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BDNF–TrkB signaling in striatopallidal neurons controls inhibition of locomotor behavior

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The physiology of brain-derived neurotrophic factor signaling in enkephalinergic striatopallidal neurons is poorly understood. Changes in cortical Bdnf expression levels, and/or impairment in brain-derived neurotrophic factor anterograde transport induced by mutant huntingtin (mHdh) are believed to cause striatopallidal neuron vulnerability in early-stage Huntington’s disease. Although several studies have confirmed a link between altered cortical brain-derived neurotrophic factor signaling and striatal vulnerability, it is not known whether the effects are mediated via the brain-derived neurotrophic factor receptor TrkB, and whether they are direct or indirect. Using a novel genetic mouse model, here, we show that selective removal of brain-derived neurotrophic factor–TrkB signaling from enkephalinergic striatal targets unexpectedly leads to spontaneous and drug-induced hyperlocomotion. This is associated with dopamine D2 receptor-dependent increased striatal protein kinase C and MAP kinase activation, resulting in altered intrinsic activation of striatal enkephalinergic neurons. Therefore, brain-derived neurotrophic factor/TrkB signaling in striatopallidal neurons controls inhibition of locomotor behavior by modulating neuronal activity in response to excitatory input through the protein kinase C/MAP kinase pathway.

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Distinct basal ganglia (BG) circuits involved in motor control (direct and indirect pathways), originate in the dorsal striatum. These receive convergent excitatory afferents from the cortex and thalamus. The direct striatogniral pathway activity induces disinhibition of excitatory thalamocortical projections and facilitation of movement. The indirect striatopallidal pathway induces inhibition of thalamocortical projection neurons, which decrease cortical premotor drive and inhibit movement. Preferential loss of the indirect pathway medium-sized spiny neurons (MSNs) is thought to decrease the amount of inhibitory control over unwanted movements, leading to chorea and hyperkinesia typically associated with define HD: Huntington’s disease (HD) (ref. 3). HD is an autosomal dominant and progressive neurodegenerative disorder characterized by severe deficits in motor function, cognitive and psychiatric changes. This disorder is correlated with an expansion of a CAG repeat at the N-terminal region of the huntingtin gene.

Although the exact mechanism by which mHdh causes neuronal dysfunction is not yet established, it is clear that regions of the brain such as the striatum are the first to degenerate.

Previous studies at early stages of the disease demonstrate that striatal enkephalinergic (ENK)-positive GABAergic medium spiny neurons projecting to the external part of the globus pallidus are more vulnerable than the striatal MSN subset expressing substance P, an effect also found in HD mouse models. These neurons, also known as striatopallidal neurons, coexpress the dopamine receptor 2 (D2R). Their degeneration in HD ultimately results in the loss of inhibitory input to the thalamus, enhancing involuntary movements (chorea). The mechanisms that drive selective striatal cell loss are currently unclear and still extensively investigated. A recent hypothesis suggests that mHdh drives a progressive decline in striatal trophic support that ultimately leads to MSNs degeneration, initially of ENK−MSNs. The neurotrophic brain-derived neurotrophic factor (BDNF) appears to be indispensable for striatal cell long-term survival and maintenance. Cortical neurons are the main source of striatal BDNF, which is delivered by anterograde transport through the corticostratal afferents. Mutant Hdh impairs BDNF promoter II transcriptional activity through the cytoplasmic retention of the neuron-specific transcriptional repressor REST, and can also interfere with BDNF anterograde transport from cortical neurons to striatal targets. Therefore, reduction in BDNF supply to the striatum jeopardizes the long-term survival and morphology of striatal MSNs, contributing ultimately to striatal degeneration in HD. This is consistent with the reduced cortical and striatal BDNF levels observed in both HD patients and mouse models. Forebrain-specific Bdnf mutants recapitulate many behavioral and anatomical abnormalities seen in mouse models of HD. Similarly, mice carrying mHdh transgene, with genetic reduction of BDNF levels, exhibit advanced onset and increased severity of motor dysfunction, resulting from the degeneration of striatal ENK−MSNs.

Despite extensive evidence associating cortical Bdnf levels with striatal vulnerability, there is no demonstration that these effects are mediated via TrkB. This is mainly due to any ablation or alteration of Bdnf levels, either globally or in a region-specific manner, affecting cortical physiology, and therefore confounding the action of Bdnf anterogradely transported from the cortex to other regions of the brain, such as its striatal targets.

To address the physiological role of BDNF/TrkB signaling in striatopallidal neurons, we have selectively deleted the high-affinity BDNF receptor, TrkB, from ENK+ MSNs. Surprisingly, this manipulation led to spontaneous and drug-induced hyperlocomotion associated with increased D2R-dependent MAPK/PKC phosphorylation and reduced striatopallidal activation, without obvious effects on motor coordination and gait parameters, or long-term survival and morphological defects of ENK+ MSNs. We therefore demonstrate that BDNF−TrkB signaling in striatal ENK+ MSNs contributes to the inhibitory control of locomotor behavior exerted by the indirect pathway.

