

Nicotine exploits a COPI-mediated process for chaperone-mediated up-regulation of its receptors

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Chronic exposure to nicotine up-regulates high sensitivity nicotinic acetylcholine receptors (nAChRs) in the brain. This up-regulation partially underlies addiction and may also contribute to protection against Parkinson's disease. nAChRs containing the $\alpha 6$ subunit ($\alpha 6^*$ nAChRs) are expressed in neurons in several brain regions, but comparatively little is known about the effect of chronic nicotine on these nAChRs. We report here that nicotine up-regulates $\alpha 6^*$ nAChRs in several mouse brain regions (substantia nigra pars compacta, ventral tegmental area, medial habenula, and superior colliculus) and in neuroblastoma 2a cells. We present evidence that a coat protein complex I (COPI)-mediated process mediates this up-regulation of $\alpha 6^*$ or $\alpha 4^*$ nAChRs but does not participate in basal trafficking. We show that $\alpha 6\beta 2\beta 3$ nAChR up-regulation is prevented by mutating a putative COPI-binding motif in the $\beta 3$ subunit or by inhibiting COPI. Similarly, a COPI-dependent process is required for up-regulation of $\alpha 4\beta 2$ nAChRs by chronic nicotine but not for basal trafficking. Mutation of the putative COPI-binding motif or inhibition of COPI also results in reduced normalized Förster resonance energy transfer between $\alpha 6\beta 2\beta 3$ nAChRs and ϵ COP subunits. The discovery that nicotine exploits a COPI-dependent process to chaperone high sensitivity nAChRs is novel and suggests that this may be a common mechanism in the up-regulation of nAChRs in response to chronic nicotine.

INTRODUCTION

One of the earliest discoveries of chronic nicotine exposure was the observation of an increased number of nicotinic acetylcholine receptors (nAChRs; termed up-regulation) (Marks et al., 1983; Schwartz and Kellar, 1983; Breese et al., 1997; Mamede et al., 2007; Nashmi et al., 2007). Classically, up-regulation of nAChRs has been defined as an increase in nAChR protein, as identified by increased agonist binding (Marks et al., 1983; Benwell et al., 1988; Flores et al., 1992; Peng et al., 1994), but we now define up-regulation as a change in receptor number, stoichiometry, and trafficking (Lester et al., 2009; Miwa et al., 2011). Often, an increase in total binding as opposed to cell surface binding has been reported. Therefore, up-regulation of nAChRs involves an increase in nAChR abundance throughout the cell (ER, Golgi, etc.) and not exclusively on the plasma membrane (PM). There have been several hypotheses regarding the mechanism of up-regulation, but a consensus is emerging that nicotine acts inside the cell to enhance a critical step(s) in the maturation process

of nAChRs (Salette et al., 2005). This intracellular enhancement process has been characterized as pharmacological chaperoning (Kuryatov et al., 2005; Lester et al., 2009), and it occurs at the nanomolar concentrations thought to persist in the brain for hours after a person smokes. Research from many laboratories indicates that up-regulation of $\alpha 4\beta 2$ nAChRs occurs through a similar process in human brains, animal brains, cultured neurons, and clonal transfected cell lines (Nashmi et al., 2007; Mukhin et al., 2008; Miwa et al., 2011; Srinivasan et al., 2011; Lester et al., 2012).

Up-regulation of nAChRs in response to chronic nicotine plays a major role in nicotine dependence and, perhaps, in the inverse correlation between a person's history of tobacco use and his or her susceptibility to Parkinson's disease (Ritz et al., 2007; Koob, 2009). Individual deletions of the $\alpha 4$, $\alpha 6$, or $\beta 2$ nAChR subunits are sufficient to block the self-administration of nicotine in mice (Pons et al., 2008), whereas the selective reexpression of these deleted subunits in the ventral tegmental area (VTA) is sufficient to reinstate self-administration of nicotine (Pons et al., 2008; Brunzell et al., 2010).

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Abbreviations used in this paper: COPI, coat protein complex I; ERES, ER exit sites; nAChR, nicotinic acetylcholine receptor; Neuro-2a, neuroblastoma 2a; NFRET, normalized Förster resonance energy transfer; PM, plasma membrane; PMID, PM integrated density; ROI, regions of interest; SEP, supercliptic-pHluorin; Snc, substantia nigra pars compacta; TIRFM, total internal reflection fluorescence microscopy; VTA, ventral tegmental area.

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Furthermore, nicotine self-administration can be blocked by the selective antagonism of $\alpha 6^*$ (*, IUPHAR nomenclature, “other subunits may be present”) (Jackson et al., 2009) or $\alpha 4^*$ (Yoshimura et al., 2007) nAChRs. From this, it is clear that the nAChRs mediating nicotine addiction include those that contain $\alpha 4$, $\alpha 6$, and $\beta 2$ subunits (Picciotto et al., 1998; Tapper et al., 2004; Pons et al., 2008).

Despite the clarity of $\alpha 4\beta 2$ nAChR up-regulation, there have been conflicting reports of up-regulation, down-regulation, or no change in response to chronic nicotine for $\alpha 6^*$ nAChRs (Tumkosit et al., 2006; Perez et al., 2008; Walsh et al., 2008). Here, we used mice expressing $\alpha 6$ -enhanced green fluorescence protein (eGFP) nAChR subunits to test for in vivo up-regulation of $\alpha 6^*$ nAChRs using concentrations of nicotine comparable to those produced by smoking in humans. We determined that this up-regulation occurs in all four brain areas that robustly express $\alpha 6^*$ nAChRs: the VTA, the substantia nigra pars compacta (SNc), the medial habenula, and the superior colliculus.

We then used $\alpha 6$ subunits tagged with supercliptic-pHluorin (SEP), a pH-sensitive eGFP analogue, expressed in neuroblastoma cells to analyze the mechanism of $\alpha 6^*$ nAChR up-regulation. SEP has been used to study vesicle dynamics and synaptic delivery (Miesenböck et al., 1998; Mani and Ryan, 2009) as well as the trafficking of AMPA and GABA receptors (Jacob et al., 2005; Jaskolski et al., 2009; Lin et al., 2009; Araki et al., 2010), and we have used $\alpha 4$ -SEP* to study nAChR trafficking after chronic exposure to nicotine (Richards et al., 2011). Here, we show that nicotine-induced up-regulation fails to occur with inhibition of coat protein complex I (COPI)-mediated retrograde traffic from the Golgi to the ER (either through the mutation of a putative COPI-binding motif [$\beta 3$ -KKK] or through the use of the COPI inhibitor CI-976). COPI is a protein complex that coats vesicles for retrograde transport of proteins from the cis end of the Golgi to the ER (Brandizzi and Barlowe, 2013). COPI is a heptameric protein complex, and di-lysine-traffic motifs, including KKxx and KxKxx, facilitate protein binding to COPI (Jackson et al., 2012; Spang, 2013). Inhibition of COPI-mediated trafficking resulted in a decrease of nAChR density in the ER and a complementary increase of nAChR density in the Golgi. Inhibition of COPI had no effect on basal insertion of nAChRs in the PM but prevented their up-regulation during chronic nicotine treatment. Therefore, we propose that the cycling of nAChRs between the Golgi and ER (via COPI) is necessary for the up-regulation of $\alpha 4\beta 2$ and $\alpha 6\beta 2\beta 3$ nAChRs. Our data suggest that this may be a common mechanism of nAChR up-regulation by nicotine, and that the manipulation of Golgi-ER cycling may lead to novel therapeutic strategies for nicotine cessation or neuroprotection against Parkinson’s disease.

MATERIALS AND METHODS

Reagents

The β -COP antibody (rabbit polyclonal) was obtained from Abcam (AB2899). CI-976 and (–)-nicotine hydrogen tartrate were obtained from Sigma-Aldrich.

Mice and chronic nicotine administration

The construction of $\alpha 6$ -GFP mice has been described previously (Mackey et al., 2012). All experiments were conducted in accordance with the guidelines for care and use of animals provided by the National Institutes of Health, and protocols were approved by the Institutional Animal Care and Use Committee at the California Institute of Technology.

Chronic nicotine administration

Mice were kept on a standard 12-h light/dark cycle at 22°C and given food and water ad libitum. Chronic nicotine or saline was administered to mice using miniosmotic pumps (model 2002; Alzet). On the day of pump implantation, saline or (–)-nicotine hydrogen tartrate (Sigma-Aldrich) was prepared freshly and loaded into the pump to deliver nicotine at 2 mg/kg/h or 0.4 mg/kg/h. This concentration provides maximal nAChR up-regulation and a peripheral blood concentration that is near the peak concentration found in the blood of human smokers (Marks et al., 2004). Surgical procedures for the pump implantation have been reported previously (Nashmi et al., 2007).

Cell culture

Mouse neuroblastoma 2a (Neuro-2a) cells were cultured using standard techniques (Xiao et al., 2011). Cells were plated by adding 90,000 cells to poly-D-lysine-coated 35-mm glass-bottom imaging dishes (MatTek Corporation) and cultured in a humidified incubator (37°C, 95% air, 5% CO₂). Cells were transfected with 500 ng of each nAChR subunit plasmid and 250 ng GalT-mCherry or Sec24D-mCherry for assays. Plasmids were mixed with 250 μ l Opti-MEM. Lipofectamine-2000 was separately added to 250 μ l Opti-MEM. After 5 min at 24°C, DNA and Lipofectamine solutions were mixed together and incubated for 25 min at 24°C. The solutions were then added to preplated Neuro-2a cells and incubated for 24 h. After 24 h, the Opti-MEM was removed and replaced with growth medium. 50 or 500 nM of filter-sterilized nicotine was added after replacing the Opti-MEM with standard culture medium ($\alpha 6\beta 2\beta 3$ nAChRs). For $\alpha 4\beta 2$ nAChRs, 100 nM nicotine was used for 48 h (nicotine was added at the time of transfection and then replenished when the media was changed). 20 μ M CI-976 was added with nicotine 24 h before imaging. Cells were imaged 48 h after transfection.

Patch-clamp recordings

For patch-clamp electrophysiology, 50,000 Neuro-2a cells were plated onto sterilized 12-mm glass coverslips (Deckgläser), placed in 35-mm culture dishes, and cultured in a humidified incubator (37°C, 95% air, 5% CO₂). Recorded cells were visualized with an inverted fluorescence microscope (IX71; Olympus) in either bright field or fluorescence (eGFP) mode using a high pressure Hg lamp (HB-10103AF; Nikon). Electrophysiological signals were recorded with an amplifier (Axopatch-1D; Axon Instruments), analogue-to-digital converter (Digidata 1440A; Axon Instruments), and software (pClamp 10.0; Axon Instruments). Patch pipettes were filled with solution containing (mM): 135 K gluconate, 5 KCl, 5 EGTA, 0.5 CaCl₂, 10 HEPES, 2 Mg-ATP, and 0.1 GTP (pH was adjusted to 7.2 with Tris-base, and osmolarity was adjusted to 280–300 mOsm with sucrose). The resistance of patch pipettes was 2–4 M Ω for whole-cell recordings. Junction potential was nulled just before

forming a gigaseal. All recordings were done at 24°C. Data were sampled at 10 kHz and filtered at 2 kHz for whole-cell recordings. Acetylcholine was dissolved in extracellular solution containing (mM) 140 NaCl, 5 KCl, 2 CaCl₂, 1 MgCl₂, 10 HEPES, and 10 glucose (320 mOsm, pH set to 7.3 with Tris-base), and were puffed (0.3 s, 20 psi) onto voltage-clamped Neuro-2a cells (holding potential [V_H] of -50 mV). To avoid receptor desensitization by repetitive ACh application, we applied ACh at ~3-min intervals and continually perfused the recording chamber with extracellular solution. For concentration response studies, we used a rapid superfusion system with 500-ms puffs of agonist (Octaflow II; ALA Scientific Instruments).

