

## Discrepancies in characterization of $\sigma$ sites in the mouse central nervous system

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Received 26 January 1995; revised 7 June 1995; accepted 20 June 1995

### Abstract

The characteristics of [ $^3$ H](+)-pentazocine and [ $^3$ H]1,3-di(2-tolyl)guanidine (DTG) binding to mouse whole brain, cortex, cerebellum and spinal cord membranes were investigated in radioreceptor assays. [ $^3$ H](+)-Pentazocine bound to a single, high affinity site ( $K_d = 1.2$ – $1.6$  nM) with increasing density along the neuraxis from the cortex ( $B_{max} = 543$  fmol/mg protein) to the spinal cord ( $B_{max} = 886$  fmol/mg protein). Hot saturation studies resolved the presence of one binding site for [ $^3$ H]DTG showing no tissue variations in terms of density ( $B_{max} = 1075$ – $1264$  fmol/mg protein) or affinity ( $K_d = 16.6$ – $22.3$  nM). Incubation with 100 nM (+)-pentazocine revealed two classes of high affinity [ $^3$ H]DTG labeled binding sites corresponding to  $\sigma_1$  and  $\sigma_2$  subtypes. A preponderance of  $\sigma_2$  sites was revealed in all investigated tissues. Different pharmacological profiles were demonstrated for the  $\sigma_2$  sites in mouse whole brain compared to mouse spinal cord. However, competition studies indicated that the whole brain and spinal [ $^3$ H](+)-pentazocine labeled  $\sigma_1$  binding sites exhibited similar pharmacological properties. The density of [ $^3$ H](+)-pentazocine labeled  $\sigma_1$  population was found not to match that of [ $^3$ H]DTG labeled  $\sigma_1$  site throughout the mouse central nervous system. The presence of low affinity [ $^3$ H]DTG labeled sites was demonstrated in cold saturation experiments. Equilibrium binding data for the low affinity [ $^3$ H]DTG binding site resulted in an increasing density ( $B_{max} = 1973$ – $11369$  fmol/mg protein) with a decreasing affinity ( $K_d = 242$ – $943$  nM) in mouse cortex through the spinal cord.

**Keywords:**  $\sigma$  Binding site, heterogeneity; Radioreceptor assay; DTG (1,3-di(2-tolyl)guanidine); (+)-Pentazocine; Brain; Spinal cord; (Mouse)

### 1. Introduction

Subtypes of  $\sigma$  sites have been found and characterized in the central nervous system and in a wide variety of peripheral tissues (for review see Su, 1993). Although an array of data are available to indicate their multiple functional role (for review see Su, 1993), no physiological or biochemical function has been incontrovertibly associated with  $\sigma$  receptor(s). The heterogeneity of  $\sigma$  subtypes and the lack of selective  $\sigma$  ligands have largely contributed to the poor consensus on the classification and functional role(s) of  $\sigma$  sites. Based on recent ligand-binding experiments and computer modeling, the possible existence of four pharma-

cologically distinct  $\sigma$  binding sites has been reported. These are (a)  $\sigma_1$ , (b)  $\sigma_2$  (for review see Quirion et al., 1992), (c) dextromethorphan selective (Zhou and Musacchio, 1991) and (d) low affinity  $\sigma$  sites (Karbon et al., 1991; Wu et al., 1991; Codd and Shank, 1992; Connick et al., 1992).

Particular efforts have been taken in several laboratories to develop pharmacological tools to distinguish  $\sigma_1$  from  $\sigma_2$  sites. They differ from one another primarily in their stereoselectivity for (+)-benzomorphans, tissue distribution and sensitivity to GTP analogs.  $\sigma_1$  Binding sites show a high (nanomolar) affinity for (+) isomers of benzomorphans such as pentazocine and *N*-allylnormetazocine (SKF 10,047) while only low to moderate affinities were reported for (–) isomers of these compounds (De Costa et al., 1989; Hellewell and Bowen, 1990; DeHaven-Hudkins et al., 1992; Bowen et al., 1993; Cagnotto et al., 1994).  $\sigma_1$  Sites were found mostly in the central nervous system (CNS). Although

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many findings support the possible link between  $\sigma_1$  sites and GTP-binding regulatory proteins (Itzhak and Stein, 1991; Connick et al., 1992), some data are not compatible with G-protein coupling (DeHaven-Hudkins et al., 1992; Monnet et al., 1994).  $\sigma_2$  Sites were found in high densities in several peripheral tissues (Dumont and Lemaire, 1991; Hellewell et al., 1994) and cell cultures (Hellewell and Bowen, 1990; Georg and Friedl, 1992). Their stereoselectivity profile for benzomorphanes was opposite to the one for  $\sigma_1$  sites having a significantly higher affinity for (–) isomers than for (+) isomers of pentazocine and SKF 10,047 (Hellewell and Bowen, 1990; Dumont and Lemaire, 1991; Georg and Friedl, 1992; Hellewell et al., 1994). Other important tools allowing a clear distinction between  $\sigma_1$  and  $\sigma_2$  sites include 1,3-di(2-tolyl)guanidine (DTG) and phenytoin. [ $^3\text{H}$ ]DTG binds with high, and essentially equal, affinity to both  $\sigma_1$  and  $\sigma_2$  sites (Hellewell and Bowen, 1990; Rothman et al., 1991). In the presence of  $\sigma_1$  blockers, such as (+)-pentazocine, (+)-SKF 10,047 or dextrallorphan, [ $^3\text{H}$ ]DTG has been used as a pharmacological tool for characterization of  $\sigma_2$  sites. The allosteric modulator, phenytoin, also affects  $\sigma_1$  sites without altering  $\sigma_2$  binding (Musacchio et al., 1988; McCann and Su, 1991).

