

¹²⁵I- α -Conotoxin MII Identifies a Novel Nicotinic Acetylcholine Receptor Population in Mouse Brain

PAUL WHITEAKER, J. MICHAEL MCINTOSH, SIQIN LUO, ALLAN C. COLLINS, and MICHAEL J. MARKS

Institute for Behavioral Genetics, University of Colorado, Boulder, Colorado (P.W., A.C.C., M.J.W.), and Departments of Biology (J.M.M., S.L.) and Psychiatry (J.M.M.), University of Utah, Salt Lake City, Utah.

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ABSTRACT

α -Conotoxin MII (CtxMII), a peptide toxin from the venom of the predatory cone snail *Conus magus*, displays an unusual nicotinic pharmacology. Specific binding of a radioiodinated derivative (¹²⁵I- α -CtxMII) was identified in brain region homogenates and tissue sections. Quantitative autoradiography indicated that ¹²⁵I- α -CtxMII binding sites have a unique pharmacological profile and distribution in mouse brain, being largely confined to the superficial layers of the superior colliculus, nigrostriatal pathway, optic tract, olivary pretectal, and mediolateral and dorsolateral geniculate nuclei. Expression of α -CtxMII binding sites in the nigrostriatal pathway, combined with evidence for α -CtxMII-sensitivity of nicotine-induced [³H]dopamine release in rodent striatal preparations indicates that ¹²⁵I- α -CtxMII binding nicotinic acetylcholine receptors are likely to be physiologically important. Unlabeled α -CtxMII potently ($K_i < 3$ nM) competed for a subset of [³H]epibatidine binding sites in

mouse brain homogenates, but weakly ($IC_{50} > 10$ μ M) interacted with ¹²⁵I- α -bungarotoxin and (–)-[³H]nicotine binding sites, confirming this compound's novel nicotinic pharmacology. Quantitative autoradiography revealed that α -CtxMII binds with high affinity at a subset of [³H]epibatidine binding sites with relatively low cytosine affinity ("cytosine-resistant" sites), resolving [³H]epibatidine binding into three different populations, each probably corresponding to a receptor subtype. The majority population seems to correspond to that which binds nicotine and cytosine with high affinity ("cytosine-sensitive" sites). Comparison of the cytosine-resistant population's distribution with that of $\alpha 3$ subunit mRNA expression suggests that the fractions both more and less sensitive to α -CtxMII probably contain the $\alpha 3$ subunit, perhaps in combination with different β subunits.

Molecular cloning approaches have revealed the expression of 10 nicotinic acetylcholine receptor (nAChR) subunits [$\alpha 2$ – $\alpha 7$, $\alpha 9$ (to date $\alpha 8$ has been identified only in avian neurons), and $\beta 2$ – $\beta 4$] in mammalian neuronal tissue (Lindstrom et al., 1996). Each of the subunits' mRNA has a distinct pattern of expression, suggesting the possibility that they may mediate different processes. In the central nervous system, many of these neuronal nAChRs seem to be presynaptic, where they modulate the release of neurotransmitters such as dopamine, norepinephrine, acetylcholine, and γ -aminobutyric acid with differential pharmacologies (Wonnacott, 1997).

Heterologous expression of neuronal nAChR subunits has shown different subunit combinations (or expression of $\alpha 7$ – $\alpha 9$ alone) result in functional nAChR subtypes with differing pharmacological and biophysical properties (Lind-

strom et al., 1996). The number of neuronal nAChR subunits known theoretically allows the possibility of vast numbers of potential subunit combinations and receptor subtypes. However, efforts to discover which nAChR subtypes exist in the central nervous system (and their locations) have been hindered by the lack of subtype-specific pharmacological probes to identify individual subtypes within the mixed native nAChR population. Indeed, only two subtypes of neuronal nAChRs, those which correspond to ¹²⁵I- α -Bgt and 'high-affinity agonist binding' sites, have been thoroughly characterized so far. In mammalian neuronal systems, ¹²⁵I- α -Bgt is believed to bind largely (if not exclusively) to nAChRs containing the $\alpha 7$ subunit (Schoepfer et al., 1990; Seguela et al., 1992), whereas (–)-[³H]nicotine, [³H]cytosine, [³H]acetylcholine, and [³H]methylcarbamylcholine binding is only detectable at the $\alpha 4\beta 2$ combination of subunits (Whiting and Lindstrom, 1987; Flores et al., 1992). Neuronal bungarotoxin (Bgt), a minor component of the venom of the Taiwanese banded krait *Bungarus multicinctus* (also named κ -Bgt, toxin F, and Bgt 3.1) showed initial promise as a selective antagonist of $\alpha 3\beta 2$ subtype nAChRs (Luetje et al., 1990). However,

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ABBREVIATIONS: nAChR, nicotinic acetylcholine receptor; CtxMII, conotoxin MII; Bgt, bungarotoxin; PEI, polyethylenimine; Y₀- α -CtxMII, N-terminal tyrosine-tagged α -conotoxin MII; TFA, trifluoroacetic acid; SSC, standard saline citrate.

problems of availability (*B. multicinctus* is a protected species), complex kinetics of interaction at multiple nAChR subtypes (Papke et al., 1993), and the difficulty of ensuring the toxin's purity have severely restricted its usefulness. More recently, it has become apparent that the agonist ligand [^3H]epibatidine binds with detectable affinity to other neuronal nAChRs in addition to the $\alpha 4\beta 2$ subtype (Perry and Kellar, 1995; Marks et al., 1998; Parker et al., 1998), raising hopes of identifying further native nAChR populations.

α -CtxMII was identified in the venom of *Conus magus* by sequential fractionation and testing of the isolates for inhibition of $\alpha 3\beta 2$ nAChRs expressed in *Xenopus laevis* oocytes (Cartier et al., 1996). Experiments in native systems have demonstrated potent (nanomolar IC_{50} values) and selective blockade of nAChR subpopulations in rodent striatal (Grady et al., 1997; Kulak et al., 1997; Kaiser et al., 1998) and avian ciliary ganglion (Ullian et al., 1997) preparations. These functional studies provided strong evidence that α -CtxMII was a highly selective antagonist of a novel native receptor population. The high affinity and novel subtype selectivity of α -CtxMII make it a potentially useful ligand, particularly in light of its proven ability to interact with native neuronal nAChRs and the present paucity of selective pharmacological tools.

In this study, a radiolabeled version of α -CtxMII (^{125}I - α -CtxMII) was used to identify, locate, and enumerate α -CtxMII binding nAChRs in mouse brain. The distribution and pharmacology of these receptors differs from those previously characterized, indicating that they represent a novel population. Using unlabeled α -CtxMII in combination with existing nicotinic ligands allowed identification of α -CtxMII binding nAChRs as part of the set of high-affinity [^3H]epibatidine binding nAChRs, but distinct from the traditionally recognized "high-affinity agonist binding" $\alpha 4\beta 2$ subtype. The pharmacological characteristics and distribution of α -CtxMII binding nAChRs indicate that they are likely to be composed of (at least) $\alpha 3$ and $\beta 2$ subunits.

Experimental Procedures

Animals. Male mice (C57BL/6J, 60–90 days old) were used throughout this study. Mice were bred at the Institute for Behavioral Genetics and housed five per cage. The vivarium was maintained on a 12-h/12-h light/dark cycle (lights on 7 AM to 7 PM), and mice were given free access to food and water. The Animal Care and Utilization Committee of the University of Colorado, Boulder, approved all procedures used in this study.

Materials. [^3H]Epibatidine (specific activity, 33.8 Ci/mmol), (–)-[^3H]nicotine (specific activity, 81.5 Ci/mmol), Na^{125}I (specific activity, 2200 Ci/mmol), and uridine triphosphate (α - ^{35}S ; initial specific activity, 800 Ci/mmol) were obtained from DuPont NEN (Boston, MA). ^{125}I - α -Bgt (initial specific activity, 230 Ci/mmol), Hyperfilm β -max, and Hyperfilm- ^3H were purchased from Amersham (Mt. Prospect, IL). NaCl, NaOH, KCl, MgCl_2 , MgSO_4 , CaCl_2 , chloramine T, ammonium acetate, lysozyme, Tris \cdot HCl, sodium carbonate, sodium bicarbonate, polyethylenimine (PEI; 50% w/v solution), yeast tRNA, triethanolamine, sodium citrate, dithiothreitol, Denhardt's solution, acetic anhydride, diethylpyrocarbonate, sodium phosphate, gelatin, poly-L-lysine, chromium aluminum sulfate, BSA (Fraction V), phenylmethylsulfonyl fluoride, EDTA, EGTA, aprotinin, leupeptin trifluoroacetate, and pepstatin A were obtained from Sigma Chemical Co. (St. Louis, MO). (–)-nicotine bitartrate and DPX mountant were bought from BDH Chemicals (Poole, UK). Glass fiber filters Type A/E were obtained from Gelman Sciences (Ann Arbor,

MI) and Type GB from MFS (Dublin, CA). Budget Solve scintillation fluid was purchased from RPI (Arlington Heights, IL). ATP, CTP, GTP, and RNase A were purchased from Boehringer-Mannheim (Indianapolis, IN). The enzymes SP6 RNA polymerase and *Hind*III, were obtained from Promega (Madison, WI). Dextran sulfate was purchased from Pharmacia (Uppsala, Sweden), and formamide from Fluka Chemical Corp. (Ronkonkoma, NY).

Preparation of α -CtxMII and Y_0 - α -CtxMII and Iodination of Y_0 - α -CtxMII. Unmodified α -CtxMII was synthesized as described previously (Cartier et al., 1996). To provide an iodination site, α -CtxMII was synthesized with the addition of a tyrosine at the N terminus (Y_0 - α -CtxMII). Although the two histidine residues present in native MII also provide potential iodination sites, structure-function studies suggested that modification of these sites would lead to unacceptable levels of decreased toxin potency (G. E. Cartier, unpublished observations). Synthesis of Y_0 - α -CtxMII was achieved by methods described previously (Cartier et al., 1996).