Results

Generation of Enkephalin-specific TrkB knockout mice. To address the physiological relevance of BDNF−TrkB signaling in the ENK+ MSNs, we conditionally deleted TrkB from these neurons by generating a BAC transgenic mouse line carrying Cre-recombinase under the control of the pre-proenkephalin promoter (BAC−Penk−CreEYFP+) (Supplementary Fig. S1a). We studied the recombination pattern of the BAC−Penk−Cre line using two reporter lines (ref. 18,19), which recapitulated mostly the expression of the pre-proenkephalin gene reported in literature. Namely, recombination was detected in particular in the caudate-putamen, the nucleus accumbens, scattered cells in cortical layer II and V–VI (Fig. 1a), and the granular layer of the olfactory bulb and cerebellum. The cellular specificity of Cre-recombination was examined by double immunofluorescence using antibodies (Abs) to YFP and to markers either specific for ENK+ MSNs, showing Cre-mediated EYFP expression occurring in 98% of the ENK+ MSNs (Fig. 1b), or specific for striatal interneurons, showing no colocalisation (Fig. 1c). The relative abundance of EYFP+ MSNs within the striatum was estimated using COUPtFI-interacting protein 2 (CTIP2) (ref. 21). We found 40–50% EYFP+ MSNs (Supplementary Fig. S1b), consistent with the literature. However, 4% of the total EYFP+ cells do not express enkephalin. Possibly the enkephalin-promoter was active at some point during cell differentiation or expression of enkephalin is below the minimum detectable level. This novel Cre line was crossed to a Trkb floxed line to generate TrkbPenk-KO mice. E(Y)GFP immunoreactivity was not detected in the developing mouse brain before E12.5, consistent with enkephalin expression in differentiating multipotent progenitors but not in earlier progenitors, unlike Dlx5 (refs. 20,24) (Fig. 1d–g). Western blot analysis of adult mouse brain lysate further confirmed specific striatal TrkB reduction in TrkbPenk-KO mice compared with Cre-negative TrkbPenk-WT littermates (Trkb[+/–],Penk+/+) (Fig. 1h). TrkB deletion was also validated by in situ hybridization combined with immunofluorescence double staining (Trkb/EYFP), (Supplementary Fig. S1c). These data confirmed the specificity and successful inactivation of Trkb in ENK+ neurons.

TrkB deletion in ENK MSNs results in hyperlocomotion. Recent work, by toxin ablation or functional disruption of the striatopallidal D2R neurons, has provided direct evidence that these neurons regulate inhibitory functions on locomotor activity. To understand if BDNF−TrkB signaling would be relevant to this function, we tested locomotor activity of TrkbPenk-KO mice and age-matched TrkbPenk-WT in the open field (OF) apparatus (Fig. 2a,b). We found a sharp significant increase in total distance traveled in TrkbPenk-KO mice compared with that in TrkbPenk-WT littermates by 9 M of age until later in life (Fig. 2a). The TrkbPenk-KO spontaneous hyperlocomotion observed in the OF cannot be due to increased anxiety or general activity of the mice, as analysis of time spent in the border compared with that in the center of the OF apparatus, and analysis of home-cage activity over 3 days did not reveal significant differences between genotypes (Supplementary Fig. S2a,b). Mice carrying only the Cre transgene tested at 9 M of age in an OF did not show significant changes compared with wild-type mice (P = 0.897, unpaired Student’s t-test), confirming that the spontaneous increased locomotion observed in the TrkbPenk-KO mice was due to the specific deletion of Trkb in ENK+ MSNs.
Drugs of abuse, like cocaine, are able to induce hyperlocomotion, an action that is largely mediated through the striatum. As the hyperlocomotion phenotype observed in the OF is age-dependent, it is likely that young mice already have a mild defect in motor control that is not detectable in a paradigm of spontaneous locomotion. Therefore, we tested the response of young adult (3 M old) TrkbPENK-KO and TrkbPENK-WT mice in the OF when acutely challenged with cocaine. TrkbPENK-KO mice showed significantly increased distance traveled compared with TrkbPENK-WT littermates upon cocaine treatment (Fig. 2b), demonstrating that lack of BDNF–TrkB signaling in ENK+ MSNs enhances cocaine-induced hyperlocomotion. This confirms that striatopallidal neurons lacking BDNF–TrkB signaling lose inhibitory control over locomotor activity, and that young mutants show a phenotype similar to the aged one if challenged.  

To assess if lack of TrkB signaling in ENK+ MSNs would affect motor coordination and overall motor function, we first examined TrkbPENK-KO mice and TrkbPENK-WT littermates for clasping, and found normal hind-paw clasping in both genotypes at all stages analyzed. We then assessed motor coordination performance of TrkbPENK-KO using the rotarod test. Twenty-five TrkbPENK-WT and 25 TrkbPENK-KO mice from 2 to 14 M of age were tested. Mutants showed similar performance compared with control littermates (Fig. 2c), indicating that lack of BDNF–TrkB signaling in striatopallidal ENK+ MSNs does not affect general performance in motor coordination.  

Subtle gait differences in TrkbPENK-KO mice. As neither the OF nor the rotarod test would allow for objective assessment and quantitative measures of fine motor movements, we then used the