Marked species differences have been shown in the pharmacological profile of the  $\sigma$  binding sites in neural tissues of the rat and guinea pig (Klein and Musacchio, 1990; Walker et al., 1990; Rothman et al., 1991) as well as in the proportion of the  $\sigma_1$  and  $\sigma_2$  subtypes at the same locations (Leitner et al., 1994). Although the presence of the  $\sigma$  binding sites in spinal cord of the rat has been shown in autoradiographic studies (Gundlach et al., 1986; Aanonsen and Seybold, 1989), no comparative data from quantitative radioligand binding assays have been available characterizing subtypes of  $\sigma$  binding sites in spinal cord. In our previous study, differences in the rank order of potencies for displacers of [ $^3\text{H}$ ]DTG binding were found in the mouse brain and spinal cord (Mousseau and Larson, 1994). In the present study, equilibrium bindings of [ $^3\text{H}$ ](+)-pentazocine and [ $^3\text{H}$ ]DTG to homogenates of mouse spinal cord were compared to those of whole brain, cortex and cerebellum. Subtypes of the  $\sigma$  sites were further characterized in competition experiments using various  $\sigma$  ligands. Some of these findings has been previously reported in abstract form (Kovács et al., 1994).

## 2. Materials and methods

### 2.1. Drugs and chemicals

[ $^3\text{H}$ ](+)-Pentazocine (35.3 Ci/mmol) and [ $^3\text{H}$ ]DTG (39.4 Ci/mmol) were purchased from Dupont/New England Nuclear (Boston, MA, USA). (+)-Penta-

zocine and (–)-pentazocine were provided by the National Institute on Drug Abuse. Dextromethorphan, 1,3-di(2-tolyl)guanidine (DTG), (+)-*N*-allylnormetazocine and (–)-*N*-allylnormetazocine (SKF 10,047) and (+)-3-(3-hydroxyphenyl)-*N*-1-(propyl)piperidine (3-PPP) were obtained from Research Biochemicals (Natick, MA, USA). Haloperidol was purchased from Sigma Chemical Co. (St. Louis, MO, USA).

### 2.2. Membrane preparation

Crude membranes were prepared for [ $^3\text{H}$ ](+)-pentazocine and [ $^3\text{H}$ ]DTG binding. Male Swiss-Webster mice (20–25 g, Sasco, Omaha, NE, USA) were decapitated and brains, brain sections and spinal cords were rapidly removed and homogenized in 40 volumes of 10 mM Tris-HCl buffer (pH 7.4) at 4°C with a Brinkmann Polytron (setting 8, for 5 s). The homogenate was centrifuged at  $30\,000 \times g$  for 20 min at 4°C. The resulting pellet was resuspended in the same amount of buffer and incubated at 37°C for 30 min. The suspension was then centrifuged ( $30\,000 \times g$ , 20 min, 4°C) and the final pellet was resuspended in 15 volumes (350–400  $\mu\text{g}/\text{ml}$  protein) of 50 mM Tris-HCl buffer (pH 7.7) at 37°C. The homogenate was used immediately for binding studies.

### 2.3. Receptor binding assays

Binding of [ $^3\text{H}$ ](+)-pentazocine and [ $^3\text{H}$ ]DTG to crude membranes of mouse whole brain, cortex, cerebellum and spinal cord tissues was performed in duplicate in 50 mM Tris-HCl buffer (pH 7.7) at 37°C for 210 min and 60 min respectively. For determination of equilibrium dissociation values ( $K_d$ ) and the number of binding sites ( $B_{\text{max}}$ ) saturation experiments were conducted over a concentration range of 0.05–24 nM [ $^3\text{H}$ ](+)-pentazocine and 0.75–72 nM [ $^3\text{H}$ ]DTG in the absence and presence of 100 nM (+)-pentazocine. To detect the low affinity [ $^3\text{H}$ ]DTG labeled site, 3 nM [ $^3\text{H}$ ]DTG was incubated in the presence of unlabeled DTG ranging in concentration from 0.1 nM to 1  $\mu\text{M}$ . For inhibition assays, 0.75 nM [ $^3\text{H}$ ](+)-pentazocine or 3 nM [ $^3\text{H}$ ]DTG was incubated with 11 concentrations (0.1 nM–100  $\mu\text{M}$ ) of the unlabeled test ligands. Non-specific binding was defined by addition of a final concentration of 10  $\mu\text{M}$  haloperidol. The assays were terminated by rapid filtration through Whatman GF/C glass fiber filter on a Brandel cell harvester using  $3 \times 4$  ml ice-cold 10 mM Tris-HCl buffer (pH 8.0) at 4°C. Filters were pre-soaked in 0.1% polyethylenimine for 2 h at 4°C prior to use. The filter-bound radioactivity was determined by liquid scintillation spectrometry at a 50% efficiency. Membrane protein concentrations were measured using the method of Lowry et al. (1951) with bovine serum albumin as the standard.

