



High affinity of σ_1 -binding sites for sterol isomerization inhibitors: evidence for a pharmacological relationship with the yeast sterol C_8-C_7 isomerase

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1 The σ -drug binding site of guinea-pig liver is carried by a protein which shares significant amino acid sequence similarities with the yeast sterol C_8-C_7 isomerase (ERG2 protein). Pharmacologically - but not structurally - the σ_1 -site is also related to the emopamil binding protein, the mammalian sterol C_8-C_7 isomerase. We therefore investigated if sterol C_8-C_7 isomerase inhibitors are high affinity ligands for the (+)-[³H]-pentazocine labelled σ_1 -binding site.

2 Among the compounds which bound with high affinity to native hepatic and cerebral as well as to yeast expressed σ_1 -binding sites were the agricultural fungicide fenpropimorph (K_i 0.005 nM), the antihypercholesterinaemic drugs triparanol (K_i 7.0 nM), AY-9944 (K_i 0.46 nM) and MDL28,815 (K_i 0.16 nM), the enantiomers of the ovulation inducer clomiphene (K_i 5.5 and 12 nM, respectively) and the antioestrogene tamoxifen (K_i 26 nM).

3 Except for tamoxifen these affinities are essentially identical with those for the [³H]-ifenprodil labelled sterol C_8-C_7 isomerase of *S. cerevisiae*. This demonstrates that σ_1 -binding protein and yeast isomerase are not only structurally but also pharmacologically related. Because of its affiliations with yeast and mammalian sterol isomerases we propose that the σ_1 -binding site is localized on a sterol isomerase related protein, involved in postsqualene sterol biosynthesis.

Keywords: Ergosterol; cholesterol; ERG2; sterol C_8-C_7 isomerase; σ_1 -binding site; AY-9944; fenpropimorph; triparanol; tamoxifen; clomiphene

Introduction

The so-called σ -receptor was originally postulated to account for the psychomimetic effects elicited by benzomorphan opioids such as N-allylnormetazocine and pentazocine (Martin *et al.*, 1976). Subsequent studies revealed that σ -sites were distinct from opiate and NMDA/phencyclidine-receptors (Walker *et al.*, 1990). Despite much data on *in vitro* or *in vivo* effects mediated by σ -sites no physiological or biochemical function was unequivocally shown to be associated with them. In contrast to function, the binding characteristics of σ -sites are well defined and allow the distinction of two subtypes (Quirion *et al.*, 1992). The σ_1 - is distinguished from the σ_2 -subtype by its higher affinity for the benzomorphan (+)-pentazocine. Several clinically used drugs such as the antipsychotic haloperidol (K_i 0.2 nM), the tricyclic antidepressant opipramol (K_i 0.3 nM) and the opioid analgesic pentazocine (K_i 1.7 nM) interact with σ_1 -binding sites with high affinity (Hanner *et al.*, 1996). New drugs developed for other receptor systems are also often active in σ -binding assays. However, the biological consequences of these high affinity interactions are unknown. To clarify the pharmacological significance of drug binding to σ -sites we isolated and cloned the cDNA of the high affinity (+)-[³H]-pentazocine binding protein (Hanner *et al.*, 1996). Its amino acid sequence showed significant similarities with sterol C_8-C_7 isomerases from fungi (Hanner *et al.*, 1996). This enzyme is essential for the biosynthesis of ergosterol which is the fungal equivalent of cholesterol.

In mammals cholesterol biosynthesis after cyclization of squalene and formation of lanosterol involves several enzymatic steps such as the saturation of the C24(25) double bond in the side chain, the C14-demethylation, isomerization of the

C8(9) double bond to C7(8), introduction of the C5(6) double bond and removal of the C7(8) double bond (Trzaskos *et al.*, 1982). Whereas all relevant genes of the yeast sterol pathway have been cloned from *S. cerevisiae* (Lees *et al.*, 1995) most of the cDNAs encoding the mammalian enzymes are yet unknown. The recent discovery that the biochemical defect of the inborn Smith-Lemli-Opitz syndrome which is associated with mental retardation and genital malformations results in the accumulation of 7-dehydrocholesterol (Tint *et al.*, 1994) and desmosterol (Clayton *et al.*, 1996) revealed the pivotal importance of mammalian postsqualene cholesterol biosynthesis.

