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ARTICLE

Axon targeting of the alpha 7 nicotinic receptor in developing hippocampal neurons by Gprn1 regulates growth

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Abstract

Cholinergic signaling plays an important role in regulating the growth and regeneration of axons in the nervous system. The $\alpha 7$ nicotinic receptor ($\alpha 7$) can drive synaptic development and plasticity in the hippocampus. Here, we show that activation of $\alpha 7$ significantly reduces axon growth in hippocampal neurons by coupling to G protein-regulated inducer of neurite outgrowth 1 (Gprn1), which targets it to the growth cone. Knockdown of Gprn1 expression using RNAi is found sufficient to abolish the localization and calcium signaling of $\alpha 7$ at the growth cone.

In addition, an $\alpha 7$ /Gprn1 interaction appears intimately linked to a G α , growth-associated protein 43, and CDC42 cytoskeletal regulatory pathway within the developing axon. These findings demonstrate that $\alpha 7$ regulates axon growth in hippocampal neurons, thereby likely contributing to synaptic formation in the developing brain.

Keywords: axon growth, choline, cytoskeleton, G protein-regulated inducer of neurite outgrowth, hippocampus, synaptic maturation.

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In the adult nervous system, neurotransmitters play a central role in driving communication between neurons. In the developing nervous system, neurotransmitters such as serotonin and acetylcholine (ACh) contribute to various phases of development including neurogenesis, cell differentiation, and the formation of synapses (Erskine and McCaig 1995; Rudiger and Bolz 2008). The cerebral cortex and hippocampus receive ACh innervation from the basal forebrain, medial septum, and vertical diagonal band (Bruehl-Jungerman *et al.* 2011). Beginning prenatally and continuing through early adulthood, the cholinergic signal informs neural connectivity by modulating the growth and retraction of axons (Elsas *et al.* 1995; Luo and O'Leary 2005) and the formation of local dendritic fields within hippocampal neurons (Campbell *et al.* 2010; Lozada *et al.* 2012).

Cholinergic signaling is dependent on the activation of the G protein-coupled muscarinic receptor and nAChRs (Jones *et al.* 2012). In the adult hippocampus, $\alpha 7$ is expressed pre- and post-synaptically, contributing to GABA and glutamate neurotransmission (Liu *et al.* 2006; Lozada *et al.* 2012). Although studies show that $\alpha 7$ regulates neural development in the cortex and hippocampus (Coronas *et al.* 2000; Liu

et al. 2006; Lozada *et al.* 2012), its mechanism of action is ill defined.

Several nAChRs have been found to interact with G proteins and contribute to intracellular signaling in neurons (Fischer *et al.* 2005; Kabbani 2007; Paulo *et al.* 2009; Nordman and Kabbani 2012). Recently, we have shown that coupling to a G protein complex consisting of G protein-regulated inducer of neurite outgrowth 1 (Gprn1), the heterotrimeric GTP-binding subunit G α , and growth-associated protein 43 (GAP-43) enables $\alpha 7$ to regulate neurite growth (Nordman and Kabbani 2012). To determine the role of $\alpha 7$ in brain development, we have examined the effect of $\alpha 7$ /Gprn1 interaction in cultured hippocampal neurons. We

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Abbreviations used: DMSO, dimethylsulfoxide; EB3, end binding protein 3; GAP-43, growth associated protein 43; GC, growth cone; IP, immunoprecipitation; Ptx, pertussis toxin; SA, surface area.

find that $\alpha 7$ activation contributes to calcium signaling and inhibition of axon growth via a Gprn1 pathway.

Materials and methods

Neuronal cultures, transfection, and drug treatment

Primary hippocampal neurons were obtained from post-natal day 1 (P1) male and female Sprague–Dawley rats (Charles River, Frederick, MD, USA) as described (Nunez 2008) in accordance with the Institutional Animal Care and Use Committee (IACUC) and ARRIVE guidelines. Neuronal cultures were grown in Neurobasal media with B27 supplement and 1% Pen-strep. Serum was withdrawn 12 h after plating to minimize glial growth. A knockdown in $\alpha 7$ expression ($\alpha 7^-$) was obtained by transducing high-titer lentiviral stocks of short hairpin RNAs representing $\alpha 7$ RNAi (Lozada *et al.* 2012) into hippocampal neurons. A scrambled $\alpha 7$ RNAi was used as a transduction control (Lozada *et al.* 2012). Knockdown in Gprn1 expression (Gprn1⁻) was obtained by transfecting Gprn1 siRNA in pRNAT H1.1 into hippocampal neurons. Gprn1 vectors have been described previously (Ge *et al.* 2009; Nordman and Kabbani 2012). Neurons were transfected using Lipofectamine 2000 (Invitrogen, Carlsbad, CA, USA). Unless otherwise stated, an empty plasmid corresponding to the transfected protein vector was used as a control in the assay.

Drug concentrations were determined based on published studies (Chan and Quik 1993; Strittmatter *et al.* 1994; Swarzenski *et al.* 1996; Khiroug *et al.* 2002; El Kouhen *et al.* 2009): α -Bungarotoxin (Bgtx) (50 nM, Sigma, St Louis, MO, USA); PNU282987 (PNU) (1–10 μ M, Sigma); pertussis toxin (Ptx) (1 μ M, Calbiochem, San Diego, CA, USA); mastoparan (30 μ M, Tocris, Bristol, UK); and choline (1 mM, Acros, Geel, Belgium). Drug treatment experiments were performed in triplicate and the data presented represent the average for each condition. Prior to initiation of the assay, the effect of drug concentration on cell viability and health was pre-established using trypan blue (EMD).

Immunochemical detection of proteins in brain slices and cultured cells

Hippocampal brain slice preparation and immunohistochemistry were performed as described in Lozada *et al.* (2012). Briefly, P5 and P30 rats were anesthetized using 5% isoflurane and then perfused using 4% paraformaldehyde, pH 7.2. Brains were dissected and submerged in 4% paraformaldehyde for 24 h before being transferred to 30% sucrose for cryoprotection. The tissue was then embedded in 5% agarose and sectioned into 30- μ m slices using a vibrating blade microtome (Thermo Scientific, Waltham, MA, USA). For immunohistochemistry, brain slices were permeabilized using 0.5% Triton X-100 and quenched by 50-mM ammonium chloride for 30 min at 25°C. Tissue was blocked in 10% goat serum prior to being probed with the $\alpha 7$ ligand fluorescein-conjugated Bgtx (fBgtx) (Molecular Probes, Eugene, OR, USA), a polyclonal rabbit anti-Gprn1 antibody (Ab) (Abcam, Cambridge, MA, USA), and a polyclonal mouse anti-Tau-1 Ab (Millipore, Billerica, MA, USA) overnight at 4°C. Tissue was reblocked in 10% goat serum followed by immunoprobings with a Dylight 560 secondary anti-rabbit Ab and an Alexafluor 647 secondary anti-mouse Ab for Gprn1 and Tau-1 detection, respectively (Jackson-ImmunoResearch, West Grove, PA, USA).

Background autofluorescence was accounted for using the secondary Ab alone.

Cellular immunostaining was carried out as described in He *et al.* (2005). Briefly, cultured neurons were fixed in 1 \times PEM (80 mM PIPES, 5 mM EGTA, and 1 mM MgCl₂, pH 6.8) containing 0.3% glutaraldehyde, and then permeabilized at 25°C in 0.05% Triton X-100 prior to glutaraldehyde quenching with 10 mg/mL sodium borohydride. Cells were blocked in 10 mg/mL bovine serum albumin + 10% goat serum prior to being immunostained overnight at 4°C with the following primary Abs: mouse anti-Tau-1; anti-Gprn1; pGAP-43; GTP-CDC42 (New East, King of Prussia, PA, USA). Secondary Abs used (all purchased from Jackson-ImmunoResearch): carbocyanine (Cy) 2/3; Dylight 488; Dylight 560; and AlexaFluor 647. $\alpha 7$ was visualized using fBgtx. F-actin was visualized using Rhodamine Phalloidin (Cytoskeleton). All staining was visualized using a Nikon Eclipse 80i confocal microscope (Melville, NY, USA) fitted with a Nikon C1 CCD camera and images were captured using AxioVision and EZ-C1 software (Thornwood, NY, USA).