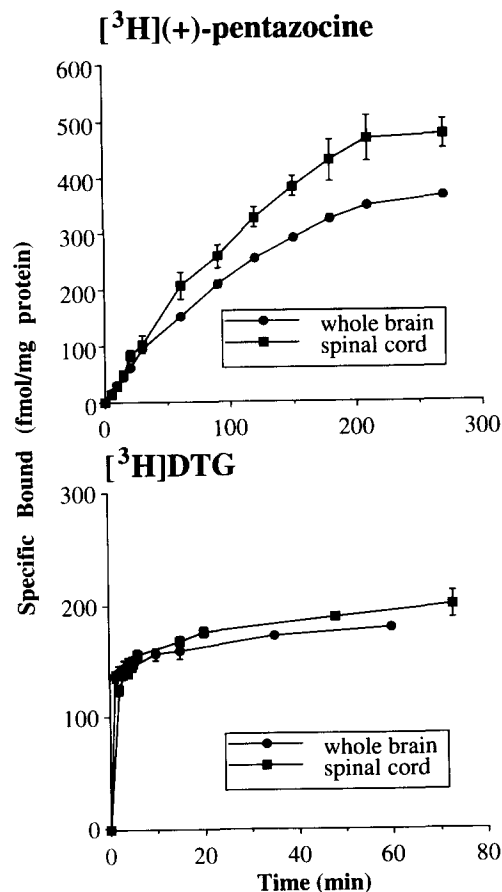


Fig. 1. Time course of binding of [<sup>3</sup>H](+)-pentazocine (1 nM) and [<sup>3</sup>H]DTG (3 nM) to mouse whole brain and spinal cord membranes. Experiments were performed in 50 mM Tris-HCl buffer (pH 7.7) at 37°C. The final assay volume was 1 ml with a tissue concentration of approximately 250 µg protein/ml. Non-specific binding was defined by addition of 10 µM haloperidol. The levels of non-specific binding for [<sup>3</sup>H](+)-pentazocine were 7 fmol/mg protein and 6 fmol/mg protein, for [<sup>3</sup>H]DTG were 17 fmol/mg protein and 50 fmol/mg protein in the whole brain and spinal cord homogenates, respectively. Each point represents the mean ± S.E.M. of 3 independent determinations performed in triplicate.

Equilibrium-saturation and inhibition binding data were analyzed with the EBDA (Munson and Rodbard, 1980) and LIGAND (McPherson, 1983) computer programs. One- and two-site models were fit to the data and an *F*-test was used to determine which model afforded the best fit to the data.  $K_i$  values, the affinity constants for inhibitors were calculated from  $IC_{50}$  values based on the Cheng-Prusoff equation (Cheng and Prusoff, 1973), assuming simple competitive interaction between radioligand and displacer.  $K_i = IC_{50}/(1 + [L]/K_d)$ , where  $[L]$  is the concentration and  $K_d$  is the equilibrium dissociation constants of the radioligand.

#### 2.4. Statistical analysis

Statistical comparison of the  $K_d$  and  $B_{max}$  mean values was performed using one-way analysis of vari-

ance (ANOVA) followed by Scheffe's *F*-test with the level of significance set at  $P < 0.05$ . Student's two-tailed, unpaired *t*-test was used to determine the level of statistical difference between  $K_i$  mean values calculated from whole brain versus spinal cord membranes.

### 3. Results

Specific binding of [<sup>3</sup>H](+)-pentazocine was monophasic, saturable and reached equilibrium slowly. At 37°C, maximal binding was not observed until 180–210 min of incubation (Fig. 1). After reaching equilibrium, the binding remained constant for at least an additional 4 h time period. Computer-assisted Scatchard analysis resulted in the selection of a one-site model as the best fit for [<sup>3</sup>H](+)-pentazocine with  $K_d$  values that did not significantly vary between regions of the CNS (Table 1). The density of [<sup>3</sup>H](+)-pentazocine binding sites ( $B_{max}$ ), however, increased along the neuraxis from  $543 \pm 34$  fmol/mg protein in the cortex to  $886 \pm 57$  fmol/mg protein in the spinal cord (Table 1).

Specific binding of [<sup>3</sup>H]DTG was markedly faster than that of [<sup>3</sup>H](+)-pentazocine reaching its equilibrium after only a 10 min incubation period at 37°C (Fig. 1) and remained stable for at least 2 h. Saturation binding of [<sup>3</sup>H]DTG was performed at increasing concentrations of the radioligand between 0.75 and 72 nM. A single high affinity binding site was observed. The best fit estimates of the  $K_d$  (16.6–22.3 nM) and  $B_{max}$  (1075–1264 fmol/mg protein) values were not significantly different in the whole brain, cortex, cerebellum or spinal cord (Table 2).

Binding sites in mouse whole brain and spinal cord were further characterized with competition studies displacing [<sup>3</sup>H](+)-pentazocine or [<sup>3</sup>H]DTG with

Table 1

Hot saturation binding parameters of [<sup>3</sup>H](+)-pentazocine in mouse whole brain, cortex, cerebellum and spinal cord

Tissue	[ <sup>3</sup> H](+)-Pentazocine binding		
	<i>n</i>	$K_d$ (nM)	$B_{max}$ (fmol/mg protein)
Whole brain	8	$1.3 \pm 0.1$	$640 \pm 37$
Cortex	3	$1.3 \pm 0.6$	$543 \pm 34$
Cerebellum	3	$1.6 \pm 0.2$	$725 \pm 32$
Spinal cord	5	$1.2 \pm 0.1$	$886 \pm 57^a$

Binding was performed in 50 mM Tris-HCl buffer (pH 7.7) at 37°C. Experiments were conducted over a concentration range of 0.05–24 nM [<sup>3</sup>H](+)-pentazocine for 210 min. Non-specific binding was defined by addition of a final concentration of 10 µM haloperidol. The values were determined using the iterative curve-fitting program LIGAND and are the means ± S.E.M. of 3–8 independent determinations performed in duplicate.

