I-DOPA–induced dyskinesia (LID), a detrimental consequence of dopamine replacement therapy for Parkinson’s disease, is associated with an alteration in dopamine D1 receptor (D1R) and glutamate receptor interactions. We hypothesized that the synaptic scaffolding protein PSD-95 plays a pivotal role in this process, as it interacts with D1R, regulates its trafficking and function, and is overexpressed in LID. Here, we demonstrate in rat and macaque models that disrupting the interaction between D1R and PSD-95 in the striatum reduces LID development and severity. Single quantum dot imaging revealed that this benefit was achieved primarily by destabilizing D1R localization, via increased lateral diffusion followed by increased internalization and diminished surface expression. These findings indicate that altering D1R trafficking via synapse-associated scaffolding proteins may be useful in the treatment of dyskinesia in Parkinson’s patients.
PSD-95 expression controls L-DOPA dyskinesia through dopamine D1 receptor trafficking

Gregory Porras,1,2 Amandine Berthet,1,2 Benjamin Dehay,1,2 Qin Li,3 Laurent Ladepuche,4,5 Elisabeth Normand,4,5 Sandra Dovero,1,2 Audrey Martinez,1,2 Evelyne Doudnikoff,1,2 Marie-Laure Martin-Négrier,1,2 Qin Chuan,3 Bertrand Bloch,1,2 Daniel Choquet,4,5 Eric Boué-Grabot,1,2 Laurent Groc,4,5 and Erwan Bezard1,2,3

1Université de Bordeaux and 2CNRS, Institut des Maladies Neurodégénératives, Bordeaux, France. 3Institute of Lab Animal Sciences, China Academy of Medical Sciences, Beijing, China. 4Université de Bordeaux and 5CNRS, Institut Interdisciplinaire des Neurosciences, Bordeaux, France.

L-DOPA–induced dyskinesia (LID), a detrimental consequence of dopamine replacement therapy for Parkinson’s disease, is associated with an alteration in dopamine D1 receptor (D1R) and glutamate receptor interactions. We hypothesized that the synaptic scaffolding protein PSD-95 plays a pivotal role in this process, as it interacts with D1R, regulates its trafficking and function, and is overexpressed in LID. Here, we demonstrate in rat and macaque models that disrupting the interaction between D1R and PSD-95 in the striatum reduces LID development and severity. Single quantum dot imaging revealed that this benefit was achieved primarily by destabilizing D1R localization, via increased lateral diffusion followed by increased internalization and diminished surface expression. These findings indicate that altering D1R trafficking via synapse-associated scaffolding proteins may be useful in the treatment of dyskinesia in Parkinson’s patients.

Introduction
In the striatum, dopamine (DA) terminals from the substantia nigra pars compacta (SNc) converge with glutamatergic signals from the cortex on dendritic spines of striatal medium spiny projecting GABAergic neurons (1, 2). The degeneration of the nigrostrial pathway in Parkinson’s disease (PD) induces complex modifications in both DA and glutamate signaling, leading to significant morphological and functional modifications in the striatal neuronal circuitry (3–5). Chronic DA replacement therapy with L-3,4-dihydroyxyphenylalanine (L-DOPA) superimposes upon these DA depletion–induced changes, resulting in debilitating motor complications known as L-DOPA–induced dyskinesia (LID) (6–8). At the molecular level, the subcellular organization of and functional interactions between glutamate and DA receptors within the striatum are crucial both in the pathogenesis of PD (9) and in the development of LID (10, 11). LID has indeed been associated with plastic changes in postsynaptic neuronal targets in the striatum, including elevated extracellular levels of glutamate (12) and DA (13) and abnormal trafficking of DA D1 receptor (D1R) (14, 15) and of NMDA and AMPA glutamate receptor subunits (5, 10, 16, 17). Such exaggerated DA and glutamate receptor expression at the plasma membrane results in abnormal activation of key signaling kinases (18–22). All these changes point to dysfunctional interactions between DA and glutamate neurotransmission in LID (5, 23, 24), although the molecular mechanisms remain elusive, despite recent progress (14, 25).

The membrane-associated guanylate kinase (MAGUK) proteins, such as postsynaptic density 95 (PSD-95), organize ionotropic glutamate receptors and their associated signaling proteins, regulating the strength of synaptic activity. Interestingly, PSD-95 might also interact with DA D1R (26), thereby potentially regulating DA D1R trafficking and function (27, 28). Increased levels of PSD-95 in the synaptic compartment have been reported in a rat model of LID (29), grounding the hypothesis that downregulation of PSD-95 levels could decrease the severity of LID by releasing D1R membrane anchorage. In order to test this challenging hypothesis, we here manipulated PSD-95 expression as well as its interaction with D1R in brain areas of rat and monkey models of dyskinesia and explored its effect at different biological scales, from behavioral outcomes to trafficking of single receptors.

Results
PSD-95 levels are massively increased in dyskinetic parkinsonian monkeys. Both expression and subcellular distribution of PSD-95 are altered in the striatum of L-DOPA–treated unilateral 6-OHDA–lesioned rats (29), with increased expression and enrichment in the synaptic compartment. To further support our hypothesis, we assessed the status of PSD-95 expression after DA denervation and after DA denervation followed by long-term L-DOPA treatment in the reference experimental model of PD and LID, the 1-methyl-4-phenyl-1,2,3,6-tetrahydropyridine–treated (MPTP-treated) macaque monkey (30, 31). Immunoblot levels were measured on striatal homogenates collected in normal and MPTP-treated macaque monkeys chronically exposed to vehicle or L-DOPA (see Supplemental Figure 1A for experimental design; supplemental material available online with this article; doi:10.1172/JCI59426DS1). Expression of PSD-95 in monkey putamen was significantly increased in the dyskinetic MPTP-lesioned monkeys compared with the various control animals (P < 0.05; Figure 1A). Interestingly, another synapse-associated protein (SAP) — SAP97, reported to be increased in the monkey putamen was significantly increased in the dyskinetic MPTP-lesioned monkeys compared with the various control animals (P < 0.05; Figure 1A). Interestingly, another synapse-associated protein (SAP) — SAP97, reported to be increased in the dyskinetic MPTP-lesioned monkeys compared with the various control animals (P < 0.05; Figure 1A). Interestingly, another synapse-associated protein (SAP) — SAP97, reported to be increased in the dyskinetic MPTP-lesioned monkeys compared with the various control animals (P < 0.05; Figure 1A). Interestingly, another synapse-associated protein (SAP) — SAP97, reported to be increased in the dyskinetic MPTP-lesioned monkeys compared with the various control animals (P < 0.05; Figure 1A). Interestingly, another synapse-associated protein (SAP) — SAP97, reported to be increased in the dyskinetic MPTP-lesioned monkeys compared with the various control animals (P < 0.05; Figure 1A). Interestingly, another synapse-associated protein (SAP) — SAP97, reported to be increased in the dyskinetic MPTP-lesioned monkeys compared with the various control animals (P < 0.05; Figure 1A). Interestingly, another synapse-associated protein (SAP) — SAP97, reported to be increased in the dyskinetic MPTP-lesioned monkeys compared with the various control animals (P < 0.05; Figure 1A). Interestingly, another synapse-associated protein (SAP) — SAP97, reported to be increased in the dyskinetic MPTP-lesioned monkeys compared with the various control animals (P < 0.05; Figure 1A). Interestingly, another synapse-associated protein (SAP) — SAP97, reported to be increased in the dyskinetic MPTP-lesioned monkeys compared with the various control animals (P < 0.05; Figure 1A). Interestingly, another synapse-associated protein (SAP) — SAP97, reported to be increased in the dyskinetic MPTP-lesioned monkeys compared with the various control animals (P < 0.05; Figure 1A). Interestingly, another synapse-associated protein (SAP) — SAP97, reported to be increased in the dyskinetic MPTP-lesioned monkeys compared with the various control animals (P < 0.05; Figure 1A). Interestingly, another synapse-associated protein (SAP) — SAP97, reported to be increased in the dyskinetic MPTP-lesioned monkeys compared with the various control animals (P < 0.05; Figure 1A). Interestingly, another synapse-associated protein (SAP) — SAP97, reported to be increased in the dyskinetic MPTP-lesioned monkeys compared with the various control animals (P < 0.05; Figure 1A). Interestingly, another synapse-associated protein (SAP) — SAP97, reported to be increased in the dyskinetic MPTP-lesioned monkeys compared with the various control animals (P < 0.05; Figure 1A). Interestingly, another synapse-associated protein (SAP) — SAP97, reported to be increased in the dyskinetic MPTP-lesioned monkeys compared with the various control animals (P < 0.05; Figure 1A). Interestingly, another synapse-associated protein (SAP) — SAP97, reported to be increased in the dyskinetic MPTP-lesioned monkeys compared with the various control animals (P < 0.05; Figure 1A).

