

## ACCELERATED COMMUNICATION

# Identification of Two H<sub>3</sub>-Histamine Receptor Subtypes

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### SUMMARY

The H<sub>3</sub>-histamine receptor provides feedback inhibition of histamine synthesis and release as well as inhibition of other neurotransmitter release. We have characterized this receptor by radioligand binding studies with the H<sub>3</sub> agonist N<sup>-</sup>[<sup>3</sup>H]methylhistamine ([<sup>3</sup>H]NAMHA). The results of [<sup>3</sup>H]NAMHA saturation binding and NAMHA inhibition of [<sup>3</sup>H]NAMHA binding were consistent with an apparently single class of receptors ( $K_D = 0.37$  nM,  $B_{max} = 73$  fmol/mg of protein) and competition assays with other agonists and the antagonists impromidine and dimaprit disclosed only a single class of sites. In contrast, inhibition of [<sup>3</sup>H]NAMHA binding by the specific high affinity H<sub>3</sub> antagonist thioperamide revealed two classes of sites ( $K_{iA} = 5$  nM,  $B_{maxA} = 30$  fmol/mg

of protein;  $K_{iB} = 68$  nM,  $B_{maxB} = 48$  fmol/mg of protein.) Buri-  
 namide, another antagonist that, like thioperamide, contains a  
 thiourea group, likewise discriminated between two classes of  
 sites. In addition to differences between some antagonist poten-  
 cies for the two receptors, there is a differential guanine nucleo-  
 tide sensitivity of the two. The affinity of the H<sub>3A</sub> receptor for [<sup>3</sup>H]  
 NAMHA was reduced less than 2-fold, whereas [<sup>3</sup>H]NAMHA  
 binding to the H<sub>3B</sub> receptor was undetectable in the presence of  
 guanosine 5'-O-(3-thiotriphosphate). The distinction between H<sub>3A</sub>  
 and H<sub>3B</sub> receptor subtypes, the former a high affinity and the  
 latter a low affinity thioperamide site, draws support from pub-  
 lished *in vitro* data.

The H<sub>3</sub>-histamine receptor is a high affinity receptor reported  
 to inhibit central nervous system histamine synthesis (1, 2)  
 and release (1, 3, 4) and serotonin release (5), as well as  
 peripheral norepinephrine (6), acetylcholine (7-9), and neuro-  
 peptide release (10) and histamine synthesis (1). Early evidence  
 for such a receptor derived from high affinity binding of [<sup>3</sup>H]  
 histamine to rat brain membranes (11). More recently, a spe-  
 cific high affinity H<sub>3</sub> agonist, RAMHA, and an H<sub>3</sub> antagonist,  
 thioperamide, have been reported (1).

The binding of [<sup>3</sup>H]RAMHA shows complex kinetics, char-  
 acteristic of ligand binding either to two discrete sites or to a  
 single site that undergoes a change in affinity subsequent to  
 ligand binding (12). We report here that the specific high  
 affinity antagonist thioperamide discriminates between two  
 discrete H<sub>3</sub> receptor subtypes at equilibrium.

### Experimental Procedures

**Materials.** [<sup>3</sup>H]NAMHA (80 Ci/mmol) was from Dupont NEN.  
 NAMHA was from Calbiochem, histamine and pyrilamine from Sigma,  
 impromidine from Smith, Kline, and French Laboratories, and GTP $\gamma$ S  
 from Boehringer Mannheim. All other compounds were from Schering-  
 Plough Research. Male Wistar rats (250 g) were from Charles River  
 Breeding Farms.

**Tissue preparation.** Freshly dissected rat brains were disrupted  
 with a Polytron in 20 volumes of ice-cold Tris·HCl (pH 7.5 at 25°).

Debris was sedimented by 10 min of centrifugation at 1000  $\times$  g, after  
 which membranes were sedimented at 50,000  $\times$  g for 10 min. Mem-  
 branes were resuspended and washed twice. They were stored as pellets  
 at -20° for up to 3 months with no apparent change in [<sup>3</sup>H]NAMHA  
 binding.