Immunostaining and antibodies

For immunostaining, cultured Neuro-2a cells were fixed with 4% paraformaldehyde (15 min), permeabilized in 0.01% Triton X-100 (5 min), and blocked with 10% goat serum (30 min). After two PBS washes, the appropriate primary antibody (1:500) in 1% goat serum was applied for 1 h at 24°C. Cells were washed three times with PBS and incubated with 1% goat serum containing secondary antibody (1:2,000) for 1 h at 24°C. Cells were washed three times with PBS and imaged immediately. Immunostaining used β -COP rabbit polyclonal antibody as the primary (antibody 2899; Abcam) and Alexa Fluor 555-labeled secondary (goat anti-rabbit; Invitrogen) antibodies.

Midbrain α 6-GFP neurons existed in a high enough density to be imaged without staining (direct/inherent fluorescence). Immunohistochemical techniques were required for observing α 6-GFP fluorescence in other regions of the mouse brain. 20- μ m brain sections on slides were rinsed twice for 10 min with PBS and then permeabilized with 0.5% Triton X-100 in PBS for 1 h. Brain sections were then blocked with 4% goat serum (Jackson Immuno-Research Laboratories, Inc.) in PBS for 45 min. The primary antibody was diluted in 4% goat serum in PBS and incubated overnight at 24°C. Brain sections were then washed three times for 15 min with PBS, and the secondary antibody was diluted in 4% goat serum in PBS and incubated for 1 h at 24°C. Finally, brain sections were washed three times for 15 min with PBS and mounted with Vectashield.

Total internal reflection fluorescence microscopy (TIRFM)

Cultured Neuro-2a cells were imaged live at 37°C in a stage-mounted culture dish incubator (Warner Instruments) using methods similar to TIRFM assays in cultured cortical neurons (Richards et al., 2011). TIRFM enables the visualization of fluorescently labeled intracellular molecules within ~250 nm of the cell-coverslip interface. TIRFM images were obtained using an inverted fluorescence microscope (IX81; Olympus) equipped with a 100 \times 1.45 NA oil objective (PlanApo; Olympus) and a stepper motor (Thorlabs) to control the position of the fiber optic and TIRFM evanescent field illumination. Just before imaging, growth medium was exchanged for extracellular solution (150 mM NaCl, 4 mM KCl, 10 mM HEPES, 2 mM MgCl₂, 2 mM CaCl₂, and 10 mM glucose), adjusted to the appropriate pH (5.4 or 7.4). SEP was excited at 488 nm with an air-cooled argon laser (IMA101040ALS; Melles Griot). Images were captured with a back-illuminated EMCCD camera (iXon DU-897; Andor). We acidified the imaging dish by perfusing the bath, normally at pH 7.4, with an otherwise identical solution adjusted to pH 5.4. The PM integrated density (PMID) was determined by taking an initial TIRFM image of each cell at pH 7.4, followed by acidification of the solution and a subsequent low pH image (pH 5.4). Low pH images were used to demarcate ER-localized nAChRs, which were subtracted from the total footprint to determine the PMID of nAChRs. The figures show a single imaging session that is representative of at least three sessions performed on separate days with similar or identical results.

Insertion events were quantified using methods similar to those published previously (Richards et al., 2011). TIRFM measurements to detect membrane insertion events were performed with consecutive 200-ms frames. Insertion events were defined as punctate regions of fluorescence appearing at the membrane. Insertion event data are represented in terms of insertion per 10 μ m² of area per minute of observation.

Spectral confocal imaging

Imaging experiments were performed in live Neuro-2a cells at 37°C in a stage-mounted culture dish incubator (Warner Instruments). For immunostaining, fixed Neuro-2a cells were imaged. A C1si laser-scanning confocal microscope (Eclipse; Nikon) equipped with a 60 \times 1.2 NA VC Plan Apo water objective and 32 photomultiplier tubes was used for confocal imaging. Before an imaging session, cell culture medium was replaced with phenol red-free CO₂-independent Leibovitz (L-15) medium (Invitrogen). eGFP and mCherry were excited at 488 and 561 nm, respectively. For each cell, we focused on a plane containing the most ER exit sites (ERES), Golgi bodies, or COPI vesicles, and sequential images of eGFP and mCherry fluorescence were obtained. eGFP and mCherry fluorescence emission spectra were captured and images were unmixed using standard spectra acquired from cells expressing eGFP, mCherry, or Alexa Fluor 555 alone. For quantification, ERES, Golgi, or COPI regions of interest (ROI) were manually demarcated using intensity-based thresholding, and the raw integrated densities were measured for each cell using ImageJ. The total Sec24D fluorescence in ERES and the total GalT fluorescence per cell were quantified. Error bars for measurements indicate the SEM, and p-values are based on a two-tailed *t* test. The figures show a single imaging session that is representative of at least three sessions performed on separate days with similar or identical results.

For direct fluorescence imaging of in vivo α 6-GFP* nAChRs, images were acquired with the 60 \times 1.2 NA VC Plan Apo water objective at wavelengths between 496 and 561 nm using a 488-nm line of an argon laser. Images were collected at 12-bit intensity over 512 \times 512 pixels, a resolution of 2.5 nm, a 61.3- μ m pinhole, and a pixel dwell time of 8.4 μ s. Autofluorescence was separated from GFP fluorescence using techniques that have been published previously (Nashmi et al., 2007). Image analysis of cell counts and mean intensities were done with ImageJ.

Normalized Förster resonance energy transfer (NFRET)

The general methods for Förster resonance energy transfer from sensitized acceptor emission have been described previously (Srinivasan et al., 2011, 2012a). For these studies, Neuro-2a cells were transfected with α 6-mCherry, β 2wt, ϵ COP-GFP, and β 3wt or β 3_{AAA} nAChR subunits. Cells transfected with either α 6-mCherry or ϵ COP-GFP alone were included in all imaging sessions to control for pixel saturation and spectral bleedthrough. Live cells were imaged with a C1si laser-scanning confocal microscope (Eclipse; Nikon). During acquisition of images, cells were focused on a plane where the ϵ COP-GFP fluorescence most resembled the pattern for the endogenous COPI staining (see Fig. 4 E). GFP and mCherry images were acquired with 488- and 561-nm laser lines, and images were linearly unmixed by using reference spectra. Reference spectra were acquired from Neuro-2a cells transfected with α 6-mCherry or ϵ COP-GFP during the same imaging session. After linear unmixing, NFRET was calculated using the PixFRET ImageJ plug-in as described previously (Moss et al., 2009; Srinivasan et al., 2011, 2012a).

Plasmid constructs

Mouse α 6 with a C-terminal fusion of an SEP tag was constructed by PCR amplification using the forward primer 5'-CATGGTTGG-CTGGTATGATCAGTAAAGGAGAAGAAGCTT-3' and the reverse primer 5'-ATGGATGAACATATACAATAGGGAATAGCGGCACCT-3',

which overlap with sequences within the C-terminal end of the $\alpha 6$ -coding sequence. This PCR product was then cloned directly into the vector containing the $\alpha 6$ gene using Pfu-Turbo polymerase. The mouse $\beta 2_{\beta 3}$ nAChR subunit was constructed by PCR amplification using the forward primer 5'-ATGGCCCGGTGCTCCAATC-3' and the reverse primer 5'-CCTGCCCTCAGACTGTGGTG-3'. The mouse $\beta 3_{\beta 2}$ nAChR subunit was constructed by PCR amplification using the forward primer 5'-ATGACAGGCTTCTACGGGT-3' and the reverse primer 5'-ACGGTCTGTTCTACTCT-3'. Mouse $\beta 3_{AAA}$ (the KKK mutated to AAA) and the $\beta 2_{\beta 3[AAA]}$ were constructed by PCR amplification using the forward primer 5'-AAATTTCCAGGGCTGCAGCCCAGACTCCTACC-3' and the reverse primer 5'-CCCCCTCACGGTTCCTTACTCTCCGT-3'.

Online supplemental material

Fig. S1 illustrates that the presence of $\beta 3$ subunits increased the PMID of $\alpha 6\beta 4$ nAChRs but is not required for the up-regulation of $\alpha 6\beta 4^*$ nAChRs. Fig. S2 presents a schematic of the various chimeras and mutant nAChR subunits that were used in this paper. Fig. S3 shows the electrophysiological characterization of $\alpha 6\beta 2\beta 3_{WT}$ and $\alpha 6\beta 2\beta 3_{AAA}$ nAChRs after treatment with nicotine. Here, we show that functional up-regulation accompanies the increase in PMID for $\alpha 6\beta 2\beta 3_{WT}$ nAChRs when treated with nicotine, but there is no functional up-regulation with $\alpha 6\beta 2\beta 3_{AAA}$ nAChRs. Fig. S4 illustrates that treatment with CI-976 produces an increase of nAChRs in the Golgi but does not increase export from the ER. Fig. S5 shows that CI-976 treatment does not affect basal PMID of nAChRs. The online supplemental material is available at <http://www.jgp.org/cgi/content/full/jgp.201311102/DC1>.

RESULTS

In vivo up-regulation of $\alpha 6^*$ nAChRs with chronic nicotine

We used a mouse line containing GFP-labeled $\alpha 6^*$ nAChRs to study the up-regulation of $\alpha 6^*$ nAChRs (Mackey et al., 2012). $\alpha 6$ -GFP nAChR subunits assemble appropriately with other nonlabeled nAChR subunits, traffic properly, and function in electrophysiological assays in a manner comparable to that of WT nAChRs (Mackey et al., 2012).

Direct fluorescence of $\alpha 6$ -GFP* nAChRs was apparent in the VTA and SNc of saline- or nicotine-treated mice (Fig. 1 A). The fluorescence intensity of $\alpha 6$ -GFP* nAChRs in the VTA and SNc of mice administered nicotine at 2 mg/kg/h for 10 d was increased twofold compared with that of saline-treated mice ($P < 0.005$; Fig. 1, B₁ and C₁). In addition, we used cumulative percentage plots to compare saline and nicotine treatments (Fig. 1, B₂ and C₂). Cumulative percentage plots are similar to a cumulative frequency analysis and allow us to compare the full range of $\alpha 6$ -GFP fluorescence intensities between saline and nicotine treatments. In all four brain regions, we can see that the percentage of low, mid, and high $\alpha 6$ -GFP fluorescence intensities are shifted to the right

