

Brain-Derived Neurotrophic Factor and *trkB* Signaling in Parasympathetic Neurons: Relevance to Regulating $\alpha 7$ -Containing Nicotinic Receptors and Synaptic Function

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Parasympathetic neurons do not require neurotrophins for survival and are thought to lack high-affinity neurotrophin receptors (i.e., *trks*). We report here, however, that mRNAs encoding both brain-derived neurotrophic factor (BDNF) and its high-affinity receptor tropomyosin-related kinase B (*trkB*) are expressed in the parasympathetic chick ciliary ganglion (CG) and that BDNF-like protein is present in the ganglion and in the iris, an important peripheral target of ciliary neurons. Moreover, CG neurons express surface *trkB* and exogenous BDNF not only initiates *trk*-dependent signaling, but also alters nicotinic acetylcholine receptor (nAChR) expression and synaptic transmission. In particular, BDNF applied to CG neurons rapidly activates cAMP-dependent response element-binding protein (CREB), and over the long-term selectively upregulates expression of $\alpha 7$ -subunit-containing, homomeric nAChRs ($\alpha 7$ -nAChRs), increasing $\alpha 7$ -subunit mRNA levels, $\alpha 7$ -nAChR surface sites, and $\alpha 7$ -nAChR-mediated whole-cell currents. At nicotinic synapses formed on CG neurons in culture, brief and long-term BDNF treatments also increase the frequency of spontaneous EPSCs, most of which are mediated by heteromeric nAChRs containing $\alpha 3$, $\alpha 5$, $\beta 4$, and $\beta 2$ subunits ($\alpha 3^*$ -nAChRs) with a minor contribution from $\alpha 7$ -nAChRs. Our findings demonstrate unexpected roles for BDNF-induced, *trk*-dependent signaling in CG neurons, both in regulating expression of $\alpha 7$ -nAChRs and in enhancing transmission at $\alpha 3^*$ -nAChR-mediated synapses. The presence of BDNF-like protein in CG and iris target coupled with that of functional *trkB* on CG neurons raise the possibility that signals generated by endogenous BDNF similarly influence $\alpha 7$ -nAChRs and nicotinic synapses *in vivo*.

Key words: ciliary ganglion; nicotinic acetylcholine receptor; BDNF; *trkB*; patch-clamp; neurotrophin; bungarotoxin; EPSC; cAMP response element-binding protein (CREB); PACAP

Introduction

Neurotrophins [nerve growth factor (NGF), brain-derived neurotrophic factor (BDNF), and neurotrophin-3 (NT-3)] act via high-affinity tyrosine kinase-containing receptors (*trkA*, *trkB*, and *trkC*, respectively) to support the survival and growth of diverse neuron populations and influence the form and function of chemical synapses (Lewin and Barde, 1996; Kaplan and Miller, 2000; Huang and Reichardt, 2001). In particular, BDNF and sometimes NT-3, exert rapid, primarily presynaptic effects at central, autonomic, and neuromuscular synapses and produce long-term presynaptic and postsynaptic changes consistent with

altered gene expression (for review, see Lewin and Barde, 1996; Schuman, 1999; Poo, 2001). Thus, in addition to providing trophic support, neurotrophins also induce *trk*-dependent acute and long-term changes that coordinately influence synaptic interactions.

Parasympathetic neurons typified by those in the chicken ciliary ganglion (CG) do not require neurotrophins for survival (Helfand et al., 1976; Rohrer and Sommer, 1982; Lindsay et al., 1985; Kriegstein et al., 1998). Instead, CG neurons rely on other growth factors, notably ciliary neurotrophic factor (CNTF) (Leung et al., 1992; Finn et al., 1998) and glial-derived neurotrophic factor (GDNF) (Hashino et al., 2001) for trophic support. Moreover, studies using Northern and RNase protection assays failed to detect *trk* mRNA in ciliary ganglia (Dechant et al., 1993; Hallbook et al., 1995). These observations have led to the presumption that CG neurons lack *trks* (Huang and Reichardt, 2001).

As with sympathetic ganglion neurons and skeletal muscle fibers, fast chemical synapses on ciliary and other parasympathetic ganglion neurons are mediated by nicotinic acetylcholine receptors (nAChRs). In sympathetic neurons, NGF supports the expression of $\alpha 3$ -nAChR subunit protein (Yeh et al., 2001), an effect mirrored in pheochromocytoma cell line (PC12) cells, where NGF increases $\alpha 3$, $\alpha 5$, $\alpha 7$, $\beta 2$, and $\beta 4$ nAChR subunit mRNAs as well as

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nAChR function (Henderson et al., 1994; Takahashi et al., 1999). Also, sympathetic neurons overexpressing BDNF display increased preganglionic innervation density (Causing et al., 1997), indicative of long-term presynaptic effects. BDNF acting through trkB also regulates neuromuscular junction form and function. For example, BDNF rapidly enhances presynaptic release to increase the frequency and amplitude of spontaneous nAChR-mediated synaptic currents in nerve–muscle cultures (Lohof et al., 1993; Stoop and Poo, 1996). Over the long term, BDNF restores neuregulin levels, restricts axon sprouting, and maintains postsynaptic architecture in muscle disrupted by activity blockade (Loeb et al., 2002), whereas sustained trkB-mediated signaling is likely required to maintain postsynaptic nAChR clusters (Gonzalez et al., 1999). These findings prompted us to speculate that previous assays were perhaps insufficiently sensitive to detect trks expressed in ciliary ganglia and that neurotrophin–trk signaling, although not required for trophic support, might influence the components and function of nAChR-mediated synapses on CG neurons. We focused on BDNF–trkB signaling, and we have demonstrated expression of BDNF-like protein in ciliary ganglia and functional trkB on CG neurons. To explore synaptic relevance, the impact of BDNF–trkB signaling on $\alpha 7$ -nAChRs and $\alpha 3^*$ -nAChR-mediated synapses was assessed using CG neurons grown in cell culture. BDNF treatment upregulated expression of $\alpha 7$ -nAChRs after several days and increased the frequency of spontaneous synaptic currents within minutes. The results reveal an unanticipated relevance for BDNF–trkB signaling in parasympathetic CG neurons.

Materials and Methods

Neurons. CG neuron cultures were prepared under sterile conditions from embryonic day 8 (E8) chick embryos. Dissociated neurons were plated at one or two ganglion equivalents in 15 mm diameter polystyrene tissue culture wells or on 12-mm-diameter glass coverslips; both substrates were precoated with poly-DL-ornithine and laminin (Pugh and Margiotta, 2000; Chen et al., 2001). The standard culture medium consisted of minimum essential medium (MEM) containing 100 U/ml penicillin, 100 μ g/ml streptomycin, 2 mM glutamine, and 10% heat inactivated horse serum (MEM^{hs}; all components from Invitrogen, Rockville, MD) and was supplemented with 3% embryonic eye extract (Nishi and Berg, 1981). Neurons were maintained at 37°C in 95% air and 5% CO₂ for 4–7 d and received fresh culture medium every 2–3 d, conditions that support 100% survival of CG neurons for at least 7 d (Nishi and Berg, 1981). In test cultures the medium was further supplemented with BDNF (50 ng/ml, unless indicated otherwise) sometimes in conjunction with other reagents as described for individual experiments in Results. For some studies, CG neurons were acutely dissociated from E8 or E14 ganglia, as previously described (McNerney et al., 2000; Nai et al., 2003). Neurons were plated on acid-washed, poly-D-lysine-coated glass coverslips in electrophysiological recording solution (RS) containing (in mM): 145.0 NaCl, 5.3 KCl, 5.4 CaCl₂, 0.8 MgSO₄, 5.6 glucose, and 5.0 HEPES, pH 7.4 (Dichter and Fischbach, 1977) that was supplemented with 10% heat-inactivated horse serum (RS^{hs}). Acutely dissociated neurons were maintained in RS^{hs} at 37°C for 2–4 hr before use.