Some of the compounds interfering with the late steps of cholesterol biosynthesis impair sterol C_8-C_7 isomerization resulting in the accumulation of zymosterol. The striking structural similarities between mammalian σ_1 -binding protein and yeast sterol isomerases (Hanner *et al.*, 1996) suggested a pharmacological and functional relationship. We now demonstrate that drugs which were shown to interfere with sterol isomerization inhibit (+)-[³H]-pentazocine binding to liver, brain and yeast expressed σ_1 -sites with high affinities which are similar to those for the sterol C_8-C_7 isomerase of *S. cerevisiae*. A pharmacological relationship between σ_1 -binding site and the mammalian sterol C_8-C_7 isomerase (which is identical with the emopamil binding protein (Moebius *et al.*, 1993; Silve *et al.*, 1996a)) has already been suggested previously (Moebius *et al.*, 1994). Our findings insinuate an affiliation of σ_1 -sites with postsqualene cholesterol biosynthesis.

Methods

Binding assays

(±)-[³H]-pentazocine (0.6 nM) or [³H]-ifenprodil (0.6 nM) were incubated in 0.5 or 2.5 ml 25 mM Tris-HCl (pH 9 at 4°C, pH 8.3 at 22°C) for 12 h at 22°C with 2 to 35 μ g ml⁻¹ microsomal protein. Non-specific binding was measured in the presence of

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1 μM of unlabelled drug. Serial dilutions of competing drugs were prepared in dimethylsulphoxide (Boer *et al.*, 1989) and added directly to the assay. The final dimethylsulphoxide (DMSO) concentration was $\leq 1\%$, which did not affect specific binding. For the separation of bound and free ligand we used Whatman GF/C filters presoaked in 0.3% (w/v) polyethyleneimine. Filters were washed with icecold 10 mM Tris-HCl (pH 9 at 4°C). Binding parameters were calculated by non linear curve fitting to a rectangular hyperbola (K_d (dissociation constant), B_{max} (maximal density of binding sites)) or the general dose-response equation (IC_{50} (concentration causing half-maximal inhibition), slope factors, (DeLean *et al.*, 1978)). K_i (inhibition constants) values were calculated according to Linden (1982).

Membrane preparation

Microsomes from yeast strain WA0 (a his7-2 leu2-3,112 ura3-52 erg2-3) overexpressing the σ_1 -binding site ($6 \times \text{HIS-}\lambda\text{GP8-ORF}$, (Hanner *et al.*, 1996)) and from strain JB811 (ade2-1 leu2-3,112 pep4-3 trp1-289 ura3-52) were prepared as described by Moebius *et al.* (1996). Guinea-pig liver and whole brain microsomes were prepared by homogenization with a glass-Teflon homogenizer in ice-cold 0.25 M sucrose, 10 mM Tris-HEPES pH 7.4 (4°C). The homogenate was centrifuged at $8,000 \times g$ and the resulting supernatant was collected by centrifugation at $100,000 \times g$. The pellet was resuspended in 0.5 M KCl, 0.15 M Tris-HCl pH 8.0 (4°C) and centrifuged at $100,000 \times g$. The final pellet was resuspended in 5% (w/v) glycerol, 20 mM Tris-HCl pH 9 (4°C) at a protein concentration of 4 to 8 mg ml⁻¹, shock-frozen in liquid nitrogen and stored at -80°C. Protein concentrations were determined according to Bradford (1976), with bovine serum albumin (BSA) as a standard.

