Lobelane Inhibits Methamphetamine-Evoked Dopamine Release via Inhibition of the Vesicular Monoamine Transporter-2*

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ABSTRACT

Lobelane is currently being evaluated in clinical trials as a methamphetamine abuse treatment. Lobelane interacts with nicotinic receptor subtypes, dopamine transporters (DATs), and vesicular monoamine transporters (VMAT2s). Methamphetamine inhibits VMAT2 and promotes dopamine (DA) release from synaptic vesicles, resulting ultimately in increased extracellular DA. The present study generated structure-activity relationships by defunctionalizing the lobeline molecule and determining effects on [3H]dihydrotetrabenazine binding, inhibition of [3H]DA uptake into striatal synaptic vesicles and synaptosomes, the mechanism of VMAT2 inhibition, and inhibition of methamphetamine-evoked DA release. Compared with lobeline, the analogs exhibited greater potency inhibiting DA transporter (DAT) function. Saturated analogs, lobelane and norlobelane, exhibited high potency (K_i = 45 nM) inhibiting vesicular [3H]DA uptake, and lobelane competitively inhibited VMAT2 function. Lobeline and lobelane exhibited 67- and 35-fold greater potency, respectively, in inhibiting VMAT2 function compared to DAT function. Lobelane potently decreased (IC_{so} = 0.65 μM; I_{max} = 73%) methamphetamine-evoked DA overflow, and with a greater maximal effect compared with lobeline (IC_{so} = 0.42 μM, I_{max} = 56.1%). These results provide support for VMAT2 as a target for inhibition of methamphetamine effects. Both trans-isomers and demethylated analogs of lobelane had reduced or unaltered potency inhibiting VMAT2 function and lower maximal inhibition of methamphetamine-evoked DA release compared with lobelane. Thus, defunctionalization, cis-stereochemistry of the side chains, and presence of the piperidino N-methyl are structural features that afford greatest inhibition of methamphetamine-evoked DA release and enhancement of selectivity for VMAT2. The current results reveal that lobelane, a selective VMAT2 inhibitor, inhibits methamphetamine-evoked DA release and is a promising lead for the development of a pharmacotherapeutic for methamphetamine abuse.

Treatment center admissions involving methamphetamine abuse have more than doubled between 1995 and 2005 [Substance Abuse and Mental Health Services Administration, 2008], indicating the need for effective pharmacotherapies. There are currently no approved medications to treat methamphetamine abuse. Methamphetamine abuse liability stems from its reinforcing and rewarding effects demonstrated by use of animal models of intravenous methamphetamine self-administration and conditioned place preference, respectively (Yokel and Pickens, 1973; Hart et al., 2001, Xu et al., 2008). Brain dopaminergic pathways are activated by methamphetamine (Di Chiara and Imperato, 1988; Gold et al., 1989; Wise, 2002). Methamphetamine releases dopamine (DA) from synaptic vesicles into the cytosol via an interaction with the vesicular monoamine transporter (VMAT2) and by disruption of the vesicular proton gradient because of its weak basicity and high lipophilicity (Sulzer and Rayport, 1990; Pifl et al., 1995; Brown et al., 2000, 2001; Fleckenstein et al., 2007). Subsequently, available cytosolic DA is reverse-transported by the DA transporter (DAT) into the extracellular space (Sulzer et al., 1995). In addition, methamphetamine inhibits monoamine oxidase (MAO), preventing DA metabolism into dihydroxyphenylacetic acid (DOPAC) and increasing cytosolic DA (Mantle et al., 1976). VMAT2 is an...
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Fig. 1. Chemical structures of lobeline and its defunctionalized analogs. MTD is a defunctionalized, unsaturated meso-analogue of lobeline. (-)-TTD and (+)-TTD are enantiomers and trans-isomers of MTD. Lobelane is a defunctionalized, saturated meso-analogue of lobeline. (-)-trans-Lobelane and (+)-trans-lobelane are enantiomers and trans-isomers of lobeline. nor-MTD, nor-lobelane, and (±)-nor-lobelane are demethylated analogs of MTD, lobelane, and (±)-TTD, respectively.

Lobelane (Fig. 1), the major alkaloid of Lobelia inflata, attenuates methamphetamine self-administration in rats, but not by acting as a substitute reinforcer (Harrod et al., 2001, 2003), suggesting that it has potential as a low-abuse liability treatment for methamphetamine abuse. In addition, lobelane attenuates both methamphetamine-induced hyperactivity in locomotor activity assays and methamphetamine toxicity through its interaction at nAChRs and/or neurotransmitter transporters.

With respect to nAChRs, initial generation of structure-activity relationships (SARs) revealed that both C2 and C6 piperidino ring substituents of lobelane are critical for interaction with α4β2* nAChRs (* indicates putative nAChR subtype assignment), as evaluated by inhibition of [3H]cytisine binding to rat cortical membranes and stimulation of [3H]Rb+ efflux from striatal synaptosomes (Terry et al., 1998). Lobeline fragments, containing either the phenylcarboxymethyl-ene or phenylhydroxyethylene moiety, were less potent than lobelane at α4β2* nAChRs, indicating the importance of these moieties for optimal affinity at α4β2* (Flammia et al., 1999). Chemical defunctionalization (i.e., removal of the hydroxyl and ketone functionalities) of the lobelane molecule was predicted to yield compounds with low affinity at nAChRs. As expected, lobelane and meso-transdiene (MTD; Fig. 1) had little or no affinity for either α4β2* or α7* nAChRs (Miller et al., 2004; Zheng et al., 2005). It is noteworthy that lobelane and MTD exhibited higher affinity for both DAT and VMAT2.

In addition, changing the stereochemistry of the C2, C6 substituents in MTD from cis to trans afforded the isomeric analog, (-)-trans-transdiene [(−)-TTD; Flammia et al., 1999], which provides support for mechanism of inhibition of VMAT2 function was determined for isolated synaptic vesicles and synaptosomes, respectively.

The present study determined the affinity of lobelane and a wider range of defunctionalized analogs for VMAT2 and DAT, i.e., analog-induced inhibition of [3H]DA uptake into isolated synaptic vesicles and synaptosomes, respectively. Mechanism of inhibition of VMAT2 function was determined for the best analog, lobelane, in the series. In addition, analogs were evaluated for inhibition of methamphetamine-evoked endogenous DA overflow from rat striatal slices. SAR trends emerging from these studies are of value in providing new leads in the development of clinical candidates for the treatment of methamphetamine abuse.