**Radioligand binding assay.** Tissue (0.2-0.3 mg of protein) was  
 incubated 40 min in 0.5 ml of 50 mM Tris·HCl at 30°, with or without  
 added compounds, and then filtered over 0.3% polyethylenimine-pre-  
 soaked (13) Whatman GF/C filters (Brandel, Gaithersburg, MD). Fil-  
 ters were washed with three 4-ml volumes of buffer and counted at  
 48% efficiency. Nonspecific binding was determined in the presence of  
 2  $\mu$ M RAMHA or 1  $\mu$ M NAMHA. For competition binding assays, the  
 [<sup>3</sup>H]NAMHA concentration was 0.2 nM (16,000 dpm). At equilibrium,  
 less than 10% of free ligand was bound (1,100 dpm) and, of this, 6%  
 was nonspecific binding (50-70 dpm). Protein was assayed with a  
 bicinchoninic acid reagent (14) (Pierce).

**Data analysis.** Saturation and competition binding experiments  
 were individually analyzed with the LIGAND, weighted, nonlinear,  
 least-squares, curve-fitting program (15). Curves were fit to saturation  
 binding data with the equation

$$[B] = \sum_{i=1}^{i=n} \left( \frac{B_{max_i}[L]}{K_{D_i} + [L]} \right) + N[L]$$

where  $K_{D_i}$  and  $B_{max_i}$  are constants for ligand binding to site  $i$  of  $n$  sites,  
 $[L]$  is the concentration of free radioligand, and  $N$  is the ratio of  
 nonspecific to specific binding at infinite radioligand concentration.  
 Competition curves were fit to the equation modified for multiple

**ABBREVIATIONS:** RAMHA, (R)- $\alpha$ -methylhistamine; NAMHA, N<sup>-</sup>-methylhistamine; GTP $\gamma$ S, guanosine 5'-O-(3-thiotriphosphate).

ligands. Goodness of fit was tested for  $n = 1, 2$ , and  $3$  with an  $F$  test of residual sums of squares corrected for differences in degrees of freedom.

## Results

[<sup>3</sup>H]NAMHA saturation binding data (Fig. 1A) subjected to Scatchard analysis disclosed an apparently single class of sites at equilibrium ( $K_D = 0.37 \pm 0.037$  nM,  $B_{\max} = 73 \pm 3$  fmol/mg of protein, four experiments (Fig. 1B). Analysis of competition binding data for most compounds yielded Hill coefficients ( $n_H$ ) close to 1.0, consistent with an apparently single class of binding sites (Table 1). These compounds include the agonists NAMHA, RAMHA, and histamine and the antagonists impromidine and dimaprit.