To iodinate the peptide, 25 nmol of Y_0 - α -CtxMII was dissolved in 25 μl of H_2O . To this was added 40 μl of 0.3 M NH_4Ac , pH 5.3. Approximately 10 mCi of Na^{125}I (volume \sim 22 μl) was added. The iodination reaction was initiated by the addition of 40 μl of freshly prepared 0.4 mM chloramine T. The reaction proceeded at room temperature (\sim 22°C) for 10 min and was then terminated by the addition of 65 μl of 0.5 M ascorbic acid. The pH of the reaction mix was further lowered by the addition of 0.8 ml of 0.1% trifluoroacetic acid (TFA). Mono- and di-iodinated peptides were separated from unmodified peptides by reversed-phase HPLC using an analytical Vydac C18 column. Buffer A was 0.1% TFA, buffer B was 0.09% TFA, 60% acetonitrile, and the loading loop size was 5 ml. The gradient was 25%–75% B over 50 min. Flow rate was 1 ml/min and absorbance was monitored at 220 nm. A solution of sodium thiosulfate (2.5%) and potassium iodide (0.2%) in 1 N NaOH was added to the waste collection beaker to trap unreacted ^{125}I . Fractions containing peptide were collected in polypropylene tubes containing 10 μl of 20 mg/ml lysozyme to decrease adsorption to the tubes. After collection, peptide material was lyophilized and resuspended in 100 μl of 40% MeOH. Under the above chromatographic conditions, the unmodified peptide elutes at approximately 27 min with moniodo peptide eluting at approximately 29 min. Final yield (estimated from counting an aliquot in a gamma counter) was approximately 2 nmol of moniodo peptide. Monoiodination of the N-terminal tyrosine was verified by chemical sequencing and mass spectrometry.

Quantitative Autoradiography of ^{125}I - α -CtxMII and [^3H]Epibatidine Binding. Quantitative autoradiography procedures were similar to those described previously (Pauly et al., 1989; Marks et al., 1998). Three C57BL/6J mice were sacrificed by cervical dislocation. The brains were removed from the skulls and rapidly frozen by immersion in isopentane (-35°C , 10 s). The frozen brains were wrapped in aluminum foil, packed in ice, and stored at -70°C until sectioning. Tissue sections (14 μm thick) prepared using an IEC Minotome Cryostat refrigerated to -16°C were thaw mounted onto subbed microscope slides (Richard Allen, Richland, MI). Slides were subbed by incubation with gelatin (1% w/v)/chromium aluminum sulfate (0.1% w/v) for 2 min at 37°C , drying overnight at 37°C , incubation at 37°C for 30 min in 0.1% (w/v) poly-L-lysine in 25 mM Tris, pH 8.0, and drying at 37°C overnight. Mounted sections were stored desiccated at -70°C until use. Eight series of sections were collected from each mouse brain.

Before incubation with ^{125}I - α -CtxMII, three adjacent series of sections from each mouse were incubated in binding buffer (144 mM NaCl, 1.5 mM KCl, 2 mM CaCl_2 , 1 mM MgSO_4 , 20 mM HEPES, 0.1% BSA (w/v), pH 7.5) + phenylmethylsulfonyl fluoride (1 mM, to inactivate endogenous serine proteases) at 22°C for 15 min. For all ^{125}I - α -CtxMII binding reactions, the standard binding buffer was supplemented with BSA [0.1% (w/v)], 5 mM EDTA, and 10 $\mu\text{g}/\text{ml}$ each of aprotinin, leupeptin trifluoroacetate, and pepstatin A to protect the ligand from endogenous proteases. The sections were then incubated with 0.5 nM ^{125}I - α -CtxMII for 2 h at 22°C . The first

series of sections was used to determine total ^{125}I - α -CtxMII binding (no competing ligands), the second to measure cytosine-resistant ^{125}I - α -CtxMII binding (in the presence of 20 nM unlabeled cytosine), whereas the third series of sections from each mouse was used to determine nonspecific ^{125}I - α -CtxMII binding (in the presence of 1 μM unlabeled epibatidine). ^{125}I - α -CtxMII binding was further investigated by incubation of sections with varying concentrations of unlabeled ligands (cytosine, 1–300 nM; epibatidine, 10–1000 pM; α -Bgt, 1 μM ; (–)-nicotine, 30–3000 nM). After incubation with ^{125}I - α -CtxMII, the slides were washed as follows: 30 sec in binding buffer + 0.1% (w/v) BSA (22°C), 30 sec in binding buffer + 0.1% (w/v) BSA (0°C), 5 sec in 0.1 \times binding buffer + 0.01% (w/v) BSA (twice at 0°C), and twice at 0°C for 5 sec in 5 mM HEPES (pH 7.5).

Sections for use in [^3H]epibatidine binding were rehydrated in binding buffer at 22°C for 15 min, followed by incubation with 500 pM [^3H]epibatidine for 2 h at 22°C. Four series of adjacent sections were used from each mouse to measure total [^3H]epibatidine binding (no competing ligand), [^3H]epibatidine binding in the presence of 100 nM unlabeled cytosine, [^3H]epibatidine binding in the presence of 100 nM unlabeled cytosine + 50 nM unlabeled α -CtxMII, and non-specific [^3H]epibatidine binding [in the presence of 1 mM unlabeled (–)-nicotine]. Concentrations of unlabeled cytosine and α -CtxMII were chosen on the basis of results obtained in this study from [^3H]epibatidine inhibition binding studies in membrane preparations (Fig. 4, right). Slides were washed by sequential incubation in the following buffers (all steps at 0°C): 5 sec in binding buffer (twice), 5 sec in 0.1 \times binding buffer (twice), and 5 sec in 5 mM HEPES, pH 7.5 (twice).

Sections were initially dried with a stream of air, then by overnight storage (22°C) under vacuum. Mounted, desiccated sections were apposed to film (1–3 days, Amersham Hyperfilm β -Max film for ^{125}I -labeled sections; 8 weeks, Amersham Hyperfilm- ^3H for ^3H -labeled sections). To allow quantification, each film was also exposed to tissue paste standards of defined specific activity (Geary et al., 1985). For tritium, specific activities were 0.05 to 50 nCi/mg (wet weight), whereas for ^{125}I , specific activities were 0.25 to 60 nCi/mg (wet weight). The exact specific activities of the tissue paste standards were determined by measuring radioactivity in weighed aliquots.

After the films had been exposed to the sections for an appropriate length of time, they were developed and signal intensity in selected brain regions was measured by digital image analysis. Films were illuminated using a Northern Light light box, and autoradiographic images of the sections and tissue paste standards were captured using a CCD imager camera. Signal intensity was determined using NIH-Image 1.61. Where possible, six independent measurements from different tissue sections were made for each brain region, under each incubation condition, for each mouse. The absorbance measurements for each brain area were averaged, and the mean absorbance was used to calculate the degree of labeling by reference to the relevant standard curve.

Membrane Preparation. Each C57BL/6J mouse was sacrificed by cervical dislocation. The brain was removed from the skull and placed on an ice-cold platform. Brains were either dissected into 12 regions [olfactory bulbs, cerebellum, hindbrain (pons-medulla), hypothalamus, hippocampus, striatum, cerebral cortex, thalamus, mid-brain, interpeduncular nucleus, superior colliculus, and inferior colliculus] or the hindbrain, cerebellum, and olfactory bulbs were discarded without further dissection ("whole brain" preparation). Samples were homogenized in ice-cold hypotonic buffer (14.4 mM NaCl, 0.2 mM KCl, 0.2 mM CaCl_2 , 0.1 mM MgSO_4 , 2 mM HEPES, pH 7.5) using a glass-Teflon tissue grinder. The particulate fractions were obtained by centrifugation at 20,000g (15 min, 4°C; Sorval RC-2B centrifuge). The pellets were resuspended in fresh homogenization buffer, incubated at 37°C for 10 min, then harvested by centrifugation as before. Each pellet was washed twice more by resuspension/centrifugation, then stored (in pellet form under homogenization buffer) at –70°C until used. Protein concentrations in

the membrane preparations were measured using the method of Lowry et al. (1951), using BSA as the standard.

(–)-[^3H]Nicotine Binding to Membranes. The binding of (–)-[^3H]nicotine was measured using the method of Marks et al. (1986), modified for use with a 96-well plate washer. Membrane samples (200 μg of whole brain preparation) were incubated in 96-well polystyrene plates with 20 nM (–)-[^3H]nicotine in 100 μl of binding buffer for 30 min at 22°C. Binding reactions were terminated by filtration of samples onto PEI-soaked (0.5% w/v in binding buffer) glass fiber filters (types GFA/E and GB) using an Inotech Cell Harvester (Inotech, East Lansing, MI). Samples were subsequently washed six times with ice-cold binding buffer. Total and nonspecific [in the presence of 1 mM (–)-nicotine tartrate] binding were determined in triplicate. Where inhibition binding was being measured, various concentrations of competing ligands were included in triplicate wells.

^{125}I - α -Bgt Binding to Membranes. Binding of ^{125}I - α -Bgt to membrane preparations was performed using procedures similar to those used with (–)-[^3H]nicotine, except that incubation times were extended to 5 h and samples contained 1 nM ^{125}I - α -Bgt instead of 20 nM (–)-[^3H]nicotine. The GFA/E filters were treated with 0.5% PEI and the GFB filters were treated with 5% (w/v) nonfat dry milk before filtration.

[^3H]Epibatidine Binding to Membranes. Binding of [^3H]epibatidine was quantified as described previously (Marks et al., 1998). Incubations were performed in 1-ml polypropylene tubes in a 96-well format, using 50 to 200 μg of membrane protein per tube (depending on brain region). A 500- μl reaction volume was used to minimize problems of ligand depletion, and all incubations progressed for 2 h at 22°C. The concentration of [^3H]epibatidine (500 pM) used in inhibition binding experiments was chosen to maintain binding of ligand to the tissue at 5% or less of total ligand added. Saturation binding experiments were performed for membrane preparations from each brain region, using ligand concentrations in the range 10 to 800 pM. At the lower concentrations, a significant proportion of ligand bound to the tissue. Free [^3H]epibatidine concentrations were estimated by correcting for the amount of ligand bound to tissue, and these corrected concentrations were used to calculate K_d values for [^3H]epibatidine binding in each brain region and, thus, the K_i values for each compound versus [^3H]epibatidine binding.

^{125}I - α -CtxMII Binding to Membranes. Large amounts of non-specific binding were seen when using ^{125}I - α -Ctx (0.2–32 nM) in filtration binding assays. Best assay performance was produced using the following modifications to the (–)-[^3H]nicotine binding procedure. Incubation times were extended to 2 h, and incubation buffer was supplemented with BSA [0.1% (w/v)], 5 mM EDTA, 5 mM EGTA, and 10 $\mu\text{g}/\text{ml}$ each of aprotinin, leupeptin trifluoroacetate, and pepstatin A to protect the ligand from endogenous proteases. The glass fiber filters were treated with 5% (w/v) nonfat dry milk before filtration.

In Situ RNA Hybridization. The method used for in situ hybridization using riboprobes was identical with that used by Simmons et al. (1989) and Marks et al. (1992). Probes were prepared by in vitro transcription, using α - ^{35}S -UTP as the sole source of UTP. The $\alpha 3$ probe was prepared from clone PCA48E(4) cloned in pSP65, linearized using *Hind*III, and synthesized using SP6 RNA polymerase. The synthesis was designed to yield full-length antisense transcript. Immediately before hybridization, the probe was subjected to alkaline hydrolysis using the method of Cox et al. (1984) to yield products with average sizes of 500 bases.

