Behavioral/Cognitive

Disrupted Choline Clearance and Sustained Acetylcholine Release *In Vivo* by a Common Choline Transporter Coding Variant Associated with Poor Attentional Control in Humans

Eryn Donovan,¹ Cassandra Avila,¹ Sarah Klausner,¹ ©Vinay Parikh,² Cristina Fenollar-Ferrer,³ Randy D. Blakely,⁴ and ©Martin Sarter⁵

¹Department of Psychology, University of Michigan, Ann Arbor, Michigan 48109, ²Department of Psychology & Neuroscience Program, Temple University, Philadelphia, Pennsylvania 19122, ³Laboratory of Molecular Genetics, Section of Human Genetics, National Institute on Deafness and Other Communication Disorders, Bethesda, Maryland 20892, ⁴Stiles-Nicholson Brain Institute and Department of Biomedical Science, Charles E. Schmidt College of Medicine, Florida Atlantic University, Jupiter, Florida 33458, and ⁵Department of Psychology, Neuroscience Program and Department of Neurology, University of Michigan, Ann Arbor, Michigan 48109

Transport of choline via the neuronal high-affinity choline transporter (CHT; *SLC5A7*) is essential for cholinergic terminals to synthesize and release acetylcholine (ACh). In humans, we previously demonstrated an association between a common CHT coding substitution (rs1013940; Ile89Val) and reduced attentional control as well as attenuated frontal cortex activation. Here, we used a CRISPR/ Cas9 approach to generate mice expressing the I89V substitution and assessed, *in vivo*, CHT-mediated choline transport, and ACh release. Relative to wild-type (WT) mice, CHT-mediated clearance of choline in male and female mice expressing one or two Val89 alleles was reduced by over 80% in cortex and over 50% in striatum. Choline clearance in CHT Val89 mice was further reduced by neuronal inactivation. Deficits in ACh release, 5 and 10 min after repeated depolarization at a low, behaviorally relevant frequency, support an attenuated reloading capacity of cholinergic neurons in mutant mice. The density of CHTs in total synaptosomal lysates and neuronal plasma-membrane-enriched fractions was not impacted by the Val89 variant, indicating a selective impact on CHT function. When challenged with a visual disruptor to reveal attentional control mechanisms, Val89 mice failed to adopt a more conservative response bias. Structural modeling revealed that Val89 may attenuate choline transport by altering conformational changes of CHT that support normal transport rates. Our findings support the view that diminished sustained cholinergic signaling capacity underlies perturbed attentional performance in individuals expressing CHT Val89. The CHT Val89 mouse serves as a valuable model to study heritable risk for cognitive disorders arising from cholinergic dysfunction.

Key words: acetylcholine; amperometry; attention; choline transporter; s1013940; transgenic mouse

Significance Statement

Acetylcholine (ACh) signaling depends on the functional capacity of the neuronal choline transporter (CHT). Previous research demonstrated that humans expressing the common CHT coding variant Val89 exhibit attentional vulnerabilities and attenuated fronto-cortical activation during attention. Here, we find that mice engineered to express the Val89 variant exhibit reduced CHT-mediated choline clearance and a diminished capacity to sustain ACh release. Additionally, Val89 mice lack cognitive flexibility in response to an attentional challenge. These findings provide a mechanistic and cognitive framework for interpreting the attentional phenotype associated with the human Val89 variant and establish a model that permits a more invasive interrogation of CNS effects as well as the development of therapeutic strategies for those, including Val89 carriers, with presynaptic cholinergic perturbations.

Received June 27, 2021; revised Feb. 20, 2022; accepted Feb. 23, 2022.

Author contributions: E.D., C.F.-F., and M.S. designed research; E.D., C.A., S.K., V.P., and C.F.-F. performed research; R.D.B. contributed unpublished reagents/analytic tools; E.D., S.K., V.P., and C.F.-F. analyzed data; E.D., C.F.-F., R.D.B., and M.S. wrote the first draft of the paper; E.D., V.P., C.F.-F., R.D.B., and M.S. edited the paper; E.D., C.F.-F., R.D.B., and M.S. wrote the paper.

This work was supported in part by the Public Health Service Grant R01DA045063 (to M.S.), University of Michigan funds (M.S.), and the National Institute on Deafness and Other Communication Disorders National Institutes of Health

Intramural Research Funds Grant DC000039 (to Thomas B. Friedman for C.F-F.). We thank Evan Haley (Temple University) for assistance with subcellular fractionation and immunoblotting studies.

The authors declare no competing financial interests.

Correspondence should be addressed to Martin Sarter at msarter@umich.edu https://doi.org/10.1523/JNEUROSCI.1334-21.2022 Copyright © 2022 the authors

Introduction

Activation of forebrain cholinergic projection systems is necessary for a wide range of perceptual-attentional functions. In humans, disruption of acetylcholine (ACh) signaling, either by pharmacological blockade of cholinergic receptors or as a result of cholinergic cell loss, impairs the capacity to select behaviorally significant stimuli for behavioral control, to filter distractors, and to perform under dual-task conditions (Perry and Hodges, 1999; Mentis et al., 2001; Thienel et al., 2009; Yarnall et al., 2011; Albin et al., 2018; Kim et al., 2019). Results from experiments in nonhuman primates and rodents have confirmed the essential role of forebrain cholinergic activity for performing a broad range of attentional tasks (Muir et al., 1992; Voytko et al., 1994; McGaughy et al., 1996; Turchi and Sarter, 1997, 2000) and specified underlying cholinergic signaling characteristics and postsynaptic mechanisms (Parikh et al., 2007; Chen et al., 2015; Gritton et al., 2016; Howe et al., 2017; Nair et al., 2018; Laszlovszky et al., 2020; Lu et al., 2020; Sarter and Lustig, 2020; Venkatesan and Lambe, 2020; Yang et al., 2021).

The sodium-dependent, high-affinity, neuronal choline transporter (CHT) is essential for sustaining the capacity of presynaptic cholinergic neurons to synthesize and release ACh (Yamamura and Snyder, 1972; Guyenet et al., 1973; Simon et al., 1976; Ennis and Blakely, 2016). Early experiments assessing the effects of pharmacological blockade of the CHT with the competitive CHT antagonist hemicholinium-3 (HC-3), and more recent investigations into the impact of CHT heterozygosity in transgenic mice (Paolone et al., 2013b; Parikh et al., 2013), have demonstrated that reduced CHT capacity causes attenuated cholinergic signaling and broadly disrupts perceptual-attentional functions (for review, see Ferguson and Blakely, 2004; Sarter and Parikh, 2005; Bazalakova and Blakely, 2006; Sarter et al., 2016).

Okuda et al. (2002) discovered a common, single nucleotide polymorphism (SNP) within human CHT coding sequences (rs1013940) that produces the Ile89Val, or I89V, substitution in transmembrane domain 3, with \sim 8% minor allele frequency in Caucasians (1000 Genomes Project). Importantly, expression of the Val89 variant in transfected cells demonstrated reduced choline transport [40-50% of wild-type (WT) CHT] without changes in transporter surface expression or substrate affinity. We previously demonstrated that humans heterozygous for the Val89 variant exhibit attention control deficits when challenged with a distractor (Berry et al., 2014, 2015; Sarter et al., 2016). Moreover, right frontal activation during an attention-demanding task was significantly attenuated in these individuals (see also Gorka et al., 2015). Clinical significance of the expression of the CHT Val89 variant was indicated by studies showing an association with attention-deficit hyperactivity disorder (English et al., 2009) and major depression (Hahn et al., 2008).

The current experiments aimed to determine, *in vivo*, the impact of the CHT Val89 variant on cortical and striatal choline transport and cortical ACh release when introduced into mouse CHT coding sequences using CRISPR/Cas9 methods. Using these mutant mice and their Ile89 (WT) littermates, we establish a significant reduction in the capacity of Val89-expressing mice to clear choline from the extracellular space. Furthermore, successive, depolarization-evoked ACh release events, induced at a frequency mirroring cholinergic activity in attention task-performing rodents, led to a robust attenuation of ACh release in the mutant mice. These cholinergic capacity limits by the CHT Val89 variant were not associated with altered surface expression of the transporter. Consistent with attentional control deficits seen in Val89-expressing humans, a visual disruptor revealed

that WT mice, but not Val89 mice, deployed attentional control mechanisms to perform a continuous signal detection task. Structural modeling studies suggested that the Val89 substitution reduces choline transport by potentially disrupting the architecture of the CHT close to the substrate binding site. Our findings provide a neurobiological and cognitive framework for the inattentive phenotype that characterizes human Val89 carriers.

Materials and Methods

Subjects

Mice were generated to express a valine rather than the WT isoleucine residue at amino acid 89 in the endogenous Slc5a7 gene that encodes the CHT. Briefly, a CRISPR/Cas9 approach (Cong et al., 2013) was implemented through the Transgenic Mouse Embryonic Stem Cell Shared Resource facility at Vanderbilt University. An RNA guide expressing construct was developed by ligation of a 193 bp protospacer into the Bbsl sites of the pX330 plasmid (http://www.addgene.org/42230/) that also affords expression of human codon optimized Cas9. The protospacer contained an 82-nucleotide arm 5' of the codon (ATT) that encodes Ile89 in exon three and a 93-nucleotide arm 3' of the site in intron 4 (HDR oligos). An 18mer donor oligonucleotide (5'CCGTGGGCTA TTCTCTGAGTC3') was generated to encode a Val89 codon along with the introduction of a synonymous codon change in the adjacent codon (GGA to GGC), designed to reduce the chance of repeated CRISPR activity on the edited sequence. Base changes also afford diagnostic restriction digestion (novel Hpy99I site) to ascertain the presence of the sequence change introduced. Cas9/gRNA plasmid and donor oligonucleotide were injected into single cell C57BL6/J mouse embryos subsequently implanted into a pseudopregnant female mouse. After weaning, tissue from tail samples of pups were genotyped by restriction digest, followed by DNA sequencing of PCR fragments spanning the mutation, cloned into pGEMTEasy (Promega). From three different litters, three animals were identified as heterozygous founders carrying the intended codon mutation without other deletions apparent. Two of these founders produced pups carrying the Val89 allele, one of which served as the founder for the line studied. These animals were subsequently crossed to WT C57BL6/J mice for eight generations before maintaining the colony as homozygous or heterozygous breeders.

The data from a total of N = 117 mice (n = 38 CHT^{Ile/Ile} mice, henceforth termed WT, 18 females; n = 38 heterozygous CHT^{Ile/Val} mice, henceforth termed CHT I/V, 18 females; and 41 homozygous CHT^{Val/Val} mice, henceforth termed CHT V/V, 17 females) were included in the final analyses. Following arrival at the University of Michigan, the CHT Val89 colony was further crossed to C57BL/6J mice for an additional four generations. Mice were housed in a temperature and humidity-controlled environment (23°C, 45%) with a 12/12 h light/dark cycle (lights on at 7 A.M.). Data collection for all experiments began during the light phase. Litters were weaned at three weeks and group-housed with same-sex siblings for a maximum of six animals per cage. Genomic DNA from tail tissue samples was used to determine mouse genotype as either WT, CHT I/V, or CHT V/V by Transnetyx. Animals were given food (Envigo Teklad rodent diet) and water ad libitum. Mice were a minimum of eight weeks old at the beginning of experiments. All procedures were conducted in adherence with protocols approved by the Institutional Animal Care and Use Committee (IACUC) at the University of Michigan.

Mice used in individual experiments

Amperometric recordings of *in vivo* choline clearance were used to quantify the impact of the Val89 variant on CHT uptake efficiency under various conditions. Initial experiments determined exogenous choline clearance in cortex of WT (n=4, 3 females), CHT I/V (n=4, 1 female), and CHT V/V (n=5, 2 females) mice, and in striatum of WT (n=4, 2 females), CHT I/V (n=3, 2 females), and CHT V/V (n=3, 2 females) mice. A total of 15 additional mice (6 females) were used to determine choline clearance in the presence of the CHT competitive inhibitor HC-3 (n=5 WT, n=4 CHT I/V, n=6 CHT V/V; 2 females for each

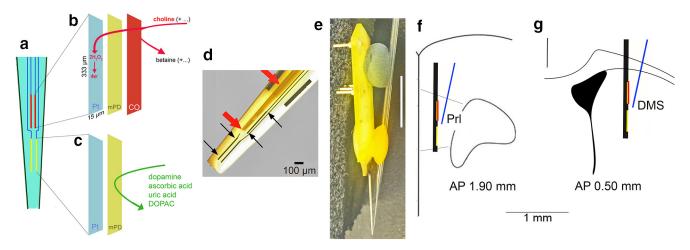


Figure 1. Diagram of the choline measurement scheme, the glass capillary attached to the electrode for pressure ejections, and placement of the assembly in the medial frontal cortex. a, Ceramic backbone with the four platinum-iridium (Pt/Ir) recording sites, each 333 μm long and 15 μm wide, organized in two pairs (see text for references). The upper pair was used to measure choline currents (red; see also b), while the lower pairs served as sentinels (yellow; see also c). The choline-sensing scheme is illustrated in abbreviated form in b, c. Choline is oxidized by choline oxidase (CO) that was immobilized onto the upper pair of recording sites. The resulting current, over background activity, was converted to choline concentration (μ M). Before immobilization of CO, the nonconducting polymer m-(1,3)-phenylenediamine (mPD) was electroplated onto the recording sites to block the transfer of small electroactive organic molecules to the Pt site. Importantly, such mPD films do not block small species such as H_2O_2 from reaching the Pt surface (Huang et al., 1993; Mitchell, 2004). The lower pair of recording sites (c) were coated with mPD but not CO and thus served to record the concentration of electroactive interferents, which, despite the mPD barrier, still reached the Pt sites. A top view of the electrode in d shows the Pt recording sites (black arrows), with the upper left Pt site situated underneath a glass capillary (red arrows) used to pressure eject compounds into the recording area (100-μm scale inserted). The entire electrode/glass capillary assembly is shown from the side in e. Note that the Pt recording sites are not visible because of low resolution (1-cm scale vertically inserted on the right) but are located in the very tip of the ceramic backbone (see also the tip of the glass capillary). For the present experiments, the electrode/glass capillary assembly was inserted into the medial frontal cortex or the dorsomedial striatum of mice. f, g Coronal section of the frontal co

genotype). In separate mice (n = 6, 2 per genotype, 1 female per genotype), the contribution of diffusion rate to overall choline clearance was quantified in the presence of lidocaine HCl to arrest transporter activity and, more broadly, neuronal firing (Malpeli and Schiller, 1979; Chu and Lee, 1994; Tehovnik and Sommer, 1997). To illustrate the efficacy of lidocaine, we also determined endogenous ACh release in the presence of lidocaine in one WT male mouse. To demonstrate attenuated ACh release following repeated potassium-evoked depolarizations, a total of 15 mice were used (WT, n = 4, 2 females; CHT I/V, n = 6, 3 females; CHT V/V, n = 5, 3 females). A group of 30 mice (equal group sizes and sex distribution across genotypes) were used to assess CHT densities in total synaptosomal lysates and synaptosomal plasma membraneenriched fractions. Data from 27 mice (WT, n = 8, 3 females; CHT I/V, n = 9, 4 females; CHT V/V, n = 10, 2 females) that reached the acquisition criterion in a continuous signal detection task were used to characterize their attentional phenotype.