However, analysis of competition by the specific high affinity

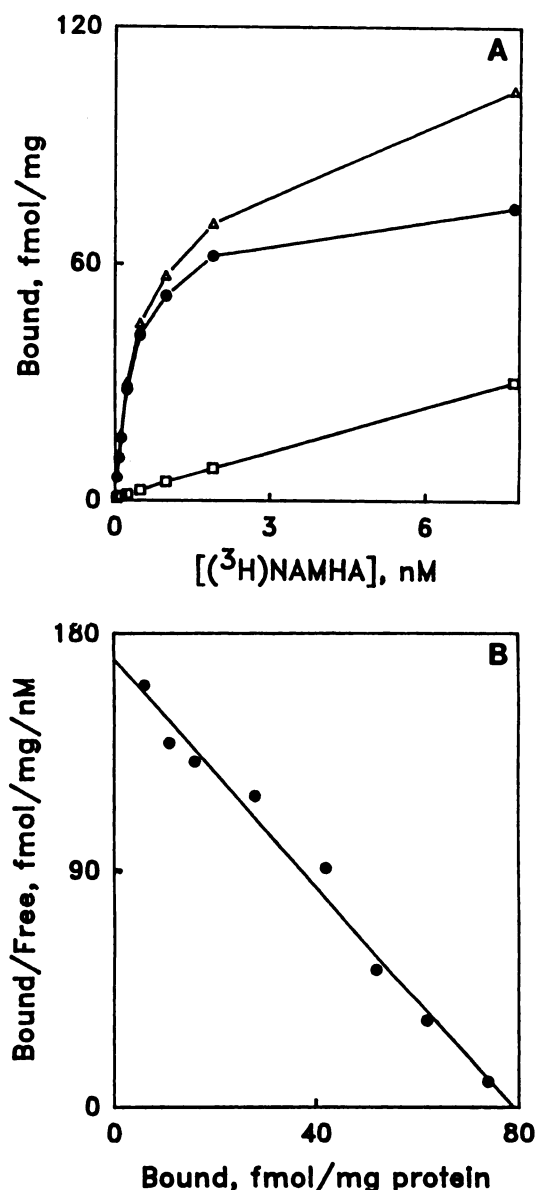


Fig. 1. [<sup>3</sup>H]NAMHA saturation binding. A, Binding isotherm. ●, Specific binding; Δ, total binding; □, nonspecific binding. Points are the averages of triplicate determinations. B, Scatchard plot. Each of four saturation binding experiments was analyzed individually with the LIGAND program (15). Shown is the monoexponential fit from one experiment.

TABLE 1

### Inhibition of [<sup>3</sup>H]NAMHA binding

[<sup>3</sup>H]NAMHA (0.2 nM) was incubated 40 min with 0.3 mg of tissue and 10 concentrations of each compound in duplicate. Nonspecific binding, 5% of total, was the same for all these drugs.  $K_i$  values were determined with the LIGAND program, based on 0.37 nM as the  $K_D$  value for [<sup>3</sup>H]NAMHA. Values are mean  $\pm$  range from two assays. The following compounds inhibited binding 20% or less at 1  $\mu$ M: cimetidine, chlorpheniramine, ranitidine, pyrilamine, tiotidine, and metiamide.

Compound	$K_i$ nM	$n_H$
NAMHA	$0.36 \pm 0.22$	1.0
RAMHA	$1.2 \pm 0$	0.9
Histamine	$4.4 \pm 0.1$	1.0
Impromidine	$45 \pm 1$	1.0
Dimaprit	$520 \pm 150$	1.0

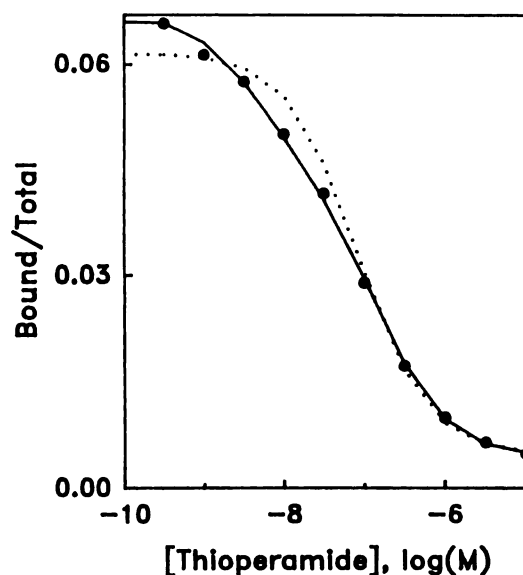


Fig. 2. Thioperamide competition for [<sup>3</sup>H]NAMHA binding sites. . . . . One-site fit; —, two-site fit. Shown is one of six experiments. In all six experiments, a two-site fit was significantly better than a one-site fit ( $p = 0.001$ ).

TABLE 2

### Inhibition of [<sup>3</sup>H]NAMHA binding by compounds that discriminate two sites

Assay details are as for Table 1. Values are mean  $\pm$  standard error (six experiments for thioperamide, four experiments for burimamide). Hill coefficients were 0.7 for thioperamide and 0.6 for burimamide. For only these two compounds was a two-site fit an improvement over one site (thioperamide,  $p = 0.001$ ; burimamide,  $p = 0.02$ ).

