

NIH Public Access

Author Manuscript

ACS Chem Neurosci. Author manuscript; available in PMC 2012 December 21.

Published in final edited form as:

ACS Chem Neurosci. 2011 December 21; 2(12): 730–742. doi:10.1021/cn200090z.

The Discovery and Characterization of ML218: A Novel, Centrally Active T-Type Calcium Channel Inhibitor with Robust Effects in STN Neurons and in a Rodent Model of Parkinson's Disease

Zixiu Xiang^{1,4,†}, Analisa D. Thompson^{1,4,†}, John T. Brogan^{2,3}, Michael L. Schulte^{2,3}, Bruce J. Melancon^{1,3,4}, Debbie Mi^{1,4}, L. Michelle Lewis^{1,4}, Bende Zou⁷, Liya Yang⁷, Ryan Morrison^{1,3,4}, Tammy Santomango^{1,4}, Frank Byers^{1,4}, Katrina Brewer^{1,4}, Jonathan S. Aldrich^{3,4}, Haibo Yu⁵, Eric S. Dawson^{1,3,4}, Min Li^{5,6}, Owen McManus^{5,6}, Carrie K. Jones^{1,4}, J. Scott Daniels^{1,3,4}, Corey R. Hopkins^{1,2,3,4}, Ximin Simon Xie⁷, P. Jeffrey Conn^{1,3,4}, C. David Weaver^{1,3,4}, and Craig W. Lindsley^{1,2,3,4,*}

¹Department of Pharmacology, Vanderbilt University Medical Center, Nashville, TN 37232-6600, USA

²Department of Chemistry, Vanderbilt University Medical Center, Nashville, TN 37232-6600, USA

³Vanderbilt Specialized Chemistry Center for Accelerated Probe Development (MLPCN), Vanderbilt University Medical Center, Nashville, TN 37232-6600, USA

⁴Vanderbilt Center for Neuroscience Drug Discovery, Vanderbilt University Medical Center, Nashville, TN 37232-6600, USA

⁵Department of Neuroscience, High Throughput Biology Center, Baltimore, MD 21205, USA

⁶Johns Hopkins Ion Channel Center (JHICC), Baltimore, MD 21205, USA

⁷AfaSci Research Laboratory, AfaSci, Inc. Redwood, CA 94063, USA

Abstract

T-type Ca^{2+} channel inhibitors hold tremendous therapeutic potential for the treatment of pain, epilepsy, sleep disorders, essential tremor and other neurological disorders; however, a lack of truly selective tools has hindered basic research, and selective tools from the pharmaceutical industry are potentially burdened with intellectual property (IP) constraints. Thus, an MLPCN

Corresponding author: Professor Craig W. Lindsley, Departments of Pharmacology and Chemistry, Vanderbilt Specialized Chemistry Center (MLPCN), Vanderbilt Center for Neuroscience Drug Discovery, Vanderbilt University Medical Center, Nashville, TN 37232-6600, USA. craig.lindsley@vanderbilt.edu; Ph: 615-322-8700; FAX: 615-343-6532. ^T These authors contributed equally to this work.

These authors contributed equality

Author Contributions

Professors Lindsley, Weaver, Conn, Jones, Daniels and Dawson oversaw and designed the chemistry, HTS, molecular pharmacology, behavioral pharmacology, DMPK and modeling, respectively.

Professor Thompson performed the haloperidol-induced catalepsy studies.

Professor Xiang performed the electrophysiology studies.

Professor Hopkins oversaw chemistry.

Mr. Brogan performed synthetic chemistry.

Mr. Schulte performed synthetic chemistry.

Dr. Melancon performed synthetic chemistry and scaled-up key compounds for in vivo studies.

Dr. Xie was the assay provider, coordinated the HTS and electrophysiology.

Dr. Zou and Dr. Yang performed electrophysiology.

Mr. Aldrich performed synthetic chemistry.

Dr. Lewis and Mrs. Mi conducted the HTS.

Mrs. Santomango, Mr. Byers, Mr. Morrison and Ms. Brewer performed DMPK and bioanalytical studies.

Dr. Yu, Dr. McManus and Dr. Li performed the Cav3.2/3.3 electrophysiology

high-throughput screen (HTS) was conducted to identify novel T-type Ca^{2+} channel inhibitors free from IP constraints, and freely available through the MLPCN, for use by the biomedical community to study T-type Ca²⁺ channels. While the HTS provided numerous hits, these compounds could not be optimized to the required level of potency to be appropriate tool compounds. Therefore, a scaffold hopping approach, guided by SurflexSim, ultimately afforded ML218 (CID 45115620) a selective T-Type Ca²⁺ (Ca_v3.1, Ca_v3.2, Ca_v3.3) inhibitor (Ca_v3.2, IC₅₀ = 150 nM in Ca²⁺ flux; Ca_v3.2 IC₅₀ = 310 nM and Ca_v3.3 IC₅₀ = 270 nM, respectively in patch clamp electrophysiology) with good DMPK properties, acceptable in vivo rat PK and excellent brain levels. Electrophysiology studies in subthalamic nucleus (STN) neurons demonstrated robust effects of ML218 on the inhibition of T-Type calcium current, inhibition of low threshold spike and rebound burst activity. Based on the basal ganglia circuitry in Parkinson's disease (PD), the effects of ML218 in STN neurons suggest a therapeutic role for T-type Ca^{2+} channel inhibitors, and ML218 was found to be orally efficacious in haloperidol-induced catalepsy, a preclinical PD model, with comparable efficacy to an A_{2A} antagonist, a clinically validated PD target. ML218 proves to be a powerful new probe to study T-Type Ca²⁺ function *in vitro* and *in vivo*, and freely available.

Keywords

T-Type calcium channel; inhibitor; electrophysiology; Parkinson's disease

Introduction

Voltage-gated calcium channels (VGCCs) modulate the entry of calcium (Ca²⁺) into cells in response to depolarization of the membrane and elicit a number of Ca²⁺-dependent processes.¹ VGCCs are classified as either high-voltage-activated (L-, N-, P/Q-, and Rtypes) or low-voltage-activated (T-type).²⁻⁷ The 'T' in T-type Ca²⁺ channels refers to their characteristic transient, or fast, inactivation of the channel and small conductance.^{2–8} Molecular cloning studies have identified three genes (CACNAIG, CACNAIH, CACNAII) encoding the main pore-forming $\alpha 1$ subunit resulting in three members in the T-Type calcium channel family: $Ca_v 3.1$ ($\alpha 1G$), $Ca_v 3.2$ ($\alpha 1H$), and $Ca_v 3.3$ ($\alpha 1I$).^{2–10} All three $Ca_v 3$ family members are heterogeneously expressed in the brain and many peripheral organs such as the heart and vascular smooth muscle. A wealth of literature suggests that subtypeselective Ca_v3 modulators should produce more specific pharmacological actions for a variety of neurological, psychiatric, and cardiovascular disorders (such as epilepsy, pain, movement disorders, hearing loss, sleep/wake states, cancer and over-active bladder) with fewer side effects than pan-Ca²⁺ channel inhibitors.^{2–11} Importantly, T-type Ca²⁺ channels are strongly associated with rhythmic firing patterns in key brain regions within the CNS.^{2–11} For example, Ca_v3.1 T-Type calcium channels are present in the olivocerebellar system, and inhibitors could block tremor-rhythm generation, suggesting potential therapeutic potential for essential tremor.¹²⁻¹⁶ This postulate has been borne out in both preclinical models of essential tremor as well as in the clinic with zonisamide (ZNS), and lamotrigine, antiepileptic drugs possessing T-type calcium channel inhibitory activity in addition to blockade of voltage-gated Na⁺ channels.¹²⁻¹⁶ This could also be extended to the basal ganglia circuit, where abnormal burst firing results in over-activation of the indirect pathway, leading to Parkinsonian tremors,¹⁷⁻²⁰ and this is where our interest in this target resides. However, the therapeutic potential of T-type calcium channels has been hampered by both a lack of truly selective, central active small molecule tools and, in the case of exceptional tool compounds, intellectual property constraints.