After hybridization, slides were air dried and stored under vacuum overnight before exposure to Amersham Hyperfilm β -Max film (10 days). To allow $\alpha 3$ hybridization to be quantified, the film was also exposed to a set of dot-blotted ^{35}S standards. Serial dilutions of the [^{35}S]cRNA were made in 5 \times standard saline citrate (SSC; 1 \times SSC, 150 mM NaCl, 15 mM trisodium citrate, pH 7.0, with HCl), and 400 μl of each dilution was applied to a prewetted (5 \times SSC) nylon membrane (New England Nuclear, Beverly, MA) by vacuum filtra-

tion through a 96-well manifold (Life Technologies, Bethesda, MD). The samples were washed three times with $5\times$ SSC and allowed to dry at room temperature. One set of the dilution series was cut from the membrane and counted on a liquid scintillation counter to determine exact counts per unit area. Standards ranged from 0.1 to 60 pCi/mm². After exposure to the sections and standards, the films were developed. Autoradiographic images were captured and hybridization densities quantified as described above for autoradiographic analysis of ligand binding.

Calculations. Results for saturation binding experiments were calculated using the Hill equation: $B = B_{\max} L^n / (L^n + K_d^n)$, where B is the binding at the free ligand concentration L , B_{\max} is the maximum number of binding sites, K_d is the equilibrium binding constant, and n is the Hill coefficient. Values of B_{\max} , K_d , and n were calculated using the nonlinear least-squares fitting algorithm of Sigma Plot version 5.0 (Jandel Scientific, San Rafael, CA). Results for inhibition of (–)-[³H]nicotine and [¹²⁵I]- α -Bgt binding were calculated using a one-site fit: $B = B_o / [1 + (I / IC_{50})]$ where B is ligand bound at inhibitor concentration I , B_o is the binding in the absence of inhibitor, and IC_{50} is the concentration of inhibitor required to reduce binding to 50% of B_o . Results for inhibition of ligand binding were calculated using the formulae for either one (as above) or two binding sites: $B = B_1 / [1 + (I / IC_{50-1})] + B_2 / [1 + (I / IC_{50-2})]$, where B is ligand bound at inhibitor concentration I , and B_1 and B_2 are binding sites sensitive to inhibition with IC_{50-1} and IC_{50-2} , respectively. Values for K_i (inhibition binding constant) were derived by the method of Cheng and Prusoff (1973): $K_i = IC_{50} / 1 + (L / K_d)$.

Results

¹²⁵I- α -CtxMII Autoradiography, Mouse Brain Sections. Preliminary experiments (using monoiodinated but nonradioactive Y_o - α -CtxMII) showed that it remained a potent inhibitor at $\alpha 3\beta 2$ nAChRs expressed in *X. laevis* oocytes ($K_d = 1.9$ nM, compared with 0.35 nM for the native toxin; data not shown). In light of these data, attempts were made to identify specific, nicotinic α -CtxMII binding using the monoradioiodinated version of this ligand (¹²⁵I- α -CtxMII) in mouse brain sections.

To minimize nonspecific binding, and reduce the possibility of labeling lower-affinity receptor populations, a low (0.5 nM) concentration of [¹²⁵I]- α -CtxMII was used in autoradiography experiments. Under these conditions, a small amount of tissue mediated nonspecific binding of [¹²⁵I]- α -CtxMII binding (defined in the presence of 1 μ M unlabeled epibatidine) was seen. However, specific [¹²⁵I]- α -CtxMII binding could be clearly distinguished over the tissue background (Fig. 1). Minor variations in nonspecific binding were noted, making it necessary to measure nonspecific binding in each individual region to ensure accurate quantification of specific signal strength. Subsequent experiments showed that using higher [¹²⁵I]- α -CtxMII concentrations produced unacceptably high levels of nonspecific binding. Monoiodination produced radio-labeled toxin of high specific activity (2200 Ci/mmol), allowing short film exposures (24–72 h). Toxin was used through one half-life without any detectable increase in nonspecific signal or decrease in specific signal strength.

The highest levels of specific [¹²⁵I]- α -CtxMII binding [>5 fmol/mg (wet weight)] were detected in the dorsolateral and ventrolateral geniculate nuclei, olivary pretectal nucleus, and the zonal layer of the superior colliculus. High levels (4–5 fmol/mg) of [¹²⁵I]- α -CtxMII binding were also detected in the superficial gray of the superior colliculus and in the oculomotor nerve. Outside these highly labeled regions, binding

densities were lower and sites were mainly found in nigrostriatal and optic-tract-associated regions, as summarized in Table 1.

The distribution of specific [¹²⁵I]- α -CtxMII binding resembled a subset of the cytosine-resistant high affinity [³H]epibatidine binding population reported by Marks et al. (1998). The abilities of unlabeled epibatidine, cytosine, α -Bgt, and nicotine to compete for [¹²⁵I]- α -CtxMII-binding sites were measured by quantitative autoradiography. Competition binding was assessed in striatum and the superficial gray of the superior colliculus, as most of the other regions containing [¹²⁵I]- α -CtxMII binding are too small to allow sufficient tissue slices to be collected. Even high concentrations of α -Bgt (1 μ M) had no effect on [¹²⁵I]- α -CtxMII-binding. In contrast, all three of the remaining ligands competed effectively

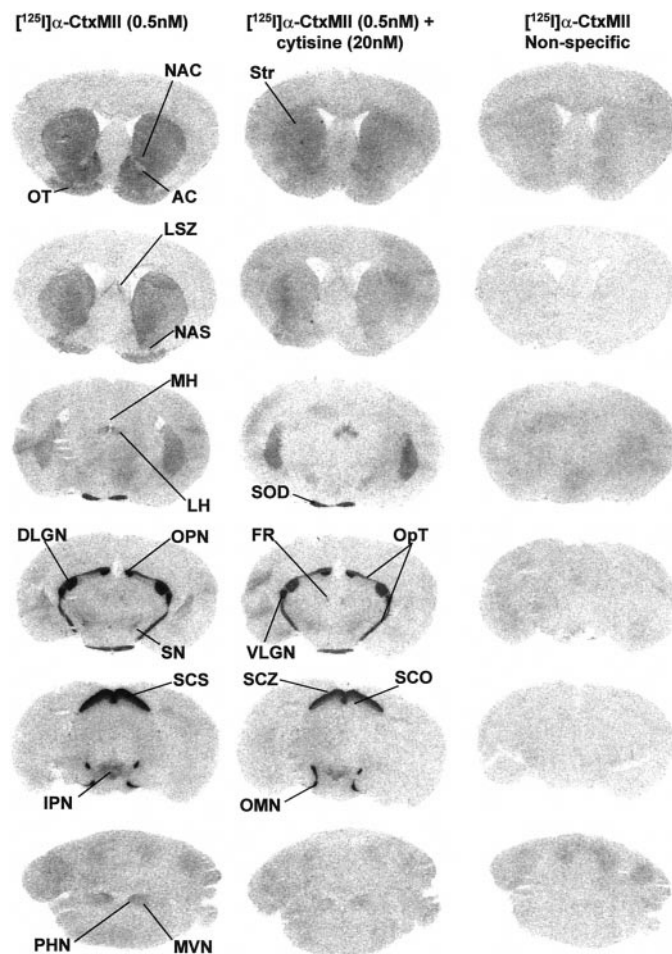


Fig. 1. Autoradiographic representation of [¹²⁵I]- α -CtxMII binding in mouse brain [total, with cytosine (20 nM), and nonspecific]. Sections (14 μ m) were incubated with 0.5 nM [¹²⁵I]- α -CtxMII alone (left column), in the presence of 20 nM cytosine (center column), and with 1 μ M epibatidine (nonspecific [¹²⁵I]- α -CtxMII; right column) as described under *Experimental Procedures*. The sections are adjacent in each row, and the panels are digital images of autoradiograms. The abbreviations used to identify brain regions are: AC, anterior commissure; DLGN, dorsolateral geniculate nucleus; FR, fasciculus retroflexus; IPN, interpeduncular nucleus; LH, lateral habenula; LSZ, lambdoid septal zone; MH, medial habenula; MVN, medial vestibular nucleus; NAC, nucleus accumbens core; NAS, nucleus accumbens shell; OMN, oculomotor nerve (or root); OPN, olivary pretectal nucleus; Opt, optic tract; OT, olfactory tubercle; PHN, prepositus hypoglossal nucleus; SCO, superior colliculus, optic nerve layer; SCS, superior colliculus, superficial gray; SCZ, superior colliculus, zonal layer; SN, substantia nigra; SOD, supraoptic decussation; Str, striatum; VLGN, ventrolateral geniculate nucleus.

To determine whether ^{125}I - α -CtxMII binding sites nAChRs are uniformly cytosine resistant, the cytosine sensitivity of ^{125}I - α -CtxMII binding was assessed by addition of 20 nM cytosine to the binding buffer (a concentration that the previous workers' data indicated would abolish binding to cytosine-sensitive [^3H]epibatidine binding nAChRs). Binding of ^{125}I - α -CtxMII (0.5 nM) was noticeably diminished by incubation with cytosine (20 nM) but was not reduced to background levels (Fig. 1; Table 1). Comparison of ^{125}I - α -CtxMII binding densities in the presence and absence of cytosine showed that across brain regions, addition of 20 nM cytosine reduced ^{125}I - α -CtxMII binding by an average of 40% (Table 1, right column; Fig. 3). In all regions where it was detectable, specific ^{125}I - α -CtxMII binding displayed the same cytosine sensitivity (correlation analysis showed $r = 0.96$; Fig. 3).

(-)-[³H]Nicotine, ¹²⁵I-α-Bgt and [³H]Epibatidine Binding, Mouse Brain Membrane Preparations. To ex-

Regional distribution of specific mouse brain ^{125}I - α -CtxMII binding in the presence and absence of 20 nM cytisine.

