
Medicinal Chemistry of σ_1 Receptor Ligands: Pharmacophore Models, Synthesis, Structure Affinity Relationships, and Pharmacological Applications

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Abstract

In the first part of this chapter, we summarize the various pharmacophore models for σ_1 receptor ligands. Common to all of them is a basic amine flanked by two hydrophobic regions, representing the pharmacophoric elements. The development of computer-based models like the 3D homology model is described as

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well as the first crystal structure of the σ_1 receptor. The second part focuses on the synthesis and biological properties of different σ_1 receptor ligands, identified as **1-9**. Monocyclic piperazines **1** and bicyclic piperazines **2** and **3** were developed as cytotoxic compounds, thus the IC_{50} values of cell growth and survival inhibition studies are given for all derivatives. The mechanism of cell survival inhibition, induction of time-dependent apoptosis, of compound **ent-2a** is discussed. Experimentally determined σ_1 affinity shows good correlation with the results from molecular dynamics simulations based on a 3D homology model. Spirocyclic compounds **4** and **5** represent well-established σ_1 receptor ligands. The homologous fluoroalkyl derivatives **4** have favorable pharmacological properties for use as fluorinated PET tracers. The (*S*)-configured fluoroethyl substituted compound (**S**)-**4b** is under investigation as PET tracer for imaging of σ_1 receptors in the brain of patients affected by major depression. 1,3-Dioxanes **6c** and **6d** display a very potent σ_1 antagonist profile and the racemic 1,3-dioxane **6c** has high anti-allodynic activity at low doses. The arylpropenylamines **7** are very potent σ_1 receptor ligands with high σ_1/σ_2 selectivity. The top compound **7g** acts as an agonist as defined by its ability to potentiate neurite outgrowth at low concentrations. Among the morpholinoethoxy-pyrazoles **8**, **8c** (known as S1RA) reveals the most promising pharmacokinetic and physicochemical properties. Due to its good safety profile, **8c** is currently being investigated in a phase II clinical trial for the treatment of neuropathic pain. The most potent ligand **9e** of 3,4-dihydro-2(1H)-quinolones **9** shows promising anti-nociceptive activity in the formalin test.

Keywords

Pharmacological data • Structure (σ_1) affinity relationships • Synthesis • σ_1 Receptor ligands • σ_1 Receptor pharmacophore models

1 Introduction

The sigma-1 (σ_1) receptor is a membrane-bound protein distributed in the central nervous system and in peripheral organs like heart, kidney, and liver (Weissman et al. 1988; Samoilova et al. 1988; Ela et al. 1994). The σ_1 receptor is mainly localized at the endoplasmic reticulum and the mitochondria-associated membranes. Ruoho et al. have shown that the σ_1 receptor consists of two transmembrane regions connected by a loop. Both C- and N-terminus are located extracellularly or in the ER lumen (Chu and Ruoho 2016). Two additional hydrophobic regions of the σ_1 receptor were identified by Fontanilla et al. named steroid binding domain-like regions (SBDL I and II). With the help of *N*-substituted photoaffinity labels it was shown that the SBDL I overlaps with one of the two transmembrane regions of the σ_1 receptor forming the ligand-binding domain together with the SBDL II (Ruoho et al. 2012). σ_1 Receptors were shown to take part in the regulation of ion channels (e.g., K^+ and Ca^{2+}) and in the modulation of neurotransmitter systems (Lupardus et al. 2000; Hong and Werling 2000; Hayashi

and Su 2007). In the brain, the σ_1 receptor is particularly well expressed in areas associated with memory and emotion (Mash and Zabetian 1992). Steroids like progesterone (Su et al. 1988; Schwarz et al. 1989) and *N,N*-dimethyltryptamine (Fontanilla et al. 2009) were previously discussed to be endogenous ligands but their σ_1 receptor binding affinities are low compared with those of sphingosines showing high affinity in the low-nanomolar range (Ruoho et al. 2012). Since many centrally active drugs show high σ_1 affinity, σ_1 receptors represent promising targets for the research and development of drugs to treat several neurological or neuropsychiatric disorders like depression, psychosis, and cocaine abuse (Hascoet et al. 1995; Matsumoto et al. 2001; Sharkey et al. 1988; Bermack and Debonnel 2001; Ishikawa et al. 2007; Skuza and Rogoz 2006). The fact that many human cancer cell lines show up-regulated levels of σ_1 receptors brought them into focus for the development of new antitumor drugs and cancer diagnostics (Hashimoto and Ishiwata 2006). Based on various studies, Chen and Pasternak postulated that the σ_1 receptor functions as an endogenous anti-opioid receptor system (Chien and Pasternak 1993). By investigation of different σ_1 receptor ligands in animal models it was shown that σ_1 agonists inhibit morphine-induced analgesia whereas σ_1 antagonists potentiate opioid induced analgesia (Chien and Pasternak 1994, 1995). The fact that σ_1 knockout mice show reduced pain response in the formalin test but not hypersensitivity after treatment with capsaicin lead to interest in σ_1 receptors as a target in the treatment of neuropathic pain (Entrena et al. 2009).

For the development of new potent σ_1 receptor ligands with high affinity several pharmacophore models have been developed and optimized. Herein the pharmacophore models reported so far are summarized and compared with respect to existing ligands.

2 Pharmacophore Models

In 1994, Glennon et al. reported a two-dimensional pharmacophore model based on deconstruction–reconstruction analysis of different flexible σ_1 receptor ligands. In this model, two hydrophobic regions flanking a basic amine represent the pharmacophoric elements required for high σ_1 affinity. A distance of 6–10 Å between the amine moiety and the primary hydrophobic region and of 2.5–3.9 Å between the amino group and the secondary hydrophobic region provides optimal binding conditions. The amine could be of primary, secondary, or tertiary nature. In case of a tertiary amine, only small substituents are allowed, whereas the amine could also be part of a ring system (e.g., piperazine ring). The primary hydrophobic region tolerates sterically demanding residues whereas the secondary region favors smaller substituents like a three-carbon chain. As the two hydrophobic binding pockets of the σ_1 receptor tolerate bulky groups, the size of substituents can vary slightly without decreasing binding affinity (Fig. 1) (Glennon et al. 1994; Glennon 2005).

Laggner et al. presented in 2005 the first computer-aided three-dimensional pharmacophore model (Fig. 2) based on 23 structurally very different σ_1 ligands.

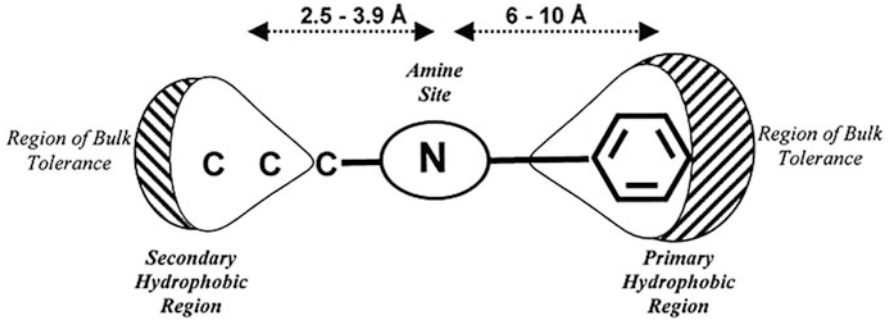


Fig. 1 Pharmacophore model of Glennon (2005)

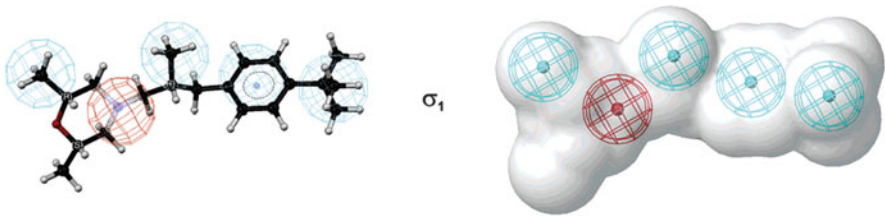


Fig. 2 3D- σ_1 -Pharmacophore model of Laggner et al. (2005) red: positive ionizable group; blue: hydrophobic regions

The pharmacophoric elements consist of a positive ionizable group like an amine and four hydrophobic features. The calculated distances between the pharmacophoric elements are in good agreement with the results obtained by Glennon et al. (1994).

In 2009, Zampieri et al. designed another computer-based model containing five pharmacophoric elements (Fig. 3) (Zampieri et al. 2009). The model included three hydrophobic areas in total (depicted in blue and pink), whereby two of them should have aromatic character (blue). A basic center (red) is located at a distance of 7.01 and 8.50 Å from the aromatic moieties and at a distance of 3.58 Å from the further hydrophobic elements. These distances are comparable to those postulated in the models of Glennon and Laggner. Additionally, the Zampieri model established an H-bond acceptor function, which was already defined in a pharmacophore model by Gilligan et al. This model was published in the early 1990s and did not differentiate between σ_1 and σ_2 ligands (Gilligan et al. 1992).

In 2011, Laurini et al. published the first computer-based 3-dimensional (3D) homology model of the σ_1 receptor. For the identification of a reliable ligand-binding domain, results of docking studies, mutagenesis studies, structure–affinity-relationship studies, and pharmacophore models were combined. The validation of the homology model was implemented by docking studies of well-known σ_1 ligands at the postulated binding site of the receptor, calculation of free binding

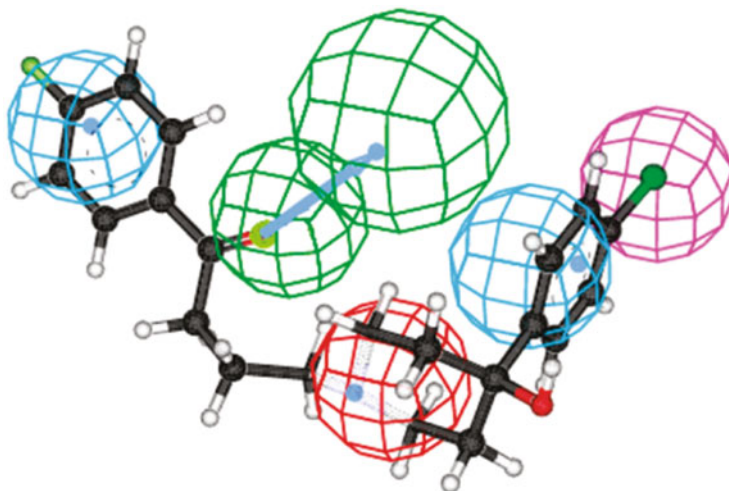


Fig. 3 Pharmacophore model of Zampieri et al. (2009) *red*: basic center; *green*: H-bond acceptor; *blue*: aromatic hydrophobic area; *pink*: hydrophobic area

energy, and comparison with the experimentally determined σ_1 affinities of these ligands (Laurini et al. 2011, 2012).