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**Figure 1 | Generation of Trkb-enkephalin knockout mice.** (a) Adult brain image of BAC–Penk–CreEgfp/::Z/EG–egfptg/+ mouse stained with G(Y)FP Abs revealing Cre-mediated GFP expression in specific brain areas, the caudate-putamen (CPu), the nucleus accumbens (NAc), scattered cells in cortical layer II and V–VI (CX). (b) Confocal images showing Cre-mediated EYFP expression in ENK striatal cells, (c) and absence of EYFP expression in striatal interneurons (arrows) identified by specific markers. (d,e) Dlx5 in situ expression at E11.5 in the striato-pallidal neural epithelium (strial-ne, pallidal-ne), and absence of Penk1 expression in the same regions at this stage (images are from the Allen Mouse Brain Atlas20). (f) E11 BAC–Penk–CreEgfp/::Z/EG–egfptg/+ transversal brain section with no GFP staining in the striato-pallidal neural epithelium. (g) PCR analysis of Trkb recombination in brains carrying the BAC–Penk–Cre transgene at E12.5. Primers were specific for un-recombined (Nr) and recombined (r) Trkblx allele; each embryo carried a Trkblx allele and TrkbWT allele. Recombination occurred only in presence of the Cre transgene (embryos 1–3). (h) Protein lysates from different brain regions of adult TrkbPENK mice either in presence (+) or in absence (−) of the Cre transgene were analyzed for TrkB protein levels. Blots were probed with Abs to p145-TrkB. Striatal TrkB levels were 50% reduced, but unchanged in the cortex. Blots were reprobed with Abs to TUJ.1 to control for loading and quantification of fold changes (values are mean ± s.d.). ChAT, choline acetyltransferase. Scale bars: a, 500 μm; b,c, 50 μm; f, 100 μm. See also Supplementary Fig. S1.
Figure 2 | Enhanced spontaneous and cocaine-induced locomotion in TrkbPENK-KO mice. (a) Spontaneous locomotor measures obtained for control and mutant mice tested as independent groups at different stages (TrkbPENK-WT, n = 9, 9, 7; TrkbPENK-KO, n = 7, 9, 7, for the three groups, respectively). Mice were individually placed into an OF arena for 30 min and the distance traveled was recorded. While no difference was observed up to 8 M of age, a sharp significant increase in total distance traveled was found in TrkbPENK-KO mice compared with that in TrkbPENK-WT littermates by 9 M of age until later in life (two-way ANOVA, $F_{(2,42)} = 8.964; P = 0.0046$, main effect of genotype; $F_{(2,42)} = 6.406; P = 0.0037$, interaction between genotype and age). TrkbPENK-KO versus TrkbPENK-WT, $P = 0.0001$); values are means ± s.e.m. (b) Cocaine-induced hyperlocomotion. After 2 days of habituation to the apparatus, mice (TrkbPENK-WT, n = 11; TrkbPENK-KO, n = 10) were i.p.-injected with either vehicle on day 0, showing no difference between mutants and controls (one-way ANOVA, $F_{(1,19)} = 0.386; P = 0.5417$), or with 10 mg kg$^{-1}$ cocaine for 3 consecutive days, immediately before being placed into the arena where locomotor activity was recorded for 30 min (one-way ANOVA, $F_{(1,19)} = 4.797; *P = 0.0412$). Cocaine values are mean across the 3 days ± s.e.m. (c) TrkbPENK-KO and TrkbPENK-WT mice were tested for motor coordination as independent groups at different stages (TrkbPENK-WT, n = 9, 8, 8, and TrkbPENK-KO mice, n = 7, 11, 7, for the three groups, respectively). On day 1, they were habituated to the rotarod apparatus and subjected to three sessions of 3 min at 4 r.p.m. constant speed, separated by 15-min rest intervals. The next day they were placed on the rod accelerating from 4 to 40 r.p.m for three consecutive sessions (separated by 5-min intervals). Latency of each animal to fall from the rod was scored. The performance was not significantly different between mutants and controls (two-way ANOVA, $F_{(2,44)} = 0.298; P = 0.5878$ main effect of genotype; $F_{(2,44)} = 0.874; P = 0.4244$, interaction between genotype and age; $F_{(2,44)} = 9.702; P = 0.0003$ effect of age). Values are means ± s.e.m.

CatWalk test; a new method of quantitative gait analysis.28,29 This method allows easy quantification of a large number of locomotion parameters during walkway crossing (Supplementary Methods). This test has also been validated to study motor impairment associated with different neurological motor disorders, including an HD model.30 Analysis of mice at 5 M of age, before mutants develop spontaneous hyperlocomotion, revealed no significant differences between genotypes in activity-related parameters (Supplementary Table S1). All parameters related to single paws, previously reported to be compromised in a transgenic rat model of HD,30 were normal in TrkbPENK-KO mice compared with that in controls (Supplementary Fig. S3a–d, Supplementary Table S2). However, a smaller print width for the left hind paw (LH) was found in mutant mice compared with that in controls (Supplementary Fig. S3e, Supplementary Table S2).

There was no difference between genotypes in front-paw parameters (Supplementary Table S2).

As for the coordination-related gait parameters, both mutant and control mice showed normal interlimb coordination based on the analysis of the step sequence patterns, and normal phase dispersion (Supplementary Table S3, Supplementary Fig. S4). However, some differences between genotypes were found in the time relationship between footfalls of two different paws (couplings), with slightly decreased timely coordination between ipsilateral (RH→RF) and girdle paws (RH→LH), but slightly increased coordination between diagonal paws (RF→LH) in TrkbPENK-KO mice compared with that in controls (Supplementary Fig. S3f–g).

Mice examined in the CatWalk test were additionally analyzed in the inverted screen test, which requires motor coordination...
and muscle strength, and in the weight-lifting test to further ascertain their strength/forepaw grasping. No significant differences were scored between the two genotypes for either test (Supplementary Fig. S5). Overall, these data reveal subtle changes in gait parameters in TrkbPENK-KO mice, which are evident before the appearance of spontaneous altered locomotion.

Reduced ENK expression in striatal MSNs upon Trkb deletion.