Conventional RT-PCR. The presence of mRNA encoding chicken trkB, BDNF, $\alpha 7$, and $\alpha 3$ -nAChR subunits, as well as β -actin (βA) or glyceraldehyde-3-phosphate dehydrogenase (GAPDH), was assessed by conventional reverse transcriptase-based PCR (RT-PCR) as previously described (Burns et al., 1997). Briefly, RNA was isolated from E8–E15 chick tissues or from CG neuron cultures using a one-step kit (RNAqueous; Ambion, Austin, TX). Total tissue RNA (1 μ g) was treated with Amplification Grade RNase-free DNase (1 U at 1 U/ μ l; Invitrogen), and then 25–200 ng of DNase-treated RNA used to synthesize cDNA using Superscript II reverse transcriptase (RT+; Invitrogen). The resulting cDNAs were then used as templates for PCR amplifications in 25 μ l reaction volumes containing 50 mM KCl, 20 mM Tris-HCl, 2.5 mM MgCl₂, 200 μ M dNTPs, 5 U/ μ l TaqDNA polymerase (Invitrogen), and 0.4 μ M forward (F)

and reverse (R) oligonucleotide primers (synthesized by Marshall University DNA Core Facility, Huntington, WV). The chicken-specific primers used were: *trkB* (Dechant et al., 1993): F, C₁₁₅₆TTCAGCTGGACAA-CCCTAC₁₁₇₅; R_K⁺, T₁₈₆₈GGAAGTCCTTGCGGGCATT₁₈₄₉; R_K⁻, GCCCTCTCTCATCTT; *BDNF* (Maisonpierre et al., 1992): F, G₂₈₇CAGTCAAGTGCCCTTG₃₀₃; R, G₇₄₈AGCCCACTATCTTCCCC₇₃₁; $\alpha 7$ -nAChR subunit (Couturier et al., 1990b): F, G₁₀₉₂GGGAAAAATGCTAAAT₁₁₀₉; R, G₁₆₁₄ACAGCCTCTACAAAGTT₁₅₉₃; $\alpha 3$ -nAChR subunit (Couturier et al., 1990a): F, A₉₈₅TGCCTGTATGGGTGAGAACT₁₀₀₅; R, T₁₂₂₆TGCCACTGAAATCGGAAAAAC₁₂₀₆; *GAPDH* (Stone et al., 1985): F, G₅₃₂CCATCACAGCCACACAGAA₅₅₁; R, A₉₈₀CCATCAAGTCCACAAACAG₉₆₁; and β -actin (GenBank accession number L08165): F, A₈₆₀TCTTTCTTGGGTATGGA₈₇₇; R, A₁₁₃₄CATCTGCTGGAAGGTCC₁₁₁₇.

The two trkB primer pairs (F/R_K⁺ and F/R_K⁻) correspond to those shown previously to amplify kinase-containing (full-length) and truncated (kinase-deleted) chicken trkB isoforms, respectively (Garner et al., 1996). The F/R_K⁺ pair is not predicted to hybridize with chicken trkA (Schropel et al., 1995) or trkC (Garner and Large, 1994) cDNAs. The $\alpha 7$ - and $\alpha 3$ -nAChR subunit primer pairs both amplify products within non-conserved regions of their respective cytoplasmic domains, located between transmembrane segments III and IV (Schoepfer et al., 1990). The trkB and AChR subunit primers were optimized for amplification, and the reactions were performed in the linear range of the assay (25–29 cycles). PCR products were separated on 1.0% agarose gels stained with ethidium bromide. Identical reactions lacking RT served as controls for possible amplification of genomic DNA and were consistently negative. Changes in the levels of $\alpha 7$ and $\alpha 3$ mRNAs in response to BDNF treatment were estimated semiquantitatively after digitizing gel images using Kodak 1D Image Analysis software (Eastman Kodak, Rochester, NY) from the ratio of PCR product intensities to those of βA from the same cultures.

Real-time PCR. Changes in $\alpha 7$ - and $\alpha 3$ -nAChR subunit mRNA levels induced by BDNF were confirmed using RT-based real-time PCR. cDNA samples corresponding to 50 ng of input RNA were combined with Taqman universal PCR master mix (Roche, Branchburg, NJ), F and R primers (0.4 μ M), and Taqman probe (0.1 μ M) [with 6-FAM (carboxyfluorescein, reporter dye) and TAMRA (tetramethylrhodamine, quencher dye) inserted at 5' and 3' ends, respectively]. Selection of the following primers and probes was optimized using Applied Biosystems (Foster City, CA) Primer-Express software, with $\alpha 7$ - and $\alpha 3$ -nAChR subunit primers chosen to amplify regions within transmembrane segments III and IV, and span intron-exon boundaries (Schoepfer et al., 1990): $\alpha 7$ -nAChR subunit (Couturier et al., 1990b): F, C₁₀₂₀CATGATTATTGTTGGCCTCTCT₁₀₄₂; R, T₁₂₁₀CGGCCCTGTTATGTTGAC₁₁₉₀; Probe, A₁₁₁₅GAGTCATCCTTCTGAATTGGTGTGCTTGGT₁₁₄₅; $\alpha 3$ -nAChR subunit (Couturier et al., 1990a): F, G₁₁₇₈CAGCTGCTGCCAGTACCA₁₁₉₆; R, A₁₃₉₈ATGACCATGGCAACATATTTC₁₃₇₆; Probe, T₁₂₁₆TCAGTGGCAATC-TACAAGAAGTTCAGC₁₂₄₅; and *GAPDH* (Stone et al., 1985): F, C₁₇₉₅CGTCTCTCTGGCAAAGTC₁₈₁₄; R, A₂₃₇₄ACATACTCAGC-ACCTGCATCTG₂₃₅₂; Probe, A₂₂₁₁TCAATGGGCACGCCATCACTATCTTCC₂₂₂₈.

Twenty-five microliter PCRs were performed in triplicate using a GeneAmp 5700 sequence detection system (Applied Biosystems). This system allows the increase in PCR product to be monitored directly based on the threshold number of cycles (CT) required to produce a detectable change in fluorescence (ΔF) resulting from the release of probe. Relative levels of $\alpha 7$ - and $\alpha 3$ -nAChR cDNA ($R_{\alpha 7}$, $R_{\alpha 3}$) in control and BDNF-treated cultures were calculated from the difference in CT values ($\Delta CT = CT_{\text{control}} - CT_{\text{BDNF}}$) for $\alpha 7$ or $\alpha 3$ amplifications ($\Delta CT_{\alpha 7}$, $\Delta CT_{\alpha 3}$) compared with those for the housekeeping gene, GAPDH (ΔCT_{GAPDH}) using:

$$R_{\alpha} = (E_{\alpha}^{\Delta CT_{\alpha}}) / (E_{\text{GAPDH}}^{\Delta CT_{\text{GAPDH}}}) \quad (1)$$

In Equation 1, E_{α} and E_{GAPDH} are the real-time PCR amplification efficiencies determined in separate studies from the slope of CT versus input log cDNA dilution, where $E = 10^{-1/\text{slope}}$. E values for amplifying $\alpha 7$, $\alpha 3$, and GAPDH cDNAs were 2.10, 2.10, and 2.23, respectively.

Immunocytochemistry. A polyclonal antibody (pAb) generated against

the extracellular domain of chicken trkB (#R22781) that does not recognize trkA or trkC (von Bartheld et al., 1996) was generously provided by Dr. Frances Lefcort (Montana State University). pAb recognizing Ser₁₃₃-phosphorylated cAMP response element (CRE) binding protein (p-CREB) was purchased from Cell Signaling Technology (Beverly, MA). Ciliary and dorsal root ganglia (DRG) were fixed for 1–4 hr in 4% paraformaldehyde prepared in 0.15 M PBS at pH 7.4 (PBS), washed in PBS, cryoprotected in PBS containing 30% sucrose, embedded in OCT (Miles Laboratories, Elkhart, IN), cryosectioned at 10 μ m, and mounted on glass slides. After rehydration, sections were blocked for 1 hr at room temperature in 30 mM Tris and 150 mM NaCl containing 0.4% Triton X-100, 1% glycine, 10% goat serum, and 3% bovine serum albumin. trkB antibody (Ab) was applied to sections in blocking solution containing 4% goat serum (1:1000, 4°C, 16 hr), and after washing, secondary Ab (AlexaFluor594-conjugated anti-rabbit IgG; Molecular Probes, Eugene, OR) was applied in the same solution (1:400, 22°C, 1 hr). Sections were then washed, dipped in distilled water, and mounted in Vectashield (Vector Laboratories, Burlingame, CA). Acutely dissociated CG neurons or CG neuron cultures, both on glass coverslips, were fixed for 0.5–1.0 hr in 2–4% paraformaldehyde and blocked in PBS containing 10% donkey or goat serum. Coverslips were then incubated in trkB Ab (1:2000, 37°C, 2 hr) treated with Cy3-conjugated anti-rabbit IgG (1:400, 1 hr, 37°C; The Jackson Laboratory, Bar Harbor, ME) in PBS containing 4% serum, washed, and mounted. A similar protocol was followed for p-CREB immunostaining except that the block and wash buffers contained 0.1% Triton X-100, p-CREB Ab was applied (1:400, 4°C, 16 hr), and the secondary Ab was AlexaFluor488-conjugated anti-rabbit IgG (1:400, 22°C, 1 hr).

Image analysis. Immunostained preparations were viewed using epifluorescence microscopy (BX50, UplanFL 40 \times , 0.75 numerical aperture objective; Olympus, Tokyo, Japan), and images were acquired and processed using a SenSys KAF-1400 cooled digital CCD camera under the control of IP Lab software (version 3.6; Scanalytics, Reading, PA) as described previously (Chen et al., 2001). Neurons were considered p-CREB-positive if the mean fluorescence intensity of pixels in an elliptical region of interest (ROI) superimposed over the nucleus exceeded that of the ROI when placed over cytoplasm by >15%.