Compound	$K_{i_A}$ nM	$K_{i_B}$ nM
Thioperamide	$4.9 \pm 1.6$	$64 \pm 11$
Burimamide	$97 \pm 38$	$3500 \pm 1800$

H<sub>3</sub> antagonist thioperamide for [<sup>3</sup>H]NAMHA binding sites (Fig. 2, Table 2) revealed two classes of H<sub>3</sub> sites ( $K_{i_A} = 4.9 \pm 1.6$  nM,  $B_{\max_A} = 30 \pm 5$  fmol/mg of protein;  $K_{i_B} = 64 \pm 11$  nM,  $B_{\max_B} = 45 \pm 6$  fmol/mg of protein; six experiments), based on  $p = 0.001$  for a two- relative to one-site fit and  $p > 0.05$  for a three- relative to a two-site fit. Burimamide, another antagonist that, like thioperamide, contains a thiourea group in the side chain, also discriminated between two classes of [<sup>3</sup>H]NAMHA binding sites (Table 2).

The possibility that one of these sites might arise from an agonist-induced conformational change in the receptor was

TABLE 3

Densities of high and low affinity thioperamide sites at different times during [ $^3$ H]NAMHA steady state binding

Experimental details are summarized in Table 1. Values are mean  $\pm$  standard error. Six experiments were performed for the 40-min assay, and three for the 60-min assay.

Length of incubation	$K_{iA}$	$B_{maxA}$	$K_{iB}$	$B_{maxB}$
min	nM	fmol/mg of protein	nM	fmol/mg of protein
40	$4.9 \pm 1.6$	$30 \pm 5$	$64 \pm 11$	$45 \pm 6$
60	$8.6 \pm 3.8$	$31 \pm 3$	$123 \pm 25$	$40 \pm 6$

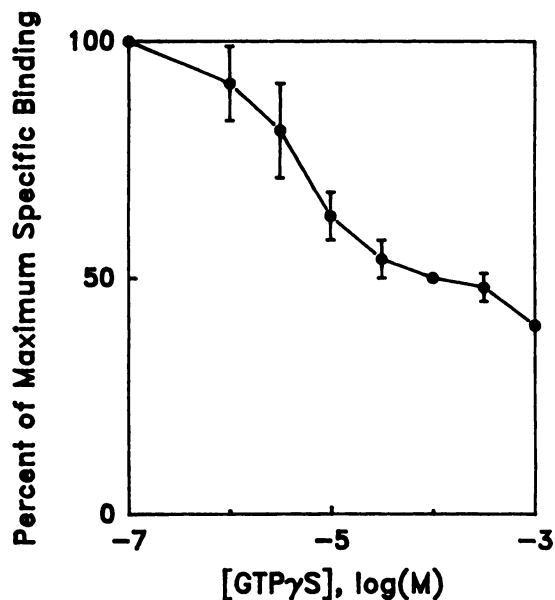


Fig. 3. GTP $\gamma$ S inhibition of specific [ $^3$ H]NAMHA binding. Binding of 0.2 nM [ $^3$ H]NAMHA in the absence and presence of  $10^{-6}$  M NAMHA was determined with various concentrations of GTP $\gamma$ S. Shown is the average  $\pm$  range of two experiments.

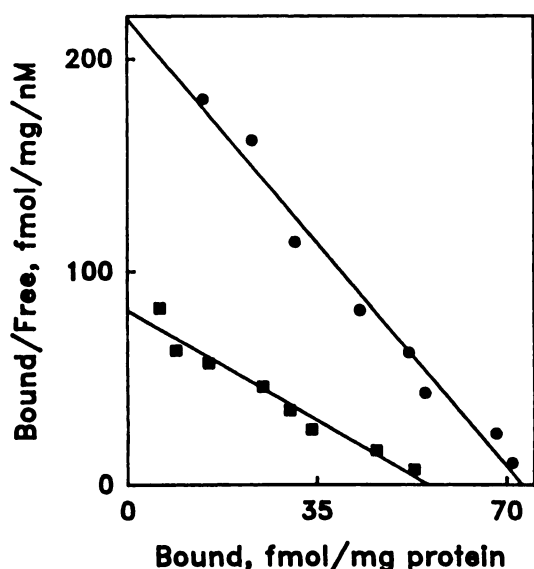


Fig. 4. Scatchard plot of [ $^3$ H]NAMHA saturation binding with and without GTP $\gamma$ S.  $\bullet$ , Control;  $\blacksquare$ , GTP $\gamma$ S. Shown is one of three paired experiments.