Schmidt et al. have just published a crystal structure of the σ_1 receptor for the first time (Schmidt et al. 2016). This structure was determined in complex with two different σ_1 receptor ligands, PD144418 and 4-IBP. Contrary to the early findings of Fontanilla and Ruoho (Ruoho et al. 2012) as well as Aydar et al. only one transmembrane domain of the σ_1 receptor was found in the crystal structure (Schmidt et al. 2016). This contradicts also the solution nuclear magnetic resonance (NMR) results of Ortega-Roldan et al. (2015) which closely match the findings of the 3D homology model of Laurini et al. (2011).

3 Ligands Introduction

In the literature, a great variety of σ_1 receptor ligand classes are reported. These classes include piperazines **1**, bicyclic compounds of type **2** and **3**, spirocyclic compounds **4** and **5**, 1,3-dioxanes **6**, arylalkenylamines **7**, morpholinoethoxypyrazoles **8**, and 3,4-dihydro-2(1*H*)-quinolones **9** (Fig. 4). The synthesis and the pharmacological properties of these ligands are presented herein.

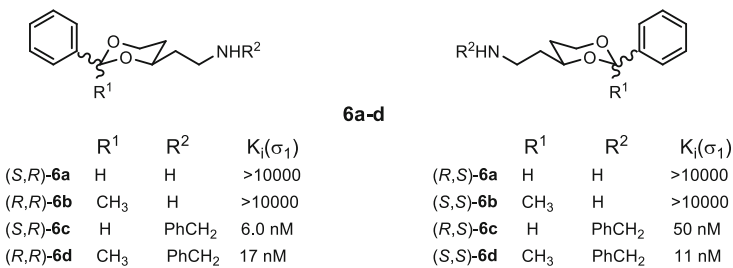
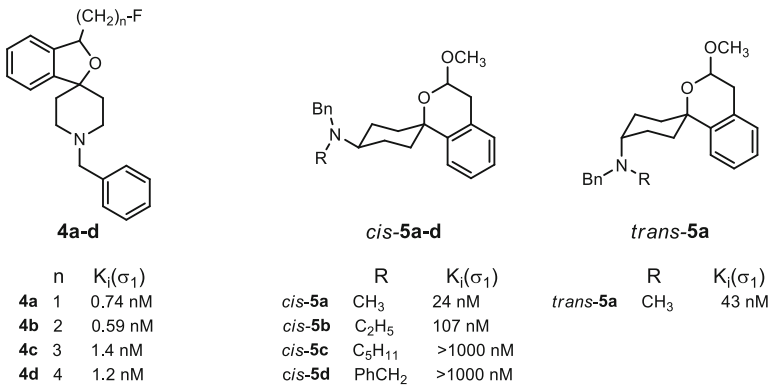
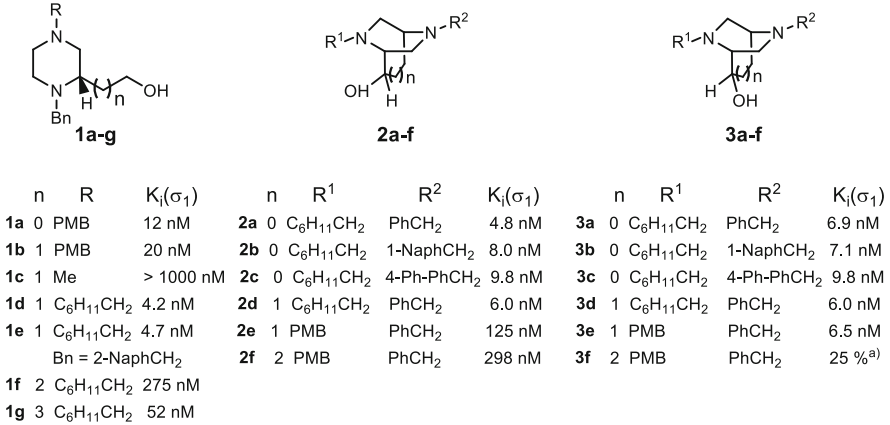
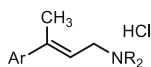
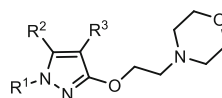


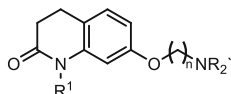
Fig. 4 Representative σ_1 receptor ligands of different compound classes. Inhibition of σ_1 receptor radioligand binding at 1 μ M concentration of test compound. *PMB* *p*-methoxybenzyl; *Naph* naphthyl

**7a-f-HCl**

	Ar	NR ₂	K _i (σ_1)
7a	4-Ph-PhCH ₂	piperidine	0.86 nM
7b	2-Naph	piperidine	0.97 nM
7c	4-Ph-PhCH ₂	4-benzylpiperidine	7.0 nM
7d	2-Naph	4-benzylpiperidine	23 nM
7e	4-Ph-PhCH ₂	morpholine	12 nM
7f	Ph	morpholine	>1000 nM

**8a-f**

	R ¹	R ²	R ³	K _i (σ_1)
8a	<i>tert</i> -butyl	H	H	>1000 nM
8b	4-chlorophenyl	H	H	18 nM
8c	2-Naph	CH ₃	H	17 nM
8d	2-Naph	CH ₃	CH ₃	139 nM
8e	3,4-dichlorophenyl	CH ₃	CH ₃	9.4 nM
8f	3,4-dichlorophenyl	CH ₃	C(C=O)CH ₃	741 nM

**9a-i**

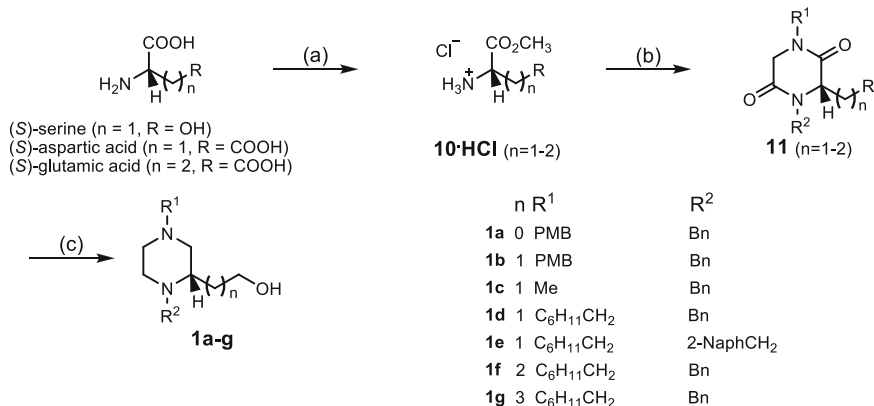
	R ¹	n	NR ₂	K _i (σ_1)
9a	PhCH ₂	2	morpholine	> 2000 nM
9b	PhCH ₂	3	morpholine	15 nM
9c	H	3	piperidine	89 nM
9d	CH ₃	3	piperidine	34 nM
9e	PhCH ₂	3	piperidine	1.84 nM
9f	4-F-Ph-CH ₂	3	piperidine	1.22 nM
9g	PhCH ₂	4	morpholine	60 nM
9h	PhCH ₂	5	morpholine	88 nM
9i	PhCH ₂	6	morpholine	682 nM

Fig. 4 (continued)

4 Homologous 2-Piperazinealkanols

Piperazines **1** with different hydroxyalkyl side chains represent well-established σ_1 receptor ligands. A broad structure affinity relationship study was performed based on 2-hydroxyethyl substituted piperazines with high σ_1 affinity and their larger and smaller homologs bearing hydroxypropyl or hydroxymethyl side chains. The cytotoxic activity against human cancer cell lines was tested by *in vitro* cell survival assays (Weber et al. 2014; Holl et al. 2012; Bedurftig and Wunsch 2004).

(*S*)-Serine, (*S*)-aspartic acid, and (*S*)-glutamic acid as enantiomerically pure amino acids of nature's chiral pool were used for the synthesis of homologous piperazinealkanols **1a-g**. The first reaction step includes the esterification of the particular amino acid. The dioxipiperazines **11** were prepared from the aminoesters **10**·HCl in a three-step reaction sequence consisting of reductive alkylation, chloroacetylation, and ring closure with different primary amines. Reduction with



Scheme 1 Synthesis of homologous 2-piperazinealkanol **1**. Reagents and reaction conditions: (a) $(\text{H}_3\text{C})_3\text{SiCl}$, CH_3OH , room temperature (rt), 16 h; (b) (1) $\text{Ph}-\text{CH}=\text{O}$, NEt_3 , CH_2Cl_2 , rt, 16 h; (2) NaBH_4 , CH_3OH , 0°C , 40 min; (3) ClCH_2COCl , NEt_3 , CH_2Cl_2 , rt, 2.5 h; (4) R^1-NH_2 , NEt_3 , CH_3CN , rt, 16 h–3 d; (c) LiAlH_4 , THF, reflux, 16 h. *PMB* *p*-methoxybenzyl, *2-NaphCH*₂ 2-naphthyl (Weber et al. 2014; Holl et al. 2012; Bedurftig and Wunsch 2004)

LiAlH_4 led to the piperazinealkanol **1a-g** (Scheme 1). Reduction of the ester moiety of **11d** ($n = 1$, $\text{R} = \text{CO}_2\text{CH}_3$) followed by Wittig reaction of the resulting aldehyde with methyl(triphenylphosphoranylidene)acetate and subsequent reduction of the α,β -unsaturated ester led to the homologous hydroxybutyl piperazine **1g** (Weber 2012).

The σ_1 and σ_2 affinity of hydroxyalkyl piperazines **1a-g** was tested with tissue membrane preparations of animal origin (guinea pig brain, rat liver). Selected ligands (**1d-g**) were also assayed with membrane preparations bearing human σ_1 receptors to evaluate ligand-binding affinity towards σ_1 receptors from different species (Table 1).