	Total ^{125}I - α -CtxMII Binding	^{125}I - α -CtxMII Binding with Cytisine (20 nM)	% Cytisine-Resistant (of total)
<i>fmol / mg wet wt.</i>			
Telencephalon			
Neocortex			
Olfactory tubercles	0.58 \pm 0.05	0.11 \pm 0.03	18.9
Basal ganglia			
Nucleus accumbens, core	1.44 \pm 0.07	0.66 \pm 0.06	45.8
Nucleus accumbens, shell	0.86 \pm 0.04	0.39 \pm 0.03	45.3
Striatum	0.81 \pm 0.02	0.38 \pm 0.07	46.9
Substantia nigra	0.75 \pm 0.19	0.39 \pm 0.13	52.0
Ventral tegmental area	1.29 \pm 0.06	0.96 \pm 0.11	74.4
Septum			
Lambdoid septal zone	0.82 \pm 0.13	0.34 \pm 0.08	41.5
Diencephalon			
Metathalamus			
Dorsolateral geniculate nucleus	5.00 \pm 0.20	2.55 \pm 0.26	51.0
Olivary pretectal nucleus	5.97 \pm 0.12	3.20 \pm 0.39	53.6
Ventrolateral geniculate nucleus	5.84 \pm 0.18	3.31 \pm 0.29	56.7
Epi- and subthalamus			
Lateral habenula	0.83 \pm 0.11	0.43 \pm 0.03	51.8
Medial habenula	1.11 \pm 0.05	0.68 \pm 0.07	61.3
Mesencephalon			
Interpeduncular nucleus	1.38 \pm 0.04	1.05 \pm 0.03	76.1
Superior colliculus, optic nerve layer	1.21 \pm 0.03	0.65 \pm 0.03	53.7
Superior colliculus, superficial grey	4.13 \pm 0.01	2.23 \pm 0.10	54.0
Superior colliculus, zonal layer	5.33 \pm 0.10	3.47 \pm 0.06	65.1
Pons			
Medial vestibular nucleus	0.50 \pm 0.04	0.15 \pm 0.02	30.0
Prepositus hypoglossal nucleus	0.56 \pm 0.05	0.39 \pm 0.06	69.6
Fiber Tracts			
Anterior commissure, anterior	0.26 \pm 0.07	0.03 \pm 0.03	11.5
Fasciculus retroflexus	0.80 \pm 0.15	0.58 \pm 0.07	72.5
Oculomotor nerve (or root)	4.71 \pm 0.56	3.52 \pm 0.25	74.7
Optic tract/brachium superior colliculus	2.66 \pm 0.15	1.66 \pm 0.19	62.4
Supraoptic decussation	2.48 \pm 0.04	1.44 \pm 0.15	58.1

pand on the data provided by ^{125}I - α -CtxMII competition binding experiments, the ability of α -CtxMII to displace (–)-[^3H]nicotine, ^{125}I - α -Bgt, and [^3H]epibatidine binding to mouse brain membrane preparations was assessed. For comparison, the ability of the nicotinic agonist cytosine to inhibit binding of the same ligands was also measured.

Inhibition of (–)-[^3H]nicotine (20 nM) and ^{125}I - α -Bgt (1 nM) binding (to $\alpha 4\beta 2$ and $\alpha 7$ nAChRs, respectively) was measured in whole-brain membrane preparations. Cytosine produced a monophasic inhibition of both (–)-[^3H]nicotine and ^{125}I - α -Bgt binding (Fig. 4) but was much more potent in its interaction with the former ($K_i = 0.36 \pm 0.04$ nM and 1.1 ± 0.5 μM , respectively). In contrast, α -CtxMII was only a weak inhibitor of (–)-[^3H]nicotine and ^{125}I - α -Bgt binding ($K_i > 10$ μM in each case).

Filtration binding was used to measure cytosine and α -CtxMII inhibition of [^3H]epibatidine binding in membrane preparations from 12 brain regions. Levels of [^3H]epibatidine binding varied widely among brain regions (Table 2). Levels were particularly high in the interpeduncular nucleus, with large amounts of [^3H]epibatidine binding also detected in superior colliculus and thalamic membranes. The lowest amounts of binding were found in the cerebellum and olfactory bulbs. As reported previously (Marks et al., 1998), saturation binding analysis provided [^3H]epibatidine binding K_d values of 20 to 40 pM in each region investigated, with no evidence for multiphasic ligand binding (data not shown). Also matching the results of Marks et al. (1998), cytosine inhibition of [^3H]epibatidine binding exhibits two phases (with $K_i = 0.27 \pm 0.05$ nM and 32 ± 6 nM: “cytosine-sensitive” and “cytosine resistant,” respectively), indicating that [^3H]epibatidine binds to receptor populations with differen-

tial cytosine affinity (but similar affinities for [^3H]epibatidine, according to the saturation binding profiles).

In the majority of regions surveyed, the more cytosine-sensitive phase of [^3H]epibatidine binding [believed to correspond to the high-affinity (–)-[^3H]nicotine binding site (Marks et al., 1998)] predominated. However, in olfactory bulb and interpeduncular nucleus preparations, the density of binding sites less sensitive to cytosine inhibition equaled or exceeded that of the cytosine-sensitive fraction (50% and 62% of [^3H]epibatidine binding in the two regions, respectively). Hippocampal membranes contained only cytosine-sensitive [^3H]epibatidine binding. Binding of [^3H]epibatidine was not detectably inhibited by α -CtxMII in interpeduncular nucleus, hindbrain, olfactory bulb, hippocampal, or cerebellar membranes. However, it is important to note that these region's small size (interpeduncular nucleus) or low overall receptor densities (hippocampus, hindbrain, olfactory bulb, and cerebellum) resulted in low numbers of total [^3H]epibatidine counts being retained, so minor populations of α -CtxMII-sensitive [^3H]epibatidine binding sites may have been overlooked. In contrast, α -CtxMII potently [$K_i = 2.7 \pm 1.3$ nM (mean \pm S.E.M. of values in the seven regions where α -CtxMII competition was observed); Table 2] inhibited a fraction of [^3H]epibatidine binding in the remaining brain regions studied. At the concentrations studied (30 pM–300 nM), α -CtxMII inhibition of [^3H]epibatidine binding appeared monophasic. The α -CtxMII-sensitive fraction of [^3H]epibatidine binding was approximately equal to the cytosine-resistant portion in the superior colliculus, striatum, and olfactory tubercles but was smaller in the remaining brain regions. Statistical analysis showed no significant differences in K_i values between regions for α -CtxMII inhibition

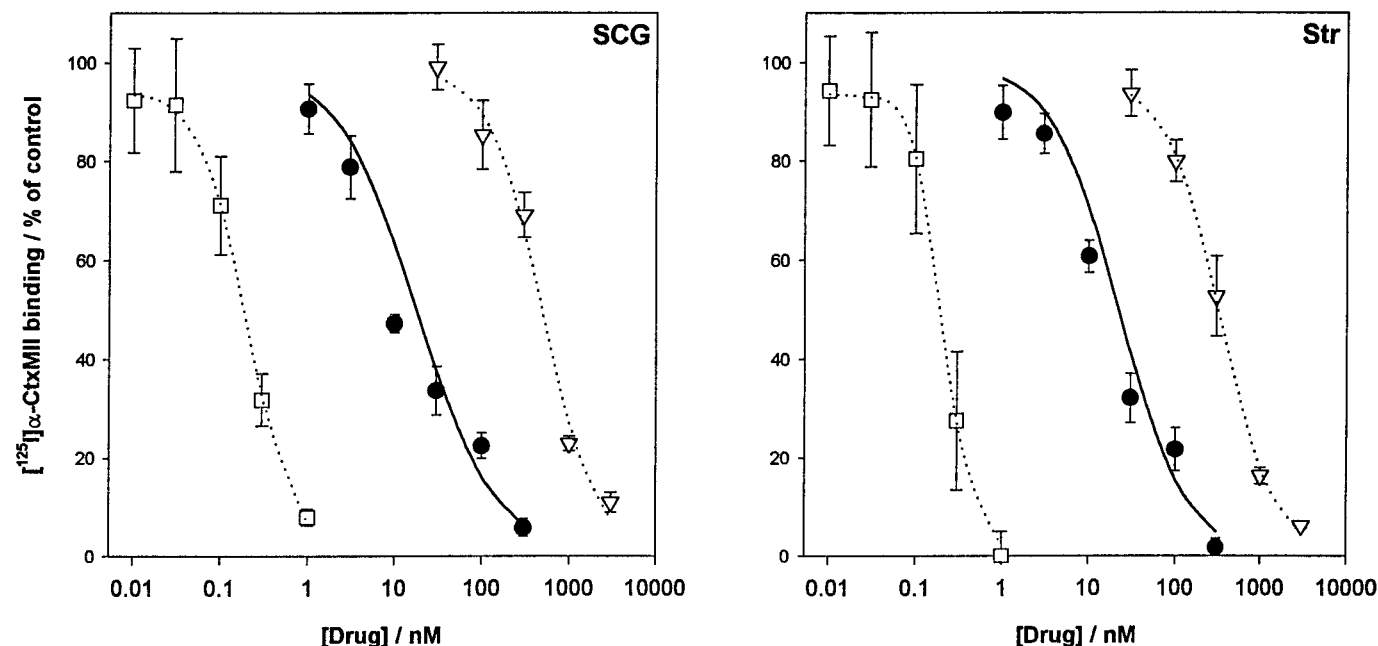


Fig. 2. Competition by cytosine, epibatidine, (–)-nicotine, and α -Bgt for ^{125}I - α -CtxMII binding sites in mouse brain. Sections (14 μm) were incubated with 0.5 nM ^{125}I - α -CtxMII in the presence of varying unlabeled ligand concentrations as described in the Methods section. Left, competition for specific ^{125}I - α -CtxMII binding in the superficial gray of the superior colliculus by epibatidine (\square , $\text{IC}_{50} = 103 \pm 41$ pM, $n_H = -0.87 \pm 0.09$), cytosine (\bullet , $\text{IC}_{50} = 18 \pm 8$ nM, $n_H = -0.94 \pm 0.12$), and (–)-nicotine (\triangle , $\text{IC}_{50} = 481 \pm 54$ nM, $n_H = -1.47 \pm 0.20$). Right, competition for specific ^{125}I - α -CtxMII binding in the striatum by epibatidine (\square , $\text{IC}_{50} = 112 \pm 38$ pM, $n_H = -0.80 \pm 0.23$), cytosine (\bullet , $\text{IC}_{50} = 23 \pm 9$ nM, $n_H = -1.12 \pm 0.17$), and (–)-nicotine (\triangle , $\text{IC}_{50} = 348 \pm 84$ nM, $n_H = -1.05 \pm 0.24$). α -Bgt (1 μM) did not displace ^{125}I - α -CtxMII binding in either region. Nonspecific binding was determined in the presence of 1 μM unlabeled epibatidine. Each point and value is the mean \pm S.E.M. of four or five separate determinations. Data were fitted to a one-site Hill inhibition equation (see *Experimental Procedures*).

of [^3H]epibatidine binding (one-way ANOVA; $F(6,13) = 2.64$, $P > .05$). Displacement of [^3H]epibatidine binding to superior colliculus membranes by cytosine and α -CtxMII is shown in Fig. 4, right, whereas the regional distribution of total, cy-

tisine-resistant, and α -CtxMII-sensitive specific [^3H]epibatidine binding is summarized in Table 2.