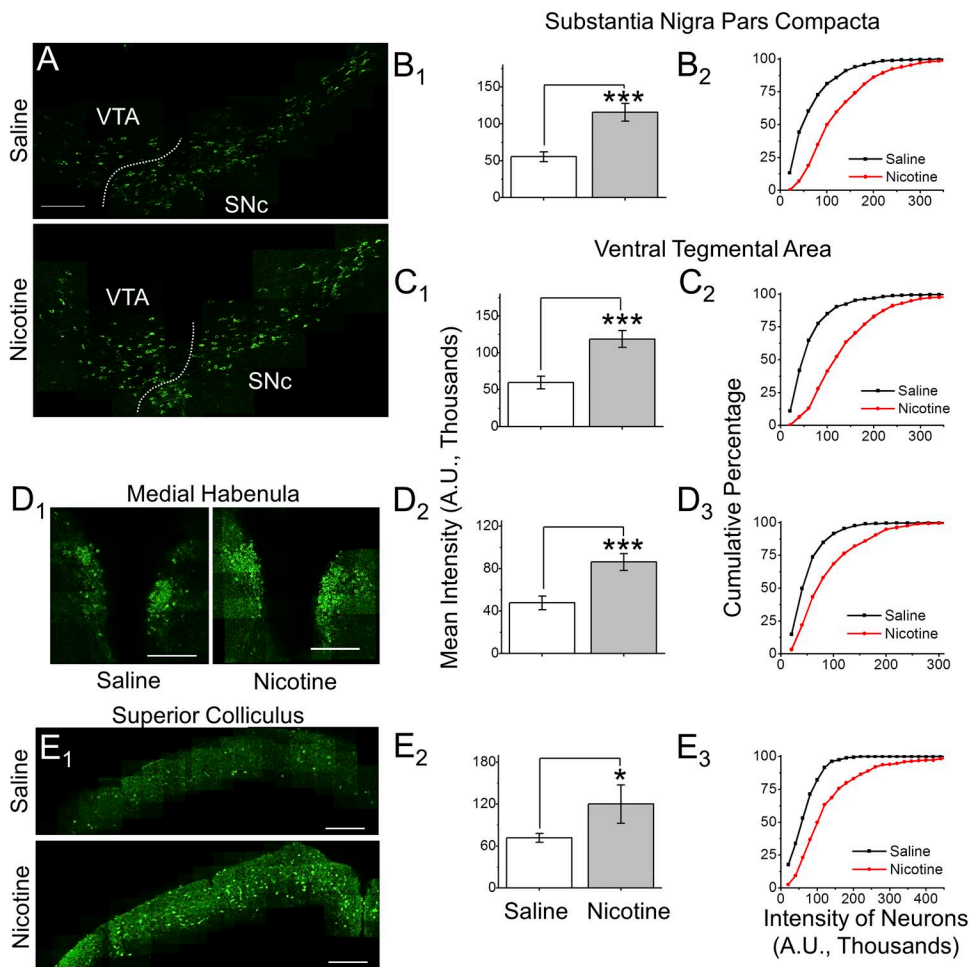


Figure 1. Chronic nicotine up-regulates $\alpha 6$ -GFP* nAChRs in four brain regions. (A) Direct fluorescence imaging of SNc and VTA in mice expressing $\alpha 6$ -GFP subunits after treatment with saline or nicotine (2 mg/kg/h, 10 d). (D₁ and E₁) Immunohistochemical imaging of medial habenula and superior colliculus in mice expressing $\alpha 6$ -GFP subunits after treatment with saline or nicotine. For all images, montages were compiled from individual 60 \times images. (A and E₁) Bars, 100 μ m. (D₁) Bars, 20 μ m. (B₁ and C₁) Quantification of $\alpha 6$ -GFP intensities in SNc and VTA of saline- and nicotine-treated mice ($n = 6$ mice). (D₂ and E₂) Quantification of $\alpha 6$ -GFP intensities in medial habenula and superior colliculus of saline- and nicotine-treated mice ($n = 6$ mice). (D₃ and E₃) Quantification of $\alpha 6$ -GFP immunofluorescence of medial habenula and superior colliculus neurons in saline- and nicotine-treated mice ($n = 4-5$ mice). (B₂, C₂, D₃, and E₃) Cumulative percentage plots show changes in $\alpha 6$ -GFP fluorescence in all brain regions with chronic nicotine. ***, $P < 0.005$; *, $P < 0.05$.

(higher intensity) after chronic nicotine treatment (Fig. 1, B₂ and C₂). Nicotine at 0.4 mg/kg/h also increased $\alpha 6$ -GFP fluorescence intensity in VTA ($60.8 \pm 10.9\%$ increase) and SNc ($70.0 \pm 1.4\%$ increase) neurons ($P < 0.05$ compared with saline; Fig. 2). Similar to 2 mg/kg/h nicotine treatments, the 0.4-mg/kg/h nicotine treatment shifted the full range of $\alpha 6$ -GFP fluorescence intensities to the right in cumulative percentage plots (Fig. 2, D₁ and D₂).

2 mg/kg/h nicotine also increased the fluorescence intensity of medial habenula and superior colliculus $\alpha 6$ -GFP* nAChRs (81% [$P < 0.005$] and 68% [$P < 0.05$], respectively; Fig. 1, D and E). In cumulative percentage plots, we can also see that the full range of $\alpha 6$ -GFP fluorescence intensities is shifted to the left after chronic nicotine treatment (Fig. 1, D₁ and D₂).

$\alpha 6$ -SEP* nAChRs assemble functional receptors, and $\beta 3$ subunits increase $\alpha 6$ * nAChR function

To perform a systematic study on $\alpha 6$ * nAChR up-regulation, we used an expression system, Neuro-2a cells, which has been optimized to avoid many artifacts of “overexpressing” PM proteins. The appearance of functional PM proteins in this system is linear and unsaturated as a function of parameters such as level of transfected cDNAs, trafficking machinery, oligomerization, and anchoring (Moss et al., 2009).

SEP undergoes 488-nm excitation at neutral pH (~ 7.4) but not under acidic conditions (Richards et al., 2011). $\alpha 6$ -SEP* nAChRs in the Golgi and secretory vesicles (~ 6.7 and < 6.5 pH, respectively) are not fluorescent

(Asokan and Cho, 2002; Jaskolski et al., 2009). When exposed to a neutral buffer (pH ~ 7.4) both ER- and PM-resident nAChRs are fluorescent. Upon perfusing an acidic buffer (pH 5.4), PM-resident nAChR fluorescence is quenched, whereas the ER component remains fluorescent. Thus, SEP allows the differentiation of receptors within the ER from those in the PM. As in a previous study using SEP tagged to $\alpha 4$ nAChRs (Richards et al., 2011), we fused SEP to the C terminus of $\alpha 6$ subunits so that it was positioned on the luminal side of organelles in the secretory pathway.

nAChRs with eGFP subunits have little or no functional difference (i.e., peak current amplitude, decay time) from WT subunits (Drenan et al., 2008; Richards et al., 2011; Srinivasan et al., 2011, 2012b; Xiao et al., 2011) and, indeed, we found no detectable functional difference between $\alpha 6$ -SEP $\beta 2\beta 3$ nAChRs and $\alpha 6$ -eGFP $\beta 2\beta 3$ nAChRs (Fig. 3). Inclusion of the $\beta 3$ subunit increased ACh-induced peak currents by threefold compared with those obtained with $\alpha 6$ -eGFP $\beta 2$ nAChRs (Fig. 3 B; $P < 0.05$). Inclusion of the $\beta 3$ subunit also reduced the concentration of ACh at which we observed maximal peak amplitude from 300 μ M ($\alpha 6$ -eGFP $\beta 2$) to 3 μ M ($\alpha 6$ -eGFP $\beta 2\beta 3$) (Fig. 3). Concentration–response studies of ACh and (–)-nicotine on $\alpha 6$ -eGFP $\beta 2\beta 3$ nAChRs yielded EC₅₀ values of 0.31 ± 0.10 μ M and 0.12 ± 0.08 μ M, respectively (Fig. 3, C and D), which agree with those obtained with linked concatamers containing ($\alpha 6\beta 2$)₂ $\beta 3$ nAChRs (Kuryatov and Lindstrom, 2011).

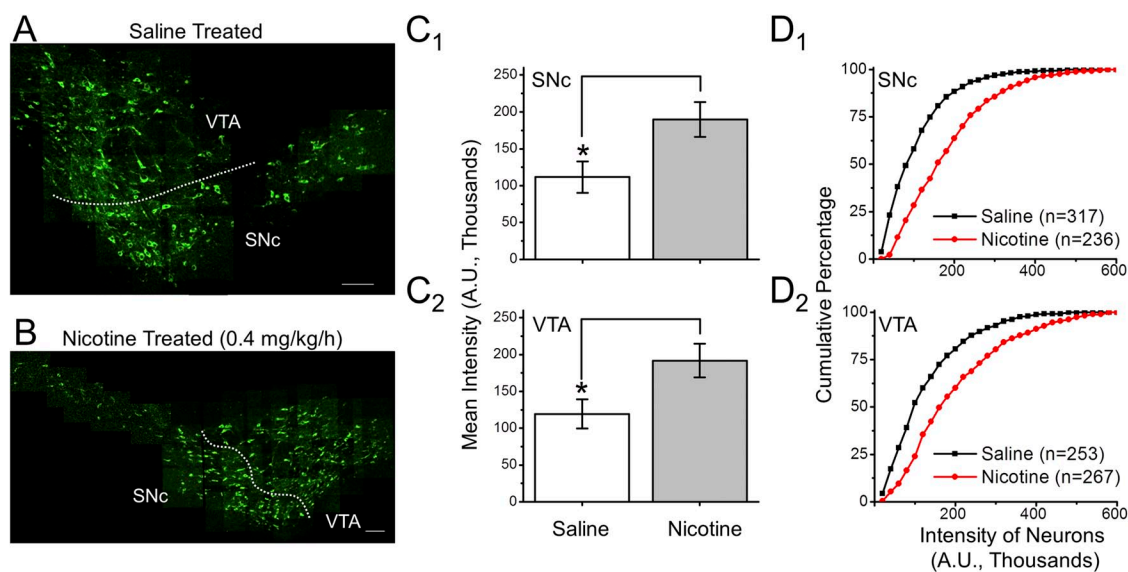


Figure 2. Low concentrations of nicotine (0.4 mg/kg/h) up-regulate $\alpha 6$ -GFP* nAChRs in midbrain neurons. (A and B) Direct fluorescence imaging of VTA and SNc neurons in mice expressing $\alpha 6$ -GFP subunits after treatment with saline or nicotine (0.4 mg/kg/h, 10 d). Bars, 100 μ m. (C₁ and C₂) Quantification of $\alpha 6$ -GFP intensities of neurons in saline- or nicotine-treated mice ($n = 4$ mice). (D₁ and D₂) Cumulative percentage plots show changes in $\alpha 6$ -GFP fluorescence in the SNc and VTA with chronic nicotine (0.4 mg/kg/h; $n = 4$ mice). *, $P < 0.05$.

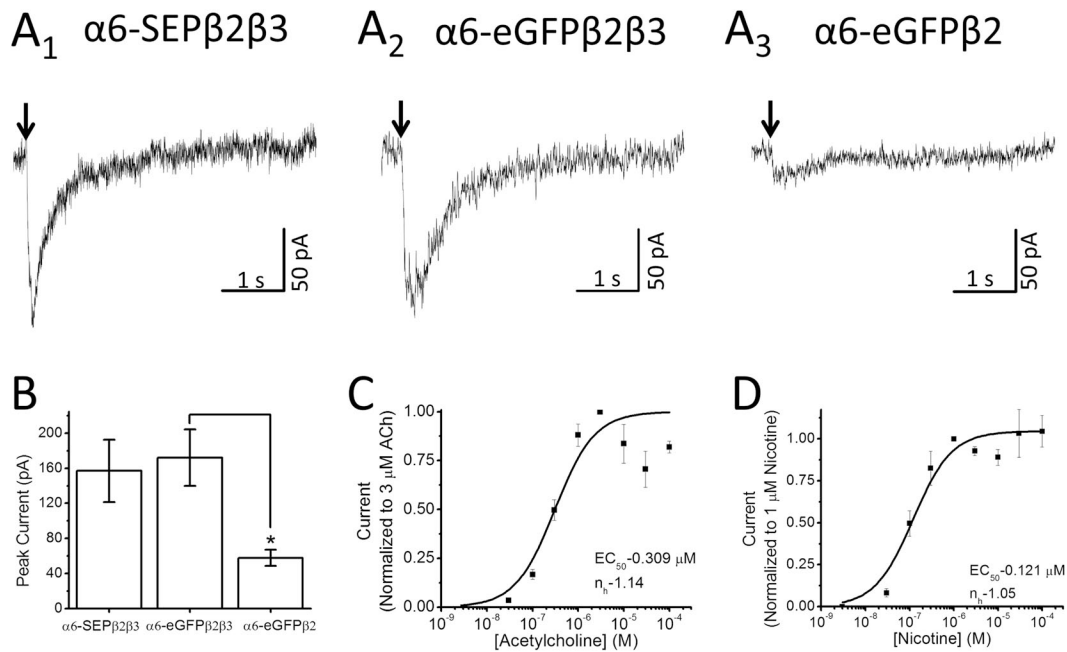


Figure 3. The α6-SEP nAChRs are functional. Representative whole-cell ACh-induced currents of (A₁) α6-SEPβ2β3 nAChRs (3 μM ACh), (A₂) α6-eGFPβ2β3 nAChRs (3 μM ACh) and (A₃) α6-eGFPβ2 nAChRs (300 μM ACh). (B) Comparison of whole-cell ACh-induced current amplitudes of α6-SEPβ2β3, α6-eGFPβ2β3, and α6-eGFPβ2 nAChRs (*n* = 7–11). (C) Concentration–response of ACh on α6-eGFPβ2β3 nAChRs (*n* = 15). (D) Concentration–response of (–)-nicotine on α-eGFPβ2β3 nAChRs (*n* = 14). *, *P* < 0.05.