The first generation of T-Type Ca^{2+} inhibitors were discovered based on drugs originally developed for other targets (Figure 1). The antiepileptic drugs zonisamide (ZNS) **1** and

ethosuximide **2** were found to inhibit T-type calcium channel at therapeutic concentrations.^{20,21} Roche later identified the dual T-/L-type calcium channel antagonist mibefradil **3**, which was briefly marketed as an antihypertensive agent.^{22,23} Studies have shown that **3** is a potent T-type inhibitor with 10- to 30-fold selectivity for T- over L-type Ca^{2+} channels. Therefore, it has been widely used as a pharmacological tool for studying T-type Ca^{2+} channels.^{22,23} However, mibefradil has poor penetration into CNS. Similarly, efonidipine **4** was also shown to preferentially inhibit T- over L-type Ca^{2+} channels.²⁴ Shortly thereafter, the neuroleptic agent flunazirine **5**²⁵ and the antipsychotic pimozide **6**²⁶ were shown to inhibit (Ca_v3 IC₅₀s ~600 nM) or bind (K_i ~40 nM), to T-type Ca²⁺ channels; however, both were far more potent against biogenic amine targets. While these tools provided modest advances in our understanding of T-type Ca²⁺ channels, they did provide evidence that inhibition of T-type Ca²⁺ channels in man was well tolerated upon both acute and chronic dosing with **1–6**.^{20–26}

The majority of the second generation T-Type Ca^{2+} inhibitors were derived from HTS campaigns in pharmaceutical companies aimed at developing selective T-Type Ca^{2+} inhibitors and are covered by patents (Figure 2).²⁷ Here, the major players of published efforts are Merck, inhibitor series **7–10**, and Korea Institute of Science and Technology, inhibitor series **11**, **13** and **14**.^{28–40} Of these, the Merck series are the most potent, possessing good DMPK profiles and *in vivo* activity in multiple preclinical models. Interestingly, **7** and **8** are voltage independent, while **9** displays voltage-dependent inhibition. Importantly, there are no reports of inhibitors selective amongst the Ca_v3 family (all are equipotent on $Ca_v3.1$, $Ca_v3.2$ and $Ca_v3.3$). While compounds **7–15** represent a major advance, they are covered by patents and/or published patent applications that can restrict their use by the broader biomedical research community.^{27–40}

Results & Discussion

In 2005, the NIH Molecular Libraries Roadmap launched the Molecular Libraries Screening Center Network (MLSCN), a nationwide consortium of facilities that provide highthroughput small molecule screening of biological targets/pathways for which small molecule tools are unavailable.⁴¹ One such target that lacked the appropriate small molecule tools was, at that time, T-Type Ca^{2+} channels, as only **1–6** were known.^{20–26} Xie and his colleagues have developed and validated an HTS assay using a FLIPR (Fluorescent Imaging Plate Reader).⁴² The target and the assay were assigned to the Vanderbilt Specialized Screening Center for GPCRs, Ion Channels and Transporters for primary drug screen to identify novel selective T-type Channel inhibitors from the MLSCN compound library.⁴¹ During the course of this initiative, the MLSCN became the Molecular Libraries Production Center Network (MLPCN), with increased medicinal chemistry expertise to develop high quality probes for the biomedical community, of which the Vanderbilt Specialized Chemistry Center and the Johns Hopkins Ion Channel Center are members.⁴¹ Also during this time, the pharmaceutical industry disclosed 7–15, which not only raised the bar for an impactful T-Type Ca²⁺ channel inhibitor probe, but also required that our Centers develop a T-Type Ca²⁺ channel inhibitor probe free of potential IP constraints, of comparable quality to 7-15, and to be freely available through the MLPCN mechanism for use by the biomedical research community.²⁷⁻⁴⁰

High-Throughput Screening and Identification of T-Type Ca²⁺ channel inhibitors

For the discovery of novel T-Type Ca^{2+} channel inhibitors, we completed a primary HTS using the 2008 collection of MLSCN library containing 110,720 compounds screened at 10 μ M in a single measurement.⁴¹ The primary assay was run against the Ca_v3.2 channel expressed in HEK293 cells, and performed in a 384-well format using an FDSS 6000 kinetic imaging plate measuring Ca²⁺ fluorescence. Data quality was controlled by the known T-

Type Ca²⁺ channel inhibitor **8**²⁹ as the positive control and the 0.1% DMSO vehicle as the negative control. The assay quality control used a screening window coefficient "Z-factor". The Z' values were between 0.6–0.8, indicating excellent assay quality. Primary data analysis was performed by comparing the activity of any test compound with all the other test compounds on the same plate, taking the ratio max in the kinetic time window of 12–40 sec with 0–1 sec as the baseline. Primary hits were defined as any outliers that were different from all the others based on a *z*-score threshold of 3, which resulted in 4,246 hits. For the confirmation screen, 890 hits were available from Biofocus-DPI and tested with full concentration-response curves (CRCs), validating only seven hits as Ca_v3.2 inhibitors from the primary screen. A counter-screen with the parental HEK line, with native Ca²⁺ channels, presumably both L- and N-Type Ca²⁺ channels, indicated that only one confirmed hit was selective for T-Type Ca²⁺ channels, **16** (CID3373841), with an IC₅₀ of 2.5 μ M (Figure 3). Moreover, this represented a new T-Type Ca²⁺ inhibitor chemotype, and was only active in 23/440 MLPCN assays (never for an ion channel), suggesting minimal promiscuous pharmacology and a good starting point for chemical lead optimization.⁴³

Chemical Optimization of 16 (CID3373841)

The modular nature of **16**, coupled with the commercial availability of 2-amino oxadiazole monomers led to a two-dimensional library (Figure 4) of 56 analogs, wherein the western aryl moiety and the eastern amide moiety of **16** were varied. All analogs were screened in full CRC in the kinetic Ca²⁺ fluorescence assay. Structure-activity-relationships (SAR) was rather flat, with the best analog **17** possessing an IC₅₀ of 1.1 μ M, ~10-fold less potent than the best known inhibitors **7–15**. Evaluation of **17** in Ca_v3 patch electrophysiology (EP) experiments demonstrated an ~10-fold right-shift in potency (Ca_v3.2 IC₅₀ = 13.5 μ M and Ca_v3.3 IC₅₀ = 11.5 μ M), affording weak inhibition of T-Type Ca²⁺ channels and far from a high quality MLPCN probe for this target.

While studying a molecular model of **17** and comparing it to a model of Merck's **8**, it appeared as though the oxadiazole moiety of **17** could be mimicking the piperidine spacer element in **8**, although, poorly in this case due to the presence of western aryl ring. Therefore, we prepared a second generation library of analogs (Figure 5) in which the western aryl group in **17** was replaced with branched aliphatic moieties to more closely align with Merck's **8**.²⁹ A sub-micromolar inhibitor 18 (CID46943243, IC₅₀ = 810 nM) resulted, but was still not potent enough to meet target criteria. A SurflexSim⁴⁴ morphological similarity algorithm was used to perform multiple flexible alignments of **18** with Merck's **8**, which suggested that the oxadioazole moiety was in fact mimicking the piperidine ring, but the hydrogen bond donors/acceptors of the amide moieties were not properly aligned (Fig. 5D). Moreover, patch EP with **18** still displayed a significant right-shift in potency that decreased enthusiasm for continued optimization efforts within this scaffold.