For a high σ_1 affinity, the length of hydroxyalkyl side chain and the size of the residues at both *N*-atoms are of particular importance. Short side chains like hydroxymethyl and hydroxyethyl are well-tolerated by the σ_1 receptor leading to K_i values in the range of 4–20 nM. The σ_1 affinity of the hydroxypropyl piperazines is more than tenfold lower (e.g., **1f**, $K_i = 275$ nM). The extension of the side chain by another methylene moiety in case of hydroxybutyl piperazine **1f** leads to an increased σ_1 affinity, but the K_i value of 52 nM remained higher than the K_i value measured for hydroxymethyl and hydroxyethyl derivatives. In accordance with the pharmacophore model of Glennon postulating two hydrophobic regions, the *N*-methyl substituted piperazine **1c** does not show high σ_1 receptor affinity. The affinity increases by the introduction of a larger residue such as cyclohexylmethyl or *p*-methoxybenzyl group.

The hydroxyethyl derivatives show almost the same σ_1 affinity as the hydroxymethyl derivatives, but show reduced σ_2 affinity than hydroxyethyl piperazines. Regarding the σ_1/σ_2 selectivity, it becomes clear that the hydroxyethyl

Table 1 Inhibition of cancer cell growth by piperazinealkanoles **1a-g** compared with σ_1 and σ_2 binding affinity

	σ_1 (gp) ^b K_i [nM] \pm SEM	σ_2 (rat) ^b K_i [nM] \pm SEM	σ_1 (hu) ^c K_i [nM] \pm SEM	σ_1/σ_2 selectivity (ratio)	$\sigma_1 K_i$ (calc) [nM]	IC ₅₀ [μ M] RT4 ^d	IC ₅₀ [μ M] 5637 ^e	IC ₅₀ [μ M] A-427 ^f	IC ₅₀ [μM] LCLC-103H ^g	IC ₅₀ [μM] DAN-G ^h	IC ₅₀ [μM] MCF-7 ⁱ	IC ₅₀ [μM] RPMI 8226 ^j
1a	12 \pm 1.4	70 \pm 10	–	6	–	–	–	–	–	–	–	–
1b	20 \pm 6.0	>1,000	–	50	–	–	–	–	–	–	–	–
1c	28% ^k	27% ^k	–	–	–	>20	>20	>20	>20	>20	>20	>20
1d	4.2 \pm 1.1	116 ^l	21 \pm 4	28	35	>20	>20	14 \pm 5	>20	>20	6.8 \pm 8.4	–
(ent)- 1d	1.9 \pm 0.4	60 \pm 11	23 \pm 8	32	–	–	>20	>20	>20	>20	–	>20
1e	4.7 \pm 1.8	69 \pm 24	6.8 \pm 2.3	15	11	>20	2.1 \pm 1.9	2.5 \pm 2.6	>20	>20	1.5 \pm 2	7.3 \pm 3.9
1f	275 ^l	690 ^l	71 ^l	3	–	>20	8.0 \pm 5.6	>20	>20	>20	>20	>20
1 g	52 ^l	348 ^l	54 \pm 26	7	–	>20	6.0 \pm 4.0	9.6 \pm 5.1	>20	>20	>20	>20

^aGuinea pig brain (gp)

^bRat liver (rat)

^cRPMI 8226 human cancer cell line (hu)

^dRT4 bladder cancer cell line

^eBladder cancer

^fSmall cell lung cancer

^gLarge cell lung cancer

^hPancreas cancer

ⁱBreast cancer

^jMultiple myeloma

^kInhibition of radioligand binding at 1 μ M concentration of test compound (% inhibition)

^lResult from one measurement. All other results are from three independent experiments

derivatives are more than tenfold selective for the σ_1 over the σ_2 receptor. The PMB-substituted compound **1b** provides the best selectivity in this set of compounds with a σ_2 affinity of $>1,000$ nM. As a result of further chain extension, the σ_2 affinity increases leading to decreased selectivity (**1f**, $K_i(\sigma_2) = 690$ nM, **1g**, $K_i(\sigma_2) = 348$ nM).

The σ_1 affinity of **1d-g** measured with membrane preparations from a human cancer cell line (RPMI 8226) is slightly reduced compared to the affinity measured with the guinea pig brain membrane preparations. Because the same trend was found for the reference compounds haloperidol and (+)-pentazocine it can be assumed that the results of both assays are well comparable.

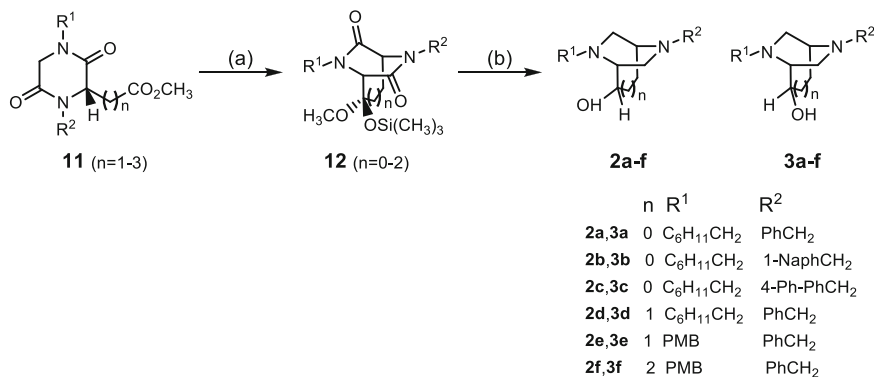
As the enantiomer of **1d** prepared in the same manner from (*R*)-aspartic acid shows the same σ_1 affinity, it could be assumed that the stereochemistry has only negligible influence on σ_1 receptor affinity and selectivity over the σ_2 subtype ($K_i(\sigma_1) = 1.9$ nM, $\sigma_1/\sigma_2 = 32$).

The σ_1 affinity obtained with human receptor preparations was supported by docking of the ligands in the putative binding site of the 3D σ_1 receptor homology model. The calculated free binding energies are in good accordance with their recorded affinities towards the σ_1 receptor. For the most potent human σ_1 receptor ligand **1g** ($K_{i,\text{exp.}} = 6.8$ nM) a ΔG_{bind} of -10.85 ± 0.36 kcal/mol was calculated which corresponds to an estimated $K_i(\sigma_1)_{\text{calcd}}$ value of 11.2 nM, consistent with the experimentally determined K_i values (Weber et al. 2014).

The cell growth inhibition potential of piperazinealkanol **1c-g** was tested in seven human tumor cell lines. The potent σ_1 receptor ligand **1e** inhibited the growth and survival of the bladder cancer cell line 5637, the small cell lung cancer cell line A427, and the multiple myeloma cell line RPMI 8226 in the low micromolar range. Even at high concentrations (20 μM) of **1e**, a growth inhibition activity could not be found for the bladder cancer cell line RT4, the large cell lung cancer LCLC-103H, and the pancreas cancer cell line DAN-G. Additionally only low activity was found for the breast cancer cell line MCF-7, indicating a selective mechanism of growth and survival inhibition. Further investigation of the mechanism associated with the inhibitory activity of **1e** was performed with RPMI 8226 cells and revealed an increase in the amount of early apoptotic cells after 48 h compared to the untreated control.

5 Bicyclic Piperazines

In order to investigate the influence of conformational restriction on σ_1 receptor affinity bicyclic compounds of type **2** and **3** with diazabicycloalkan scaffold were designed by intramolecular connection of the 2-hydroxyalkyl side chain of piperazines **1** with C-5 of the piperazine ring. Propano- and butano-bridged homologs of **2** and **3** with diazabicyclo[3.2.2]nonane and diazabicyclo[4.2.2]decane scaffold were synthesized by an expansion of the ethano-bridge by one or two



Scheme 2 Synthesis of bicyclic diazabicycloalkanols **2** and **3**. Reagents and reaction conditions: (a) $n = 1$: NaHMDS, THF, $-78^\circ C$, 40 min, then $(H_3C)_3SiCl$, $-78^\circ C$, 1 h, then rt, 2 h (Weber et al. 2016); $n = 2$: LiHMDS, THF, $-78^\circ C$, 30 min, then $(H_3C)_3SiCl$, $-78^\circ C$, 0.5 h, then rt, 3 h (Geiger et al. 2007); $n = 3$: LiHMDS, THF, $-78^\circ C$, 30 min, then $(H_3C)_3SiCl$, $-78^\circ C$, 2 h, then rt, 0.5 h (Sunnam 2010); (b) $n = 0$: 1. 0.5 M HCl, THF, rt, 16 h; 2. $LiAlH_4$, THF, reflux, 16 h (Weber et al. 2016); $n = 1$: 1. *p*-TosOH, THF, H_2O , rt, 16 h; 2. $LiBH_4$, THF, $-90^\circ C$, 3.5 h; 3. $LiAlH_4$, THF, reflux, **16 h** (Geiger et al. 2007) $n = 2$: 1. *p*-TosOH, THF, H_2O , rt, 16 h; 2. $LiBH_4$, THF, $-90^\circ C$, 2.5 h; 3. $LiAlH_4$, THF, reflux, 15 h (Sunnam 2010)

methylene moieties to elucidate the effect of bridge size on σ_1 receptor affinity (Geiger et al. 2007; Weber et al. 2016; Sunnam 2010).

Homologous dioxopiperazines **11** ($n = 1-3$) with different substitution pattern of the *N*-atoms were used as starting material for the synthesis of bicyclic compounds **2** and **3** (Scheme 2). The dioxopiperazine **11** ($n = 3$) was synthesized from racemic-2-aminoadipic acid in the same manner as explained for dioxopiperazines **11** ($n = 1,2$) in Scheme 1. The mixed methyl silyl ketals **12** were obtained by Dieckmann analogous cyclization of **11**. The Dieckmann analogous cyclization gave only low yields for the dioxopiperazines **11** ($n = 1$) with acetate side chain due to the rigidity of the resulting products **12** ($n = 0$). The (*R*)-configuration of the ketalic center of **12** was shown by X-ray crystal structure analysis (Holl et al. 2008). Hydrolysis and reduction of **12** led to the bicyclic alcohols **2** and **3**. The enantiomers *ent*-**2** and *ent*-**3** were obtained starting with (*R*)-configured amino acids.