Incorporation of β3 subunits increases the density of α6* nAChRs at the PM

We used TIRFM and α6-SEP* nAChRs transfected in Neuro-2A cells to quantify the density of receptors in the peripheral ER versus the PM. Fig. 4 A shows TIRFM images of Neuro-2a cells expressing α6-SEPβ2 or α6-SEPβ2β3 nAChRs at neutral (7.4) and acidic (5.4) pH.

As reported previously (Tumkosit et al., 2006), α6β2β3 nAChRs had a higher density on the PM than α6β2 nAChRs; the PMID of α6β2β3 nAChRs was three times that of α6β2 nAChRs (*P* < 0.0001; Fig. 4 B). We also found a 1.5-fold increase in the PMID of α6β4β3 nAChRs compared with that of α6-SEPβ4* nAChRs (Fig. S1). We found that, when expressed in Neuro-2a

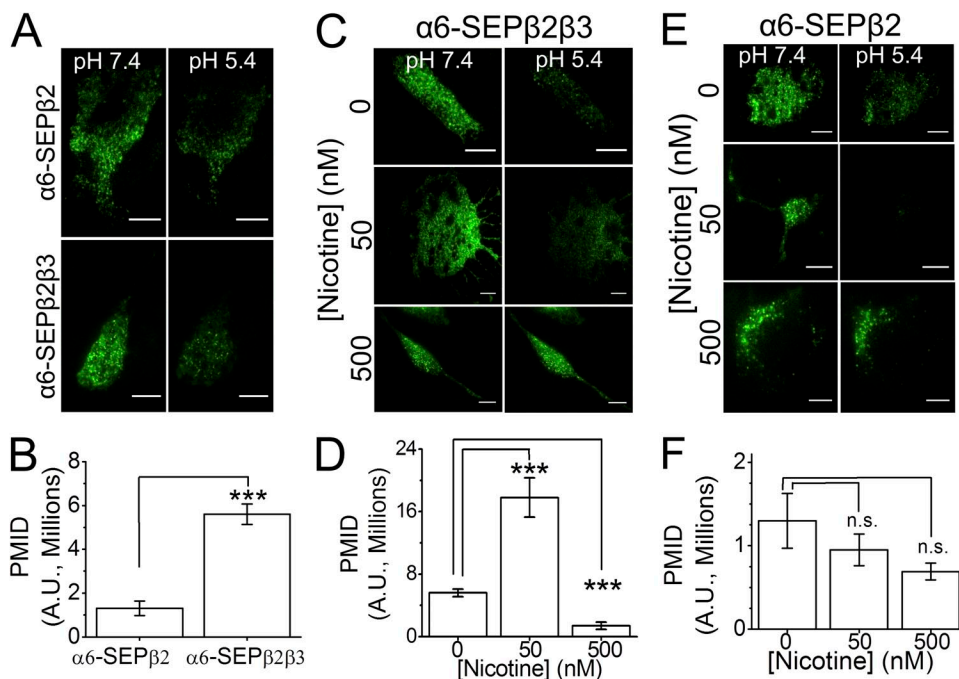


Figure 4. The β3 subunit increases the density of α6* nAChRs at the PM and is necessary for the nicotine-induced up-regulation of α6β2* nAChRs. (A, C, and E) Representative TIRFM images of Neuro-2a cells transfected with α6-SEP and a combination of β2 with or without β3 nAChR subunits at basic (pH 7.4) and acidic (pH 5.4) conditions. Nicotine was added at listed concentrations 24 h before imaging. (B, D, and F) PMID for α6-SEP* nAChRs. Bars, 10 μm (*n* = 19–23). Data are mean values ± SEM. n.s., not significant; ***, *P* < 0.0001.

cells, $\alpha 6$ -SEP $\beta 3$ nAChRs resided almost entirely in the ER ($94.5 \pm 2.2\%$ ER localization; not depicted). Collectively, these data show that the $\beta 3$ subunits, combined with $\beta 2$ or $\beta 4$ nAChR subunits, increase the density of $\alpha 6^*$ nAChRs on the PM.

Confirmation that $\beta 3$ subunits enhance nicotine-induced up-regulation of $\alpha 6\beta 2^*$ nAChRs

We next assessed the effect of chronic nicotine on $\alpha 6\beta 2^*$ nAChRs. We examined $\alpha 6^*$ nAChRs exposed to 50 or 500 nM nicotine for 24 h. 50- and 500-nM nicotine concentrations, respectively, mimic the steady-state and peak concentrations of nicotine found in the brains of smokers (Marks et al., 2004). 50 nM nicotine also equals the plasma concentration produced by 14–21-mg nicotine patches (Schnoll et al., 2013). We observed a significant ($P < 0.0001$) increase in the PMID of $\alpha 6$ -SEP $\beta 2\beta 3$ nAChRs after treatment with 50 nM nicotine, but a significant ($P < 0.0001$) decrease in PMID after 500-nM nicotine treatment (Fig. 4, C and D). $\alpha 6$ -SEP $\beta 2$ nAChRs showed no change in PMID after 50-nM nicotine treatment and a decrease in PMID after 500-nM nicotine treatment (Fig. 4, E and F). 50 or 500 nM nicotine induced a greater than threefold increase in $\alpha 6$ -SEP $\beta 4^*$ nAChRs PMIDs in the presence or absence of the $\beta 3$ subunit (Fig. S1).

Dynamic visualization of SEP nAChRs enables us to see when nAChRs arrive at the PM, as they appear as a region of punctate fluorescence (Richards et al., 2011; Fig. 5, A and B_{1–3}). We quantified the frequency of these vesicular insertion events at the PM (Fig. 5). After exposure to 50 nM nicotine, we observed a threefold increase in the insertion events of $\alpha 6$ -SEP $\beta 2\beta 3$ nAChRs ($P < 0.05$) and no change in the insertion events of $\alpha 6$ -SEP $\beta 2$

nAChRs (Fig. 5, C and D). These data correlated with the observed nicotine-induced changes of PM intensities for $\alpha 6$ -SEP $\beta 2\beta 3$ and $\alpha 6$ -SEP $\beta 2$ nAChRs. Specifically, we note a similarity between the increase in PMID and rate of insertion of $\alpha 6$ -SEP $\beta 2\beta 3$ nAChRs (both are an approximate threefold increase) (Fig. 5, C and D). This suggests that the up-regulation of $\alpha 6\beta 2\beta 3$ nAChRs is caused primarily by increased insertion into the PM, with no major contribution from decreased internalization of receptors from the PM (Lomazzo et al., 2011).

Despite the different PMIDs between $\alpha 6$ -SEP $\beta 2\beta 3$ and $\alpha 6$ -SEP $\beta 2$ nAChRs, they showed the same rate of insertion under basal conditions (~ 12 ins/min). Therefore, the difference in the PM density of $\alpha 6$ -SEP $\beta 2\beta 3$ and $\alpha 6$ -SEP $\beta 2$ nAChRs may be caused by differential rates of internalization or stability of these nAChR subtypes at the PM, suggesting that the $\beta 3$ nAChR subunit may provide a stabilizing effect.

A KKK motif within the $\beta 3$ M3–M4 loop governs the up-regulation of $\alpha 6\beta 2\beta 3$ nAChRs

Our data suggest that $\beta 3$ subunits markedly influence the trafficking and dynamics of $\alpha 6\beta 2^*$ nAChRs. We next sought to identify specific motifs within $\beta 3$ subunits that may play a role in the trafficking and dynamics of $\alpha 6\beta 2^*$ nAChRs.

We engineered a series of chimeras: (a) a chimera with a $\beta 3$ M3–M4 loop replacing the $\beta 2$ M3–M4 loop in a $\beta 2$ subunit ($\beta 2_{\beta 3}$) and (b) a chimera with a $\beta 2$ M3–M4 loop replacing the $\beta 3$ M3–M4 loop in a $\beta 3$ subunit ($\beta 3_{\beta 2}$) (see Fig. S2). $\alpha 6$ -SEP $\beta 2\beta 2_{\beta 3}$, $\alpha 6$ -SEP $\beta 2_{\beta 3}$, $\alpha 6$ -SEP $\beta 2\beta 3_{\beta 2}$, and $\alpha 6$ -SEP $\beta 3_{\beta 2}$ nAChRs exhibited typical TIRFM footprints and PM-ER fluorescence that were similar to $\alpha 6$ -SEP $\beta 2^*$ nAChRs (Fig. 6). $\alpha 6$ -SEP $\beta 2\beta 2_{\beta 3}$ and $\alpha 6$ -SEP $\beta 2_{\beta 3}$

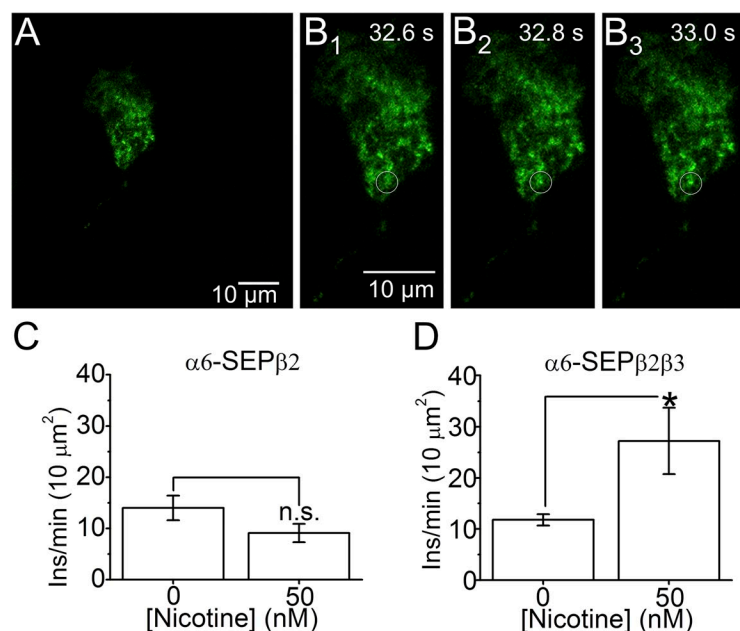


Figure 5. Visualization of membrane insertion events. (A) TIRFM image of a Neuro-2a cell transfected with $\alpha 6$ -SEP $\beta 2\beta 3$ nAChRs and excited with a 488-nm laser. Images were taken with a frame rate of 200 ms. (B₁) Frame directly preceding an insertion event at the PM. (B₂) Frame showing the insertion and (B₃) the frame directly after the observation of the insertion event. The white circle marks the location of the insertion. (C and D) Frequency of insertion events for $\alpha 6$ -SEP $\beta 2^*$ nAChRs without or with nicotine treatment (50 nM, 24 h) calculated as insertion events per minute ($n = 4$). Bars, 10 μm . Data are mean values \pm SEM. n.s., not significant; *, $P < 0.05$.

nAChRs up-regulated with 50 nM nicotine, whereas $\alpha 6$ -SEP $\beta 3_{\beta 2}$ and $\alpha 6$ -SEP $\beta 2\beta 3_{\beta 2}$ nAChRs showed either no change or a decrease in the PMID with 50 nM nicotine, suggesting that the $\beta 3$ subunit M3–M4 domain specifically regulates nicotine-induced up-regulation but may not control the basal trafficking of $\alpha 6^*$ nAChRs (Fig. 6, A₂ and B₂).