Based on the overall profile of Merck's **8** and the overlay with **18**, we decided to 'scaffold hop' around Merck's **8** to access patent-free space and hopefully a more potent T-Type Ca²⁺ channel inhibitor probe.²⁹ Thus, we explored patent databases, and in conjunction with molecular modeling, identified two potential scaffolds to explore that would be free of IP issues, if we could engender good T-Type Ca²⁺ channel activity. The first was based on a 3amino piperdine scaffold (Figure 6). Employing solution phase parallel synthesis, we quickly constructed two, 12-member libraries (six with the (*R*)-enantiomer **19** and six with the (*S*)-enantiomer **20**) wherein the secondary piperdinyl nitrogen was substituted with either a benzyl group or a *tert*-butyl ethyl moiety and the exo-cyclic primary amine was acylated with a variety of acid chlorides to deliver analogs libraries **21** and **22**. This effort provided only one active compound, **23** (CID45115608), the (*S*)-enantiomer with an IC₅₀ of 2.4 μ M. Interestingly, the (*R*)-enantiomer (CID45115614) was devoid of activity.

In parallel, we were advancing a [3.1.0] scaffold, wherein modeling suggested we need the (1R,5S,6S) stereochemistry about the 3-azabicylco[3.1.0]hexan-6ylmenthanamine core 24. As in the Merck series containing 8^{29} the 3,5-dichlorobenzamide was optimal, thus we held that moiety constant in our first generation library, and surveyed 8 different alkyl groups on the secondary nitrogen. Thus, acylation of 24 with 3,5-dichlorobenzoyl chloride produced 25. Deprotection of the Boc group and standard reductive amination delivered analogs 26 (Scheme 1). This library proved far more productive than the 3-aminopiperidine series. Here, four of eight analogs (Figure 7) had potent T-type Ca²⁺ channel inhibitory activity, with **26b** (CID45115620) possessing activity comparable to Merck's 8^{29} with an IC₅₀ of 150 nM. Now, a SurflexSim⁴⁴ flexible alignment of **26b** with Merck's **8** and **18** provides almost perfect placement of key groups due to the conformational lock of the boat orientation of the enantiopure [3.1.0] ring system (Figure 7B), which is clearer in Figure 7C, showing overlay of 8 and 26b alone. The potency of 26b confirmed in patch EP, with a Ca_v3.2 IC₅₀ of 310 nM and a $Ca_v 3.3 IC_{50}$ of 274 nM (Figure 8), and in excellent agreement with the Ca^{2+} fluorescence data. Thus, 26b, derived via 'scaffold hopping', was declared an MLPCN probe and assigned ML218.⁴¹ Importantly, ML218 and related structures are not covered under any issued patent or patent application; therefore, the MLPCN probe, ML218, is free of any use restrictions.

Ancillary Pharmacology and DMPK Profiling of ML218

Ancillary pharmacology is a major concern and an issue that plagued both the first and second generation of T-type Ca²⁺ channel inhibitors.^{20–40} The original Merck HTS hit was a des-fluoro analog of 8, which was a potent T-type Ca²⁺ channel inhibitor, but the basic p K_a (8.7) of the piperdinyl nitrogen, in combination with lipophillic moieties at both termini, led to promiscuous ion channel and GPCR pharmacology.²⁹ To circumvent this, the installation of the *cis* β -fluoroamine in **8** maintained T-type Ca²⁺ channel inhibition, but attenuated amine bascitiy ($pK_a = 7.9$) to a degree where ancillary pharmacology significantly improved.²⁹ Due to modest inductive effects of the π -character of the cyclopropyl ring of the [3.1.0] system, the pK_a (8.1) of the bicyclic nitrogen was also diminished, relative to a piperidine nitrogen, but to a lesser degree than the *cis* β -fluoroamine in **8**.²⁹ However, even this subtle diminution in pK_a translated to a clean ancillary pharmacology profile for ML218. To more fully characterize this novel Ca_v^3 inhibitor probe molecule, ML218 was tested in Ricerca's Lead Profiling Screen (binding assay panel of 68 GPCRs, ion channels and transporters screened at 10 µM), and was found to only significantly bind two targets (sodium channel site 2 and sigma 1) of the 68 assays conducted (no inhibition of radio ligand binding > 50% at 10 μ M).⁴⁵ Importantly, ML218 had no significant inhibition of Land N-type calcium channels (17-49% inhibition @10 µM, respectively), KATP potassium channel (4% inhibition @10 µM) or hERG (48% inhibition @10 µM).⁴⁵

ML218 was evaluated in our tier 1 *in vitro* DMPK panel.^{46–49} In plasma protein binding studies (equilibrium dialysis), ML218 possessed good free fraction in both rat ($f_u = 9.1\%$) and human ($f_u = 3.3\%$). ML218 displayed minimal inhibition of P450 enzymes, including 3A4 (IC₅₀ >30 µM), 2C9 (IC₅₀ >30 µM), 1A2 (IC₅₀ = 10.8 µM) and 2D6 (IC₅₀ = 1.7 µM). Intrinsic clearance experiments in liver microsomes indicated that ML218 was highly cleared in rat (CL_{int} = 115 ml/min/kg), but low to moderately cleared in human liver microsomes (CL_{int} = 12.7 ml/min/kg). Results from an *in vivo* pharmacokinetic (PK) study in rats (1 mg/kg, IV) were consistent with the in vitro assessment, specifically with regards to an observed plasma clearance value (CL_P = 56 mL/min/kg) obtained from rat liver microsomes. Noncompartmental PK analysis indicated ML218 had a mean residence time (MRT) of nearly 7 hours, a value which was consistent with its terminal half life ($t_{1/2} = 7$ hours). Importantly, ML218 was found to be highly brain penetrant with a Brain_{AUC}/

Plasma_{AUC} of 7.4, implicating ML218's potential as an exceptional probe for CNS studies. The profile of ML218 is virtually identical to Merck's **8**, which showed modest rat PK parameters, but excellent dog, rhesus and predicted human PK parameters.²⁹ Thus, the highly selective probe, ML218 (CID 45115620), possesses the requisite attributes (PK and CNS penetration) necessary to dissect the role of T-type Ca²⁺ channel inhibition *in vitro* and *in vivo*.

T-Type Ca²⁺ Channel Inhibitors and Parkinson's Disease

As mentioned earlier, T-type Ca²⁺ channels are strongly associated with rhythmic firing patterns in key brain regions within the CNS, such as in the olivocerebellar system, and inhibitors do block tremor-rhythm generation, affording therapeutic potential for essential tremor.^{13–16,28,29} This concept could also be extended to the basal ganglia circuit, where abnormal burst firing in the STN results in over-activation of the indirect pathway, leading to Parkinsonian tremors.^{17–19} Almost two decades ago, Hassani and co-workers demonstrated that 6-hydroxydopamine (6-OHDA) lessioned rats showed increased incidence of burst firing in the STN than control animals.¹⁷ It is now well established that modification from a single-spike activity to a mixed bursting mode discharge pattern in STN neurons is one of the major hallmarks of Parkinsonism in rats and primates. Moreover, both high-frequency stimulation (HFS) and/or surgical resection of the STN ameliorates motor disturbances in PD patients, and significantly, the burst-firing mode of STN neurons can be attributed to both T- and L-type Ca²⁺ currents.^{17–19} In a series of papers from Merck, both 7 and 8 were efficacious in dose-response on reversing harmaline-induced essential tremor, which led them to evaluate **8** in the haloperidol-induced catalepsy model of PD.^{28,29} Here, a 10 mg/kg oral dose significantly reduced haloperidol-induced catalepsy, but there was no internal control by which to gauge the efficacy. To date, little has been done to thoroughly study T-type Ca²⁺ channel inhibitors as potential therapeutics for PD.