For protein distribution analysis in immunostainings, brain slices and cells in culture were analyzed for co-localization and distribution of $\alpha 7$ and Gprn1 by first thresholding the individual signals to reduce background (Lozada *et al.* 2012) and then merging the two signals using ImageJ (NIH, Bethesda, MD, USA). For brain slices, the colocalized $\alpha 7$ /Gprn1 signal was measured in various compartments according to overlap with DIC. Distribution analysis was conducted by measuring the signal within a 1 μ m² area for neurons or a 25 μ m² area for brain slices sufficient for analysis at the single neuron level. Analysis was conducted in Tau-1 positive (Tau-1+) cultured hippocampal neurons. All data are based on averages from three separate independent experiments.

Protein isolation and detection

Solubilized membrane protein fractions (membrane proteins) of hippocampal cells were obtained using a published protocol optimized for nAChRs (Kabbani *et al.* 2007). Immunoprecipitation (IP) of receptor–protein complexes for $\alpha 7$ was optimized and described previously (Nordman and Kabbani 2012). In brief, proteins were solubilized overnight at 4°C using a solution of 1% Triton X-100, 137 mM NaCl, 2 mM EDTA, and 20 mM Tris-HCl (pH 8) with a 1 \times protease inhibitor cocktail (Complete, Roche, Indianapolis, IN, USA) and a 1 \times phosphatase inhibitor cocktail (Sigma). Protein concentrations were obtained using the Coomassie Protein Assay reagent (Thermo Scientific). IP experiments were performed by first incubating membrane proteins with an anti- $\alpha 7$ monoclonal Ab (mAb306; Lindstrom *et al.* 1990) or an anti-Gprn1 rabbit polyclonal Ab (Abcam) followed by extraction using protein G Dynabeads (Invitrogen). Control IP experiments were performed by incubating the same amount of membrane proteins (MP, 100 μ g) with an equal amount of a pure polyclonal rabbit IgG Ab (5 μ g) (Cell Signaling Technology, Beverly, MA, USA).

Western blot detection was obtained using nitrocellulose membranes blocked with 5% non-fat milk (or 1% bovine serum albumin for biotinylation experiments). Membranes were probed with the following primary Abs overnight at 4°C: $\alpha 7$ nAChR (SC); Gprn1; HCN1 (SC); GAP-43 (Abcam); pGAP-43; CDC42 (SC); GTP-CDC42; glyceraldehyde 3-phosphate dehydrogenase (GAPDH) (Cell Signaling). Species-specific peroxidase-conjugated secondary Abs were purchased from Jackson-ImmunoResearch. Signals were

detected using the SuperSignal West Pico Chemiluminescent Substrate (Thermo Scientific) with SeeBlue and MagicMark (Invitrogen) as molecular weight standards. Blots were imaged using the Gel Doc Imaging system (Bio-Rad Laboratories, Hercules, CA, USA). Band density analysis was performed using ImageJ version 10.2 (NIH). Western blot values are based on averages from three separate independent experiments.

Growth cone purification

The growth cone (GC) fraction was prepared as described in Lohse *et al.* (1996). In brief, hippocampi pooled from a litter of P0 rat pups were homogenized in 1-mM TES buffer (50 mM Tris-HCl, pH 8.0, 5 mM EDTA, 51 mM NaCl) supplemented with 1-mM MgCl₂, 0.32-M sucrose, and protease (Complete) and phosphatase (Sigma) inhibitors. The homogenate was spun at low-speed centrifugation (1660 g) and the supernatant was collected then layered over a sucrose gradient of 0.83 M and 1.0 M. Layered sample was then centrifuged at 242 000 g for 40–60 min at 4°C. The GC fraction was found at the 0.83-M interface, whereas the cell fraction was found at the 0.83–1.0 M fraction. GC proteins were solubilized for membrane protein enrichment using the method described earlier in this study.

Biotinylation of cell surface receptors

The quantification of proteins at the cell surface was conducted using a biotinylation protocol (Kabbani *et al.* 2002; Hannan *et al.* 2008). In these experiments, 300 µg/mL EZ-Link Sulfo-NHS-LC-Biotin (Pierce, Rockford, IL, USA) was used to label cell surface proteins in living cells for 30 min at 4°C. The biotin reaction was quenched using tris-buffered saline, and membrane proteins were prepared as described previously (Kabbani *et al.* 2007). A neutravidin agarose bead matrix (Pierce) was used to pull down biotinylated proteins over a 30-µm pore size Snap Cap spin column (Pierce). Proteins were detected using western blot as described earlier. Biotin labeling was detected using horseradish peroxidase-conjugated streptavidin (Cell Signaling). Experiments were performed in triplicate and the data presented are the group averages.

Morphological and statistical analysis

Morphological reconstruction was performed using Neuromantic software (Myatt *et al.* 2012). Tau-1+ axons were measured for surface area (SA) and branch number to assess size and complexity under various conditions. Only the primary axon was analyzed in each neuron. Neurons were readily visualized by Tau-1+ staining, and GCs were visualized by reactivity to rhodamine phalloidin. Group averages were derived from three separate experiments with 20–30 cells in each set. Statistical analysis was conducted on raw data of image tracing using Neuromantic. Statistical values have been obtained using a Student's *t*-test or one-way ANOVA. Asterisks indicate statistical significance in a paired Student's *t*-test, two-tailed *p* value, * < 0.05; ** < 0.01; *** < 0.001. Error bars indicate standard error of the mean (SEM).

EB3 comet imaging

End-binding protein 3 (EB3) is a microtubule capping protein that binds to the plus end of polymerized microtubules during periods of assembly (Stepanova *et al.* 2003). Hippocampal neuronal cultures were transfected with red fluorescent protein (RFP) conjugated EB3 and the indicated cDNA plasmids using Lipofectamine 2000 at 60 h

after plating (Liu *et al.* 2010). EB3 comet imaging was performed 24 h later. Images were captured 1 frame per sec for 2 min at 2 × 2 binning using AxioVision. The images were quantified for the velocity of the comets that enter the growth cone filopodia (Nadar *et al.* 2008). Cells were treated with PNU (1–10 µM), a combination of PNU 10 µM and 50 nM Bgtx, 0.1% dimethylsulfoxide (DMSO) as vehicle control, or an empty vector as a transfection control. The comet trajectories were traced in these experiments using ImageJ software (NIH). Experiments were performed in triplicate and the data presented are the group averages.

Calcium imaging

Calcium imaging was carried out as described previously (Del Negro *et al.* 2011) with minor modifications. P0 hippocampal neurons were cultured for 3 DIV before loading with the calcium dye Fluo-8 AM (SC). Cultures were washed with Hank's Balanced Salt Solution (HBSS) and then incubated for 20 min at 25°C in a dye-loading solution containing HBSS with 10-mM HEPES, 2.5-µM Fluo-8 AM, and 0.2% Pluronic acid F-127. Following incubation, the cultures were again washed in fresh HBSS and then immediately placed into a recording chamber perfused with artificial cerebrospinal fluid at a rate of 1 mL/min. We imaged Fluo-8 AM-labeled neurons using a fixed-stage Zeiss Axio Examiner A.1 with a 63×/1.0 NA objective and a Rolera EMC² camera (Q-Imaging, Surrey, BC, Canada) imaged at 10 Hz. Phototoxicity was minimized through the use of low-wavelength polarized light filters. Image stacks were acquired at 70 Hz using a CCD camera (Andor Technology, South Windsor, CT, USA). The following drugs and their concentrations were diluted in artificial cerebrospinal fluid and administered via a perfusion system at 1 mL/min: PNU (10 µM); PNU and Bgtx (50 nM); choline (1 mM); choline and Bgtx. 0.1% DMSO was used as a vehicle control. An empty RFP vector was used as a transfection control. Regions of interest were analyzed using ImageJ (NIH) and normalized as $\Delta F/F_0$. Regions of interest were averaged over conditions. The data presented have been normalized to controls.

Mass spectrometry

Protein analysis was conducted using liquid chromatography electro spray ionization (LC-ESI) mass spectrometry (MS). Protein complexes were prepared as described (Kaiser *et al.* 2008) and mass spectrometry was carried out using a published protocol (de Luca *et al.* 2009). Tandem mass spectra collected by Xcalibur (version 2.0.2, Thermo Scientific) were searched against the NCBI rat protein database using SEQUEST (Bioworks software from Thermo Scientific, version 3.3.1). The SEQUEST search results were filtered using the following criteria: minimum X correlation (XC) of 1.9, 2.2, and 3.5 for 1+, 2+, and 3+ ions, respectively, and $\Delta C_n > 0.1$. The protein score represents the XC where scores < 0.1 were excluded from the analysis and does not reflect the quantity of a protein in the sample.