Amperometric recordings of choline currents

The experiments described herein measured choline concentrations in the extracellular space. We determined the capacity of the CHT by assessing the effects of pressure ejections of choline on CHT-mediated choline clearance. The capacity of cholinergic neurons to release ACh was determined by assessing the effects of repeated KCl-induced depolarization on extracellular choline concentrations. Prior experiments validated the use of choline-sensitive microelectrodes to monitor ACh release in real time, in part by demonstrating that, *in vivo*, endogenous acetylcholinesterase (AChE) completely hydrolyzes new released ACh (Parikh et al., 2004, 2007, 2013; Parikh and Sarter, 2006; Giuliano et al., 2008).

Choline concentrations were measured using an amperometric measurement scheme, illustrated in Figure 1. In this approach, choline oxidase (CO) is immobilized onto the surface of an electrode to oxidize choline, resulting in a by-product of hydrogen peroxide. Electrochemical oxidation of hydrogen peroxide then results in the release of two electrons, which generate the current that is translated into the concentration of choline at the recording site based on prior *in vitro* calibrations (Burmeister et al., 2000, 2003).

The construction of choline sensors and the amperometric measurement scheme were detailed previously (references above). Briefly, four platinum-iridium (Pt/Ir) recording sites (15 × 333 μ m) were arranged in two pairs positioned linearly along the shank of the ceramic backbone probe (Quanteon). The members of a pair of recording sites were separated by 30 μ m, and pairs were separated along the vertical axis by 100 μ m (Fig. 1).

An exclusion layer of m-(1,3)-phenylenediamine (mPD) was electroplated onto the surface of each recording site by applying a constant voltage of 0.85 V versus an Ag/AgCl reference electrode (Bioanalytical Systems) for 5 min to prevent sensing of electroactive interferents such as ascorbic acid (AA) and catecholamines. After allowing 24 h for sites to dry, the top pair of electrodes, further named choline sensors, were coated with a solution of CO (Sigma-Aldrich) cross-linked with a BSA and glutaraldehyde mixture by manual application of microdroplets using a 1- μ l Hamilton syringe. The lower pair of sites were coated with only the BSA and glutaraldehyde mixture to identify currents unrelated to the presence of choline (hence named "sentinels"). Following a 48-h minimum incubation period to allow for optimal adherence of the enzyme layer, recording sites were calibrated to determine the sensitivity for choline, selectivity for choline versus interferents, and limit of detection of choline. Calibrations were performed using a FAST-16 electrochemical system (Quanteon) to apply a constant voltage of 0.7 V versus an Ag/AgCl reference electrode in a heated bath of 40 ml 0.05 M PBS. After recording a 30-min baseline, 20 mm AA (interferent), 20 mm choline (analyte), and 2 mm dopamine were added to the bath to generate the following concentrations: 250 μ M AA, 20, 40, and 60 μ M choline, and 2 μM dopamine. Changes in amperometric current recorded at the individual electrode sites following the addition of each solution were used to calculate site sensitivity, selectivity, and limit of detection. In order to qualify for use in in vivo experiments, choline sensors were required to have a sensitivity of >5 pA/ μ M, choline:AA selectivity ratio >80:1, and a limit of detection $< 0.9 \mu M$ choline.

Probes that passed the above calibration criteria were prepared for *in vivo* experiments. Glass capillaries $(1.0 \times 0.58 \text{ mm})$, with filament; A-M Systems) were pulled using a micropipette puller (Model 51 210, Stoelting), and the tips were clipped until the inner diameter was ~15 μ m. Pipettes were immobilized to the probe

surface using clay and wax such that the tip of the pipette was positioned on the mid-line between the upper and lower site pairs, $\sim\!50\,\mu\mathrm{m}$ from the probe surface (Fig. 1e-g). Micropipettes were filled with one of the following solutions immediately before surgery: 5 mm choline, 5 mm choline and 10 $\mu\mathrm{m}$ HC-3, 20 $\mu\mathrm{g}/\mu\mathrm{l}$ lidocaine HCl, or either 70 or 120 mm KCl. All drug solutions were prepared in DI water.

In vivo amperometric recordings

Mice were anesthetized using isoflurane gas (4% induction and 1-2% maintenance) and mounted in a stereotaxic frame on an isothermal heating pad to maintain a 37°C body temperature. Eyes were lubricated with ophthalmic ointment. A craniotomy and durotomy were performed above the right medial frontal cortex (AP: +1.9 mm; ML: -0.5 mm from bregma), or above the right dorsomedial striatum (AP: +0.50 mm; ML: -2.00 mm from bregma), using a bent 27-gauge needle for dura removal. The prepared microelectrode probe was then lowered 2.0 or 2.5 mm dorsoventrally from the top of the brain into cortex or striatum, respectively, while an Ag/AgCl reference electrode was implanted at a remote site in the contralateral hemisphere. Amperometric recordings were achieved using a FAST-16 electrochemical system and digitized at a sampling rate of 10 Hz. Pressure ejections began following a 60-min baseline period. Choline sensing was accomplished through CO catalyzation of the conversion of choline into hydrogen peroxide and glycine betaine. The hydrogen peroxide by-product was oxidized electrochemically via FAST application of 0.7 V versus the Ag/AgCl reference electrode. The resulting current, over background activity, was previously demonstrated to reflect either choline derived from newly released ACh and hydrolyzed by endogenous AChE (references above), or choline concentrations resulting from pressure ejections of choline into the recording area. In the latter case, diminishing choline currents were shown to largely reflect clearance by the CHT, as indicated by the attenuation of clearance efficiency following CHT blockade with HC-3 (Parikh et al., 2006, 2013; Parikh and Sarter, 2006).

Choline clearance

We determined the impact of the Val89 variant on CHT-mediated clearance of choline pressure-ejected into the recording field. Micropipettes were filled with a 5 mm choline solution and pressure ejections of 10–35 nl (corresponding to a recorded amplitude of 10–35 $\mu{\rm M}$ choline) were administered at least 1 min apart to limit intertrial interference on choline clearance. The first three current traces within the 10–35 $\mu{\rm M}$ range were used for analysis.

In separate mice, exogenous choline clearance in the presence of HC-3 (Sigma-Aldrich) was determined to test the relative contribution of heterozygous and homozygous CHT Val89 expression, as compared with the expression of the CHT Ile89, to choline clearance. Micropipettes were filled with a cocktail of 5 mm choline and 10 μ m HC-3. Pressure ejections were separated by a minimum of 2 min, and the first three ejection traces with a peak amplitude between 10 and 35 μ m were used in subsequent analyses to avoid aggregating effects of serial HC-3 ejections. Clearance data from previous sessions in which only 5 mm choline was ejected were used as a baseline measure of comparison for all analyses regarding the inhibitory effects of HC-3.

In a third experiment, to quantify total voltage-dependent choline clearance, exogenous choline clearance was tested before and after administrating the voltage-gated sodium channel blocker, lidocaine HCl (Hospira Inc.). In addition to inhibiting neuronal activity (Malpeli and Schiller, 1979; Tehovnik and Sommer, 1997) that supports CHT surface trafficking (Ferguson et al., 2003), lidocaine-induced increases in sodium concentrations (Onizuka et al., 2008), and intracellular proton levels disrupt CHT function (Chu and Lee, 1994; Iwamoto et al., 2006). For this experiment, two micropipettes, one filled with 5 mm choline and the other filled with $20\,\mu\text{g}/\mu\text{l}$ lidocaine HCl, were immobilized to the probe. Recording sessions consisted of two baseline pressure ejections of 5 mm choline, followed by a single 0.5- μ l infusion of lidocaine HCl given over 3 min, a 2-min rest period for drug diffusion, and finally, a second series of two postlidocaine choline ejections (Malpeli and Schiller, 1979).

These parameters were also used in an experiment designed to verify the efficacy of lidocaine by measuring effects on depolarization-induced release of endogenous ACh. In one male WT mouse, prelidocaine and postlidocaine pressure ejections of KCl (70 mm; 40–60 nl) were used to generate ACh release events.

ACh release capacity following repeated depolarization

A final amperometric experiment determined the impact of the Val89 variant on ACh release following repeated depolarization. Repeated series of pressure ejections of potassium chloride (KCl; 120 mm; 30–40 nl per pressure ejection) were used to induce low-frequency, behaviorally-relevant episodes of ACh release. Choline currents following KCl-induced depolarization have previously been shown to reflect the hydrolysis of newly released ACh (Parikh et al., 2004; Parikh and Sarter, 2006; for further justification and additional references, see Results). Each depolarization series consisted of five evenly distributed KCl ejections delivered over 2 min, with an ejection delivered every 24 s. Two depolarization series, separated by 15–20 min, were followed by a 5- and a 10-min pause, after which a single KCl-induced ACh release was measured (see timeline in Fig. 5a).

Analysis of choline currents

Choline currents were calculated by subtracting the background current recorded on sentinel sites from the choline-specific current recorded on choline sensors. Current was then converted to choline concentration based on calibration data. Choline clearance capacity was analyzed by calculating the slope of current decline from 40% to 80% of the peak choline concentration (Slope₄₀₋₈₀). Slope₄₀₋₈₀ was previously shown to be maximally sensitive to CHT inhibition by HC-3, indicating that this measure primarily reflects choline clearance via CHT-mediated transport (Parikh and Sarter, 2006). To enhance the illustration of the effects of HC-3 and lidocaine across the three genotypes, choline clearance traces were normalized by dividing individual values by the peak concentration obtained from the individual trace. Statistical analyses of clearance rates (Slope₄₀₋₈₀) were always based on values extracted from raw traces.

Determination of synaptosomal CHT density

Synaptosome preparation

Urethane-anesthetized mice were decapitated, and frontal cortices were dissected on ice immediately. Isolated tissues were homogenized in ice-cold 0.32 M sucrose and centrifuged at $1000 \times g$ for 4 min at 4°C to remove cellular debris. Next, the supernatant was centrifuged at $12,500 \times g$ for 15 min to yield a crude synaptosomal pellet. Pellets were stored at -80°C until fractioned.

Isolation and quantification of plasma membrane-enriched fractions In addition to the determination of CHT density in total synaptosomal lysates, subcellular fractionation was conducted to isolate CHTs in a plasma membrane-enriched fraction (Ferguson et al., 2003; Parikh et al., 2006, 2013). Briefly, the synaptosomal pellet was lysed in 5 mm HEPES-NaOH, pH 7.4, containing a cocktail of protease inhibitors (1.0 μ g/ml leupeptin, 1.0 μ g/ml aprotinin, 1.0 μ g/ml pepstatin, and 250 μ g/ml phenylmethylsulfonyl fluoride). The fraction was collected by spinning the lysate at $15,000 \times g$ for 20 min. Proteins were extracted from each fraction with 5 mm HEPES-KOH (pH 7.2) solution containing 1% SDS, 1 mm EDTA, 1 mm EGTA, and protease inhibitor cocktail. Protein concentrations were determined by using a modified Lowry Protein Assay (Pierce). Equal quantities (25 μ g) of protein from each fraction were subjected to immunoblot analysis. Proteins were separated on 4-15% Tris HCl polyacrylamide gels and transferred to PVDF membranes. Immunodetection of CHT bands was accomplished by incubating the membraned overnight with 1:2000 diluted rabbit anti-CHT polyclonal antibody (ABN458; Millipore). The membranes were then exposed to peroxidase-conjugated anti-rabbit secondary antibody and SuperSignal West Femto Maximum Sensitivity Substrate (ThermoFisher Scientific). The resulting chemiluminescent signal was acquired with a Chemidoc Touch Imaging System (Bio-Rad). Densitometric analysis of CHT-immunoreactive bands was performed by calculating the integrated pixel densities

using NIH ImageJ software. The membranes were stripped for the detection of β -actin in all samples that served as a control to accommodate any differences in the sample loading during gel electrophoresis. CHT densities were normalized to the levels of β -actin-immunoreactive bands for each sample analyzed. Total synaptosomal CHT density was determined in frontal cortical tissue from six mice (three females) per genotype. CHT density in the plasma membrane-enriched fraction was measured in tissues from 10 mice (five females) per genotype.

Disruptor-challenged attentional performance

Subjects

A total of $N\!=\!45$ mice, aged 12 weeks at the beginning of experiments, began acquisition training of the task. The final analyses were based on mice that reached asymptotic performance in the unchallenged version of the task and completed an additional sequence of sessions that included sessions during which the disruptor was presented.