Electrophysiology Studies with ML218 in STN Neurons

Previous studies suggest that the burst-firing mode of STN neurons can be attributed to both T- and L-type Ca²⁺ currents.^{17–19} Thus, we examined the effect of ML218 on the T-type calcium current, low threshold spike (LTS) and rebound burst activity in STN neurons (Figures 9–11). In a voltage clamp experiment, bath application of 3 μ M ML218 significantly reduced the T-type Ca²⁺ current by ~45% (Figure 9). Conversely, in a current clamp mode, 3 μ M ML218 inhibited >50% of LTS amplitude, which was triggered by returning of a hyperpolarizing current pulse (-160 pA) in STN neurons (Figure 10). Similarly, 3 μ M ML218 depressed >60% of rebound burst activity induced by intracellular injection of a hyperpolarizing current (-100 pA) followed by a depolarizing current (-20 pA) in STN neurons (Figure 11). Together, these data suggest T-type Ca²⁺ channel inhibitors are an attractive therapeutic target for PD to regulate the abnormal burst firing in the STN, and therefore able to reduce signaling in the indirect pathway of the basal ganglia circuit.

In Vivo Efficacy of ML218 in the Haloperidol-Induced Catalepsy Model of PD

In our PD programs, haloperidol-induced catalepsy is our first tier pharmacodynamic model to access therapeutic utility for PD.^{49–51} In this model, a cataleptic state is induced by the administration of the dopamine antagonist haloperidol. Test compounds are then added, and potential anti-Parkisonian efficacy is based on a reversal of the cataleptic state.^{49–51} As a control, we employ an A_{2A} antagonist, as previous studies have shown that selective A_{2A} antagonists produce robust anti-Parkinsonian-like effects in animal models of dopamine depletion as well as in recent clinical trials with PD patients, when given alone or in combination with L-DOPA.⁵² As shown in Figure 12A, Merck's **8**²⁹ at 10, 30 and 56.6 mg/kg i.p. reversed cataleptic behavior in rats induced by a 0.75 mg/kg dose of haloperidol, and this is comparable to a 56.6 mg/kg oral dose of an A_{2A} inhibitor. By way of comparison,

ML218 at doses of 1, 3, 10 and 30 mg/kg p.o. reversed cataleptic behavior in rats induced by a 0.75 mg/kg dose of haloperidol, and this is comparable to a 56.6 mg/kg p.o. dose of an A_{2A} inhibitor (Figure 12B). Notably, the reversal with orally dosed ML218 approached the efficacy of a 56.6 mg/kg p.o. dose of the A_{2A} inhibitor.⁵² Taken together with the electrophysiology data in the STN, these experiments confirmed and validated inhibition of T-type Ca²⁺ channels as a therapeutic target for PD. Moreover, free brain and plasma concentrations of ML218 increased in a dose proportional manner across the dose range (3 mg/kg: [plasma] = 98 nM, [brain] = 1.66 μ M; 10 mg/kg: [plasma] = 282 nM, [brain] = 5.03 μ M; 30 mg/kg: 1.2 μ M, [brain] = 17.7 μ M)⁵³ with no alterations in brain/plasma ratios (average Brain_{AUC}:Plasma_{AUC} of 16.9).

In summary, an MLPCN high-throughput screen (HTS) was conducted to identify novel Ttype Ca²⁺ channel inhibitors free from IP constraints for use by the biomedical community to study T-type Ca²⁺ channels. While the HTS provided numerous hits, these compounds could not be optimized to the required level of potency to be appropriate tool compounds. Therefore, a scaffold hopping approach, guided by SurflexSim,⁴⁴ ultimately afforded ML218 (CID 45115620) a selective T-Type Ca^{2+} (Ca_v3.1, Ca_v3.2, Ca_v3.3) inhibitor (Ca_v3.2, IC₅₀ = 150 nM in Ca²⁺ flux; Ca_v3.2 IC₅₀ = 310 nM and Ca_v3.3 IC₅₀ = 270 nM in patch clamp electrophysiology) with good DMPK properties, acceptable in vivo rat PK and excellent brain levels. Electrophysiology studies in STN neurons demonstrated robust effects of ML218 on the inhibition of T-Type calcium current, inhibition of low threshold spike and rebound burst activity. ML218 was found to be orally efficacious in a dosedependent manner in a preclinical PD model, haloperidol-induced catalepsy, and comparable to clinically validated A2A antagonism. Based on the over-active, indirect pathway in the basal ganglia circuitry in Parkinson's disease (PD), the effects of ML218 in STN neurons and in an in vivo PD model suggest a therapeutic role for T-type Ca²⁺ channel inhibitors. Thus, ML218 is a powerful new probe to study T-Type Ca²⁺ function *in vitro* and in vivo, and, as all MLPCN probes, is freely available upon request. Current efforts are focused on the exploring the efficacy of ML218 in additional preclinical PD models, comparing the effects of voltage-dependent and voltage-independent T-type Ca^{2+} channel inhibitors, and these data will be reported in due course.

Methods

Chemical Synthesis and Purification

All ¹H & ¹³C NMR spectra were recorded on Bruker AV-400 (400 MHz) or Bruker AV-NMR (600 MHz) instrument. Chemical shifts are reported in ppm relative to residual solvent peaks as an internal standard set to δ H 7.26 or δ C 77.0 (CDCl₃) and δ H 3.31 or δ C 49.0 (CD₃OD). Data are reported as follows: chemical shift, multiplicity (s = singlet, d =doublet, t = triplet, q = quartet, br = broad, m = multiplet), integration, coupling constant (Hz). IR spectra were recorded as thin films and are reported in wavenumbers (cm-1). Low resolution mass spectra were obtained on an Agilent 1200 LCMS with electrospray ionization. High resolution mass spectra were recorded on a Waters Qtof-API-US plus Acquity system. The value \varDelta is the error in the measurement (in ppm) given by the equation $\Delta = [(ME - MT)/MT] \times 106$, where ME is the experimental mass and MT is the theoretical mass. The HRMS results were obtained with ES as the ion source and leucine enkephalin as the reference. Optical rotations were measured on a Perkin Elmer-341 polarimeter. Analytical thin layer chromatography was performed on 250 µM silica gel 60 F254 plates. Visualization was accomplished with UV light, and/or the use of ninhydrin, anisaldehyde and ceric ammonium molybdate solutions followed by charring on a hot-plate. Chromatography on silica gel was performed using Silica Gel 60 (230-400 mesh) from Sorbent Technologies. Analytical HPLC was performed on an Agilent 1200 analytical LCMS with UV detection at 214 nm and 254 nm along with ELSD detection. Solvents for

extraction, washing and chromatography were HPLC grade. All reagents were purchased from Aldrich Chemical Co. and were used without purification. All polymer-supported reagents were purchased from Biotage, Inc. Flame-dried (under vacuum) glassware was used for all reactions. All reagents and solvents were commercial grade and purified prior to use when necessary. Mass spectra were obtained on a Micromass Q-Tof API-US mass spectrometer was used to acquire high-resolution mass spectrometry (HRMS) data.