Materials

(+)-[³H]-pentazocine (32 Ci mmol⁻¹) and [³H]-ifenprodil (44 Ci mmol⁻¹) were obtained from NEN (Vienna, Austria). Chemicals were obtained from the following sources: zuclophene, enclomiphene, triparanol and MDL28,815 (M-

[1,5,9)-trimethyldecyl]-4,10-dimethyl-8-aza-trans-decal-3 β -ol) Hoechst Marion Roussel Research Institute (Cincinnati, OH); L-690,404 (1-butyl-3,4-dihydrospiro[naphthalene-1-(2H),4'-piperidine]) Merck Sharp & Dohme (Harlow, U.K.); fenpropimorph and tridemorph, BASF (Limburgerhof, Germany); AY-9944 (1,4-bis-(2-chlorobenzylaminomethyl)cyclohexane) Dr P. Benveniste (Strasbourg, France); trifluoperidol, RBI (Natick, MA); Bradford Protein Reagent, Bio-Rad (Vienna, Austria). All other chemicals were obtained from Sigma (Vienna, Austria). *S. cerevisiae* strain WA0 was kindly provided by Dr M. Bard (Indianapolis, IN, U.S.A.); strain JB811 was from Dr K. Nasmyth (Vienna, Austria).

Results

We examined the interaction of inhibitors of sterol isomerization with native σ_1 -binding sites in guinea-pig brain and liver microsomes as well as with the σ_1 -binding site expressed in *S. cerevisiae* strain WA0 (Ashman *et al.*, 1991) which is devoid of sterol C₈-C₇ isomerase activity. Although (+)-[³H]-pentazocine is a σ_1 -selective ligand which has low affinity for the endogenous σ_1 -ligand binding site of *S. cerevisiae* associated with the sterol C₈-C₇ isomerase (ERG2 gene product) (Moebius *et al.*, 1996) we wanted to rule out any interference with endogenous isomerase. Transformation with the cDNA encoding the σ_1 -receptor ($6 \times \text{HIS-}\lambda\text{GP8-ORF}$, (Hanner *et al.*, 1996)) induced the formation of high affinity (+)-[³H]-pentazocine binding sites (Figure 1a) which were sensitive to haloperidol and ditolylguanidine (Figure 1b). The expressed (+)-[³H]-pentazocine binding sites also showed high affinities for L-609,404, a spiro-piperidine, (Table 1, Figure 1b) (Chambers *et al.*, 1992) and amiodarone, an antiarrhythmic drug which is also a σ_1 -ligand (Cagnotto *et al.*, 1994) (K_i 1.7 ± 0.5 nM ($n=3$)).

Inhibitors of fungal ergosterol biosynthesis

The haloperidol analogue trifluoperidol is clinically used as an antipsychotic. Besides its prominent antidopaminergic activity trifluoperidol also inhibits the yeast sterol C₈-C₇ iso-