PSD-95 expression controls L-DOPA dyskinesia through dopamine D1 receptor trafficking
Western blot analysis revealed that anti-D1 antibody immunoprecipitated PSD-95 and, conversely, anti–PSD-95 immunoprecipitated DA D1R (Figure 1B), supporting relevant D1R/PSD complex formation in vivo.

PSD-95 downregulation interferes with behavioral sensitization to l-DOPA in 6-OHDA–lesioned rats. The above findings indicate that increased PSD-95 levels are associated with dyskinesia. We next explored the consequences of overexpressing or downregulating PSD-95 using lentivirus (LV) in the rat analog of dyskinesia, known as the abnormal involuntary movements (AIMs) model (25, 32–34).

All LV transfer vector constructs were modified from the original FUGW vector backbone (35) and encoded GFP (36) carrying PSD-95 or shRNA to PSD-95 (referred to as PSD and shPSD groups, respectively; see Methods). Control vectors were a LV encoding GFP (mock group) and a LV encoding for shRNA to PSD-95 plus exogenous shRNA-resistant PSD-95 (SH+PSD group) (36).

We first investigated the effect of regulating PSD-95 expression levels on AIM development (Supplemental Figure 1B). Drug-naive LV-injected 6-OHDA–lesioned rats were treated once daily with l-DOPA, and the frequency of AIMs was scored daily. Repeated administration of l-DOPA to the mock group progressively increased AIM score (Figure 2A) and refs. 25, 32). Knockdown of PSD-95 markedly reduced this score compared with the mock group from day 5 onward (P < 0.01; Figure 2A), whereas both the PSD and SH+PSD groups were not different from mock rats, suggestive of a ceiling effect for PSD-95. The highlight of AIM time course at day 9 further revealed that AIMs were significantly decreased for the whole time course after l-DOPA administration in the SH+PSD versus the mock group (P < 0.01; Figure 2B). Interestingly, shPSD did not modify SAP97 levels (Supplemental Figure 2B), which suggests that the shPSD-mediated behavioral effect does not involve compensation for reduced PSD-95 levels by increasing expression of this other SAP. These behavioral data were further supported by the significant decrease in pERK immunoblot levels, a signaling marker classically associated with AIMs (20, 21, 37), in the shPSD versus the mock group (pERK/total ERK; mock, 125.9 ± 2.9 AU; shPSD, 104.0 ± 8.8 AU; P > 0.05, unpaired t test). Similarly, the mRNA level of cytochrome oxidase subunit I (COX I; a marker whose expression correlates with electrical activity; refs. 38, 39), as determined by qPCR in the substantia nigra pars reticulata (SNr; the output structure of the basal ganglia) was higher in the shPSD than in the mock group (21% increase; P = 0.032). This suggests that improvement in behavior correlates with normalization of the overdecreased SNr activity in dyskinetic animals (40, 41).

We examined the expression level of endogenous PSD-95 as well as exogenous PSD-95–GFP by Western blotting using anti–PSD-95 antibodies in rat striatum of animals infected with the different LVs. Detection of 2 bands (95 and 125 kDa) corresponding to endogenous and exogenous GFP-tagged PSD-95 in animals infected with PSD-95–GFP alone or shPSD+PSD-95–GFP (Figure 2C) was indicative of the efficiency of LV-driven expression in both set of experiments. Importantly, the expression of endogenous PSD-95 was significantly diminished in presence of shPSD. Endogenous PSD-95 represented about 45% (P < 0.01) and 65% (P < 0.05) of the control mock group in experiments 1 and 2, respectively (Figure 2, A and B). Biochemical demonstration of PSD-95 downregulation was further confirmed in the striatum by double immunohistochemistry for GFP and PSD-95 (Supplemental Figure 3A) across the 4 groups, with medium spiny-like neurons being clearly labeled (Supplemental Figure 3B). The shPSD group displayed a severe
reduction of PSD-95 immunoreactivity, with most medium spiny-like neurons being immunopositive only for GFP (Supplemental Figure 3B). 3-dimensional reconstruction of LV-transfected rat brain showed that the striatum was largely transfected (>50%) in the 4 groups (Supplemental Movie 1). Together, these controls demonstrated the efficiency of the targeting and of the reduction in PSD-95 levels in the shPSD group, a result compatible with the marked reduction in AIM severity in these animals.

Cell-permeable peptide disrupting the PSD-95–D1R interaction reduces established AIMs in 6-OHDA–lesioned rats An alternate option to lowering PSD-95 levels for preventing the interaction of PSD-95 with D1R is to design (and administrate) competitive peptides corresponding to the domain of rat D1R interacting with PSD-95 (42) that would disrupt D1R/PSD-95 interaction. We first examined the ability of the distal region of the C-terminal tail of D1R to bind to PSD-95 by performing pulldown assays with active (D1-CT) and control (D1-SCR) peptides (Supplemental Figure 4). Extracts of monkey or rat brain tissues as well as HEK293 cells expressing PSD-95 or PSD-95–GFP were incubated in the presence of immobilized biotinylated peptides. PSD-95 coprecipitated with the active D1-CT peptide (Supplemental Figure 4), which indicates that D1-CT interacts with native or recombinant PSD-95 from different species. In addition, detection of PSD-95–GFP indicated that fusion of the GFP does not alter function of PSD-95 related to D1R.