Water restrictions and body weights

Training and testing took place Monday–Friday between 11 A.M. and 3 P.M. Daily training of this task was previously shown to invoke a diurnal activity pattern in rodents (Gritton et al., 2012, 2013; Paolone et al., 2012). Mice were weighed weekly. Before water deprivation, body weights ranged from 20 to 29 g (mean: 24.96 g). Beginning 5 d before the onset of training, water access was progressively reduced from *ad libitum* to 4 min/d. Water restrictions were adjusted individually throughout the experiment to maintain 90% of the mice's original body weight, resulting in 3- to 6-min access to water after completion of the session. During weekends, mice were given unrestricted water access until Sunday 12 P.M.

Task training and testing

The continuous signal detection task was previously characterized for the assessment of sustained attention in rats and humans (McGaughy and Sarter, 1995; Demeter et al., 2008) and subsequently adopted for testing mice (St Peters et al., 2011). Training and testing took place in eight operant chambers equipped with two center panel lights- the bottom light serving as the signal source-a water dispenser, two moveable nose poke devices designed and built in-house (MICARPs; St Peters et al., 2011), and a house light (for technical specifications, including chamber dimensions, exact position and types of lights, illuminance measures, and the position of MICARPs, see St Peters et al., 2011).

Mice were trained to perform this task using the following steps. (1) Mice were initially trained to retrieve water from the water port (water was dispensed every 30 s for a single 20-min session). (2) Both MICARPs were extended into the operant chamber and remained extended throughout the entire 40-min session. All nose-pokes resulted in the reward of water (6 µl per reward). To pass this stage, mice were required to generate 20 rewards per session for three consecutive sessions. (3) Both MICARPs extended into the operant chamber and remained extended for 20 s, or until a nose-poke triggered retraction of both, with an intertrial interval (ITI) of 20 s. To pass this stage, mice must have retrieved 20 rewards per session for three consecutive sessions. (4) Both MICARPs extended into the operant chamber and remained extended for 4 s, or until a nose-poke triggered retraction, with an ITI of 12 ± 3 s (criterion as in 3). (5) Mice were trained to discriminate signals (illumination of central panel light) from nonsignals (no illumination). Half of the mice were trained to report the presence of a signal with a left nose-poke, while the other half with a right nosepoke. Each 40-min session consisted of 160 trials (80 signals and 80 nonsignals). One second after an event (signal or nonsignal), MICARPs were extended into the chamber. In signal trials, the light remained illuminated until a nose-poke occurred or for 4 s (omission). Correct responses (hits, correct rejections) trigger water delivery, incorrect responses (misses, false alarms) triggered the onset of the ITI. This training stage also included forced correction trials. An incorrect response forced repetition of the trial. Three consecutive errors triggered a trial in which only the correct MICARP was extended, to suppress the manifestation of a side bias. To pass this stage, mice were required to score >60% hits and correct rejections. (6) Signal duration was shortened to 1 s, and the delay between signal/nonsignal onset and MICARP extension was reduced to 0.5 s (criterion as in 5). (7) Correction trials were removed, and variable signal durations were implemented (500, 50, 25 ms) to prevent the manifestation of a fixed discrimination threshold (pass criterion: >60% hits to longest signals, >60% correct rejections, and <33% omissions for three consecutive sessions). (8) The house light in the back panel of the chamber was illuminated throughout the session. This is a crucial step for assessing sustained attention as it requires mice to continuously orient to and monitor the signal source (criterion as in 7).

Visual disruptor

Presentation of a visual disruptor disrupts performance and causes reward loss, thereby potentially triggering the implementation of top-down attentional control mechanisms to stabilize and recover performance (Demeter et al., 2008, 2013; Demeter and Woldorff, 2016). Performance of the 40-min sessions was analyzed based on five 8 min trial blocks, and the disruptor (house light and panel light above the signal light flashing asynchronous at $0.5 \, \text{Hz}$) was presented during the second and third 8-min block. Following a disruptor session, mice were required to regain unchallenged criterion performance for three consecutive sessions before being eligible for additional disruptor sessions. Sessions in which predisruptor performance reached chance level were excluded from final analysis. Neither the number of disruptor sessions excluded in the final analyses (median: 1.00, range: 3.00; Kruskal–Wallis ANOVA; $H_{(2)} = 0.84$, p = 0.66) nor the number of sessions included (median: 2.00, range: 4.00; $H_{(2)} = 0.34$, p = 0.95) differed by genotype.

Measures of performance

The percentage of hits, misses, correct rejections, false alarms, and omissions, for each 8 min trial block, were calculated. Hit and false alarm rates were used to compute signal detection theory-derived measures of perceptual sensitivity (d') and response bias (B" $_D$; Green and Swets, 1974; Donaldson, 1992), as detailed previously (Demeter et al., 2008; Phillips and Sarter, 2020). The bias scores range from -1.0 to +1.0 with a score of zero depicting no bias, negative scores a liberal bias, that is, a bias toward reporting the presence of a signal, and a positive score indicating a more conservative bias, that is, a bias toward reporting the absence of a signal.

Structural modeling of the mSLC5A7 WT and I89V substitution

To identify a suitable structural template for mouse CHT we used Hidden Markov model (HMM) profiles as a descriptor for the transporter's amino acid sequence. The HMM profile obtained after a three-iteration sequence scanning performed using UniRef30 sequence database was subsequently scanned against the HMMs of the sequences corresponding to each of the Protein Data Bank (PDB) x-ray structures (pdb70 database) as per standard procedure in HHpred server (Hildebrand et al., 2009; Zimmermann et al., 2018). This method has been shown to increase the successful identification of templates by 30%, especially for distantly related sequences. The Vibrio parahaemolyticus sodium/sugar symporter vSGLT (PDB id: 3DH4; Faham et al., 2008) had the highest coverage (422 residues), the highest sequence identity (19%), and better correspondence between secondary structural elements and, as consequence, was selected as template during the structural modeling procedure. To further determine the architectural similarity of CHT and vSGLT we analyzed their hydrophobicity profiles. The hydrophobicity profiles for these proteins were constructed and aligned using AlignMe server (Khafizov et al., 2010; Stamm et al., 2013, 2014) and applying the Hessa et al. (2005) hydrophobicity scale and a 13-residue-long window.

The initial sequence alignment between vSGLT and CHT obtained from HHpred was modified to align the first transmembrane helix (TM1) of vSGLT with that predicted in CHT by Topcons server (Tsirigos et al., 2015). The resulted alignment was then refined in an iterative process that used conservation scores obtained with Consurf server (Ashkenazy et al., 2016) as a guide. This procedure positions the most conserved residues packing toward the core of the protein and removes

gaps within secondary structural elements when needed. The final alignment was used during the modeling production run where 2000 modeling iterations were performed with MODELLER (Webb and Sali, 2016). The model with the highest MODELLER probability distribution function score (molPDF), best ProQM score (Ray et al., 2010) and Procheck analysis (Laskowski et al., 1993) was selected as the final WT CHT model. A similar procedure was subsequently used to obtain the structural model of CHT Val89.

Experimental design and statistical analysis

Choline clearance capacity was assessed using the rate (μ M/s) of choline concentration decline from 40% to 80% of the choline transient peak (Slope₄₀₋₈₀). An initial ANOVA of the effects of genotype was conducted to demonstrate that pressure ejections of choline produced similar peak amplitudes in the three strains. A two-factor mixed model ANOVA was used to determine genotype differences, and the effects of repeated (3) pressure ejections of choline, on choline clearance rate for choline-only ejection experiments. Two-way ANOVAs were used to test the effect of CHT inhibitor, HC-3, and lidocaine administration, on choline clearance rates (Slope₄₀₋₈₀).

The effects of repeated depolarizations on evoked ACh release were analyzed using one-way and mixed model ANOVAs (for experimental manipulations and timeline, see Fig. 5a). In each case, results from Shapiro-Wilks tests of normality and Levene's equality of variance supported the use of parametric analyses. Effects of the very first pressure ejection of potassium on absolute peak amplitudes of choline currents in the three strains were analyzed using one-way ANOVA. The effects of subsequent KCl ejections were expressed as a ratio of the effects of the first ejection. The effects of genotype on the peak amplitudes of currents evoked by subsequent depolarizations (second to fifth depolarization) were analyzed first by mixed model ANOVAs on the effects of genotype and KCl series (first vs second) separately for the second and fifth KCl pressure ejection (with α set at 0.05/2). Based on the absence of significant effects from these two analyses, individual means over the two KCl series were analyzed for the effects of genotype and KCl ejection number (second vs fifth). Peak amplitudes of currents recorded during the recovery tests, 5 and 10 min after the depolarization series, were expressed relative to the peak amplitudes of the first KCl-induced choline current of that series. Effects of genotype and series were analyzed using mixed model ANOVA, once again supported by the results of Shapiro-Wilk and Levene's equality tests. CHT densities in total synaptosomal lysates and plasma membrane-enriched fractions were compared between the genotypes using one-way ANOVA. To determine effects on the performance in the continuous signal detection task, signal detection theory-derived parameters of perceptual sensitivity and bias were computed (see above). Mixed model ANOVAs determined effects of genotype and signal duration on unchallenged performance, and effects of trial block and genotype on the performance in the presence of a disruptor. To determine changes in bias in response to the disruptor, the difference between scores obtained during disruptor and postdisruptor trial blocks (2-5) were subtracted from scores indicating bias during the first predisruptor trial block. For each mouse, these differences were averaged across disruptor sessions and analyzed to determine effects of genotype and trial block using ANOVA. Finally, effects of task block and genotype on errors of omissions were determined for both task conditions.

For parametric mixed-model ANOVAs, in case of violation of the sphericity assumption, Huynh–Feldt-corrected F values, along with uncorrected degrees of freedom, are given. Tests used for $post\ hoc$ multiple comparisons are identified in figure legends. Data graphs depict individual values, means and 95% confidence intervals (CIs). Statistical analyses were performed using SPSS for Windows (version 17.0; SPSS) and GraphPad Prism version 8.2.1. with α set to 0.05. Exact p values were reported (Greenwald et al., 1996; Sarter and Fritschy, 2008; Michel et al., 2020). For major results derived from parametric tests, effect sizes (Cohen's d) were indicated (Cohen, 1988).

Results

Impact of Val89 on CHT-mediated choline clearance

Reduced clearance of exogenous choline

These experiments were designed to determine, *in vivo*, the capacity of the CHT in the cortex and striatum of CHT I/I, I/V, and V/V genotypes (see Fig. 1 for an illustration of the measurement scheme and the placement of the electrode/ glass capillary in cortex and striatum). The clearance of exogenous choline was determined by analyzing the component of the clearance curve, $Slope_{40-80}$, that was previously demonstrated to reflect primarily CHT-mediated transport (for details and references, see Materials and Methods).

Following pressure ejections of 5 mm choline into the prelimbic recording region, peak choline currents were reached within 1–2 s, followed by rapid decreases in current (Fig. 2a). Peak currents did not differ between genotypes ($F_{(2,36)} = 1.58$, p = 0.22; Fig. 2b; note that while currents were recorded at 10 Hz, and analyses were based on 10-Hz data, for clarity, the figures show data at a 2-Hz sampling rate).

A mixed-model ANOVA on the effects of genotype and repeated pressure ejections on normalized choline clearance (Slope₄₀₋₈₀) indicated a main effect of genotype ($F_{(2,36)} = 6.91$, p = 0.01, Cohen's d = 2.35) and neither an effect of repeated pressure ejections of choline nor a significant interaction between the two factors (main effect: $F_{(2,36)} = 0.22$; interaction: $F_{(4,36)} = 0.62$; both p > 0.41). Multiple comparisons (LSD) indicated significantly slower decay rates in CHT I/V and CHT V/V mice when compared with WT mice (p < 0.012), but no gene dosage effect (p = 0.83). Figure 2c illustrates that decay rates in CHT Val89 mice on average were reduced by over 77% (CHT I/V) and 83% (CHT V/V) relative to WT mice. The low residual decay rates in CHT I/V (25th percentile: 0.7 μ M/s; 75th percentile: 1.6 μ M/s) may have prevented the demonstration of even lower decay rates in CHT V/V mice. This issue was further addressed by the experiments which assessed effects of HC-3 and lidocaine (below).

Similarly, striatal choline clearance rates in CHT Val89 mice were significantly lower than in WT mice ($F_{(2,27)} = 8.94$, p = 0.001, d = 1.63). Furthermore, as also was the case in cortex, multiple comparisons (Fig. 2f) did not indicate a significant gene dose effect.