3,5-dichloro-*N*-((3-(3,3-dimethylbutyl)-3-azabicyclo[3.1.0]hexan-6yl)methyl)benzamide (ML218))

tert-butyl 6-((3,5-dichlorobenzamido)methyl)-3 azabicyclo[3.1.0]hexane-3-carboxylate (16): To a solution of exo-3-Boc-6-aminomethyl-3-azabicyclo[3.1.0]hexane (1.0 eq.) and triethylamine (1.0 eq.) in CH₂Cl₂ at 0 °C was added a solution of 3,5-dichlorobenzoyl chloride (1.0 eq.) in CH₂Cl₂. The reaction stirred at 0 °C for approximately 4 hrs. The mixture was then concentrated in vacuo and purified by reverse phase chromatography (MeCN/H₂O/0.1% TFA) to afford the product as a white solid (93% yield). ¹H NMR (400 MHz, CD₃OD) δ (ppm): 7.83 (d, J = 1.92 Hz, 2H); 7.66 (t, J = 1.88 Hz, 1H); 3.57 (m, 2H); 3.35 (m, 4H); 1.60 (m, 2H); 1.46 (s,9H); 0.90 (m, 1H). ¹³C NMR (125 MHz, CD₃OD) δ (ppm): 167.96, 157.61, 139.75, 137.33, 133.02, 127.99, 81.86, 43.65, 29.56, 24.46, 24.02, 23.23; HRMS calc'd for C₁₈H₂₂Cl₂N₂O₃, 385.1087 [M+H]; found 385.1089. To a vial containing tert-butyl 6-((3,5-dichlorobenzamido)methyl)-3-azabicyclo[3.1.0]hexane-3carboxylate 16, was added 4.0 M HCl in dioxane. This mixture was allowed to stir at room temp for 2 hrs. The reaction was then concentrated *in vacuo* and purified by reverse phase chromatography (MeCN/H₂O/0.1% TFA) to afford the product as a white solid (95% yield). ¹H NMR (400 MHz, CD₃OD) δ (ppm): 7.82 (d, J = 1.92 Hz, 2H); 7.68 (m, 1H); 3.45 (s, 4H); 3.34 (m, 2H); 1.90 (m, 2H); 1.21 (m,1H). ¹³C NMR (100 MHz, CD₃OD) δ (ppm): 168.04, 163.95, 139.57, 137.40, 133.17, 127.97, 43.02, 23.35, 22.48; HRMS calc'd for C₁₃H₁₄Cl₂N₂O, 285.0563 [M+H]; found 285.0566.

3,5-dichloro-N-((3-(3,3-dimethylbutyl)-3-azabicyclo[3.1.0]hexan-6-

yl)methyl)benzamide (ML218): To a solution of 3,3-dimethylbutyraldehyde (1.0 eq.) in CH₂Cl₂ was added a solution of *N*-(3-azabicyclo[3.1.0]hexan-6-ylmethyl)-3,5dichlorobenzamide (1.0 eq.) in CH₂Cl₂ followed by addition of polymer-supported sodium triacetoxyborohydride (1.2 eq.). The reaction was left to stir overnight at room temperature after which the mixture was filtered through a pad of Celite eluting with CH₂Cl₂ and concentrated *in vacuo* resulting in a crude oil which was then purified by reverse phase chromatography (MeCN/H₂O/0.1% TFA) to afford the product as an off-white solid (88% yield). ¹H NMR (400 MHz, CDCl₃) δ (ppm): 7.78 (d, *J* = 1.82 Hz, 2H); 7.57 (br, 1H); 7.42 (ap. t, *J* = 1.76 Hz, 1H); 3.64 (d, *J* = 10.56 Hz, 2H); 3.32 (t, *J* = 5.98 Hz, 2H); 2.95-2.83 (m, 4H); 1.95 (m, 1H); 1.74 (s, 2H); 1.62 (m, 2H); 0.88 (s, 9H). ¹³C NMR (100 MHz, CD₃OD) δ (ppm): 165.41, 137.46, 135.50, 131.46, 126.36, 55.69, 53.11, 41.94, 39.42, 30.07, 29.44, 21.71; HRMS calc'd for C₁₉H₂₆Cl₂N₂O, 369.1500 [M+H]; found 369.1501.

Calcium Mobilization Assays (T-type calcium channel Cell-based FDSS Ca²⁺ flux Assay)

This protocol was employed for both the high-throughput screen as well as the primary assay. Cells were routinely cultured in D-MEM/F12 (Gibco 11330-032) supplemented with 10% heat-inactivated FBS (Sigma F2442), 0.5% Sodium Pyruvate (Gibco 11360), and 1mg/ml G418 (Cellgro 30-234-CR) and then plated by the addition of 20 µl/well of 1000 cells/µl suspended in D-MEM/F12 (Gibco 11330-032) supplemented with 10% heat-inactivated FBS (Sigma F2442) and 0.5% Sodium Pyruvate (Gibco 11360). Celles were then incubated overnight at 37 °C and 5% CO₂. Medium was then removed with ELx405 and replace with 20 µl/well of Assay Buffer (140 mM NaCl, 5 mM KCl, 1 mM MgCl2, 0.5 mM CaCl2, 10 mM glucose, 10 mM HEPES pH 7.3). Add 20 µl/well Dye Loading Solution (4.6 µM Fluo-4

AM with 0.02% w/v Pluronic F-127 in Assay Buffer) with Combi, and incubated 60 minutes at room temperature (RT). Prepared 2X compound plates by transferring 80 nl/well from stock plates containing compounds at 10 mM in DMSO to compound daughter plates and add 40 µl/well Assay Buffer using Combi. Negative (vehicle) control and positive control wells are 0.2% DMSO in Assay Buffer and EC_{80} of 8 in assay buffer, respectively in every other well of columns 1–2, 23–24. Replaced dye loading solution with 20 µl/well of assay buffer using ELx405, and loaded cell plates onto Hamamatsu FDSS 6000 kinetic imaging plate reader. Then, fluorescence was measured for 10 seconds at 1Hz to establish baseline, followed by the addition of 20 µl/well test compounds from daughter plates and continue to collect data for 50 seconds at 1Hz. Plates were then removed from FDSS and incubate for 20 min at RT. Reload plates to FDSS and resume collecting data at 1Hz. After 10 seconds add 10 µl/well 5 × Stimulus Buffer (140 mM NaCl, 5 mM KCl, 1 mM MgCl2, 50 mM CaCl2, 5 mM glucose, 10 mM HEPES pH 7.3) and continue collecting data for 2 min. Calculate ratio readout as F/F_0 to normalize the data and reduce the waveform data to reflect the maximum normalized fluorescence/well. Calculate the average and standard deviation for negative and positive controls in each plate, as well as Z'.