ELISAs. The presence of BDNF in chicken tissue homogenates and tissue culture medium components was assessed using a commercial BDNF sandwich ELISA kit having no significant cross-reactivity with NGF, NT4/5, or NT3 (Chemikine; Chemicon, Temecula, CA). The ELISA uses rabbit polyclonal antibodies (raised against human BDNF) to capture BDNF from the sample, and a biotinylated mouse monoclonal antibody to detect the captured BDNF. Because mammalian and chicken BDNF share all but seven amino acids, with the mismatches distributed along the entire length of the peptide (Isackson et al., 1991), the kit antibodies likely recognize chicken BDNF. Nevertheless, we refer here to detection of “BDNF-like protein,” with levels quantified within the linear range of the assay (7.8–500 pg/ml) using recombinant human BDNF as standard.

α -Bungarotoxin binding. CG neurons were plated at one or two ganglion equivalents per well and grown in culture wells for 4–5 d. Neurons in triplicate culture wells were washed twice in MEM^{hs}, incubated in MEM^{hs} containing 10 nM [¹²⁵I]- α -Bungarotoxin (α -Bgt) (specific activity = 130–140 Ci/mmol; PerkinElmer, Boston, MA) for 1 hr at 37°C, and then washed three times with MEM^{hs}. We previously showed that these conditions are sufficient to saturate surface α -Bgt sites on dissociated CG neurons (McNerney et al., 2000). Nonspecific binding was determined in parallel wells by including 100 μ M D-tubocurarine with 10 nM [¹²⁵I]- α -Bgt. After labeling and washing, the wells were scraped in 500 μ l 0.6 N NaOH, the solution was collected, and [¹²⁵I]- α -Bgt radioactivity was determined using a Beckman Instruments (Fullerton, CA) G-5500 gamma counter.

Electrophysiology. Whole-cell recordings were obtained at 21–23°C from CG neurons after 3–5 d in culture. Patch pipettes were fabricated from Corning 8161 glass tubing (WPI, Sarasota, FL), filled with (in mM) 145.6 CsCl, 1.2 CaCl₂, 2.0 EGTA, 15.4 glucose, and 5.0 Na-HEPES, pH 7.3, and had tip impedances of 2–3 M Ω . To induce nAChR currents, neurons were bathed in RS^{hs}, held at –70 mV, and 20 μ M nicotine (Nic)

applied in RS by rapid pressure microperfusion (at 10–12 psi) from a delivery pipette (4–6 μ m tip diameter) positioned \approx 5–10 μ m from the neuron soma. We previously showed that fast-onset, rapidly desensitizing α 7-nAChR-mediated whole-cell currents induced by 20 μ M Nic in this manner are indistinguishable in amplitude from those obtained using fast piezoelectric switching (Nai et al., 2003). The fast (α 7-nAChR-mediated) and slower (α 3*-nAChR-mediated) decaying current components induced by 20 μ M Nic were identified and analyzed using Clampfit (pClamp 6.0 or 8.0; Axon Instruments, Foster City, CA) as previously described (Nai et al., 2003). For analysis, peak Nic-induced response component amplitudes (in picoamperes) were normalized to neuron soma membrane capacitance (in picofarads). To quantify BDNF effects, whole-cell Nic responses (in picoamperes per picofarad) obtained from treated neurons were normalized to those for control neurons from the same cultures. To assess synaptic function, spontaneous EPSCs (sEPSCs) were acquired at –70 mV for 2–5 min, without stimulation, as previously described (Chen et al., 2001). For these experiments, horse serum was sometimes omitted from the recording solution, without discernible effect on the results. Synaptic current frequency and amplitude analyses were subsequently performed using either BASIC-23 programs written in-house, or commercially available software (Mini Analysis 5.6.12; Synaptosoft Inc., Decatur, GA). Briefly, sEPSC frequency values obtained from BDNF-treated neurons were normalized to those from control neurons from the same culture platings. In addition, sEPSCs were extracted from selected records displaying >50 non-overlapping events, and the component amplitude and decay time constant values were pooled for control and BDNF-treated neurons.

Statistics. All parameter values are expressed as mean \pm SEM. Unless indicated otherwise, the statistical significance of paired and unpaired numerical comparisons was determined using the appropriate two-tailed *t* test (*p* < 0.05).

Results

Expression of trkB mRNA and protein

PCR primers specific for kinase-containing (K⁺) full-length trkB (Garner et al., 1996) amplified a \approx 700 bp product from both E8 and E14 CG cDNA templates (Fig. 1*a,b*). The CG product size was consistent with that predicted for chicken trkB (713 bp) (Garner et al., 1996) and indistinguishable from that obtained in amplifications from E15 DRG, previously shown to express abundant trkB mRNA (Hallbook et al., 1995) and protein (Anderson, 1999; Rifkin et al., 2000). In addition, trkB products from E14 CG and E15 DRG yielded identical restriction profiles after digestion with *Bam*HI (data not shown) or *Hba*II (Fig. 1*c*), with fragment sizes as predicted for digestion of K⁺ trkB cDNA (Dechant et al., 1993). Truncated trkB isoforms lacking the kinase domain (K[–]) but containing variable juxtamembrane insertions are also expressed in the chicken nervous system (Garner et al., 1996), and K[–]-specific primers amplified products of expected sizes (\approx 400, 500, and 600 bp) from both DRG and CG (Fig. 1*a,b*). In each case, the PCR amplifications from DRG and CG sources were specific for cDNA in the sense that they were absent when the synthesis reaction lacked RT (data not shown). Although the significance of the truncated trkB transcripts was not studied here, the results demonstrate that both truncated and full-length trkB transcripts are expressed in CG during E8–E14, a developmental window when nicotinic synapses formed on the neurons undergo substantial structural and functional maturation (Landmesser and Pilar, 1972, 1974a).

The presence of trkB protein on CG neurons was demonstrated by fluorescence immunolabeling (Fig. 2) using an Ab that recognizes the extracellular domain of chicken trkB (but not trkA or trkC; von Bartheld et al., 1996). Specific trkB labeling, similar to but somewhat less intense than that seen for E15 DRG sections, was evident in both E8 and E14 CG sections (Fig. 2*a,b,c,f*) and was localized to the neuron surface where it increased between

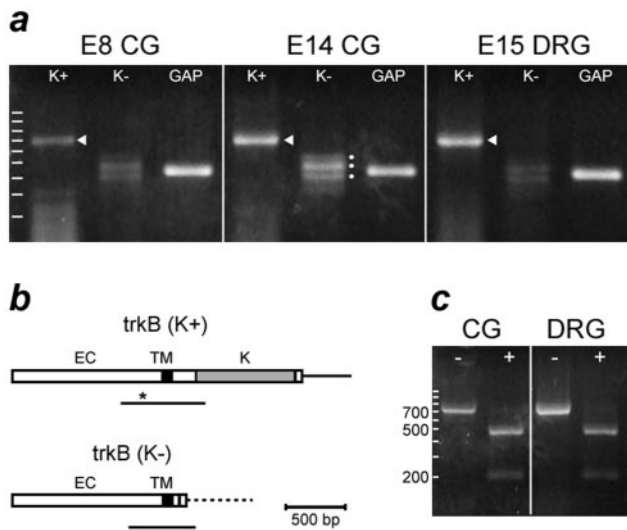


Figure 1. Detection of *trkB* transcripts in ciliary ganglia. *a*, PCR amplifications were conducted on cDNAs from E8 CG, E14 CG, and E15 DRG using primer pairs specific for chicken full-length *trkB* (K^+), kinase-deleted *trkB* (K^-), or GAPDH (GAP), and the resulting products were separated by gel electrophoresis. Arrowheads at ≈ 700 bp and dots at ≈ 400 , 500, and 600 bp (E14 CG shown) mark product sizes expected for K^+ *trkB* and for the truncated K^- *trkB* isoforms, respectively. *b*, Schematics of K^+ and K^- *trkB* isoforms showing extracellular (EC), transmembrane (TM), and kinase (K) domains. The striped bar in the bottom schematic depicts the variable-length juxtamembrane region responsible for the multiple amplification products found for K^- *trkB* in *a*. Horizontal lines indicate the *trkB* products expected for the K^+ (713 bp) and largest K^- (≈ 600 bp) *trkB* isoforms. *c*, *Hpa*II digestion (+) of CG and DRG K^+ *trkB* amplification products (site marked by * in *b*) yielded restriction fragments of identical and predicted sizes (499 and 214 bp). Undigested K^+ *trkB* (-). Lane markers in *a* and *c* depict a 100 bp DNA ladder (200–1000 bp).