Results

$\alpha 7$ and Gprin1 interact in the developing hippocampus

$\alpha 7$ is expressed in the adult hippocampus where it can modulate the activity of excitatory and inhibitory neurons

(Albuquerque *et al.* 1996, 1998). In hippocampal development, however, less is known about the expression and role of $\alpha 7$. We examined the distribution of $\alpha 7$ in the hippocampus at various stages of growth. Brains from post-natal day (P) 5 and 30 rats were probed for $\alpha 7$ expression using fluorescent-conjugated α -bungarotoxin (fBgtx). In addition, we co-immunolabeled the brain slices with an anti-Gprn1 Ab, which selectively recognizes Gprn1 (Nordman and Kabbani 2012). As shown in Fig. 1a, co-expression of fBgtx and Gprn1 was detected in the developing hippocampus. This expression, although widespread, was not ubiquitous. In particular, the Gprn1 signal was found in cells that were immunoreactive for the axon-specific marker Tau-1, whereas fBgtx labeling was detected in Tau-1+ cells (Fig. 1b) as well as those that exhibited immunoreactivity to the astrocytic marker glial fibrillary acidic protein (data not shown). These findings imply that whereas $\alpha 7$ expression is not limited to neurons, Gprn1 expression is predominantly neuronal.

We compared the expression of the two proteins during development. A fluorescence double signal (co-signal) for fBgtx and anti-Gprn1 within the cell was measured as previously described (Nordman and Kabbani 2012). In preliminary experiments, the fBgtx signal was diminished

by the addition of PNU (1 and 10 μ M), confirming the specificity of the ligands on $\alpha 7$ (Figure S1). As shown in Fig. 1a, the co-signal was more abundant during the early P5 stage of development. A quantitative assessment of the fluorescence co-signal was used to determine $\alpha 7$ /Gprn1 co-expression in the hippocampus. As shown in Fig. 1a, a noticeable $\alpha 7$ /Gprn1 co-signal was detected in CA3 and CA1 regions at P5. Immunolabeling with an anti-Tau-1 Ab confirmed that over 95% of the cells that displayed an $\alpha 7$ /Gprn1 co-signal are neurons (Fig. 1b). Hippocampal neurons are known to undergo apoptosis, maturation, and synaptic pruning during early post-natal development (Jordan *et al.* 1997; Riccomagno *et al.* 2012). $\alpha 7$ /Gprn1 co-expression was also detected in other regions of the hippocampus as well as the cortex and striatum (Fig. 1a). Single-labeling experiments using fBgtx, anti-Tau-1, or anti-Gprn1 antibodies produced a signal consistent with the double- and triple-labeling results (data not shown).

Immunofluorescent findings indicate that $\alpha 7$ /Gprn1 expression is highest after birth. To explore this, we examined the levels of the two proteins in hippocampal tissue by western blot. As indicated in Fig. 1c, the expression of $\alpha 7$, within membrane fractions of the developing

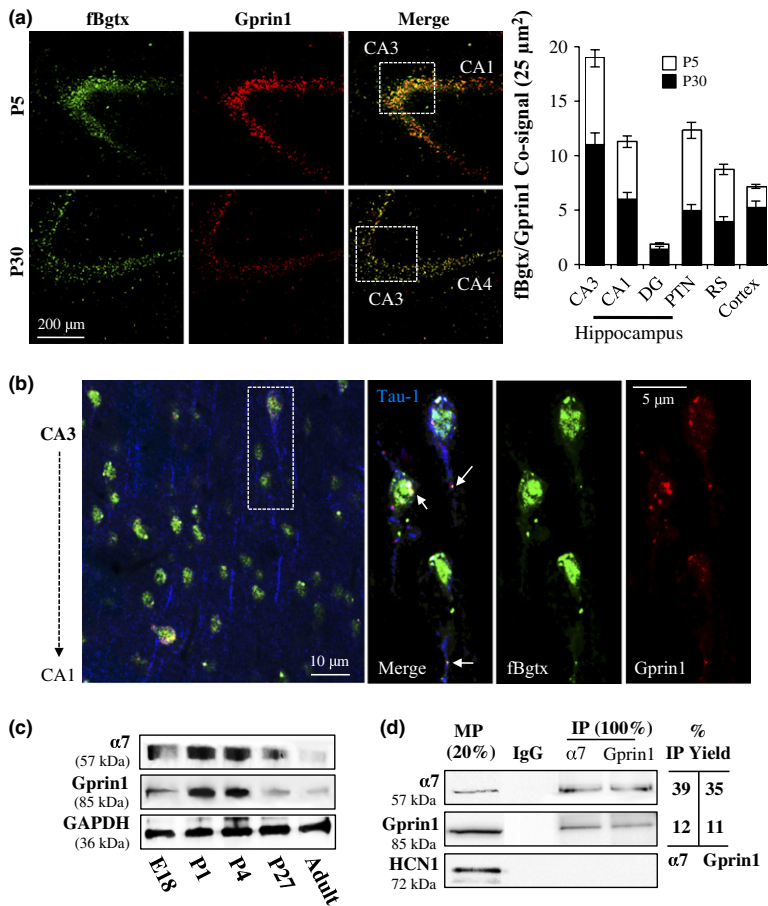


Fig. 1 $\alpha 7$ and G protein-regulated inducer of neurite outgrowth 1 (Gprn1) interact in the developing hippocampus. (a) Coronal slices of brains obtained from P5 and P30 rats. Slices were co-labeled with fBgtx and anti-Gprn1 Abs and visualized throughout the hippocampus, with prominent expression found in the CA3 region. Quantification of the co-signal in various brain regions is shown in the histogram. (b) Triple labeling for Tau-1 (blue), fBgtx (green), and Gprn1 (red) in CA3 at P5. Arrows point to colocalization of fBgtx and Gprn1 in soma and axons. (c) Western blot detection of $\alpha 7$ and Gprn1 in membrane protein (MP) fractions of the hippocampus. The same blot was used to probe for $\alpha 7$, Gprn1, and GAPDH as a loading control. (d) Western blot detection of $\alpha 7$ and Gprn1 interactions from P0 pups within immunoprecipitation (IP) experiments. In-gel digest confirms the identity of $\alpha 7$ and Gprn1 in the IP (Table S1). Top blot: $\alpha 7$; middle blot: Gprn1; bottom blot: HCN1. MP 20%: membrane protein as 20% of IP load (100 μ g). IgG: IP control. IP yield: amount of IP protein obtained as determined by the equation: optical density (O.D.) of IP bands/OD of 100% MP \times 100.

hippocampus, appeared to peak at 1 and 4 days after birth. Similarly, the expression of Gprn1 displayed a consistent expression profile, i.e., peaking after birth then sharply declining to barely detectable levels in adulthood (Fig. 1c). These results suggest a role for the two proteins in hippocampal development.

Immunoprecipitation was used to validate interactions between $\alpha 7$ and Gprn1 during hippocampal development. MP derived from P5 rat pups were used to immunoprecipitate Gprn1 and $\alpha 7$ proteins. The $\alpha 7$ -specific Ab mAb306 and an anti-Gprn1 Ab were then used to immunoprobe for Gprn1 and $\alpha 7$ in the IP, respectively. As shown in Fig. 1d, immunoreactive bands for $\alpha 7$ and Gprn1 proteins were detected in the IP experiments. The identity of the proteins within the IP assay was also confirmed in parallel experiments utilizing MS analysis (Table S1) and previously (Nordman and Kabbani 2012). An hyperpolarization-activated cyclic nucleotide-gated channel type 1 (HCN1)-specific Ab was used as a negative control in the IP experiment. As shown, neither of the two IP Abs was found to co-IP HCN1 in the assay (Fig. 1d), thus, supporting the specificity of the interaction between $\alpha 7$ and Gprn1 at the plasma membrane.