In piperazinealkanols **1** the hydroxyalkyl side chain can adopt several conformations and, moreover, the piperazine ring can adopt two conformations, leading to an axial or equatorial orientation of the side chain resulting in different distances between the pharmacophoric elements. In the bicyclic alcohols **2** and **3** the additional bridge over the piperazine ring reduces the flexibility of the ring system and its hydroxyalkyl side chain. As a result of conformational restriction, the pharmacophoric elements are fixed in a defined arrangement minimizing the loss of entropy during binding and thus increasing the overall free binding energy.

The σ_1 and σ_2 receptor affinity of the bicyclic alcohols **2** and **3** was determined with tissue membrane preparations from guinea pig brain (for σ_1) and rat liver (for

σ_2). Compounds **2a-d** and **3a-d** were also tested against human σ_1 receptors from multiple myeloma RPMI 8226 cell line membrane preparations (Geiger et al. 2007; Weber et al. 2016).

Almost all cyclohexylmethyl substituted compounds **2a-d** and **3a-d** show high σ_1 receptor affinity with K_i values in the low-nanomolar range. The only exceptions are *ent*-**2a** and **3c**, both with a K_i value of 23 nM. The extension of the ethano-bridge of **2a-c** and **3a-c** by a methylene moiety does not influence σ_1 receptor affinity since the propane-bridged homologs **2d,e** and **3d,e** show approximately the same affinity. Only **2e** and *ent*-**2e** show K_i values in the three-digit nanomolar range ($K_i(\sigma_1) = 125$ and 118 nM). However, the introduction of a second methylene moiety leads to a salient decrease in σ_1 receptor affinity, which implies that butano-bridged diazabicycloalkanol **2f** and **3f** are not tolerated by the σ_1 receptor.

The stereochemistry has only low impact on σ_1 receptor affinity since all four stereoisomers **2a**, *ent*-**2a**, **3a**, and *ent*-**3a** show the same σ_1 receptor affinity. However, in case of PMB-substituted derivatives, **2e** and *ent*-**2e** show lower σ_1 receptor affinity than **3e** and *ent*-**3e**.

The σ_1 affinity determined with human σ_1 receptor material is in good accordance with the σ_1 affinity obtained with σ_1 receptors from guinea pig brain.

Compared with the flexible hydroxyethyl piperazines **1b**, **1d**, and **1g**, the corresponding ethano-bridged piperazines **2a-c** and **3a-c** reveal the same σ_1 receptor affinity. However, the conformational restriction of the hydroxypropyl piperazines **1d** led to increased σ_1 receptor affinity of **2e** and **3e**. That is due to the higher flexibility of the hydroxypropyl piperazines **1d,e** compared to their shorter hydroxyethyl homologs **1a-d**. This is not valid for the butano-bridged piperazines **2f** and **3f** which display very low σ_1 affinity, indicating that the bridge size is too bulky for the binding pocket. Obviously the size of the butano-bridge outweighed its positive effect of conformational restriction.

The σ_1/σ_2 selectivity varies from low preference for the σ_1 receptor (**2c**: $\sigma_1/\sigma_2 = 2$) up to high selectivity for the σ_1 receptor (**3d**: $\sigma_1/\sigma_2 = 178$). The PMB-substituted derivative *ent*-**3e** ($\sigma_1/\sigma_2 = 227$) showed the highest σ_1/σ_2 selectivity.

The bicyclic compounds **2a-d** and **3a-d** were docked into the binding site of the 3D homology model to determine the free binding energies. Figure 5 illustrates the identified interactions between the high affinity compound *ent*-**3a** ($K_i(\sigma_{1\text{human}}) = 1.6$ nM) and the human σ_1 receptor (Weber et al. 2016).

Fig. 5 Interactions between *ent*-**3a** and amino acids of the binding site in the 3D homology model of the human σ_1 receptor

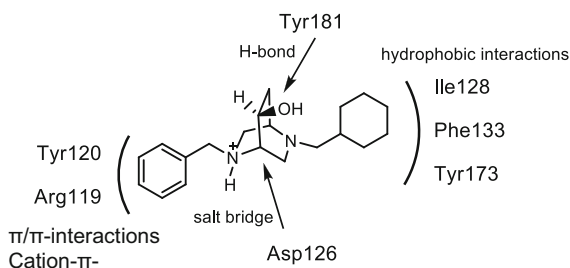


Table 2 Pharmacological data for diazabicycloalkanolis **2** and **3**

	σ_1 (gp) ^a K_i [nM] \pm SEM	σ_2 (rat) ^b K_i [nM] \pm SEM	σ_1 (hu) ^c K_i [nM] \pm SEM	σ_1/σ_2 selectivity	$\sigma_1 K_i$ (calc) [nM]	IC ₅₀ [μ M] RT4 ^d	IC ₅₀ [μ M] 5637 ^e	IC ₅₀ [μ M] A-427 ^f	IC ₅₀ [μ M] LCLC-103H ^g	IC ₅₀ [μ M] MCF-7 ^h
2a	4.8 \pm 0.7	36 \pm 9.0	3.2 \pm 0.4	7	19	>20	>20	16.5 \pm 6.2	>20	>20
3a	6.9 \pm 1.6	60 \pm 26	2.4 \pm 0.2	9	22	>20	9.2 \pm 6.3	9.8 \pm 4.3	>20	>20
<i>ent</i> -2a	23 \pm 13	197 \pm 18	2.8 \pm 1.0	9	12	>20	4.8 \pm 3.1	2.8 \pm 1.7	>20	>20
<i>ent</i> -3a	5.7 \pm 2.6	501 \pm 21	1.6 \pm 0.4	88	9.7	>20	>20	11.2 \pm 4.8	>20	>20
2b	8.0 \pm 2.0	51 \pm 16	13 \pm 5.0	6	6.3	12.9 \pm 7.3	4.3 \pm 1.8	2.3 \pm 0.9	10.4 \pm 0.9	7.0 \pm 4.1
3b	7.1 \pm 1.8	157 \pm 21	7.2 \pm 3.9	22	8.5	11.3 \pm 5.9	4.9 \pm 1.2	6.0 \pm 3.8	8.8 \pm 2.0	6.8 \pm 1.5
2c	8.7 \pm 1.2	20 \pm 7.0	27 \pm 9.0	2	15	–	4.9 \pm 4.1	1.6 \pm 1.2	3.2	–
3c	23 \pm 6.0	334 \pm 18	73 \pm 6.0	15	18	–	>10	4.5 \pm 5.7	>10	–
2d	6.0 \pm 0.2	65 \pm 7.0	6.4 \pm 0.9	11	20	>20	>20	7.6 \pm 4.7	>20	14 \pm 2.8
3d	1.6 \pm 0.1	284 \pm 72	2.2 \pm 1.1	178	17	>20	>20	10.3 \pm 2.9	>20	16 \pm 2.8
2e	125 \pm 18	705 ⁱ	–	6	–	95 \pm 14 ^j	103 \pm 7.7 ^j	46 \pm 11 ^j	96 \pm 11 ^j	96 \pm 14 ^j
3e	6.5 \pm 0.7	806 ⁱ	–	124	–	93 \pm 10 ^j	88 \pm 10 ^j	46 \pm 12 ^j	94 \pm 6.9 ^j	91 \pm 9.4 ^j
<i>ent</i> -2e	118 \pm 5.0	441 ⁱ	–	4	–	67 \pm 4.2 ^j	88 \pm 4.5 ^j	33 \pm 7.8 ^j	77 \pm 18 ^j	73 \pm 9.4 ^j
<i>ent</i> -3e	7.5 \pm 2.1	1,700 ⁱ	–	227	–	72 \pm 1.2 ^j	72 \pm 3.3 ^j	46 \pm 8.6 ^j	80 \pm 3.3 ^j	76 \pm 1.9 ^j
2f	298 \pm 28	4,800 ⁱ	–	162	–	–	–	–	–	–
3f	25% ^k	640 ^j	–	–	–	–	–	–	–	–

^aGuinea pig brain (gp)

^bRat liver (rat)

^cRPMI 8226 cells (hu)

^dBladder cancer

^eBladder cancer

^fSmall cell lung cancer

^gLarge cell lung cancer

^hBreast cancer

ⁱResult from one measurement. All other results are from three independent experiments

^jRelative cell growth (%) in relation to an untreated control

^kInhibition of radioligand binding at 1 μ M concentration of the test compound

The calculated free binding energies of all docked compounds are in good accordance with the experimentally determined receptor binding data. For *ent-3a* the calculated ΔG_{bind} is -10.93 ± 0.34 kcal/mol corresponding to a calculated K_i value of 9.7 nM. This K_i value is in good agreement with the K_i values recorded with σ_1 receptors from guinea pig brain ($K_i = 5.7$ nM) and from human RPMI 8226 cell ($K_i = 1.6$ nM) membrane preparations.

The cell growth inhibition potential of compounds **2a-e** and **3a-e** was evaluated in five different cancer cell lines (Table 2). The naphthylmethyl substituted derivatives **2b** and **3b** similarly inhibited the growth and survival of all tested cell lines. The benzyl substituted derivatives **3a** and *ent-2a* and the biphenylmethyl substituted compound **2c** show moderate inhibition of cell growth of the bladder cell line 5637. A clear correlation between σ_1 receptor affinity and growth and survival inhibition could not be determined, however, we did discern a trend revealing sensitivity of A-427 cell line against all tested bicyclic compounds. With the exception of **2b** and **3b**, growth and survival of the other cell lines were not inhibited at compound concentrations up to 10 or 20 μM .

The bridge size does not show additional influence on σ_1 receptor affinity. The K_i values of **2d** and **3d** are in the same range as the K_i values of the ethano-bridged compounds **2a-c** and **3a-c**. Although IC_{50} values are not available for compounds **2e** and **3e**, cell growth of only 33–46 % could be detected for the A-427 cell line, whereas the growth of the other cell lines was not inhibited (Geiger et al. 2007).