Following pressure ejections of choline into the dorsomedial striatum, inspection of the averaged choline current traces (Fig. 2d) suggested lower peak currents in CHT V/V than in WT and I/V mice. However, peak amplitudes did not differ across genotypes ($F_{(2,27)} = 2.37$, p = 0.11; Fig. 2e). The averaged traces shown in Figure 2d suggested longer lasting peak levels in CHT Val89 mice when compared with WT mice. We therefore conducted a post hoc analysis of the timing of peak currents. Currents peaked between 1.8 and 2.1 s after pressure ejections and peak times did not differ across genotypes ($F_{(2.27)} = 0.39$, p = 0.68). However, peak times were considerably more variable in CHT Val89 when compared with WT mice (SD: WT: 0.38 s I/V: 0.98 s; V/V:1.46 s). Multiple F tests for the equality of two variances (at $\alpha/3$) confirmed that peak current variances in CHT Val89 mice were significantly larger than in WT mice (WT vs I/V: $F_{(11,8)}$ = 6.70, p = 0.005; WT vs V/V: $F_{(11,8)} = 14.97$, p < 0.001; I/V vs V/V: $F_{(8.8)} = 2.23$, p = 0.28). Thus, the seemingly lower levels of averaged peak currents in CHT Val89 mice seen in Figure 2d did not reflect lower peak levels (Fig. 2e), but that the timing of current peaks was distributed over significantly longer periods of time, therefore lowering averaged values. The reason for this effect of genotype in the striatum is unknown. Because of our focus on

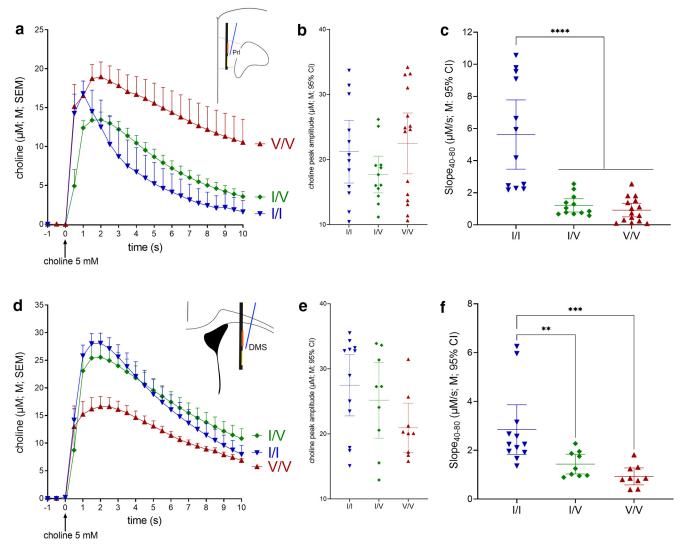


Figure 2. Impact of CHT Val89 on clearance of exogenous choline in mouse prelimbic (Prl) cortex (a–c) and dorsomedial striatum (DMS) (d–f, for illustrative clarity, choline currents, recorded at 10 Hz, are shown at 2 Hz in a and d). Following insertion of the electrode and glass capillary into the brain (Fig. 1) and a 60-min baseline period, repeated pressure ejections of 5 mm choline (10–35 nl) were given at 1- to 2-min intervals. a, Choline ejections occurred at time 0; a and d show mean (M) and standard error of the mean (SEM). b, Peak choline currents did not differ across genotypes (ANOVAs are described in Results; b,c,e and f show individual values, means and 95% confidence intervals, Cl). e, Cortical choline clearance rates were calculated for the periods of current decay from 40% to 80% (Slope_{40–80}) of peak amplitudes (μ m/s; WT, inverted blue triangles, 12 ejections from 4 mice; CHT I/V, green diamonds, 12 ejections from 4 mice; CHT V/V, red triangles, 15 ejections from 5 mice). Clearance rates in both I89V-expressing strains were markedly reduced relative to WT mice (LSD test). Choline clearance in the striatum (d–f) likewise was significantly suppressed in I89V-expressing mice (WT, 4 mice with 3 ejections each; CHT I/V and CHT V/V mice, 3 mice with 3 ejections each). As averaged peak levels in d suggested relatively lower peak currents in Val89 mice, contrasting with the absence of differences between absolute peak currents (e), an additional analysis revealed a significantly longer time period over which peak currents were distributed in Val89 mice when compared with WT mice, leading to relatively lower averages shown in d (see Results; note that, in f, the main genotype effect and multiple comparisons remain significant after removal of the two highest I/I data points; stars in e, f and in subsequent figures depict results from post hoc multiple comparisons; e0,001, ****e0.001, ****e0.0001).

CHT capacity in the cortex (see Introduction), subsequent experiments assessed effects solely in that brain region.

Impact of HC-3

We co-pressure ejected choline and the competitive CHT inhibitor HC-3 to demonstrate that the choline clearance measure used above (Slope $_{40-80}$) indeed reflects primarily CHT-mediated choline clearance (Parikh and Sarter, 2006), and to explore whether CHT-mediated choline transport can be further reduced in Val89 mice by pharmacological blocking of the transporter (for details about the ejection parameters, see Materials and Methods). Figure 3a illustrates the marked slowing of choline clearance in the presence of HC-3 in WT CHT I/I mice, contrasting with the absence of a change in the slopes of the decay curves in CHT I/V (Fig. 3b) and CHT V/V (Fig. 3c) mice. The resulting Slope $_{40-80}$ values are shown in Figure 3d.

ANOVA of the effects of genotype and HC-3 on choline clearance (Slope₄₀₋₈₀) indicated main effects and a significant interaction between the two factors (genotype: $F_{(2,167)}$ = 83.46, p < 0.0001, d = 1.9; HC-3: $F_{(1,167)} = 74.99$, p < 0.0001, d = 1.3; interaction: $F_{(2,167)} = 57.11$, p < 0.0001, d = 1.5). Post hoc multiple comparisons indicated a HC-3-induced significant decline in the clearance rate in WT mice (p < 0.0001), but not in CHT I/V or CHT V/V mice (both p > 0.81; Fig. 3d). Indeed, choline clearance rates did not differ significantly between the genotypes in the presence of HC-3 (all p > 0.06). A final post hoc comparison between absolute peak choline currents recorded following pressure ejections of solutions including and not including HC-3 $(t_{(25)} = 0.09, p = 0.93; data not shown)$ confirmed that effects on choline clearance were not confounded by differences in the volume and constitution of the ejection solution.

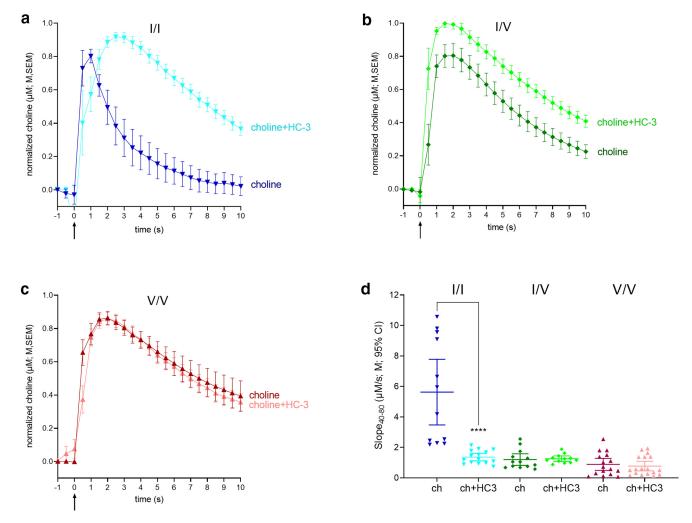


Figure 3. Impact of the CHT competitive inhibitor hemicholinium-3 (HC-3) on choline clearance in WT and CHT Val89 mice. a–c, Averaged traces of exogenous choline clearance with and without HC-3 (for visual clarity, traces shown here and in Fig. 4 were normalized to the peak current values; shown are means ,M, and standard errors of the means, SEM). Traces and slopes in the absence of HC-3 were obtained from a separate group of mice (Fig. 2). Choline clearance in the presence of HC-3 was determined from 15 traces in WT (n = 5), 12 traces in CHT I/V (n = 4), and 18 traces in CHT V/V (n = 6), with 3 traces selected from each mouse. Darker colored lines represent choline clearance traces following pressure ejections of 5 mm choline (ch) alone, while lighter colors show clearance following ejections of a solution containing 5 mm choline and 10 μ m HC-3, with each solution ejected at time 0. d, Individual choline clearance rates (Slope₄₀₋₈₀) following ejections of 5 mm choline (ch) only and a 5 mm choline and 10 μ m HC-3 cocktail (ch + HC3). Only clearance rates recorded in WT mice were significantly slowed by HC-3. Following HC-3 ejections, residual clearance rates did not differ across genotypes (LSD test; d shows individual values, means, M, and 95% confidence intervals, CI).

Together, the findings from the experiments on the effects of HC-3 confirm that the measure of choline clearance (Slope_{40–80}) reflects CHT-mediated clearance, and they indicate that the low clearance rates in both variants were not measurably reduced by HC-3-induced blockade. As the affinity of HC-3, in the absence of extracellular choline, is unaffected by the Val89 variant (Okuda et al., 2002), the unexpectedly large impact of both variants on choline clearance, indicated by residual clearance rates below 1 μ M/s, may have limited the demonstration of any further inhibition by HC-3. However, as the CHT supports vital functions (Ferguson et al., 2004) and thus was expected to remain functional in Val89 mice, we conducted an experiment designed to demonstrate such residual functionality.

Inactivation by lidocaine

Inactivation of neuronal activity and suspension of CHT activity by lidocaine (Tehovnik and Sommer, 1997; Malpeli and Schiller, 1979; Iwamoto et al., 2006; Onizuka et al., 2008) was implemented to determine whether voltage-dependent choline clearance can be detected in CHT Val89 mice. In the presence of

lidocaine, residual choline clearance rates are thought to reflect diffusion (Cass et al., 1993) and, potentially, choline uptake via low-affinity, electrogenic transporters other than CHT (Michel et al., 2006). Therefore, lidocaine was expected to suppress the already low choline clearance rates in Val89 mice.

We confirmed the efficacy of lidocaine to suspend neuronal depolarization in the recording area by measuring K⁺-evoked, depolarization-induced ACh release in one WT mouse. Three traces each prior and $\sim\!2\text{--}4\,\mathrm{min}$ after the lidocaine infusions (0.5 μ l over 3 min) are illustrated in Figure 4a. Peak amplitudes reached $\sim\!10~\mu\mathrm{M}$ before inactivation and 2 $\mu\mathrm{M}$ following lidocaine (Fig. 4b), consistent with prior results showing less than complete inactivation following intracranial infusions of relatively small volumes of lidocaine (Tehovnik and Sommer, 1997).

Choline clearance rates were assessed following the timeline of choline pressure ejections and lidocaine infusions illustrated in Figure 4c. First, we compared choline clearance rates (Slope_{40–80}) measured before lidocaine infusions with those measured in our initial experiment (Fig. 2b). This analysis reproduced the main effect of genotype ($F_{(2,45)} = 12.23$, p < 0.001), reflecting that CHT

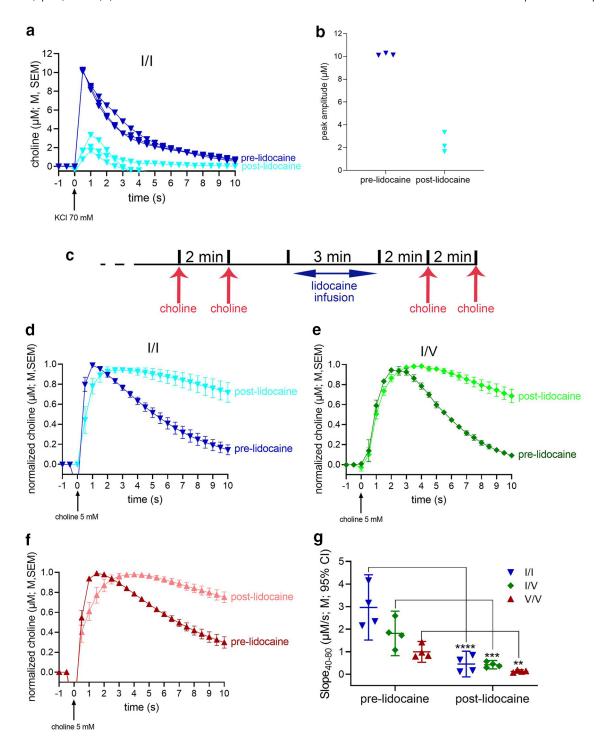


Figure 4. Reduced choline clearance rate following inactivation of neuronal activity and CHT function with lidocaine. An initial experiment, using one WT mouse, was conducted to verify the efficacy of lidocaine (0.5 μ l of 20 μ g/ μ l infused over a consistent rate of 0.17 μ l/min) to inactivate the recording area. Before lidocaine infusions, potassium chloride (KCl; 70 mm) pressure ejections reliably generated ACh release events, peaking at ~10 μ m (see dark blue traces in a). Two to 3 min following lidocaine infusions, peak choline currents were markedly reduced (b), yet not completely abolished, as would be expected from the effects of a relatively small volume of lidocaine (Tehovnik and Sommer, 1997). c, Timeline of choline pressure ejections and lidocaine infusion used to test choline dearance in the three strains. Following a 60-min baseline period, two initial pressure ejections of choline (5 mm), separated by 2 min, were followed by the infusion of lidocaine and, 2 min later, by another two pressure ejections of choline. d-f, Averaged (over 8 traces from 2 mice per genotype) prelidocaine and postlidocaine currents evoked by exogenous choline in WT (d), CHT I/V (e), and CHT V/V (f) mice. g, ANOVA indicated main effects of genotype and lidocaine as well as a significant interaction (Results). LSD multiple comparisons tests indicated significant reductions of choline clearance following inactivation in all three genotypes (g). The significant interaction between the effects of genotype and lidocaine also reflected that choline clearance rates no longer differed between the genotypes (all p > 0.10; a,d,e and f show means, M, and standard errors of the means, SEM; g shows individual values, means, M, and 95% confidence intervals, Cl).

I/I clearance rates were higher than in CHT I/V (p = 0.0004) or CHT V/V mice (p < 0.0001). Furthermore, choline clearance rates across the three strains did not differ between the two experiments ($F_{(1.45)} = 1.26$, p = 0.27).

As illustrated in Figures 4d–g, lidocaine-induced inactivation reduced choline clearance in all three genotypes (main effect of lidocaine: $F_{(1,27)} = 104.40$, p < 0.0001, d = 3.9), with relatively greater effects seen in WT than in Val89 mice (main effect of

genotype: $F_{(2,27)} = 21.12$, p < 0.0001, d = 2.5; interaction: $F_{(2,27)} = 13.40$, p < 0.0001, d = 2.0). Moreover, these results are consistent with an absence of genotype-related differences in clearance rates when transport by CHT and other electrogenic transporters is suppressed (all p > 0.10; Fig. 4g).