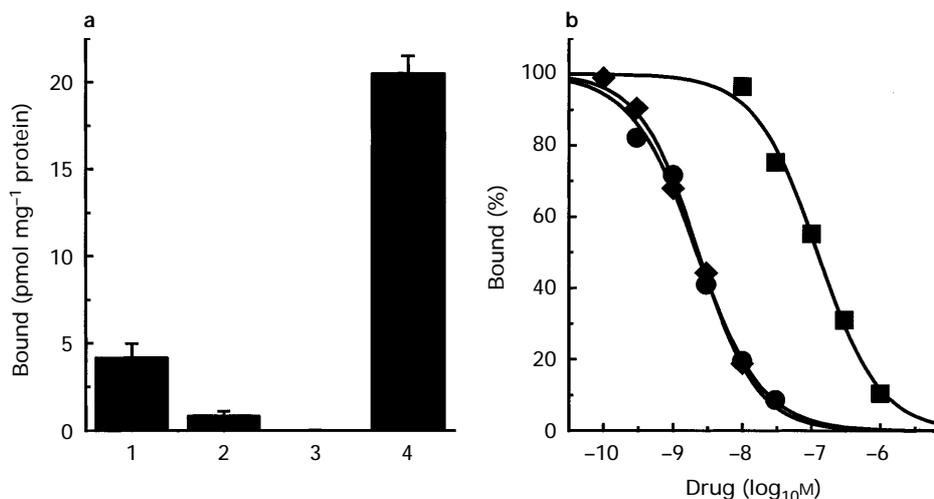


Figure 1 Characterization of (+)-[³H]-pentazocine binding to the recombinant σ_1 -binding site. (a) Binding of 0.6 nM (+)-[³H]-pentazocine to microsomes from guinea-pig liver (1), brain (2) and yeast cells (strain WA0, (Ashman *et al.*, 1991)) transformed with the vector (3) or the $6 \times \text{HIS-}\lambda\text{GP8-ORF}$ DNA (Hanner *et al.*, 1996) (4). Data shown are the mean \pm s.d. of three experiments. (b) Inhibition of (+)-[³H]-pentazocine (0.6 nM) binding to the yeast expressed σ_1 -binding site (4 μg microsomal protein ml⁻¹) by (+)-pentazocine (●, IC_{50} 2.2 nM), ditolylguanidine (■, IC_{50} 122 nM) and L-609,404 (◆, IC_{50} 2.3 nM). Data shown are the mean of duplicate determinations.

merase (Sobus *et al.*, 1977). It differs from haloperidol by a trifluoromethyl group instead of a chlorine atom and has modestly lower affinity (K_i 0.83 ± 0.24 nM ($n=3$)) than haloperidol (K_i 0.2 nM (Hanner *et al.*, 1996)) for the liver (+)-[³H]-pentazocine binding site (Table 1, Figure 2a). Fenpropimorph is an agricultural fungicide which inhibits the yeast sterol isomerase (Marcireau *et al.*, 1990). With K_i values for the liver and brain (+)-[³H]-pentazocine binding sites of 0.011 ± 0.006 nM ($n=3$) and 0.005 ± 0.004 nM ($n=3$), respectively, fenpropimorph is a sigma₁-ligand with the highest affinity found so far. Its analogue tridemorph displayed similarly high affinity for the brain binding site (K_i 0.023 ± 0.005 nM ($n=3$)). Because of this extraordinarily high affinity the K_i determination critically depends on the receptor concentration (Knaus *et al.*, 1995). Indeed when the IC₅₀ values were determined under standard conditions for liver, brain and yeast expressed (+)-[³H]-pentazocine binding sites (IC₅₀ = 0.10 ± 0.01 nM ($n=3$), 0.10 ± 0.03 nM ($n=3$) and 0.31 ± 0.11 nM ($n=3$), respectively) they corresponded to half of the receptor concentration (0.23 ± 0.04 , 0.18 ± 0.03 and 0.54 ± 0.11 nM, respectively). Apparent Hill slopes significantly different from unity (data not shown) also suggested that the number of receptors exceeded the number of drug molecules (Knaus *et al.*, 1995). Reduction of receptor

concentration and an increased assay volume (2.5 instead of 0.5 ml) remedied these problems (Figure 2a, Table 1).

Inhibitors of postsqualene cholesterol biosynthesis

MDL28,815, a rationally designed inhibitor of sterol biosynthesis, resembles cholesterol and mimics the high energy reaction intermediates of several of the late steps of sterol modification (Gerst *et al.*, 1988). Despite a chemical structure different from that of morpholines its affinity for the (+)-[³H]-pentazocine labelled sigma₁-site of guinea-pig brain microsomes was close (K_i 0.16 ± 0.04 nM ($n=3$)) to that of tridemorph (Table 1). The structure of the antihypercholesterolaemic drug AY-9944 differs from fenpropimorph and MDL28,815. AY-9944 was therefore employed to confirm further that the high affinity interaction of sterol isomerase inhibitors with the sigma₁-site did not simply reflect a chemical relationship. Indeed AY-9944, which is a symmetric molecule with two nitrogen atoms, displayed high affinity for the (+)-[³H]-pentazocine labelled sigma₁-binding site in brain (K_i 0.46 ± 0.10 nM). Triparanol was developed for the treatment of hypercholesterinaemia. It showed high affinity for the (+)-[³H]-pentazocine labelled brain sigma₁-binding site (K_i 7.0 ± 0.8 nM ($n=3$)). Several other compounds which are

Table 1 Comparison of the inhibition of (+)-[³H]-pentazocine binding to native and heterologously expressed sigma₁-sites and to the [³H]-ifenprodil binding site of the yeast sterol C₈-C₇ isomerase