Unique among nAChR subunits, the $\beta 3$ nAChR subunit possesses a KKK motif within its M3–M4 loop. Dilysine trafficking motifs, including KKxx and KxKxx, facilitate protein binding to COPI, a component of the Golgi–ER retrograde transport machinery (Jackson et al., 2012). Although these motifs typically occur at cytoplasmic C termini, the KKK found in the $\beta 3$ subunit satisfies both of these consensus motifs. Thus, we hypothesized that the KKK may act as a COPI recognition site. We mutated the ³⁸¹KKK³⁸³ motif in the $\beta 3$ M3–M4 loop to ³⁸¹AAA³⁸³ to create $\beta 2_{\beta 3[AAA]}$ and $\beta 3_{AAA}$ nAChR subunits. Mutant nAChRs exhibited a typical TIRFM footprint and showed comparable overall nAChR expression to WT receptors (Fig. 6, C and D). We observed a significant decrease in the number of ER resident $\alpha 6$ -SEP $\beta 2\beta 2_{\beta 3[AAA]}$ and $\alpha 6$ -SEP $\beta 2\beta 3_{AAA}$ nAChRs (Fig. 6, C₁ and D₁). In addition, the $\alpha 6$ -SEP $\beta 2\beta 2_{\beta 3[AAA]}$ and $\alpha 6$ -SEP $\beta 2\beta 3_{AAA}$ nAChRs did not increase the PMID when treated with 50 nM nicotine (Fig. 5, C and D), suggesting that the KKK motif was essential for the nicotine-induced up-regulation of $\alpha 6$ -SEP $\beta 2^*$ nAChRs. Similarly, we found a 44% increase in peak ACh-induced currents for $\alpha 6$ -eGFP $\beta 2\beta 3$ nAChRs after treatment with 50 nM nicotine (24 h) but no increase for $\alpha 6$ -eGFP $\beta 2\beta 3_{AAA}$

nAChRs (Fig. S3). $\alpha 6$ -eGFP $\beta 2\beta 3$ or $\alpha 6$ -eGFP $\beta 2\beta 3_{AAA}$ nAChRs did not show a difference in ACh-induced currents under basal conditions. These results correlated with the observed PMIDs of $\alpha 6$ -SEP $\beta 2\beta 3$ and $\alpha 6$ -SEP $\beta 2\beta 3_{AAA}$ nAChRs (Fig. 6).

Nicotine up-regulates nAChRs by binding to and chaperoning ER-localized receptors, which are then trafficked to the PM (Srinivasan et al., 2011). The formation of additional ERES with nicotine is a direct indicator of an event immediately after the nAChR chaperoning process (Srinivasan et al., 2011). To assess the effect of nicotine on $\alpha 6\beta 2\beta 3_{AAA}$ nAChR chaperoning, we quantified the formation of condensed ERES (Fig. 7, A and B). Neuro-2a cells were cotransfected with either $\alpha 6$ -eGFP $\beta 2\beta 3$ or $\alpha 6$ -eGFP $\beta 2\beta 3_{AAA}$ nAChRs and Sec24D-mCherry (a marker for active ERES) (Srinivasan et al., 2011). 50 nM nicotine caused a significant ($P < 0.005$) threefold increase of ERES in $\alpha 6$ -eGFP $\beta 2\beta 3$ nAChRs but no change in ERES with $\alpha 6$ -eGFP $\beta 2\beta 3_{AAA}$ nAChRs (Fig. 7 B).

The $\beta 3$ KKK motif inhibits up-regulation by preventing retrograde Golgi to ER nAChR transport

We observed a decrease in the ER density of $\beta 3$ -AAA nAChRs compared with WT nAChRs (Fig. 6, C and D) and hypothesized that loss of the KKK (COPI-binding motif) disrupted trafficking between the Golgi and ER. To test this hypothesis, we studied how WT and $\beta 3$ -AAA nAChRs localized within the Golgi. Fig. 7 (C and D) shows colocalization of $\alpha 6$ -eGFP $\beta 2\beta 3$ and $\alpha 6$ -eGFP $\beta 2\beta 3_{AAA}$ nAChRs with the Golgi marker GalT-mCherry. $\beta 3_{AAA}$ subunits caused a significant increase in the Pearson

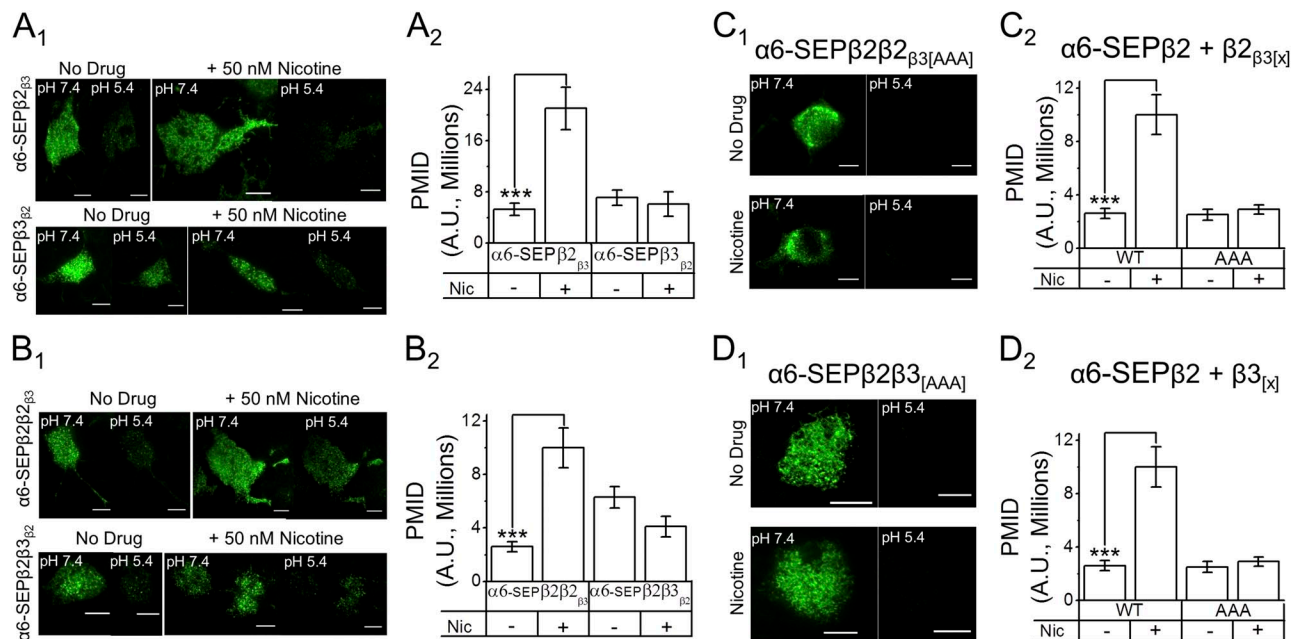


Figure 6. The $\beta 3$ KKK motif participates in nAChR up-regulation. (A₁, B₁, C₁, and D₁) Representative TIRFM images of Neuro-2a cells transfected with $\alpha 6$ -SEP and cotransfected with combinations of $\beta 2$, $\beta 2_{\beta 3}$, $\beta 3_{\beta 2}$, $\beta 2_{\beta 3[AAA]}$, and $\beta 3_{AAA}$ subunits. 50 nM nicotine was added 24 h after transfection. Bars, 10 μ m. (A₂, B₂, C₂, and D₂) PMID for nAChRs ($n = 11$ –27). ***, $P < 0.0001$ (ANOVA).

correlation coefficient for $\alpha 6$ -eGFP and GalT-mCherry colocalization (Fig. 7 D₁; $P < 0.0001$) and a significant ($P < 0.01$) increase in the $\alpha 6$ -eGFP $\beta 2\beta 3_{AAA}$ nAChR density when compared with $\alpha 6$ -eGFP $\beta 2\beta 3$ nAChRs, within the Golgi (Fig. 7 D₂). Therefore, the removal of the KKK motif increased the density of $\alpha 6\beta 2\beta 3$ nAChRs in the Golgi compartment.

To determine whether COPI Golgi-ER trafficking is deficient, we assessed colocalization by immunostaining for endogenous COPI in Neuro-2a cells transfected with $\alpha 6$ -eGFP $\beta 2\beta 3$ and $\alpha 6$ -eGFP $\beta 2\beta 3_{AAA}$ nAChRs (Fig. 7, E, F₁, and F₂). We found that $\alpha 6$ -eGFP $\beta 2\beta 3_{AAA}$ nAChRs exhibited a significant, twofold decrease in density in COPI vesicles when compared with $\alpha 6$ -eGFP $\beta 2\beta 3$ nAChRs (Fig. 7 F₁; $P < 0.01$). This was accompanied by a significant decrease in the pixel-based Pearson correlation coefficient among WT and $\beta 3$ -AAA nAChRs ($P < 0.0001$; Fig. 7 F₂). Despite this decrease, we observed no difference in the $\alpha 6$ -eGFP* whole-cell fluorescence intensity between $\alpha 6$ -eGFP $\beta 2\beta 3$ nAChRs and $\alpha 6$ -eGFP $\beta 2\beta 3_{AAA}$ nAChRs (20.2 ± 3.1 and 18.3 ± 4.8 RFUs [millions], respectively). Thus, the potential contribution from degradation is unlikely.

Upon treatment with 50 nM nicotine, we observed a twofold increase in the $\alpha 6$ -eGFP $\beta 2\beta 3$ nAChR density in COPI vesicles when compared with non-nicotine-treated conditions. In contrast, there was no increase in $\alpha 6$ -eGFP $\beta 2\beta 3_{AAA}$ nAChRs (Fig. 7 F₁). This suggests that after chronic nicotine treatment, an increase of $\alpha 6^*$ nAChRs in COPI vesicles accompanies the up-regulation observed on the PM. Collectively, these data show that a defect in the recognition between COPI vesicles and $\alpha 6$ -eGFP $\beta 2\beta 3_{AAA}$ nAChRs results in decreased retrograde movement of these nAChRs back into the ER.

The COPI inhibitor CI-976 blocks up-regulation but not basal trafficking of $\alpha 6\beta 2\beta 3$ and $\alpha 4\beta 2$ nAChRs

Our data suggest that Golgi-ER cycling of $\alpha 6\beta 2\beta 3$ nAChRs is necessary for up-regulation during chronic exposure to nicotine. Next, we investigated whether this also applies to other subtypes. For these experiments, we used CI-976, which inhibits COPI-mediated retrograde trafficking at concentrations $\geq 20 \mu\text{M}$ (Yang et al., 2005, 2011). Inhibition of COPII vesicle budding occurs at a somewhat higher concentration (50 μM) (Brown et al., 2008).