TABLE 4

Inhibition of [ $^3$ H]NAMHA binding in the presence of  $10^{-4}$  M GTP $\gamma$ S

Details are as for Table 1, except total and nonspecific binding were determined in the presence of  $10^{-4}$  M GTP $\gamma$ S.

Compound	$K_i$	$n_H$
	nM	
NAMHA	$0.97 \pm 0.33$	0.9
RAMHA	$1.6 \pm 0.2$	0.9
Histamine	$6.7 \pm 1.7$	1.0
Thioperamide	$10 \pm 2$	0.9
Impromidine	$25 \pm 5$	1.0
Burimamide	$240 \pm 90$	0.8
Dimaprit	$370 \pm 30$	0.9

tested by running a thioperamide competition assay sufficiently beyond equilibrium to permit substantial conversion of an initial to a later conformation, if it were going to occur. Clearly, no change in the relative site numbers of high and low affinity sites occurred in the 20 min after steady state binding was reached, although the affinity of thioperamide for both sites appeared to decrease during this time (Table 3).

The specific guanine nucleotide sensitivity of  $H_3$  receptor binding has already been demonstrated (11). We chose to characterize the nature of these effects with GTP $\gamma$ S. Maximum inhibition of [ $^3$ H]NAMHA binding (50% of control) occurred at  $3.2 \times 10^{-5}$  M GTP $\gamma$ S (Fig. 3). A concentration of  $10^{-4}$  M GTP $\gamma$ S was used for subsequent experiments. Two effects of GTP $\gamma$ S were apparent in Scatchard plots of saturation binding (Fig. 4). One was a reduction of apparent site density, an average of  $22 \pm 4\%$  in three experiments; the other was a  $1.7 \pm 0.1$ -fold decrease in apparent affinity of the remaining sites.

To determine whether these effects of GTP $\gamma$ S distinguished between high and low affinity thioperamide sites, competition binding assays were run in the presence of GTP $\gamma$ S (Table 4). Inhibition of [ $^3$ H]NAMHA binding in the presence of GTP $\gamma$ S was best characterized for all competitors by a one-site fit. For thioperamide and burimamide, this corresponded to the high affinity site. The average  $B_{max}$  from these experiments was  $37 \pm 2$  fmol/mg of protein (14 experiments).

## Discussion

We have found with the specific high affinity agonist [ $^3$ H]-RAMHA that binding to rat brain membranes is characterized by biexponential kinetics (12). Additionally, we found that the affinity of NAMHA for the  $H_3$ -histamine receptor was higher in a [ $^3$ H]RAMHA competition binding assay than had been reported for a [ $^3$ H]histamine binding assay (11). In our hands, NAMHA was a more potent competitor than RAMHA itself, so we tested the utility of [ $^3$ H]NAMHA as an  $H_3$ -histamine receptor radioligand. Studies in the guinea pig showed that [ $^3$ H]NAMHA bound with high affinity to a class of sites typical of an  $H_3$ -histamine receptor (16). In the rat, it binds to the same density of sites (Fig. 1B) as [ $^3$ H]RAMHA (12). The inhibitor profile of this binding is characteristic of an  $H_3$  receptor. The  $H_2$  compounds that compete with it for binding are those that are known to bind to an  $H_3$  receptor. Although NAMHA may be less selective than RAMHA for the  $H_3$  receptor (1, 3), the affinities of  $H_1$  and other  $H_2$  compounds make it clear that, at the concentrations of radioligand employed in this assay, binding is specifically to  $H_3$  receptors (Tables 1 and 2). An advantage of [ $^3$ H]NAMHA (Fig. 1A) (16) over [ $^3$ H]-



RAMHA (12) as an H<sub>3</sub> radioligand is the much lower nonspecific binding with [<sup>3</sup>H]NAMHA.

