40 (s, 2), 3.43–3.60 (m, 3), 3.83 (s, 3), 3.92 (s, 3), 5.52 (br s, 2, exchangeable with D₂O), 6.88–6.93 (m, 2), 7.31 (s, 4). Anal. (C₂₈H₄₂Cl₂N₆O₃) C, H, N.

N-{6-[(4-Amino-6,7-dimethoxyquinazolin-2-yl)methylamino]hexyl}-N-methyl-4-thiazolidin-4-ylmethylbenzamide Dihydrochloride (19). This was synthesized from **17** (0.1 g, 0.19 mmol) following the procedure described for **15** and purified by chromatography. Eluting with methylene chloride–methanol–aqueous 30% ammonia (9.5:0.5:0.04) afforded 0.06 g (50% yield) of **19** as the free base that was transformed into the dihydrochloride salt and crystallized: mp 162 °C (from EtOH/ether); ¹H NMR (free base; CDCl₃) δ 1.18–1.73 (m, 8), 2.92–3.30 (m, 10), 3.49–3.70 (m, 6), 3.89 (s, 3), 3.95 (s, 3), 4.03 (s, 2), 5.50 (br s, 1, exchangeable with D₂O), 6.86 (s, 1), 6.90–7.11 (m, 1), 7.38 (s, 4). Anal. (C₂₈H₄₂Cl₂N₆O₃) C, H, N.

N-{6-[(4-Amino-6,7-dimethoxyquinazolin-2-yl)methylamino]hexyl}-N-methyl-4-{[methyl-(6-methylamino)hexyl]amino}methylbenzamide Trioxalate (20). This was synthesized from **17** (0.14 g, 0.28 mmol) and *N,N*-dimethyl-1,6-hexanediamine²⁶ following the procedure described for **16** and purified by chromatography. Eluting with methylene chloride–methanol–aqueous 30% ammonia (9:1:0.1) afforded 0.09 g (57% yield) of **20** as the free base that was transformed into the trioxalate salt: mp < 60 °C; ¹H NMR (free base; CDCl₃) δ 1.04–1.79 (m, 16), 2.10–2.33 (m, 5), 2.51 (s, 3), 2.58–2.68 (m, 2), 2.87–3.09 (m, 3), 3.13–3.32 (m, 4), 3.40–3.78 (m, 5), 3.93 (s, 3), 3.96 (s, 3), 5.60 (br s, 2, exchangeable with D₂O), 6.91–7.04 (m, 2), 7.30–7.41 (m, 5). Anal. (C₄₀H₅₉N₇O₁₅) C, H, N.

Biology. Functional Antagonism in Isolated Rat Tissues. Male Sprague Dawley rats (Charles River, Italy) were killed by cervical dislocation under ketamine anesthesia and the organs required were isolated, freed from adhering connective tissue, and set up rapidly under a suitable resting tension in 15 mL organ baths containing physiological salt solution kept at appropriate temperature (see below) and aerated with 5% CO₂:95% O₂ at pH 7.4. Concentration–response curves were constructed by cumulative addition of agonist. The concentration of agonist in the organ bath was increased approximately 5-fold at each step, with each addition being made only after the response to the previous addition

had attained a maximal level and remained steady. Contractions were recorded by means of a force displacement transducer (FT.03 Grass and 7003 Basile) connected to a four-channel pen recorder (Battaglia-Rangoni KV 380). In addition, parallel experiments in which tissues did not receive any antagonist were run in order to check any variation in sensitivity.

Vas Deferens Prostatic Portion. This tissue (from rats of 200–230 g) was used to assess α_{1A}-adrenergic antagonism.²⁷ Prostatic portions of 2 cm length were mounted under 400–450 g of tension at 37 °C in Tyrode solution of the following composition (mM): NaCl, 130.0; KCl, 2.0; CaCl₂·2H₂O, 1.8; MgCl₂, 0.89; NaHCO₃, 25.0; NaH₂PO₄·2H₂O, 0.42; glucose, 5.6. Desipramine hydrochloride (0.01 μM) was added to prevent the neuronal uptake of (–)-noradrenaline. The preparations were equilibrated for 45–60 min. During this time the bathing solution was changed every 10 min. Concentration–response curves for isotonic contractions in response to (–)-noradrenaline were obtained at 30 min intervals: the first one being discarded and the second one was taken as control. The antagonist was allowed to equilibrate with the tissue for 30 min, and then a new concentration–response curve to the agonist was obtained. (–)-Noradrenaline solutions contained 0.05% Na₂S₂O₅ to prevent oxidation.