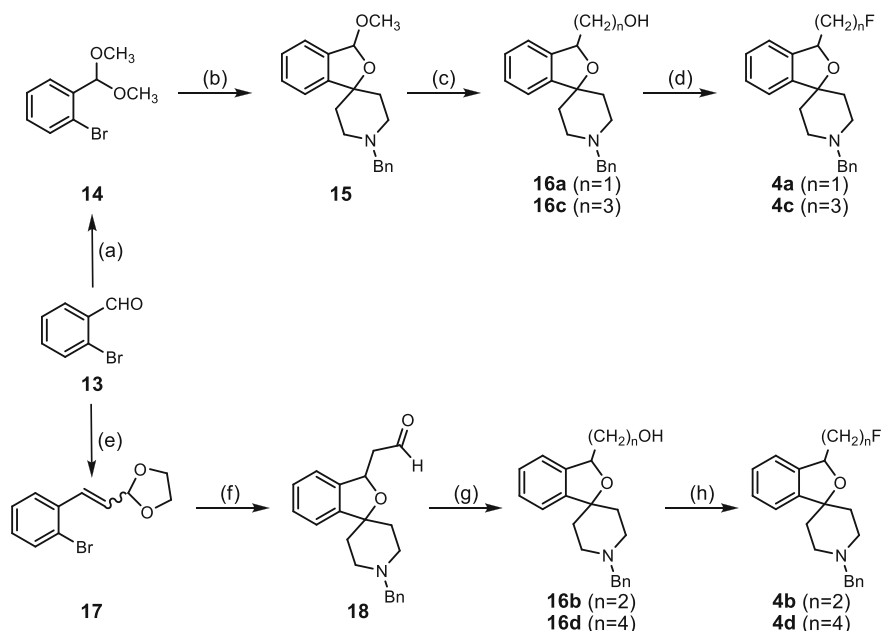
Further experiments directed to elucidate the mechanism of cell growth inhibition showed that *ent-2a* induced apoptosis in A-427 cells in a time-dependent manner (Weber et al. 2016).

6 Spirocyclic σ_1 Receptor Ligands

Spirocyclic compounds **4** represent high affinity σ_1 receptor ligands with a favorable pharmacological profile for use as fluorinated PET tracers. The homologous fluoroalkyl derivatives **4a-d** bind σ_1 receptor with K_i values in the low-nanomolar range and show high selectivity over the σ_2 receptor. The (*S*)-configured fluoroethyl substituted compound (*S*)-**4b** is currently investigated as PET tracer for imaging of σ_1 receptors in the CNS of patients suffering from major depression (Fischer et al. 2011; Wang et al. 2013; James et al. 2012). The spirocyclic σ_1 receptor ligands **5** bearing an exocyclic amino moiety allow diverse modifications by the introduction of two *N*-substituents. Furthermore, the existence of *cis/trans* isomerism increases the diversity of this compound class (Rack et al. 2011).

6.1 Homologous Fluoroalkyl Derivatives

The homologous fluoroalkyl derivatives **4a-d** were developed from the 2-benzofuran **15**, a ligand with high σ_1 receptor affinity ($K_i = 1.1$ nM) and high selectivity over the σ_2 receptor ($K_i = 1,280$ nM) and over 60 other receptors and ion



Scheme 3 Synthesis of homologous fluoroalkyl derivatives **4**. Reagents and reaction conditions (a) $\text{HC}(\text{OCH}_3)_3$, *p*-TosOH, CH_3OH , reflux, 16 h; (b) (1) *n*-BuLi, 1-benzylpiperidin-4-one, THF, -95°C , 2 h, rt, 4 h; (2) TosOH, CH_3OH , rt, 7 d; (Maier and Wunsch 2002a) (c) $n = 1$: (1) trimethylsilyl cyanide, tetracyanoethylene, CH_3CN , reflux, 4 h; (2) H_2SO_4 , EtOH, reflux, 7.5 h; (3) LiAlH_4 , Et_2O , -15°C , 30 min; (Maier and Wunsch 2002a) $n = 3$: (1) allyltrimethylsilane, $\text{BF}_3\cdot\text{OEt}_2$, CH_2Cl_2 , -25°C , 20 min, 0°C , 4 h; (2) 9-BBN, THF, rt, 16 h; (3) H_2O_2 , NaOH, -25°C , 45 min, rt 1 h; (Mastrup et al. 2009) (d) DAST, CH_2Cl_2 , -78°C to rt., 17 h; (Mastrup et al. 2009) (e) $[(\text{CH}_2\text{O})_2\text{CHCH}_2\text{PPh}_3\text{Br}]$, K_2CO_3 , TDA-1, CH_2Cl_2 , reflux, 6 d; (f) (1) *n*-BuLi, THF, 1-benzylpiperidin-4-one, -78°C , 1 h, rt, 16 h; (2) HCl, THF, rt, 2 h; (g) $n = 2$: NaBH_4 , CH_3CN , 0°C , 15 min, rt, 16 h; (Mastrup et al. 2011) $n = 4$: (1) ethoxycarbonylmethyltriphenylphosphorane, K_2CO_3 , THF, reflux, 23 h; (2) H_2 , Pd/C, EtOH, 1 bar, rt, 15 min; (3) LiAlH_4 , THF, -20°C , 30 min; (Grosse Mastrup 2010) (h) DAST, CH_2Cl_2 , -78°C to rt, 18–21 h (Mastrup et al. 2011)

channels like the hERG K^+ -channel. To eliminate the metabolically unstable acetalic function and to open up the possibility to introduce a fluorine atom into the molecule, a fluoroalkyl residue was installed instead of the acetalic moiety.

The synthesis of fluoroalkyl derivatives **4a–d** started with the acetalization of 2-bromobenzaldehyde **13** to yield the dimethyl acetal **14** (Scheme 3). Homologation of **13** with a Wittig reagent provided the α,β -unsaturated acetal **17**. Halogen-metal-exchange of the acetals **14** and **17** with *n*-BuLi followed by addition of 1-benzylpiperidin-4-one and subsequent transacetalization afforded the spirocyclic 2-benzofurans **15** and **18**. The 2-benzofurans **15** and **18** served as key intermediates for the synthesis of alcohols **16**. The alcohols were reacted with diethylaminosulfur trifluoride (DAST) to provide the homologous fluoroalkyl

Table 3 σ_1 and σ_2 binding affinities of homologous fluoroalkyl derivatives **4**

	<i>n</i>	σ_1 (gp) ^a K_i [nM] \pm SEM	σ_2 (rat) ^b K_i [nM]	σ_1/σ_2 selectivity
4a	1	0.74 \pm 0.34	550 ^c	743
4b	2	0.59 \pm 0.20	785 ^c	1,331
4c	3	1.4 \pm 0.26	837 ^c	620
4d	4	1.2 \pm 0.46	489 ^c	422
(<i>S</i>)-4b	2	2.3 \pm 0.2	897 ^c	390
(<i>R</i>)-4b	2	0.57 \pm 0.06	1,650 ^c	2,895

^aGuinea pig brain (gp)^bRat liver (rat)^cResult from one measurement. All other results are from three independent experiments, and data presented as mean $K_i \pm$ SEM (standard error of the mean)

derivatives **4** (Maestrup et al. 2009, 2011; Grosse Maestrup 2010; Maier and Wunsch 2002a, b).

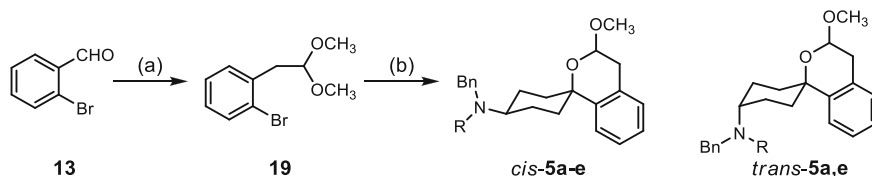
The σ_1 and σ_2 receptor affinity of the homologous fluoroalkyl derivatives **4** was determined with receptor material from guinea pig brain (σ_1) and rat liver (σ_2).

All four fluoroalkyl homologs **4a-d** bind with very high affinity to the σ_1 receptor ($K_i(\sigma_1) = 0.59\text{--}1.4$ nM) with high selectivity over the σ_2 subtype (Table 3). The fluoroethyl derivative **4b**, termed fluspidine, shows the most promising ligand-binding profile [$K_i(\sigma_1) = 0.59$ nM, $K_i(\sigma_2) = 785$ nM].

All four compounds **4a-d** were also synthesized in their [¹⁸F]-labeled form for the use as PET tracers. For radiosynthesis, the alcohols **14** were transformed into the corresponding tosylates. Nucleophilic substitution of the tosylates with K[¹⁸F]F complexed with the cryptand Kryptofix 2.2.2 led to the [¹⁸F] labeled spirocyclic σ_1 receptor ligands [¹⁸F]**4a-d** with high radiochemical purity (>98%) and radiochemical yield (40–50%) with reaction times <30 min (Fischer et al. 2011; Maestrup et al. 2009; Maisonial et al. 2011, 2012). Because of the high target affinity and selectivity, [¹⁸F]**4b** was further evaluated in animal studies with female CD-1 mice. [¹⁸F]**4b** showed fast and sufficient brain uptake (3.9 and 4.7%ID/g) and high metabolic stability in vivo (>94% parent compound in plasma samples after 30 min, only one metabolite was found). Good concordance between expression of σ_1 receptors and binding site occupancy with [¹⁸F]**4b** was found by ex vivo brain section imaging (Fischer et al. 2011). Due to the promising properties of the racemic compound [¹⁸F]**4b**, the enantiomers (*R*)- and (*S*)-[¹⁸F]**4b** were separated by chiral HPLC of the tosylate **13b** (Holl et al. 2013). The σ_1 receptor affinity was 0.57 nM for (*R*)-[¹⁸F]**4b** and 2.3 nM for (*S*)-[¹⁸F]**4b**. Thus, the (*R*)-enantiomer is the eutomer.