<sup>a</sup> Significantly different from corresponding value obtained from whole brain or cortex ( $P < 0.05$ , ANOVA followed by Scheffe's *F* test).

Table 2

Hot saturation binding parameters for [<sup>3</sup>H]DTG in the absence and the presence of 100 nM (+)-pentazocine in mouse whole brain, cortex, cerebellum and spinal cord

Tissue	[ <sup>3</sup> H]DTG binding					
	No (+)-pentazocine added			+ 100 nM (+)-pentazocine <sup>a</sup>		
	<i>n</i>	<i>K<sub>d</sub></i> (nM)	<i>B<sub>max</sub></i> (fmol/mg protein)	<i>n</i>	<i>K<sub>d</sub></i> (nM)	<i>B<sub>max</sub></i> (fmol/mg protein)
Whole brain	3	16.6 ± 0.7	1,108 ± 54	3	17.8 ± 0.8	823 ± 30 <sup>b</sup>
Cortex	4	18.5 ± 1.6	1,075 ± 65	4	21.3 ± 1.8	851 ± 41 <sup>b</sup>
Cerebellum	3	19.0 ± 1.8	1,120 ± 88	4	23.1 ± 0.8	795 ± 48 <sup>b</sup>
Spinal cord	4	22.3 ± 0.9	1,264 ± 116	4	30.0 ± 2.3 <sup>c,d</sup>	1 148 ± 78

Binding was performed in 50 mM Tris-HCl buffer (pH 7.7) at 37°C. Experiments were conducted over a concentration range of 0.75–72 nM [<sup>3</sup>H]DTG in the absence and presence of 100 nM (+)-pentazocine for 60 min. Non-specific binding was defined by addition of a final concentration of 10 μM haloperidol. The values were determined using the iterative curve-fitting program LIGAND and are the means ± S.E.M. of 3–4 independent determinations performed in duplicate.

<sup>a</sup> Note, *B<sub>max</sub>* in the presence of 100 nM (+)-pentazocine represents  $\sigma_2$  binding. Estimated *B<sub>max</sub>* of  $\sigma_1$  site = *B<sub>max</sub>* in the absence of (+)-pentazocine – *B<sub>max</sub>* in the presence of 100 nM (+)-pentazocine. <sup>b,c</sup> Significantly different from value measured in the same tissue in the absence of (+)-pentazocine (<sup>b</sup>*P* < 0.05, <sup>c</sup>*P* < 0.01; unpaired Student's *t*-test). <sup>d</sup> Significantly different from values measured in the presence of (+)-pentazocine in the whole brain or the cortex (*P* < 0.05, ANOVA followed by Scheffe's *F*-test)

various  $\sigma$  ligands. The results of these studies are summarized in Table 3. All  $\sigma$  ligands tested inhibited [<sup>3</sup>H](+)-pentazocine binding at a single site with high affinity. The calculated *K<sub>i</sub>* values were not significantly different in homogenates of whole brain and spinal cord. Suggesting the involvement of  $\sigma_1$  sites, the

rank order of potency of  $\sigma$  ligands in displacing [<sup>3</sup>H](+)-pentazocine was established as follows: haloperidol > (+)-pentazocine > (+)-SKF 10,047 > (–)-pentazocine > (+)-3-(3-hydroxyphenyl)-*N*-1-(propyl)piperidine ((+)-3-PPP) > dextromethorphan > DTG > (–)-SKF 10,047. Our finding indicates that

Table 3

Potencies (*K<sub>i</sub>* (nM)) of various  $\sigma$  compounds to inhibit [<sup>3</sup>H](+)-pentazocine and [<sup>3</sup>H]DTG binding in mouse whole brain and spinal cord

	<sup>[3]H</sup> (+)-Pentazocine binding	<sup>[3]H</sup> DTG binding			+ 100 nM (+)-pentazocine
		No (+)-pentazocine added			
		High affinity <sup>a</sup>	Low affinity	Not distinguished as high or low	
<i>Brain</i>					
(+)-Pentazocine	1.3 ± 0.1	1.3 ± 0.6	1 212 ± 77		
DTG	44.9 ± 6.4	9.6 ± 1.2	468 ± 41		12.7 ± 1.6
(+)-SKF 10,047	5.1 ± 1.1	1.5 ± 0.1	11 385 ± 970		9 771 ± 225
(-)-Pentazocine	15.0 ± 2.0			33.7 ± 3.9	63.6 ± 1.1
(-)-SKF 10,047	935 ± 28			1 284 ± 108	2 159 ± 155
(+)-3-PPP	22.7 ± 2.7			85.1 ± 2.8	302 ± 38
Haloperidol	0.9 ± 0.1			28.2 ± 0.8	53.6 ± 1.3
Dextromethorphan	27.5 ± 0.6			547 ± 29	1 740 ± 122
<i>Spinal cord</i>					
(+)-Pentazocine	1.2 ± 0.1	1.5 ± 0.2	1 842 ± 49 <sup>b</sup>		
DTG	44.1 ± 2.2	8.1 ± 0.7	934 ± 69 <sup>c</sup>		83.6 ± 2.4 <sup>c</sup>
(+)-SKF 10,047	4.1 ± 0.8	2.4 ± 0.6	12 100 ± 107		9 685 ± 103
(-)-Pentazocine	14.1 ± 1.7			31.9 ± 4.3	67.8 ± 3.8
(-)-SKF 10,047	740 ± 34			1 212 ± 83	2 544 ± 258
(+)-3-PPP	24.2 ± 0.6			71.5 ± 7.4	267 ± 50
Haloperidol	1.0 ± 0.1			23.9 ± 4.2	93.4 ± 9.7 <sup>b</sup>
Dextromethorphan	32.1 ± 1.6			585 ± 99	2 015 ± 213