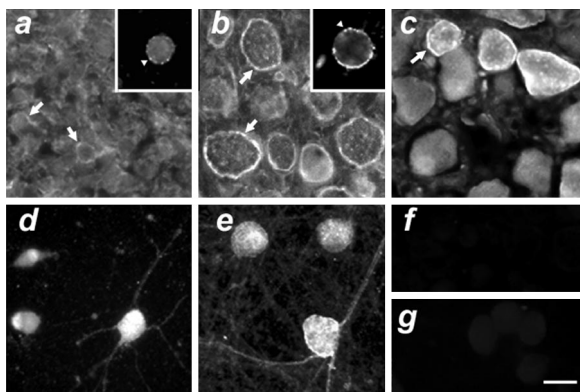


Figure 2. Localization of *trkB* protein to ciliary ganglion neurons. *a–c*, Ganglion sections obtained from E8 CG, E14 CG, and E15 DRG, displayed neuronal immunoreactivity after labeling with primary antiserum recognizing an extracellular epitope of chicken *trkB*. Both intracellular and cell-surface *trkB* labeling (arrows) was evident. Insets in *a* and *b* show freshly dissociated E8 and E14 CG neurons displaying punctate cell-surface *trkB* labeling (arrowheads). *d, e*, *trkB* labeling of E8 CG neuron somata and processes after 8 hr (*d*) and on day 4 (d4) in culture (*e*). *f, g*, To demonstrate specificity, E14 CG sections (*f*) and d4 CG cultures (*g*) processed without the *trkB* primary antiserum were unlabeled, as were E8 CG, E15 DRG, acutely dissociated CG neurons, and 8 hr CG cultures (data not shown). Scale bar: (in *g*), *a–g*, 20 μ m.

the two developmental ages. In CG neuron cultures, the somata and processes of neurons displayed specific *trkB* labeling that became more extensive and intense between 8 hr and 4 d in culture (Fig. 2*d,e,g*), a period when functional synapses are formed and increase in activity (Chen et al., 2001). At 4 d in culture, $\sim 80\%$ of CG neurons scored positive for *trkB* immunoreactivity. These findings demonstrate that *trkB* protein is ex-

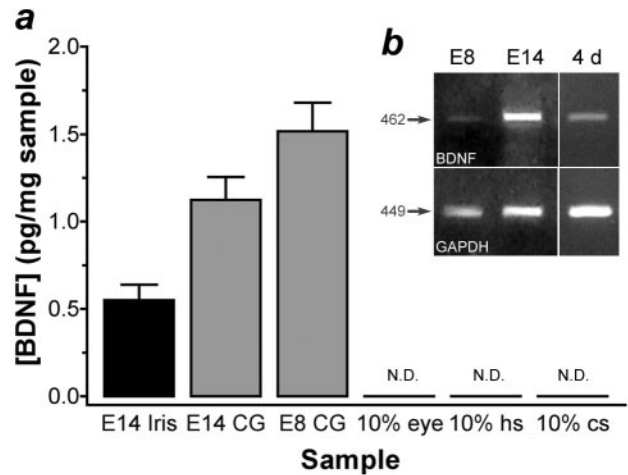


Figure 3. Detection of BDNF protein and mRNA. *a*, ELISAs demonstrate the presence of BDNF-like protein in tissue homogenates prepared from a ciliary neuron target, the iris constrictor muscle (Iris, black bar), as well as from E8 and E14 CG (gray bars). BDNF was not present at detectable levels (ND) in 10% eye extract (10% eye), 10% heat inactivated horse serum (10% hs), or 10% chicken serum (10% cs). *b*, PCR primers specific for chicken BDNF (top row) amplified products consistent with the expected size of 462 bp (arrow) from cDNA derived from E8 and E14 CG, and in separate experiments from E8 CG neurons after 4 d growth in culture. GAPDH amplifications (bottom row, 449 bp) were performed as positive controls in parallel reactions from paired experiments.

pressed by CG neurons and, because dissociated neurons in acute and culture preparations were not permeabilized, indicate that a substantial fraction is localized on the cell surface. Taken together, the mRNA and protein studies further suggest that catalytically competent, high-affinity BDNF receptors are present in the ganglion and that their expression on CG neurons increases during periods of synaptic differentiation both *in vivo* and in cell culture.

Functional relevance of BDNF and *trkB*

To be relevant *in vivo*, endogenous BDNF should both be present in the CG and be able to elicit *trk*-dependent signaling in the neurons. Because BDNF detected in spinal cord ventral horn results from both local synthesis and retrograde transport to motor neurons from striated muscle (Koliastzos et al., 1993), we tested for the presence of BDNF both in CG and in the iris, a ciliary neuron target that like the ciliary body is primarily striated muscle in birds (Marwitt et al., 1971). Using a commercial ELISA, we detected BDNF-like protein in E14 iris muscle as well as in E14 and in E8 CG (Fig. 3*a*). The assay failed to detect BDNF in 10% heat inactivated horse serum or whole eye extract, routinely used at 10 and 3%, respectively, as supplements to CG culture medium. The assay also failed to detect BDNF-like protein in intact chicken serum. Relevant to our culture experiments, we presume that dilution of BDNF derived from the iris and ciliary muscle during eye extract preparation reduces its concentration below the detection limit of the assay (7.8 pg/ml). BDNF-like protein present in E8 and E14 CG may be the source of the strong BDNF immunoreactivity previously reported at the same developmental ages for accessory oculomotor neurons (Steljes et al., 1999) which provide preganglionic input to the CG. In addition to arriving by retrograde transport from the intraocular muscle targets, however, the BDNF-like protein present in the CG may also result from local synthesis, because BDNF mRNA was detectable by RT-PCR in E8 and E14 ganglia and in CG neurons maintained in standard culture medium for 4 d (Fig. 3*b*).

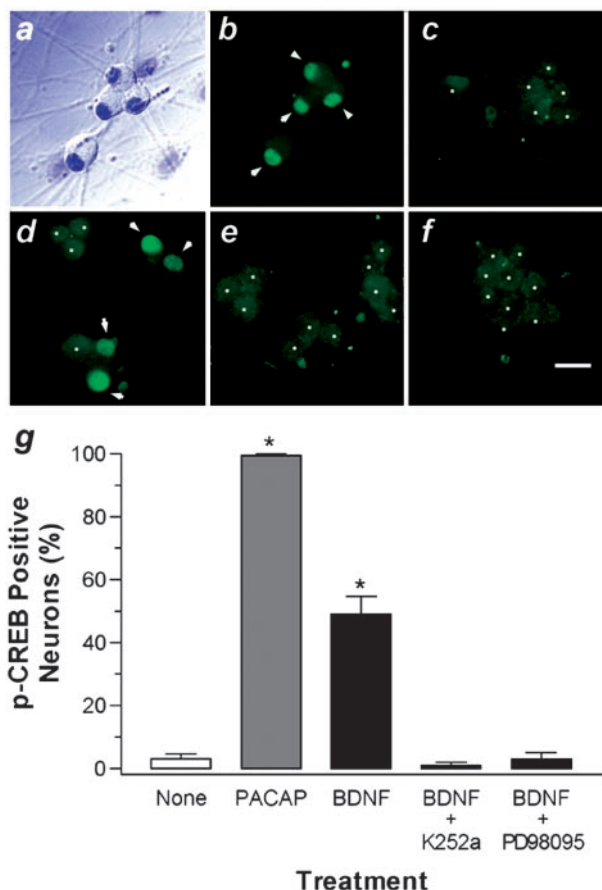


Figure 4. BDNF induction of CREB phosphorylation indicates trkB receptors on CG neurons are functional. *a–c*, Nearly all CG neuron nuclei (labeled by DAPI staining in *a*) displayed detectable p-CREB immunoreactivity after treatment with 100 nM PACAP (*b*), whereas p-CREB-immunoreactive neuronal nuclei were virtually absent in untreated cultures (*c*). *d–f*, After incubation with BDNF, many neuron nuclei displayed detectable p-CREB immunoreactivity (*d*), which was absent in cultures cotreated with 100 ng/ml BDNF and 200 nM K252a (*e*) or 50 μ M PD98059 (*f*). Scale bar: (in *f*), *a–f*, 20 μ m. Arrowheads and dots mark neuronal nuclei scoring as p-CREB-positive and -negative, respectively. *g*, Summary of treatment results. Bars represent the mean percentage of neurons per field with p-CREB-immunoreactive nuclei after the indicated treatments ($N = 85–321$ neurons in 6–18 fields from 2 experiments). Asterisks indicate a significant difference ($p < 0.001$, unpaired *t* test) from untreated cultures tested in parallel.