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Materials and Methods

Animals. Male Sprague-Dawley rats (200–250 g upon arrival) were purchased from Harlan (Indianapolis, IN). Rats were housed in the Division of Laboratory Animal Resources at the College of Pharmacy at the University of Kentucky (Lexington, KY) and had ad libitum access to food and water. Experimental protocols involving the animals were in accordance with the National Institutes of Health Guide for the Care and Use of Laboratory Animals and were approved by the Institutional Animal Care and Use Committee at the University of Kentucky.

Materials. [3H]DA (specific activity, 28.0 Ci/mmol) was purchased from PerkinElmer Life and Analytical Sciences (Boston, MA). [3H]DBTZ (specific activity, 79.0 Ci/mmol) was a generous gift from Dr. Michael R. Kilbourn (Department of Internal Medicine and Neurology, University of Michigan, Ann Arbor, MI). Bovine serum albumin, EDTA, EGTA, L-(+)-tartaric acid, succrose, magnesium sulfate, polyethyleneimine, adenosine 5'-triphosphate magnesium salt, HEPES, S(+)-nicotine ditartrate (nicotine), 3-hydroxytryptamine (dopamine, DA), DOPAC, d-methamphetamine hydrochloride (methamphetamine), sodium chloride, magnesium sulfate and ascorbate oxidase were purchased from Sigma-Aldrich (St. Louis, MO). α-D-Glucose, L-ascorbic acid, and monobasic potassium phosphate were purchased from Aldrich Chemical Co. (Milwaukee, WI), AnalR-BHD Ltd. (Poole, UK) and Mallinckrodt (Phillipsburg, NJ). Ro4-1284 was obtained from Hoffman-La Roche Ltd. (Basel, Switzerland). Lobeline hemisulfate was purchased from ICN Biomedicals Inc. (Costa Mesa, CA). All other commercial chemicals were purchased from Fisher Scientific Co. (Pittsburgh, PA).

The lobeline analogs, MTD, (−)-TDD, (+)-trans-transdiene ([+]-TDD), lobeline, (−)-trans-lolbolene, (−)-trans-lolbolene, nor-MTD, (±)-nor-TDD, and nor-lolbelane were synthesized according to reported methods (Zheng et al., 2005) and their structures are illustrated in Fig. 1. The structures of the lobeline analogs were verified by 1H and 13C NMR spectroscopy, mass spectrometry, and, in some instances, X-ray crystallography.

Synaptosomal [3H]DA Uptake Assay. Inhibition of [3H]DA uptake was conducted by use of a preparation of isolated synaptic vesicles as described previously (Teng et al., 1997). In brief, rat striata were homogenized with 10 up-and-down strokes of a Teflon pestle homogenizer (clearance = 0.003 inch) in 14 ml of 0.32 M sucrose solution. Homogenates were centrifuged (2000g for 10 min at 4°C), and the resulting supernatants were centrifuged again (10,000g for 30 min at 4°C). Pellets were resuspended in 2 ml of 0.32 M sucrose solution and subjected to osmotic shock by adding 7 ml of ice-cold water to the preparation, followed by the immediate restoration of osmolarity by adding 900 μl of 0.25 M HEPES buffer and 900 μl of 1.0 M potassium tartrate solution. Samples were centrifuged (20,000g for 20 min at 4°C), and the resulting supernatants were centrifuged again (55,000g for 1 h at 4°C), followed by the addition of 100 μl of 10 mM MgSO4, 100 μl of 0.25 M HEPES, and 100 μl of 1.0 M potassium tartrate suspension before the final centrifugation (100,000g for 45 min at 4°C). Final pellets were resuspended in 2.4 ml of assay buffer (125 mM NaCl, 5 mM KCl, 1.5 mM MgSO4, 1.25 mM CaCl2, 1.5 mM KH2PO4, 10 mM d-glucose, 25 mM HEPES, 0.1 mM EDTA, 0.1 mM pargyline, 0.1 mM ascorbic acid, saturated with 95% O2/5% CO2, pH 7.4). The assay was performed in duplicate in a total volume of 500 μl. Aliquots of the vesicular suspension (25 μl) were added to tubes containing assay buffer and various concentrations of analog (100 μM to 1 nM), and incubated at 34°C for 5 min. Nonspecific uptake was determined in the presence of nomifensine (0.1 μM). Samples were placed on ice, and 50 μl of 0.1 μM [3H]DA was added to each tube, after which accumulation was permitted to proceed for 10 min at 34°C. The reaction was terminated by the addition of 3 ml of ice-cold assay buffer and subsequent filtration. Radioactivity retained by the GF/B filters (presoaked for 2 h in assay buffer) was determined by liquid β-scintillation spectrometry.

[3H]DBTZ Binding Assay. Lobeline- and analog-induced inhibition of [3H]DBTZ binding was determined by use of modifications of a method described previously (Teng et al., 1997). Rat whole brain (excluding cerebellum) or striatum was homogenized in 20 ml of ice-cold 0.22 M sucrose solution with seven up-and-down strokes of a Teflon pestle homogenizer (clearance = 0.003 inch). Homogenates were centrifuged at 1000g for 12 min at 4°C, and the resulting supernatants were again centrifuged at 22,000g for 10 min at 4°C. Resulting pellets were incubated in 18 ml of ice-cold water for 5 min, and 2 ml of HEPES (25 mM) and potassium tartrate (100 mM) solution were subsequently added. Samples were centrifuged (20,000g for 20 min at 4°C), and 20 μl of MgSO4 (1 mM) solution was then added to the supernatants. Solutions were centrifuged (100,000g for 45 min at 4°C) and pellets resuspended in ice-cold assay buffer (25 mM HEPES, 100 mM potassium tartrate, 5 mM MgSO4, 0.1 mM EDTA, and 0.05 mM EGTA, pH 7.5). Assays were performed in duplicate by use of 96-well plates. Aliquots of vesicular suspension (15 μg of protein in 100 μl) were added to wells containing 5 mM [3H]DBTZ, 50 μl of analog (1 mM to 1 μM), and 50 μl of buffer. Nonspecific binding was determined in the presence of Ro4-1284 (20 μM). Reactions were terminated by filtration (Packard Filtermate harvester; PerkinElmer Life and Analytical Sciences) onto Unifilter-96 GF/B filter plates (presoaked in 0.5% polyethyleneimine). Filters were subsequently washed five times with 350 μl of ice-cold buffer (25 mM HEPES, 100 mM potassium tartrate, 5 mM MgSO4, and 10 mM NaCl, pH 7.5). Filter plates were dried and bottom-sealed, and each well was filled with 40 μl of scintillation cocktail (Microscint 20; PerkinElmer Life and Analytical Sciences). Radioactivity on the filters was determined by liquid β-scintillation spectrometry (TopCount NXT; PerkinElmer Life and Analytical Sciences).