Figure 2
Downregulating PSD-95 in the DA-deprived striatum reduces AIM severity in the hemiparkinsonian rat. (A) Both the peak AIM scores (rated at 60 minutes) from day 5 as well as the time course AIM (bottom right inset) scores were significantly reduced with shPSD LV compared with mock control. *P < 0.05, shPSD vs. mock. (B) Rats displayed identical AIMs after 9 l-DOPA treatment days before receiving LV at day 12 (red arrow). Peak AIM score in the shPSD group was significantly reduced compared with all other groups. This reduction was further highlighted in a time course experiment at day 27 (bottom right insets), with a significant decrease 60 and 90 minutes after l-DOPA administration. In both sets, animals had similar loss of dopaminergic innervation in the striatum as detected by TH immunohistochemistry (top left insets). *P < 0.05, shPSD vs. all other groups. (C) Detection of the endogenous PSD-95 and exogenous PSD-95–GFP transgene by Western blot in striata extracts from 2 sets of animals. Both lesioned (L) and unlesioned (UL) sides are shown. In both experiments, shPSD significantly reduced the expression of endogenous PSD-95, whereas PSD significantly increased PSD-95, compared with the mock group. Lanes were run on the same gel but were noncontiguous (white lines). *P < 0.05 vs. mock. (D) While TAT-D1-SCR peptide had no effect on AIM score, TAT-D1-CT (0.25 and 0.5 nmol) dose-dependently reduced AIM score at peak and at 60 and 90 minutes after l-DOPA administration, resulting in overall improvement. *P < 0.05 vs. vehicle.
Both peptides were fused to the cell membrane transduction domain of the human immunodeficiency virus 1 (i.e., the TAT signal) to gain cell permeability (herein, we refer to the control as TAT-D1-SCR and the active form as TAT-D1-CT). We tested whether administration of TAT-D1-CT would reduce the severity of established AIMS (Supplemental Figure 1B), as in the shPSD experiment. 6-OHDA-lesioned rats were treated with 1-DOPI for 21 days. TAT peptides were injected through intra-striatal cannulae at the same time as 1-DOPI administration. While TAT-D1-SCR did not affect AIM score (Figure 2D), TAT-D1-CT dose-dependently reduced AIM scores (P < 0.05; Figure 2D). The AIM time course showed that AIMS were significantly decreased at peak dyskinesia, i.e., at 60 and 90 minutes (P < 0.05) after 1-DOPI administration, in the highest dose of the TAT-D1/PSD condition.

Decreasing PSD-95 levels alleviates LID in MPTP-lesioned macaques. The rat experiments clearly suggested that reducing PSD-95 expression or competing with D1R/PSD-95 interaction interferes with the development (Figure 2A) and reduces the severity (Figure 2, B–D) of AIMS. When drug treatment was resumed on day 22, however, animal LID had decreased 40%–50%. At the end of day 27, all but the shPSD group was back to the same level of LID as on day 9, while shPSD LID remained low. However, since LID in the 6-OHDA rat model might not have fully consolidated, final preclinical proof-of-concept evidence for an antidyskinetic strategy should be obtained in the fully stable MPTP-lesioned macaque model of PD, as clinical manifestations resemble the severe peak dose LID in human PD (30, 31). We therefore investigated the effectiveness of the shPSD strategy in the stably dyskinetic MPTP-lesioned macaque monkey as well as the preservation of 1-DOPI-mediated antiparkinsonian action.

First, we selected the shRNA plasmids against human PSD-95 with the maximum efficiency of silencing endogenous PSD-95 in the BE-M17 human cell line (Figure 3A). We concluded that sh628 was the best candidate, with a 50% decrease in expression (P < 0.05). The ability of the LV encoding sh628 to silence human PSD-95 was tested in the same human cell line (Figure 3A).

MPTP-treated macaques rendered dyskinetic by chronic 1-DOPI treatment (Supplemental Figure 1A) received mock (encoding GFP; n = 3) or shPSD (n = 4) LV in the motor striatum. Upon completion of the behavioral experiments, all monkeys were tested to evaluate the extent of lesion and transduction efficacy. Both groups had similarly extensive nigrostriatal cell loss, as evidenced by the dramatic decrease in the number of tyrosine hydroxylase–immunopositive (TH-immunopositive) neurons in the SNc (unbiased stereological counting; mock, 13,700 ± 2,729; shPSD, 12,800 ± 1,658; unlesioned controls, 126,300 ± 3,241). PSD-95 immunohistochemistry revealed a significant decrease in PSD-95 levels of the shPSD versus the control group (Figure 3B), a decrease that covered 67.9% ± 12.8% of the postcommissural motor putamen (Figure 3B). Confirmation of the decrease in endogenous PSD-95 after shPSD LV was obtained by Western blot of homogenized striatal tissues collected on sections in the motor striatum (Figure 3C).

Prior to LV administration, parkinsonian disability scores in both the Off and On states (before and after 1-DOPI administration, respectively; Figure 4A), LID scores in the Off state (Figure 4C), and the time course of 1-DOPI–induced locomotor activity (Figure 4E) were comparable. Starting at 4 weeks postsurgery, when behavioral experiments resumed, PD and LID scores as well as locomotor activity counts in the mock group were fully comparable to those prior to surgery (Figure 4, B, D, and F). In test animals, the antiparkinsonian efficacy of 1-DOPI remained intact in mock versus shPSD animals (Figure 4, A and B). However, shPSD monkeys had significantly less severe LID (Figure 4, C and D) and a significant decrease in 1-DOPI–induced locomotor hyperactivity in the On state (Figure 4, E and F). Since locomotor activity counts capture both the antiparkinsonian and prodyskinetic actions of 1-DOPI, an antidyskinetic strategy would reduce...
the activity counts, reflecting the reduction of hyperkinetic movements, but would not suppress them, or it would indicate an anti-L-DOPA effect. Measurement of On-time also showed that shPSD animals were comparable to the mock group (Figure 4G). In fact, while On-time per se was not affected, shPSD animals benefited from shPSD LV, with a significant increase in Good On-time (Figure 4H) and a concomitant significant decrease in Bad On-time.

Figure 4
Reducing PSD-95 in the MPTP-treated macaque markedly alleviates dyskinesia. (A, C, and E) Time course of clinical ratings (median) and locomotor activity counts (mean). Data are shown without SEM for readability. L-DOPA or amantadine administration began at 0 minutes. (A) shPSD had no effect on L-DOPA–induced parkinsonian disability score at any time point. (B) Area under the curve (AUC) analysis of parkinsonian disability scores confirmed this lack of effect. Orange bar denotes the presurgery score of the entire population. *P < 0.05 vs. L-DOPA–treated mock (bracket colors correspond with respective groups to show time frame of significance). (D) The overall positive effect of shPSD and/or pharmacological treatment on LID severity was further demonstrated by the AUC data. *P < 0.05 vs. L-DOPA–treated mock; #P < 0.05 vs. L-DOPA–treated shPSD. Orange bar denotes the presurgery score of the entire population. (E and F) Consequently, locomotor activity was lower in shPSD animals receiving L-DOPA alone and in both groups when receiving L-DOPA plus amantadine. Orange bar denotes the presurgery count for the entire population. *P < 0.05 vs. L-DOPA–treated mock; #P < 0.05 vs. L-DOPA–treated shPSD. (G) On-time was not affected by any treatment. (H) shPSD significantly increased Good On-time. *P < 0.05 vs. L-DOPA–treated mock. (I) Bad On-time was reduced in shPSD animals receiving L-DOPA alone and in both groups when receiving L-DOPA plus amantadine. *P < 0.05 vs. L-DOPA–treated mock.