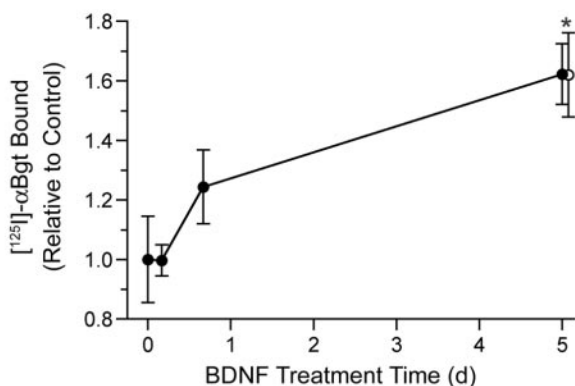
To determine if the trkB protein expressed on CG neurons represents functional receptor, we tested the ability of applied BDNF to cause phosphorylation of CREB, a cAMP- and Ca^{2+} -regulated transcription factor (for review, see Impey et al., 1996; Greenberg and Ziff, 2001; Deisseroth et al., 2003), whose activation is a hallmark of trk-dependent neurotrophin signaling (Finkbeiner et al., 1997) (Fig. 4). For this purpose, CG neurons grown in culture for 4–5 d were challenged with BDNF (100–200 ng/ml; 10–15 min) or, as a positive control, with pituitary adenylate cyclase activating polypeptide (PACAP; 100 nM, 10–15 min), previously shown to cause robust increases in intracellular cAMP and Ca^{2+} (Margiotta and Pardi, 1995; Pardi and Margiotta, 1999), and then tested the cultures for p-CREB immunoreactivity. A similar immunocytochemical approach was previously shown to provide a convenient all-or-none assay for CREB activation in single hippocampal neurons (Hu et al., 2002). After treatment with BDNF, 49 \pm 6% of 321 CG neurons from 18 fields ($N = 321$, 18) scored as p-CREB-positive, compared with 99 \pm 2% ($N = 132$, 9) after PACAP treatment and 3 \pm 2% ($N = 179$, 10) in untreated control cultures assayed in parallel (Fig. 4*a–d, g*).

Consistent with a requirement for trkB signaling, the proportion of p-CREB positive neurons induced by BDNF dropped to control levels ($1 \pm 1\%$; $N = 114$, 6) after 1 hr preincubation and 15 min cotreatment with K252a (Fig. 4*e, g*), a trk-selective tyrosine kinase inhibitor (Pizzorusso et al., 2000). Preincubation (1 hr) and 15 min cotreatment with PD98059, an MEK1 inhibitor that blocks the neurotrophin-activated, RAS-dependent signaling pathway leading to CREB activation (Ying et al., 2002) similarly reduced the proportion of p-CREB positive neurons induced by BDNF to 3 \pm 2% ($N = 85$, 6) (Fig. 4*f, g*). The observation that 51% of neurons showed no detectable response to BDNF in this assay cannot be explained by limited CREB availability because nearly all nuclei were immunoreactive after PACAP treatment. The difference might instead reflect suboptimal BDNF treatment times or heterogeneity of functional trkB expression levels. In either case, the protein and p-CREB assays (Figs. 3, 4) demonstrate that an endogenous source of BDNF-like protein is present in the parasympathetic CG, where it is poised to activate functionally competent trkB receptors present on the neurons and thereby recruit appropriate signal pathways leading to CREB activation.

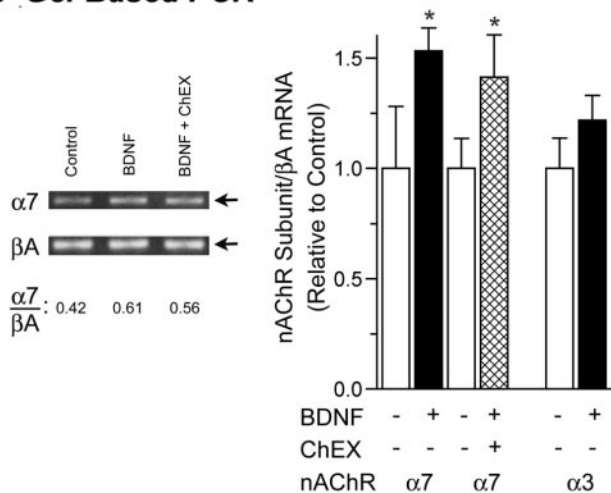
BDNF upregulates $\alpha 7$ -nAChRs

Because neurotrophins are not required for full survival of CG neurons, we examined possible roles for BDNF–trkB signaling in regulating the components and function of nicotinic synapses on the neurons. $\alpha 7$ -nAChRs were assessed as potential targets of BDNF–trkB signaling because they can rapidly modulate transmission (McGehee et al., 1995; Gray et al., 1996; Wonncott, 1997; Margiotta and Pugh, 2004), an action resembling the increased synaptic efficacy produced by BDNF (for review, see McAllister et al., 1999; Schneider and Poo, 2000; Poo, 2001). α -Bgt binds with high affinity to $\alpha 7$ -nAChRs (Couturier et al., 1990b) and [^{125}I]- α -Bgt was therefore used to quantify surface $\alpha 7$ -nAChRs on CG neurons (Fig. 5*a*), as previously described (McNerney et al., 2000). In CG neuron cultures grown with BDNF for varying times before assay at 5 d, [^{125}I]- α -Bgt binding was unchanged relative to control cultures after 4 hr exposure, but increased nominally (24%) after 16 hr. Extending the treatment time to the full 5 d culture period resulted in levels of α -Bgt binding that were significantly higher ($62 \pm 10\%$; $p < 0.01$) in CG cultures exposed to BDNF relative to control cultures assayed in parallel ($N = 10$ for both). Actual levels of [^{125}I]- α -Bgt bound in control cultures and cultures treated with BDNF for 5 d were 3.9 ± 0.2 and 6.0 ± 0.4 fmol/CG equivalent, respectively. In principle, the increased levels of $\alpha 7$ -nAChRs seen after exposure to BDNF could have been caused by activation of either trkB or the low-affinity neurotrophin receptor (p75^{NTR}) that is also present on CG neurons (Lee et al., 2002). The latter possibility is unlikely, however, because an identical elevation ($62 \pm 14\%$; $N = 3$) was observed when BDNF was coapplied for 5 d with ChEX, a pAb that recognizes and blocks chicken p75^{NTR} but not trk function (Wescamp and Reichardt, 1991). The ability of long-term BDNF exposure to upregulate $\alpha 7$ -nAChRs may reflect increased $\alpha 7$ -nAChR subunit gene expression because levels of $\alpha 7$ -nAChR subunit relative to βA mRNA, determined by semiquantitative RT-PCR, were elevated significantly (by $53 \pm 10\%$) in cultures treated with BDNF for 4–5 d compared with untreated control cultures, tested in parallel ($N = 6$ each) (Fig. 5*b*). As with the protein assays, BDNF also increased $\alpha 7$ -nAChR subunit mRNA in cultures with p75^{NTR} blocked by coapplication with ChEX ($41 \pm 15\%$; $N = 3$). Using real-time PCR, a 98 \pm 26% ($N = 5$) increase in $\alpha 7$ -nAChR subunit relative to GAPDH mRNA was

a α Bgt Binding



b Gel-Based PCR



c Real-Time PCR

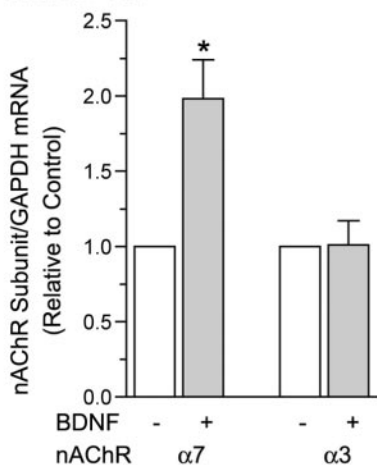


Figure 5. BDNF increases $\alpha 7$ -nAChR surface sites and $\alpha 7$ -nAChR subunit mRNA. *a*, $\alpha 7$ -nAChR levels were determined by quantifying $[\text{}^{125}\text{I}]\text{-}\alpha\text{Bgt}$ surface binding sites in CG neuron cultures maintained for 5 d. Data points indicate mean (\pm SEM) femtomoles of $[\text{}^{125}\text{I}]\text{-}\alpha\text{Bgt}$ bound per ganglion equivalent relative to control cultures after exposure to BDNF for 0 hr ($N = 12$), 4 hr (from day 5; $N = 3$), 16 hr (from day 4; $N = 6$), or 5 d (from day 0; $N = 10$). The open circle depicts relative $[\text{}^{125}\text{I}]\text{-}\alpha\text{Bgt}$ binding levels for cultures treated with both BDNF and ChEX ($N = 3$). Asterisk indicates a significant increase in $[\text{}^{125}\text{I}]\text{-}\alpha\text{Bgt}$ binding after 5 d exposure compared with untreated control cultures from the same platings ($p < 0.05$) and applies to BDNF incubations with and without ChEX. *b*, Separate PCR amplifications were conducted on cDNAs isolated from control or BDNF-treated (4–5 d) CG neuron cultures (with or without ChEX) on d 4 or d 5 using primer pairs specific for the indicated chicken nAChR subunit or βA . *b*, Left,

observed (Fig. 5c). In both cases, the BDNF-induced increases in $\alpha 7$ -nAChR subunit mRNA were selective in the sense that identical treatments failed to significantly alter $\alpha 3$ -nAChR subunit mRNA levels (Fig. 5b,c).