	Liver		(+)-[³ H]-pentazocine Brain		Yeast Expressed		[³ H]-ifenprodil <i>S. cerevisiae</i> K _i (nM)
	K _i (nM)	Slope	K _i (nM)	Slope	K _i (nM)	Slope	
<i>σ-Ligands</i>							
L-609,404	0.63 ± 0.08	1.02 ± 0.12	0.60 ± 0.11	0.93 ± 0.02	1.3 ± 0.4	0.93 ± 0.08	4.7 ± 0.6
Amiodarone	1.4 ± 0.2	1.07 ± 0.06	2.1 ± 0.7	0.95 ± 0.15	1.7 ± 0.5	0.90 ± 0.03	62 ^a
<i>Isomerase inhibitors</i>							
Trifluoperidol	0.83 ± 0.24	0.82 ± 0.04	1.3 ± 0.1	0.93 ± 0.17	3.6 ± 1.3	0.94 ± 0.09	0.15 ^a
Fenpropimorph	0.011 ± 0.006	1.07 ± 0.25	0.005 ± 0.004	1.01 ± 0.05	0.08 ± 0.02	0.98 ± 0.10	0.05 ^a
Tridemorph	0.039 ± 0.023	1.07 ± 0.17	0.023 ± 0.005	1.13 ± 0.15	0.10 ± 0.04	1.11 ± 0.03	0.09 ^a
MDL28,815	0.48 ± 0.17	1.65 ± 0.28	0.16 ± 0.04	1.46 ± 0.32	0.49 ± 0.11	1.27 ± 0.23	0.44 ^a
AY-9944	0.50 ± 0.10	1.39 ± 0.14	0.46 ± 0.10	1.27 ± 0.22	0.99 ± 0.12	1.20 ± 0.09	5.8 ^a
Triparanol	8.2 ± 0.6	1.09 ± 0.02	7.0 ± 0.8	0.92 ± 0.10	15 ± 2	1.07 ± 0.06	1.5 ^a
Tamoxifen	34 ± 3	1.27 ± 0.08	26 ± 5	0.84 ± 0.05	54 ± 12	1.16 ± 0.20	1,470 ± 95
Zuclomiphene	4.7 ± 0.6	1.06 ± 0.15	5.5 ± 0.4	1.05 ± 0.07	12 ± 2	1.08 ± 0.03	1.6 ± 0.4
Enclomiphene	7.7 ± 0.5	0.93 ± 0.13	12 ± 2	0.87 ± 0.03	17 ± 4	0.99 ± 0.03	164 ± 26

Binding conditions are given in Methods. Data shown are the mean ± s.d. ($n=3$). ^aData taken from Moebius *et al.* (1996).