6.2 Spirocyclic Ligands with Exocyclic Amino Moiety

For the synthesis of spirocyclic ligands with exocyclic amino moiety, the dimethyl acetal **19** was reacted in a bromine lithium exchange to give an aryllithium



Scheme 4 Synthesis of spirocyclic ligands **5** with exocyclic amino moiety. Reagents and reaction conditions (a) (1) (methoxymethyl)triphenylphosphonium chloride, KO^tBu, THF, $-10\text{ }^{\circ}\text{C}$, then rt, 16 h; (2) *p*TosOH·H₂O, MeOH, reflux, 72 h; (b) (1) *n*-BuLi, THF, $-78\text{ }^{\circ}\text{C}$, 20 min; (2) cyclohexane-1,4-dione, $-78\text{ }^{\circ}\text{C}$, 2 h, rt, 1 h; (3) CHCl₃, HCl, rt, 1.5 h; (4) benzylamine, THF, HOAc, NaBH(OAc)₃, rt, 2 h; (5) R-CHO, NaBH(OAc)₃, CH₂Cl₂, rt, 23 h (Rack et al. 2011)

Table 4 σ_1 and σ_2 binding affinity of spirocyclic ligands **5** with exocyclic amino moiety

	R	σ_1 (gp) ^a K_i [nM] \pm SEM	σ_2 (rat) ^b K_i [nM] \pm SEM	σ_1/σ_2 selectivity
<i>cis</i> -5a	CH ₃	24 \pm 4.7	329 ^c	14
<i>trans</i> -5a	CH ₃	43 \pm 18	>1,000	23
<i>cis</i> -5b	C ₂ H ₅	107 \pm 25	666 \pm 106	6
<i>cis</i> -5c	C ₅ H ₁₁	>1,000	719 ^c	–
<i>cis</i> -5d	PhCH ₂	>1,000	>1,000	–

^aGuinea pig brain (gp)

^bRat liver (rat)

^cResult from one measurement. All other results are from three independent experiments, and data presented as mean $K_i \pm$ SEM (standard error of the mean)

intermediate. After addition to cyclohexane-1,4-dione followed by transacetalization under acidic conditions, reductive amination with benzylamine and NaBH(OAc)₃ led to the diastereomeric benzylamines *cis*-**5e** and *trans*-**5e** (R = H, Scheme 4). In order to investigate the influence of the second *N*-substituent, the benzylamines **5e** were transformed into different tertiary amines. Each isomer can adopt different conformations with axially or equatorially oriented amino substituents.

The σ_1 and σ_2 receptor affinity of spirocyclic ligands with exocyclic amino moiety **5** was determined with membrane preparations obtained from guinea pig brain (for σ_1) and rat liver (for σ_2) (Table 4).

The shift of the basic amino group to a position outside of the spirocyclic ring was envisaged to come closer to the required distances between the pharmacophoric elements (benzene ring and amino moiety) according to the models of Glennon and Laggner. The benzylpiperidin **15** (Fig. 6) shows high σ_1 receptor affinity and high selectivity over the σ_2 subtype and over other receptors and ion channels. It was found that small residues at the *N*-atom resulted in low σ_1 affinity whereas a benzyl group turned out to be optimal. The important role of the *N*-benzyl moiety can be explained by the pharmacophore model of Glennon et al. The benzene ring of the annulated pyrane ring interacts with the primary

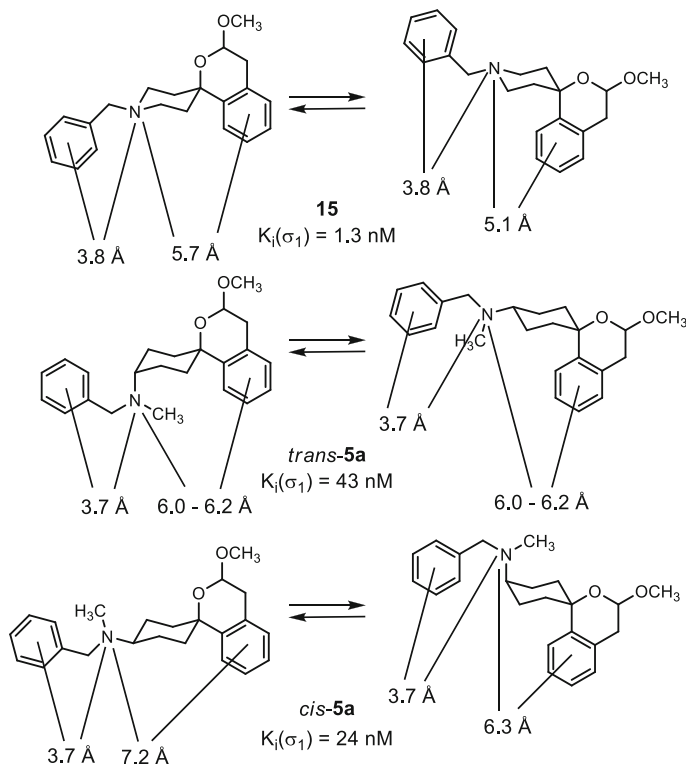


Fig. 6 Distance calculation of spirocyclic σ_1 receptor ligands with endocyclic *N* (**15**) and exocyclic amino moiety (*trans-5a* and *cis-5a*)

hydrophobic region and the benzyl moiety of the *N*-atom interacts with the secondary hydrophobic region. However, the distance between the *N*-atom and the primary hydrophobic region was too small for both conformers with axially and equatorially oriented phenyl ring. Ideally, the distance should be 6–10 Å due to the pharmacophore model of Glennon et al. In case of **15** the distance was found to be 5.7 and 5.1 Å for the equatorial and axial conformer, respectively (Fig. 6). Therefore it was decided to extend the distance between the *N*-atom and the *O*-heterocycle-annulated benzene ring by exclusion of the *N*-atom from the piperidine ring, resulting in spirocyclic compounds **5** with exocyclic amino group. Another advantage of an exocyclic amino moiety is the possibility to install and to modify two different residues at the *N*-atom.

As a result of the shift of the basic group, the distances between the *N*-atom and the *O*-heterocycle-annulated benzene ring of *cis-5a* and *trans-5a* are in good concordance with the distances postulated by Glennon et al. However, the decrease in σ_1 affinity ($K_i(\sigma_1) = 24$ and 43 nM) (Rack et al. 2011) provides an example of how receptor binding affinity does not strictly correspond with pharmacophore

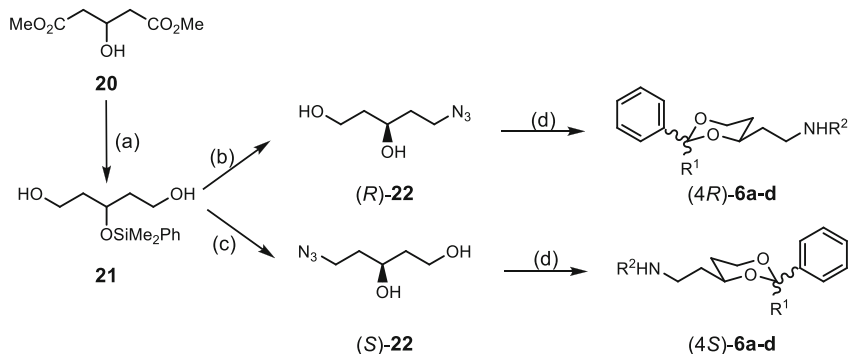
models. Other considerations like entropic factors should be noted. The introduction of the benzylamino moiety leads to an increased flexibility of the *N*-substituent.

The σ_1 receptor affinity of **5** also depends on the second *N*-substituent. Only small groups are tolerated. For small methyl and ethyl groups, the K_i values are 24 and 107 nM, respectively. Bulky residues like pentyl- or benzyl substituents lead to a salient decrease in σ_1 binding affinity ($K_i(\sigma_1) > 1,000$ nM). Generally, *cis*-configured diastereomers show higher σ_1 binding affinity than their *trans*-configured diastereomers.

7 1,3-Dioxanes

Racemic 1,3-dioxane **6c** represents a very potent σ_1 receptor antagonist (Utech et al. 2011). With these compounds σ_1 binding affinity depends on the relative configuration of the substituents at the 2- and 4-position, size of the oxygen containing heterocycle, and length of the aminoalkyl side chain. Since the racemic compound **6c**, consisting of a six-membered *O*-heterocycle combined with an aminoethyl side chain, was found to be a promising candidate as σ_1 receptor ligand, the enantiomers were synthesized and their pharmacology evaluated.

For the synthesis of enantiomerically pure 1,3-dioxanes **6**, the enantiomeric azidodiols (*S*)-**22** and (*R*)-**22** were prepared from diester **20** in high enantiomeric excess (Scheme 5). After silylation of **20** and subsequent reduction, the resulting diol **21** was converted into the azidodiols (*S*)-**22** and (*R*)-**22** following two different pathways using lipases as chiral catalysts. Stereoselective acetalization of (*S*)-**22** and (*R*)-**22** with benzaldehyde or acetophenone led to enantiomerically pure azido-



Scheme 5 Synthesis of enantiomerically pure 1,3-dioxanes **6**. Reagents and reaction conditions. (a) (1) Me_2PhSiCl , imidazole, CH_2Cl_2 ; (2) LiBH_4 , Et_2O ; (b) (1) IPA, lipase *Candida Antarctica* B, MTBE; (2) lipase *Burkholderia cepacia*, NaHCO_3 ; (3) $\text{Zn}(\text{N}_3)_2$ (pyridine) $_2$, DIAD, PPh_3 , toluene; (4) K_2CO_3 , CH_3OH ; (5) HCl ; (c) (1) IPA, lipase *Burkholderia cepacia*, MTBE; (2) $\text{Zn}(\text{N}_3)_2$ (pyridine) $_2$, DIAD, PPh_3 , toluene; (3) K_2CO_3 , CH_3OH ; (4) HCl ; (Kohler and Wunsch 2006, 2012) (d) (1) Ph-C(=O)R^1 , *p*TosOH, toluene, Dean Stark apparatus, 4 h; (2) H_2 , Pd/C, rt, 5 h; $\text{R}^2 = \text{PhCH}_2$; (3) benzaldehyde, $\text{NaBH}(\text{OAc})_3$, CH_2Cl_2 , rt, 16 h (Kohler et al. 2012). IPA isopropenyl acetate, MTBE methyl *tert*-butyl ether, DIAD diisopropyl azodicarboxylate

Table 5 σ_1 and NMDA receptor binding affinities of 1,3-dioxanes **6a-d**

	R ¹	R ²	σ_1 (gp) ^a K_i [nM] \pm SEM	NMDA (p) ^b K_i [nM] \pm SEM
(<i>S,R</i>)- 6a	H	H	>10,000	>10,000
(<i>R,S</i>)- 6a	H	H	>10,000	>10,000
(<i>R,R</i>)- 6b	CH ₃	H	>10,000	46 \pm 17
(<i>S,S</i>)- 6b	CH ₃	H	>10,000	6,120 \pm 630
(<i>S,R</i>)- 6c	H	PhCH ₂	6.0 \pm 1.0	>10,000
(<i>R,S</i>)- 6c	H	PhCH ₂	50 \pm 19	>10,000
(<i>R,R</i>)- 6d	CH ₃	PhCH ₂	17 \pm 2	>10,000
(<i>S,S</i>)- 6d	CH ₃	PhCH ₂	11 \pm 3	>10,000

^aGuinea pig brain (gp)^bPig brain cortex (p)

1,3-dioxanes, which were subsequently reduced with H₂ and Pd/C to obtain the primary amines **6a** and **6b**. Further functionalization of the amino moiety was performed by reductive monobenylation with benzaldehyde and NaB(OAc)₃ to yield the benzylamines **6c** and **6d**.