[^3H]Epibatidine Autoradiography, Mouse Brain Sections. Competition binding experiments demonstrated that α -CtxMII was able to potently displace a fraction of [^3H]epibatidine binding to mouse brain membrane preparations. In addition, the same data showed that the density of α -CtxMII-sensitive [^3H]epibatidine binding sites varied widely among regions. Because the proportion of α -CtxMII-sensitive [^3H]epibatidine binding never exceeded that of the cytosine-resistant population (and in many regions was lower), it seemed possible that α -CtxMII was selectively interacting with a subpopulation of cytosine-resistant [^3H]epibatidine binding nAChRs, as suggested by the earlier ^{125}I - α -CtxMII autoradiography experiments. Crude regional dissection of the mouse brain provided limited anatomical resolution, so the relationship between cytosine-resistant and α -CtxMII-sensitive [^3H]epibatidine binding sites was explored using an autoradiographic approach. Inhibition binding experiments conducted using filtration binding indicated that 100 nM cytosine would essentially eliminate [^3H]epibatidine binding to cytosine-sensitive sites (at a [^3H]epibatidine concentration of 500 pM) but would leave the cytosine-resistant [^3H]epibatidine binding largely unaffected (Fig. 4, right). For this reason, cytosine-resistant [^3H]epibatidine binding was visualized using 500 pM [^3H]epibatidine, in combination with 100 nM cytosine. The same series of experiments also suggested that 50 nM α -CtxMII would be sufficient to displace α -CtxMII-sensitive [^3H]epibatidine binding, without affecting binding to other types of [^3H]epibatidine binding sites.

[^3H]Epibatidine proved to be an excellent autoradiographic probe, producing nonspecific binding that was indistinguishable from the film background. As would be anticipated from

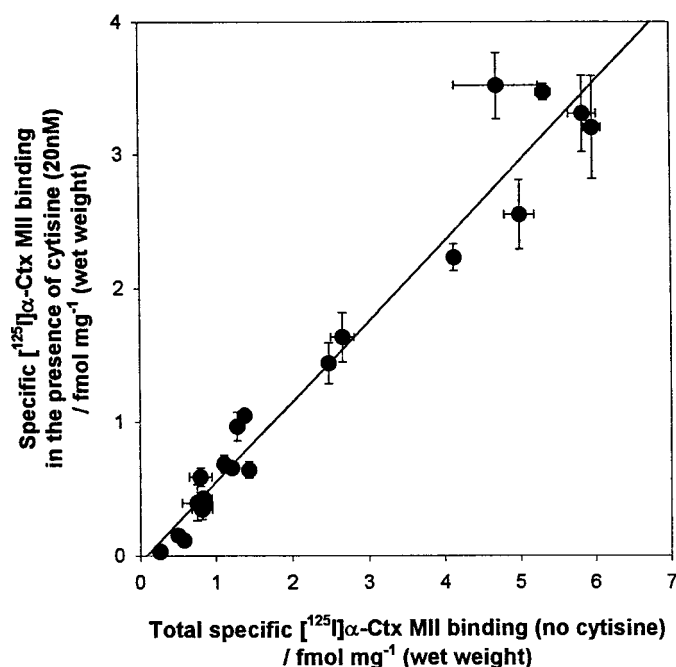


Fig. 3. Regional comparison of mouse brain-specific ^{125}I - α -CtxMII binding in the presence and absence of cytosine (20 nM). Levels of specific ^{125}I - α -CtxMII (0.5 nM) binding with and without the addition of 20 nM cytosine were determined by quantitative autoradiography and compared in 23 different brain regions. Each point is the mean \pm S.E.M. of data gathered from three different animals. Linear regression showed $r = 0.96$, slope = 0.600.

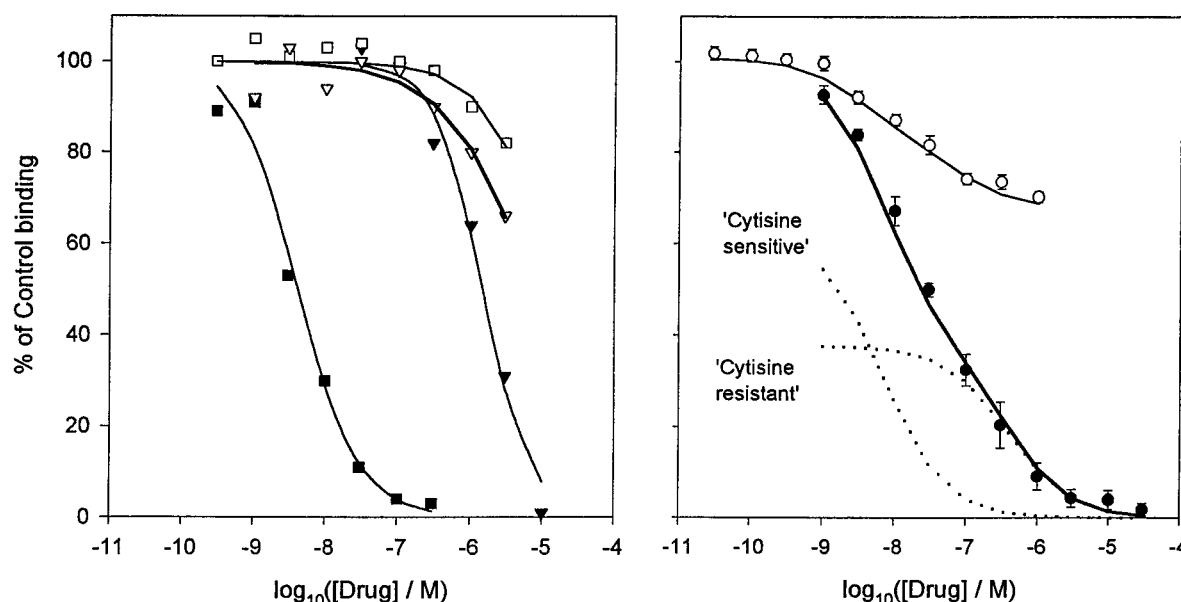


Fig. 4. Competition by cytosine and α -CtxMII for (—) [^3H]nicotine, ^{125}I - α -Bgt and [^3H]epibatidine binding sites in mouse brain membranes. Left, mouse whole brain particulate fractions were incubated at 22°C with (—) [^3H]nicotine (20 nM, 30 min; squares) or ^{125}I - α -Bgt (1 nM, 5 h; triangles) in the presence of unlabeled cytosine (1 nM - 10 μM ; filled symbols) or α -CtxMII (1 nM - 10 μM ; open symbols). Right, mouse superior colliculus particulate fractions were incubated at 22°C with [^3H]epibatidine (500 pM, 2 h; circles) in the presence of unlabeled cytosine (1 nM - 30 μM ; filled symbols) or α -CtxMII (30 pM - 3 μM ; open symbols). Nonspecific binding was determined in the presence of 1 mM unlabeled (—) nicotine. Each point represents the mean \pm S.E.M. of three separate determinations. Data were fitted to either one- or two-site (for cytosine inhibition of [^3H]epibatidine binding) Hill inhibition equations (see *Experimental Procedures*). Both cytosine-sensitive and cytosine-resistant components of [^3H]epibatidine binding are illustrated on the right (dotted lines).

Levels of specific ^{125}I - α -CtxMII binding were compared with those of α -CtxMII-sensitive $[^3\text{H}]$ epibatidine binding

The pattern of $\alpha 3$ hybridization is shown in Fig. 5 (right column). Expression of $\alpha 3$ mRNA was restricted to a number of small, well-defined nuclei distributed throughout the brain. By far the highest level of hybridization was detected in the medial habenula (48 pCi/mm^2), the next highest amount (10 pCi/mm^2) being found in the mitral layer of the accessory olfactory bulbs. Where detected, hybridization in other brain regions was much weaker than in these two regions ($3.2\text{--}0.6 \text{ pCi/mm}^2$). The distribution of $\alpha 3$ hybridization is summarized in Table 4. The highest densities of cytosine-resistant [^3H]epibatidine binding were found in the medial habenula and accessory olfactory bulbs, matching the data for $\alpha 3$ mRNA expression. In addition, patterns of $\alpha 3$ hybridization and cytosine-resistant [^3H]epibatidine binding were superimposed in many brain regions (including the dorsal cortex of the inferior colliculus, medial habenula, medial geniculate nucleus, superficial layers of the superior colliculus, and the medial vestibular and prepositus hypoglossal nuclei). In other cases, cytosine-resistant [^3H]epibati-

Particulate fractions were prepared from the indicated brain regions of C57BL/6 mice. The K_d and B_{max} values for each region were calculated from saturation binding experiments, as described under *Experimental Procedures*. Inhibition of [3 H]epibatidine (500 pM) binding by cytosine and α -CtxMII was determined as illustrated in Figure 4. Cytosine inhibition was fitted to a two-site model, with IC_{50} and B_{max} values for the less-sensitive binding site being used to calculate the inhibition binding constant (K_i) and size of the cytosine-resistant [3 H]epibatidine binding site population in each region. A one-site model was used to determine the K_i values and numbers of α -CtxMII sensitive binding sites in each region. The K_i value of α -CtxMII inhibition of [3 H]epibatidine binding did not vary significantly between regions (one-way ANOVA; $F(6,13) = 2.64$, $P > .05$). All values are the mean \pm S.E. of values derived from three to four independent experiments.