Binding was performed in 50 mM Tris-HCl buffer (pH 7.7) at 37°C. 0.75 nM [<sup>3</sup>H](+)-pentazocine was incubated with 11 concentrations (0.1 nM–100 μM) of the unlabeled test ligands for 210 min. For [<sup>3</sup>H]DTG inhibition assays 3 nM radioligand was incubated with various concentrations of unlabeled test ligands ranging from 0.1 nM to 100 μM (to 1 μM for DTG) in the absence or in the presence of 100 nM (+)-pentazocine. Non-specific binding was defined using 10 μM haloperidol. The *K<sub>i</sub>* values were determined using the iterative curve-fitting program LIGAND and are the means ± S.E.M. of 3–4 independent determinations performed in duplicate.

<sup>a</sup> Note, because DTG labels both  $\sigma_1$  and  $\sigma_2$  sites with equally high affinity, 'high affinity' [<sup>3</sup>H]DTG binding means  $\sigma_1$  and  $\sigma_2$  sites together for DTG. As (+)-pentazocine and (+)-SKF 10,047 have a markedly higher affinity for  $\sigma_1$  than  $\sigma_2$  sites, the term 'high affinity' is associated only with  $\sigma_1$  binding for those ligands. <sup>b,c</sup> Significantly different from corresponding value obtained from brain (<sup>b</sup>*P* < 0.05; <sup>c</sup>*P* < 0.01; unpaired Student's *t*-test).

(+)-pentazocine and (+)-SKF 10,047 were 10 to 200 times more potent displacers of [ $^3$ H](+)-pentazocine than their (–) isomers providing further evidence for stereospecific  $\sigma_1$  binding in mouse brain and spinal cord.

Of several  $\sigma$  ligands tested in competition experiments, (+)-pentazocine and (+)-SKF 10,047 revealed two distinct, high affinity [ $^3$ H]DTG binding sites in either tissues (Fig. 2; Table 2). As both (+)-pentazocine and (+)-SKF 10,047 compete primarily for  $\sigma_1$  sites (Quirion et al., 1992) and DTG binds to both  $\sigma_1$  and  $\sigma_2$  sites with essentially equal affinities, the two high affinity [ $^3$ H]DTG labeled sites in our study presumably corresponded to the  $\sigma_1$  and  $\sigma_2$  population of  $\sigma$  sites. Competition experiments revealed that 100 nM (+)-pentazocine was sufficient to inhibit approximately 30% of the high affinity [ $^3$ H]DTG binding (Fig. 3) yielding a  $K_i$  of 1.3–1.5 nM (Table 2). This portion of the high affinity [ $^3$ H]DTG binding was considered as

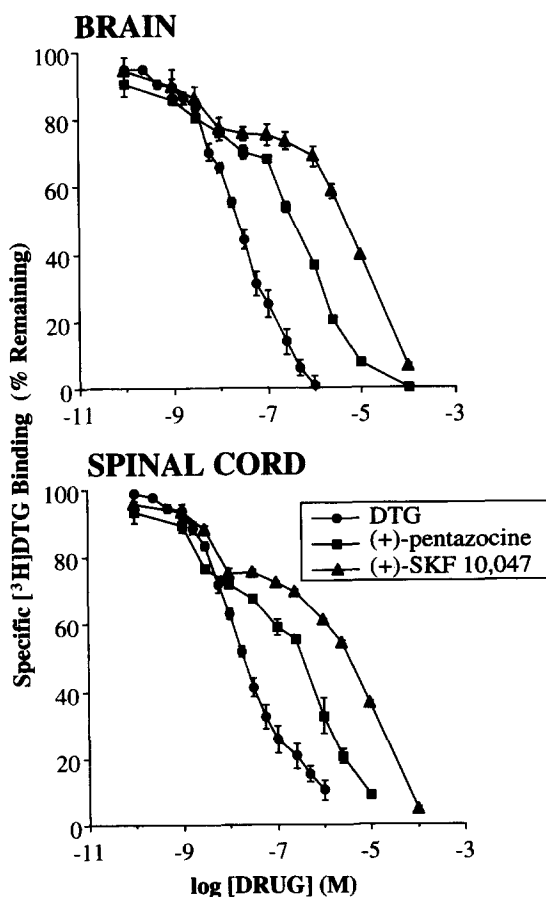


Fig. 2. Inhibition of [ $^3$ H]DTG binding by DTG, (+)-pentazocine and (+)-SKF 10,047 in mouse whole brain and spinal cord. [ $^3$ H]DTG (3 nM) was incubated in 50 mM Tris-HCl buffer (pH 7.7) at 37°C for 60 min in the presence of unlabeled ligands ranging in concentration from 0.1 nM to 100  $\mu$ M. For DTG a concentration range from 0.1 nM to 1  $\mu$ M was used to calculate the equilibrium binding data. Non-specific binding was defined by addition of a final concentration of 10  $\mu$ M haloperidol. Each point represents the mean  $\pm$  S.E.M. of 3–4 independent determinations performed in duplicate.