Having demonstrated that BDNF induces trk-dependent increases in levels of $\alpha 7$ -nAChR protein and mRNA in CG neurons, we next sought to determine if the same treatments also enhanced $\alpha 7$ -nAChR-mediated currents. Rapid application of nicotine (Nic; 20 μM) to CG neurons grown in culture typically induces a whole-cell current response featuring an initial fast-desensitizing component that is blocked by α -Bgt (Fig. 6a) and hence mediated by $\alpha 7$ -nAChRs (Pardi and Margiotta, 1999; McNerney et al., 2000; Nai et al., 2003). Whereas $65 \pm 5\%$ ($N = 80$ neurons; 5 platings) of CG neurons grown in standard culture medium displayed rapidly decaying, $\alpha 7$ -nAChR-mediated currents, BDNF treatment for 3–4 d increased the proportion to $83 \pm 4\%$ ($N = 76, 5$). In such cases, the peak $\alpha 7$ -nAChR current values relative to membrane capacitance (I_{fast}/C_m , pA/pF) were $49 \pm 14\%$ ($N = 63, 5$) larger for neurons from cultures treated with BDNF compared with untreated controls ($N = 52, 5$) tested in parallel (Fig. 6b,c). Consistent with the time course for upregulation of surface $\alpha 7$ -nAChRs seen in the $[\text{}^{125}\text{I}]\text{-}\alpha\text{Bgt}$ binding studies, BDNF treatments for 10–30 min or 16–24 hr produced only nominal increase in I_{fast}/C_m relative to untreated controls tested in parallel (Fig. 6b) (data not shown, $p > 0.05$ for both). The slowly decaying component of the Nic-induced current is mediated primarily by heteromeric $\alpha 3^*$ -nAChRs (Nai et al., 2003) that contain $\alpha 3$, $\alpha 5$, $\beta 4$, and occasionally $\beta 2$ subunits, but lack $\alpha 7$ subunits (Vernallis et al., 1993; Conroy and Berg, 1995) and are insensitive to α -Bgt (Fig. 6a). The ability of BDNF to increase I_{fast}/C_m was specific for $\alpha 7$ -nAChRs because slow currents (I_{slow}/C_m) attributable to $\alpha 3^*$ -nAChRs and present in all neurons, were unchanged after exposure to BDNF for 10–30 min, 16–24 hr, or 4–5 d (Fig. 6b). In addition, the 4–5 d BDNF treatments had no discernible effects on membrane capacitance or the voltage sensitivity or maximal values of voltage-activated Na^+ or Ca^{2+} currents (data not shown). In summary, the size, latency, and specificity of the increased $\alpha 7$ -nAChR current responses seen after chronic BDNF treatment are consistent with the BDNF-activated trkB-dependent upregulation of $\alpha 7$ -nAChR mRNA and protein that occur over a similar time course.

BDNF increases activity at nicotinic synapses

Functional synapses form between CG neurons in culture (Margiotta and Berg, 1982) and display spontaneous, impulse-driven

←
The resulting products were separated by gel electrophoresis and had sizes appropriate for βA (275 bp), $\alpha 7$ -nAChR subunit (522 bp) (arrows), and $\alpha 3$ -nAChR subunit (252 bp, data not shown). In the example, note that the intensity of $\alpha 7$ -nAChR subunit relative to βA product was higher for BDNF-treated cultures (with or without ChEX) than for controls. *b*, Right, Summary of mean (\pm SEM) relative $\alpha 7$ -nAChR and $\alpha 3$ -nAChR subunit product fluorescence intensity under different treatment conditions. In this gel-based assay, the relative levels of $\alpha 7$ -nAChR subunit mRNA were $53 \pm 10\%$ higher in BDNF-treated (black bar) compared with control cultures (white bar) within the same experiments ($*p < 0.01$; $N = 6$; paired *t* test), and a similar increase ($41 \pm 15\%$) persisted when BDNF was applied in the presence of ChEX (hatched bar) to block p75^{NTR} ($p < 0.05$; $N = 3$). In contrast, the BDNF treatments failed to significantly change mRNA levels for $\alpha 3$ -nAChR subunit ($p > 0.05$; $N = 5$). *c*, For confirmation, real-time PCR amplifications were conducted on cDNAs isolated from control or BDNF-treated (4–5 d) CG neuron cultures using primer pairs specific for the indicated chicken nAChR subunit or GAPDH. Using this approach, the levels of $\alpha 7$ -nAChR subunit mRNA were $98 \pm 26\%$ higher in BDNF-treated (gray bar) compared with control cultures (white bar) ($*p < 0.02$; $N = 5$; unpaired *t* test), whereas mRNA levels for $\alpha 3$ -nAChR subunit were unchanged ($p > 0.1$; $N = 4$).

nicotinic EPSCs (sEPSCs) (Fig. 7). We previously demonstrated that although $\alpha 7$ -nAChRs contribute to the sEPSCs, the vast majority require $\alpha 3^*$ -nAChRs because α -Conotoxin-MII, which blocks $\alpha 3^*$ - but not $\alpha 7$ -nAChRs on CG neurons (Nai et al., 2003) reduced sEPSC frequency by 95% (Chen et al., 2001). Exposure to BDNF substantially increased the overall frequency of sEPSCs (Fig. 7), most of which display slow decay kinetics indicative of a major contribution from $\alpha 3^*$ -nAChRs (Chen et al., 2001). After 16–24 hr BDNF treatment, sEPSC frequency increased approximately threefold (2.69 ± 0.35 ; $N = 46$) relative to untreated control neurons from the same five cultures tested in parallel (1.00 ± 0.17 ; $N = 39$), with a smaller yet significant increase seen after 4–5 d treatment (Fig. 7A,B). This effect is reminiscent of that seen at other synapses, where BDNF increases EPSC frequency by a presumed presynaptic mechanism (McAllister et al., 1999; Schneider and Poo, 2000; Poo, 2001) (see below). To determine if BDNF also altered sEPSC amplitudes, well separated individual synaptic currents were extracted from selected records, and components mediated by $\alpha 7$ - and $\alpha 3^*$ -nAChRs were identified by their diagnostic fast and slow decay kinetics, as previously described (Chen et al., 2001). The amplitudes of fast, $\alpha 7$ -nAChR-mediated sEPSCs identified in this manner increased after 16–24 hr of BDNF treatment (Fig. 7D), shifting by 32% from a median value of -9.6 pA in controls to -12.7 pA in BDNF-treated cultures ($N = 4$ neurons each; $p < 0.0004$; Mann–Whitney U and Kolmogorov–Smirnov tests). The effect was selective for $\alpha 7$ -nAChR-mediated sEPSCs because, despite increasing in frequency, slow $\alpha 3^*$ -nAChR-mediated sEPSCs displayed amplitudes that were unchanged by BDNF treatment (Fig. 7E). Recent studies indicate that chronic exposure to BDNF increases the proportion of postsynaptic $\alpha 7$ -nAChR clusters on hippocampal neurons (Kawai et al., 2002). Because $\alpha 3^*$ -nAChR mediated sEPSC amplitudes were unchanged, a similar postsynaptic accumulation of $\alpha 7$ -nAChRs may also explain the larger amplitude fast sEPSCs seen here after 16–24 hr exposure to BDNF.

Although significant and $\alpha 7$ -nAChR-selective, the changes in fast sEPSC amplitudes after 16–24 hr BDNF treatment were small in comparison to the accompanying threefold increase in the frequency of (primarily) $\alpha 3^*$ -nAChR-mediated sEPSCs. Studies in other systems suggest that this latter, more dramatic effect is likely to be presynaptic in origin, possibly resulting from changes in intracellular Ca^{2+} dynamics that alter quantal release (Pozzo-Miller et al., 1999; Tyler et al., 2002). Interestingly, presynaptic $\alpha 7$ -nAChRs enhance neurotransmitter release and are known to do so by elevating terminal Ca^{2+} levels (Gray et al., 1996; Coggan et al., 1997), possibly through Ca^{2+} -induced Ca^{2+} -release (CICR) recently shown to increase EPSC frequency (Sharma and Vijayaraghavan, 2003). Because CG neurons in culture express Ca^{2+} -permeable $\alpha 7$ -nAChRs on neurite tips (Pugh and Berg, 1994), and activation of CICR markedly enhances sEPSC frequency (M. Chen and J. Margiotta, unpublished observations), we wondered if upregulation of presynaptic $\alpha 7$ -nAChRs might underlie the ability of BDNF to increase sEPSC