Vesicular [3H]HDA Uptake Assay. Inhibition of [3H]HDA uptake was conducted by use of a preparation of isolated synaptic vesicles as described previously (Teng et al., 1997). In brief, rat striata were homogenized with 10 up-and-down strokes of a Teflon pestle homogenizer (clearance = 0.003 inch) in 14 ml of 0.32 M sucrose solution. Homogenates were centrifuged (2000g for 10 min at 4°C), and the resulting supernatants were centrifuged again (10,000g for 30 min at 4°C). Pellets were resuspended in 2 ml of 0.32 M sucrose solution and subjected to osmotic shock by adding 7 ml of ice-cold water to the preparation, followed by the immediate restoration of osmolarity by adding 900 μl of 0.25 M HEPES buffer and 900 μl of 1.0 M potassium tartrate solution. Samples were centrifuged (20,000g for 20 min at 4°C), and the resulting supernatants were centrifuged again (55,000g for 1 h at 4°C), followed by the addition of 100 μl of 10 mM MgSO4, 100 μl of 0.25 M HEPES, and 100 μl of 1.0 M potassium tartrate suspension before the final centrifugation (100,000g for 45 min at 4°C). Final pellets were resuspended in 2.4 ml of assay buffer (25 mM HEPES, 100 mM potassium tartrate, 50 μM EGTA, 100 μM EDTA, 1.7 mM ascorbic acid, 2 mM ATP-Mg2+, pH 7.4). Aliquots of the vesicular suspension (100 μl) were added to tubes containing assay buffer, various concentrations of analog (0.1 nM to 10 mM) and 0.1 μM [3H]HDA to produce a final volume of 500 μl. Nonspecific uptake was determined in the presence of Ro4-1284 (10 μM). Reactions were terminated by filtration, and radioactivity retained by the filters was determined as described previously.

To determine the mechanism of inhibition of [3H]HDA uptake for lobeline and lobeline, kinetic analyses were performed. Concentrations of lobeline (0.25 μM) and lobeline (0.024 μM) were chosen based on 0.5 × the K values obtained in the [3H]HDA uptake inhibition assays. Experiments were conducted in the absence (control) and presence of analog. Incubations were initiated by the addition of 50 μl of the vesicular suspension to 150 μl of assay buffer, 25 μl of analog, and 25 μl of a range of concentrations of [3H]HDA (0.001–5.0 μM). Nonspecific uptake was determined in the presence of Ro4-1284 (10 μM). After an incubation period of 8 min, [3H]HDA uptake was terminated by filtration, and radioactivity retained by the filters was determined as described previously.

Endogenous DA and DOPAC Overflow Assay. Coronal stratial slices were prepared and superfused as described previously (Teng et al., 1997). Slices were incubated in Krebs buffer (118 mM NaCl, 4.7 mM KCl, 1.2 mM MgCl2, 1.0 mM NaH2PO4, 1.3 mM CaCl2, 11.1 mM α-d-glucose, 25 mM NaHCO3, 0.11 mM L-ascorbic acid, and 0.004 mM EDTA, pH 7.4, saturated with 95% O2/5% CO2) at 34°C for 60 min in a metabolic shaker. Each slice was then transferred to a
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Inhibition of [\textsuperscript{3}H]DA Uptake into Rat Striatal Synap
tosomal Preparations. Lobeline- and analog-mediated inhibi
tion of [\textsuperscript{3}H]DA uptake in synaptosomal preparations as
essed inhibition of DAT function (Fig. 2). With the excep
tion of lobeline (\(I\textsubscript{\text{max}} = 80.7\%\)), all compounds inhibi
t[\textsuperscript{3}H]DA uptake at DAT with a maximal inhibition >95\%. Lobeline was 20-fold more potent in the inhibition of DAT than lobeline (\(K\textsubscript{i} = 1.57\) and 31.6 \(\mu M\), respectively; \(p < 0.05\)). MTD exhibited 40-fold greater potency (\(K\textsubscript{i} = 0.039\) \(\mu M\); \(p < 0.05\)) than lobeline and was equipotent with (+)-TDD (\(K\textsubscript{i} = 0.22\) \(\mu M\)), which had greater potency (\(K\textsubscript{i} = 1.89\) \(\mu M\); \(p < 0.05\)) than (+)-TDD. The (−) and (+)-\textit{trans}-
lobelane enantiomers possessed similar potency (\(K\textsubscript{i} = 1.12\) and 4.12 \(\mu M\), respectively) in inhibition of DAT function, and were not different from lobeline. N-Demethylation of MTD and (−)-TDD resulted in a 5- to 16-fold reduction in potency (\(p < 0.05\)), whereas N-demethylation of lobeline resulted in a 5-fold increase in potency (\(p < 0.05\)). Methamphetamine inhibited (\(K\textsubscript{i} = 0.11\) \(\mu M\)) [\textsuperscript{3}H]DA uptake in the synaptosomal preparation.

Inhibition of [\textsuperscript{3}H]DTBZ Binding. [\textsuperscript{3}H]DTBZ binding assays were conducted using rat whole brain to maximize protein yield. However, vesicular [\textsuperscript{3}H]DA uptake assays that assess VMAT2 function were performed with use of striatal preparations. Because two different tissue preparations were

glass superfusion chamber and superfused at 1 ml/min for 60 min. Three 5-min samples (1 ml collected into 100 \(\mu l\) of 0.1 M perchloric acid) determined basal outflow of DA or DOPAC. Slices were superfused for 30 min in the absence or presence of a single concentration of analog (0.1 nM to 3 \(\mu M\)). Subsequently, methamphetamine (5 \(\mu M\)) was added to the superfusion buffer and slices were superfused for 15 min, followed by superfusion for another 20 min with analog in the absence of methamphetamine. Methamphetamine concentration and exposure time were chosen based on previously reported concentration-response curves (Liang and Rutledge, 1982; Bowyer et al., 1991), and concentrations of lobeline (0.1–3.0 \(\mu M\)) were chosen that did not deplete striatal DA content (Teng et al., 1997) and inhibited amphetamine-evoked DA overflow (Miller et al., 2000). In each experiment, a striatal slice was superfused for 80 min in the absence of analog or methamphetamine, and served as the buffer control condition. Duplicate slices were superfused with methamphetamine in the absence of analog, and served as the methamphetamine control condition.