Attenuated ACh release capacity following repeated depolarization *in vivo*

The results described above indicate a highly significant reduction in choline clearance in CHT Val89 mice when compared with mice expressing the WT CHT Ile89. Such a reduction is expected to impact the capacity of cholinergic neurons to synthesize and release ACh, particularly when cholinergic activity is relatively high and presynaptic "reloading" of newly synthesized ACh is taxed. Basal forebrain cholinergic neurons exhibit highly diverse firing patterns, with firing bursts of subpopulations of these neurons reaching >80 Hz (Duque et al., 2000; Manns et al., 2000; Laszlovszky et al., 2020). However, given our focus on cholinergic activity during attention, we employed a markedly lower frequency and degree of cholinergic taxation to foster an interpretation of results in the context of the inattentive phenotype in humans expressing the Val89 variant. Attention tasks such as those used in humans and rodents to reveal attentional impairments typically consist of trial frequencies of \sim 1–0.1 Hz (Demeter et al., 2008; Berry et al., 2014). In such tasks, ACh release events occur during successful cue detection (Gritton et al., 2016). We tested ACh release capacity in CHT Val89 mice following a relatively small number of potassium-induced depolarizations, delivered at a relatively low rate (see Materials and Methods; Fig. 5a). ACh release capacity was assessed 5 and 10 min after such depolarization trains (Fig. 5a), mirroring recovery periods that previously revealed ACh release deficiencies in heterozygous CHT knock-out mice (Parikh et al., 2013).

The very first pressure ejection of KCl produced statistically similar choline current peaks in the three genotypes $[F_{(2,12)}]$ = 0.79, p = 0.48; M, SEM (μ M choline): WT: 11.65 \pm 2.06; CHT I/V: 10.47 ± 1.55 ; CHT V/V: 8.15 ± 2.24 ; data not shown]. The peak amplitudes of the choline currents evoked by the subsequent second, third, fourth, and fifth pressure ejections of KCl were expressed as percent of the peaks of the first ejection from the respective depolarization series. Irrespective of genotype and series, the repeated depolarizations during either series appeared to yield increasingly smaller peak amplitudes. Figures 5 b,c show the relative peak amplitudes evoked by the second and fifth ejection of both series. There were no effects of genotype or KCl series (first vs second; see Fig. 5a) on peak amplitudes evoked by the second or fifth KCl ejections (main effects of genotype, series and interactions, analyzed separately for the second and fifth ejection, with $\alpha = 0.05/2$: all F < 2.53, all p > 0.12). Therefore, the mean relative peak amplitude for each mouse was used to compare the effects of the second and fifth ejection. This analysis indicated that regardless of genotype, peak amplitudes were markedly reduced following the fifth ejection (main effect of ejection: $F_{(1,12)} = 88.89$, p < 0.001; d = 5.08; overall M, SEM: second ejection: $81.54 \pm 2.56\%$; fifth: $56.03 \pm 2.11\%$; Fig. 5b,c). These findings indicate the efficiency of the relatively low-frequency depolarization trains with KCl to reveal the limits of cholinergic terminals to synthesize and release ACh.

Five minutes after the first and 10 min after the second KClinduced depolarization series, one additional depolarization was administered to test the "reloading" capacity of cholinergic neurons in the three genotypes. Representative traces of absolute KCl-induced currents (peaks matching the means of the genotype) are shown in Figure 5d and suggested reduced amplitudes in CHT I/V and CHT V/V mice, relative to CHT I/I mice, at both time points. To indicate the degree of synthesis and release recovery, peak amplitudes of choline currents were expressed as a ratio to the peak amplitudes of currents evoked by the first depolarization from the respective series (Fig. 5e,f). Irrespective of genotype, relative peak amplitudes recorded after the 10-min pause were higher than those recorded after the 5-min pause (main effect: $F_{(1,12)} = 6.11$, p = 0.03; d = 1.42; M, SEM: 5 min: $80.86 \pm$ 5.18%; 10 min: $102.38 \pm 5.69\%$; Fig. 5e). Moreover, a main effect of genotype ($F_{(2,12)} = 7.16$, p = 0.009; d = 2.18; Fig. 5f) reflected a relatively greater recovery of ACh release in CHT I/I mice than either of the CHT Val89-expressing mice (no difference between CHT I/V and CHT V/V; genotype x series: $F_{(2,12)} = 0.56$, p = 0.59, d = 0.61; multiple comparisons, using LSD, are shown in Fig. 5f). Taken together, this experiment indicated the efficiency of the depolarization series to reduce ACh output and an attenuated capacity of cholinergic neurons of CHT I/V and CHT V/V mice to regain baseline ACh release levels after relatively long, 5and 10-min pauses.

CHT densities do not differ between genotypes

Because of our overall focus on the role of fronto-cortical cholinergic signaling in attention (Gritton et al., 2016; Howe et al., 2017) and that CHT I/V humans fail to activate right frontal cortex during attention (Berry et al., 2015), frontal cortices were harvested for the analysis of CHT densities in total synaptosomal lysates and synaptosomal plasma membrane-enriched fractions. We previously demonstrated that CHT density in the latter relatively precisely predicts choline clearance capacity (Ferguson et al., 2003; Apparsundaram et al., 2005; Parikh et al., 2013; Koshy Cherian et al., 2019). As illustrated in Figure 6, CHT densities in neither total synaptosomal lysates nor plasma membraneenriched fractions differed by genotype (both F < 0.58, both p > 0.56). These findings are consistent with the view that attenuated choline clearance and ACh release following repeated depolarizations likely reflects a conformational modification of the protein induced by the Val89 variant that impacts transporter function rather than translation, stability or surface trafficking of CHT protein.

Attentional control deficit in Val89 mice

Mice were trained to perform a continuous signal detection task that was previously used to measure sustained attention in rats and humans (Demeter et al., 2008) and modified for use in mice (St Peters et al., 2011). Subjects were trained to report the presence or absence of a signal in trials that began either with a signal (variable duration) or a blank. Responses were hits and misses, and correct rejections and false alarms, respectively, with hits and correct rejections yielding water reward (Fig. 7a). Hits and false alarms were used to compute signal detection theoryderived measures of perceptual sensitivity (d') and response bias (B"D; Swets et al., 1961; Green and Swets, 1974; Donaldson, 1992). Val89 mice were not expected to exhibit deficits in perceptual sensitivity (Berry et al., 2015), but to alter their bias in response to a performance challenge. This prediction was derived in part from evidence in unscreened humans who performed a human version of the present signal detection task and found to adopt a relatively more conservative criterion in response to the presentation of a visual disruptor. In other words, the disruptor evoked a relatively greater

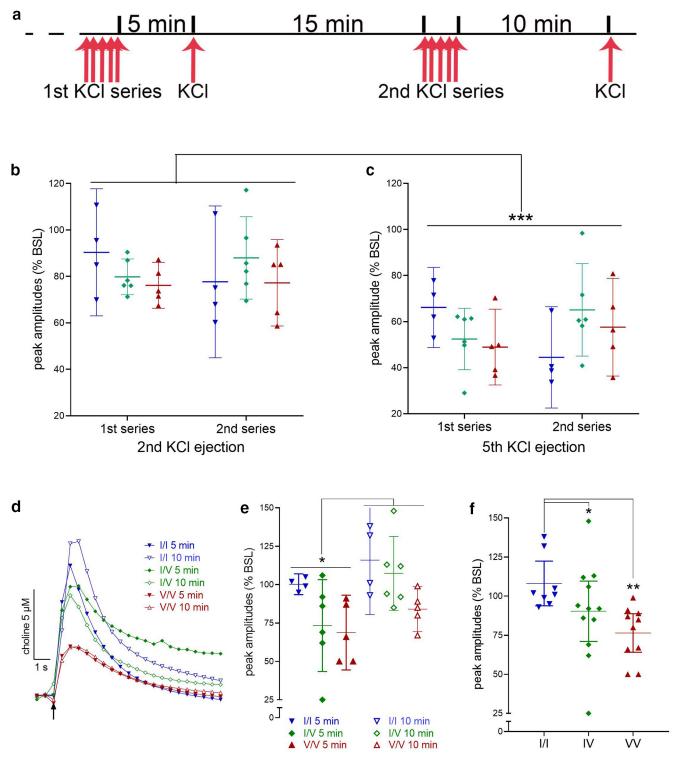


Figure 5. Impact of CHT Val89 on ACh release capacity following repeated depolarization. a, Schematic illustration of the timeline of potassium chloride (KCl) pressure ejections. Following a period of calibration, an initial train of five KCl ejections were administered over 2 min and followed, 5 min later, by a first release capacity test. After a 15-min pause, a second series of depolarizations was followed, 10 min later, by a second release capacity test. Peak amplitude currents following the first KCl ejection of either series did not vary by genotype (data not shown). b, c, Peak amplitudes of currents evoked by the second and fifth ejection, respectively, of either series, expressed as a ratio of the amplitudes of the first ejection (BSL; n = 4 WT, 6 CHT I/V, and 5 CHT V/V mice; lines and bars in b, c, e, f depict means and 95% Cl). Compared with the second depolarization, the fifth resulted in significantly smaller amplitudes across genotypes and the two series. d, Representative traces of currents evoked at the 5- and 10-min recovery tests, illustrating attenuated ACh release in CHT I/V and CHT V/V mice. e, Peak amplitudes of the recovery currents, expressed as the percent of the amplitude of the first depolarization of the respective series (baseline, BSL), indicated significantly greater recovery after the longer 10-min pause compared with peak currents recorded 5 min after the depolarization train (see Results for ANOVA). The analysis of the effects of series and genotype also revealed a main effect of genotype, illustrated in f, that reflected significantly lower peak amplitudes in CHT I/V (12 ejections from n = 6) and CHT V/V (10 ejections from n = 5) mice relative to peak amplitudes in WT mice (8 ejections from n = 4; LSD test; see f for post hoc multiple comparisons; b, c, e, f show means and 95% confidence intervals).

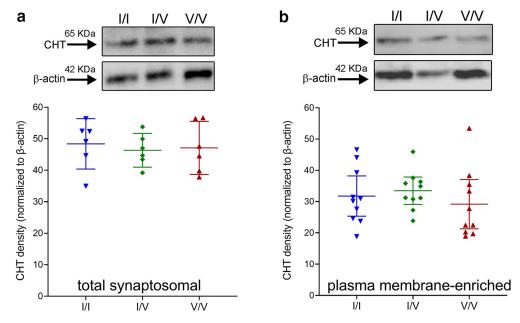


Figure 6. Top, Western blottings showing the presence of the CHT in frontal cortical total synaptosomal lysates (a, top) and synaptosomal plasma-membrane-enriched fractions (b, top; normalized against β-actin; each lane is based on tissue from one individual; 25-μg protein loaded in each well; total and plasma membrane-enriched fractions were derived from separate groups of mice). CHT densities in neither total lysates (a) nor plasma membrane-enriched fractions (b) differed between the genotypes (graphs show individual values, means and 95% confidence intervals (CI)).

propensity to report the absence of a signal (Demeter et al., 2008), thereby maintaining and recovering above-chance levels of response accuracy and reward rates. Thus, WT mice were expected to respond to a visual disrupter (see Materials and Methods) by shifting to a more conservative bias ($B^{"}_{D}$ scores between 0 and 1), whereas Val89 mice were predicted to not change their bias, reflecting that bottomup, or signal-driven mechanisms predominantly supported their performance.

Unchallenged task performance

Upon reaching asymptotic performance in the unchallenged task, mice reliably reported the presence of signals in a signal duration-dependent manner (mean percent hits to 500-, 50-, and 25-ms signals: 72.39%, 47.05%, and 38.45%) and accurately reported the absence of signals (mean percent correct rejections: 70.20%). As was expected given signal duration-dependent hit rates, perceptual sensitivity, d', varied significantly by signal duration ($F_{(2.48)} = 146.40$, p < 0.001; 500 ms signals (mean): d' = 1.07; 50 ms: -0.28; 25 ms: -0.74; multiple comparisons indicated significant differences between all three durations; data not shown). However, there were no effects of genotype or interactions between genotype and signal duration on this measure (both F < 2.80, all p > 0.07; data not shown).

As performance measures calculated over hit rates to 50- and 25-ms signals approached chancel level, B_D^* scores were computed only over hits to longest signal durations. Performing the unchallenged task, mice exhibited a mildly liberal bias ($B_D^* = -0.15$ (mean). B_D^* scores did not differ by genotype, and there were not interactions involving the factor genotype on bias (all F < 1.44; all p > 0.21; data not shown).

The relative number of omissions remained relatively low (grand mean: 17.85%) but increased across blocks of trials from 9.31% in block 1–32.06% in block 5 ($F_{(1.94,46.58)} = 15.84$, p < 0.001). There were no significant effects of genotype and no interactions between genotype and block on omissions (both F < 1.79, both p > 0.18; data not shown).

Bias changes evoked by a disruptor challenge

Performance was challenged using a visual distractor, presented in the second and third blocks of trials. The robust and genotype-unrelated efficacy of the disruptor was indicated by a significant reduction of perceptual sensitivity, calculated over hits to longest signals (main effect of block on d'; $F_{(4,96)} = 8.47$, p < 0.0001; Fig. 7b). Perceptual sensitivity during disruptor challenges was not affected by genotype (main effects and interactions with block: both F < 2.42, both p > 0.11).

Similar to the response biases derived from unchallenged task performance, during the first (predisruptor) trial block all mice exhibited a relatively liberal response bias, that is, a propensity to report the presence of a signal [no effect of genotype: $F_{(2,24)}$ = 1.71; p = 0.21; B''_D (M, SEM): -0.29, 0.09]. As a result of the disrupter presentation WT, but not Val89, mice exhibited a shift in bias, toward a greater propensity for reporting the absence of a signal, relative to the first predistractor block of trials (main effect of genotype on B''_D ($F_{(2,24)} = 4.68$, p = 0.02, d = 0.78; Fig. 7c). This genotype-specific response to the distractor remained stable throughout the remainder of the test session (no effect of block and no interaction between the effects of genotype and block; both F < 1.80, both p > 0.15). As was the case performing the unchallenged version of the task, all mice steadily omitted more trials in the course of the disruptor session (block 1 mean: 8.22%; block 5: 23.81%; main effect of block: $F_{(1.82,43.75)} = 11.59$, p = 0.0001; effects of genotype and interaction with block: both F < 3.10, both p > 0.06; data not shown).