To rule out that the phenotype observed in locomotor activity was not the result of loss or morphological alteration of the ENK+ MSNs but rather a primary effect of the lack of TrkB signaling in ENK+ MSNs, we first examined the relative abundance of striatal MSNs in aged TrkbPENK-KO mice and TrkbPENK-WT littermates. Counts of neurons labeled with Abs against an MSN-specific marker, like dopamine- and cAMP-regulated phosphoprotein of 32 kDa (DARPP-32), did not reveal loss of striatal MSNs in TrkbPENK-KO mice compared with TrkbPENK-WT littermates (Fig. 3a). Stereological analysis of histological sections revealed normal striatal volume in TrkbPENK-KO compared with that in TrkbPENK-WT mice at 12 M of age (Supplementary Fig. S6a,b). Computer-based morphometric analysis of Golgi-impregnated

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**Figure 3 | Reduced striatal enkephalin expression in TrkbPENK-KO mice in the absence of striatal loss.** (a) Striatal images from 8-M-old mice immunostained for DARPP-32. DARPP-32 cell counts at 8 M (TrkbPENK-KO n = 3, versus TrkbPENK-WT n = 3, P = 0.741, unpaired Student’s t-test). (b) Striatal images of 8-M-old TrkbPENK-WT and TrkbPENK-KO mice showing enkephalin in situ hybridization. Counts of ENK+ MSNs at 8 M (TrkbPENK-KO n = 3, versus TrkbPENK-WT n = 3, *P = 0.036, unpaired Student’s t-test). (c) Confocal stacks of Golgi-impregnated MSNs from 12-M-old mice analyzed for spine density. (d) Sholl analysis of striatal MSNs dendritic complexity. For both panels c and d (n = 30, 10 randomly selected neurons per mouse/over three mice). One-way ANOVA, F(1,30) = 0.341; P = 0.5615. See also Supplementary Figs S6 and S7. (e) Representative western blot of 12-M-old striatal tissue lysates of TrkbPENK-WT (n = 3) and TrkbPENK-KO (n = 3) mice. Membranes were blotted with the indicated Abs and densitometry analysis was used for quantification of the proteins relative abundance (TrkbPENK-KO versus TrkbPENK-WT for Syn, P = 0.46; PSD-95, P = 0.15; Neuroligin-2, P = 0.09; GAD67, P = 0.40; DARPP-32, P = 0.21; D2R, P = 0.52; D1R, P = 0.33. P-values were generated by unpaired Student’s t-test). GAPDH was used to control for protein loading. Values are means ± s.e.m. See also Supplementary Fig. S8a. CPu, caudate-putamen. Scale bars: a-b, 50 μm. Syn, synaptophysin; D1R and D2R, dopamine receptor type 1 and 2.
striatal MSNs showed no difference in dendritic spine density (Fig. 3c, and Supplementary Fig. S6c) and thickness (Supplementary Fig. S6d) between 12-M old TrkbPENK-KO mice and TrkbPENK-WT litters. Sholl analysis further supported this observation, revealing no difference in dendritic arbor complexity (Fig. 3d).

Although these morphological data apply to the total MSN population, there was no evidence for a bimodal distribution, making unlikely the possibility that defects in ENK+ MSNs were masked by the striatonigral subpopulation. However, striatal enkephalin expression was reduced in TrkbPENK-KO mice in the absence of striatal MSN loss or DARPP-32 reduction (Fig. 3b,e, and Supplementary Fig. S6e). Further analysis of synaptic markers and dopamine receptors levels in the striatum revealed no significant changes between mutants and controls at 12-M of age (Fig. 3e). These results demonstrate that long-term survival and morphology of striatopallidal MSNs are not affected by BDNF–TrkB signaling-specific inactivation. In contrast, depletion of TrkB signaling affects striatal enkephalin expression and function of these neurons, suggesting that signaling activated by BDNF–TrkB is required in these cells to perform normal inhibitory function on locomotor activity.

**Intact corticostriatal synaptic properties in TrkbPENK-KO mice.** As alterations of the corticostriatal tract are known to affect MSN physiology and, therefore, locomotor behavior, we examined the electrophysiological properties of this tract in TrkbPENK-WT and TrkbPENK-KO mice. Basal glutamatergic synaptic transmission in slices containing the dorsolateral striatum of TrkbPENK-WT and TrkbPENK-KO mice was analyzed by extracellular field potential recordings. Input–output relationships were constructed by plotting the mean amplitude of field excitatory postsynaptic potential (fEPSP) against the increasing stimulation intensities. There was no significant difference in the input–output curves or in the amplitude of the fEPSPs between TrkbPENK-WT and mutant mice (Fig. 4a). To determine whether ablation of Trkb from striatopallidal neurons would affect neurotransmitter release from presynaptic terminals at corticostriatal synapses, we measured responses to paired-pulse stimulation at 40 and 60 ms interpulse intervals, whose ratio is considered as an index of release probability, a form of short-term synaptic plasticity. Paired-pulse responses had similar amplitude between TrkbPENK-WT and TrkbPENK-KO mice, and similar ratio values (Fig. 4b). We also examined another crucial parameter of basal glutamatergic transmission, namely, the spontaneous activity of MSNs. Interestingly, no significant differences in frequency or amplitude of the spontaneous postsynaptic currents were found in ENK+ MSNs between controls and mutants (Fig. 4c). Collectively, these findings indicate that basal excitatory synaptic transmission and short-term plasticity at the corticostriatal synapses are similar in the two genotypes. However, these data do not rule out the possibility that lack of BDNF/TrkB signaling in striatopallidal neurons would lead to other electrophysiological changes.