These are key translational parameters, as they quantitatively reflect the possible qualitative improvement to be reported by dyskinetic patients enrolled in a clinical trial. The combination of the clinically assessed decrease in LID scores, reduced locomotor activity, and stable On-time with a marked increase in Good On-time supports our hypothesis that striatal downregulation of PSD-95 diminishes LID severity without reducing the positive effects of L-DOPA on parkinsonian motor scores.

Although the LID-related rise in PSD-95 levels (Figure 1A) was only reduced 40% by the shPSD LV (Figure 3B), i.e., PSD-95 levels were not brought back to normal levels, one cannot rule out a possible impairment of the glutamate corticostriatal transmission. At present, amantadine, an antagonist at the N-methyl-d-aspartate (NMDA) subtype of glutamate receptor, is the drug most often used for the treatment of LID (43). Mock animals displayed the expected reduction in LID severity after acute pharmacological challenge with amantadine (20 mg/kg orally; ref. 44) in combination with L-DOPA (Figure 4, C and D) without reduction in antiparkinsonian response to L-DOPA, as evidenced by parkinsonian disability score (Figure 4, A and B), locomotor activity profile (Figure 4, E and F), and On-time (Figure 4G). However, shPSD animals further benefited from amantadine treatment: LID severity of shPSD animals was significantly different not only from mock animals treated with L-DOPA only, but also from shPSD animals treated with L-DOPA only (Figure 4, C and D) Such additive effect was also significant for the locomotor activity counts (Figure 4, E and F). The combination of L-DOPA and amantadine in shPSD animals resulted into an almost complete suppression of Bad On-time (Figure 4I). Together, these data suggest that NMDA-mediated corticostriatal transmission is not blunted in shPSD animals, as they additively benefited from combination L-DOPA and amantadine therapy.

PSD-95 downregulation alters D1R surface content and trafficking. The binding of D1R and glutamate receptors to the scaffold protein PSD-95 (28, 42) has provided a molecular mechanism by which these receptors are anchored in synaptic areas and by which these 2
Figure 5
PSD-95 knockdown or peptide disrupting PSD-95–D1R interaction alters D1R surface content and dynamics on cultured striatal neurons. (A and B) Immunostaining of surface D1R (A) showed a significant decrease in surface D1R in shPSD neurons (B). Scale bar: 16 μm. *P < 0.05. (C) Representative trajectories of surface D1Rs labeled with a single QD-antibody complex (inset) at the surface of a PSD-95–expressing neuron. A fraction rapidly diffused along dendrites and explored large areas (arrows), whereas others diffused in a confined area (arrowheads). Scale bars: 4 μm (left); 200 nm (right). (D) Representative trajectories of surface D1R on a Homer1c-GFP–expressing neuron. Also shown are distributions (median and 25%–75% IQR) of the instantaneous diffusion coefficient of surface D1Rs inside and outside the PSD area. Scale bar: 2 μm. *P < 0.05. (E) Representative trajectories of surface D1Rs on a shPSD-expressing striatal neuron. Scale bar: 4 μm. (F) Cumulative distribution of the instantaneous diffusion coefficient of surface D1Rs, starting from the percentage of immobile receptors. (G) Corresponding distributions of the instantaneous diffusion coefficient of surface D1Rs. *P < 0.05. (H) The relationship between mean square displacement (MSD) and time of surface D1R trajectories was not linear (dotted line denotes free diffusion), adopting negative curvature, characteristic of confined behavior (Kolmogorov-Smirnov test). (I and J) Representative trajectories (I) and distributions of the instantaneous diffusion coefficient (J) of surface D1Rs in the presence of TAT-D1-SCR or TAT-D1-CT. Scale bar: 5 μm. *P < 0.05.
different pathways may communicate (27, 45). The above data provide clear evidence that PSD-95 expression affects the emergence of LID, which is greatly dependent on DA receptor–mediated signaling (30, 31). Because postmortem assessment of DA, metabotropic, and ionotropic glutamate receptors in the rat striatum did not show any dramatic regulation of the receptor content after over- or downregulation of PSD-95 (Supplemental Figure 5), we decided to identify, at the molecular level, the relationship between PSD-95 and surface D1R signaling using a combination of single nanoparticle tracking and immunocytochemical approaches in cultured striatal neurons. In this in vitro preparation, PSD-95 knockdown (i.e., shPSD) significantly reduced the cellular content of PSD-95 by approximately 30% (Supplemental Figure 6), a value equivalent to those observed in rat and monkey striata (Figure 2C and Figure 3, B and C). We then investigated the effect of PSD-95 knockdown on the cellular distribution of D1R. Consistent with the effect observed in the rat brain, the global content of D1R was not significantly altered in shPSD-transfected neurons compared with neighboring untransfected ones (Supplemental Figure 6). However, the surface content of D1R was drastically reduced (4-fold decrease) in the shPSD condition (Figure 5, A and B), in both the dendritic and the somatic compartments. Because PSD-95 strongly interacts with NMDAR and anchors the receptor to the synapse (46, 47), we investigated the effect of PSD-95 knockdown on the surface expression of GluN2B-NMDAR (Supplemental Figure 6). In contrast to the effect observed with D1R, the surface content was similar in shPSD-transfected and neighboring untransfected neurons (Supplemental Figure 6), which indicates that PSD-95 altered the surface content of D1R, but not of NMDAR, in striatal neurons. Whether the surface dynamics of different NMDAR subtypes are affected in these conditions, however, remains to be explored. Together, these data demonstrate that PSD-95 knockdown affects D1R surface content without any significant effect on its global one, suggestive of specific modulation of receptor surface trafficking.