Region	Total [³ H]Epibatidine Binding Population	α -CtxMII-Sensitive Population	K_i (α -CtxMII)	% α -CtxMII-Sensitive	Cytisine-Resistant Population	% Cytisine-Resistant
	<i>fmol/mg of protein</i>		<i>nM</i>		<i>fmol/mg of protein</i>	
Superior colliculus	259 \pm 5	67 \pm 6	0.96 \pm 0.3	25.9	84 \pm 14	32.1
Thalamus	226 \pm 5	26 \pm 4	5.3 \pm 2.8	11.5	39.5 \pm 2.6	17.5
Striatum	118 \pm 4	16 \pm 1	0.83 \pm 0.01	13.6	16 \pm 2	13.6
Inferior colliculus	123 \pm 5	15 \pm 1	7.0 \pm 1.3	12.2	39 \pm 1	31.7
Olfactory tubercles	64 \pm 5	14 \pm 4	1.1 \pm 0.1	21.9	11 \pm 1	17.2
Midbrain	159 \pm 8	13 \pm 1	2.4 \pm 1.0	8.2	14 \pm 1	8.8
Cortex	62 \pm 5	5 \pm 2	1.3 \pm 0.4	8.1	8 \pm 1	12.8
Hippocampus	68 \pm 4	N.D.	N/A	None	4 \pm 1	5.9
Cerebellum	20 \pm 2	N.D.	N/A	None	7 \pm 3	35.0
Olfactory bulbs	26 \pm 8	N.D.	N/A	None	13 \pm 4	50.0
Hindbrain	91 \pm 4	N.D.	N/A	None	18 \pm 9	19.8
Interpeduncular nucleus	973 \pm 154	N.D.	N/A	None	607 \pm 34	62.4

N.D., not detectable; N/A, not applicable.

dine binding was found in nuclei innervated by regions where $\alpha 3$ hybridization can be detected (nAChRs are known to be transported from the ventral tegmental area and substantia nigra to the striatum, frontal cortex, olfactory tubercles, and nucleus accumbens (Schwartz et al., 1984), from the retina to the superior colliculus and the geniculate nuclei (Swanson et al., 1987), and from the medial habenula to the interpeduncular nucleus (Clarke et al., 1986). In addition, cytosine-resistant [3 H]epibatidine binding was detectable on fiber tracts in the fasciculus retroflexus, oculomotor nerve, optic

tract, brachium of the superior colliculus, and supraoptic decussation. Small amounts of $\alpha 3$ hybridization were also found in a number of regions that have no apparent cytosine-resistant [3 H]epibatidine binding, most notably the motor trigeminal nucleus and somatosensory cortex.

Discussion

α -CtXMII, originally isolated from the venom of the predatory cone snail, *C. magus* (Cartier et al., 1996), has been used to investigate the diversity of nicotinic receptor binding sites in mouse brain. The data presented here demonstrate that mouse brain high-affinity epibatidine binding has three components: 1) a component that corresponds to the (–)-[3 H]nicotine- and [3 H]cytosine-binding sites [“cytosine-sensitive” sites, probably the $\alpha 4\beta 2$ subtype (Whiting and Lindstrom, 1987; Flores et al., 1992)] and two sites with lower cytosine affinity (‘‘cytosine-resistant’’ sites); 2) a cytosine-resistant component that displays high affinity for α -CtXMII and that has been visualized here using 125 I- α -CtXMII; 3) a cytosine-resistant component that displays lower affinity for α -CtXMII. This subdivision of mouse brain [3 H]epibatidine-binding nAChRs is illustrated in Fig. 7.

125 I- α -CtXMII Binds to a Novel nAChR Population.

Competition binding experiments demonstrated that α -CtXMII has a low affinity ($K_i > 10 \mu\text{M}$) at mouse brain (–)-[3 H]nicotine and 125 I- α -Bgt binding sites (Fig. 4). This shows that it binds to a novel neuronal nAChR population, distinct from the well-characterized (–)-[3 H]nicotine- and 125 I- α -Bgt-binding sites (corresponding to $\alpha 4\beta 2$ and $\alpha 7$ -containing subtypes in mammalian neurons; Whiting and Lindstrom, 1987; Schoepfer et al., 1990; Flores et al., 1992; Seguela et al., 1992).

Competition binding experiments using 125 I- α -CtXMII in tissue slices yielded K_i values versus 125 I- α -CtXMII for unlabeled epibatidine and cytosine of 80 to 90 pM and 14 to 18 nM, respectively. These values are similar to those reported by Marks et al. (1998) for cytosine-resistant [3 H]epibatidine-binding sites. As shown in Fig. 3, the cytosine sensitivity of 125 I- α -CtXMII-binding sites was constant across brain regions, suggesting that they represent a single population (an alternate but less likely explanation is that multiple sites with a mean cytosine K_i of 20 nM exist in all 125 I- α -CtXMII-binding regions). Additionally, 125 I- α -CtXMII binding sites have no appreciable affinity for α -Bgt (no displacement by 1 μM α -Bgt) and have a relatively low affinity for (–)-nicotine [$K_i = 280\text{--}380 \text{ nM}$, compared with 8.9 nM at the mouse brain $\alpha 4\beta 2$ high affinity [3 H]nicotine binding subtype (Marks et al., 1998)].

Quantitative autoradiographic analysis showed that specific 125 I- α -CtXMII binding occurs in discrete nuclei distributed throughout the mouse brain. The sites’ regional distribution is unlike previously reported nicotinic binding patterns and seemed to represent a subset of the cytosine-resistant [3 H]epibatidine binding sites reported by Marks et al. (1998). The unusual distribution and nicotinic pharmacology of 125 I- α -CtXMII-binding sites confirms that they represent a novel native neuronal nAChR subtype.

α -CtXMII-Binding nAChRs are a Subset of Cytosine-Resistant [3 H]Epibatidine Binding Sites. Confirming the data provided by the 125 I- α -CtXMII autoradiography experiments, α -CtXMII was a potent inhibitor (mean $K_i = 2.7 \text{ nM}$)

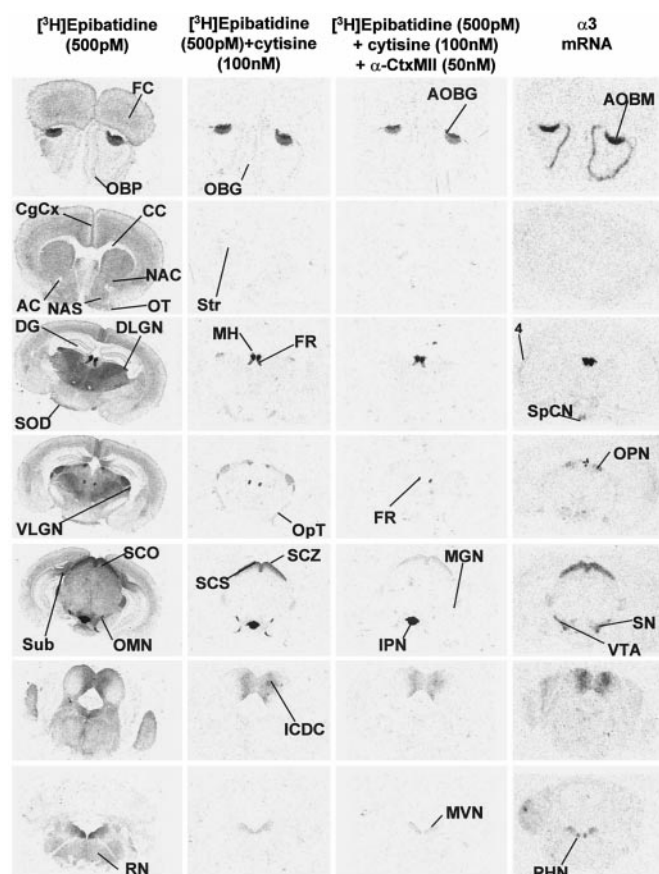


Fig. 5. Autoradiographic comparison of total [3 H]epibatidine (500 pM) binding with that in the presence of cytosine (100 nM) or cytosine (100 nM) + α -CtXMII (50 nM) and $\alpha 3$ mRNA expression in mouse brain. Sections (14 μm) were incubated in the presence of 500 pM [3 H]epibatidine (left column), 500 pM [3 H]epibatidine + 100 nM cytosine (center-left column), 500 pM [3 H]epibatidine + 100 nM cytosine + 50 nM α -CtXMII (center-right column), or were subjected to $\alpha 3$ mRNA in situ hybridization ($\alpha 3$ mRNA; right column) as described under *Experimental Procedures*. Non-specific labeling was indistinguishable from film background. Panels are digital images of autoradiograms. The sections in each row are adjacent. Abbreviations for the indicated brain regions are: 4, somatosensory cortex layer four; AC, anterior commissure; AOBG, accessory olfactory bulb glomerular layer; AOBM, accessory olfactory bulb mitral layer; CC, corpus callosum; CgCx, cingulate cortex; DG, dentate gyrus; DLGN, dorsolateral geniculate nucleus; FR, fasciculus retroflexus; FC, frontal cortex; ICDC, inferior colliculus, dorsal cortex; IPN, interpeduncular nucleus; LH, lateral habenula; MGN, medial geniculate nucleus; MH, medial habenula; MVN, medial vestibular nucleus; NAC, nucleus accumbens core; NAS, nucleus accumbens shell; OBG, olfactory bulb glomerular layer; OBP, olfactory bulb internal plexiform layer; OMN, oculomotor nerve (or root); OPN, olivary pretectal nucleus; Opt, optic tract; OT, olfactory tubercle; PHN, prepositus hypoglossal nucleus; RN, reticular nuclei; SCO, superior colliculus, optic nerve layer; SCS, superior colliculus, superficial gray; SCZ, superior colliculus, zonal layer; SN, substantia nigra; SOD, supraoptic decussation; SpCN, suprachiasmatic nucleus; Str, striatum; Sub, subiculum; VLGN, ventrolateral geniculate nucleus; VTA, ventral tegmental area.

Quantitative autoradiography showed that the majority of [³H]epibatidine binding sites in mouse brain were highly cytosine-sensitive (Fig. 5, column 1 versus column 2). The distribution of the cytosine-sensitive sites mirrored that reported previously (Marks et al., 1998) and corresponded to the distribution of high affinity nicotine or cytosine binding (Perry and Kellar, 1995; Marks et al., 1998). The autoradiograms demonstrated that these sites were confined to a lim-

Regional distribution of specific [³H]epibatidine binding [total, + cytisine (100 nM), and + cytisine (100 nM) + α-CtxMII (50 nM)]

Sections (14 μ m) were incubated in the presence of 500 pM [3 H]epibatidine (total [3 H]epibatidine), 500 pM [3 H]epibatidine + 100 nM cytosine or 500 pM [3 H]epibatidine + 100 nM cytosine + 50 nM α -CtXmII. Binding of [3 H]epibatidine was detected by exposing β max 3 H-sensitive film to the sections, followed by digital densitometry of the resultant autoradiograms. Specific binding was calculated by reference to 3 H-containing tissue paste standards, and is presented in terms of femtomoles of [3 H]epibatidine bound per milligram of tissue weight. The amount of α -CtXmII-sensitive [3 H]epibatidine binding in each region was calculated as the difference between the [3 H]epibatidine binding in the presence of cytosine (100 nM) alone and that with both cytosine (100 nM) and α -CtXmII (50 nM) present. Six individual measurements were made from each region, under each condition in each mouse where possible. Each value represents the mean \pm S.E. of binding measured in three different mice.