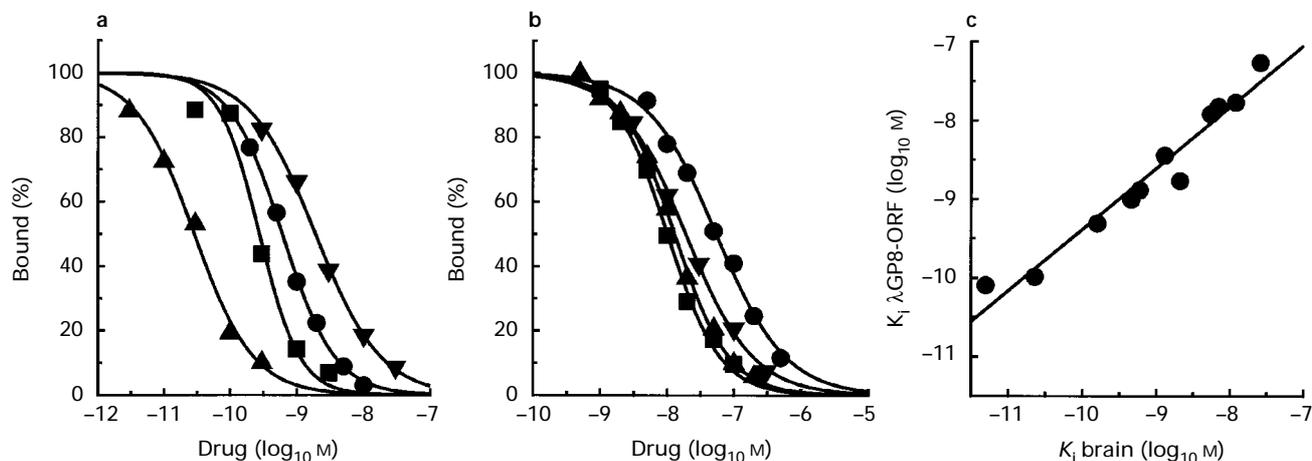


Figure 2 Inhibition of (+)-[³H]-pentazocine binding by sterol isomerase inhibitors. (a) (+)-[³H]-pentazocine 0.6 nM was incubated with brain microsomes in a final volume of 0.5 ml (fenpropimorph 2.5 ml) and the inhibition by fenpropimorph (▲), MDL28,815 (■), AY-9944 (●) and trifluoperidol (▼), and (b) triparanol (▲), zuclomiphene (■), enclomiphene (●) was measured. Data shown are the mean of duplicate determinations. (c) Comparison of the K_i values (Table 1) obtained for the brain (x-axis) and the yeast expressed (y-axis) (+)-[³H]-pentazocine binding sites ($r=0.98$).

structurally related to triparanol were also tested. The racemic mixture of the oestrogen receptor antagonist tamoxifen showed high affinity (K_i 26 ± 5 nM ($n=3$)). This binding was not stereoselective as shown by the essentially identical affinities of the *cis*- and *trans*-stereoisomers of the tamoxifen analogue clomiphene, zuclomiphene and enclomiphene (K_i 5.5 ± 0.4 nM ($n=3$) and 12 ± 2 nM ($n=3$), respectively).

The affinities for the native and expressed (+)-[³H]-pentazocine binding site correlate

The cDNA 6×HIS-λGP8-ORF (Hanner *et al.*, 1996) induces the formation of a high affinity binding site with all characteristics of the native brain and liver sigma₁-binding sites. Therefore, we compared the affinities of the isomerase inhibitors for native and recombinant (+)-[³H]-pentazocine labelled sites. In agreement with previous findings (Hanner *et al.*, 1996) the affinity of the heterologously expressed protein for most compounds is slightly reduced. This might be due to differences in the lipid composition between yeast and guinea-pig microsomes (Hanner *et al.*, 1996). However, we found an excellent correlation between the K_i values for the guinea-pig brain and the yeast expressed (+)-[³H]-pentazocine binding sites ($r=0.98$, Figure 2c). This strengthens our claim that the recently cloned cDNA (Hanner *et al.*, 1996) encodes the sigma₁-site of guinea-pig liver and brain.

Discussion

Pharmacological similarities of the sigma₁-site with the yeast sterol isomerase

The guinea-pig sigma₁-binding protein shares striking amino acid sequence similarities with the ERG2 gene product of *S. cerevisiae* (Hanner *et al.*, 1996) which encodes a sterol C₈-C₇ isomerase (Ashman *et al.*, 1991). The ERG2 protein also carries a high affinity binding site for the sigma-ligands [³H]-ifenprodil and [³H]-haloperidol but not for (+)-[³H]-pentazocine (Moebius *et al.*, 1996) in line with the inhibition of the yeast sterol C₈-C₇ isomerase by the sigma-ligand SR31747 (Silve *et al.*, 1996b). We now demonstrate that both proteins not only share structural but also pharmacological properties. Several structurally diverse drugs show only minor differences in their affinity for the [³H]-ifenprodil labelled sterol C₈-C₇ isomerase of *S. cerevisiae* and the mammalian (+)-[³H]-pentazocine binding site (Table 1). For tridemorph, MDL28,815 and zuclomiphene this difference was <5 fold, for L-690,404, AY-9944, fenpropimorph, triparanol and enclomiphene it was <15 fold and only tamoxifen showed a >50 fold reduced affinity for the yeast [³H]-ifenprodil binding site as compared to the sigma₁-site. Interestingly the yeast sterol C₈-C₇ isomerase preferred zuclomiphene (K_i 1.6 nM) to enclomiphene (K_i 164 nM) whereas the sigma₁-site was essentially not stereoselective for clomiphene. The structural and pharmacological similarities of mammalian sigma₁-binding protein and yeast sterol C₈-C₇ isomerase suggests that both proteins originated from a common precursor.