α₂-Adrenoreceptor antagonist potency was determined also on vas deferens prostatic portions of 1.5–2 cm length which were set up in an organ bath containing Krebs solution of the following composition (mM): NaCl, 118.4; KCl, 4.7; CaCl₂, 2.52; MgSO₄, 0.6; KH₂PO₄, 1.2; NaHCO₃, 25.0; glucose, 11.1. (±)-Propranolol hydrochloride (1 μM) and desipramine (0.01 μM) were present in the Krebs solution throughout the experiments to block β-adrenoreceptor and neuronal uptake mechanisms, respectively. The medium was maintained at 37 °C. Field stimulation of the tissues was carried out by means of two platinum electrodes placed near the top and bottom of the vas deferens using 0.1 Hz square pulses of 3 ms duration at voltage of 20–40 V. The tissues were allowed to equilibrate for at least 1 h under a resting tension of 0.35 g before addition of any drug. A first clonidine concentration–response curve, taken as control, was obtained cumulatively, avoiding the inhibition of more than 90% of twitch. Under these conditions it was possible to obtain a second concentration–response curve not significantly different from the first one. The antagonist was allowed to equilibrate with the tissue for 30 min before a second cumulative concentration–response curve with agonist was made. Parallel experiments without any antagonist were run in order to determine the concentration of agonist causing 100% inhibition of twitch response.

Spleen. This tissue (from rats of 250–300 g) was employed to determine α_{1B}-adrenoreceptor antagonist potency.²⁸ The spleen was removed and bisected longitudinally into two strips which were suspended in tissue baths containing Krebs solution of the following composition (mM): NaCl, 118.4; KCl, 4.7; CaCl₂, 1.9; MgSO₄, 1.2; NaHCO₃, 25.0; NaH₂PO₄, 1.2; glucose, 11.7. Desipramine hydrochloride (0.01 μM) and (±)-propranolol hydrochloride (1 μM) were added to prevent the neuronal uptake of (–)-phenylephrine and to block β-adrenoreceptors, respectively. The spleen strips were placed under 1 g of resting tension and equilibrated for 1 h. The cumulative concentration–response curves to phenylephrine were measured isometrically and obtained at 30 min intervals, the first one being discarded and the second one taken as control. The antagonist was allowed to equilibrate with the tissue for 30 min, and then a new concentration–response curve to the agonist was constructed.

Aorta. This tissue (from rats of 250–300 g) was used to assess α_{1D}-adrenoreceptor antagonist potency.²⁸ Thoracic aorta was cleaned from extraneous connective tissue and placed in Krebs solution of the following composition (mM): NaCl, 118.4; KCl, 4.7; CaCl₂, 1.9; MgSO₄, 1.2; NaHCO₃, 25.0; NaH₂PO₄, 1.2; glucose, 11.7. Desipramine hydrochloride (0.01 μM) and (±)-propranolol hydrochloride (1 μM) were added to prevent the neuronal uptake of (–)-noradrenaline and to block β-adrenoreceptors, respectively. Two helical strips (15 mm

× 3 mm) were cut from each aorta beginning from the end most proximal to the heart. The endothelium was removed by rubbing with filter paper: the absence of acetylcholine-induced relaxation was taken as an indicator that vessel was denuded successfully. Vascular strips were then tied with surgical thread and suspended in a jacketed tissue bath containing Tyrode solution. Strips were secured at one end to Plexiglas hooks and connected to a transducer for monitoring changes in isometric contraction. After at least a 1 h equilibration period under an optimal tension of 1 g, cumulative (–)-noradrenaline concentration–response curves were recorded at 30 min intervals, the first two being discarded the third one taken as control. The antagonist was allowed to equilibrate with the tissue for 30 min before the generation of the fourth cumulative concentration–response curve to (–)-noradrenaline.

Radioligand Binding Assays. Native Receptors. Binding studies on native α_2 -adrenoreceptors and D_2 receptors were carried out in membranes of rat cerebral cortex and striatum, respectively. Male Sprague Dawley rats (200–300 g, SD Harlan/Nossan, Italy) were killed by cervical dislocation, and different tissues were excised and immediately frozen and stored at -70°C until use. For α_2 membrane preparations, cerebral cortices were homogenized (2×20 s) in 50 volumes of cold Tris-HCl buffer, pH 7.4, using a Politron homogenizer (speed 7). Homogenates were centrifuged at $49000g$ for 10 min, resuspended in 50 volumes of the same buffer, incubated at 37°C for 15 min, and centrifuged and resuspended 2 more times. The final pellets were suspended in 100 volumes of Tris-HCl buffer, pH 7.4, containing $10\ \mu\text{M}$ pargiline and 0.1% ascorbic acid. Membranes were incubated in a final volume of 1 mL for 30 min at 25°C with $0.5\text{--}1.5\ \text{nM}$ [^3H]rauwolscine, in absence or presence of competing drugs. For D_2 membrane preparations, rat striata were homogenized (2×20 s) in 30 volumes of cold Tris-HCl buffer, pH 7.4, using a Politron homogenizer (speed 7) and centrifuged at $49000g$ for 10 min. The final pellets were suspended in 200 volumes of Tris-HCl incubation buffer containing $10\ \mu\text{M}$ pargiline, 0.1% ascorbic acid, and the following saline concentrations: NaCl, 120 mM; KCl, 5 mM; CaCl_2 , 2 mM; MgCl_2 , 1 mM. The membranes were then incubated for 15 min at 37°C with $0.2\text{--}0.6\ \text{nM}$ [^3H]spiperone. Nonspecific binding was determined in the presence of $10\ \mu\text{M}$ phentolamine (α_2 -adrenoreceptors) and $1\ \mu\text{M}$ (+)-butaclamol (D_2 receptors). The incubation was stopped by addition of ice-cold Tris-HCl buffer and rapid filtration through 0.2% polyethyleneimine pretreated Whatman GF/B or Schleicher & Schuell GF52 filters. The filters were then washed with ice-cold buffer, and the radioactivity retained on the filters was counted by liquid scintillation spectrometry.