Ca_v3.2 and Ca_v3.3 calcium channel lonWorks Electrophysiology Assays

Cav3 channel activity was examined in electrophysiological assays using the population patch clamp mode on the Ionworks Quattro (MDC, Sunnyvale, CA), an automated patch clamp instrument. The HEK293 cells stably expressing Cav3.2 and Cav3.3 channels were freshly dislodged from flasks and dispensed into a 384-well population patch clamp (PPC) plate, separately. The cell plating density was 7,000 cells/well suspended in the extracellular solution, composed of (in mM): 137 NaCl, 4 KCl, 1 MgCl₂, 1.8 CaCl₂, 10 HEPES, and 10 glucose, pH 7.4 adjusted with NaOH. After dispensing, seal resistance of cells was measured for each well and cells were perforated by incubation with 50µg/mL amphotericin B (Sigma, St. Louis, MO) on the intracellular side, which was dissolved in the internal solution composed of (in mM): 40 KCl, 100 K-Gluconate, 1 MgCl₂, 5 HEPES, 2 mM CaCl₂ pH 7.2 adjusted with KOH. The channel activity was then measured with the recording protocol as described below. Leak currents were linearly subtracted by extrapolating the current elicited by a 100-ms step to -110 mV from a holding potential of -100 mV. During the voltage pulse protocol, cells were held at -100 mV, followed by a depolarization step to -30 mV for 1 s, and then back to -90 mV. The currents were measured at the peak of the inward currents before and after the application of compounds for 3 min. Only cells with a current amplitude greater than 200 pA at -30 mV and a seal resistance >30 M Ω were included in the data analysis. Compound effects were assessed by the percentage changes in the inward currents, which were calculated by dividing the difference of current amplitude between pre- and post-compound recordings by the respective pre-compound currents in the same well. No corrections for liquid junction potentials were applied. The current signal was sampled at 0.625 kHz.

Brain slice preparation

Sprague-Dawley rats (P16-P22; Charles River, Wilmington, MA) were anesthetized with isoflorane and brains were rapidly removed from skulls and submerged in ice-cold cutting solution composed of the following (in mM), 200 glucose, 2.5 KCl, 8 MgSO₄, 0.5 CaCl₂, 1.25 NaH₂PO₄, 26 NaHCO₃, and 10 D-glucose, oxygenated with 95% O₂/5% CO₂. Parasagital brain slices (290–300 μ m) containing the STN were cut using a Vibratome 3000 (Vibratome, St. Louis, MO). Slices were incubated in oxygenated artificial cerebrospinal fluid (ACSF) at 32°C for 30 min and then maintained at room temperature afterward until transferred to a recording chamber. The recording chamber was continuously perfused with oxygenated ACSF containing (mM) 126 NaCl, 2.5 KCl, 2.0 CaCl₂, 1.3 MgSO₄, 1.25 NaH₂PO₄, 26 NaHCO₃, and 10 D-glucose.

Brain slice electrophysiology

Whole cell recordings were made from visually identified STN neurons under an Olympus BX50WI upright microscope (Olympus, Lake Success, NY). A low-power objective (4 X) was used to identify the brain region and a 40 X water immersion objective coupled with Hoffman optics was used to visualize the individual STN neurons. Patch pipettes were prepared from borosilicate glass (Sutter Instruments, Novato, CA) using a Narishige puller (model PP-830; Narishige International USA, East Meadow, NY) and had resistance of 3-4 $M\Omega$ when filled with the following intracellular solutions. For current-clamp recordings, the intracellular solution contained (in mM), 120 K-MeSO₄, 1 MgCl₂, 0.1 CaCl₂, 10 HEPES, 1 EGTA, 12 phosphocreatine, 2 Mg-ATP, and 0.4 Na₂-GTP. The pH of the pipette solution was adjusted to 7.3 with 1 M KOH, and osmolarity was adjusted to 290-295. For the voltage clamp recordings, the intracellular solution contained the following (in mM), 123 Csmethanesulphonate, 10 TEA-Cl, 2 NaCl, 10 HEPES, 0.1 EGTA, 5 QX-315, 4 Mg-ATP and 0.3 Na₂-GTP. The pH was adjusted to 7.3 with 1 M CsOH, and osmolarity was adjusted to 290–295. Whole cell current and voltage signals were amplified using an Axon Multiclamp 700B amplifier (Molecular Devices, Sunnyvale, CA). Voltage or current commands, the digitization and acquisition of membrane potentials or currents were done using Clampex 9.2/Digidata 1332 system (Molecular Devices, Sunnyvale, CA). In voltage clamp experiments, T-type calcium currents in STN neurons were isolated from voltage-gated sodium currents and high-voltage-gated calcium currents by including 0.5 uM tetrodotoxin (TTX) in the perfusate and applying the following voltage clamp protocol, a 400ms hyperpolarization prepulse to -100 mV followed by a 300 ms depolarization test pulse from a holding potential of -60 mV. The amplitude of the test pulse was determined for each cell using depolarizing voltage steps in 5 mV increments from -100 mV until maximal T-type current amplitude was observed (typically -35 to -30 mV). Leakage subtraction was performed on-line using a P/4 protocol. In current clamp experiments, low threshold spike (LTS) was elicited by the termination of a hyperpolarizing current pulse (typically -150 to -160pA, 1.5s in duration) from zero current injection. Rebound burst activity was evoked by a hyperpolarizing current pulse (typically -100pA, 1.5s in duration) followed by a depolarizing current push-pulse (+20 pA, 0.5s in duration) from zero current injection. ClampFit (Molecular Devices, Sunnyvale, CA) and Origin (OriginLab Corp, Northampton, MA) were used for data analysis. Group data were presented as mean \pm SEM. The t-test was used for statistical analysis and p < 0.05 was considered to be significant.

Drug Metabolism Methods

In vitro—Protein binding of ML218 was determined in rat plasma via equilibrium dialysis employing Single-Use RED Plates with inserts (ThermoFisher Scientific, Rochester, NY). Briefly plasma (220 μ L) was added to the 96 well plate containing test article (5 μ L) and mixed thoroughly. Subsequently, 200 µL of the plasma-test article mixture was transferred to the *cis* chamber (red) of the RED plate, with an accompanying 350 μ L of phosphate buffer (25 mM, pH 7.4) in the *trans* chamber. The RED plate was sealed and incubated 4 h at 37 °C with shaking. At completion, 50 µL aliquots from each chamber were diluted 1:1 $(50 \ \mu L)$ with either plasma (*cis*) or buffer (*trans*) and transferred to a new 96 well plate, at which time ice-cold acetonitrile (2 volumes) was added to extract the matrices. The plate was centrifuged (3000 rpm, 10 min) and supernatants transferred to a new 96 well plate. The sealed plate was stored at -20° C until LC/MS/MS analysis. The metabolism of ML218 was investigated in rat hepatic microsomes (BD Biosciences, Billerica, MA) using substrate depletion methodology (% test article remaining). A potassium phosphate-buffered reaction mixture (0.1 M, pH 7.4) of test article (1 µM) and microsomes (0.5 mg/mL) was preincubated (5 min) at 37°C prior to the addition of NADPH (1 mM). The incubations, performed in 96-well plates, were continued at 37 °C under ambient oxygenation and aliquots (80 µL) were removed at selected time intervals (0, 3, 7, 15, 25 and 45 min). Protein

was precipitated by the addition of chilled acetonitrile (160 μ L), containing glyburide as an internal standard (50 ng/mL), and centrifuged at 3000 rpm (4°C) for 10 min. Resulting supernatants were transferred to new 96-well plates in preparation for LC/MS/MS analysis. The in vitro half-life ($t_{1/2}$, min, Eq. 1), intrinsic clearance (CL_{int}, mL/min/kg, Eq. 2) and subsequent predicted hepatic clearance (CL_{hep}, mL/min/kg, Eq. 3) was determined employing the following equations:

$$t_{1/2} = \text{Ln}(2)/k;$$
 (1)

where k represents the slope from linear regression analysis (% test article remaining)