	Total [³ H]Epibatidine Binding	[³ H]Epibatidine Binding with Cytisine (100 nM)	% Total Binding in Presence of Cytisine	[³ H]Epibatidine Binding with Cy- tisine (100 nM) + α-CtxMII-(50 nM)	% Total Binding in Presence of Cytisine + α-CtxMII	α-CtxMII (50 nM) Sensitive [³ H]Epi- batidine Binding	% α-CtxMII- Resistant (of Cytisine- Resistant)
	fmol/mg protein wet wt.			fmol/mg protein wet wt.		fmol/mg protein wet wt.	
Telencephalon							
Neocortex							
Cingulate cortex	18.4 ± 1.8	N.D.	None	N.D.	N/A	None	N/A
Frontal cortex	16.0 ± 1.6	0.2 ± 0.1	1.0	N.D.	0None	0.2 ± 0.1	100
Olfactory tubercles	13.7 ± 0.8	2.7 ± 0.3	19.7	0.5 ± 0.3	18.5	2.2 ± 0.01	81.5
Olfactory bulbs							
Accessory olfactory bulbs, glomerular layer	82.2 ± 8.7	66.7 ± 2.7	81.1	48.6 ± 1.7	72.8	18.1 ± 2.5	27.2
Accessory olfactory bulbs, mitral cell layer	37.6 ± 2.7	30.1 ± 0.9	80.0	21.5 ± 1.8	71.6	8.5 ± 1.3	28.4
Olfactory bulbs, glomerular layer	5.7 ± 0.5	2.2 ± 0.4	38.8	2.0 ± 0.3	88.9	0.2 ± 0.1	11.1
Olfactory bulbs, internal plexiform layer	6.6 ± 0.5	2.1 ± 0.3	31.3	1.8 ± 0.3	85.1	0.3 ± 0.1	14.9
Basal ganglia							
Nucleus accumbens, core	15.7 ± 0.8	2.3 ± 0.3	14.4	0.6 ± 0.3	25.9	1.7 ± 0.1	74.1
Nucleus accumbens, shell	13.4 ± 0.5	1.4 ± 0.4	10.7	0.4 ± 0.2	26.8	1.0 ± 0.2	73.2
Striatum	17.2 ± 0.7	1.4 ± 0.3	8.0	0.4 ± 0.2	30.0	1.0 ± 0.1	70.0
Substantia nigra	29.8 ± 1.7	3.3 ± 0.1	10.9	1.8 ± 0.9	56.3	1.4 ± 0.9	43.8
Ventral tegmental area	32.3 ± 2.3	3.5 ± 0.4	10.9	1.1 ± 0.5	33.1	2.3 ± 0.9	66.9
Septum							
Lambdoid septal zone	12.7 ± 0.3	1.2 ± 0.2	9.1	0.2 ± 0.1	18.7	0.8 ± 0.2	81.3
Hippocampus							
Dentate gyrus	14.1 ± 0.8	N.D.	None	N.D.	N/A	None	N/A
Subiculum	43.5 ± 1.3	0.7 ± 0.3	1.6	N.D.	None	0.7 ± 0.3	100
Diencephalon							
Thalamus							
Anteroventral thalamic nucleus	71.7 ± 4.0	4.5 ± 0.4	6.3	3.7 ± 0.1	82.5	0.8 ± 0.5	17.5
Metathalamus							
Dorsolateral geniculate nucleus	69.4 ± 6.1	13.3 ± 0.6	19.3	3.1 ± 0.3	23.6	10.2 ± 0.6	76.4
Medial geniculate nucleus	39.3 ± 2.2	6.6 ± 0.8	16.7	3.5 ± 0.7	53.9	3.1 ± 0.5	46.1
Olivary pretectal nucleus	79.0 ± 6.1	21.0 ± 1.1	26.6	3.4 ± 0.2	16.2	17.6 ± 0.9	83.8
Ventrolateral geniculate nucleus	59.4 ± 5.1	15.5 ± 0.3	26.0	3.1 ± 0.2	20.2	12.3 ± 0.5	79.8
Epi- and subthalamus							
Lateral habenula	69.6 ± 7.0	41.8 ± 5.4	60.0	31.9 ± 1.7	76.4	9.9 ± 3.8	23.6
Medial habenula	380.2 ± 14.5	251.5 ± 20.7	66.1	214.3 ± 4.0	85.2	37.2 ± 16.9	14.8
Mesencephalon							
Inferior colliculus, dorsal cortex	28.3 ± 0.1	10.8 ± 1.0	38.3	7.6 ± 0.6	70.5	3.2 ± 0.6	29.5
Interpeduncular nucleus	308.1 ± 8.9	185.1 ± 15.2	60.1	139.9 ± 4.9	75.6	45.2 ± 10.3	24.4
Superior colliculus, optic nerve layer	28.4 ± 1.0	3.1 ± 0.42	10.9	1.4 ± 0.3	44.6	1.7 ± 0.4	55.4
Superior colliculus, superficial grey	49.1 ± 1.9	16.3 ± 2.0	33.2	3.9 ± 0.5	24.2	12.3 ± 2.2	75.8
Superior colliculus, zonal layer	70.9 ± 2.6	33.8 ± 1.9	47.7	7.4 ± 0.3	22.0	26.4 ± 2.0	78.0
Pons							
Medial vestibular nucleus	23.1 ± 1.3	7.4 ± 0.6	31.9	5.1 ± 0.8	68.8	2.3 ± 0.2	31.2
Prepositus hypoglossal nucleus	25.2 ± 1.7	8.2 ± 1.0	32.5	6.3 ± 0.8	76.8	1.9 ± 0.4	23.2
Pontine nuclei	22.3 ± 2.1	N.D.	None	N.D.	N/A	None	N/A
Reticular nuclei	8.1 ± 0.3	0.3 ± 0.2	3.7	N.D.	None	0.3 ± 0.2	100
Fiber tracts							
Anterior commissure, anterior	2.7 ± 0.6	N.D.	None	N.D.	N/A	None	N/A
Corpus callosum	2.4 ± 0.2	N.D.	None	N.D.	N/A	None	N/A
Fasciculus retroflexus	98.6 ± 7.7	64.8 ± 4.3	65.7	48.4 ± 7.0	74.7	16.4 ± 10.2	25.3
Oculomotor nerve (or root)	74.6 ± 6.4	43.4 ± 4.4	58.2	6.6 ± 0.5	15.2	36.8 ± 4.0	84.8
Optic tract/brachium superior colliculus	15.7 ± 1.2	4.8 ± 0.5	30.6	1.0 ± 0.4	21.8	3.7 ± 0.4	78.2
Supraoptic decussation	17.5 ± 1.9	5.5 ± 0.5	31.3	0.8 ± 0.2	14.7	4.7 ± 0.3	85.3

N.D., not detectable; N/A, not applicable.

ited set of nuclei, scattered throughout the brain. Some (but not all) cytosine-resistant [^3H]epibatidine binding was highly α -CtxMII-sensitive (Fig. 5, column 3 versus column 2). The ability of unlabeled α -CtxMII (50 nM) to selectively displace [^3H]epibatidine from some cytosine-resistant populations, but not others, further confirms that α -CtxMII interacts only with a subset of cytosine-resistant [^3H]epibatidine-binding sites.

The regional densities of ^{125}I - α -CtxMII-binding sites and α -CtxMII-sensitive [^3H]epibatidine-binding sites were compared directly (Fig. 6). The distributions of specific ^{125}I - α -CtxMII and α -CtxMII-sensitive [^3H]epibatidine-binding sites largely coincided, but numbers of ^{125}I - α -CtxMII-binding sites were consistently lower than those α -CtxMII-sensitive [^3H]epibatidine-binding sites. This occurred because a saturating [^3H]epibatidine concentration was used, whereas [^{125}I] α -CtxMII was used at a concentration below its K_d value. However, in six regions more α -CtxMII-sensitive [^3H]epibatidine binding was found than would be predicted from the density of ^{125}I - α -CtxMII binding sites (Fig. 6). This finding needs to be interpreted with some caution: all of these regions contained high levels of [^3H]epibatidine binding, and the additional α -CtxMII-sensitive [^3H]epibatidine binding sites represent a small portion of total binding. However, if the discrepancy is real, it may represent evidence for a sec-

ond α -CtxMII-sensitive [^3H]epibatidine-binding nAChR population in a small number of nuclei. A relatively high unlabeled α -CtxMII concentration (50 nM) was used to displace [^3H]epibatidine binding, so these putative sites could have a relatively low affinity for α -CtxMII (making it undetectable by direct binding of 0.5 nM ^{125}I - α -CtxMII). Thus, although α -CtxMII displays good selectivity for its primary site of action versus [^3H]nicotine and ^{125}I - α -Bgt binding sites, high concentrations may distinguish less well between subtypes of cytosine-resistant [^3H]epibatidine binding. As a result, it is uncertain whether ^{125}I - α -CtxMII binding to the medial habenula/interpeduncular nucleus tract reflects faint cross-labeling to the high density of other cytosine-resistant [^3H]epibatidine binding sites, or the expression of a minor ^{125}I - α -CtxMII binding population.

Physiological Relevance of α -CtxMII-Binding nAChRs. Although α -CtxMII binding nAChRs are relatively rare, they probably exert important physiological effects. α -CtxMII inhibits a component of nicotine-evoked mouse striatal synaptosomal [^3H]dopamine release with an IC_{50} value of 2 nM (Grady et al., 1997), similar to the binding affinity reported here (K_i versus [^3H]epibatidine = 2.7 nM). Similar IC_{50} values have also been reported for α -CtxMII inhibition

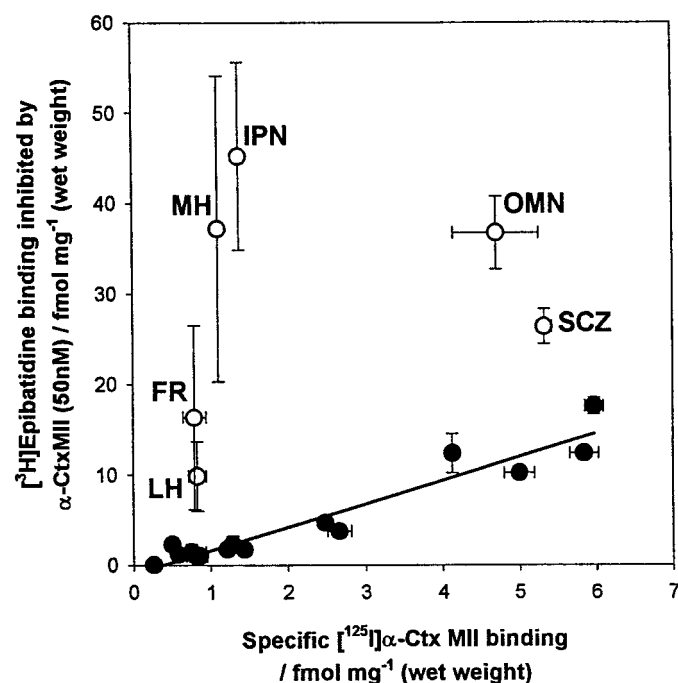


Fig. 6. Comparison of specific ^{125}I - α -CtxMII binding and [^3H]epibatidine binding sensitive to α -CtxMII (50 nM) in mouse brain. Amounts of specific ^{125}I - α -CtxMII (0.5 nM) binding were determined in 23 different brain regions by quantitative autoradiography. The amount of α -CtxMII-sensitive [^3H]epibatidine binding was quantified in the same regions as the difference between [^3H]epibatidine binding in the presence of cytosine (100 nM) and cytosine (100 nM) + α -CtxMII (50 nM). Each point is the mean \pm S.E.M. of data gathered from three different animals. In regions with small or zero cytosine- + α -CtxMII-resistant [^3H]epibatidine binding populations (filled symbols), linear regression showed $r = 0.96$, slope = 2.6. Those regions containing large amounts of α -CtxMII-resistant [^3H]epibatidine binding (open symbols) deviated markedly above the regression line. Abbreviations for the indicated brain regions are: FR, fasciculus retroflexus; IPN, interpeduncular nucleus; LH, lateral habenula; MH, medial habenula; OMN, oculomotor nerve (or root); SCZ, superior colliculus zonal layer.