An argument for multiple receptor subtypes based only on the kinetics of radiolabeled agonist binding is bedeviled by the possibility that rapid agonist binding might induce a slow conformational change or enzymatic modification that is recognized as a second component in kinetic studies. Indeed, neither [<sup>3</sup>H]NAMHA equilibrium saturation binding nor competition by NAMHA or other agonists for [<sup>3</sup>H]NAMHA binding sites gave any evidence for multiple agonist-differentiated sites at equilibrium. However, competition by certain antagonists, notably thioperamide, for [<sup>3</sup>H]NAMHA binding sites makes it clear that agonists bind to two H<sub>3</sub> receptor subtypes at equilibrium. Thus, a distinction between an H<sub>3A</sub> site with high affinity for thioperamide ( $K_i = 5$  nM) and an H<sub>3B</sub> site with lower affinity for thioperamide ( $K_i = 64$  nM) can be drawn.

Another recent study (17), as well as this one (Fig. 4), describes two effects of guanine nucleotides on H<sub>3</sub> receptor agonist binding. The dual effect of GTP $\gamma$ S on [<sup>3</sup>H]NAMHA binding is consistent with two receptor subtypes. Agonist binding to the H<sub>3A</sub> receptor is only slightly affected by GTP $\gamma$ S, whereas binding to the H<sub>3B</sub> receptor is undetectable in the presence of GTP $\gamma$ S (Table 4).

Ample evidence already exists for functional distinctions between H<sub>3</sub> receptor subtypes, although it has not been interpreted as such. An 8-fold difference between the potencies of thioperamide to block RAMHA inhibition of histamine synthesis and release in rat brain has been ascribed to an artifact (1, 2). Because the duration of the release assay was 2 min and that of the synthesis assay 30 min, it was reasoned that accumulation of endogenous released histamine in the synthesis assay would result in an actual agonist concentration much greater than the nominal exogenous concentration. Because an antagonist IC<sub>50</sub> value, from which the  $K_i$  is determined, is a function of agonist concentration in the assay (18), a substantial underestimation of thioperamide potency in the histamine synthesis assay would then result (1, 2). However, the same conditions should similarly affect tiotidine antagonism in the two assays, and yet tiotidine was reported to be more potent in the synthesis than in the release assay (2). Our data substantiate the difference between thioperamide potencies in the two assays as real. An H<sub>3A</sub> receptor, one with high affinity for thioperamide, would have been detected in the release assay, whereas an H<sub>3B</sub> receptor, one with low affinity for thioperamide, would have been detected in the synthesis assay.

Insofar as they have been characterized, the guinea pig ileum and mesenteric artery preparations contain H<sub>3</sub> histamine receptors. Of interest in the mesenteric artery preparation is a lack of metiamide antagonist effect, because in the rat brain histamine release assay, which we posit to involve an H<sub>3A</sub> subtype, the  $K_i$  for metiamide is reported as 2.5  $\mu$ M (2). As one of the compounds that contains a thiourea group in its side chain, metiamide might be expected to discriminate between

two classes of sites. However, its very low potency (IC<sub>50</sub>  $\geq 10^{-4}$  M) precluded determination of a full dose-response curve in the binding assay. So it remains a possibility that the H<sub>3B</sub> receptor is assayed in the guinea pig mesenteric artery. More detailed pharmacological characterization of the mesenteric artery and the ileum, with thioperamide and burimamide in particular, may distinguish between the H<sub>3</sub> receptors assayed in these tissues and, as such, provide receptor subtype-specific bioassays. Meanwhile, the rat brain [<sup>3</sup>H]NAMHA binding assay may be useful for developing receptor subtype-specific drugs.

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