To further identify the pathways involved in PSD-95–dependent regulation of D1R trafficking, we used a single-particle tracking approach (quantum dot [QD]; ref. 48) to follow over time single D1R-QDs at the surface of striatal neurons (Figure 5C). A small fraction of surface D1R-CFP was labeled with anti-GFP antibody coupled to a single QD in live striatal neurons. In PSD-95–expressing neurons, surface D1Rs were highly mobile and diffusive (Figure 5C). Representative trajectories from a 60-second imaging session revealed that most D1Rs diffused (median instantaneous diffusion coefficient, 0.04 μm²/s; interquartile range [IQR], 0.01–0.08 μm²/s; n = 2,695 trajectories), although a confined pattern of diffusion was observed (Figure 5C). Because PSD-95 itself can affect the surface dynamics of D1R, we measured the surface diffusion of D1R in neurons expressing Homer1c, another protein of the postsynaptic density that does not interact with D1R. The diffusion of D1R in these neurons was not significantly different from the one in neurons expressing PSD-95 (median instantaneous diffusion coefficient, 0.03 μm²/s; IQR, 0.001–0.07 μm²/s; n = 312 synaptic and 1,695 extrasynaptic trajectories; Figure 5D). We took advantage of the presence of Homer1c clusters, indicative of postsynaptic density, to investigate D1R diffusion characteristics in glutamate synapses. D1Rs were transitorily retained in the PSD area. D1Rs were thus less diffusive in the vicinity and within PSD (Figure 5D), consistent with the presence of anchoring mechanisms for surface D1R in glutamate synapses.

In the basal condition, more than 80% of surface D1Rs were mobile at the surface of striatal neurons (expressing wild-type PSD-95 or Homer1c). In contrast, PSD-95 knockdown produced a substantial change in D1R surface trafficking (800-frame duration; Figure 5E). Indeed, the overall surface diffusion of D1Rs was significantly increased, with more than 94% mobile receptors (bin size, 0.005 μm²/s; n = 121–2,695 trajectories; Figure 5F). The D1R median instantaneous diffusion coefficient was significantly increased, from 0.04 to 0.09 μm²/s (IQR, 0.04–0.16 μm²/s), in PSD-95 and shPSD conditions, respectively (n = 121–2,695 trajectories; Figure 5G). In addition, the D1R diffusion pattern was examined by plotting the mean squared displacement (surface explored by the receptor) versus time interval (49). In PSD-95–expressing neurons, the D1R surface diffusion pattern was not purely Brownian (comparison of 56 trajectories per group; Figure 5H), suggestive of the presence of regulatory mechanisms at the neuronal surface. In shPSD conditions, the diffusion pattern abruptly changed toward free diffusion (Figure 5H), which indicates that the surface trafficking of D1R was highly regulated by the presence of PSD-95. To investigate the role of the direct interaction between D1R and PSD-95, we incubated neurons with the TAT-D1-CT peptide, which disrupted interaction between D1R and PSD-95, and measured its acute effect on D1R surface trafficking. TAT-D1-CT (5 μM for 10 minutes) produced an increase in the diffusion of surface D1R (Figure 5I). The D1R median instantaneous diffusion coefficient was significantly increased, from 0.08 μm²/s for TAT-D1-SCR (IQR, 0.03–0.16 μm²/s; n = 2,619 trajectories) to 0.125 μm²/s for TAT-D1-CT (IQR, 0.05–0.215 μm²/s; n = 3,924 trajectories; P < 0.01 between groups; Figure 5I). Since the peptide described may interfere with the interaction between GluN2A and D1R, we confirmed that TAT-D1-CT (5 μM, 10 minutes) did not alter the content or distribution of GluN2A or GluN2B subunits (Supplemental Figure 7). Together, these data indicated that PSD-95 plays an important role in the surface content and dynamics of D1R in striatal neurons.

Discussion
We demonstrated here that disrupting D1R and PSD-95 interaction in the striatum of dyskinetic rats and monkeys, through either lowering PSD-95 levels or blocking direct interaction using competing peptides, reduced dyskinesia severity. This outcome was achieved by destabilizing D1R localization through increased lateral diffusion, possibly leading to increased internalization and thus less surface expression. PSD-95 is at the crossroad of glutamate and DA signal modulation (47), especially as glutamatergic and DA axons form a synaptic triad at postsynaptic dendritic spines in the striatum, a close spatial association that is required for the converging and coordinating actions of the 2 afferent pathways (50). LID is associated with abnormal trafficking of DA D1Rs (14, 15) and of NMDA and AMPA glutamate receptor subunits (5, 10, 16, 17). Such exaggerated DA and glutamate receptor expression at the plasma membrane results in abnormal activation of key signaling kinases (18–22, 37). The molecular mechanisms mediating such lack of trafficking that causes enhanced anchorage and function remain the subject of debate. Alteration of the DA D1R homologous desensitization process is partly responsible for such anchoring (14, 51), as increasing the capacity of the desensitization machinery (52, 53) in the parkinsonian striatum ameliorates LID (14, 25). In the same way, the redistribution of NMDA glut-
mate receptor subunits with higher levels of GluN2A and lower levels of GluN2B (16, 54) is paralleled by profound modifications of NMDA receptor subunit association with interacting elements (47), i.e., MAGUK protein family members PSD-95 and SAP (29, 54). Treatment of nondyskinetic animals with TAT2A, a synthetic peptide able to affect GluN2B binding to MAGUK proteins as well as synaptic localization of this subunit in nondyskinetic rats, was sufficient to induce a shift of treated rats toward dyskinetic motor behavior (54). Comparably, another peptide directly targeting PSD-95–GluN2A subunit interactions, TAT2A, could reverse motor and synaptic plasticity abnormalities in early parkinsonism (55). These data indicate that interaction of NMDA subunits with PSD-95 governs their redistribution between synaptic and extrasynaptic membranes.

To date, the lateral diffusion at the neuronal surface has been demonstrated for glutamatergic metabotropic mGlu5, AMPA, and NMDA receptors (56–60), glycine receptor (61, 62), GABA-A receptor (63), β2-adrenergic receptor (64), μ-opioid receptor (65), and, finally, D1R (66). This initial demonstration of D1R mobility showed that the majority of dendritic D1Rs diffuse in the plane of the membrane and that their mobility is confined by interaction with NMDAR (66). Exposure to NMDA reduced the diffusion coefficient for D1R and caused an increase in the number of D1R-positive spines (66), a situation highly reminiscent of the elevated extracellular levels of glutamate (12) and DA (13, 41) associated with LID. Such interaction could be direct (66) or indirect through the PSD-95 protein that interacts and regulates D1R trafficking and anchorage (26, 27, 42). Psychostimulants in wild-type mice or genetic models of hyperdopaminergia have shown reduced PSD-95 levels in the striatum (42, 50). In this hyperdopaminergic context, PSD-95 inhibits the physical association between D1 and NMDAR and functionally uncouples D1R trafficking and signaling from modulation by NMDAR (28). In the l-DOPA–exposed dopamine-depleted experimental condition, the situation was rather different, with anchorage of D1R under the control of PSD-95.

To our knowledge, however, the role of surface receptor lateral diffusion in such a pathological condition had not previously been determined. We therefore addressed the dynamics of surface synaptic D1R by single QD tracking (48) to discern the mechanisms underlying PSD-95–dependent D1R trafficking. In keeping with the initial report (66), surface D1Rs were mobile and diffusive in PSD-95–expressing neurons. However, lowering PSD-95 levels using shPSD further increased the overall surface diffusion of D1R and decreased D1R surface membrane localization, but did not affect GluN2B-NMDAR surface expression. Comparable effects were obtained with the TAT-D1/PSD-95 peptide, which prevents direct interaction between D1R and PSD-95. However, it is important to note that knocking down PSD-95 had a broader effect on D1R surface dynamics than did TAT-D1-CT. It is thus likely that other interacting partners of PSD-95 participate in the membrane organization of D1R.