TABLE 4

Regional quantification of $\alpha 3$ mRNA

Data show in situ hybridization of ^{35}S -cRNA for $\alpha 3$ in mouse brain regions. Hybridization was performed as described under *Experimental Procedures*. Hybridized probe was detected by exposing β max film to the sections, followed by digital densitometry of the autoradiographic images. Specific hybridization was quantified by reference to dot-blotted samples of ^{35}S -cRNA probe of known activity and is presented in terms of pCi/mm^2 . Where possible, measurements were made six times in each region, in each mouse. Values are the mean \pm S.E. of binding measured in three different mice.

	$\alpha 3$ Hybridization Signal/ pCi/mm^2
Telencephalon	
Neocortex	
Somatosensory cortex, layer IV	2.6 ± 0.4
Olfactory bulbs	
Accessory olfactory bulbs, mitral cell layer	10.0 ± 0.3
Olfactory bulbs, internal plexiform layer	2.4 ± 0.2
Basal ganglia	
Substantia nigra	3.2 ± 0.2
Ventral tegmental area	1.4 ± 0.1
Diencephalon	
Thalamus	
Anteroventral thalamic nucleus	1.5 ± 0.2
Mediodorsal thalamic nucleus	2.0 ± 0.6
Parataenial thalamic nucleus	1.8 ± 0.4
Rostral interstitial nucleus	0.7 ± 0.2
Metathalamus	
Medial geniculate nucleus	2.2 ± 0.3
Olivary pretectal nucleus	1.4 ± 0.1
Epi- and subthalamus	
Medial habenula	47.8 ± 7.7
Parasubthalamus	0.6 ± 0.2
Hypothalamus	
Arcuate hypothalamic nucleus	1.9 ± 0.4
Medial mamillary nucleus	0.8 ± 0.1
Suprachiasmatic nucleus	1.7 ± 0.3
Mesencephalon	
Central grey, α	0.8 ± 0.2
Central grey, β	2.4 ± 0.3
Inferior colliculus, dorsal cortex	1.8 ± 0.1
Interpeduncular nucleus	1.6 ± 0.3
Superior colliculus, superficial layers	2.8 ± 0.1
Pons	
Locus ceruleus	0.7 ± 0.1
Medial vestibular nucleus	1.2 ± 0.2
Motor trigeminal nucleus	2.9 ± 0.2
Prepositus hypoglossal nucleus	1.4 ± 0.4

of functional responses in rat and avian preparations (Kulak et al., 1997; Ullian et al., 1997; Kaiser et al., 1998). The similar affinities for binding and functional measures strongly imply a competitive mode of antagonism for α -CtxMII in these preparations. Although less than 15% of [3 H]epibatidine binding in mouse striatum is α -CtxMII-sensitive, about 50% of nicotine-evoked [3 H]dopamine release is inhibited by α -CtxMII (Grady et al., 1997). This disproportionate α -CtxMII sensitivity may arise because of preferential α -CtxMII-sensitive nAChR location on dopaminergic termini.

Identity of α -CtxMII-Binding nAChRs. Marks et al. (1998) noted a resemblance between the patterns of $\alpha 3$ nAChR subunit mRNA expression and cytosine-resistant [3 H]epibatidine binding in mouse brain. Direct comparison of the two measures in brain slices prepared from the same animals reinforced this initial impression (Fig. 5). The highest densities of both $\alpha 3$ hybridization and cytosine-resistant [3 H]epibatidine binding are found in the medial habenula and accessory olfactory bulbs, and in many nuclei $\alpha 3$ hybridization and cytosine-resistant binding are completely superimposed. The majority of the remaining cytosine-resistant binding sites are found in regions known to be innervated by $\alpha 3$ mRNA expressing regions. Cytosine-resistant nAChRs may also contain other subunits: for instance, the $\alpha 6$ subunit is extensively coexpressed with $\alpha 3$ in rat brain (LeNovere et al., 1996), and Vailati et al. (1999) have shown that $\alpha 6$ -

containing nAChRs also represent a class of cytosine-resistant nAChRs (K_i values versus [3 H]epibatidine binding for cytosine and epibatidine = 11 nM and 20 pM, respectively). nAChRs containing $\alpha 6$ also bind α -CtxMII with moderate affinity (K_i versus [3 H]epibatidine = 66 nM), making them possible candidates for the putative second, lower affinity α -CtxMII-binding site discussed previously.

Further evidence supports the involvement of $\alpha 3$ subunits in α -CtxMII-binding sites. α -CtxMII is a selective antagonist of heterologously expressed rat $\alpha 3\beta 2$ nAChRs in *X. laevis* oocytes (Cartier et al., 1996) and human $\alpha 3\beta 2$ in human embryonic kidney 293 cells (Crona et al., 1997). Parker et al. (1998) showed that *X. laevis* oocyte-expressed rat $\alpha 3\beta 2$ nAChRs bind [3 H]epibatidine with high affinity and have a low cytosine affinity. Furthermore, the pattern of α -CtxMII-sensitive [3 H]epibatidine binding was similar to that described by Schultz et al. (1991) for α -Bgt-insensitive [125 I]neuronal Bgt binding. Although neuronal Bgt exhibits complex kinetics of interaction at a variety of nAChR subtypes (Papke et al., 1993), it is a comparatively selective antagonist of heterologously expressed $\alpha 3\beta 2$ nAChRs (Luetje et al., 1990). Together, these data suggest that mouse brain α -CtxMII-sensitive [3 H]epibatidine binding occurs at receptors containing $\alpha 3$ and $\beta 2$ subunits.

The remaining cytosine-resistant [3 H]epibatidine binding was found in regions expressing high levels of both the $\alpha 3$ and $\beta 4$ nAChR subunits (Dinelly-Miller and Patrick, 1992). Again, Parker et al. (1998) report that $\alpha 3\beta 4$ -subtype nAChRs bind [3 H]epibatidine with detectable affinity and exhibit low cytosine affinity. Thus, it is possible that in mouse brain, the sites less sensitive to α -CtxMII cytosine-resistant binding are a combination of (minimally) $\alpha 3$ and $\beta 4$ subunits.

In conclusion, this study shows that α -CtxMII is a potent, selective, competitive antagonist at a novel population of mouse brain nAChRs. [125 I]- α -CtxMII was used in this study to quantify the high α -CtxMII affinity population and map its distribution. Selective inhibition with cytosine and α -CtxMII revealed high affinity [3 H]epibatidine binding at three nAChR pharmacological subtypes. The largest [3 H]epibatidine binding population was highly cytosine-sensitive and corresponds to the high affinity (-)-[3 H]nicotine binding, $\alpha 4\beta 2$ nAChR subtype. Cytosine-resistant sites are likely to be $\alpha 3$ subunit-containing and exhibited differential α -CtxMII sensitivity that may be caused by differing β subunit composition.

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Mouse CNS subunits: $\alpha 2$ - $\alpha 7$, $\beta 2$ - $\beta 4$.

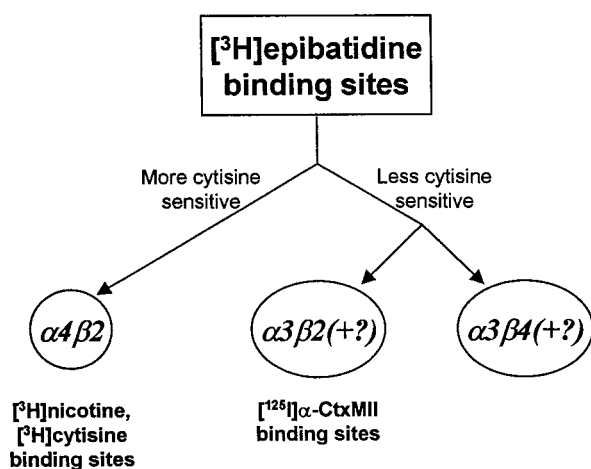


Fig. 7. Summary of [3 H]epibatidine binding sites in mouse brain. Using [3 H]epibatidine in combination with cytosine reveals that [3 H]epibatidine binds specifically and with high affinity in mouse brain to both the well-established (-)-[3 H]nicotine and [3 H]cytosine binding site of $\alpha 4\beta 2$ composition (Whiting and Lindstrom, 1987; Flores et al., 1992), and a smaller population of nAChRs that are less cytosine sensitive (Marks et al., 1998). This allows the use of cytosine to selectively remove [3 H]epibatidine binding at the (-)-[3 H]nicotine binding site, revealing sites less sensitive to cytosine binding. The [3 H]epibatidine sites that are less cytosine-sensitive are likely to contain the $\alpha 3$ nAChR subunit, and may be divided into two groups according to their α -CtxMII sensitivity. [125 I]- α -CtxMII may be used to identify and quantify the population that is more α -CtxMII-sensitive and was used in this study to map the distribution of mouse brain high-affinity α -CtxMII binding nAChRs. The population that is more α -CtxMII-sensitive is likely to contain (minimally) $\alpha 3$ and $\beta 2$ nAChR subunits, whereas the subpopulation that is less α -CtxMII-sensitive can tentatively be assigned a minimal $\alpha 3$ and $\beta 4$ nAChR subunit composition.

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Send reprint requests to: Dr. A.C. Collins, Institute for Behavioural Genetics, University of Colorado, Campus Box 447, Boulder, CO 80303-0447.