**TrkB signaling modulates striatopallidal MSN activation.** As Trkb deletion in ENK+ MSNs results in spontaneous and drug-induced hyperlocomotion, we next asked if TrkB signaling in these neurons would interfere with dopamine/D2R function that represses striatopallidal MSNs excitability in response to glutamatergic cortical input. Therefore, we combined precise mouse genetics with a pharmacological approach, and determined whether loss of Trkb specifically from D2R/ENK-positive MSNs would impair the response of these cells upon acute excitatory effect induced by D2R blockade. We performed D2 receptor blockade using a D2R-like antagonist (haloperidol)31, which triggers an acute excitatory effect of striatopallidal MSNs. This is followed by neuronal activation and c-Fos induction predominantly in D2R neurons31. Accordingly, TrkbPENK-KO and TrkbPENK-WT animals were injected either with 1 mg kg⁻¹ haloperidol or vehicle (DMSO), and the density of c-Fos-immunoreactive (c-Fos-ir) cells in the caudate-putamen was determined after 2 h by immunohistochemistry. The results showed a significant reduction in the density of haloperidol-induced c-Fos-ir cells in TrkbPENK-KO mice compared with that in TrkbPENK-WT (Fig. 5a,c). These data indicate that Trkb-depleted striatopallidal MSNs are not able to respond efficiently to the excitatory effect induced by D2R blockade, suggesting that TrkB signaling modulates activation of striatopallidal MSNs integrating cortical glutamatergic input.

To further support the observations obtained in the dorsal striatum, we also assessed the ability of haloperidol to activate D2R/ENK+ (PV−) MSNs in the LGP of TrkbPENK-KO and TrkbPENK-WT by similar analysis of c-Fos-ir expression in response to D2R blockade. These cells project to the striatum and form the pallidostriatral pathway32. The results obtained 2 h after haloperidol injection showed a significant reduction in the density of haloperidol-induced c-Fos-ir cells in TrkbPENK-KO mice compared with that in TrkbPENK-WT (Fig. 5b,d,e,f). These data indicate impaired feedback of PV-negative LGP cells to striatal neurons. This effect was similar to that observed in the striatopallidal MSNs of the indirect pathway, and indicates that TrkB signaling is required in these cells for their normal activation.

**Intact striatopallidal projections in the absence of TrkB.** To analyze if reduced activation of the striatopallidal neurons would affect targeting of striatopallidal projections in the LGP, we analyzed the density of GABAergic synapses on PV+ LGP neurons, which are part of the indirect pathway and receive direct striatopallidal projections33. Densities of pre- and postsynaptic GABAergic components were analyzed by immunostaining of the presynaptic enzymes glutamic acid decarboxylase 67 (GAD67) and the postsynaptic scaffolding protein gephyrin, respectively. The imaging at single-cell level of PV-expressing pallidal cells paired with 3D computer-based reconstruction of the immunostaining allowed detailed analysis of the GABAergic striatopallidal terminations. Quantification of either gephyrin- or GAD67-containing synapses on PV-labeled neurons in the LGP of haloperidol-treated TrkbPENK-WT and TrkbPENK-KO mice revealed no significant difference in the somatodendritic density of both GAD67 and gephyrin (Fig. 6a–d). These data indicate that absence of TrkB signaling does not affect targeting of striatopallidal projections and/or synaptic density on LGP PV+. TrkB modulates D2R-dependent striatal PKC and MAPK activation. It has been shown that D2 receptor agonists lead to MAPK phosphorylation in neurons through increased intracellular Ca²⁺ and PKC activity34. Striatal activation of ERK is necessary for the expression of D2 receptor-mediated locomotor hyperactivity35. Therefore, to provide a possible mechanistic insight into our novel findings, we have tested the effect of TrkB signaling depletion on the activation of these kinases in 13-M-old TrkbPENK-KO striatum, when mutants already show spontaneous hyperlocomotion, and in 5-M-old, a presymptomatic stage. We found significantly enhanced phosphorylation of both PKC and ERK1/2 in TrkbPENK-KO mice compared with that in TrkbPENK-WT at 13 M of age, but no significant difference in AKT or DARPP-32 (Thr34) phosphorylation (Fig. 7a,c). The enhanced phosphorylation of PKC and MAPKs was not evident in TrkbPENK-WT mice at 5 M of age, suggesting that aging is a critical factor (Fig. 7b,d). To
test whether the increased phosphorylation state of PKC
and ERK1/2 was D2R-dependent, we challenged 13-M-old
TrkbPENK-KO mice and TrkbPENK-WT with a more selective
antagonist of the D2R (L-741,626) (ref. 36). This treatment
blocked the increase in phosphorylation of PKC and ERK1/2 in
TrkbPENK-KO mice (Fig. 7a,c), indicating that increased PKC/MAPK
phosphorylation is the result of enhanced D2R-signaling in the
absence of TrkB signaling.

Discussion
The specific depletion of TrkB signaling from mouse striatopallidal
neurons, while maintaining cerebral cortex TrkB expression, has
allowed us to investigate the physiological role of BDNF–TrkB
signaling in ENK⁺ MSNs and to determine whether the known
effects of altered cortical BDNF signaling on striatal vulnerability
are mediated via TrkB. Here, we show that TrkB signaling is
required to control inhibition of locomotor behavior in ENK⁺
MSNs. Ablation of Trkb from striatopallidal MSNs results in
spontaneous and drug-induced hyperlocomotion associated with
reduced enkephalin expression and increased D2R-dependent
striatal PKC and MAPKs phosphorylation, leading to altered
response of ENK⁺ MSNs to glutamate-mediated excitation.
However, long-term survival and morphology of ENK⁺ MSNs
were not affected. Finally, these data identify a critical role for