This latter point further supports the claim of a direct role of PSD-95 in governing the surface stabilization of D1R in the extrasynaptic compartment, as well as in the vicinity of corticostriatal synapses. Both diminishing PSD-95 expression levels and competing with direct interaction between D1R and PSD-95 clearly favored D1R lateral diffusion, possibly by unanchoring D1R in dendritic spines. In addition, PSD-95 can alter D1R intracellular trafficking (exocytotic or endocytotic rate) and surface content (26, 27, 42), which consequently modulates the dynamics of the surface receptors. For instance, D1Rs form dimers with GluN1/ GluN2A containing NMDA receptors (67), an association that is regulated by PSD-95 (28) and reduced after chronic l-DOPA treatment (68). Diminishing PSD-95 expression levels or competing with direct interaction between D1R and PSD-95 unanchors D1R, but would thus favor reformation of D1/NMDA complexes, a dynamic that warrants investigation.

We posited that releasing pathologically anchored D1R from dendritic spines would contribute to reduced LID. Our rat and primate experiments support this claim, with dramatic reductions in LID severity and concomitant increases in Good On-time (i.e., time without disabling dyskinesia). Development of such strategy should not, however, interfere with the antiparkinsonian action of the gold standard therapy, l-DOPA. In that respect, our nonhuman primate experiments clearly showed that lowering PSD-95 levels had no negative effect on therapeutic response to l-DOPA. Glutamatergic corticostriatal transmission seemed to be preserved, as further supported by our pharmacological experiments with amantadine. At present, the noncompetitive NMDA receptor antagonist amantadine is frequently used as adjunctive therapy for LID (43, 69, 70). Combination of l-DOPA and amantadine in shPSD animals caused near-complete suppression of dyskinesia and Bad On-time, dramatically increasing Good On-time. Together, these data suggest that NMDA-mediated corticostriatal transmission is not blunted in the shPSD animals, as they additively benefited from combination therapy with l-DOPA and amantadine. Rather, combined knockdown of striatal PSD-95 expression and amantadine therapy may be superior to amantadine alone as an adjunct to l-DOPA to treat dyskinetic side effects in PD patients.

Our data set identified a pathological mechanism for D1R anchoring at synapses. Pathological overexpression of PSD-95 anchors D1R in the dopamine-depleted brain, reducing its mobility in and out of synapses. Easing such mobility through enhanced lateral diffusion is achieved by (a) lowering PSD-95 expression levels and (b) administration of cell-permeable peptide affecting D1/PSD-95 interaction, with a magnitude compatible with preservation of corticostriatal communication. Because alterations in D1R function contribute to the clinical features of PD and underlie the development of dyskinesia, therapies aiming at interrupting or regulating its trafficking via synapse-associated scaffolding protein organization, such as the cell-permeable peptide tested in the present study, may be useful in the treatment of dyskinesia in PD patients. Thus, understanding the rules that govern D1R surface distribution and, most importantly, their dynamic retention in postsynaptic density may shed new light on the fine-tuning of D1R in pathological conditions such as PD and LID.

Methods

Rat experiments

LV construction. We used 4 LVs, provided and validated by O.M. Schluter (Stanford University School of Medicine, Palo Alto, California, USA; ref. 36). The different sequences were cloned into the LV vector FUGW (35), used as control (mock). The 3 other vectors were FUGW plus PSD-95, FUG plus shRNA for PSD-95, and FUGW plus shRNA for PSD-95 plus PSD-95 (referred to herein as PSD, shPSD, and SH+PSD, respectively). LV production was performed in the IFR 66 Vectorology Platform (University of Bordeaux 2, Bordeaux, France) by transfection with a viral vector.
system — mock (with the different inserts), pCMV-A8-9 (encapsidation plasmid), and VSV-G (cDNA encoding the envelope glycoprotein of vesicular stomatitis virus) — in FT-HEK293 cells. LV supernatants were concentrated by centrifugation concentration filter (centricon) to obtain a final titer of 2.5 × 10^9 pfu/ml.

**Peptide preparation and validation.** We used active peptides (D1-CT) corresponding to the last 46-amino acid sequence (from Leu401 to Thr446) of rat D1R and a scrambled version of the peptide as negative control (D1-SCR) (GenScript). Biotinylated forms of peptides were used for pull-down assays. For functional experiments, we used FITC-tagged D1 peptides fused to the cell membrane transduction domain of the human immunodeficiency virus 1 (VGRKKRRQRRR) to gain cell permeability (71). Peptides were dissolved immediately before used in water (10 μg/ml) for pull-down assays and in PBS at their final concentration for in vivo intracerebral administration.

For pulldown assay, 500 g of protein extracts from rat or monkey brain tissues as well as HEK293 cells transfected with PSD-95 or PSD-95–GFP were incubated with immobilized streptavidin bound to the indicated biotinylated peptides (100 μg) overnight at 4°C, as previously described (72). Beads (40 μl) were washed 4 times with 150 mM NaCl buffer and eluted with 1 volume of SDS sample buffer. Eluates (20 μl) and input samples (10 μg) were subjected to SDS-PAGE and Western blot using mouse anti–PSD-95 (UC Davis/NIH NeuroMab Facility) as described below.

**Stereotactic procedures.** Adult male Sprague Dawley rats (Charles River Laboratories), weighing 175–200 g at the beginning of the experiment, were used. They were maintained under standard laboratory conditions in a 12-hour light/12-hour dark cycle with free access to food and water. The 6-OHDA lesion was performed as previously described (14). Unilateral DA deprivation of the striatum was obtained by injection with 6-OHDA (3 μg/μl; Sigma-Aldrich) in the right medial forebrain bundle (2.5 μl at anterior-posterior –3.7 mm, medial-lateral 1.7 mm, and dorsoventral –8 mm relative to bregma; refs. 41, 73) in rats pretreated with citalopram (1 mg/kg i.p.; Lundbeck), an inhibitor of serotonin reuptake, and with noradrenergic reuptake (74). 5 μl of the concentrated LVs (mock, PSD-95, SH+PSD, and shPSD) or 3 μl of peptide solution was injected into the caudal lesioned putamen at the following coordinates: anterior-posterior, 0.2; medial-lateral, 3.5; dorsoventral, 5.7. LVs were injected either at the time of the 6-OHDA lesion (set 1) or after behavioral pretesting (set 2) (Supplemental Figure 1), while the peptides were injected concomitantly with t-DOPA on testing days.