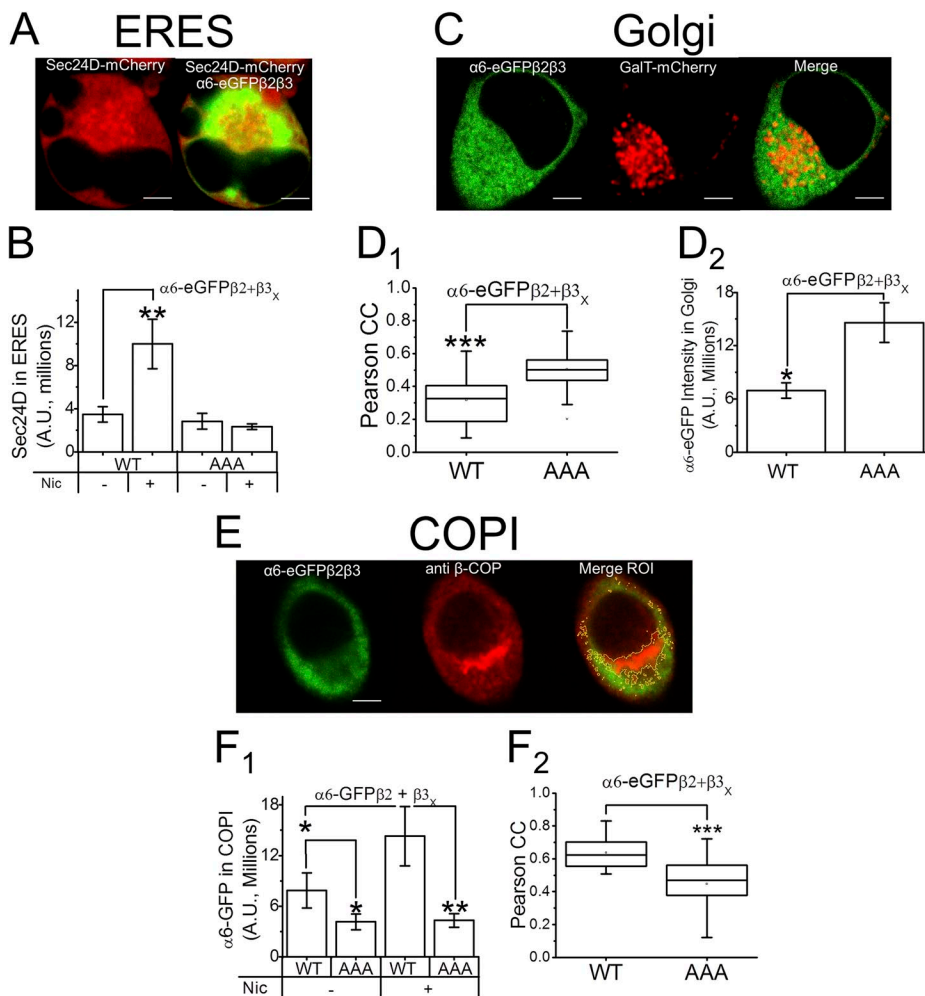


Figure 7. $\beta 3$ -AAA nAChRs show increased density in Golgi despite no change in ER export. Confocal image of $\alpha 6$ -eGFP $\beta 2\beta 3$ and Sec24D-mCherry (A), $\alpha 6$ -eGFP $\beta 2\beta 3$ with Golgi marker GalT-mCherry (C), and $\alpha 6$ -eGFP $\beta 2\beta 3$ nAChRs stained for endogenous β -COP (E). Quantification of: Sec24D fluorescence in ERES per cell for $\alpha 6$ -eGFP $\beta 2\beta 3_{WT}$ and $\alpha 6$ -eGFP $\beta 2\beta 3_{AAA}$ nAChRs (B) ($n = 21$ – 31); Pearson correlation between $\alpha 6$ -eGFP* and GalT-mCherry fluorescence for $\alpha 6$ -eGFP $\beta 2\beta 3_{WT}$ and $\alpha 6$ -eGFP $\beta 2\beta 3_{AAA}$ nAChRs (D₁) ($n = 30$ – 35); $\alpha 6$ -eGFP* intensity in Golgi for $\alpha 6$ -eGFP $\beta 2\beta 3_{WT}$ and $\alpha 6$ -eGFP $\beta 2\beta 3_{AAA}$ nAChRs (D₂) ($n = 30$ – 35); $\alpha 6$ -eGFP fluorescence in COPI vesicles for $\alpha 6$ -eGFP $\beta 2\beta 3_{WT}$ and $\alpha 6$ -eGFP $\beta 2\beta 3_{AAA}$ nAChRs (F₁) ($n = 23$ – 25); and Pearson correlation for the colocalization of $\alpha 6$ -eGFP* fluorescence with COPI fluorescence in $\alpha 6$ -eGFP $\beta 2\beta 3_{WT}$ and $\alpha 6$ -eGFP $\beta 2\beta 3_{AAA}$ nAChRs (F₂) ($n = 23$ – 24). Bars, 5 μm . *, $P < 0.01$; **, $P < 0.005$; ***, $P < 0.0001$ (ANOVA).

First, we assessed the effects of 20 μM CI-976 on ERES (Fig. S4 A). CI-976 did not affect the basal number of ERES in $\alpha 6\text{-GFP}\beta 2\beta 3$ nAChRs but did prevent the increase in ERES with chronic nicotine treatment (50 nM). This is the same trend that was observed with $\alpha 6\text{-GFP}\beta 2\beta 3_{\text{AAA}}$ nAChRs (see Fig. 6 D₂). With CI-976 treatment, we also noted a significant ($P < 0.005$) increase in the $\alpha 6\text{-GFP}$ density in the Golgi (Fig. S4 B) and a significant ($P < 0.05$) increase in the Pearson correlation coefficient for colocalization of $\alpha 6\text{-eGFP}$ and GalT-mCherry (Fig. S4 C). These data mimic the results seen with $\alpha 6\text{-GFP}\beta 2\beta 3_{\text{AAA}}$ nAChRs (Fig. 7) and suggest that CI-976 can be used to study the inhibition of Golgi-ER cycling in nAChRs.

COPI may play a role in anterograde transport in addition to its role in retrograde transport (Yang et al., 2011). We conducted experiments to determine whether the concentration of CI-976 we used inhibited COPI-mediated anterograde traffic of nAChRs. We studied $\alpha 4\beta 4$ and $\alpha 4\beta 2_{\text{enhanced-ER-export}}$ (also called $\alpha 4\beta 2\text{-DM}$ nAChRs), which both have a high rate of passage from ER to the PM (Richards et al., 2011; Srinivasan et al., 2011). We reasoned that if CI-976 reduced the PMID of

either $\alpha 4\beta 4$ or $\alpha 4\beta 2\text{-DM}$ nAChRs, it would indicate an inhibitory effect on COPI-mediated anterograde traffic. We observed no change in the PMID of $\alpha 4\beta 4$ or $\alpha 4\beta 2\text{-DM}$ nAChRs after CI-976 treatment (Fig. S5). We also note (again) that COPI did not change the level of basal ERES (Fig. S4 A), arguing against the idea that exit from the ER is affected by CI-976. Collectively, these data suggest that CI-976 effects are exerted on COPI's retrograde pathway only.

$\alpha 6\beta 2\beta 3$ nAChRs treated with CI-976 produced similar trends compared to $\alpha 6\beta 2\beta 3_{\text{AAA}}$ nAChRs (compare Figs. 6 D₂ to 8 B₁). As usual, 50 nM nicotine treatment produced a significant ($P < 0.005$) increase in the PMID of $\alpha 6\text{-SEP}\beta 2\beta 3$ nAChRs (Fig. 8 B₁). Treatment with 20 μM CI-976 alone did not differ from the basal conditions seen in $\alpha 6\text{-SEP}\beta 2\beta 3$ nAChRs (no change in PMID), but when cells were coexposed to CI-976 and nicotine, up-regulation of $\alpha 6\text{-SEP}\beta 2\beta 3$ nAChRs was blocked (Fig. 8 B₁). Previous analyses show that 100 nM nicotine is an appropriate concentration for the study of $\alpha 4\text{-SEP}\beta 2$ nAChR up-regulation (Richards et al., 2011; Srinivasan et al., 2011). We observed a significant ($P < 0.05$) increase in the $\alpha 4\text{-SEP}\beta 2$ nAChR PMID with

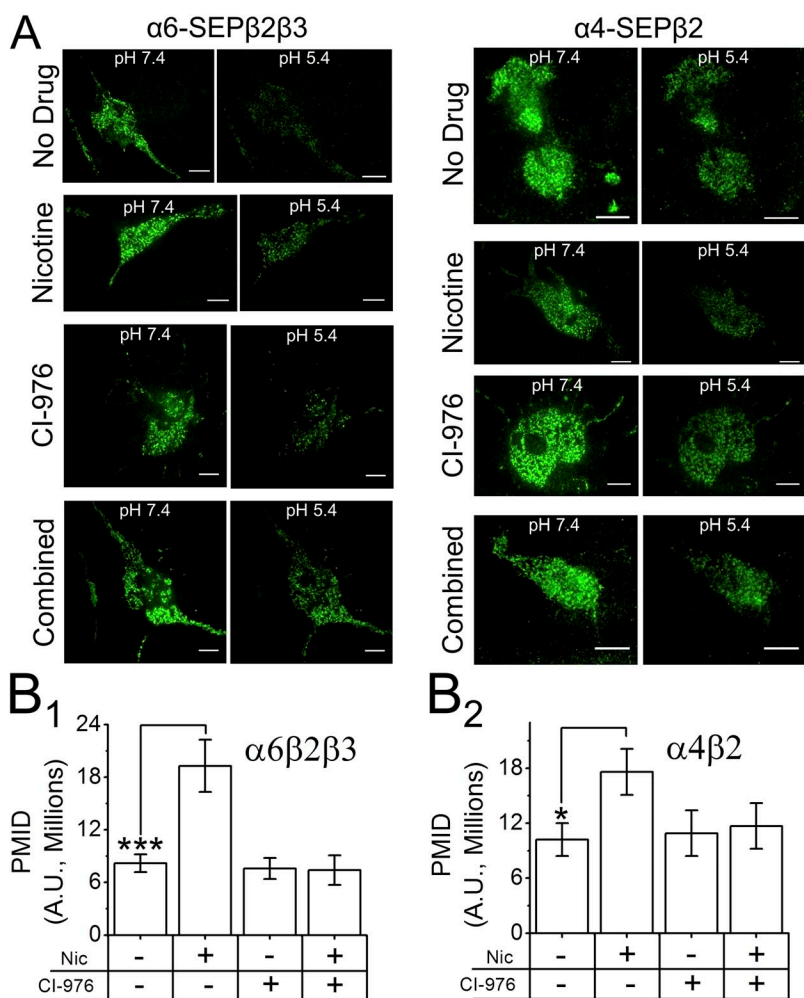


Figure 8. Blocking Golgi-ER cycling prevents up-regulation of nAChRs. (A) Representative TIRFM images of $\alpha 4\text{-SEP}\beta 2$ (right) or $\alpha 6\text{-SEP}\beta 2\beta 3$ (left) nAChR subunits. Nicotine (50 nM $\alpha 6\text{-SEP}\beta 2\beta 3$ nAChRs or 100 nM $\alpha 4\text{-SEP}\beta 2$ nAChRs) and/or 20 μM CI-976 were added 24 h before imaging. Bars, 10 μm . Quantification of PMID for $\alpha 6\text{-SEP}\beta 2\beta 3$ nAChRs (B₁) or $\alpha 4\text{-SEP}\beta 2$ nAChRs (B₂) in Neuro-2a cells with drug treatments: no drug, nicotine, CI-976, or combined ($n = 17\text{--}23$). *, $P < 0.05$; ***, $P < 0.0001$ (ANOVA).