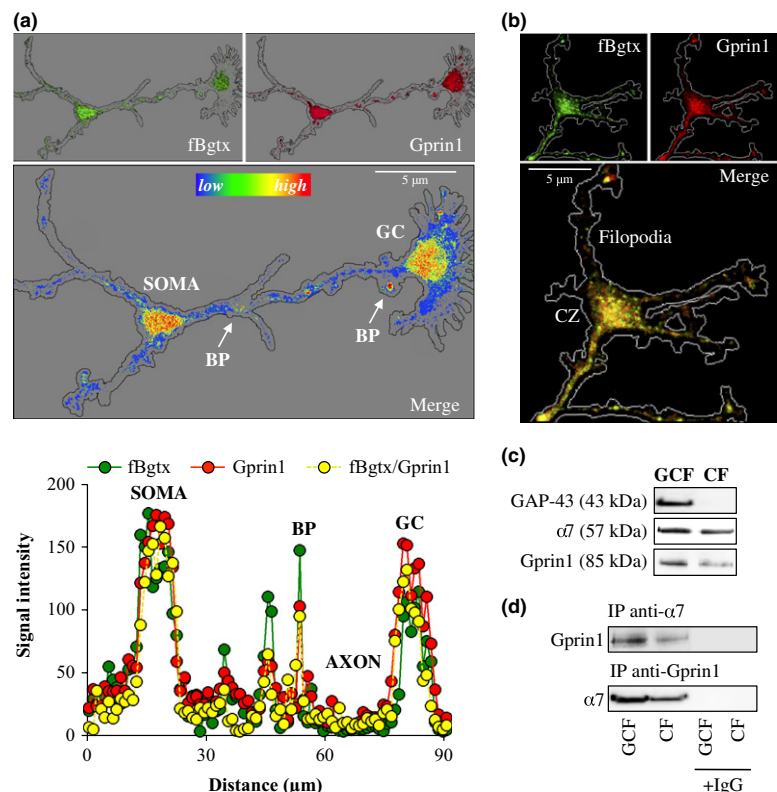
Localization of $\alpha 7$ and Gprn1 in axons and growth cones

We examined expression of endogenous $\alpha 7$ and Gprn1 proteins in developing hippocampal neurons. At 3 days

in vitro (DIV) hippocampal neurons extend axons capable of detecting and reacting to various external growth signals (Kater and Mills 1991). As shown in Fig. 2, $\alpha 7$ and Gprn1 proteins were detected within developing hippocampal neurons using fBgtx and anti-Gprn1 Abs, respectively. The specificity of the Gprn1 Ab was verified in preliminary studies that showed a proportional decrease in the Gprn1 Ab signal following transfection with Gprn1 siRNA (data not shown). In particular, strong levels of $\alpha 7$ and Gprn1 co-expression were observed in the soma, at neurite branch points, and in the GC (Fig. 2a). Growth cones and branch points are known to be active regions of growth and retraction (Lowery and Van Vactor 2009). We find that $\alpha 7$ /Gprn1 co-localization is highest at these regions of active growth suggestive of a role for the interaction in cytoskeletal remodeling and axon development.

The axon growth cone consists of three main zones: peripheral, transitional, and central. The central zone (CZ) is located at the base of the GC, nearest to the axon, and is composed of microtubules as well as various receptors and organelles (Lowery and Van Vactor 2009). We find strong co-localization of $\alpha 7$ and Gprn1 within the CZ and, in some cases, the two proteins were detected in the filopodia of the GC (Fig. 2b). A GC purification strategy (described in *Materials and Methods*) (Lohse *et al.* 1996) was utilized to confirm expression and interaction of $\alpha 7$ and Gprn1 in the

Fig. 2 $\alpha 7$ and G protein-regulated inducer of neurite outgrowth 1 (Gprn1) associate in the growth cone. (a) Hippocampal neurons from P0 pups were cultured for 3 DIV. Neurons were probed with fBgtx (green) and anti-Gprn1 Abs (red) (top images). A heat map measure of the co-signal (bottom image) shows the distribution and co-localization of the two proteins in the growing axon and GC. Co-localization was highest in the soma, growth cone (GC), and branch points (BP) (arrows). (b) Localization of the fBgtx and Gprn1 signals in the GC. CZ: central zone. (c) Protein detection of $\alpha 7$ and Gprn1 within GC fraction (GCF) obtained from P0 pups as described in Materials and methods. Growth-associated protein 43 (GAP-43) is used as a marker for the GCF. Cell fraction (CF). (d) An anti- $\alpha 7$ and anti-Gprn1 Ab was used to immunoprecipitate (IP) $\alpha 7$ and Gprn1 proteins from the GCF. Western blot detection confirms interaction of the two proteins. Protein identity was also determined by in-gel digest mass spectrometry (Table S2). IgG was used as an Ab control.



GC of the developing hippocampus. As shown in Fig. 2c, an anti-GAP-43 Ab was used to confirm the isolation of the GC compartment from hippocampal tissue. A western blot analysis reveals that $\alpha 7$ as well as Gpr1 are present within the GC consistent with our immunocytochemical finding. The interaction between the two proteins within the IP was also confirmed in parallel using MS analysis (Table S2). The above experiments demonstrate an $\alpha 7$ /Gpr1 interaction within the GC of developing hippocampal neurons.

$\alpha 7$ regulates the structure of the growth cone and axon

$\alpha 7$ has been shown to regulate neurogenesis and dendritic as well as axonal synaptic formation in the nervous system (Bromberg *et al.* 2008b; Lozada *et al.* 2012). To examine the role of this receptor in early axon growth of hippocampal neurons, we examined the effect of the $\alpha 7$ -specific agonist PNU282987 (PNU) on axon size and structure in hippocampal neurons. PNU is a known agonist at the $\alpha 7$ receptor site found to bind other nAChR ligands including Bgtx (El Kouhen *et al.* 2009). The stability of PNU and its resistance to neurotransmitter reuptake mechanisms make it

an ideal experimental tool when attempting to selectively target $\alpha 7$. Newly formed axons were visualized using an antibody for the axon-specific microtubule protein Tau-1. We assessed the effects of PNU (1–10 μ M) or a combination of PNU (10 μ M) and the $\alpha 7$ -specific antagonist α -Bungarotoxin (Bgtx) (50 nM) on axon growth. PNU treatment was associated with a significant, dose-dependent (10 μ M) reduction in axon growth [–42% (\pm 6%)], measured as total SA and branching number [47% (\pm 7%)] (Fig. 3a). In the presence of Bgtx, however, PNU did not significantly alter the growth and complexity of the primary axon relative to control-treated cells (Fig. 3a). Group significance was observed between controls and PNU doses (1–10 μ M) (SA: $F(2,30) = 4.85, p < 0.01$; branch: $F(2,30) = 6.12, p < 0.01$). These results reveal that pharmacological activation of $\alpha 7$ reduces axon growth and branching in hippocampal neurons.

GC dynamics are instrumental in guiding the developing axon toward its final destination and have been shown to be a useful measure of real-time growth (Zheng and Poo 2007). In particular, the innervation of microtubules into filopodia