Ascorbate oxidase (20 \(\mu l\)) was added to each superfusate sample (500 \(\mu l\)) and 100 \(\mu l\) of the resulting solution was injected onto the high-performance liquid chromatography-electrochemical detection unit, which consisted of a model 116 pump, model 508 autosampler (Beckman Coulter, Inc., Fullerton, CA), an ODS ultrasphere C18 reverse-phase 80 \(\times\) 4.6 mm, 3-\(\mu m\) column, and a coulometric-II detector with model 5020 guard cell (+0.60 V) and model 5011 cell (0.9 V, 0.32 V) (ESA, Inc., Chelmsford, MA). The eluent was 0.07 M citrate/0.1 M acetate buffer (pH 4) containing: 175 mg/l octylsulfonic acid-sodium salt, 650 mg/l NaCl, and 7% methanol. Separations were performed at room temperature at a flow rate of 1.5 ml/min, and required 5 to 6 min per sample. Reten
tion times of DA and DOPAC standards were used to identify peaks. Peak heights were used to calculate the detected amounts of analyte based on standard curves. Detection limits for DA and DOPAC were 1 and 2 pg/100 \(\mu l\), respectively.

To determine whether the high concentration of analog altered tissue DA content, slices were homogenized by sonication in 500 \(\mu l\) of perchloric acid immediately after perfusion of striatal slices with 3 \(M\) perchloric acid immediately after perfusion of striatal slices with 3 \(M\) each analog. In each experiment, two slices were perfused in the absence of analog and served as the control condition. The homoge
nate was centrifuged at 30,000g at 4°C for 15 min, and the superna
tant was separated and filtered (0.22 \(\mu M\)). Filtrate (100 \(\mu l\)) was 
processed via high-performance liquid chromatography-electrochemi
cal detection as described previously.

Data Analyses. For inhibition of [\textsuperscript{3}H]DTBZ binding, specific bind
ing was determined by subtracting nonspecific binding from total binding. For inhibition of [\textsuperscript{3}H]DA uptake, specific uptake was deter
ded by subtracting nonspecific uptake from total uptake. Concen
trations of inhibitor that produced 50% inhibition of binding or uptake (\(IC_{50}\) values) were determined from the concentration-effect curves via an iterative curve-fitting program (Prism 4.0; GraphPad Software Inc.). \(IC_{50}\) values were determined by use of an iterative curve-fitting program (Prism 4.0; GraphPad Software Inc.). ANOVAs and post hoc tests were conducted with use of SPSS version 9.0 (SPSS Science, Chicago, IL). Statistical significance was defined as \(p < 0.05\).

Results

Inhibition of [\textsuperscript{3}H]DA Uptake into Rat Striatal Synap
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lobelane enantiomers possessed similar potency (\(K\textsubscript{\text{i}} = 1.12\) and 4.12 \(\mu M\), respectively) in inhibition of DAT function, and were not different from lobeline. N-Demethylation of MTD and (−)-TDD resulted in a 5- to 16-fold reduction in potency (\(p < 0.05\)), whereas N-demethylation of lobeline resulted in a 5-fold increase in potency (\(p < 0.05\)). Methamphetamine inhibited (\(K\textsubscript{\text{i}} = 0.11\) \(\mu M\)) [\textsuperscript{3}H]DA uptake in the synaptosomal preparation.

Inhibition of [\textsuperscript{3}H]DTBZ Binding. [\textsuperscript{3}H]DTBZ binding assays were conducted using rat whole brain to maximize protein yield. However, vesicular [\textsuperscript{3}H]DA uptake assays that assess VMAT2 function were performed with use of striatal preparations. Because two different tissue preparations were
used, the ability of two standard compounds, TBZ and Ro4-1284, to inhibit $[3H]$DTBZ binding in these preparations was compared (Fig. 3). $K_i$ values for both TBZ and Ro4-1284 were not different between whole brain and striatum. Thus, whole brain and striatal preparations provide comparable data in the $[3H]$DTBZ binding assay.

Lobeline and its analogs inhibited $[3H]$DTBZ binding to vesicle membranes obtained from rat whole brain (Fig. 4). With the exception of MTD ($I_{\text{max}} = 51.4\%$) and methamphetamine ($I_{\text{max}} = 25.1\%$), all compounds exhibited a maximal inhibition of $85\%$. Lobeline inhibited ($K_i = 2.04 \mu M$) $[3H]$DTBZ binding to VMAT2, and MTD had a lower affinity ($K_i = 9.88 \mu M$) in comparison ($p < 0.05$). Lobeline had 2-fold greater affinity ($K_i = 0.97 \mu M$; $p < 0.05$) than lobeline at VMAT2. The 2,6-trans-analogs of MTD, (+)-TTD and (-)-TTD, had $K_i$ values of 7.09 and 19.4 $\mu M$, respectively. Methamphetamine had a low affinity ($K_i = 80.1 \mu M$) for the $[3H]$DTBZ binding site.
compounds exhibited a maximal inhibition of [³H]DA uptake of >90%. Lobeline inhibited [³H]DA uptake with a $K_i$ value of 0.47 μM, whereas lobelane and nor-lobelane exhibited a 10-fold increase in potency ($K_i = 0.045$ and 0.044 μM, respectively; $p < 0.05$) compared with lobeline. (+)- and (-)-trans-Lobelane exhibited a 5- and 9-fold reduction in potency ($K_i = 2.22$ and 3.83 μM, respectively; $p < 0.05$) compared with lobeline, and were equipotent with methamphetamine ($K_i = 2.46$ μM). MTD, (-)-TTD, and (+)-TTD all exhibited potencies ($K_i = 0.46, 0.38,$ and 0.72 μM, respectively) not different from each other and not different from lobeline. (+)-nor-TTD exhibited a potency ($K_i = 0.36$ μM) similar to lobeline. In contrast, nor-MTD had the lowest potency ($K_i = 7.99$ μM) for inhibiting [³H]DA uptake into synaptic vesicles.