Sterol isomerase inhibitors: a new class of sigma-ligands

Fenpropimorph and tridemorph are agricultural fungicides which inhibit sterol C₈-C₇ isomerases of fungi and plants (Schmitt *et al.*, 1981; Rahier *et al.*, 1986; Marcireau *et al.*, 1990). The ultrahigh affinity of fenpropimorph for the brain α₁-site (K_i 5 pM) suggests that this morpholine might be a valuable tool for studies on sigma₁-mediated effects. The haloperidol congener trifluoperidol (K_i 1.3 nM) inhibits the sterol C₈-C₇ isomerase of yeast (Sobus *et al.*, 1977) whereas the azadecalin MDL28,815 (K_i 0.16 nM) inhibits the mammalian sterol C₈-C₇ isomerase in 3T3 mouse fibroblasts (Gerst *et al.*, 1988) as well as in the human hepatoma cell

line HepG2 (van Sickle *et al.*, 1993). The anti-hypercholesterolaemic drug AY-9944 (K_i 0.46 nM) reduces sterol C₈-C₇ isomerization in yeast (Pereira *et al.*, 1983) and rat adrenal gland (Givner *et al.*, 1967). It also inhibits the rat sterol C₈-C₇ isomerase *in vitro* (Paik *et al.*, 1986). Triparanol (K_i 7.0 nM) was used in man for the treatment of hypercholesterinaemia but was withdrawn from marketing because of cataract induction (Kirby *et al.*, 1962). In yeast and rat hepatoma cells triparanol is an inhibitor of sterol C₈-C₇ isomerization (Campagnoni *et al.*, 1977; Popják *et al.*, 1989). Clomiphene induces ovulation and is used to treat female sterility. Its stereoisomer zuclomiphene (K_i 5.5 nM) inhibits the sterol C₈-C₇ isomerase of the developing rat brain (Ramsey *et al.*, 1977). Tamoxifen (K_i 26 nM) is an antioestrogenic drug which is used to treat hormone-dependent breast cancer. Sterol C₈-C₇ isomerase activity is compromised in tamoxifen treated women (Gylling *et al.*, 1995).

The propensity of sigma-sites to bind chemically diverse drugs from various pharmacological classes such as antidepressants, opioids and antipsychotics is well known. However, α-sites fail to bind all representatives of the respective pharmacological class of drugs. For example their affinity for the dopamine D₂-antagonists haloperidol and chlorpromazine is high whereas it is low for risperidone (Leysen *et al.*, 1992), sulpiride and spiperone (Walker *et al.*, 1990). We therefore consider sterol isomerase inhibitors to be the first class of sigma-ligands with diverse chemical structures but a common biochemical effect. It remains to be clarified whether *vice versa* all sigma₁-ligands are inhibitors of sterol C₈-C₇ isomerization in brain which contains high densities of α₁-sites.

Conclusion

Blockers of eukaryotic sterol C₈-C₇ isomerization potently inhibit (+)-[³H]-pentazocine binding to sigma-sites. Their affinities are similar to those for the sterol C₈-C₇ isomerase of *S. cerevisiae* except for tamoxifen. Despite its structural and pharmacological relationship with the yeast isomerase the overexpressed α₁-binding protein failed to complement the sterol C₈-C₇ isomerization deficiency of yeast mutants (Hanner *et al.*, 1996). It is therefore unclear whether the protein carrying the sigma₁-site is not only a structural homologue and a pharmacological relative but also a functional equivalent of the yeast isomerase. A mammalian sterol C₈-C₇ isomerase cloned by complementation of an isomerase deficient yeast strain (Silve *et al.*, 1996a) is identical with the emopamil binding protein (Zech *et al.*, 1991; Moebius *et al.*, 1993; 1994; Hanner *et al.*, 1995). The latter is pharmacologically related to the sigma₁-site although it bears no structural similarities. The pharmacological and structural affiliations of sigma₁-sites with yeast and mammalian sterol C₈-C₇ isomerases imply a related function. We therefore propose that the α₁-binding site is carried by a sterol isomerase-related protein involved in post-squalene cholesterol biosynthesis. Its precise biochemical function remains to be determined by targeted disruption of the murine gene.

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