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**Figure 4 | Functional integrity of the corticostriatal pathway in TrkbPENK-KO mice.** (a) Averaged fEPSP traces (upper panel) evoked by a stimulation of 12, 16, 22, 28, 32 and 36 V. Input–output best-fit curves (lower panel) revealed no significant difference between TrkbPENK-WT (n = 7 slices/4 mice) and mutant mice (n = 10 slices/4 mice) (P > 0.05, unpaired Student’s t-test). Values are means ± s.e.m. of fEPSP amplitude evoked by a given stimulation. (b) Histograms of group data showing similar paired-pulse ratio (PPR) of responses at 40 and 60 ms of interpulse intervals stimulation between TrkbPENK-WT (n = 8 slices/4 mice) and TrkbPENK-KO mice (n = 9 slices/4 mice), and similar ratio values (P > 0.05, unpaired Student’s t-test). Values are means ± s.e.m. of changes in the respective cell populations. (c) Graphs showing mean frequency and amplitude of the spontaneous postsynaptic currents (sEPSC) recorded from A2A-positive MSNs (TrkbPENK-KO, frequency: 2.67 ± 0.56 (n = 9/5 mice) versus TrkbPENK-WT: 2.94 ± 0.32 (n = 14/9 mice), P = 0.64, unpaired Student’s t-test; TrkbPENK-KO, amplitude: 13.3 ± 0.87 (n = 9/5 mice) versus TrkbPENK-WT: 13.8 ± 1.2 (n = 14/9 mice), P = 0.76, unpaired Student’s t-test. Values are means ± s.e.m.
TrkB signaling in locomotor behavior exerted by ENK striatopallidal neurons and demonstrate that these neurons are not dependent on BDNF–TrkB signaling for long-term survival and morphology in vivo.

BDNF is a major regulator of neuronal survival and morphology. One current main hypothesis suggests that a reduction in cortical BDNF and/or impairment in BDNF anterograde transport from cortex to striatal targets contribute to the degeneration of striatal neurons observed both in patients and in HD animal models. This suggestion, however, is based mainly on genetic models that compromise cerebral cortex physiology, and do not allow direct evaluation of the contribution of BDNF–TrkB signaling to striatal vulnerability. Therefore, to address the physiological role of BDNF–TrkB signaling in striatopallidal neurons, the most vulnerable neuronal subtype in HD, we have selectively deleted the high-affinity BDNF receptor, TrkB, from these neurons. Our data indicate that BDNF–TrkB signaling is required to control inhibition of locomotor behavior, but is not an essential factor for long-term survival and maintenance of ENK MSNs, and does not have a direct role in the striatal vulnerability observed upon cortical deletion of Bdnf. These contradictory roles for BDNF may be explained by its necessity for the maintenance of cortical neuron morphology. In particular, forebrain-specific conditional deletion of Bdnf leads to a reduction in cellular size and dendritic complexity of pyramidal neurons in cortical layer II–III at just 3 weeks of age. This, and possible alteration of other cortical layers, could compromise cortical input to striatal targets leading to the observed morphological changes in striatal MSNs.

Therefore, in contrast to the current belief, our data support the theory that complex alteration of the corticostriatal tract, and/or cortical cell-intrinsic properties ultimately contribute to striatal vulnerability. Indeed, dysfunction of the corticostriatal tract seems to occur during the development of the HD phenotype before any significant neuronal cell loss is apparent in the striatum. In our mouse model, we found functional integrity of the corticostriatal tract analyzed by electrophysiology. The procedure we have used (IPSP) has been shown to be sensitive and reliable for detecting alterations in the electrophysiological properties of MSN subpopulations. Unsurprisingly, disruption of TrkB signaling in ENK+ MSNs induces spontaneous hyperlocomotion, suggesting that TrkB in striatopallidal neurons contributes to the inhibitory control of locomotor behavior exerted by the indirect pathway. Notably ablation of D2R neurons in the entire striatum produced a bigger hyperlocomotor response than when ablation was limited to ventral striatum only. This is in agreement with the current hypothesis of the BG identifying striatopallidal MSNs as an inhibitory component of the circuit controlling motor activity. Particularly noteworthy is that conditional inactivation of TrkB

Figure 5 | Reduced neuronal activation in Trkb-depleted striatopallidal cells in response to D2R-blockade. (a–b) Representative images of CPu (a) and LGP (b) showing c-fos expression 2 h after systemic i.p. injection of 1 mg kg−1 haloperidol in 3-month-old TrkbPENK-WT and TrkbPENK-KO mice (n = 5, each genotype). In (b) tissues are co-labeled with an Ab against parvalbumin. These cells are the target of the striatopallidal MSNs and do not express D2R/ENK. Therefore, these cells are insensitive to haloperidol in contrast to the other cell type within the LGP that is D2R/ENK+. (c,d) Quantification of density of haloperidol-induced c-fos-ir cells in the CPu (c) (TrkbPENK-WT, 100 ± 8.81 n = 5, TrkbPENK-KO, 53.03 ± 9.40 n = 5, P = 0.006, unpaired Student’s t-test), and LGP (d) (TrkbPENK-WT, 100 ± 15.86 n = 5, TrkbPENK-KO, 46.52 ± 13.68 n = 5, P = 0.034, unpaired Student’s t-test); values are means ± s.d. (e,f) Schematic representation of TrkB signaling requirement in striatopallidal MSNs, and relative neuronal populations in the CPu and the LGP. Red and green cells in the dorsal striatum represent respectively striatopallidal (indirect pathway) and striatonigral (direct pathway) MSNs. Brown and orange cells in the LGP represent respectively PV+ and PV− neurons, respectively. Scale bars: a, 50 μm; b, 20 μm. See also Supplementary Methods.
signaling in striatopallidal MSNs leads to a reduction of enkephalin, one of the earliest morphological changes observed in HD, without affecting MSN viability. The decrease in enkephalin expression presumably reflects a reduction in neuronal activity of the striatopallidal efferents projecting to the external globus pallidus. This could affect the identity of MSN belonging to the indirect pathway, compromising their functionality and consequently causing an increase of spontaneous and cocaine-associated activity by disinhibition of the striatopallidal projections. Indeed, a similar phenotype occurs by conditional deletion of DARPP-32 in striatopallidal MSNs. This leads to increased basal and cocaine-induced locomotor activity in the presence of a compromised corticostriatal LTP without any cell loss. In our model, however, since cortical TrkB has been left mainly intact, the functional integrity of the corticostriatal pathway is not affected and the inhibitory function observed is therefore mainly confined to the striatopallidal pathway. Recent studies though, showed that mice carrying deletion of TrkB mediated by a less-restricted Cre line (D2R-Cre), have decreased locomotor activity upon cocaine administration without significant changes of c-Fos-ir in the dorsal striatum upon acute cocaine treatment. This inconsistency with our study may be due to D2R expression in both striatopallidal MSNs and striatal cholinergic interneurons, unlike enkephalin. Therefore, considering the modulatory activity of striatal cholinergic cells over the striatopallidal pathway, it is not surprising that the opposite result was obtained in the activation of striatopallidal neurons and, therefore, in the behavioral output.