100 nM nicotine and, similar to our observations in $\alpha 6$ -SEP $\beta 2\beta 3$ nAChRs, treatment of $\alpha 4$ -SEP $\beta 2$ nAChRs with CI-976 alone did not differ from basal conditions (no change in PMID) (Fig. 8 B₂). When both CI-976 and nicotine were present with $\alpha 4$ -SEP $\beta 2$ nAChRs, up-regulation was prevented (Fig. 8 B₂). These results suggest that a COPI-dependent process, related to Golgi-ER cycling, is necessary for the nicotine-induced up-regulation of $\alpha 4\beta 2$ as well as $\alpha 6\beta 2\beta 3$ nAChRs.

NFRET measurements of the $\alpha 6\beta 2\beta 3$ - ϵ COP interaction

We asked whether nAChRs and COPI display NFRET. Neuro-2a cells were transfected with $\alpha 6$ -mCherry, $\beta 2$, and $\beta 3$ (WT and $\beta 3$ -AAA) nAChR subunits along with ϵ COP-eGFP (Fig. 9 A). Fig. 9 A shows that NFRET between $\alpha 6$ -mCherry and ϵ COP-eGFP exhibited patterns similar to endogenous β COP staining (see Fig. 7 E). Evidently, $\alpha 6$ -mCherry and ϵ COP-eGFP colocalize within ~ 100 Å.

We asked whether the $\beta 3$ -AAA mutation or CI-976 can inhibit the interaction between $\alpha 6\beta 2\beta 3$ and ϵ COP. nAChRs with the $\beta 3$ -AAA mutation consistently shifted pixel-based NFRET amplitude distributions to lower values (Fig. 9, A-C). Treatment with CI-976 also consistently shifted pixel-based NFRET amplitude distributions to lower values (Fig. 9, A-C). Similarly, when we assigned an NFRET value to each cell by calculating the average

NFRET of the cell's nonzero FRET pixels, the NFRET among cells was significantly decreased in the presence of nAChRs with the $\beta 3$ -AAA mutation or treatment with CI-976 (Fig. 9 C). In addition to NFRET, we used ϵ COP-GFP fluorescence to demarcate ROI for COPI vesicles, and then examined the density of $\alpha 6^*$ nAChRs in the COPI ROI (as in Fig. 7 F₁). $\alpha 6$ -eGFP $\beta 2\beta 3_{AAA}$ nAChRs (no drug) and WT $\alpha 6$ -eGFP $\beta 2\beta 3$ nAChRs (treated with CI-976) exhibited a significant decrease in their density within COPI vesicles (Fig. 9 D; $P < 0.005$). This was accompanied by a significant decrease in the pixel-based Pearson correlation coefficient (Fig. 9 E). Collectively, these results demonstrate that disruption of the KKK motif or inhibition of COPI destabilizes the interaction between COPI and $\alpha 6\beta 2\beta 3$ nAChRs.

DISCUSSION

$\alpha 6^*$ nAChRs up-regulate after chronic nicotine treatment

This paper documents the *in vivo* up-regulation of $\alpha 6^*$ nAChR protein at a sustained nicotine concentration comparable to that in human smokers. We found that up-regulation occurs in the four major brain regions that express $\alpha 6$ -GFP nAChRs: VTA, SNc, medial habenula, and superior colliculus (see Figs. 1 and 2).

The $\alpha 6$ and $\beta 3$ nAChR subunits have nearly identical expression patterns in the brain, as reported by analysis

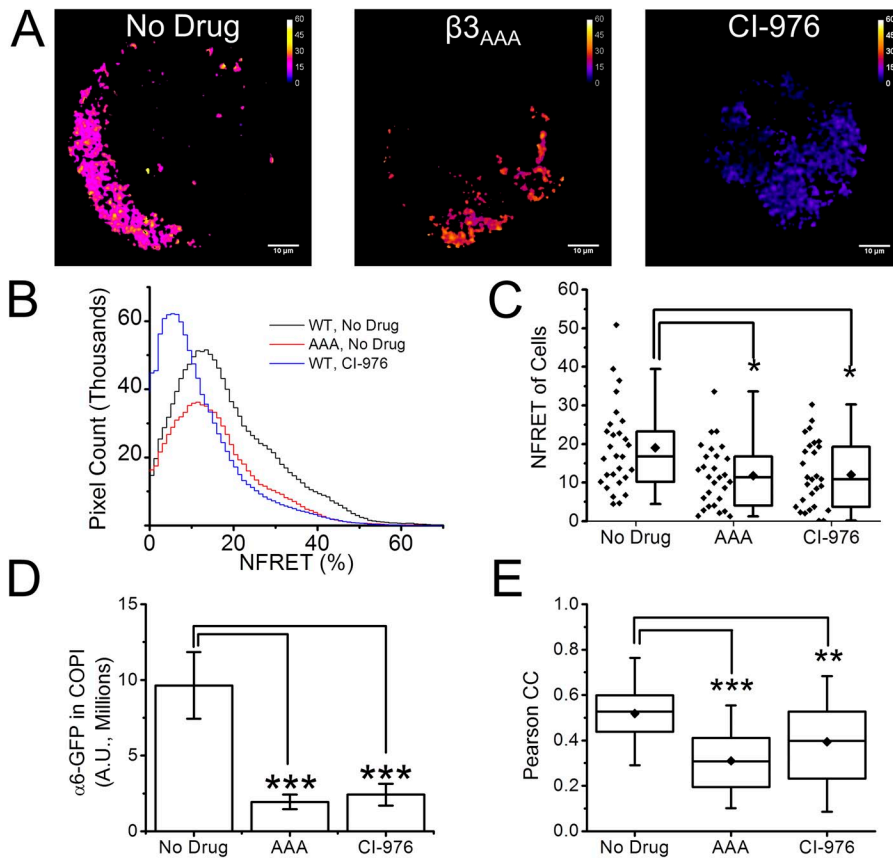


Figure 9. $\beta 3$ -AAA nAChRs and CI-976 treatment decrease interaction between $\alpha 6$ and COPI. NFRET measurements were made using $\alpha 6$ -mCherry and ϵ COP-eGFP transfected in Neuro-2a cells (along with $\beta 2$ and $\beta 3_{wt}$ or $\beta 3_{AAA}$ subunits). (A) Representative NFRET images between $\alpha 6$ -mCherry and ϵ COP-eGFP in Neuro-2a cells. Bars, 10 μ m. (B) Summed histograms of pixel-based NFRET measurements, representative of four independent NFRET assays ($n = 20$ cells). C-E show analysis of the same group of cells ($n = 28$ cells). (C) Another of the four experiments. Scatter-box plot for the mean NFRET values of individual cells. The larger black diamond (\blacklozenge) in the box plot represents the cell-based mean of each condition. (D) $\alpha 6$ -eGFP fluorescence in COPI vesicles for $\alpha 6$ -eGFP $\beta 2\beta 3_{AAA}$ nAChRs and $\alpha 6$ -eGFP $\beta 2\beta 3_{WT}$ nAChRs (without and with 20 μ M CI-976). (E) Pearson correlation for the colocalization of $\alpha 6$ -eGFP* fluorescence with ϵ COP-GFP fluorescence ($n = 28$ cells). *, $P < 0.05$; **, $P < 0.01$; ***, $P < 0.005$ (ANOVA).

of mRNA and autoradiography (Le Novère et al., 1996; Azam et al., 2002; Baddick and Marks, 2011). In heterologous systems, $\alpha 6\beta 2\beta 3$ nAChRs show better function (higher density on the PM, larger peak current response) than $\alpha 6\beta 2$. Therefore, it is appropriate to study effects of chronic nicotine on $\alpha 6\beta 2^*$ nAChRs in the context of coexpressed $\beta 3$ subunits. We found that, at concentrations mimicking the steady-state levels found in human smokers and during nicotine replacement therapy (50 nM nicotine), $\alpha 6$ -SEP $\beta 2\beta 3$ nAChRs up-regulate whereas $\alpha 6$ -SEP $\beta 2$ nAChRs failed to up-regulate. The $\beta 3$ subunit thus appears necessary for the up-regulation of $\alpha 6$ -SEP $\beta 2^*$ nAChRs. This agrees well with Tumkosit et al. (2006), who showed that $\beta 3$ nAChR subunits promote nicotine-induced up-regulation of $\alpha 6^*$ nAChRs.

Our observations of $\alpha 6^*$ up-regulation agree with some previously published work (Parker et al., 2004; Perez et al., 2008; Walsh et al., 2008). Other studies found that chronic nicotine treatment leads to a decrease

in the contribution of $\alpha 6^*$ nAChRs, as identified by binding or dopamine release assays (Lai et al., 2005; Perry et al., 2007; Exley et al., 2013). Several factors may contribute to these disparate findings. First, the up-regulation of $\alpha 6^*$ nAChRs we observed was in midbrain (VTA and SNc) dopaminergic neurons, whereas the studies showing a decrease in $\alpha 6^*$ nAChR contribution were completed in the striatal regions. We have confirmed previously that $\alpha 6$ -GFP * nAChRs in the VTA and SNc are localized exclusively in dopaminergic neurons (Mackey et al., 2012). The difference in striatal (down-regulation) and midbrain (up-regulation) may be explained by a different population of nAChRs ($\alpha 6\alpha 4^*$, $\alpha 6[\text{non}\alpha 4]^*$, $\alpha 6\beta 3^*$, and/or $\alpha 6[\text{non}\beta 3]^*$). Perez et al. (2008) have shown that although $\alpha 6(\text{non}\alpha 4)^*$ nAChRs up-regulate, $\alpha 6\alpha 4^*$ nAChRs down-regulate. We, along with others (Tumkosit et al., 2006), have shown that $\alpha 6^*$ nAChRs may require the $\beta 3$ subunit to up-regulate. Therefore, the population of nAChRs residing in the somata of midbrain dopaminergic neurons may be composed of