To elucidate the mechanism of lobeline and lobelane interaction with VMAT2, kinetic analyses of [³H]DA uptake into synaptic vesicles were conducted. Compared with control ($K_m = 0.11 \pm 0.0081$ μM; $V_{max} = 46.7 \pm 7.92$ pmol/min/mg) both lobeline ($K_m = 0.29 \pm 0.064$ μM; $V_{max} = 50.2 \pm 7.80$ pmol/min/mg) and lobelane ($K_m = 0.38 \pm 0.051$ μM; $V_{max} = 57.1 \pm 12.1$ pmol/min/mg) significantly ($p < 0.05$) increased the $K_m$ without altering $V_{max}$, indicating that these compounds inhibit [³H]DA uptake in a competitive manner.

**DA and DOPAC Overflow from Superfused Rat Striatal Slices.** The ability of lobeline and its analogs (0.1–3.0 μM) to evoke DA and DOPAC overflow was assessed during the 30-min period of superfusion before the addition of methamphetamine (5 μM) to the superfusion buffer. At the highest concentration evaluated, lobeline and its analogs did not deplete striatal DA content (Table 1). In addition, these compounds alone did not evoke DA overflow (Table 2). In contrast, each compound alone increased DOPAC overflow (Fig. 6), with the exception of (-)-nor-TTD. Analysis of the effect of lobeline on DOPAC overflow revealed a concentration-dependent effect ($F_{4,16} = 4.14, p < 0.05$). Post hoc analysis indicated that 3 μM lobeline increased DOPAC overflow (Fig. 6, top). Analysis of DOPAC overflow also revealed a concentration effect for MTD ($F_{4,13} = 19.94, p < 0.001$), (-)-TTD ($F_{4,14} = 147.61, p < 0.001$), and (+)-TTD ($F_{4,20} = 8.59, p < 0.001$). The lowest concentration of MTD

**TABLE 1**
DA content in striatal slices exposed to lobeline and its analogs

<table>
<thead>
<tr>
<th>Compound (3 μM)</th>
<th>Total Tissue DA Content (ng/mg)</th>
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<tbody>
<tr>
<td>Control</td>
<td>16.4 ± 1.20</td>
</tr>
<tr>
<td>Lobeline</td>
<td>19.7 ± 5.50</td>
</tr>
<tr>
<td>MTD</td>
<td>23.3 ± 1.50</td>
</tr>
<tr>
<td>(-)-TTD</td>
<td>17.7 ± 1.90</td>
</tr>
<tr>
<td>(+)-TTD</td>
<td>15.2 ± 2.40</td>
</tr>
<tr>
<td>Lobelane</td>
<td>15.0 ± 2.17</td>
</tr>
<tr>
<td>(-)-trans-Lobelane</td>
<td>13.9 ± 1.58</td>
</tr>
<tr>
<td>nor-MTD</td>
<td>21.7 ± 5.00</td>
</tr>
<tr>
<td>(-)-nor-TTD</td>
<td>22.2 ± 3.90</td>
</tr>
<tr>
<td>nor-lobelane</td>
<td>20.2 ± 2.43</td>
</tr>
</tbody>
</table>

Data are mean ± S.E.M.; n = 3 experiments/compound.

**TABLE 2**
Lobeline and its analogs do not increase DA overflow from rat striatal slices

<table>
<thead>
<tr>
<th>n</th>
<th>Concentration (μM)</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>0 *</td>
</tr>
<tr>
<td>Lobeline</td>
<td>4.5 ± 2.06</td>
</tr>
<tr>
<td>MTD</td>
<td>7.0 ± 3.5</td>
</tr>
<tr>
<td>(-)-TTD</td>
<td>3.5 ± 2.5</td>
</tr>
<tr>
<td>(+)-TTD</td>
<td>2.5 ± 2.0</td>
</tr>
<tr>
<td>Lobelane</td>
<td>2.5 ± 1.5</td>
</tr>
<tr>
<td>(-)-trans-Lobelane</td>
<td>2.0 ± 1.5</td>
</tr>
<tr>
<td>(+)-trans-Lobelane</td>
<td>1.0 ± 0.5</td>
</tr>
<tr>
<td>nor-MTD</td>
<td>0.0 ± 0.0</td>
</tr>
<tr>
<td>(-)-nor-TTD</td>
<td>0.0 ± 0.0</td>
</tr>
<tr>
<td>nor-lobelane</td>
<td>0.0 ± 0.0</td>
</tr>
</tbody>
</table>

*0* indicates buffer control condition (superfusion with buffer only, in the absence of lobeline or analog).

*Data are expressed as pg/30 ml/mg (mean ± S.E.M.). n indicates number of rats/compound.*
and (-)-TTD to increase \( p < 0.05 \) DOPAC overflow was 1 \( \mu M \), and the lowest concentration of (+)-TVD was 0.3 \( \mu M \) (Fig. 6, bottom). A concentration effect \( p < 0.001 \) was found for lobeline \((F_{4,11} = 9.89; \text{Fig. 6, top})\), (-)-trans lobelane \((F_{4,11} = 14.21)\) and (+)-trans lobelane \((F_{4,19} = 15.71)\) to increase DOPAC overflow (Fig. 6, bottom). The lowest concentration of lobeline to increase \( p < 0.05 \) DOPAC overflow was 1.0 \( \mu M \), and for both (-)-trans lobelane and (+)-trans lobelane, the lowest concentration was 3.0 \( \mu M \). A concentration effect \( p < 0.001 \) for nor-MTD \((F_{4,18} = 13.1)\) and nor-lobelane \((F_{4,19} = 26.93)\) was found, whereas \((-\)-nor-TTD \((F_{4,11} = 1.797, p = 0.194)\) did not increase DOPAC overflow (Fig. 6, bottom). The lowest concentration of both nor-MTD and nor-lobelane to increase \( p < 0.05 \) DOPAC overflow was 1 \( \mu M \).