It is evident from the classical BG model that normal locomotor behavior results from the balance between direct and indirect BG pathways. Recently, this model has also been supported by the use of optogenetics. Moreover, D2R activation reduces indirect pathway excitability specifically induced by cortical afferent activation and subsequent Ca\(^{2+}\) inflow. Diminished D2R or a D2R blockade leads to an excess of Ca\(^{2+}\) inflow, hence increasing indirect pathway activation. Essentially, D2R activation represses striatopallidal MSNs excitability by inhibiting calcium entry in these neurons through modulation of NMDAR and R-type Ca\(^{2+}\) channels. This effect is, in part, compensated by A2AR activation, which enhances Ca\(^{2+}\) entry. In our model, striatopallidal MSNs carrying TrkB depletion are not able to respond efficiently to the acute excitatory effect induced by D2R-like antagonist haloperidol (measured by c-Fos induction). The observed reduced enkephalin expression in these neurons also supports this finding. This suggests that TrkB signaling modulates activation of striatopallidal MSNs, integrating cortical glutamatergic input, most likely by enhancing Ca\(^{2+}\) entry. In addition, we found that major manifestation of phenotypes, such as hyperlocomotion, is noticeable with age or by challenges such as cocaine administration in young mutant animals. One possible explanation for this is that during aging there is a progressive decline in brain production of neurotrophic factors and their cognate receptors. In the BG of young animals, the presence of a significant amount of trophic factors may be able to compensate for the deficit caused by the absence of TrkB signaling. With aging, the availability of trophic factors declines, exacerbating the deficit caused by the absence of TrkB. Epigenetic changes occurring with aging could also explain the appearance of the phenotype only in older animals.
Finally, TrkB-specific deletion in ENK\(^+\) MSNs demonstrates that in physiological conditions, BDNF signaling is not required for the long-term survival and morphology of these striatal neurons. It does, however, affect enkephalin expression and neuronal activity resulting in altered locomotion. As reduced levels of BDNF in a diseased brain exacerbate the severity of motor dysfunction\(^5\), its replacement should maintain normal levels of BDNF in a diseased brain exacerbate the severity of neuronal activity resulting in altered locomotion. As reduced

**Methods**

**Ethics statement.** All animal procedures at the European Molecular Biology Laboratory (Mouse Biology Unit) conformed to National and International laws and policies (EEC Council Directive 86/609, OJ L 358, 1, 12 December 1987; NIH Guide for the Care and Use of Laboratory Animals, NIH Publication No. 85-23, 1985, revised in 1995); at the Centre for Neuroregeneration and at the Department of Pharmacology, animal procedures conformed to UK legislation (Scientific procedures) Act 1986, the University of Edinburgh and the University of Oxford ethical review committee policy, respectively.

**Immunohistochemistry.** Immunohistochemistry was performed as previously described\(^\text{23}\). Primary Abs were used at the following dilution: Ctip2 (1:1,000, ab18465, Abcam), DARPP-32 (1:500, AB1656, Chemicon), EGFP (1:500, ab290, Abcam), c-Fos (1:500, sc-52, Santa Cruz Biotechnology), enkephalin (1:200, RA14124, Neuroics), ChAt (1:1,000, ab14149, Chemicon), Calretinin (1:1,000, 7699/4, Swant), Somatostatin (1:200, ab5494, Chemicon), Parvalbumin (1:1,000, P3088, Sigma, rabbit polyclonal), GAD67 (1:2,000, Millipore) and gephyrin (1:500, Synaptic System). Alexa-labeled secondary Abs (1:1,000 Invitrogen), or biotin-conjugated anti-rabbit Ab (1:500 Jackson Lab). All fluorescent images were taken on LSM710 Meta confocal microscope (Carl Zeiss) mounting 40x/0.75 NA, and ×63/1.4 NA planapochromat objectives (Carl Zeiss). Bright field images were taken using Axio Scope (Carl Zeiss) mounting ×20/0.8 NA and ×40/0.75 NA planapochromat objectives (Carl Zeiss), coupled with an Axio Cam ICC1 color camera (Carl Zeiss).