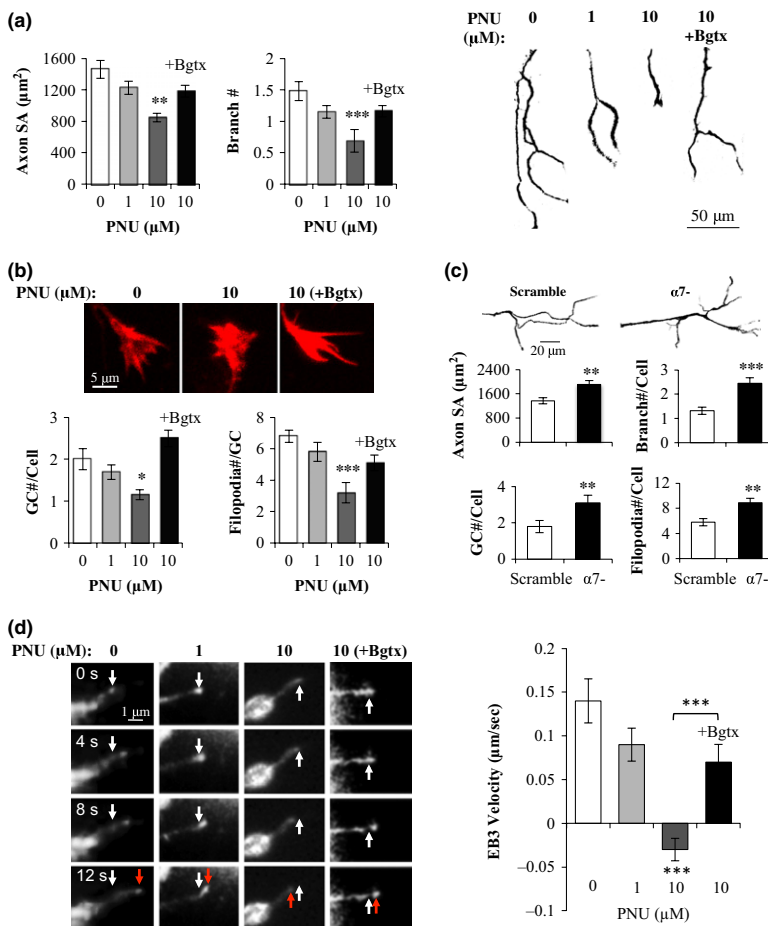


Fig. 3 Activation of $\alpha 7$ inhibits axon growth. Hippocampal neurons were treated with PNU (1 and 10 μM) or a 0.1% dimethylsulfoxide (DMSO) control (PNU = 0). (a) Analysis of surface area (SA) and branch number of Tau-1+ axons (left) represented as phase contrast images (right). (b) Images: growth cones (GCs) stained with rhodamine phalloidin. Histogram: an analysis of the GC. (c) Morphometric analysis of axons and GCs in $\alpha 7$ or scramble cells in the absence of drug treatment. (d) Images: RFP-end-binding protein 3 (EB3) comet movement (red arrow) from the starting position (white arrow) within the GC filopodia. Histogram: Changes in EB3 comet velocity with drug treatment. Bgtx (50 nM); Control (0.1% DMSO). * $p < 0.05$; ** $p < 0.01$; *** $p < 0.001$.

correlate with the directionality of axonal growth (Lowery and Van Vactor 2009). We find high levels of α 7 expression in the GC formation at the end of the primary axon branch (Fig. 2a). We hypothesize a role for α 7 in regulating axon growth via the GC. To test this, we analyzed the structure of the GC in hippocampal neurons treated with the α 7 agonist PNU (1–10 μ M). We find a dose-dependent reduction in the number of GCs for each neuron [–44% (\pm 10%)] and a reduction in filopodia number and size in each GC [–53% (\pm 11%)] following PNU treatment and compared with vehicle-treated controls (Fig. 3b). Group significance was observed between controls and PNU doses (1–10 μ M) (GC#/cell: $F(2,30) = 5.13$, $p < 0.01$; Filopodia#/GC: $F(2,30) = 6.95$, $p < 0.01$). In PNU (10 μ M)-treated cells, GC and axon morphology were highly correlated (Table S3), suggesting that α 7 activation inhibits growth uniformly throughout the cell. Similar to axon growth, Bgtx was found to abolish the effects of PNU (10 μ M) on GC structure and number in a highly correlative manner (Fig. 3b, Table S3), underscoring the specificity of the α 7 signal in these developmental effects.

To confirm the role of α 7 in axon growth, hippocampal neurons were transduced with α 7 RNAi or a scramble vector then analyzed for morphological changes in axon growth in the absence of drug treatment. As shown in Fig. 3c, cells expressing α 7 RNAi (α 7-) displayed significantly enhanced growth of axons [SA: +39.5% (\pm 9%); branch number: +85.6% (\pm 10%)] and growth cones [GC number/cell: +72.2% (\pm 16%); filopodia number/cell: 53.4% (\pm 11%)] compared with both the scramble controls and non-transfected cells (Fig. 3a–c). The scramble vector was found to have no effect on axon growth. Similar findings were observed in hippocampal neurons treated with 50-nM Bgtx showing that inhibition of α 7 promotes axon growth (data not shown). In light of earlier findings on PNU-mediated inhibition of axon growth, the current data supports the hypothesis that inactivation of α 7 can augment neurite growth (Nordman and Kabbani 2012).

We observed the effects of PNU treatment on cytoskeletal growth using the microtubule + end capping protein RFP-EB3 (EB3 comets), which has been utilized in analyzing real-time axon growth (Stepanova *et al.* 2003; Liu *et al.* 2010). As shown in Fig. 3d, at 1 μ M of PNU treatment, EB3 comets velocity was noticeably diminished relative to controls. At 10 μ M of PNU, however, the velocity of the EB3 comets was not only significantly attenuated but also moved in a retrograde direction suggestive of growth cone collapse. Group significance was observed between controls and PNU doses (1–10 μ M) ($F(2,30) = 5.80$, $p < 0.01$). The addition of Bgtx was found to significantly attenuate the effects of PNU treatment (10 μ M) on EB3 comet velocity, and restored the directionality of comet movement to anterograde (Fig. 3d).

Gprin1 is critical for α 7-mediated growth

Gprin1 can regulate neurite growth via its ability to bind and regulate large G proteins such as $G\alpha_o$, as well as small G proteins such as CDC42 (Chen *et al.* 1999; Nakata and Kozasa 2005). We determined a role for Gprin1 in α 7-mediated axon growth in hippocampal neurons by transfecting neurons with plasmids encoding short interfering RNA (siRNA) (Gprin1-), which has been shown to reduce Gprin1 expression in neural cells (Ge *et al.* 2009). Consistent with earlier findings, reduction in Gprin1 expression was associated with an overall decrease in neurite growth (Ge *et al.* 2009). When examining the effects of PNU on axon growth, we found that PNU has no effect on axon morphology in Gprin1- cells (Fig. 4a), suggesting that Gprin1 expression is necessary for α 7 function. We confirmed the involvement of $G\alpha_o$ in the α 7/Gprin1 pathway. As shown in Fig. 4b, treatment of cells with the $G\alpha_o$ activator mastoparan (MSP) was found to significantly enhance axon growth [axon SA: 31% (\pm 11%); branch #: 47% (\pm 4%); GC #/cell: 58% (\pm 5%); filopodia #/cell: 45% (\pm 4%)]. The effect of MSP was found sufficient to override PNU-mediated inhibition of axon growth suggesting that $G\alpha_o$ operates downstream of α 7. We confirmed this by also examining the role of the $G\alpha_i/o$ inhibitor PTX on axon development. Ptx exposure was associated with a significant reduction in axon growth [axon SA: –34% (\pm 10%); branch #: –41% (\pm 7%); GC #/cell: –40% (\pm 6%); filopodia #/cell: –47% (\pm 8%)] and this effect was not enhanced by PNU. Group significance was observed between controls, MSP, MSP+PNU, Ptx, and Ptx+PNU treated cells [SA: $F(4,50) = 5.37$, $p < 0.01$; Branch#: $F(4,50) = 12.36$, $p < 0.001$; GC#/Cell: $F(4,50) = 12.79$, $p < 0.001$; Filopodia#/Cell: $F(4,50) = 14.01$, $p < 0.001$], suggesting that α 7 operates in a $G\alpha_i/o$ pathway.

α 7 signaling at the GC regulates GAP-43 and CDC42

A number of signaling pathways have been shown to regulate axon growth or direct its retraction by directing the assembly of the cytoskeleton (Lowery and Van Vactor 2009). Studies demonstrate that activation of GAP-43 via its phosphorylation at Ser41 is critical for axon growth and branching (Kozma *et al.* 1997; Dent and Meiri 1998; Leu *et al.* 2010). Similarly, the GTP activation of CDC42 (GTP-CDC42) functions as an important indicator of GC function (Kozma *et al.* 1997). Recently, we demonstrated that α 7 activation regulates GAP-43 phosphorylation in neurites (Nordman and Kabbani 2012), while other studies have shown a role for Gprin1/ $G\alpha_o$ in the activation of CDC42 in the growth cone (Nakata and Kozasa 2005). To test the role of α 7 signaling in GAP-43 and CDC42 activation, we examined the effect of PNU (60-min treatment) on the phosphorylation of GAP-43 and the levels of GTP-CDC42 in hippocampal neurons. We find a significant reduction in phospho-GAP-43 (pGAP-43) (Zakharov and Mosevitsky 2007) and active (GTP bound) CDC42 (Elbediwy *et al.*