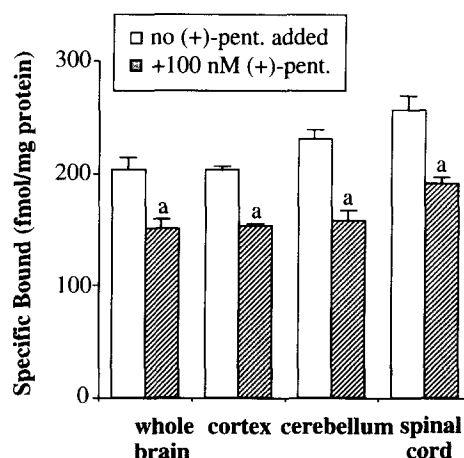


Fig. 3. Specific binding of [ $^3$ H]DTG to mouse whole brain, cortex, cerebellum and spinal cord homogenates. Membranes were incubated with 3 nM [ $^3$ H]DTG in 50 mM Tris-HCl buffer (pH 7.7) at 37°C for 60 min, in the absence and the presence of 100 nM (+)-pentazocine. Non-specific binding was defined by addition of a final concentration of 10  $\mu$ M haloperidol. Each value represents the mean  $\pm$  S.E.M. of 3 independent determinations performed in triplicate. <sup>a</sup> Statistically different from bound [ $^3$ H]DTG values calculated in the absence of (+)-pentazocine,  $P < 0.05$ , paired Student's  $t$ -test.

$\sigma_1$  binding. Hence, the presence of 100 nM (+)-pentazocine could in fact mask  $\sigma_1$  sites providing an effective pharmacological tool to study heterogeneity of the  $\sigma$  sites. In the presence of 100 nM (+)-pentazocine, competition between [ $^3$ H]DTG and a variety of  $\sigma$  ligands produced a rank order of potency of ligands that closely resembled that of the  $\sigma_2$  site. The absence of stereoselectivity of [ $^3$ H]DTG binding for (–)-pentazocine and (–)-SKF 10,047 also indicated that the mouse whole brain and spinal cord homogenates contain binding sites with characteristics of the  $\sigma_2$  site. At  $\sigma_2$  sites, studied in the presence of 100 nM (+)-pentazocine, inhibition of [ $^3$ H]DTG binding by DTG, (+)-pentazocine or haloperidol was found to have significantly higher potencies in the brain than the spinal cord membranes (Table 3). Also, significant regional differences were found in the rank order of potencies for  $\sigma$  ligands in displacing [ $^3$ H]DTG in the presence of 100 nM (+)-pentazocine in the spinal cord versus whole brain. The rank order of potency was in whole brain: DTG > haloperidol > (–)-pentazocine > (+)-3-PPP > (+)-pentazocine > dextromethorphan > (–)-SKF 10,047 > (+)-SKF 10,047, whereas in spinal cord: (–)-pentazocine > DTG > haloperidol > (+)-3-PPP > (+)-pentazocine > dextromethorphan > (–)-SKF 10,047 > (+)-SKF 10,047.

After establishing the presence of  $\sigma_1$  and  $\sigma_2$  sites in tissues of the mouse CNS, equilibrium binding parameters for these two high affinity [ $^3$ H]DTG labeled  $\sigma$  sites were assessed. When the  $\sigma_1$  site was covered by 100 nM (+)-pentazocine, equal densities and same affinities for the  $\sigma_2$  sites were found in the whole

Table 4

Cold saturation binding parameters for [<sup>3</sup>H]DTG in mouse whole brain, cortex, cerebellum and spinal cord

Tissue	[ <sup>3</sup> H]DTG binding				
	High affinity			Low affinity	
	<i>n</i>	<i>K<sub>d</sub></i> (nM)	<i>B<sub>max</sub></i> (fmol/mg protein)	<i>K<sub>d</sub></i> (nM)	<i>B<sub>max</sub></i> (fmol/mg protein)
Whole brain	8	9.6 ± 1.2	637 ± 95	468 ± 41	5590 ± 281
Cortex	3	8.7 ± 1.2	640 ± 124	242 ± 35	1973 ± 78
Cerebellum	3	9.4 ± 1.1	687 ± 54	667 ± 109	3485 ± 206
Spinal cord	5	8.1 ± 0.7	617 ± 53	943 ± 69 <sup>a</sup>	11369 ± 1345 <sup>a,b</sup>

Binding was performed in 50 mM Tris-HCl buffer (pH 7.7) at 37°C. For determination of equilibrium dissociation values (*K<sub>d</sub>*) and the number of binding sites (*B<sub>max</sub>*), 3 nM [<sup>3</sup>H]DTG was incubated for 60 min in the presence of unlabeled ligand ranging in concentration from 0.1 nM to 1 μM. Non-specific binding was defined by addition of a final concentration of 10 μM haloperidol. The values were determined using the iterative curve-fitting program LIGAND and are the means ± S.E.M. of 3–8 independent determinations performed in duplicate.