**Lobeline, MTD, and Lobelane Inhibition of Methamphetamine-Evoked DA Overflow.** Lobeline-mediated inhibition of methamphetamine (5 \( \mu M \))-evoked DA release and fractional DA release is illustrated in Fig. 7. Lobeline \((0.3–3 \mu M)\) inhibited methamphetamine-evoked DA overflow \((IC_{50} = 0.42 \pm 0.10 \mu M, \text{max} = 56.1 \pm 4.18\%\)), and a concentration effect \((F_{5,19} = 41.24, p < 0.001)\) was found. A significant concentration × time interaction \((F_{9,180} = 22.07, p < 0.01)\) was found. Compared with the buffer control, methamphetamine evoked an increase in fractional DA release from 20 to 45 min after its addition to the superfusion buffer. A separate two-way repeated-measures ANOVA on the time course data revealed a concentration × time interaction \((F_{36,180} = 3.46, p < 0.01)\). Lobeline \((1.0–3.0 \mu M)\) decreased \( p < 0.05 \) methamphetamine-evoked fractional DA release from 30 to 35 min.

Figure 7 also illustrates the concentration effect of MTD \((F_{5,19} = 49.62, p < 0.001)\) on methamphetamine-evoked DA overflow. The lowest MTD concentration to produce inhibition was 0.1 \( \mu M \) \((IC_{50} = 0.44 \mu M; \text{max} = 76\%\)). The time course for MTD inhibition of methamphetamine-evoked fractional DA release is illustrated in Fig. 8 (middle). Methamphetamine evoked a significant increase in fractional DA release from 15 to 30 min. Analysis of the time course for MTD to inhibit methamphetamine-evoked fractional DA release revealed a concentration × time interaction \((F_{36,180} = 6.75, p < 0.001)\). MTD \((1.0–3.0 \mu M)\) decreased \( p < 0.05 \) methamphetamine-evoked fractional DA release from 25 to 35 min.

Figure 7 also illustrates the concentration-dependent inhibition of methamphetamine-evoked DA overflow by lobeline \((F_{5,14} = 11.91, p < 0.001)\). The lowest lobeline concentration to inhibit methamphetamine-evoked DA overflow was 1 \( \mu M \).
The IC$_{50}$ value for lobeline was 0.65 µM, and the $I_{\text{max}}$ value was 73%. The time course for lobeline to inhibit methamphetamine-evoked fractional DA release is shown in Fig. 8 (bottom), and a concentration × time interaction (F$_{5,125} = 4.93$, p < 0.01) was found. Lobeline at 1.0 µM significantly decreased methamphetamine-evoked fractional DA release from 30 to 35 min, and at 3.0 µM, lobeline decreased methamphetamine-evoked fractional DA release from 25 to 35 min.

(→)-trans-Lobelane, (+)-trans-Lobelane (→)-TTD, and (+)-TTD Inhibition of Methamphetamine-Evoked DA Overflow and the Time Course of Inhibition. Supplemental Fig. 1 (top) illustrates the inhibition of methamphetamine-evoked DA overflow by (→)-trans-lobelane and (+)-trans-lobelane. (→)-trans-Lobelane did not inhibit methamphetamine-evoked DA overflow. With respect to inhibition of methamphetamine-evoked fractional DA release, neither the main effect of concentration nor the concentration × time interaction were significant. Analysis of the inhibition of methamphetamine-evoked DA overflow by (→)-trans-lobelane showed a concentration effect (F$_{5,25} = 15.39$, p < 0.001). Subsequent post hoc analysis revealed that the lowest concentration of (→)-trans-lobelane to inhibit methamphetamine-evoked DA overflow was 3 µM. The IC$_{50}$ value for (→)-trans-lobelane was >3.0 µM, whereas the $I_{\text{max}}$ value was 34%. The $I_{\text{max}}$ value for lobeline was greater (p < 0.05) than that for (→)-trans-lobelane. A concentration × time interaction (F$_{36,225} = 1.84$, p < 0.01) was found for (→)-trans-lobelane inhibition of methamphetamine-evoked DA release (data not shown). For (+)-trans-lobelane, 0.3 to 3.0 µM decreased (p < 0.05) methamphetamine-evoked fractional DA release only at 35 min.

Supplemental Fig. 1 also illustrates the inhibitory effect of the enantiomers of TTD on methamphetamine-evoked DA overflow. Concentration effects for (→)-TTD (F$_{5,27} = 20.518$, p < 0.001) and (+)-TTD (F$_{5,18} = 20.48$, p < 0.001) were found. The lowest concentration to inhibit (p < 0.05) methamphetamine-evoked DA overflow for (→)-TTD was 0.1 µM; the IC$_{50}$ value was >3.0 µM, and the $I_{\text{max}}$ value was 42%. The $I_{\text{max}}$ value for MTD was greater (p < 0.05) than that for (→)-TTD. The lowest concentration of (+)-TTD to inhibit methamphetamine was 1 µM. The IC$_{50}$ value was 0.22 µM, and the $I_{\text{max}}$ value was 50%.

The time course of (→)-TTD- and (+)-TTD-induced inhibition of methamphetamine-evoked fractional DA release is illustrated in Supplemental Fig. 2. Inhibition of methamphetamine-evoked fractional DA release by (→)-TTD and (+)-TTD revealed concentration × time interactions (F$_{36,250} = 3.20$, p < 0.01; F$_{36,170} = 4.15$, p < 0.01, respectively). (→)-TTD (1.0–3.0 µM) decreased (p < 0.05) methamphetamine-evolved fractional DA release only at 30 min, whereas the inhibitory effect of (+)-TTD was more prolonged. At 3.0 µM, (+)-TTD decreased methamphetamine-evolved fractional DA release from 30 to 45 min. At 0.3 and 1.0 µM, (+)-TTD decreased fractional DA release from 35 to 45 min.

nor-MTD, (±)-nor-TTD, and nor-Lobelane Inhibition of Methamphetamine-Evoked DA Overflow. The inhibitory effect of nor-MTD, (±)-nor-TTD, and nor-lobelane on methamphetamine-evoked DA overflow is illustrated in Supplemental Fig. 3. A concentration effect for nor-MTD (F$_{5,19} = 49.62$, p < 0.001), (±)-nor-TTD (F$_{5,18} = 20.48$, p < 0.001), and nor-lobelane (F$_{5,27} = 20.518$, p < 0.001) was found. The lowest concentration to inhibit methamphetamine-evoked DA overflow was 0.3 µM, 0.1 µM, and 1 µM for nor-MTD, nor-lobelane, and (±)-nor-TTD, respectively. The IC$_{50}$ value for nor-MTD was 0.40 µM, and the $I_{\text{max}}$ value was 64%. Although IC$_{50}$ values were greater than 3 µM for nor-lobelane and (±)-nor-TTD, $I_{\text{max}}$ values of 47% and 34%, respectively, were obtained for the highest concentration of these analogs.