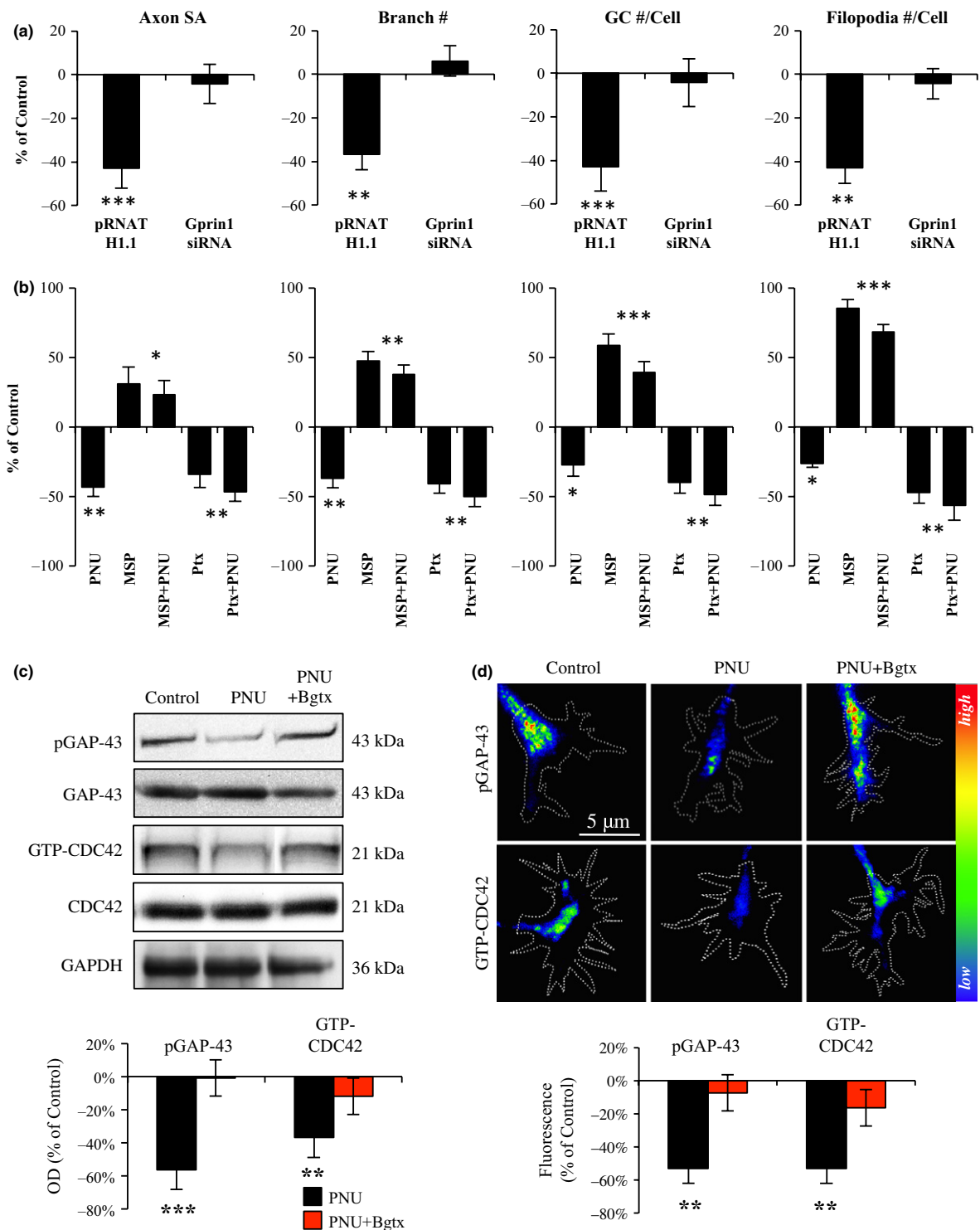


Fig. 4 $\alpha 7$ -mediated inhibition of axon growth is dependent on a G protein-regulated inducer of neurite outgrowth 1 (Gpr1) pathway. (a) An analysis of axon and growth cone (GC) morphology in cells treated with 10- μ M PNU. Values are based on average percent change in axon growth from control cells [cells transfected with the same plasmid, but treated with 0.1% dimethylsulfoxide (DMSO)]. (b) An analysis of axon and GC morphology in cells treated with: PNU (10 μ M); pertussis

toxin (Ptx) (1 μ M), mastoparan (30 μ M). Values are based on average change in axon growth from control cells which were treated with 0.1% DMSO alone. (c and d) Analysis of pGAP-43 and GTP-CDC42 expression within hippocampal neurons treated with PNU (10 μ M), PNU+Bgtx (50 nM), or control (0.1% DMSO) for 60 min in membrane protein (MP) (c) and GCs (d). Immunofluorescence signal in (d) represented as a heat map. * $p < 0.05$; ** $p < 0.01$; *** $p < 0.001$.

2012) levels in PNU-treated cells (Fig. 4c). Decrease in pGAP-43 and GTP-CDC42 expression was especially pronounced in the GC (Fig. 4d). The effect of PNU on GAP-43 phosphorylation and CDC42 activation was abolished in the presence of Bgtx, suggesting that $\alpha 7$ can inhibit the function of these two growth regulatory proteins.

Gprn1 localizes $\alpha 7$ to the growth cone

Gprn1 is a membrane anchored signaling protein that localizes to regions where cytoskeletal remodeling demands are high. In neural cells, Gprn1 can direct the formation of neurites by regulating the function of CDC42 in the growth cone (Nakata and Kozasa 2005). Because recent studies suggest that Gprn1 can also contribute to receptor localization at the cell surface (Ge *et al.* 2009; Nordman and Kabbani 2012), we examined the role of Gprn1 in trafficking and targeting $\alpha 7$ to the plasma membrane. As shown in Fig. 5a and b, Gprn1- cells displayed a significant reduction in the axonal structure, which was accompanied by a noticeable reduction in fBgtx labeling in the GC. To determine the role of Gprn1 in directing $\alpha 7$ to the plasma membrane, cell surface biotinylation experiments were performed in Gprn1- cells and empty vector (pRNAT H1.1) transfected controls. As shown in Fig. 5c, on the basis of the detection of biotinylated $\alpha 7$ proteins, we find a significant reduction in $\alpha 7$ at the cell surface in Gprn1- relative to controls. In both the GC and the MP fraction, levels of biotinylated $\alpha 7$ proteins was reduced in Gprn1- cells suggesting that Gprn1 plays an essential role in localizing $\alpha 7$ to sites important for axon growth. In these experiments, HCN1 cell surface expression was found unaffected.

Detection of an $\alpha 7$ calcium signal in the growth cone

Multiple ion channels and receptors contribute to calcium entry and signaling within the GC (Zheng and Poo 2007). The temporal and spatial dynamics of calcium within the GC contribute to the elongation, retraction, and turning of the axon via the actions of molecules such as GAP-43 and Gprn1, which regulate the cytoskeleton (Nakata and Kozasa 2005; Zheng and Poo 2007). We hypothesize that $\alpha 7$ /Gprn1 interaction enables calcium signaling leading to cytoskeletal remodeling in the developing axon. To test this we analyzed $\alpha 7$ -mediated calcium changes in hippocampal neurons using the calcium indicator Fluo-8 (Hayes *et al.* 2012). 0.1% DMSO was used as a vehicle control. The control by itself did not appear to impact calcium levels in the cell (data not shown). As shown in Fig. 6, PNU treatment was found to promote an overall increase in calcium relative to the control within the soma [+37% ($\pm 11\%$)], the axon [+52% ($\pm 14\%$)], and the central zone of the GC [+68% ($\pm 17\%$)]. In particular, the intracellular calcium rise was strongest within the CZ of the GC, a region that expresses an abundant amount of $\alpha 7$ and Gprn1 (Fig. 2). In filopodia,