The σ_1 and σ_2 receptor affinity of 1,3-dioxanes **6** was determined with tissue membrane preparations from guinea pig brain (for σ_1) and rat liver (for σ_2). The 1,3-dioxanes **6** were also tested against the PCP binding site of the NMDA receptor using pig brain cortex membrane preparations (Table 5).

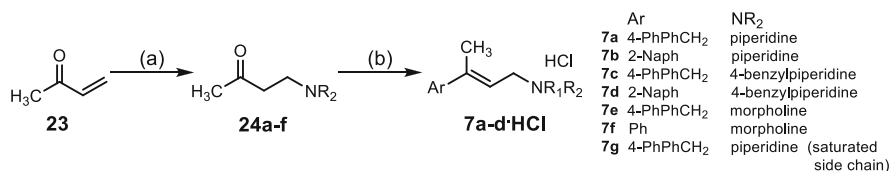
We found that both enantiomers of the primary amines **6a** and **6b** do not bind σ_1 ($K_i(\sigma_1) > 10,000$ nM) in this assay. According to the pharmacophore model of Glennon et al. (Glennon 2005; Glennon et al. 1994) affinity for σ_1 should increase by introducing an *N*-benzyl group as a second hydrophobic residue flanking the basic amino moiety as it is shown for the secondary amines **6c** and **6d**. In the case of benzylamines the orientation of the phenyl ring at the 1,3-dioxane ring has minimal influence on σ_1 affinity as **6a** and **6b** show comparable K_i values of 6.0 and 17 nM for the (4*R*)-configured enantiomers and 50 and 11 nM for the (4*S*)-configured enantiomers. Compound (*S,R*)-**6c** with equatorially oriented 2-phenyl moiety shows high σ_1 affinity ($K_i(\sigma_1) = 6.0$ nM).

Regarding σ_1/σ_2 selectivity, we found that primary amines **6a** and **6b** do not bind σ_1 and the benzyl amines **6c** and **6d** bind σ_2 with low affinity ($K_i(\sigma_2) > 200$ nM).

Depending on the absolute configuration the primary amines **6b** with axially oriented phenyl moiety reveal high affinity to the PCP binding site of the NMDA receptor with K_i values of 46 nM ((*R,R*)-**6b**) but only 6,120 nM for (*S,S*)-**6b**, respectively. The equatorial orientation of the phenyl ring (**6a**) as well as the introduction of a benzyl group at the amino moiety (**6c** and **6d**) led to complete loss of NMDA affinity ($K_i(\text{NMDA}) > 10,000$ nM).

The benzyl substituted 1,3-dioxane (*S,R*)-**6c** represents the most potent candidate among the secondary amines with high σ_1 affinity ($K_i(\sigma_1) = 6.0$ nM) and high selectivity over the σ_2 subtype and the NMDA receptor.

In further studies performed with racemic **6c** ($K_i(\sigma_1) = 19$ nM), promising results were obtained in a capsaicin-induced pain assay with mice. In these studies,



Scheme 6 Synthesis of arylalkenylamines **7a-f**. Reagents and reaction conditions (a) HNR₂, PEG 400, rt; (b) (1) Ar-Br, t-BuLi, Et₂O, -78 °C to rt; (2) 37% HCl, rt; (3) 1 M NaOH; (4) 37% HCl, rt; (5) crystallization from acetone (Rossi et al. 2011). *Naph* naphthyl

Table 6 σ_1 and σ_2 affinity of arylalkenylamines **7a-g**

	σ_1 (gp) ^a K_i [nM] \pm SEM	σ_2 (rat) ^b K_i [nM] \pm SEM	σ_1/σ_2 selectivity	clogP	clogD	MW
7a	0.86 \pm 0.4	111 \pm 21	129	5.32	3.82	291.43
7b	0.97 \pm 0.3	35 \pm 9	36	4.66	3.07	265.39
7c	7.0 \pm 0.9	18 \pm 1.7	3	7.19	5.60	381.55
7d	23 \pm 2.6	16 \pm 1.1	1	6.53	4.86	355.52
7e	12 \pm 2.0	386 \pm 27	33	4.25	4.17	293.40
7f	>1,000	>1,000	–	2.61	2.51	217.31
7g	0.70 \pm 0.3	103 \pm 10	147	5.43	3.00	293.45

^aGuinea pig brain

^bRat liver

even a very low dose of 0.25 mg/kg, *rac*-**6c** has high anti-allodynic activity (Utech et al. 2011).

8 Arylalkenylamines

Arylpropenylamines of type **7** show high σ_1 binding affinity and high σ_1/σ_2 selectivity. The influence of the novel σ_1 ligands on nerve growth factor (NGF)-induced neurite outgrowth was evaluated in the in vitro PC12 cell neurite sprouting assay.

Michael addition of cyclic amines to unsaturated ketone **23** led to the β -aminoketones **24**. Subsequent nucleophilic aryllithium addition followed by dehydration with HCl provided the arylalkenylamines **7**, which were crystallized as HCl salts (*E*)-**7a-f** (Scheme 6). The racemic arylalkylamine **7g** was obtained by catalytic hydrogenation of **7a**.

The σ_1 and σ_2 receptor binding affinity of compounds **7a-g**·HCl was tested using guinea pig brain (for σ_1) and rat liver (for σ_2) membrane preparations. Additionally, the selectivity towards the PCP binding site of the NMDA receptor and against μ - and κ -opioid receptors was determined (Table 6).

Piperidinyl substituted compounds **7a** and **7b** reveal high σ_1 receptor affinity independent of the aromatic residue ($K_i(\sigma_1) = 0.86$ and 0.97 nM). Interestingly, naphthalen-2-yl- or biphenyl-4-yl residues appear to be important for high σ_1

binding affinity of morpholinyl substituted compounds since only low σ_1 affinity is found when a phenyl substituent is present as aromatic moiety (**7f**, $K_1(\sigma_1) > 1,000$ nM). The tested compounds show high selectivity over the σ_2 receptor subtype, opioid receptors, and NMDA receptors (the PCP binding site). High σ_2 binding affinity was found only for 4-benzylpiperidinyl substituted derivatives **7c** and **7d**, with K_1 values of 18 and 16 nM, respectively. **7a** represents the most potent and selective σ_1 receptor ligand of this set of compounds ($K_1(\sigma_1) = 0.86$ nM, $\sigma_1/\sigma_2 = 129$). Therefore the corresponding arylalkylamine **7g** was included in this study. Receptor binding studies revealed similar σ_1 binding affinity ($K_1(\sigma_1) = 0.70$ nM) and selectivity ($\sigma_1/\sigma_2 = 147$) (Rossi et al. 2011).

$\text{clog}P$ and $\text{clog}D$ values were calculated for **7a-g**. Their drug-like properties were confirmed according to Lipinski's "rule of five." With the exception of **7c** ($\text{log}D > 5$) all compounds fulfill the "rule of five," i.e., $\text{clog}D < 5$, molecular weight < 500 , H-bond acceptors < 10 , and H-bond donors < 5 .

In order to determine whether the top compounds **7a** and **7g** function as agonists or antagonists their influence on nerve growth factor (NGF)-induced neurite outgrowth in PC12 cells was evaluated. **7g** potentiated the neurite outgrowth at lower concentrations, consistent with agonist activity. This effect was blocked by co-treatment with the σ_1 receptor antagonist BD-1063, demonstrating the participation of σ_1 receptors. In contrast, (*E*)-**7a** did not significantly increase neurite sprouting (Rossi et al. 2011).

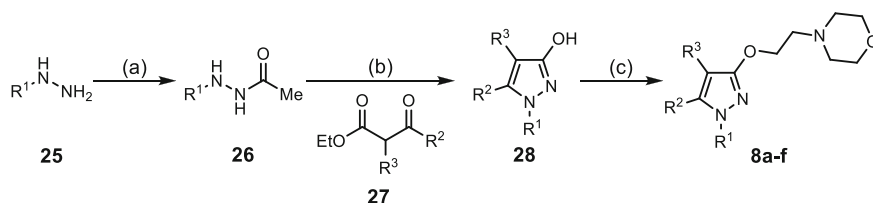
9 Morpholinoethoxypyrazoles

Substituted 1-arylpyrazoles with a basic amino function represent a promising class of σ_1 receptor antagonists. For high σ_1 binding affinity, the distance between the basic amino moiety and the pyrazole ring is of major importance. In previous studies an ethylenoxy spacer and a morpholino residue as the *N*-component resulted in high σ_1 affinity and excellent selectivity over the σ_2 subtype.

For the synthesis of morpholinoethoxypyrazoles **8a-f**, the 3-hydroxypyrazole **28** was prepared in a two-step reaction sequence starting from arylhydrazines **25**. At first the terminal amino group of hydrazines **25** was protected by acetylation (Scheme 7). Reaction of **26** with β -ketoesters **27** led to the 3-hydroxypyrazoles **28** with high regioselectivity. Subsequent reaction with 4-(2-chloroethyl)morpholine provided the morpholinoethoxypyrazoles **8a-f** (Diaz et al. 2012).

The σ_1 and σ_2 binding affinity of morpholinoethoxypyrazoles **8a-f** was determined with human σ_1 receptors (from transfected HEK-293 cell membrane preparations) and membrane preparations from guinea pig brain (for σ_2).