The time course for inhibition of methamphetamine-evoked fractional DA release by the nor-analogs is illustrated in Supplemental Fig. 4. A concentration × time interaction (F$_{36,195} = 3.66$, p < 0.01) was found for nor-MTD, and the highest concentration tested (3.0 µM) decreased (p < 0.05) methamphetamine-evoked fractional DA release from 30 to 35 min. (±)-nor-TTD inhibition of methamphetamine displayed a concentration × time interaction (F$_{36,125} = 3.39$, p < 0.01). (±)-nor-TTD at 1.0 µM decreased (p < 0.05) methamphetamine-evoked fractional DA release at 30 min, and at 3.0 µM from 30 to 35 min. A concentration × time interaction (F$_{36,215} = 3.30$, p < 0.01) was also found for nor-lobelane, and 3.0 µM decreased (p < 0.05) methamphetamine-evoked fractional DA release at 35 min.

Lobelane and (+)-trans-Lobelane Increase DOPAC Overflow in the Presence of Methamphetamine. A concentration effect was found on DOPAC overflow for only lobeline and (+)-trans-lobelane (F$_{5,19} = 49.62$, p < 0.001 and F$_{5,22} = 4.49$, p = 0.005, respectively; data not shown). Lobeline (0.1–3.0 µM) and (+)-trans-lobelane (0.3 and 3.0 µM) increased (p < 0.05) DOPAC overflow compared with the methamphetamine control condition (data not shown).

Discussion

Lobeline has high affinity for VMAT2 and low affinity for DAT (Teng et al., 1997, 1998; Dwoskin and Crooks, 2002). Lobeline increased DOPAC overflow, but did not increase DA overflow (Teng et al., 1997; Miller at al., 2001), indicating alterations in presynaptic DA storage and lack of MAO inhibition (Dwoskin and Crooks, 2002). In the current study, low concentrations of lobeline inhibited methamphetamine-evoked DA release from rat striatal slices at concentrations that inhibited amphetamine-evoked DA release (Miller et al., 2001). The current study provides SAR for a larger group of defunctionalized lobeline analogs to identify selective, high-affinity inhibitors of VMAT2 that inhibit methamphetamine-evoked DA release.

In the synaptosomal [3H]DA uptake assay assessing DAT function, lobeline markedly increased inhibitory potency, whereas MTD resulted in a further increase in potency compared to lobeline. An enantioselective effect was observed with the stereoisomers of MTD; (+)-TTD exhibited reduced potency at DAT relative to (→)-TTD, and both enantiomers exhibited decreased potency compared with MTD. This trend was not observed with the lobeline stereoisomers; (−) and (+)-trans-lobelane were equipotent with each other and with lobeline. N-Demethylation of MTD and (→)-TTD resulted in decreased potency, whereas N-demethylation of lobeline increased potency. Collectively, these data demonstrate that defunctionalization of lobeline increases inhibitory potency at DAT, whereas effects of N-demethylation and C2,C6 stereochemistry on potency depend on the nature of the C$_2$ linker units.

With respect to analog interaction with VMAT2, MTD was
~5-fold less potent than lobeline at inhibiting [3H]DTBZ binding, whereas (−)-TTD and (+)-TTD were equipotent with MTD inhibiting [3H]DTBZ binding, indicating that piperidino C2,6 stereochemistry is not a critical factor. Because both (−)-TTD and (+)-TTD lack affinity for α4β2* and α7* nAChRs (Zheng et al., 2005), these analogs are more selective for VMAT2 than either lobeline or MTD. trans-Isomers of lobeline were less potent than lobeline and lobeline inhibiting [3H]DTBZ binding to VMAT2; (−)-trans-lobelane and (+)-trans-lobelane were equipotent and had 5-fold lower affinity at VMAT2 compared with lobeline, consistent with the premise that C2,6 piperidino stereochemistry is not critical for VMAT2 interaction. Because (−)-trans-lobelane and (−)-trans-lobelane were inactive at α4β2* and α7* nAChRs (Zheng et al., 2005), these enantiomers are more selective for VMAT2 than lobeline, MTD, or lobelane. nor-MTD was 5-fold more potent than MTD at the [3H]DTBZ binding site, and because nor-MTD did not interact with α4β2* and α7* nAChRs (Zheng et al., 2005), it had greater selectivity for VMAT2 than MTD. Likewise, nor-lobelane and lobelane were equipotent at VMAT2, but nor-lobelane was more selective for VMAT2. (±)-nor-MTD was equipotent with MTD at the VMAT2 binding site, but (±)-nor-lobelane was more selective for VMAT2. Taken together, N-demethylation does not seem to alter affinity at the [3H]DTBZ binding site, but improves VMAT2 selectivity. A high degree of [3H]DTBZ binding site affinity does not necessarily translate into functional inhibition of VMAT2. Therefore, the ability of these compounds to inhibit [3H]DA uptake into synaptic vesicles was evaluated. Lobelane displayed a 10-fold increase in potency compared with lobeline inhibiting [3H]DA uptake at VMAT2, and nor-lobelane was equipotent with lobeline. In contrast, (+)- and (−)-trans-lobelane exhibited markedly reduced inhibition of [3H]DA uptake, indicating that whereas the N-methyl group is not critical, cis-stereochemistry of the defunctionalized, saturated compounds is a contributing structural characteristic for potent VMAT2 inhibition. MTD was equipotent with lobeline inhibiting [3H]DA uptake, and was 10-fold less potent than lobeline. (+)-TTD, (−)-TTD, and (±)-nor-TTD were equipotent with MTD. However, nor-MTD had the lowest potency, whereas nor-lobelane and lobelane were equipotent in the VMAT2 functional assay. Thus, demethylation-induced changes in potency were most evident in the unsaturated MTD series of analogs.