Structural modeling of WT mouse CHT and the CHT Val89 variant

To gain insights into the structural underpinnings of the effect of the I89V substitution on the CHT transport mechanism, we pursued a template-based molecular modeling approach for WT mCHT and the Val89 variant. Our approach involved two key steps: identifying a suitable template and determining a high-quality sequence alignment between the template and query used

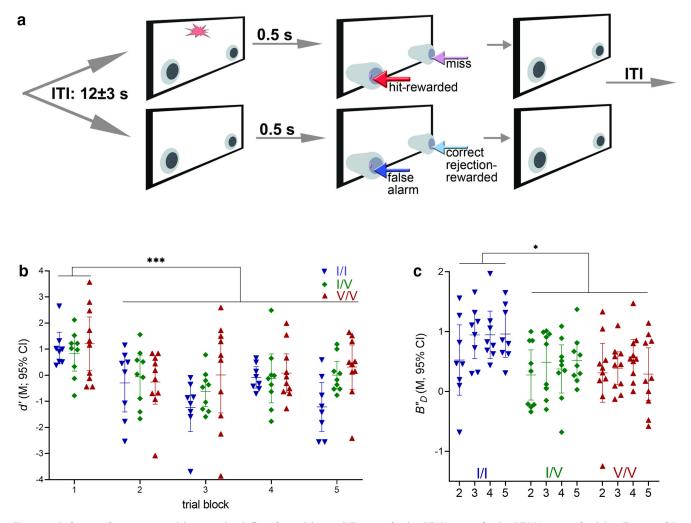


Figure 7. Performance of a continuous signal detection task and effects of a visual disruptor (WT: n = 8, 3 females; CHT I/V: n = 9, 4 females; CHT V/V: n = 10, 2 females). **a**, Illustration of the two trial types and main events of the signal detection task employed to assess sustained attention in mice. Briefly, trials began either with a light signal of variable duration (top) or a blank (bottom), followed by the extension of nose-poking device into the chamber. Half of the mice were trained to nose-poke on the left to report the presence of a signal and on the right to report the absence of a signal. Correct responses (hits in signal trials and correct rejections in nonsignal trials) were rewarded with water delivery while incorrect responses (misses and false alarms, respectively) triggered the onset if the intertrial interval (ITI; for more details, see Materials and Methods). A test session lasted 40 min. Performance was challenged by the presentation of a visual disruptor (house lights flashing at 0.5 Hz) during the second and third 8 min block of trials (out of 5 blocks total). Hits and false alarms were used to calculate signal detection theory-derived measures of perceptual sensitivity d' and response bias B''_D (references in Materials and Methods). Perceptual sensitivity depicts the detectability of signals; in contrast, response bias indicates the individual's propensity to report the presence or absence of a signal. **b**, The visual disruptor was highly effective in reducing the detection of signals in all genotypes, as indicated by near or at chance level performance during and after the presentation of the disruptor (trial blocks 2–5; main effect of trial block; see Results; **b**, **c** depict measures calculated over longest signal durations, as scores to shorter signals indicated chance level performance). **c**, Effects of the disruptor (main effect of genotype, see Results). In other words, WT mice were less likely to report the presence of a signal when challenged with a disruptor.

during the production run. The use of HMM-based profiles in fold-recognition algorithms, such as implemented in HHpred, allows for detecting distantly related homolog sequences. This approach identified the *Vibrio parahaemolyticus* sodium/sugar symporter (vSGLT, PDB id: 3DH4; Faham et al., 2008) as the best template when compared with other SLC5 family members. The initial alignment of vSGLT and CHT covers 80% of the sequences with a sequence identity of 19%. Despite relatively low sequence similarity, the hydrophobicity profiles of the two transporters, used as indicators for proteins that share the same fold (Lolkema and Slotboom, 1998; Sarkar and Kellogg, 2010), showed visibly similar widths and distribution of hydrophobicity peaks, indicating that both proteins have a similar fold. It is noteworthy that TM14 in vSGLT has no counterpart in mouse (or

human) CHT, which is predicted to have 13 transmembrane helices (Fig. 8*a*,*b*), extracellular N and cytoplasmic C termini (Fig. 8*d*).

The initial alignment was refined using an iterative procedure in which the gaps within secondary structural elements are removed, and the conservation scores from Consurf were used as a guide to position highly conserved residues within the core of the protein (Ashkenazy et al., 2016). The refined sequence alignment shows an overall correspondence between secondary structural elements, except in three of the loops connecting transmembrane helices. This matching was particularly striking for the first 13 TM helices (Fig. 8b). The broad sequence coverage between mCHT and vSGLT, similarity of the hydrophobicity profiles and matching between the TM helices, further substantiate the view that these two proteins share a common fold. The

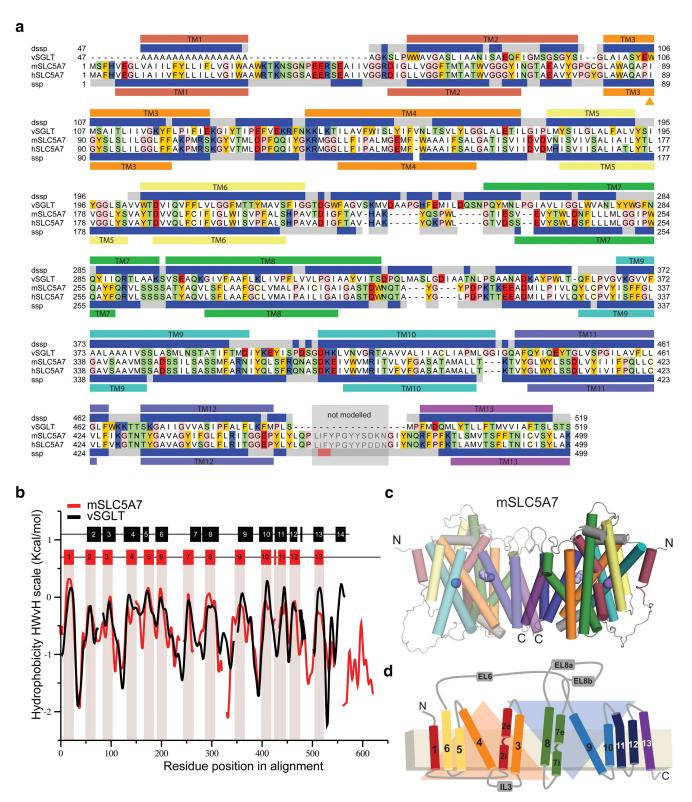


Figure 8. Structural model of a mouse CHT dimer. a, Sequence alignment between mouse and human CHT (SLC5A7) and the template vSGLT used for structural modeling studies. The secondary structure prediction (SSP) averaged over the SLC5A7 profile from HHpred server (Hildebrand et al., 2009) is shown below hSLC5A7 sequence as gray (coil), blue (helix) and red (β -strand) bars. The transmembrane topology prediction from Topcons server (Tsirigos et al., 2015) is also shown as bars below SSP colored and numbered according to the topology proposed by Faham et al. (2008). The secondary structure and transmembrane helices based on the structure of the template vSGLT are also shown on top of vSGLT sequence, similarly to that of CHT. The region between TM12 and TM13 shaded in gray was not modeled. Residues shown in the model are highlighted according to their properties; neutral in white, aromatic in yellow, polar in green, basic in blue and acidic in red. b, Hydrophobicity profiles for mSLC5A7 (red) and vSGLT (black) revealing conserved hydrophobic regions. Residues corresponding to the TM helices in vSGLT or those predicted for mSLC5A7 are indicated as black and red bars at the top of the graph and in the case of mSLC5A7 also as red shading. c, d, Final structural model of the dimeric form of mSLC5A7 and the correspondent topology. In both, TM helices are colored according to Faham et al. (2008) and the inverted repeat units are shown as orange and blue triangles.

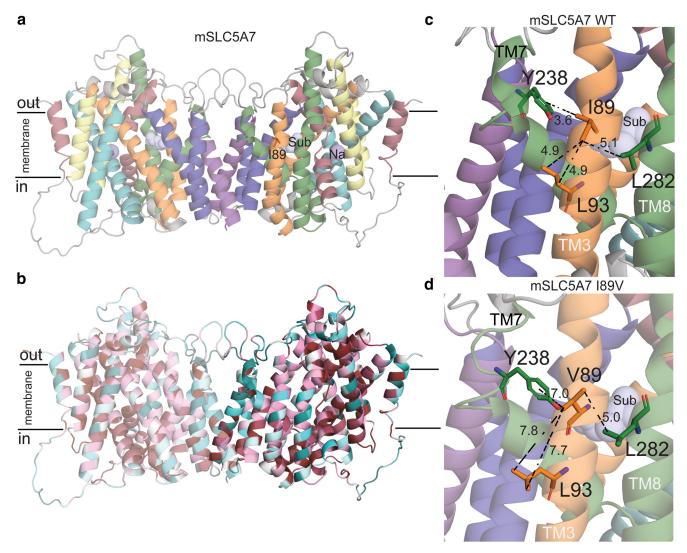


Figure 9. Structural model of mouse CHT Val89. *a*, Structural model of mSLC5A7 in dimeric form colored according to the topology. 189, the sodium ion and substrate are highlighted as orange sticks and purple and light blue spheres, respectively. The position of the substrate and sodium ions where those obtained after structural superimposition of CHT model and vSGLT structure. *b*, Final model of mSLC5A7 where each residue is colored by its conservation score obtained with Consurf server (Ashkenazy et al., 2016). *c*, *d*, Close-up views of lle89 and Val89 in their corresponding mSLC5A7 structures. Residues interacting with lle89 are shown as sticks and the distances between sidechains are shown as dashed lines. The helices are colored as in *a*.

analysis of the stereochemistry for our model of CHT shows only one residue, located in a loop, in the disallowed region of the Ramachandran plot obtained with PROCHECK (see Materials and Methods). In addition to the excellent stereochemistry, the model has a ProQM score averaged by the number of residues (global ProQM score) of 0.67, comparable to that of the template (0.7), indicating the good quality of the model obtained.

Our work indicates that CHT has a similar global architecture as that of vSGLT (Fig. 8c,d). mCHT was modeled as a dimer, similar to the template, where transmembrane helices TM12 and TM13 constitute the dimerization interface with highly conserved residues taking part in the dimerization interacting network (Figs. 8c,d, 9a,b), supporting the possibility that mouse and human CHT assemble as a dimer. Each of the protomers has an inverted repeat topology, in which TMs 2–6 and TMs 7–11 constitute repeat units 1 and 2 (RU1, RU2) with a similar topology to the core of LeuT (TM1–TM10; Faham et al., 2008), and a central cavity formed by TM2, TM3, TM7, TM8, and TM11 exposed to the cytoplasm, and helices 1, 12, and 13 as peripheral.

In our model, we found Ile89 to be located in TM3 of CHT, facing the membrane and stabilizing the packing between TM3,

TM8, and TM7 by interacting with residues Leu93 (TM3), Tyr238 (TM7), and Leu282 (TM8; Fig. 9c). This network of interactions is disrupted when Ile89 is substituted by Val89 (Fig. 9d), suggesting that the variant may destabilize helical packing in this region. It is worth noting that these three TMs are predicted to participate in shaping the cavity where choline binds, and thus Val89 may disrupt the exposure of the CHT substrate binding site. In addition, TM3, TM7, and TM8 are part of the "bundle" of the protein, which is believed to move with respect to the scaffold and the membrane plane on the conformational changes responsible for transitions of the substrate binding site between states open to either extracellular or cytoplasmic spaces (alternating-access mechanism of transport). Destabilization of the fold next to position 89 by the Ile89Val substitution could also affect the ability of the transporter to adopt canonical conformations that the protein needs to reach during a transport cycle.

Discussion

The current study aimed to determine the cholinergic impact of the common human CHT coding variant, Val89, that was

previously demonstrated to be associated with an inattentive phenotype in humans, along with attenuated frontal activation during attentional performance (Berry et al., 2014, 2015; Sarter et al., 2016). The results indicate a dominant-negative reduction in choline clearance capacity in Val89 mice of ~80% in cortex and over 50% in striatum. Furthermore, although the initial depolarization-induced release of ACh from cholinergic terminals was similar across all mice, we found that 5 or 10 min following repeated depolarization, the capacity to release ACh was significantly attenuated in Val89-expressing mice. The effects of an attentional performance challenge indicated that, in contrast to WT mice, Val89 mice performed a continuous signal detection task largely through bottom-up, signal-driven processes. Immunoblotting indicated that surface expression of the CHT did not differ between the genotypes, suggesting that Val89-associated impairments were not because of aberrant surface trafficking. Structural modeling suggested potential cellular mechanisms underlying attenuated choline transport by the Val89 variant.

Results from the depolarization studies (Fig. 4) demonstrated that, in response to the initial depolarization, choline transport by the variant remains sufficient to support ACh release at a level that is comparable to WT mice. However, the impact of attenuated CHT function on ACh release capacity manifests following repeated depolarizations. Importantly, the paradigm we used to demonstrate a diminished "reloading" capacity for subsequent ACh release involved a relatively small number of depolarizations at a relatively low frequency, mirroring ACh release events seen during hits in attentional task-performing rodents. An even greater impact of the Val89 variant would be expected when cholinergic neurons exhibit burst firing (Manns et al., 2003).