Generally, the morpholinoethoxypyrazoles **8a-f** show only very low affinity for the σ_2 receptor. Affinity for σ_1 depends on the substitution pattern of the pyrazole ring. Substitution at position 1 with aromatic residues (**8b-f**) produces high σ_1 binding affinity. Non-aromatic residues (*tert*-butyl, **8a**) produce a salient decrease in σ_1 binding affinity (Table 7). Only small residues (e.g., CH_3 , H) are tolerated at 4- and 5-position of the pyrazole ring. In the naphthyl series even a methyl group in



Scheme 7 Synthesis of morpholinoethoxy pyrazoles **8a-f**. Reagents and reaction conditions (a) Ac_2O , toluene, rt; (b) PCl_3 , $50\text{ }^\circ\text{C}$, 2 h; NaH , DMF, $60\text{ }^\circ\text{C}$, 4 h; (c) 4-(2-chloroethyl)morpholine, K_2CO_3 , NaI, DMF, $95\text{ }^\circ\text{C}$, 18 h (Diaz et al. 2012)

Table 7 σ_1 and σ_2 binding affinity of morpholinoethoxy pyrazoles **8a-f**

	R^1	R^2	R^3	σ_1 (h) ^a K_i [nM] \pm SEM	σ_2 (gp) ^b K_i [nM] \pm SEM	σ_1/σ_2 selectivity
8a	<i>tert</i> -butyl	H	H	>1,000	>1,000	–
8b	4-Chlorophenyl	H	H	18 ± 1.5	357 ± 357	20
8c	2-Naph	CH_3	H	17 ± 7.0	>1,000	–
8d	2-Naph	CH_3	CH_3	139 ± 9	>1,000	–
8e	3,4-Dichlorophenyl	CH_3	CH_3	9.4 ± 1.8	351 ± 400	37
8f	3,4-Dichlorophenyl	CH_3	$C(=O)CH_3$	741 ± 134	>1,000	–

^aHuman σ_1 receptor from transfected HEK-293 cell membrane preparations (h)

^bGuinea pig brain membrane preparations (gp)

position 4 seems to be detrimental for high σ_1 affinity (**8d**, $K_i(\sigma_1) = 139$ nM). The introduction of larger moieties in position 4 (e.g., **8f**, $C(=O)Me$) produces a salient decrease in σ_1 binding affinity ($K_i(\sigma_1) = 741$ nM). The 5-alkoxy regioisomer of the most promising ligand **8c** ($K_i(\sigma_1) = 17$ nM) shows a complete loss of σ_1 affinity ($K_i > 1,000$ nM) (Diaz et al. 2012).

The high σ_1 binding affinity of naphthylpyrazole **8c** (also known as S1RA and E-52862) cannot be explained completely by the common pharmacophore models. The 2-[1-(2-naphthyl)pyrazol-3-yloxy]ethyl moiety fits well into the primary hydrophobic region of the Glennon model, tolerating sterically demanding residues. However, the morpholine ring does not fulfill the requirements to address the second hydrophobic region.

Ligands **8** with excellent σ_1 receptor binding affinity and selectivity were further evaluated for their activity at the hERG channel and for efficacy in mouse models of neuropathic pain. Naphthylpyrazole **8c** proved to be the most promising candidate with regard to metabolic stability, interaction with the hERG channel ($IC_{50} > 10\text{ }\mu\text{M}$), and analgesic activity in different pain models (Diaz et al. 2012). It was found that **8c** shows dose-dependent analgesic effects in both the capsaicin-induced hypersensitivity and the formalin-induced pain model. In the partial sciatic nerve ligation mouse model, **8c** shows dose-dependent inhibition of

thermal hypersensitivity and mechanical allodynia, comparable to the effects of pregabalin, the gold-standard for the treatment of neuropathic pain.

The selectivity of **8c** towards 170 other targets including various receptors and ion channels was shown. With the exception of moderate affinity for the human serotonin 5-HT_{2B} receptor ($K_i(5\text{-HT}_{2B}) = 328$ nM) other targets were not engaged by **8c**. The antagonist profile of **8c** was verified using phenytoin as an allosteric modulator of σ_1 .

The chemical properties of **8c** meet Lipinski's "rule of five." The pharmacokinetic properties were evaluated in mice. Due to the acceptable solubility and high metabolic stability, a good oral bioavailability can be assumed.

In light of all of the aforementioned properties, **8c** entered clinical trials. Passing the single and multiple dose phase I clinical study provided proof-of-concept that **8c** is safe and well tolerated by healthy humans. Thus, the development of **8c** will be continued into phase II clinical trials (Abadias et al. 2013).

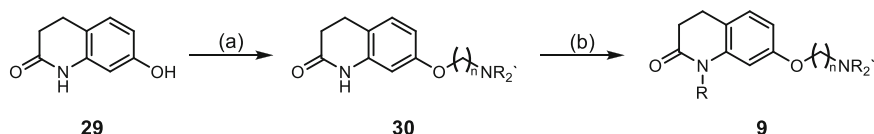
10 3,4-Dihydro-2(1H)-quinolones

3,4-Dihydro-2(1H)-quinolones **9** were developed following the idea of combining a piperidine or morpholine basic element (as realized in the σ_1 receptor antagonist **8c**) with the quinolone scaffold, which was identified as the interacting element at the σ_1 receptor (Oshiro et al. 2000). The aim was to obtain compounds with high affinity for the σ_1 receptor and potent anti-nociceptive properties as shown for S1RA.

For the synthesis of quinolones **9**, 7-hydroxyquinolone **29** was alkylated with dibromoalkans with various length of the alkyl group (Scheme 8). Subsequent reaction with morpholine or piperidine led to the amines **30**, which were converted into 1-alkylated quinolones **9** by reaction with benzyl bromides or iodomethane in presence of NaH (Lan et al. 2014).

The σ_1 and σ_2 affinity of quinolones **9** was determined in competition experiments using guinea pig brain membrane preparations (Table 8).

The distance between the quinolone scaffold and the morpholine ring has a strong impact on σ_1 binding affinity. Whereas **9a** with an ethylene spacer has negligible affinity for σ_1 ($K_1(\sigma_1) > 2,000$ nM), the corresponding homolog **9b** bearing three CH₂ moieties in the side chain binds σ_1 with high affinity ($K_1(\sigma_1) = 14.8$ nM). Elongation by introduction of additional methylene moieties



Scheme 8 Synthesis of quinolones **9**. Reagents and reaction conditions (a) (1) Br(CH₂)_nBr, K₂CO₃, acetone, reflux; (2) HNR₂, K₂CO₃, KI, CH₃CN, reflux; (b) NaH, DMF or THF, 0–50 °C (Lan et al. 2014)

Table 8 σ_1 and σ_2 binding affinity of 3,4-dihydro-2(1*H*)-quinolinones **9**

	R ¹	<i>n</i>	NR ₂ '	σ_1 (gp) ^a K_i [nM] \pm SEM	σ_2 (gp) ^a K_i [nM] \pm SEM	σ_1/σ_2 selectivity
9a	PhCH ₂	2	Morpholine	>2,000	>2,000	–
9b	PhCH ₂	3	Morpholine	14.8 \pm 0.8	471 \pm 38	32
9c	H	3	Piperidine	89 \pm 14	288 \pm 52	3
9d	CH ₃	3	Piperidine	34 \pm 3	357 \pm 131	10
9e	PhCH ₂	3	Piperidine	1.84 \pm 0.33	662 \pm 42	360
9f	4-F-Ph- CH ₂	3	Piperidine	1.22 \pm 0.45	1,301 \pm 204	1,066
9 g	PhCH ₂	4	Morpholine	60 \pm 4.4	530 \pm 46	9
9 h	PhCH ₂	5	Morpholine	88 \pm 5.1	1,811 \pm 37	20
9i	PhCH ₂	6	Morpholine	682 \pm 31	2,000	–

^aGuinea pig brain (gp)

decreased σ_1 affinity in the order $n = 3 > n = 4 > n = 5 > n = 6$. Compound **9i**, with a hexamethylene linker, had the lowest σ_1 binding affinity of this set of compounds with a K_i value of 682 nM. Replacement of the morpholine ring of the most potent ligand **9b** of this series by a piperidine ring led to an almost tenfold increase in σ_1 binding affinity ($K_i(\sigma_1) = 1.84$ nM). For the promising piperidine derivatives, the effect of the quinolinone *N*-substituent on σ_1 receptor binding was evaluated by synthesizing different substituted analogs **9c**, **9d**, and **9f**. It has been found that small residues led to decreased σ_1 affinity compared with the *N*-benzyl substituted derivative **9e** ($K_i(\sigma_1) = 1.84$ nM). In the case of substitution with a proton or a methyl group, the K_i values were only 89 nM (**9c**) or 34 nM (**9d**). The substitution of the phenyl ring with an electron-withdrawing fluorine atom led to a slight increase in σ_1 binding affinity (**9f**, $K_i(\sigma_1) = 1.22$ nM).

Regarding σ_1/σ_2 selectivity, it was found that the most potent σ_1 ligand **9f** has also the highest selectivity with a $\sigma_2 K_i > 1,000$ nM. For the piperidine derivatives **9c-f**, the σ_2 affinity increased with decreased size of substituents. The secondary lactam **9c** shows the highest σ_2 affinity and the lowest σ_1/σ_2 selectivity of this set of compounds ($K_i(\sigma_2) = 288$ nM, $\sigma_1/\sigma_2 = 3$). The chain length between the quinolone scaffold and the morpholine residue also influences affinity for the σ_2 receptor. Compounds **9a** ($n = 2$), **9h** ($n = 5$), and **9i** ($n = 6$) do not show σ_2 affinity. However, compounds **9b** and **9g** with a trimethylene or tetramethylene linker displayed moderate σ_2 affinity, with a K_i of approximately 500 nM.

The most potent ligand **9f** was further evaluated for its anti-nociceptive activity in the formalin-induced pain assay. It was found that **9f** dose-dependently reduced both phases of the pain response with ED₅₀ values of 49.4 \pm 4.1 and 50.5 \pm 2.5 mg/kg for the acute phase I and the longer-lasting tonic phase II, respectively. The σ_1 antagonist activity of **9e** was shown using phenytoin as an allosteric modulator of σ_1 .

11 Conclusion

During the past several years, the fields of σ_1 receptor chemistry and pharmacology have made remarkable progress. Various pharmacophore models of σ_1 ligands, a 3D homology model of the σ_1 receptor, its structure in solution (NMR), and its structure in the solid state (X-ray crystallography) have been reported, allowing a closer look at the binding properties of σ_1 receptors to their ligands. Evidence of σ_1 as a promising target for the development of new therapeutic approaches has been demonstrated. The σ_1 antagonist S1RA (**8c**) is currently in clinical trials for the treatment of neuropathic pain. Bicyclic piperazines **2** and **3** inhibit the growth of small cell lung cancer cells (A-427 cells) in a dose-dependent manner, demonstrating their potential as new tumor therapeutics. The (*S*)-configured spirocyclic σ_1 antagonist **4b** (fluspidine) with a fluoroethyl side chain has been developed as tracer for positron emission tomography (PET) and is currently in clinical trials for imaging and analysis of the brain of patients suffering from major depression.

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