In the methamphetamine-evoked DA overflow assay, MTD and lobelane had the best profile of inhibition. MTD inhibited methamphetamine-evoked DA overflow and was equipotent with lobeline; however, maximal inhibition was greater with MTD. (−)-TTD was equipotent with both MTD and lobeline inhibiting methamphetamine-evoked DA release; however, maximal inhibition produced by (−)-TTD was only 50%. Although maximal inhibition produced by the TTD enantiomers was not different, enantioselective effects on potency were observed. Lobelane was equipotent with lobeline, although lobelane exhibited greater maximal inhibition of methamphetamine-evoked DA release. (−)-trans-Lobelane was less potent than lobeline, and exhibited only 34% inhibition. (−)-trans-Lobelane did not inhibit methamphetamine. Thus, the combination of chemical defunctionalization and C2,6-cis-stereochemistry affords analogs with a better profile for inhibition of methamphetamine effects.

nor-MTD was equipotent with MTD and exhibited similar maximal inhibition of methamphetamine-evoked DA release. Thus, removal of the methyl group in MTD did not alter its inhibitory profile. Because nor-MTD had no activity at α4β2* and α7* nAChRs, it is more selective for VMAT2 than MTD. nor-Lobelane had both a lower potency and lower maximal inhibition of the effect of methamphetamine compared with lobelane, which contrasts with its equipotent and complete inhibition of VMAT2 function. Thus, with respect to methamphetamine-evoked DA release, the N-methyl group improves the inhibitory profile in the saturated lobelane compounds, but has no impact in the unsaturated MTD series.

A comparison of results from the three assays reveals SAR inconsistencies and a different rank order of potencies. Inconsistencies in [3H]DTBZ binding and vesicular [3H]DA uptake to predict the ability of the analogs to inhibit methamphetamine were noted. For example, MTD was 5-fold less potent inhibiting [3H]DTBZ binding than was lobeline, whereas MTD and lobeline had similar potencies inhibiting methamphetamine. Although both trans-lobelane enantiomers inhibited [3H]DTBZ binding to VMAT2 equipotent, (−)-trans-lobelane showed only ~30% inhibition of methamphetamine-evoked DA release, whereas (−)-trans-lobelane exhibited no inhibition. In addition, (−)-trans-lobelane was a false positive, i.e., active in VMAT2 binding and uptake assays, but inactive in the methamphetamine-evoked DA release assay. Although both (−)-TTD and (−)-TTD inhibited [3H]DTBZ binding equipotently, (−)-TTD had lower potency inhibiting methamphetamine. Potential interpretations of these results are that an additional site on VMAT2 or alternative vesicle protein sites are being targeted.

Lobelane exhibited the best overall profile, having high affinity at the VMAT2 binding site, high potency inhibiting VMAT2 function, low potency inhibiting DAT function, and potently and almost completely inhibiting methamphetamine-evoked DA release. Taking into account the greater selectivity for VMAT2 exhibited by lobelane compared with lobeline, lobelane has greater potential as a lead compound for the development of treatments for methamphetamine abuse. Lobelane-induced inhibition of VMAT2 function was 10-fold greater than lobeline, despite lobeline having only 2-fold greater affinity than lobeline for the [3H]DTBZ binding site, suggesting that lobelane is acting at alternate sites on VMAT2 to inhibit function or that the mechanism of inhibition (i.e., competitive versus noncompetitive) is different between these two compounds. To elucidate the mechanism of action, saturation kinetic assays were conducted by use of the synaptic vesicle preparation. Results showed that both compounds inhibited vesicular [3H]DA uptake competitively, supporting the contention that lobelane and lobeline act at different sites on VMAT2.

All analogs studied inhibited VMAT2 function. However, several potential underlying mechanisms by which these analogs inhibit methamphetamine are possible. Some of these analogs are probably interacting with DAT to inhibit the effects of methamphetamine to evoke DA release. Four analogs were equipotent in inhibiting DAT and VMAT2 function, including (−)-TTD, (−)-TTD, (−)-trans-lobelane, and (−)-trans-lobelane. Furthermore, MTD and nor-MTD inhibited DAT with 12- and 42-fold greater potency, respectively, than at VMAT2. Thus, the underlying mechanism for these analogs to inhibit methamphetamine-evoked DA release may be via inhibition of DAT function.
In contrast, lobeline and lobelane exhibited 67- and 35-fold greater potency, respectively, in inhibiting VMAT2 function than DAT function. Thus, there is a high likelihood that lobelane and lobeline are acting at VMAT2 to inhibit the effect of methamphetamine.

Another potential mechanism for analog-induced inhibition of methamphetamine-evoked DA release may be through facilitation of DA metabolism to DOPAC via reverse transport. Therefore, the presence of presynaptic DA from the vesicles to the cytosol. Once in the cytosol, DA would be available for metabolism to DOPAC by MAO. Although not considered "release" (i.e., exocytotic or transporter-mediated), extracellular DOPAC concentration is altered in response to neuronal activity and drugs, and essentially reflects increased dopaminergic neuron activity. These analogs produce no reduction in striatal DA content. Previous studies determining the mechanism of action of lobeline revealed increased [3H]DA overflow from rat striatal slices in the presence of nomifensine and pargyline (DAT and MAO inhibitors, respectively), whereas in their absence, lobeline increased DOPAC overflow. In addition, lobeline inhibited both [3H]DA uptake and elicited [3H]DA release from synaptic vesicles. It is noteworthy that lobeline reduced methamphetamine-evoked DA overflow from striatal slices regardless of the presence or absence of pargyline (Dowski and Crooks, 2002).

Together, these results indicate that lobeline redistributes presynaptic DA from synaptic vesicles to the cytosol, where it is metabolized to DOPAC by MAO, thus reducing the DA pool available for DAT-mediated reverse-transport after methamphetamine exposure. After interaction with VMAT2, altered DA redistribution and metabolism may contribute to the underlying mechanism responsible for lobeline-mediated inhibition of methamphetamine.

In summary, lobeline inhibits methamphetamine-evoked DA release and is more selective inhibiting VMAT2 than lobeline. Lobelane is 35-fold more potent as an inhibitor of VMAT2 function than DAT function, consistent with the interpretation that lobeline acts at VMAT2 to inhibit methamphetamine effects. Thus, lobeline is a promising lead for interpretation that lobelane acts at VMAT2 to inhibit methamphetamine. Thus, there is a high likelihood that lobeline and lobeline are acting at VMAT2 to inhibit the effect of methamphetamine.

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References


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