Our behavioral assessments indicated that in response to a visual disrupter, WT mice, but not Val89 mice, adopted a relatively more conservative criterion. Such a shift in bias is thought to reflect the deployment of a set of psychological processes that are conceptualized as voluntary top-down attentional control or model-based decision-making (Clark et al., 2012). Specifically, in the present context, WT mice may have monitored disruptor-induced reward losses, triggering an increase in their threshold for reporting the presence of a signal. In contrast, Val89 mice may have relied primarily on the bottom-up detection of cues and model-free processing to perform this task. Consistent with the present evidence, such processing preserves unchallenged signal detection performance but limits the flexibility to respond to contextual alterations, such as disruptor-induced performance loss

The finding that WT mice adopted a more conservative bias in response to a disruptor was not associated with improvement in performance. This finding mirrors the absence of relatively greater performance deficits in heterozygous Val89 humans, compared with WT humans, using a comparable task and visual disruptor (Berry et al., 2015). We hypothesize that performing such a signal detection task may not sufficiently tax top-down control to produce performance loss arising from poor attentional control. To reveal performance benefits of a conservative criterion shift, and performance deficits in Val89 subjects, a rodent task that more severely taxes top-down processes and utilizes a true distractor, as opposed to a sensory disruptor, is needed, similar to the task used to demonstrate distractor vulnerability in Val89 humans (Berry et al., 2014).

The hypothesis that relatively poor attentional control results from attenuated cholinergic activity, is supported by evidence from rats with a spontaneous CHT intracellular trafficking deficit (Paolone et al., 2013a; Koshy Cherian et al., 2017; Pitchers et al.,

2017a, b; Sarter and Phillips, 2018; Phillips and Sarter, 2020). Furthermore, in patients with Parkinson's disease (PD), residual attentional performance, assessed by using a continuous temporal expectancy test (O'Connell et al., 2009) and challenged with a content-rich distractor, correlated with residual cholinergic innervation of cortex (Kim et al., 2019). Using the same test as in PD patients with cholinergic losses, humans heterozygous for the Val89 variant showed reduced resistance against such a distractor when compared with WT humans (Berry et al., 2014). In addition, Val89 humans encoded more distractor content than WT humans, further supporting relatively poor top-down attentional control as an essential phenotypic characteristic.

Our *in vivo* method indicated a relatively large reduction of CHT-mediated choline transport rates by the minor variant when compared with prior data obtained *in vitro* (Okuda et al., 2002). This finding warranted additional experiments to reveal the functionality of the CHT variant. The effects of HC-3 and lidocaine, while validating our *in vivo* choline transport measure in terms of a reduction in CHT-mediated transport, suggest either a residual, voltage-dependent choline transport capacity of CHT Val89, or a contribution to choline clearance by other electrogenic transporters (Onizuka et al., 2008). As interactions of choline, sodium, chloride and HC-3 with CHT Val89 have been reported to be unaffected in transfected cells (Okuda et al., 2002), our results are consistent with a diminished capacity for the high-affinity choline transport needed to sustain high-demand ACh release.

Our structural models of WT mouse CHT and the Val89 encoding transporter point to a structural importance of Ile89 in local packing of substrate transport-critical helices. We show that Ile89 is predicted to participate in an interacting network that stabilizes the fold of helices that contribute to the "bundle" domain of the "rocking bundle" model of transport (Forrest and Rudnick, 2009). This network may also contribute to the shaping of the binding site by interacting with TM7 and TM8. Because Ile89 is not predicted to contribute to the substrate binding site (see also Okuda et al., 2002), it seems most likely that any impact on choline recognition by Val89 must be indirect. Therefore, we favor a model whereby conformational transitions required to support alternating access of the choline binding site to extracellular and intracellular compartments are compromised. Additionally, our models were well fit to a dimeric structure solved for vSGLT, including TMs position to support the dimeric interface.

Such a structure may underlie dominant-negative mode of action, which has been previously shown for human disease-associated CHT coding variants (Barwick et al., 2012). Although choline clearance data from CHT I/V mice showed compression toward zero, thereby limiting potentially the demonstration of greater effects in CHT V/V mice, measures of potassium-evoked ACh release and response bias might have readily revealed genedose effects. In humans expressing Val89 variant, the presence of a gene-dose effect on attentional control remains unknown but appears unlikely given the already relatively large effect size seen in heterozygous Val89 humans (Berry et al., 2014, their Fig. 2). Further studies, including biophysical analyses of choline uptake and choline-induced currents (Iwamoto et al., 2006), are needed to characterize the conformational dynamics of the Val89 variant.

Here, we focused on measures of presynaptic cholinergic capacity in the cortex because of the association of the Val89 variant with heightened distractor vulnerability in humans and based on evidence indicating an essential role of cortical

cholinergic activity for distractor filtering (references in Introduction). Although the demonstration of attenuated clearance rates in a second brain region, the striatum, is consistent with a systemic impact of the variant, the striatal alteration is less striking, indicating a region-dependence to the functional impact of Val89 that may also underlie why CHT Val89 mice show no overt behavioral, autonomic or motor deficits. Motor deficits would have drastically interfered with the ability to perform the basic operant response requirements of the signal detection task, including executing nose-pokes and retrieving water reward across relatively long test sessions and rapidly sequenced trials. We suspect that motor (and other peripheral) cholinergic terminals possess greater ACh synthesis reserve than those we queried in the CNS, though further efforts are needed to explore this possibility.

The present findings establish that the Val89 variant, which has previously been linked to attentional impairment in humans (English et al., 2009; Berry et al., 2014, 2015; Sarter et al., 2016), as well as major depression (Hahn et al., 2008), greatly impacts the capacity of cortical cholinergic presynaptic terminals to import choline and to sustain ACh release, even following relatively moderate depolarization rates. As such, the present evidence provides a mechanistic framework to interpret the increased distraction vulnerability and frontal cortical activation deficits in Val89-expressing humans. As impairments in cholinergically mediated, attentional-perceptual functions and frontal cortical abnormalities are essential variables in the manifestation of major psychiatric and neurologic disorders (Bohnen and Albin, 2009; Bohnen et al., 2009; Lustig and Sarter, 2016; Berry et al., 2017; Albin et al., 2018; Kim et al., 2019; Sarter et al., 2021), and Val89 allele frequency is surprisingly common for a transporter functional coding variant, further investigation of the impact of the CHT Val89 variant in humans and CHT Val89 mice is warranted.

References

- Albin RL, Bohnen NI, Muller M, Dauer WT, Sarter M, Frey KA, Koeppe RA (2018) Regional vesicular acetylcholine transporter distribution in human brain: a [(18) F]fluoroethoxybenzovesamicol positron emission tomography study. J Comp Neurol 526:2884–2897.
- Apparsundaram S, Martinez V, Parikh V, Kozak R, Sarter M (2005) Increased capacity and density of choline transporters situated in synaptic membranes of the right medial prefrontal cortex of attentional task-performing rats. J Neurosci 25:3851–3856.
- Ashkenazy H, Abadi S, Martz E, Chay O, Mayrose I, Pupko T, Ben-Tal N (2016) ConSurf 2016: an improved methodology to estimate and visualize evolutionary conservation in macromolecules. Nucleic Acids Res 44: W344–W350.
- Barwick KE, Wright J, Al-Turki S, McEntagart MM, Nair A, Chioza B, Al-Memar A, Modarres H, Reilly MM, Dick KJ, Ruggiero AM, Blakely RD, Hurles ME, Crosby AH (2012) Defective presynaptic choline transport underlies hereditary motor neuropathy. Am J Hum Genet 91:1103–1107.
- Bazalakova MH, Blakely RD (2006) The high-affinity choline transporter: a critical protein for sustaining cholinergic signaling as revealed in studies of genetically altered mice. Handb Exp Pharmacol (175):525–544.
- Berry AS, Demeter E, Sabhapathy S, English BA, Blakely RD, Sarter M, Lustig C (2014) Disposed to distraction: genetic variation in the cholinergic system influences distractibility but not time-on-task effects. J Cogn Neurosci 26:1981–1991.
- Berry AS, Blakely RD, Sarter M, Lustig C (2015) Cholinergic capacity mediates prefrontal engagement during challenges to attention: evidence from imaging genetics. Neuroimage 108:386–395.
- Berry AS, Sarter M, Lustig C (2017) Distinct frontoparietal networks underlying attentional effort and cognitive control. J Cogn Neurosci 29:1212–1225

- Bohnen NI, Albin RL (2009) Cholinergic denervation occurs early in Parkinson disease. Neurology 73:256–257.
- Bohnen NI, Müller MLTM, Koeppe RA, Studenski SA, Kilbourn MA, Frey KA, Albin RL (2009) History of falls in Parkinson disease is associated with reduced cholinergic activity. Neurology 73:1670–1676.
- Burmeister JJ, Moxon K, Gerhardt GA (2000) Ceramic-based multisite microelectrodes for electrochemical recordings. Anal Chem 72:187–192.
- Burmeister JJ, Palmer M, Gerhardt GA (2003) Ceramic-based multisite microelectrode array for rapid choline measures in brain tissue. Analytica chimica acta 481:65–74.
- Cass WA, Zahniser NR, Flach KA, Gerhardt GA (1993) Clearance of exogenous dopamine in rat dorsal striatum and nucleus accumbens: role of metabolism and effects of locally applied uptake inhibitors. J Neurochem 61:2269–2278.
- Chen N, Sugihara H, Sur M (2015) An acetylcholine-activated microcircuit drives temporal dynamics of cortical activity. Nat Neurosci 18:892–902.
- Chu AJ, Lee JM (1994) Lidocaine inhibits choline uptake and phosphatidylcholine biosynthesis in human leukemic monocyte-like U937 cells. Cell Biochem Funct 12:89–98.
- Clark JJ, Hollon NG, Phillips PE (2012) Pavlovian valuation systems in learning and decision making. Curr Opin Neurobiol 22:1054–1061.
- Cohen J (1988) Statistical power analysis for the behavioral sciences, Ed 2. Hillsdale: L. Erlbaum Associates.
- Cong L, Ran FA, Cox D, Lin S, Barretto R, Habib N, Hsu PD, Wu X, Jiang W, Marraffini LA, Zhang F (2013) Multiplex genome engineering using CRISPR/Cas systems. Science 339:819–823.
- Demeter E, Woldorff MG (2016) Transient distraction and attentional control during a sustained selective attention task. J Cogn Neurosci 28:935–947.
- Demeter E, Sarter M, Lustig C (2008) Rats and humans paying attention: cross-species task development for translational research. Neuropsychology 22:787–799.
- Demeter E, Guthrie SK, Taylor SF, Sarter M, Lustig C (2013) Increased distractor vulnerability but preserved vigilance in patients with schizophrenia: evidence from a translational sustained attention task. Schizophr Res 144:136–141.
- Donaldson W (1992) Measuring recognition memory. J Exp Psychol Gen 121:275–277.
- Duque A, Balatoni B, Detari L, Zaborszky L (2000) EEG correlation of the discharge properties of identified neurons in the basal forebrain. J Neurophysiol 84:1627–1635.
- English BA, Hahn MK, Gizer IR, Mazei-Robison M, Steele A, Kurnik DM, Stein MA, Waldman ID, Blakely RD (2009) Choline transporter gene variation is associated with attention-deficit hyperactivity disorder. J Neurodev Disord 1:252–263.
- Ennis EA, Blakely RD (2016) Choline on the move: perspectives on the molecular physiology and pharmacology of the presynaptic choline transporter. Adv Pharmacol 76:175–213.
- Faham S, Watanabe A, Besserer GM, Cascio D, Specht A, Hirayama BA, Wright EM, Abramson J (2008) The crystal structure of a sodium galactose transporter reveals mechanistic insights into Na+/sugar symport. Science 321:810–814.
- Ferguson SM, Blakely RD (2004) The choline transporter resurfaces: new roles for synaptic vesicles? Mol Interv 4:22–37.
- Ferguson SM, Savchenko V, Apparsundaram S, Zwick M, Wright J, Heilman CJ, Yi H, Levey AI, Blakely RD (2003) Vesicular localization and activitydependent trafficking of presynaptic choline transporters. J Neurosci 23:9697–9709.
- Ferguson SM, Bazalakova M, Savchenko V, Tapia JC, Wright J, Blakely RD (2004) Lethal impairment of cholinergic neurotransmission in hemicholinium-3-sensitive choline transporter knockout mice. Proc Natl Acad Sci U S A 101:8762–8767.
- Forrest LR, Rudnick G (2009) The rocking bundle: a mechanism for ion-coupled solute flux by symmetrical transporters. Physiology (Bethesda) 24:377–386.
- Giuliano C, Parikh V, Ward JR, Chiamulera C, Sarter M (2008) Increases in cholinergic neurotransmission measured by using choline-sensitive microelectrodes: enhanced detection by hydrolysis of acetylcholine on recording sites? Neurochem Int 52:1343–1350.
- Gorka AX, Knodt AR, Hariri AR (2015) Basal forebrain moderates the magnitude of task-dependent amygdala functional connectivity. Soc Cogn Affect Neurosci 10:501–507.