**Behavioral assessment.** Animals displaying both an impaired stepping test measured on days 18–20 (75–77) and >95% loss of TH-immunopositive fibers in the striatum (78, 79), assessed after completion of all experiments, were retained for final analysis (Supplemental Figure 1). 3 weeks after 6-OHDA injection, rats were treated once daily with benzerazide (15 mg/kg i.p.; Sigma-Aldrich) or t-DOPA (6 mg/kg i.p.; Sigma-Aldrich) and benzerazide (15 mg/kg i.p.) for 9 days in order to induce gradual AIM development (n = 9 [mock]; 12 [PSD-95]; 8 [SH+PSD and shPSD]). In the second set of experiments, the rats were first treated with t-DOPA for 10 days with daily testing for AIMS. LV was injected on day 12, and rats were allowed to recover for 9 days before resuming the injection schedule and then tested for AIMS for the next 7 days (n = 9 [mock and shPSD]; 13 [PSD-95]; 11 [SH+PSD]). Peptide experiments were conducted on 11 dyskinetic rats. The 4 AIM categories (limb, axial, orolingual, and locomotive) were scored using a validated rating scale (33, 34) for 1 minute with each observation; maximal score for each observation, 16; maximal total score per session, 96) by a trained investigator as described previously (41, 80). Figures display the sum of the axial, limb, and orolingual AIMS.

Upon completion of the behavioral testing schedule, rats were deeply anesthetized with sodium chloral hydrate (400 mg/kg) 1 hour after t-DOPA injection and killed either by decapitation (for Western blotting experiments), with their brains being rapidly removed and frozen in a Snap Frost 80 Specimen Freezing System (Alphelys), or by transcardiac perfusion (for immunohistochemistry experiments) with 50–100 ml 0.9% NaCl followed by 250 ml 4% PFA in 0.1 M phosphate buffer (pH 7.4), with their brains being removed and postfixed overnight in 4% PFA at 4°C.

**Western blot.** Patches of tissue were collected directly around the injection site from cryostar-cut 200-μm-thick sections after GFP expression signal was observed immediately before and after on adjacent 20-μm-thick sections under fluorescence microscopy to delineate the transfected area. Mirror contralateral patches were collected as controls. Patches were individually sonicated in a lysis buffer RIPA (R0278; Sigma Aldrich) containing 0.1% SDS and centrifuged at 20,817 g at 4°C for 15 minutes. The resulting supernatants were collected, and the proteins were reduced with a buffer containing 25 mM DTT and 1% SDS until quantification of the whole protein by the Lowry method (Bio-Rad DC Protein Assay). Samples containing 10 μg of proteins were heated to 100°C for 5 minutes, and electrophoresis was performed onto a 8% polyacrylamide gel. Proteins were blotted on polyvinylidene fluoride membranes (Immbolon P; Millipore) in Miniprotein III (BioRad). Membranes were saturated for 90 minutes by incubation with 5% milk in TBS-Tween (Tris HCL, pH 7.5; 100 mM NaCl; 0.1% Tween 20) and incubated overnight with the following primary monoclonal antibodies diluted in TBS-Tween 0.1% supplemented with 1% BSA: anti–PSD-95 (UC Davis/NIH NeuroMab Facility), anti-tubulin (Sigma-Aldrich), anti-GFP (Invitrogen), anti-phosphoERK1/2 (Sigma-Aldrich) and anti-ERK1/2 (Cell Signalling Technology). Membranes were then incubated for 2 hours at room temperature with secondary anti-mouse or anti-rabbit (both 1:5,000; Jackson ImmunoResearch) HRP-coupled antibody diluted in TBS-Tween 0.1% and supplemented with 5% milk. The complexes were detected with enhanced chemiluminescence reagents (Eurobio). Images were acquired on a computer with a CCD camera coupled with an optical bench. Optical density (OD) was measured with Mercator software (Explorer Nova). Results were derived from 3 independent experiments.

**Immunohistochemistry.** Cryostar-cut 50-μm-thick free-floating sections were processed for PSD-95/GFP double immunohistochemistry. Block step in 5% milk powder in PBS for 30 minutes was followed by overnight incubation at room temperature with a mixture of mouse monoclonal antibody to PSD-95 (73 028, UC Davis/NIH NeuroMab Facility) and rabbit polyclonal antibody to GFP (A11122; Invitrogen) diluted at 1:2,000 and 1:1,000, respectively, in 0.2% BSA, 0.3% triton X100, and PBS. PSD-95 revelation was conducted first, and sections were incubated with anti-mouse HRP envision system (K4001, Dako) for 30 minutes at room temperature, rinsed in PBS, and incubated in anti-HRP 549 dye light conjugate diluted 1:1,000 in PBS for 60 minutes at room temperature. Sections were then incubated in anti-rabbit Alexa Fluor 488 diluted 1:500 in PBS for 60 min at room temperature for revealing GFP signal. Sections were mounted on gelatin-coated slides in PBS and coveredslipped with Vectashield HardSet mounting medium (H1400; Vector Laboratories). Fluorescent images were captured on a confocal Leica microscope DM 2500 TCS SPE. Digital images were acquired separately for each wavelength. Adobe Photoshop CS5 was used to further process digital images. Any adjustments to brightness and contrast were made uniformly to all parts of the image.

**Cytochrome oxidase subunit I (COX I) qPCR.** RNA extracts from isolated SNr were purified with RNeasy Plus Micro Kit (Qiagen), and 10 ng was retrotranscribed to cDNA at 42°C with reverse transcriptase (Fermentas) in the presence of 1 μl random primer (150 ng/μl) and 1 μl oligodt (200 ng/μl). Real-time PCR was performed on a Light Cycler 480 (Roche) in a total volume of 10 μl using 1 ng cDNA as tem-
plates, 3 μl mixed forward (GCTGGCTTCGTCCACTGATT) and reverse (AAGTGGGCTTTTGGCTCATGGT) primers, and 5 μl of mix 2 (Roche) containing Fast Start Taq DNA polymerase reaction buffer dNTP mix (with dUTP instead of dTTP), Sybr Green I dye, and MgCl2. Reliability of reference genes (ef1a1 and Sdha) was evaluated with GeNorm. The cycle point (Cp) value of COX I was normalized against that of Sdha and Ef1a1, and the relative level of expression was calculated using the comparative 2^ΔΔCt method (81).

**Monkey experiments**

Animals. 37 F2-bred female rhesus monkeys (Macaca mulatta; Xierxin; mean weight, 5.3 ± 0.8 kg; mean age, 5 ± 1 years) were used. Animals were housed in individual primate cages under controlled conditions of humidity (50% ± 5%), temperature (24°C ± 1°C), and light (12-hour light/12-hour dark cycles, lights on at 08:00 AM), with food and water ad libitum. Veterinarians skilled in the healthcare and maintenance of nonhuman primates supervised animal care. All reasonable efforts were made to minimize animal suffering. The use of primates was minimized by using an experimental design that permits statistically significant changes to be demonstrated with the smallest number of animals per group and the smallest number of groups, consistent with scientific rigor. Experiments followed previously published procedures (18, 22, 25, 37, 82).