<sup>a</sup> Significantly different from corresponding value obtained from whole brain or cortex (*P* < 0.05, ANOVA followed by Scheffe's *F*-test).

<sup>b</sup> Significantly different from corresponding value obtained from cerebellum (*P* < 0.05, ANOVA followed by Scheffe's *F*-test).

brain, cortex and cerebellum membranes (Table 2). The density of  $\sigma_1$  sites was calculated by subtracting the number of [<sup>3</sup>H]DTG labeled sites (*B<sub>max</sub>*) detected in the presence of 100 nM (+)-pentazocine from that of measured in the absence of (+)-pentazocine. For whole brain, cortex and cerebellum we found  $\sigma_1$  sites densities of 285, 224 and 325 fmol/mg protein, respectively. In spinal cord, however, hot saturation experiments revealed an increased *K<sub>d</sub>* of DTG which was accompanied by an unchanged *B<sub>max</sub>* in the presence of 100 nM (+)-pentazocine (Table 2). This suggests a very low density of  $\sigma_1$  sites in the spinal cord of the mouse.

To detect the low affinity [<sup>3</sup>H]DTG labeled site, cold saturation experiments were performed. Scatchard analysis of these data indicated the existence of two classes of binding sites as the two-site model yielded a significantly better fit (*F*-test, *P* < 0.001) than fitting experimental data to a one-site model. The calculated numerical values are reported in Table 4. The best fit estimates of the *K<sub>dH</sub>* (8.1–9.6 nM) and *B<sub>maxH</sub>* (617–687 fmol/mg protein) values for the high affinity [<sup>3</sup>H]DTG labeled binding were not significantly different in the whole brain, cortex, cerebellum or spinal cord. [<sup>3</sup>H]DTG also labeled another, bigger population of binding sites with lower affinity. The affinity of DTG to the low affinity  $\sigma$  site was significantly lower in the spinal cord than in the whole brain or the cortex. The density of [<sup>3</sup>H]DTG labeled low affinity binding site in mouse spinal cord was approximately 6 times greater than the amount observed in cortex and approximately 3 times greater compared to the cerebellum.

#### 4. Discussion

The present study provides the first equilibrium binding data and pharmacological characterization of  $\sigma$  binding sites in mouse whole brain, cortex, cerebellum and spinal cord labeled by [<sup>3</sup>H](+)-pentazocine and [<sup>3</sup>H]DTG.

[<sup>3</sup>H](+)-Pentazocine has been proposed to bind to a single, high affinity site in the CNS (De Costa et al., 1989; DeHaven-Hudkins et al., 1992; Bowen et al., 1993; Cagnotto et al., 1994). The presence of another  $\sigma$  site in the human cerebellum recognized by [<sup>3</sup>H](+)-pentazocine with affinity in the nanomolar range has been recently reported (Zabetian et al., 1994). In the present study [<sup>3</sup>H](+)-pentazocine labeled a single class of  $\sigma$  sites with similar *K<sub>d</sub>* values throughout the CNS. Regional differences in the density of  $\sigma_1$  receptor subtype in rat (Leitner et al., 1994; McCann et al., 1994) or guinea pig brain (Walker et al., 1992) have previously been reported. In our present experiments, a significantly larger population of [<sup>3</sup>H](+)-pentazocine labeled sites was found in the spinal cord as compared to the cortex or the whole brain. This suggested a differential distribution of [<sup>3</sup>H](+)-pentazocine labeled  $\sigma_1$  sites. *K<sub>i</sub>* data from competition studies using various  $\sigma$  ligands revealed that [<sup>3</sup>H](+)-pentazocine labeled binding sites in the whole brain and spinal cord were identical.

The present data confirm that [<sup>3</sup>H]DTG interacts with more than one class of binding sites (Hellewell and Bowen, 1990; Karbon et al., 1991; Rothman et al., 1991; Codd and Shank, 1992; Connick et al., 1992). It labeled two sites,  $\sigma_1$  and  $\sigma_2$  sites, with high affinity, and an additional, large population of binding sites with low affinity. The presence of the  $\sigma_2$  receptor subtype in mouse brain and spinal cord was confirmed by the rank order of potency and lack of selectivity of  $\sigma$  ligands when the  $\sigma_1$  site was masked with 100 nM (+)-pentazocine. Similar densities and affinities for  $\sigma_2$  sites were detected in whole brain, cortex and cerebellum. However, we found several discrepancies in the characterization of spinal  $\sigma_2$  binding sites: (a) In the spinal cord, [<sup>3</sup>H]DTG labeled a larger population of the  $\sigma_2$  sites with lower affinity. (b) Potency of (+)-pentazocine, haloperidol and DTG displacing [<sup>3</sup>H]DTG was 1.5-, 1.7-, and 6.6-fold higher in the brain than in the spinal cord. (c) The rank order of potency for (–)-pentazocine, haloperidol and DTG was different

in spinal cord from that found in the brain. These discrepancies raise the possibility that the  $\sigma_2$  sites located in the spinal cord may not be identical with those found in various brain structures.