**Striatal cell counts.** The total number of striatal MSNs defined by fluorescent DARPP-32 expression and in situ labeled enkephalin-expressing MSNs

**Figure 7 | ENK-specific TrkB deletion affects striatal D2R-dependent PKC and ERK1/2 phosphorylation.** (a,b) Representative immunoblots of proteins extracted from dorsal striatum of 13-M- and 5-M-old Trkb\(^{\text{PENK-WT}}\) and Trkb\(^{\text{PENK-KO}}\) mice. Animals were either treated with vehicle or with a D2R-specific antagonist L-741,626 (20 mg/kg), and killed 30 min later. Phosphorylation states of ERK1/2, PKC, AKT and DARPP-32 were determined using phospho-specific Abs. See also Supplementary Fig. S8b–c. (c) Graphs represent the phosphorylation state of each protein normalized to its total amount. Fold changes were determined against vehicle-treated Trkb\(^{\text{PENK-KO}}\) (set as 1), (vehicle: Trkb\(^{\text{PENK-WT}}\) versus Trkb\(^{\text{PENK-KO}}\), pERK1/2, P = 0.020; p-PKC, P = 0.028; p-DARPP32, P = 0.065; pAKT, P = 0.17). (L-741,626 treated Trkb\(^{\text{PENK-KO}}\) versus vehicle treated Trkb\(^{\text{PENK-KO}}\), pERK1/2, P = 0.016; p-PKC, P = 0.026). Values were generated by unpaired Student’s t-test. Values are means ± s.e.m., n = 6 per genotype (three per condition) analyzed. (d) Graphs represent the phosphorylation state of each protein normalized to its total amount at 5 M of age. Total DARPP-32 was normalized to ERK1. Fold changes were determined against vehicle-treated Trkb\(^{\text{PENK-WT}}\) (set as 1), (vehicle: Trkb\(^{\text{PENK-KO}}\) versus Trkb\(^{\text{PENK-WT}}\), pERK1/2, P = 0.06; p-PKC, P = 0.90; DARPP-32, P = 0.33). Values were generated by unpaired Student’s t-test. Values are means ± s.e.m., n = 5 per group analyzed.
(Supplementary Methods) were estimated using design-based stereology with the optical fractionator method as described by Gundersen et al.53 Sequential sagittal sections with a thickness of 0.3 mm spanning the entire medio-lateral extension of one brain hemisphere were obtained at the cryostat (Leica Microsystems). One in every 12 sections were counted with an unbiased counting frame size 50 × 10 μm for DARPP-32 and 100 × 100 μm for Enk. The frames were randomly positioned with the ‘Unbiased counting frame’ macro of Image (v1.46s, NIH) with a regular distance over 200 μm. For DARPP-32 and x = 330 μm, y = 300 μm for Enk. The heights of the dissectors were of 8 and 15 μm with a guard of 4 and 2 μm on either side, respectively (Supplementary Fig. 7). Cells in each frame were counted complying with the optical fractionator counting rules. Striatal boundaries were defined using the following morphological references: the corpus callosum, the external capsule, the lateral ventricle and the anterior commissure.

Morphological analysis of MSNs including spine density. The morphological analysis of striatal neurons was based on previously described Gólig-Cox impregnation method reported to be highly sensitive for detecting morphological alteration of striatal cells6. More details are reported in Supplementary Methods.

Haloperidol treatment. Three-month-old TrkB-PENK-KO and TrkB-PENK-WT mice were intraperitoneally (i.p.) injected with 1 mg kg\(^{-1}\) haloperidol (H1512, Sigma) or vehicle (DMSO), and 2 h later, were euthanized, perfused transcardially and processed for immunohistochemistry as described above.

Immunoblotting. Protein lysates from adult controls and mutant mouse brains were obtained by dissection of the cortex and striatum. Tissues were homogenized in 10 volumes of (vol/wt) lysis buffer (150 mM NaCl, 50 mM Tris pH 8.0, 1% (vol/vol) NP-40, 0.1% (wt/vol) SDS plus a cocktail of protease inhibitors, Sigma) by a glass dounce (VWR), followed by mild sonication. Total protein lysate (30 μg) were then subjected to SDS–PAGE. After transfer onto nitrocellulose membranes (GE healthcare), membranes were probed with Ab against TrkB (1:500, 80E3, Cell Signaling), Synaptophysin (1:10,000, Syb-Sy), D2R (1:200, Millipore), DARPP-32 (1:1,000, Millipore), D1R (1:200, Millipore), D1R (1:200, Millipore), D1R (1:200, Millipore), ERK (1:1,000, Zymed), PKC\(_{\text{epsilon}}\) (1:1,000, Calbiochem), Akt (1:1,000, Cell Signaling), GAD67 (1:200, Millipore), Neuroligin-2 (1:500, Syb-Sy), GAD67 (1:1,000, Millipore), DARPP-32 (1:2,000, Millipore). The body weights of mice involved in the behavioral tests were noted before every task.

Electrophysiological experiments. The electrophysiological recordings were performed according to Calabresi et al.55 Adult male mice (5–9 months) were used for all the experiments. Details are reported in the Supplementary Methods.

Statistical analysis. Two-way ANOVA using either age and genotype, or dendritic order and genotype as independent variables, was used for the data analysis of FDR corrected, and spontaneous locomotion or spine density and dendritic thickness, respectively. One-way ANOVA was used to determine significance between different genotypes in Sholl analysis and for cocaine-induced locomotion analysis. In case of significant difference among the means, a post hoc Fischer PLSD test was performed and the P-value was reported. Cell counts, synaptic density, WB analysis, localization of BAC–PENK and BAC–PENK-WT, t-data, and behavioral data for the CatWalk, inverted screen and grip strength were analyzed by two-tailed unpaired Student’s t-test. All tests of significance were performed at \(p = 0.05\) using Unistat 5.6 (Unistat Ltd, London, UK).

References


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Author contributions

D.B. planned and performed most of the experiments. M.G. conducted the in situ experiments. D.K. contributed to the characterization of the transgenic BAC-Cre line used in this study. B.P., V.P. and P.C. provided the expertise for and conducted the electrophysiological experiments. T.S. and D.M.R. provided the expertise for and conducted the fine motor behavioral analysis. L.M. is the principal investigator and contributed to the experimental plans, supervised the project, provided theoretical input and wrote the manuscript.

Additional information

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