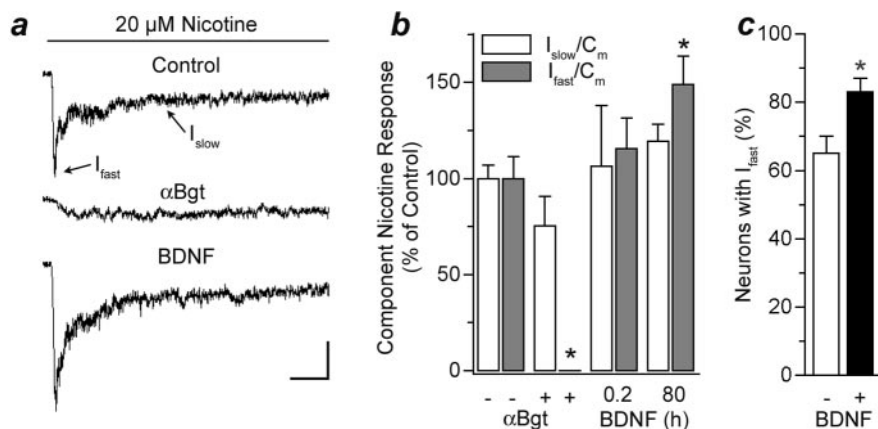


Figure 6. BDNF enhances $\alpha 7$ -nAChR currents. CG neurons were held at -70 mV, and whole-cell nAChR currents were induced by pressure microperfusion with $20 \mu M$ nicotine (2 sec, 10 – 15 psi). *a*, Example records showing that $\alpha 7$ -nAChRs mediate the initial fast current component (I_{fast} , top trace) because this component was absent in neurons incubated with α -Bgt (60 nM, 30 min, middle trace). I_{fast} was enhanced after treating CG neuron cultures with BDNF (4 d, 50 ng/ml, bottom trace). Calibration: 100 msec, 200 pA. *b*, Summary of α -Bgt and BDNF effects on $\alpha 7$ -nAChR currents. Left, I_{fast} normalized for membrane capacitance (I_{fast}/C_m , black bars) was absent, but I_{slow}/C_m reduced only slightly (white bars) in cultures treated with α -Bgt (+; $N = 8$) relative to control neurons tested in parallel (–; $N = 12$). Right, In neurons with detectable I_{fast} , 3–4 d BDNF treatment significantly increased I_{fast}/C_m by $49 \pm 14\%$ ($N = 63$) relative to untreated controls tested in parallel ($N = 52$). In the same records, the $\alpha 3^*$ -nAChR-mediated current component (I_{slow}/C_m) was not significantly different ($p = 0.08$) for control and BDNF-treated neurons ($N = 80$ and 76 , respectively). Shorter treatment times of 10 – 30 min (0.2 hr; $N = 9$) or 16 – 24 hr (data not shown) failed to detectably alter either fast, $\alpha 7$ -, or slow, $\alpha 3^*$ -nAChR mediated currents. *c*, Three or four day exposure to BDNF increased the proportion of neurons with detectable I_{fast} from $65 \pm 5\%$ ($N = 80$) in control cultures to $83 \pm 4\%$ ($N = 76$) in treated cultures. Asterisks indicate a significant difference from untreated controls tested in parallel ($p < 0.05$).

frequency. This hypothesis predicts that BDNF applied for 10 – 30 min should not increase sEPSC frequency because brief exposures were insufficient to increase surface $\alpha 7$ -nAChR levels or somatic $\alpha 7$ -nAChR currents (Figs. 5, 6). In accord with results from other systems (McAllister et al., 1999; Schneider and Poo, 2000; Poo, 2001) however, brief exposure to BDNF induced a significant, K252a-sensitive increase in sEPSC frequency (Fig. 7C, left), thereby demonstrating an expected trk dependence, but arguing against a requirement for rapid $\alpha 7$ -nAChR modulation. Because $\alpha 7$ -nAChRs at somatic and presynaptic sites could differ in their acute responsiveness to BDNF, we devised a more direct test, blocking $\alpha 7$ -nAChRs with α -Bgt and comparing sEPSCs in cultures treated with or without coapplied BDNF. Even with α -Bgt present to block $\alpha 7$ -nAChRs, however, BDNF applied for 16 – 24 hr was still able to increase sEPSC frequency (Fig. 7C, right), with all events now displaying slow decay kinetics indicative of $\alpha 3^*$ -nAChRs. These results indicate that brief- (10 – 30 min) and intermediate-duration exposures to BDNF (16 – 24 hr) can increase sEPSC frequency and do so without a requirement for $\alpha 7$ -nAChRs. Nevertheless, $\alpha 7$ -nAChRs are strongly implicated in long-term synaptic regulation (Role and Berg, 1996; MacDermott et al., 1999; Liu et al., 2001; Kawai et al., 2002). Thus, because 4 – 5 d BDNF treatments also increased sEPSC frequency and were required to detect significant changes in $\alpha 7$ -nAChRs, we cannot exclude the possibility that chronic neurotrophin exposure sustains the long-term function of neuronal nicotinic synapses in ways that somehow depend on $\alpha 7$ -nAChRs.

Discussion

Detection of $trkB$ and BDNF

We have shown that $trkB$ and BDNF-like proteins are present in the chick CG, a model parasympathetic system, where both $trks$ and neurotrophins were presumed irrelevant. No other studies have attempted to detect $trkB$ protein in CG, however, previous Northern and RNase protection analyses failed to detect $trkB$

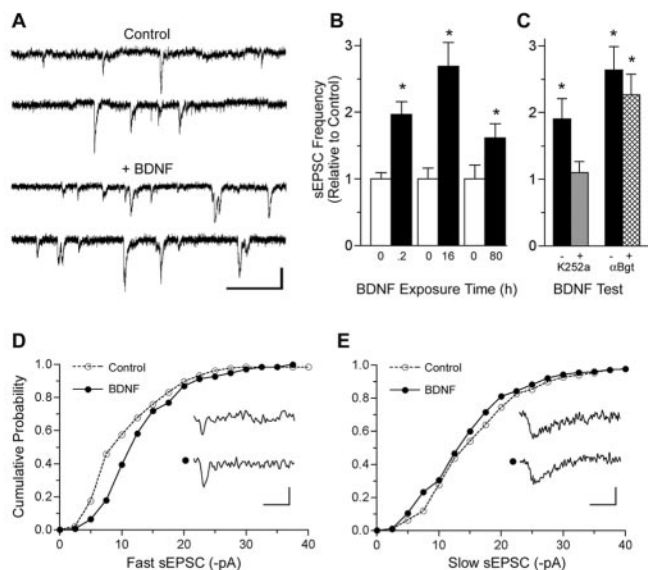


Figure 7. BDNF enhances function at nicotinic synapses. *A*, Example records of sEPSCs obtained on day 5 from control and BDNF-treated (16 hr, 50 ng/ml BDNF) CG neurons from the same cultures. Calibration: 25 pA, 100 msec. *B*, BDNF treatment effects on sEPSC frequency. Values indicate mean (\pm SEM) sEPSC frequency in neurons assessed on day 4 or day 5 after exposure to BDNF (black bars) for 10–30 min (0.2 hr; $N = 71$), 16–24 hr (16 hr; $N = 46$), or 4–5 d (80 hr; $N = 29$) relative to that for control neurons not exposed to BDNF (white bars; $N = 74, 37, 21$, respectively) tested in parallel. *C*, Tests for trkB and $\alpha 7$ -nAChR involvement in BDNF-enhanced synaptic function. The ability of BDNF to increase sEPSC frequency (black bars) is abolished by 10–30 min coincubation with 200 nM K252a to block trks (gray bar, left) but unaffected by 16–24 hr coincubation with α -Bgt to block $\alpha 7$ -nAChRs (hatched bar, right). In both *B* and *C*, asterisks indicate a significant increase sEPSC frequency for the indicated conditions compared with untreated control cultures from the same platings ($p < 0.05$). *D*, *E*, Summary of BDNF treatment effects on $\alpha 7$ - and $\alpha 3^*$ -nAChR-mediated sEPSC amplitudes. Cumulative amplitude distribution histograms are shown for fast-decaying sEPSCs mediated by $\alpha 7$ -nAChRs (*D*) and slow-decaying sEPSCs mediated by $\alpha 3^*$ -nAChRs (*E*) in control neurons (open circles) and in neurons treated with 50 ng/ml BDNF for 16–24 hr (filled circles). BDNF treatment resulted in a significant shift in fast sEPSC amplitudes (*D*) from a median of -9.6 pA ($N = 137$) for control neurons to -12.7 pA ($N = 139$) for BDNF-treated neurons ($p < 0.0004$; Mann–Whitney U and Kolmogorov–Smirnov tests). In contrast, slow sEPSC amplitudes (*E*) were unaffected by the treatment, with median amplitudes of -15.0 pA ($N = 220$) and -14.5 pA ($N = 180$) for control and BDNF-treated neurons, respectively ($p > 0.06$, same tests). Insets show examples of fast and slow sEPSCs with amplitudes close to the median values for control (top) and BDNF treatment (bottom, with dot) conditions. Calibration: 10 pA, 2 msec.

mRNA (Dechant et al., 1993; Hallbook et al., 1995), possibly because of the lower sensitivity of these assays compared with RT-PCR. Standard criteria for trkB primer (Garner and Large, 1994; Garner et al., 1996) and antiserum specificity (von Bartheld et al., 1996), as well as controls involving primer and primary antiserum omission (this study) support our detection of trkB mRNA in CG, and cell surface trkB protein on CG neurons. In addition, the observations that BDNF elicits trkB-dependent signaling leading to CREB activation and trkB-dependent changes in $\alpha 7$ -nAChRs and synaptic function further indicate that CG neurons express functional trkB. Because the trkB antiserum used recognizes an extracellular epitope (von Bartheld et al., 1996), and PCR amplifications using F/R_K primers revealed the presence of isoforms lacking an intracellular kinase domain, however, some trkB immunoreactivity may represent truncated receptor. Whereas the role of kinase-deficient trkB isoforms is poorly understood, the notion they are expressed on CG neurons is strengthened by the observation that in 50% of neurons BDNF application failed to induce p-CREB, a process expected to require trkB kinase activity (Finkbeiner et al., 1997; Huang and Reichardt,

2003). In such cases, full-length trkB receptors may still be present but either rendered functionally incompetent or expressed at levels insufficient to activate CREB because truncated trkB isoforms have been reported to inhibit both the function and expression of full-length receptors (for review, see Huang and Reichardt, 2003).