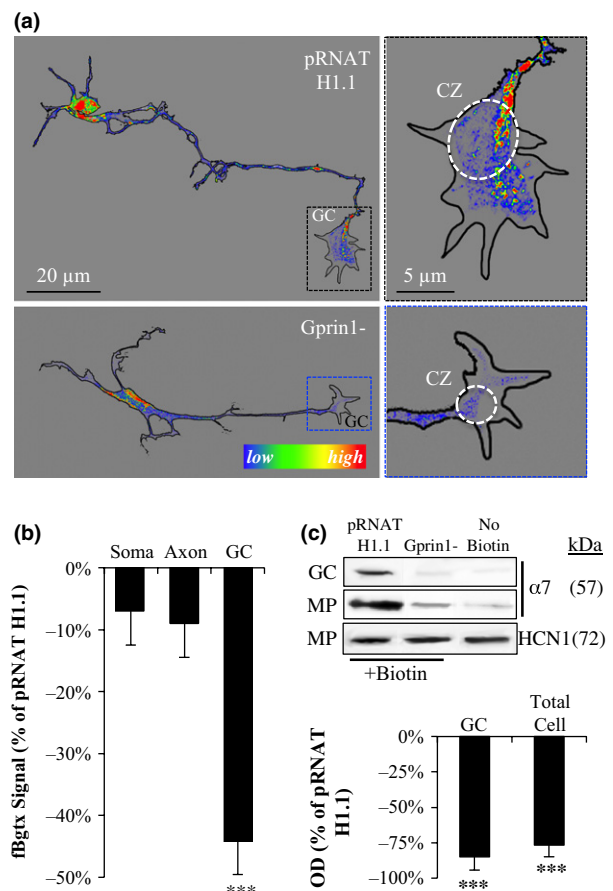


Fig. 5 G protein-regulated inducer of neurite outgrowth 1 (Gprn1) directs the localization of $\alpha 7$ to the cell surface of the growth cone. (a) pRNAT H1.1/Gprn1- cells were labeled with fBgtx for the visualization of $\alpha 7$ expression. Right panels: magnified images of the growth cone (GC) showing changes in $\alpha 7$ expression within the central zone and filopodia between the two conditions. The fBgtx signal is shown as heat map. (b) A quantification of the fBgtx signal in Gprn1- cells relative to empty pRNAT H1.1 controls. (c) Cell surface biotinylation was used to determine $\alpha 7$ expression at the cell surface in the GC and membrane protein (MP) fraction. HCN1 was used as a negative control. *** $p < 0.001$.

PNU treatment was also found to augment calcium levels by 24% ($\pm 7\%$). The effect of PNU on intracellular calcium levels was recovered to baseline at ~ 0.5 s, consistent with data on $\alpha 7$ channel kinetics (Fayuk and Yakel 2004) (Fig. 6).

Choline can endogenously activate $\alpha 7$, thereby critically contributing to brain development (Khiroug *et al.* 2002). We confirmed the effects of $\alpha 7$ activation in hippocampal neurons using (1 mM) choline, a concentration previously found to physiologically activate the receptor (Khiroug *et al.* 2002). We found that choline increased intracellular calcium levels but most significantly in the GC of the neuron (Fig. 7). The effect of choline was blocked by Bgtx, underscoring the specificity of this transmitter for $\alpha 7$. Interestingly choline was found to have little to no effect on calcium levels in the soma (Fig. 7a). The findings suggest that the effects of

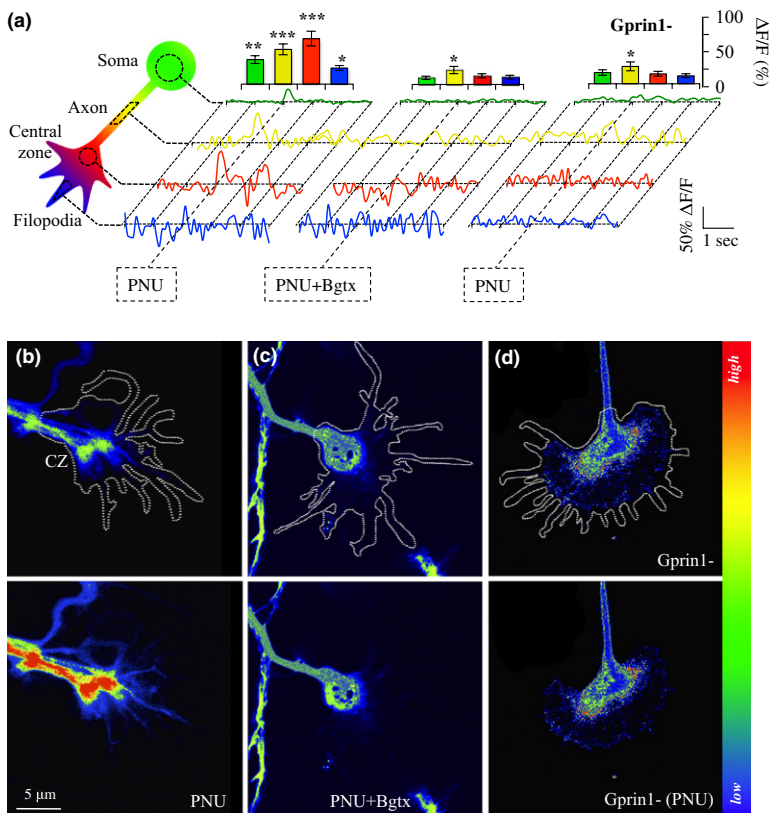


Fig. 6 Detection of an $\alpha 7$ calcium current in the growth cone. Cells were analyzed for intracellular calcium changes using Fluo-8AM. (a) Calcium detection in the soma, axon, central zone, and filopodia of hippocampal neurons. Normalized traces were obtained from calcium reading in 10 cells ($n = 10$) where 0.1% dimethylsulfoxide (DMSO) alone was used as a vehicle control. An empty pRNAT H1.1 vector was used as a transfection control for G protein-regulated inducer of neurite outgrowth 1 (Gprn1-) traces. Histograms showing relative changes in calcium peaks at time of drug treatment. (b–d) Drug-induced calcium changes in the growth cone (GC) (white trace). Top row: before drug application; bottom row: after drug application. Central zone (CZ); PNU (10 μ M); Bgtx (50 nM). * $p < 0.05$; ** $p < 0.01$; *** $p < 0.001$.

choline on $\alpha 7$ may be altered by differential mechanisms of transmitter reuptake in various neuronal compartments (Guermonez *et al.* 2002). To confirm that choline and PNU produce similar effects on axon growth, an analysis of axon morphology and pGAP-43 levels was also determined in response to choline. Axon development, GC branching, and pGAP-43 levels were all significantly attenuated by choline and this effect was also blocked by the addition of Bgtx (Fig. 7). Taken together, the findings indicate that $\alpha 7$ activation mediates intracellular calcium signaling leading to an inhibition in axon development.

To confirm the contribution of Gprn1 in $\alpha 7$ -mediated calcium entry into the growing hippocampal neurons, Gprn1- cells were analyzed for the effect of PNU (Fig. 6) and choline (Fig. 7). As shown in Fig. 6, knockdown of Gprn1 expression is associated with a significant reduction in the cell's calcium responsiveness to PNU treatment ($-19\% (\pm 9\%)$ in the soma; $-25\% (\pm 11\%)$ in the axon; and $-53\% (\pm 8\%)$ in the CZ) relative to an empty plasmid (pRNAT H1.1) control. Similar data was obtained with choline treatment (Fig. 7), confirming the role of Gprn1 in $\alpha 7$ -mediated calcium signaling in the developing axon. These findings also present a first ever demonstration of an $\alpha 7$ calcium signal in the GC, and suggest that this cholinergic signal contributes to the maturation of the axon.

Discussion

A new role for $\alpha 7$ in axon growth

In this study we define an interaction between $\alpha 7$ and Gprn1 and demonstrate that this interaction regulates axon growth in hippocampal neurons. We present evidence on the localization and signaling of $\alpha 7$ in GCs as supported by histochemical and western blot detection of the receptor and its interacting protein Gprn1 in developing hippocampal neurons. By interacting with Gprn1, the localization of $\alpha 7$ in the GC appears to enable cholinergic signals to regulate the growth of the axon. Based on this evidence, the interaction of the two proteins and their coupling to a downstream G protein-signaling pathway is proposed to contribute to brain development.