A large population of [ $^3\text{H}$ ]DTG labeled low affinity binding has been described in the guinea pig (Karbon et al., 1991; Zhou and Musacchio, 1991) and rat brains (Codd and Shank, 1992). The existence of the low affinity [ $^3\text{H}$ ]DTG binding site in the mouse CNS was confirmed in our cold saturation experiments using 3 nM [ $^3\text{H}$ ]DTG. The density of low affinity  $\sigma$  sites was increased while their affinity was decreased along the neuraxis from cortex to spinal cord. The pharmacological characteristics and function of the low affinity site are presently unclear. The possibility that a substantial component of [ $^3\text{H}$ ]DTG binding described as binding to low affinity sites may represent nonspecific binding has also been raised (Basile et al., 1994).

Variation in the ratio of  $\sigma_1$  and  $\sigma_2$  binding across brain regions is subject of intense research. Survey of autoradiographic distribution of [ $^3\text{H}$ ](+)-pentazocine and [ $^3\text{H}$ ]DTG binding in various brain regions of the guinea pig showed significant correlations, but in certain nuclei different ratios were found (Walker et al., 1992). The different distribution of  $\sigma_1$  and  $\sigma_2$  binding among brain regions is consistent with receptor binding assays using rat brain (Leitner et al., 1994; McCann et al., 1994). Enrichment of  $\sigma_2$  sites in rat cortex, and a ratio of  $\sigma_1$  and  $\sigma_2$  sites close to 1.0 in rat cerebellum were reported by McCann and coworkers (1994). However, the amount of ligand bound to  $\sigma_2$  sites exceeded the amount bound to  $\sigma_1$  sites by 300% in both cortex and cerebellum in an other study (Leitner et al., 1994). The presence of larger densities of [ $^3\text{H}$ ]DTG labeled  $\sigma_2$  sites compared to [ $^3\text{H}$ ](+)-pentazocine labeled  $\sigma_1$  sites in the cortex and cerebellum was also confirmed by the present data. The preponderance of  $\sigma_2$  sites was revealed in the mouse spinal cord as well. However, the density of the [ $^3\text{H}$ ](+)-pentazocine labeled  $\sigma_1$  population was found not to match that of the [ $^3\text{H}$ ]DTG labeled  $\sigma_1$  sites throughout the mouse CNS. Proportions of (+)-pentazocine labeled  $\sigma_1$  to [ $^3\text{H}$ ]DTG labeled  $\sigma_1$  site varied by anatomical regions producing ratios ranging from 2.2, 2.4, 2.2 to 7.6 in the whole brain, cortex, cerebellum and spinal cord, respectively. This is in agreement with the proposal that in rat brain [ $^3\text{H}$ ]DTG labels primarily the  $\sigma_2$  site under physiological conditions (Codd and Shank, 1992; Connor and Chavkin, 1992).

Marked species differences have been described among the characteristics of  $\sigma$  binding sites (Klein and Musacchio, 1990; Walker et al., 1990; Rothman et al., 1991; Cagnotto et al., 1994; Leitner et al., 1994). In our study, the potency of (+)-SKF 10,047 relative to (+)-pentazocine, in displacing bound [ $^3\text{H}$ ](+)-pentazocine was 5–22 times higher in mouse brain as compared to

guinea pig brain (De Costa et al., 1989; DeHaven-Hudkins et al., 1992). Also, DTG has been characterized as a good inhibitor of [ $^3\text{H}$ ](+)-pentazocine binding in rat (Cagnotto et al., 1994) or guinea pig brain (De Costa et al., 1989; DeHaven-Hudkins et al., 1992). But we found it to be second to the last in the rank order of potency, followed only by (–)-SKF 10,047. Recent findings suggest a significant species difference in the proportions of the  $\sigma$  receptor subtypes as well. Varying ratios of the [ $^3\text{H}$ ]DTG labeled  $\sigma_2$  to [ $^3\text{H}$ ](+)-pentazocine labeled  $\sigma_1$  binding have been reported; 1.63–3.51 for rat brain regions and 0.67 for guinea pig whole brain (Leitner et al., 1994). Here we report ratios of 1.29, 1.57, 1.01 and 1.30 for whole brain, cortex, cerebellum and spinal cord of the mouse, respectively. Our findings confirmed the presence of species differences in the properties of  $\sigma$  receptor subtypes.

In conclusion, the present study illustrates the complexity of the use of [ $^3\text{H}$ ]DTG as a ligand for  $\sigma$  receptors (Connick et al., 1992) as the following discrepancies were found comparing the binding of [ $^3\text{H}$ ](+)-pentazocine and [ $^3\text{H}$ ]DTG. (a) The density and distribution of  $\sigma_1$  sites labeled by [ $^3\text{H}$ ]DTG and calculated either from hot or cold saturation experiments, were not paralleled by those of labeled by [ $^3\text{H}$ ](+)-pentazocine. (b) The brain and spinal [ $^3\text{H}$ ](+)-pentazocine labeled  $\sigma_1$  binding sites exhibited similar pharmacological properties whereas different pharmacological profiles were found for the  $\sigma_2$  sites in these two tissues. Introduction of ligands showing selectivity for  $\sigma_2$  receptor subtypes (Bonhaus et al., 1993; Bertha et al., 1994; De Costa et al., 1994) can provide additional evidence for the distinct nature of the spinal  $\sigma_2$  site which we observed when we compared it to the  $\sigma_2$  site of the brain.

## Acknowledgements

This work was supported by U.S. Public Health Service Grant DA 04090. We thank Dr. Dénes Budai for his careful review of the manuscript and Drs. Virginia M. Goettl and Julie S. Kreeger for their helpful editorial suggestions.

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