- Green DM, Swets JA (1974) Signal detection theory and psychophysics. Huntington: R. E. Krieger Publ Co.
- Greenwald AG, Gonzalez R, Harris RJ, Guthrie D (1996) Effect sizes and p values: what should be reported and what should be replicated? Psychophysiology 33:175–183.
- Gritton HJ, Kantorowski A, Sarter M, Lee TM (2012) Bidirectional interactions between circadian entrainment and cognitive performance. Learn Mem 19:126–141.
- Gritton HJ, Stasiak AM, Sarter M, Lee TM (2013) Cognitive performance as a zeitgeber: cognitive oscillators and cholinergic modulation of the SCN entrain circadian rhythms. PLoS One 8:e56206.
- Gritton HJ, Howe WM, Mallory CS, Hetrick VL, Berke JD, Sarter M (2016) Cortical cholinergic signaling controls the detection of cues. Proc Natl Acad Sci U S A 113:E1089–E1097.
- Guyenet P, Lefresne P, Rossier J, Beauquan JC, Glowinski J (1973) Inhibition by hemicholinium-3 of [14C] acetylcholine synthesis and [3H] choline high-affinity uptake in rat striatal synaptosomes. Mol Pharmacol 9:630–639.
- Hahn MK, Blackford JU, Haman K, Mazei-Robison M, English BA, Prasad HC, Steele A, Hazelwood L, Fentress HM, Myers R, Blakely RD, Sanders-Bush E, Shelton R (2008) Multivariate permutation analysis associates multiple polymorphisms with subphenotypes of major depression. Genes Brain Behav 7:487–495.
- Hessa T, Kim H, Bihlmaier K, Lundin C, Boekel J, Andersson H, Nilsson I, White SH, von Heijne G (2005) Recognition of transmembrane helices by the endoplasmic reticulum translocon. Nature 433:377–381.
- Hildebrand A, Remmert M, Biegert A, Söding J (2009) Fast and accurate automatic structure prediction with HHpred. Proteins 77 [Suppl 9]:128–132.
- Howe WM, Gritton HJ, Lusk NA, Roberts EA, Hetrick VL, Berke JD, Sarter M (2017) Acetylcholine release in prefrontal cortex promotes gamma oscillations and theta-gamma coupling during cue detection. J Neurosci 37:3215–3230.
- Huang Z, Villarta-Snow R, Lubrano GJ, Guilbault GG (1993) Development of choline and acetylcholine Pt microelectrodes. Anal Biochem 215:31–37.
- Iwamoto H, Blakely RD, De Felice LJ (2006) Na+, Cl-, and pH dependence of the human choline transporter (hCHT) in *Xenopus* oocytes: the proton inactivation hypothesis of hCHT in synaptic vesicles. J Neurosci 26:9851–9859
- Khafizov K, Staritzbichler R, Stamm M, Forrest LR (2010) A study of the evolution of inverted-topology repeats from LeuT-fold transporters using AlignMe. Biochemistry 49:10702–10713.
- Kim K, Müller M, Bohnen NI, Sarter M, Lustig C (2019) The cortical cholinergic system contributes to the top-down control of distraction: evidence from patients with Parkinson's disease. Neuroimage 190:107–117.
- Koshy Cherian A, Kucinski A, Pitchers KK, Yegla B, Parikh V, Kim Y, Valuskova P, Gurnani S, Lindsley CW, Blakely RD, Sarter M (2017) Unresponsive choline transporter as a trait neuromarker and a causal mediator of bottom-up attentional biases. J Neurosci 37:2947–2959.
- Koshy Cherian A, Tronson NC, Parikh V, Kucinski A, Blakely RD, Sarter M (2019) Repetitive mild concussion in subjects with a vulnerable cholinergic system: lasting cholinergic-attentional impairments in CHT+/- mice. Behav Neurosci 133:448–459.
- Laskowski RA, MacArthur MW, Moss DS, Thornton JM (1993) PROCHECK: a program to check the stereochemical quality of protein structures. J Appl Crystallogr 26:283–291.
- Laszlovszky T, Schlingloff D, Hegedüs P, Freund TF, Gulyás A, Kepecs A, Hangya B (2020) Distinct synchronization, cortical coupling and behavioral function of two basal forebrain cholinergic neuron types. Nat Neurosci 23:992–1003.
- Lolkema JS, Slotboom DJ (1998) Estimation of structural similarity of membrane proteins by hydropathy profile alignment. Mol Membr Biol 15:33–42.
- Lu Y, Sarter M, Zochowski M, Booth V (2020) Phasic cholinergic signaling promotes emergence of local gamma rhythms in excitatory-inhibitory networks. Eur J Neurosci 52:3545–3560.
- Lustig C, Sarter M (2016) Attention and the cholinergic system: relevance to schizophrenia. Curr Top Behav Neurosci 28:327–362.
- Malpeli JG, Schiller PH (1979) A method of reversible inactivation of small regions of brain tissue. J Neurosci Methods 1:143–151.

- Manns ID, Alonso A, Jones BE (2000) Discharge profiles of juxtacellularly labeled and immunohistochemically identified GABAergic basal forebrain neurons recorded in association with the electroencephalogram in anesthetized rats. J Neurosci 20:9252–9263.
- Manns ID, Alonso A, Jones BE (2003) Rhythmically discharging basal forebrain units comprise cholinergic, GABAergic, and putative glutamatergic cells. J Neurophysiol 89:1057–1066.
- McGaughy J, Sarter M (1995) Behavioral vigilance in rats: task validation and effects of age, amphetamine, and benzodiazepine receptor ligands. Psychopharmacology (Berl) 117:340–357.
- McGaughy J, Kaiser T, Sarter M (1996) Behavioral vigilance following infusions of 192 IgG-saporin into the basal forebrain: selectivity of the behavioral impairment and relation to cortical AChE-positive fiber density. Behav Neurosci 110:247–265.
- Mentis MJ, Sunderland T, Lai J, Connolly C, Krasuski J, Levine B, Friz J, Sobti S, Schapiro M, Rapoport SI (2001) Muscarinic versus nicotinic modulation of a visual task. a PET study using drug probes. Neuropsychopharmacology 25:555–564.
- Michel MC, Murphy TJ, Motulsky HJ (2020) New author guidelines for displaying data and reporting data analysis and statistical methods in experimental biology. Mol Pharmacol 97:49–60.
- Michel V, Yuan Z, Ramsubir S, Bakovic M (2006) Choline transport for phospholipid synthesis. Exp Biol Med (Maywood) 231:490–504.
- Mitchell KM (2004) Acetylcholine and choline amperometric enzyme sensors characterized in vitro and in vivo. Anal Chem 76:1098–1106.
- Muir JL, Dunnett SB, Robbins TW, Everitt BJ (1992) Attentional functions of the forebrain cholinergic systems: effects of intraventricular hemicholinium, physostigmine, basal forebrain lesions and intracortical grafts on a multiple-choice serial reaction time task. Exp Brain Res 89:611–622.
- Nair J, Klaassen AL, Arato J, Vyssotski AL, Harvey M, Rainer G (2018) Basal forebrain contributes to default mode network regulation. Proc Natl Acad Sci U S A 115:1352–1357.
- O'Connell RG, Dockree PM, Robertson IH, Bellgrove MA, Foxe JJ, Kelly SP (2009) Uncovering the neural signature of lapsing attention: electrophysiological signals predict errors up to 20 s before they occur. J Neurosci 29:8604–8611.
- Okuda T, Okamura M, Kaitsuka C, Haga T, Gurwitz D (2002) Single nucleotide polymorphism of the human high affinity choline transporter alters transport rate. J Biol Chem 277:45315–45322.
- Onizuka S, Kasaba T, Tamura R, Takasaki M (2008) Lidocaine increases intracellular sodium concentration through a Na+-H+ exchanger in an identified Lymnaea neuron. Anesth Analg 106:1465–1472, table of contents.
- Paolone G, Lee TM, Sarter M (2012) Time to pay attention: attentional performance time-stamped prefrontal cholinergic activation, diurnality, and performance. J Neurosci 32:12115–12128.
- Paolone G, Angelakos CC, Meyer PJ, Robinson TE, Sarter M (2013a) Cholinergic control over attention in rats prone to attribute incentive salience to reward cues. J Neurosci 33:8321–8335.
- Paolone G, Mallory CS, Koshy Cherian A, Miller TR, Blakely RD, Sarter M (2013b) Monitoring cholinergic activity during attentional performance in mice heterozygous for the choline transporter: a model of cholinergic capacity limits. Neuropharmacology 75:274–285.
- Parikh V, Sarter M (2006) Cortical choline transporter function measured in vivo using choline-sensitive microelectrodes: clearance of endogenous and exogenous choline and effects of removal of cholinergic terminals. J Neurochem 97:488–503.
- Parikh V, Pomerleau F, Huettl P, Gerhardt GA, Sarter M, Bruno JP (2004) Rapid assessment of in vivo cholinergic transmission by amperometric detection of changes in extracellular choline levels. Eur J Neurosci 20:1545–1554.
- Parikh V, Apparsundaram S, Kozak R, Richards JB, Sarter M (2006) Reduced expression and capacity of the striatal high-affinity choline transporter in hyperdopaminergic mice. Neuroscience 141:379–389.
- Parikh V, Kozak R, Martinez V, Sarter M (2007) Prefrontal acetylcholine release controls cue detection on multiple timescales. Neuron 56:141– 154.
- Parikh V, St. Peters M, Blakely RD, Sarter M (2013) The presynaptic choline transporter imposes limits on sustained cortical acetylcholine release and attention. J Neurosci 33:2326–2337.
- Perry RJ, Hodges JR (1999) Attention and executive deficits in Alzheimer's disease. A critical review. Brain 122:383–404.

- Phillips KB, Sarter M (2020) Addiction vulnerability and the processing of significant cues: sign-, but not goal-, tracker perceptual sensitivity relies on cue salience. Behav Neurosci 134:133–143.
- Pitchers KK, Phillips KB, Jones JL, Robinson TE, Sarter M (2017a) Diverse roads to relapse: a discriminative cue signaling cocaine availability is more effective in renewing cocaine seeking in goal trackers than sign trackers and depends on basal forebrain cholinergic activity. J Neurosci 37:7198–7208.
- Pitchers KK, Kane LF, Kim Y, Robinson TE, Sarter M (2017b) 'Hot' vs. 'cold' behavioural-cognitive styles: motivational-dopaminergic vs. cognitive-cholinergic processing of a Pavlovian cocaine cue in sign- and goal-tracking rats. Eur J Neurosci 46:2768–2781.
- Ray A, Lindahl E, Wallner B (2010) Model quality assessment for membrane proteins. Bioinformatics 26:3067–3074.
- Sarkar A, Kellogg GE (2010) Hydrophobicity-shake flasks, protein folding and drug discovery. Curr Top Med Chem 10:67–83.
- Sarter M, Parikh V (2005) Choline transporters, cholinergic transmission and cognition. Nat Rev Neurosci 6:48–56.
- Sarter M, Fritschy JM (2008) Reporting statistical methods and statistical results in EJN. Eur J Neurosci 28:2363–2364.
- Sarter M, Phillips KB (2018) The neuroscience of cognitive-motivational styles: sign- and goal-trackers as animal models. Behav Neurosci 132:1–12.
- Sarter M, Lustig C (2020) Forebrain cholinergic signaling: wired and phasic, not tonic, and causing behavior. J Neurosci 40:712–719.
- Sarter M, Lustig C, Blakely RD, Koshy Cherian A (2016) Cholinergic genetics of visual attention: human and mouse choline transporter capacity variants influence distractibility. J Physiol Paris 110:10–18.
- Sarter M, Avila C, Kucinski A, Donovan E (2021) Make a left turn: corticostriatal circuitry mediating the attentional control of complex movements. Mov Disord 36:535–546.
- Simon JR, Atweh S, Kuhar MJ (1976) Sodium-dependent high affinity choline uptake: a regulatory step in the synthesis of acetylcholine. J Neurochem 26:909–922.
- St Peters M, Cherian AK, Bradshaw M, Sarter M (2011) Sustained attention in mice: expanding the translational utility of the SAT by incorporating the Michigan controlled access response port (MICARP). Behav Brain Res 225:574–583.
- Stamm M, Staritzbichler R, Khafizov K, Forrest LR (2013) Alignment of helical membrane protein sequences using AlignMe. PLoS One 8:e57731.

- Stamm M, Staritzbichler R, Khafizov K, Forrest LR (2014) AlignMe–a membrane protein sequence alignment web server. Nucleic Acids Res 42: W246–W251.
- Swets J, Tanner WP Jr, Birdsall TG (1961) Decision processes in perception. Psychol Rev 68:301–340.
- Tehovnik EJ, Sommer MA (1997) Effective spread and timecourse of neural inactivation caused by lidocaine injection in monkey cerebral cortex. J Neurosci Methods 74:17–26.
- Thienel R, Kellermann T, Schall U, Voss B, Reske M, Halfter S, Sheldrick AJ, Radenbach K, Habel U, Shah NJ, Kircher T (2009) Muscarinic antagonist effects on executive control of attention. Int J Neuropsychopharmacol 12:1307–1317.
- Tsirigos KD, Peters C, Shu N, Käll L, Elofsson A (2015) The TOPCONS web server for consensus prediction of membrane protein topology and signal peptides. Nucleic Acids Res 43:W401–W407.
- Turchi J, Sarter M (1997) Cortical acetylcholine and processing capacity: effects of cortical cholinergic deafferentation on crossmodal divided attention in rats. Brain Res Cogn Brain Res 6:147–158.
- Turchi J, Sarter M (2000) Cortical cholinergic inputs mediate processing capacity: effects of 192 IgG-saporin-induced lesions on olfactory span performance. Eur J Neurosci 12:4505–4514.
- Venkatesan S, Lambe EK (2020) Chrna5 is essential for a rapid and protected response to optogenetic release of endogenous acetylcholine in prefrontal cortex. J Neurosci 40:7255–7268.
- Voytko ML, Olton DS, Richardson RT, Gorman LK, Tobin JR, Price DL (1994) Basal forebrain lesions in monkeys disrupt attention but not learning and memory. J Neurosci 14:167–186.
- Webb B, Sali A (2016) Comparative protein structure modeling using MODELLER. Curr Protoc Protein Sci 86:2.9.1–2.9.37.
- Yamamura HI, Snyder SH (1972) Choline: high-affinity uptake by rat brain synaptosomes. Science 178:626–628.
- Yang Y, Gritton H, Sarter M, Aton SJ, Booth V, Zochowski M (2021) Thetagamma coupling emerges from spatially heterogeneous cholinergic neuromodulation. PLoS Comput Biol 17:e1009235.
- Yarnall A, Rochester L, Burn DJ (2011) The interplay of cholinergic function, attention, and falls in Parkinson's disease. Mov Disord 26:2496–2503.
- Zimmermann L, Stephens A, Nam SZ, Rau D, Kübler J, Lozajic M, Gabler F, Söding J, Lupas AN, Alva V (2018) A completely reimplemented MPI bioinformatics toolkit with a new HHpred server at its core. J Mol Biol 430:2237–2243.