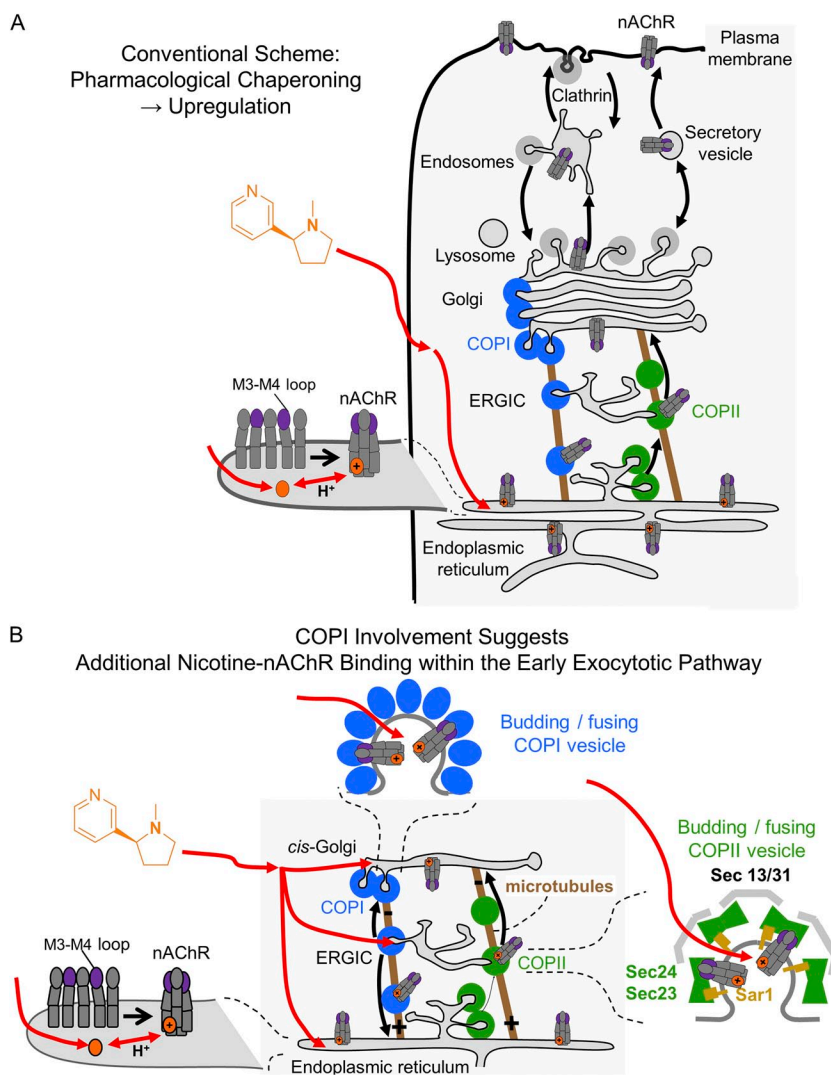


Figure 10. A schematic representation of selective pharmacological chaperoning of nAChRs in response to chronic nicotine. (A) Nicotine interconverts within milliseconds between the protonated membrane-impermeant form and the neutral membrane-permeant form. The latter enters cells and enters the ER (red arrow). Here, the charged form pharmacologically chaperones nAChRs so that increased numbers exit toward the Golgi via the COPII pathway, eventually resulting in receptor up-regulation at the PM. Thus, pharmacological chaperoning by nicotine is thought to underlie the process of nAChR up-regulation by chronic nicotine. (B) This study finds that nicotine-mediated up-regulation also depends on the COPI machinery involved in ER to Golgi and Golgi to ER transport of nAChRs. One possible explanation for the COPI dependence is that nicotine enters additional organelles and binds more extensively than previously thought within the early exocytotic pathway, shown by the additional red arrows leading to both COPI and COPII vesicles, ERGIC, and cis-Golgi. See also the animation, “Nicotine Up-regulates nAChRs”: <http://www.jgp.org/cgi/content/full/jgp.201311102/DC2>.

more $\alpha 6(\text{non}\alpha 4)\beta 3^*$ nAChRs than $\alpha 6\alpha 4^*$ or $\alpha 6(\text{non}\beta 3)^*$ nAChRs. However, there may be more $\alpha 6\alpha 4^*$ or $\alpha 6(\text{non}\beta 3)^*$ nAChRs on these neurons' dopaminergic terminals in the striatum and accumbens. This is supported by a report that there are more $\alpha 6\alpha 4^*$ nAChRs on dopaminergic terminals (in striatum and accumbens) than there are on dopaminergic cell bodies (in VTA and SNC) (Champtiaux et al., 2003).

Second, differing concentrations of nicotine were used in the different studies. The doses of nicotine we used in mice (2 mg/kg/h and 0.4 mg/kg/h) produce plasma concentrations of <500 and <100 nM nicotine, respectively (Marks et al., 2004). Perry et al. (2007) used a dose (6 mg/kg/day) in rats that produces >500 nM plasma nicotine (Nguyen et al., 2004). Our data show that 500 nM nicotine decreased the density of $\alpha 6^*$ nAChRs on the PM (Fig. 4, D and F). Therefore, the differences observed in our current work and in that of others (Lai et al., 2005; Perry et al., 2007; Exley et al., 2013) may be caused by the concentration of nicotine used.

The KKK motif of $\beta 3$ nAChR subunit suggests that a COPI-mediated retrograde pathway participates in up-regulation. A KKK motif appears in the M3–M4 intracellular loop of the $\beta 3$ nAChR subunit. COPI binds to di-lysine motifs (KKxx or KxKxx) (Jackson et al., 2012; Ma and Goldberg, 2013), and the KKK of the $\beta 3$ subunit satisfies both motifs. Measurements of FRET between $\alpha 6\beta 2\beta 3$ nAChRs and ϵ COP indicate that the two molecules approach with ~ 100 Å, and this interaction is decreased by mutating the $\beta 3$ -KKK sequence to $\beta 3$ -AAA (Fig. 9). This suggests that one of these di-lysine motifs in $\beta 3$ mediates a direct interaction between nAChRs and COPI.

We hypothesized that COPI likely plays a role in the retrograde transport of $\alpha 6\beta 2\beta 3$ nAChRs from the Golgi back to the ER. $\alpha 6^*$ nAChRs with a $\beta 3$ -AAA in place of $\beta 3$ -KKK failed to up-regulate on the PM in the presence of nicotine. We found that the ER localization of the $\alpha 6$ -SEP $\beta 2\beta 3_{\text{AAA}}$ nAChRs was decreased compared with $\alpha 6$ -SEP $\beta 2\beta 3$ nAChRs, suggesting that the $\beta 3$ -AAA motif changed the distribution of nAChRs among organelles in the early secretory pathway. Indeed, we found that the $\alpha 6$ -SEP $\beta 2\beta 3_{\text{AAA}}$ nAChRs had a significantly higher localization within the Golgi than the $\alpha 6$ -SEP $\beta 2\beta 3$ nAChRs (Fig. 7, C, D₁, and D₂).

When we add the fact that the number of ERES does not differ between the $\alpha 6\beta 2\beta 3$ and $\alpha 6\beta 2\beta 3_{\text{AAA}}$ nAChRs in basal conditions (e.g., no nicotine) (Fig. 7, A and B), we can conclude that the export (via COPII vesicles) of nAChRs is not affected. Thus, $\alpha 6\beta 2\beta 3_{\text{AAA}}$ nAChRs are exported from the ER at a rate similar to that of $\alpha 6\beta 2\beta 3$ nAChRs. Because export (COPII mediated) is not affected, this suggests that retrieval (COPI mediated) is responsible for the disrupted trafficking. This was confirmed when we observed significantly fewer $\alpha 6\beta 2\beta 3_{\text{AAA}}$ nAChRs in COPI vesicles when compared with $\alpha 6\beta 2\beta 3$ nAChRs

(see Fig. 7, E, F₁, and F₂). Furthermore, $\beta 3$ -AAA nAChRs shifted the NFRET distribution to lower values compared with WT nAChRs (see Fig. 9). Collectively, these data suggest that the disruption of the KKK motif results in deficient retrograde transport of nAChRs between the Golgi and ER. Although one cannot draw specific conclusions about the locus of interference in the cyclic processes involving ER, COPII, cis-Golgi, and COPI (Orci et al., 1997), the data suggest that this disrupted cycling from the Golgi back to the ER prevents up-regulation during exposure to nicotine.

A COPI-dependent process, such as Golgi–ER cycling, is necessary for the up-regulation of nAChRs

We asked whether Golgi–ER cycling plays a role in the other case of nAChRs ($\alpha 4\beta 2$) that up-regulate at smoking-relevant concentrations of nicotine. Because the $\beta 3$ KKK motif is unique in nAChR subunits, we exploited a more general inhibitor approach, CI-976, to block COPI function. On $\alpha 6\beta 2\beta 3$ nAChRs, CI-976 produced results that resembled the pattern observed for $\beta 3$ -WT and $\beta 3$ -AAA nAChRs (compare Figs. 8 B₁ with 6 D₂). Inhibiting Golgi–ER cycling prevented up-regulation of $\alpha 6\beta 2\beta 3$ and $\alpha 4\beta 2$ nAChRs after nicotine treatment. This suggests that the cycling of nAChRs between the Golgi and ER is necessary for the up-regulation of nAChRs on the PM.

Previous reports indicate that nAChR up-regulation is initiated within intracellular compartments and that the locus is partially in the ER (see Fig. 10) (Kuryatov et al., 2005; Sallette et al., 2005; Lester et al., 2009; Miwa et al., 2011; Srinivasan et al., 2011). The data presented here suggest novel aspects of the up-regulation of nAChRs, related to general observations that secretory pathways have both regulated and constitutive components. The increased nAChR density on the PM results from increased ER export of $\alpha 6\beta 2\beta 3$ nAChRs (increase in ERES/COPII, leading to increased insertion events). Previous data show that this also occurs in $\alpha 4\beta 2$ nAChRs (Richards et al., 2011; Srinivasan et al., 2011) (depicted in Fig. 10). We show here that chronic nicotine also increases the number of nAChRs in COPI vesicles (Fig. 7 F₁). This suggests that in addition to an increase in ER export, there is also an increase of retrograde movement of nAChRs from the Golgi to the ER. Possibly, $\alpha 6\beta 2\beta 3$ and $\alpha 4\beta 2$ nAChRs are continually cycled between the ER and Golgi through COPI and COPII. This cycling maintains a population of nAChRs in the ER that may be necessary for up-regulation. In the absence of nicotine, disruption of Golgi–ER cycling (either through the $\beta 3$ -AAA mutants or use of CI-976) does not affect PM nAChR density (Figs. 8 B₁ and 6 D₂). However, when Golgi–ER cycling is inhibited and nicotine is present, nicotine cannot produce up-regulated PM nAChRs.

How does nicotine exploit a COPI process to up-regulate PM nAChRs? There are three nonmutually exclusive possibilities. (1) Decreased retrieval of nAChRs

from the Golgi produces an increased number of Golgi-resident nAChRs but a reduced number of ER-resident nAChRs. Individual nAChRs therefore spend a smaller fraction of time in the ER. Perhaps this renders them less available for pharmacological chaperoning by nicotine (Fig. 10 A). Such chaperoning seems to be relatively inefficient, even though it takes place at nicotine concentrations 10–100 times lower than that required for activation of PM nAChRs. This inefficiency could occur because binding of nicotine must await the dissociation of a competing endogenous chaperone protein such as lynx1 (Miwa et al., 2012). (2) Pharmacological chaperoning by nicotine does stabilize pentameric nAChRs in the ER (Whiteaker et al., 1998; Sallette et al., 2005) but is nonetheless a pathological process that could block or prevent one or more modifications required for efficient exit from the Golgi to the PM. If the nAChRs chaperoned by nicotine fail to pass the quality control checks in the Golgi, they would be targeted for retrieval back to the ER via a COPI-dependent process. (3) An intriguing possibility is that nicotine may enter additional organelles within the early exocytotic pathway. Nicotine might remain bound to nAChRs while the nAChRs reside in COPI and/or COPII vesicles (Fig. 10 B). Processes related to pharmacological chaperoning, termed “matchmaking,” “escorting,” or “abduction,” might occur in these vesicles (Lester et al., 2012). Because the ER and cis-Golgi have nearly neutral luminal pH, acid trapping of nicotine (Jia et al., 2003; Lester et al., 2009) probably plays little role in the processes studied here.

Golgi–ER cycling as a new mechanistic target for drug discovery

Discovery of the role played by Golgi–ER cycling in both $\alpha 6\beta 2\beta 3$ and $\alpha 4\beta 2$ nAChRs may eventually provide additional avenues for nicotine addiction therapies. A hypothetical small molecule designed to bind to the KKK and disrupt COPI binding would prevent the Golgi–ER cycling in a manner similar to observations in this work. This inhibition of Golgi–ER cycling would prevent the up-regulation of $\beta 3$ -containing nAChRs and may produce effects similar to many antagonists that have shown clinical promise (Watkins et al., 1999; Rauhut et al., 2003; Yoshimura et al., 2007). In the search for potential therapeutics, the selective expression pattern of $\alpha 6\beta 2\beta 3$ nAChRs to just a few neuronal populations would render the targeting of $\beta 3$ more selective than a drug that inhibits Golgi–ER cycling in general.

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