 $CL_{int} = (0.693/t_{1/2}) (rxn volume/mg of microsomes) (45 mg microsomes/gram go liver) (20^a gm of liver/kg body weight);^a scale – up factors of 20 (human) and 45 (rat) ⁽²⁾$

$$CLhep = \frac{Q \cdot CLint}{Q + CLint}$$
(3)

In vivo—Male Sprague-Dawley rats (n=2) weighing around 300g were purchased from Harlon laboratories (Indianapolis, IN) and implanted with catheters in the carotid artery and jugular vein. The cannulated animals were acclimated to their surroundings for approximately one week before dosing and provided food and water ad libitum. ML218 was administered intravenously (IV) to rats via the jugular vein catheter in 20% DMSO/80% saline at a dose of 1 mg/kg and a dose volume of 1 mL/kg. Blood collections via the carotid artery were performed at pre-dose, and at 2 min, 7 min, 15 min, 30 min, and 1, 2, 4, 7 and 24 hrs post dose. Samples were collected into chilled, EDTA-fortified tubes, centrifuged for 10 minutes at 3000 rpm (4 °C), and resulting plasma aliquoted into 96-well plates for LC/MS/ MS analysis. All pharmacokinetic analysis was performed employing noncompartmental analysis. For oral exposure studies, measuring both systemic plasma and CNS tissue exposure, ML218 was administered (oral gavage) to fasted rats (n=2) as suspensions in 10% tween 80/0.5% methylcellulose at a dose of 10 mg/kg and in a dosing volume of 10 mL/kg; blood and whole brain samples were collected at 1.5h post dose. Blood was collected into chilled, EDTA-fortified tubes, centrifuged for 10 minutes at 3000 rpm (4 °C) and stored at -80 °C until LC/MS/MS analysis. The brain samples were rinsed in PBS, snap frozen and stored at -80 °C. Prior to LC/MS/MS analysis, brain samples were thawed to room temperature and subjected to mechanical homogenation employing a Mini-Beadbeater™ and 1.0 mm Zirconia/Silica Beads (BioSpec Products). All animal studies were approved by the Vanderbilt University Medical Center Institutional Animal Care and Use Committee. The animal care and use program is fully accredited by the Association for Assessment and Accreditation of Laboratory Animal Care, International.

Liquid Chromatography/Mass Spectrometry Analysis

In vivo experiments: ML218 was analyzed via electrospray ionization (ESI) on an AB Sciex API-4000 (Foster City, CA) triple-quadrupole instrument that was coupled with Shimadzu LC-10AD pumps (Columbia, MD) and a Leap Technologies CTC PAL autosampler (Carrboro, NC). Analytes were separated by gradient elution using a Fortis C18 2.1 \times 50 mm, 3.5 µm column (Fortis Technologies Ltd, Cheshire, UK) thermostated at 40°C. HPLC mobile phase A was 0.1% NH₄OH (pH unadjusted), mobile phase B was acetonitrile. The gradient started at 30% B after a 0.2 min hold and was linearly increased to 90% B over 0.8 min; held at 90% B for 0.5 min and returned to 30% B in 0.1 min followed by a reequilibration (0.9 min). The total run time was 2.5 min and the HPLC flow rate was 0.5 mL/

min. The source temperature was set at 500°C and mass spectral analyses were performed using multiple reaction monitoring (MRM) utilizing a Turbo-Ionspray® source in positive ionization mode (5.0 kV spray voltage). All data were analyzed using AB Sciex Analyst 1.4.2 software.

In vitro experiments: ML218 was analyzed similarly to that described above (*In vivo*) with the following exceptions: LC/MS/MS analysis was performed employing a TSQ Quantum^{ULTRA} that was coupled to a ThermoSurveyor LC system (Thermoelectron Corp., San Jose, CA) and a Leap Technologies CTC PAL auto-sampler (Carrboro, NC). Chromatographic separation of analytes was achieved with an Acquity BEH C18 2.1×50 mm, 1.7 µm column (Waters, Taunton, MA).

Plasma-Brain exposure Profile PK study

ML218 was dissolved in sterile water at the concentration of 5 mg/ml (base form) and administered intraperitoneally to male Sprague-Dawley rats weighing 225 to 250 g (Harlan Sprague-Dawley, Inc., Indianapolis, IN) at the dose of 10 mg/kg. The rat blood and brain were collected at 0.5, 1, 2, 4 and 8 h. Animals were euthanized and decapitated, and the brains were removed, thoroughly washed with cold phosphate buffered saline and immediately frozen on dry ice. Trunk blood was collected in EDTA Vacutainer tubes, and plasma was separated by centrifugation and stored at -80° C until analysis. Three animals were used for each time point.

On the day of analysis, frozen whole-rat brains were weighed and homogenized in 1:3 (w/w) volumes of ice-cold phosphate buffered saline (pH 7.4). The sample extraction of plasma (100 μ l) and brain homogenate (250 μ l) was performed by a method based on protein precipitation, using three volumes of cold acetonitrile containing 0.1% formic acid and an internal standard (VU-178) having final concentration of 50 ng/ml. Extracts were vortex mixed for 5 min. followed by centrifugation at 14000 rpm for 10 min.

The supernatants of plasma and brain homogenate extracts were analyzed by means of HPLC/MS/MS, using a ThermoFinnigan TSQ Quantum Access (Thermo Fisher Scientific, Waltham, MA) mass spectrometer in positive ion mode. The chromatographic separation was achieved on an Acquity UPLC BEH C18 column (1.7um; 2.1×50mm) at a flow rate of 0.8 ml/min. The gradient program was used with the mobile phase, combining solvent A (95: 5: 0.1% formic acid in water: acetonitrile) and solvent B (95: 5: acetonitrile: 0.1% formic acid in water) as follows: 20% B for 0-0.5 min, 20-95% B for 0.5-1 min, 95% B for 1–2 min, 95–20% B for 2- 2.2 min and finally equilibration for 2.8 min before injection of next sample. The column temperature was set at 50°C. The software X calibur version 2.0 was used to control the instrument and collect data. The electrospray ionization source was fitted with a stainless steel capillary (100 µm i.d.). Nitrogen was used as both the sheath gas and the auxiliary gas. The ion transfer tube temperature was 300°C. The spray voltage, tube lens voltage, and pressure of sheath gas and auxiliary gas were optimized to achieve maximal response using the test compounds mixing with the mobile phase A (50%) and B (50%) at a flow rate of 0.8 ml/min. Collision-induced dissociation was performed on ML218 and internal standard under 1.0 mTorr of argon. Selected reaction monitoring was carried out using the transitions from m/z 320 to 148 for ML218 and m/z 310 to 223 for VU-178 (internal standard). The calibration curves were constructed and linear response was obtained in the range of 10- 2000 ng/ml by spiking known amounts of analytes in blank brain homogenates and plasma. The final PK parameters were calculated by noncompartmental analysis using WinNonlin software (version 5.1, Pharsight Inc.). Data analysis was carried out utilizing the GraphPad Prism 4.0 software package. Statistical analyses and nonlinear curve fits were carried out utilizing built in functions. Data are plotted as mean ± S.E.M. unless otherwise specified. Experiments were conducted in

accordance with the US National Institutes of Health regulations of animal care and approved by the Institutional Animal Care and Use Committee, Vanderbilt University Medical Center. Subjects were housed in groups of two per cage in a large colony room under a 12-h light-dark cycle (lights on at 6:00 a.m.) with food and water provided *ad libitum*.