Interestingly, whereas this study demonstrates a role for $\alpha 7$ in axon development, earlier work has shown that knockdown of the $\alpha 7$ can decrease dendritic length and branching in newborn neurons of the dentate gyrus (Campbell *et al.* 2010; Liu *et al.* 2010). This suggests that $\alpha 7$ may differentially contribute to axon and dendrite development by interacting with scaffolding proteins such as Gprn1, which can guide the expression of the receptor to specific neuronal compartments. In this regard, the function of Gprn1 in localizing $\alpha 7$ activity in the developing

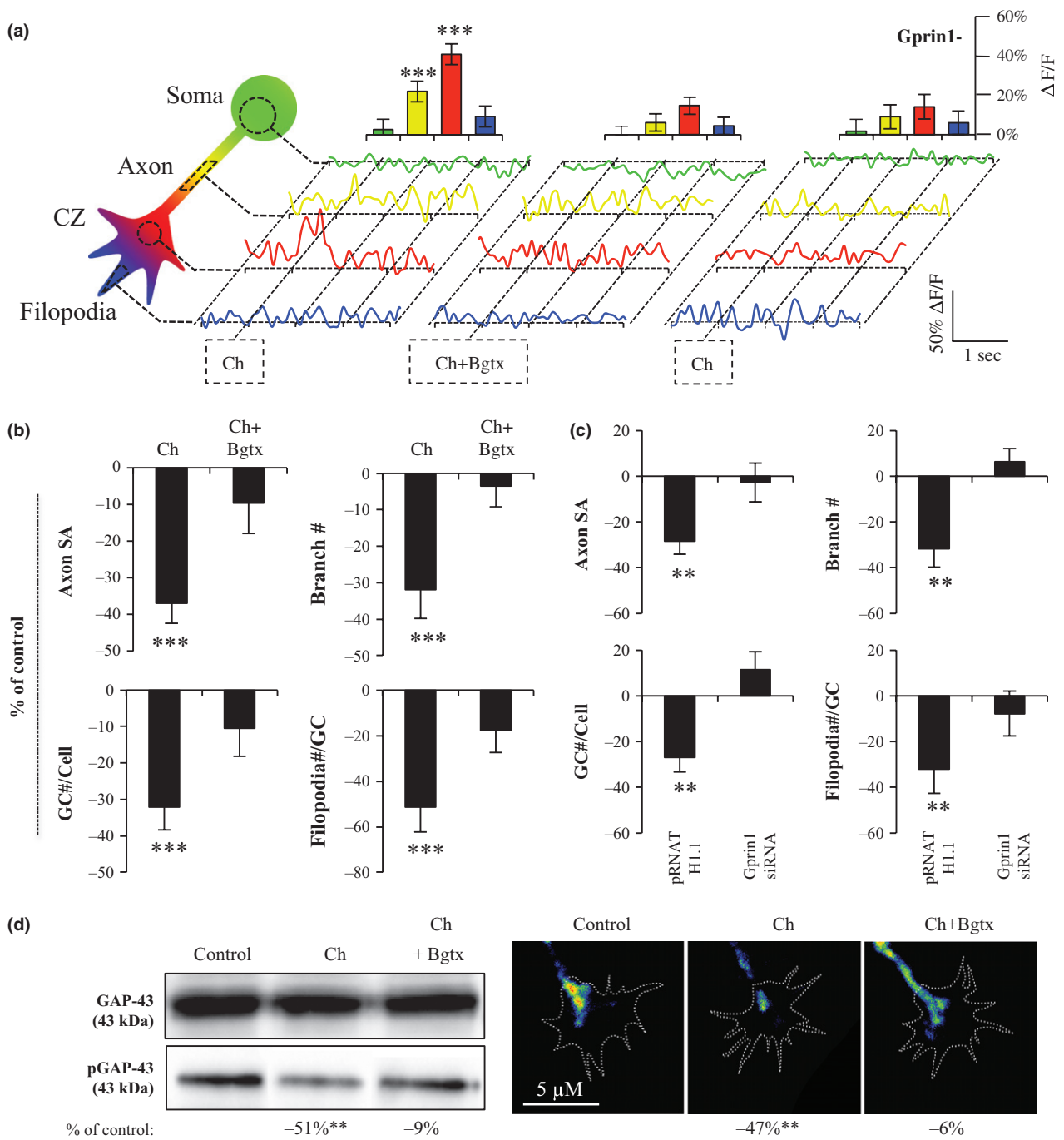


Fig. 7 Choline activation of $\alpha 7$ in the growth cone. Cells were analyzed for intracellular calcium changes using Fluo-8AM in response to Choline (Ch) treatment. Ch = 1 mM; Bgtx = 50 μ M. (a) Calcium detection in soma, axon, central zone, and filopodia of hippocampal neurons. Traces were averaged from seven cells ($n = 7$). Histograms show the relative change in calcium of peaks at the time of drug treatment. Water was used as a vehicle control. (b) Morphometric analysis of growth in choline-treated cells. The data are contiguous with analysis presented in Fig. 3 confirming the effect of $\alpha 7$ on axon growth. Water was used as a vehicle control. (c) Morphometric

analysis in G protein-regulated inducer of neurite outgrowth 1 (Gprin1) cells treated with choline. The data are contiguous with analysis presented in Fig. 4 supporting the role of Gprin1 in $\alpha 7$ -mediated inhibition of growth. Values are based on average per cent change in axon growth from control cells [cells transfected with the same plasmid, but treated with 0.1% dimethylsulfoxide (DMSO)]. (d) Detection of growth-associated protein 43 (GAP-43) and pGAP-43 in neurons following 60-min treatment with choline or choline and Bgtx. Changes in pGAP-43 expression were detected in the GC. Water was used as a vehicle control. ** $p < 0.01$; *** $p < 0.001$.

axon, may be paralleled by another, still unknown, protein that can direct the receptor to dendrites (Campbell *et al.* 2010). In future studies, it will be important to examine the proteomic interactions of $\alpha 7$ in maturing dendrites.

Activation of $\alpha 7$ causes local calcium elevation and a collapse of the GC

The synaptic localization and high calcium permeability of $\alpha 7$ enables it to influence diverse events, ranging from modulation of transmitter release to synaptic plasticity (McGehee *et al.* 1995; Gu and Yakel 2011). The expression and function of $\alpha 7$ in the GC during axon maturation appears dependent on interaction with Gprin1, which directs the trafficking of the receptor to the cell surface and the central zone. This positions $\alpha 7$ at sites where developmental demands are high, and enables it to influence cytoskeletal mechanisms of growth (Lowery and Van Vactor 2009). As a consequence, $\alpha 7$ -mediated elevations in intracellular calcium can arrest axon growth as shown previously for other calcium channels (Kater and Mills 1991; Henley and Poo 2004). The $\alpha 7$ -mediated calcium rise in the GC communicates a 'stop' signal for the growing axon, which is made possible by association with Gprin1 and modulation of the $G\alpha_o$, GAP-43, and CDC42 pathway (Frey *et al.* 2000; Nakata and Kozasa 2005). This hypothesis is supported by our observation that activation by $\alpha 7$ (by PNU or choline) promotes dephosphorylation of GAP-43, GTP hydrolysis of CDC42, and negatively influences EB3 microtubule comet velocity in the GC leading to its collapse. Mechanistically this may be achieved via $\alpha 7$ -mediated calcium elevations into the GC which can inactivate GAP-43 via calcium sensitive proteins such as calpain (Zakharov and Mosevitsky 2007) and PP2B (Lyons *et al.* 1994), leading to inhibition of $G\alpha$ and CDC42 in the axon (Bromberg *et al.* 2008a). It is interesting to consider the possible effects of $\alpha 7$ on other downstream Rho-GTPases involved in growth. RhoA has previously been shown to inhibit axon growth and GC collapse, while $G\alpha_o$ can regulate RhoA activity (Bromberg *et al.* 2008a). Thus, the ability of $\alpha 7$ to inhibit axon growth may involve the regulation of other possible Rho-GTPases through interaction with the G protein complex. In the long term, activation of $\alpha 7$ is associated with a dramatic reduction in axon length and branching complexity.

A functional $\alpha 7$ signal in the growing axon can play an important role in early-life synaptic development (Luo and O'Leary 2005). Defects in processes such as neurogenesis and axon pruning can lead to pernicious axon growth implicated in a number of developmental brain disorders such as autism, epilepsy, and schizophrenia (Saugstad 2011; Zhou *et al.* 2012). Recent findings linking the expression of the $\alpha 7$ gene *CHRNA7* to developmental disorders (Yasui *et al.* 2011; Adams *et al.* 2012) highlights the role of this receptor in brain development.

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Supporting information

Additional supporting information may be found in the online version of this article at the publisher's web-site:

Figure S1. The Alex-647 Bgtx (fBgtx) signal is in the presence of the selective $\alpha 7$ agonist PNU282987.

Table S1. Confirmation of protein bands in using mass spectrometry.

Table S2. Confirmation of protein bands using mass spectrometry.

Table S3. Pearson's r-values confirming the effect of drug treatment on various parameters of axon growth as presented in Fig. 3 of the manuscript.

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