Cloned Receptors. Binding to cloned human α_1 -adrenoreceptor subtypes was performed in membranes from CHO cells transfected by electroporation with DNA expressing the gene encoding each α_1 -adrenoreceptor subtype. Cloning and stable expression of the human α_1 -adrenoreceptor gene was performed as previously described.³² CHO cell membranes (30 μg proteins) were incubated in 50 mM Tris-HCl, pH 7.4, with $0.1\text{--}0.4\ \text{nM}$ [^3H]prazosin, in a final volume of 1.02 mL for 30 min at 25°C , in absence or presence of competing drugs ($1\ \text{pM}\text{--}10\ \mu\text{M}$). Nonspecific binding was determined in the presence of $10\ \mu\text{M}$ phentolamine. The incubation was stopped by addition of ice-cold Tris-HCl buffer and rapid filtration through 0.2% polyethyleneimine pretreated Whatman GF/B or Schleicher & Schuell GF52 filters. Genomic clones G-21 coding for the human 5-HT_{1A} receptor are stably transfected in a human cell line (HeLa).³³ HeLa cells were grown as monolayers in Dulbecco's modified Eagle's medium (DMEM), supplemented with 10% fetal calf serum, gentamicin (100 $\mu\text{g}/\text{mL}$), and 5% CO_2 at 37°C . Cells were detached from the growth flask at 95% confluence by a cell scraper and were lysed in ice-cold Tris 5 mM and EDTA 5 mM buffer (pH 7.4). Homogenates were centrifuged at $40000g$ for 20 min, and pellets were resuspended in a small volume of ice-cold Tris 5 mM and EDTA 5 mM buffer (pH 7.4) and immediately frozen and stored at -70°C until use. On the experimental day, cell

membranes were resuspended in binding buffer of 50 mM Tris-HCl (pH 7.4), 2.5 mM MgCl_2 , and $10\ \mu\text{M}$ pargiline.³⁴ Membranes were incubated in a final volume of 1 mL for 30 min at 30°C with $0.7\text{--}1.4\ \text{nM}$ [^3H]8-OH-DPAT, in the absence or presence of competing drugs. Nonspecific binding was determined in the presence of $10\ \mu\text{M}$ 5-HT. The incubation was stopped by addition of ice-cold Tris-HCl buffer and rapid filtration through 0.2% polyethyleneimine pretreated Whatman GF/B or Schleicher & Schuell GF52 filters.

Data Analysis. The affinity constants (pA_2 values, Table 1) were determined by Schild plots²⁹ obtained from the dose ratios at the EC_{50} values of the agonists calculated at three antagonist concentrations. Each concentration was tested five times, and Schild plots were constrained to a slope of -1 .³⁰ The compounds were tested at only one concentration when determining α_2 -adrenoreceptor blocking activity because of their low affinity for this receptor. In these cases, pA_2 ($-\log K_B$) values were calculated according to van Rossum.³¹ Functional data were analyzed by pharmacological computer programs³⁸ and are presented as the mean \pm SE of n experiments. Differences between mean values were tested for significance by Student's t -test.

Binding data were analyzed using the nonlinear curve-fitting program Allfit.³⁹ Scatchard plots were linear in all preparations. All pseudo-Hill coefficients (nH) were not significantly different from unity ($p > 0.05$). Equilibrium inhibition constants (K_i) were derived from the Cheng–Prusoff equation,⁴⁰ $K_i = \text{IC}_{50}/(1 + L/K_d)$, where L and K_d are the concentration and the equilibrium dissociation constant of the radioligand. pK_i values (Table 2) are the mean \pm SE of three separate experiments performed in triplicate.

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