Haloperidol-Induced Catalepsy

Male Sprague-Dawley rats weighing between 275 and 299 grams (Harlan Laboratories, Inc Indianapolis, IN) were used for the behavioral studies and were housed under a 12-h light/ dark cycle (lights on at 6 AM, lights off at 6 PM) with free access to food and water. The experimental protocols, which were performed during the light cycle, were approved by the Institutional Animal Care and Use Committee of Vanderbilt University and conformed to the guidelines established by the National Research Council Guide for the Care and Use of Laboratory Animals. Rats were administered haloperidol (0.75mg/kg, i.p., dissolved in 8.5% lactic acid) 60 minutes prior to vehicle (10% Tween 80 in 0.5% methylcellulose) or ML218 (0.1–30mg/kg). After an additional 30 minute pretreatment interval, all rats were assessed in the catalepsy model. Catalepsy was measured by placing the forepaws of each rat gently onto a horizontal bar 6cm from the testing surface with the body positioned at an angle of ~45° to the testing surface. The latency in seconds required for the rat to remove one or both forepaws from the bar was measured with a testing cutoff of 30 seconds.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

Acknowledgments

The authors thank the NIH (U54MH084659-01, U54MH084691, R03NS050771-01 and R01 MH078194) for support of our programs. Vanderbilt and John Hopkins are members of the Molecular Libraries Production Center Network (MLPCN) and house the Vanderbilt Specialized Chemistry Center and Johns Hopkins Ion Channel Center, respectively. Three recombinant Ca_v3 T-type channel stable cell lines were kindly provided by Dr. Edward Perez-Reyes, the University of Virginia School of Medicine.

Funding Sources

CWL receives funding from NIH, NIDA, the Alzheimer's Association, the Michael J. Fox Foundation, Seaside Therapeutics and Johnson&Johnson.

PJC receives funding from NIH, NIMH, NIDA, the Michael J. Fox Foundation, Seaside Therapeutics and Johnson&Johnson.

Abbreviations

IP	intellectual property
STN	subthalamic nucleus
HTS	high-throughput screen
PD	Parkinson's disease
VGCC	voltage-gated calcium channels
i.p	intraperioneal
p.o	oral dosing
LTS	low threshold spike

ZNS	zonisamide
GPCR	G protein-coupled receptor

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- 53. See Supporting Information for full details.



Figure 1.

First Generation T-Type Ca^{2+} Inhibitors derived from marketed antiepileptic 1 and 2, antihypertensives 3 and 4, neuroleptic 5 and antipsychotic 6 drugs.





Second Generation T-Type Ca²⁺ inhibitors 7–15 derived from optimization of HTS hits.



Figure 3. Confirmed HTS hit 16

A) Structure of 16, CID3373841; B) Raw fluorescence-based HTS assay traces; C) CRC from fluorescence-based assay (IC₅₀ = 2.5 μ M).



Figure 4.

A) Chemical optimization plan for **16**, a two-dimensional library; B) the 8×7 member library design that generated 56 analogs of **16**; C) Structure of 17 (CID85285667) the most potent analog with an IC₅₀ of 1.1 μ M in the Ca²⁺ fluorescence assay; D) Ca_v3.2 IonWorks Quattro (patch EP) CRC IC₅₀ = 13.5 μ M; E) Ca_v3.3 IonWorks Quattro (patch EP) CRC IC₅₀ = 11.5 μ M.



Figure 5. Second generation library of analogs of 17

A) 4×2 library of branched alkyl analogs; B) structure and activity of most potent analog, **18** (CID46943243); C) manual overlay of **18** with Merck's **8**; D) SurflexSim flexible alignment of **18** (grey) with Merck's **8** (tan), affording reasonable overlap except for amide HBD/HBA.



Figure 6. Initial 'Scaffold Hopping' Library

24-membered library of 3-amino piperidine 'scaffold hopping' libraries leading to **23** (CID45115608), displaying enantiospecific T-Type Ca^{2+} channel inhibition.







Figure 8. Dose-response curves for ML218 (26b)

A) Structure of ML218 (**26b**); B) Raw fluorescence-based HTS assay traces; C) CRC from fluorescence-based assay (IC₅₀ = 150 nM); D) Inhibition time-course of ML218. Ba²⁺ currents were elicited by pulse depolarized to -20mV from a holding potential of -90mV (200ms, 0.2Hz). The bath application of ML218 caused rapid inhibition and slow partially recovery after washout (n=5); E) Ca_v3.2 IonWorks Quattro (patch EP) CRC IC₅₀ = 310±15 nM; F) Ca_v3.3 IonWorks Quattro (patch EP) CRC IC₅₀ = 274±53 nM.



Figure 9.

ML218 inhibits T-type calcium currents in STN neurons.

A. Averaged traces of T-currents (lower) elicited by a voltage clamp protocol (upper) in control (*a*) and after application of 3uM ML218 (*b*) in a voltage clamp experiment. **B.** Time course of T-current amplitude before and after application of 3 uM ML218 from the same cell as in A. *a* and *b* indicate the time points at which averaged traces were taken. **C.** Bar graph summarizes group data showing ML218 inhibits T-currents in STN neurons (45.1 \pm 5.1% of the control value, n = 7, **p < 0.005). Note that the inhibition persists 20 min after washout of ML218.





Figure 10.

ML218 inhibits low threshold spike (LTS) in STN neurons.

A. Representative voltage responses to intracellular injection of a hyperpolarizing current pulse (-160 pA) in control (*a*), in the presence of 0.5 uM TTX (*b*) and combination of 0.5 uM TTX and 3 uM ML218 (*c*) from a current clamp experiment, showing the typical rebound burst firing following the termination of hyperpolarizing current pulse in control (*a*), pharmacologically isolated LTS in the presence of TTX (*b*) and inhibition of LTS by ML218 (*c*). **B.** Time course of the effect of ML218 on amplitude of LTS obtained from the same STN neuron as in A. *b* and *c* indicate the time points at which sample traces were taken. **C.** Bar graph summarizes the group data showing ML218 inhibits the amplitude of LTS (8.7 ± 2.1 mV with ML218, compared to 18.1 ± 2.2 mV in control, n = 5, ***p < 0.0001).



Figure 11.

ML218 reduces rebound burst activity in STN neurons. **A.** Representative voltage responses (upper) to intracellular injection of hyperpolarizing (-100 pA) followed by depolarizing (+20 pA) current pulses (lower) in control and after application of 3uM ML218 from a current clamp experiment. **B.** Time course of the number of rebound spikes during the depolarizing current pulse before and after application of 3 uM ML218 from the same cell as in A. *a* and *b* indicate the time points at which sample traces were taken. **B.** Bar graph summarizes the group data showing ML218 reduces the number of rebound spikes in STN neurons (5.7 ± 0.5 spikes/burst with ML218, compared to 16.0 ± 2.8 spikes/burst in control, n = 6, *p < 0.05).



Figure 12. T-type Ca²⁺ channel antagonist 8 and ML218 produce a dose-dependent reversal of haloperidol (0.75 mg/kg, i.p.)-induced catalepsy in rats

A) For comparison, the effects of increasing dose of **8** (i.p.) were compared to a top dose (56.6 mg/kg p.o.) of a previously published A_{2A} antagonist from Neurocrine. B) Comparison of the effects of increasing dose of ML218 (p.o.) to the Neurocrine A_{2A} antagonist at 56.6 mg/kg (p.o.). Catalepsy was measured as the latency to withdraw the forepaws from a horizontal bar with a cut-off of 30 s. Bar graphs represent the means \pm S.E.M. of either 6–10 rats/treatment group. (*p < 0.05 versus the vehicle control group by Dunnett's test).



Scheme 1. Synthesis of [3.1.0] analogs 26.