BDNF–trkB signaling upregulates $\alpha 7$ -nAChRs

BDNF treatment for 4–5 d induced trkB-dependent increases in $\alpha 7$ subunit mRNA and surface $\alpha 7$ -nAChRs, and enhanced $\alpha 7$ -nAChR-mediated whole-cell currents, all without changing levels of $\alpha 3$ subunit mRNA or $\alpha 3^*$ -nAChR-mediated currents. Similarly, NGF has been shown to selectively increase expression of $\alpha 7$ - over $\alpha 3$ -nAChR subunit mRNA in sympathetic neuron-like PC12 cells (Takahashi et al., 1999; but see Henderson et al., 1994). Although alterations in receptor turnover rates and mRNA stability may contribute, a straightforward interpretation of our results is that BDNF activation of trkB leads to increased $\alpha 7$ -nAChR subunit transcription and protein synthesis, thereby increasing levels of assembled cell-surface receptor. One way BDNF–trkB signaling may influence $\alpha 7$ -nAChR subunit transcription is through activation of transcription factors including not only CREB (Finkbeiner et al., 1997), but also AP-1, or NF- κ B, which like CREB are reported to be stimulated by BDNF–trkB signaling (Gaiddon et al., 1996; Lipsky et al., 2001). CRE binding sites are present in promoter-containing regions of human and bovine $\alpha 7$ -nAChR subunit genes, although in the chicken gene a 1298 bp 5' segment with a basal promoter at -406 to -230 was previously reported to lack a strong consensus CRE binding site (Matter-Sadzinski et al., 1992; Gault et al., 1998). Within this same 5' region, however, a new search of two transcription factor databases [Transfac (<http://www.gene-regulation.com/pub/databases.html>); Heinemeyer et al., 1998) and MatInspector (http://www.genomatix.de/free_login.htm)] did reveal a potential (–) strand CRE binding site ($T_{-1060}GACcTAA_{-1067}$) upstream from the basal promoter. Potential binding sites for AP-1 ($T_{-715}TcACTCAG_{-708}$) and NF- κ B ($G_{-176}GGGgcTCCC_{-167}$) were also predicted in the 5'-flanking and basal promoter regions, respectively. These considerations suggest that BDNF–trkB signaling can regulate the chicken $\alpha 7$ -nAChR subunit gene via CRE, AP-1, or NF- κ B. Without experimental data, however, it is difficult to judge the significance of these transcription factors as direct regulators. Here, it should be noted that binding sites for Egr-1, Sp1, and Sp3, transcription factors not associated with BDNF–trkB signaling, are thought to regulate the activity of the rat $\alpha 7$ -nAChR promoter (Nagavarapu et al., 2001).

BDNF increases activity at nicotinic synapses on CG neurons

BDNF increased sEPSC frequency after acute (10–30 min), intermediate (16–24 hr), or long-term (4–5 d) treatments. The increased sEPSC frequency after acute BDNF exposure depended on trkB activation and resembled that seen at other peripheral and central synapses, where enhanced transmitter release from presynaptic terminals is implicated (McAllister et al., 1999; Schneider and Poo, 2000; Poo, 2001). The basis of the acute synaptic effects seen here and in these other systems is unknown, but seems likely to reflect BDNF actions on Ca^{2+} (Berninger et al., 1993; Stoop and Poo, 1996; Li et al., 1998) and vesicular dynamics (Pozzo-Miller et al., 1999; Tyler et al., 2002) in presynaptic terminals that enhance neurotransmission reliability. The compelling possibility that Ca^{2+} -permeable, presynaptic $\alpha 7$ -nAChRs underlie these effects (Gray et al., 1996; Coggan et al., 1997; Sharma and Vijayaraghavan, 2003) is unlikely, however, because

acute BDNF exposure failed to modulate somatic $\alpha 7$ -nAChR currents and, more telling, because coincubation with α -Bgt failed to block the ability of BDNF to increase sEPSC frequency. Also unlikely are general effects on membrane excitability as seen for PC12 cells (Rudy et al., 1987; Lesser et al., 1997) because BDNF treatments failed to detectably change the amplitude or voltage sensitivity of somatic Na^+ or Ca^{2+} currents. In addition to increasing overall sEPSC frequency threefold, 16–24 hr BDNF treatments specifically increased the amplitude of $\alpha 7$ -nAChR-mediated sEPSCs. Although we cannot exclude increased quantal release at presynaptic nerve terminals that contact only $\alpha 7$ -nAChR clusters, this effect seems more likely to be postsynaptic in origin. Unlike currents induced by rapid nicotine microperfusion, which represent nAChR function integrated over the entire soma and report only a nominal increase, sEPSCs are focal events, and an increase in their amplitude would be expected even after adding a few functional receptors in the postsynaptic membrane. The increase in $\alpha 7$ -nAChR-mediated sEPSC amplitudes agrees well with the increased postsynaptic $\alpha 7$ -nAChR clusters previously observed in hippocampal neuron cultures (Kawai et al., 2002) and with the nominal increase in [^{125}I]- α -Bgt binding seen here after 16–24 hr exposure to BDNF and the significant increase seen after 5 d. The elevated sEPSC amplitude could reflect increased $\alpha 7$ -nAChR synthesis, and/or preferential insertion at existing postsynaptic sites, but we cannot presently distinguish between these possibilities.

Long-term synaptic enhancement

Our findings indicate that acute- and intermediate-term BDNF treatments increased synaptic activity without a requirement for $\alpha 7$ -nAChRs. Chronic (4–5 d) BDNF treatments continued to enhance synaptic activity, however, and, in parallel, significantly increased $\alpha 7$ -nAChR levels and whole-cell currents. $\alpha 7$ -nAChRs have been linked to activity-dependent neurite outgrowth and other developmental processes (for review, see Margiotta and Pugh, 2004) that may normally help ensure appropriate synaptogenesis or sustain existing functional synapses once formed (for review, see Role and Berg, 1996; Broide and Leslie, 1999; Jones et al., 1999). In addition, BDNF has been shown to have potent long-term effects on synaptic development and maintenance in other systems (for review, see Poo, 2001). Thus, although further experiments are needed, it remains possible that, in contrast to short- and intermediate-term $\alpha 7$ -nAChR-independent effects, the ability of BDNF signaling to sustain long-term synaptic function is related somehow to coincident regulation of $\alpha 7$ -nAChRs.

In vivo relevance?

Our results do not directly address the relevance of signals generated by BDNF through trkB for CG neurons *in vivo*. That targets and target-derived factors influence the survival, growth, and differentiation of input neurons, however, has been recognized for decades (Berg, 1984; Levi-Montalcini, 1987). Specific to this report, previous studies demonstrated that synapses on CG neurons undergo patterned maturation between E8 and E18 and that normal neuron survival and ganglionic transmission require connection to the intraocular muscle targets (Landmesser and Pilar, 1974a,b). More recent experiments indicate that $\alpha 7$ -subunit mRNA and $\alpha 7$ -nAChR protein and currents all increase during the same developmental period (Corriveau and Berg, 1993; Blumenthal et al., 1999) and that severing peripheral target connections reduces levels of $\alpha 7$ -mRNA and protein (Brumwell et al., 2002). Given the importance of target connections in sustaining $\alpha 7$ -nAChRs and synapses, the presence of BDNF-like

protein in the iris target suggests it may influence synaptic properties of CG neurons during development *in vivo*. The precise spatial and temporal patterns of BDNF and trkB expression still need to be determined. Our PCR and ELISA results suggest, however, that BDNF is both synthesized within the CG, perhaps by the neurons themselves, and transported to the ganglion from the iris muscle. Such local BDNF expression and retrograde transport from the target are consistent with the arrangement in spinal cord (Koliastzos et al., 1993) and would concentrate BDNF in the iris and CG relative to its levels in eye extract, as was observed here. In addition to BDNF, NGF and NT-3 signaling may also be important in the CG system because recent findings indicate mRNAs for trkA and trkC are expressed in the ganglion, and CG neurons express trkA and trkC immunoreactivity (Dittus et al., 2002). Experiments are in progress to reassess the ability of NGF, NT-3, and BDNF to promote CG neuronal survival, and to identify their respective roles in regulating nAChR expression and nicotinic synaptic differentiation on these parasympathetic neurons.

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