**Experimental protocol.** See Supplemental Figure 1. 6 animals were kept as untreated controls (control group), and 6 received 20 mg/kg l-DOPA twice daily for 3 months (control chronic l-DOPA). The remaining 25 animals were treated daily at 9:00 AM with MPTP hydrochloride (0.2 mg/kg i.v.; Sigma-Aldrich) dissolved in saline according to a previously described protocol (83). Following stabilization of the MPTP-induced syndrome, animals received saline (MPTP group; n = 6) or 20 mg/kg l-DOPA twice daily for 3 months (MPTP chronic l-DOPA group; n = 19). 13 animals developed severe and reproducible dyskinesia, presenting choreic-athetoid (characterized by constant writhing and jerking motions), dystonic, and sometimes ballistic movements (large-amplitude flinging, flailing movements) (MPTP-intoxicated and chronically l-DOPA-treated, dyskinetic group), whereas 6 others did not (MPTP-intoxicated and chronically l-DOPA-treated, nondyskinetic group). The MPTP intoxication protocol, chronic l-DOPA treatment, clinical assessments, terminal procedure, and characterization of the extent of nigrostriatal denervation in the animals terminated before viral transfection were conducted as previously published (18, 22, 25, 37, 82). DAT binding autoradiography using [125I]-labeled N-(3-isodoprop-2-enyl)-2β-carboxymethyl-3β-(4′-methylphenyl)-nortropane (Chelatec) showed a dramatic and similar reduction (>95%) in all MPTP-treated groups compared with control animals, as published elsewhere (84).

7 dyskinetic chronically l-DOPA–treated MPTP-intoxicated animals were kept alive for further behavioral investigation of PSD-95 knockdown. Monkeys’ response to their tailored dose of l-DOPA/carbidopa (ranging 15–20 mg/kg for maximal reversal of PD symptoms) was defined prior to LV intracerebral injections. The improved Horsley-Clarke stereotactic technique used in these monkeys was described briefly above and expanded in methods. The shRNA sequences target transcription factor (signal) and codon (G) and were described previously (85). The shRNA LV plasmids were purchased from Sigma-Aldrich. shRNAs were validated based on the Genbank human PSD-95 sequence NM001365. All sequences were BLAST confirmed for specificity. Synthetic DNA encoding shRNA sequences targeting human PSD-95 were cloned into shRNA LV plasmids (pLKO.1-puro) under the control of the U6 promoter. shRNA LV plasmids were transfected into SH-SY5Y cells for 48 hours to determine the level of knockdown of gene expression by immunoblotting. The shRNA sequence used for this study was CCGGACGATCATCGCTCACTTCACTGATT. LV titers were 8.67 × 10^9 pfu/ml for PSD-95 shRNA, 1.03 × 10^9 pfu/ml for a scrambled shRNA, and 2.00 × 10^9 gfp for the control shRNA.

**Immunohistochemistry.** PSD-95 Western blotting was identical to the method described above for fresh-frozen rat tissue (n = 6 per experimental condition). Results were derived from 3 independent experiments. For transfected animals, free-floating sections were chosen directly in the injection site. Putamen was dissected from these sections, pretreated with proteinase K (Dako), and sonicated in 100 μl RIPA buffer (Sigma-Aldrich) with a cocktail of protease inhibitors (Roche). Homogenates (300 μg) were incubated with mouse anti–PSD-95 (UC Davis/NIH NeuroMab Facility) or rabbit anti-D1 (86) antibodies at 4°C overnight with 30 μl of protein G-Sepharose (GE Healthcare). Beads were washed 4 times with TBS (50 mM Tris and 150 mM NaCl) and eluted in SDS sample buffer. Coprecipitated proteins were then identified by SDS-PAGE and Western blot analysis using mouse anti–PSD-95 (UC Davis/NIH NeuroMab Facility) or rabbit anti-D1 (Sigma-Aldrich) as indicated. Signals were revealed with HRP-conjugated secondary antibodies and chemiluminescence. Results were derived from 3 independent experiments.

**Immunohistochemistry.** PSD-95 Western blotting was identical to the method described above for fresh-frozen rat tissue (n = 6 per experimental condition). Results were derived from 3 independent experiments. For transfected animals, free-floating sections were chosen directly in the injection site. Putamen was dissected from these sections, pretreated with proteinase K (Dako), and sonicated in 100 μl Laemml buffer (25 mM Tris-Cl, pH 6.8; 1% SDS; 250 mM DTT; 7.5% glycerol; 0.05% bromophenol blue), 20 μl of which was analyzed by PSD-95 and tubulin immunoblot as described above. Striatum (PSD-95) and Snc (TH; 1 every 12 sections) cryostat free-floating sections (50 μm) were incubated overnight in mouse monoclonal antibody to PSD-95 (1:2,000 in BSA; UC Davis/NIH NeuroMab Facility) and TH (1:10,000 in BSA; Millipore), respectively. Signals were revealed with the HRP anti-mouse Envision system (Dako) followed by DAB chromogen reaction (Vector Laboratories). OD of PSD-95 staining was measured in delimited striatum areas (Mercator, Explora Nova). Stereological unbiased counting of TH-positive neurons in delimited Snc

**research article**

http://www.jci.org Volume 122 Number 11 November 2012

The Journal of Clinical Investigation
was estimated with Mercator software using a Leica CTR6000 microscope (×40 objective). The optical fractionator option was used, with a probe size of 100 μm × 80 μm spaced by 600 μm × 400 μm.

**Rat striatal primary culture experiments**

**Neuronal preparation and cell transfection.** Rat striatal cultures were prepared from embryonic day 15 rat brains as previously described (87–89). All the reagents for the culture were purchased from Sigma-Aldrich and Invitrogen. Cells were plated on coverslips coated with poly-L-lysine (10 μg/ml) and laminin (10 μg/ml) at a density of 150,000 per dish. Striatal cultures were grown in Neurobasal medium (Invitrogen) and supplemented with B27 (Invitrogen), 0.5 mM glutamine, and 12.5 mM glutamate. Before seeding, coverslips were incubated with fetal calf serum–supplemented 20% Dulbecco modified Eagle medium/F12 for 1 hour at 37°C. After seeding, the cells were kept at 37°C in a 5% CO2 incubator for 10 days. At 4 days in vitro, part of the medium, together with nonadherent cells and cell debris, was removed, and culture medium with cytosine arabinofuranoside (0.5 μM) and without glutamate was added. Striatal neurons were transfected at 7 days in vitro with D1R-CFP and PSD-95–GFP or with shPSD-GFP alone using the Effectene transfection kit (Qiagen). For 4 wells, we mixed 1.2 μg of DNA with 24 μl effectene and 9.6 μl enhancer in 300 μl reaction buffer, then added the mixture to cultured neurons, which were transferred to new culture medium 10 minutes beforehand. After a 45-minute incubation, neurons were placed back in their original medium.

**Single QD tracking and surface diffusion.** As previously described (48, 59, 60, 90, 91), QDs 655 Goat F(ab')2 anti-Rabbit IgG (Invitrogen) were first incubated for 30 minutes with 1 μl polyclonal antibodies against GFP (Chemicon). Nonspecific binding was blocked by additional casein (Vector Laboratories) to the QD 15 minutes before use. Cultured neurons were then incubated for 10 minutes with precoated anti-GFP QD complexes (final dilution, 1:10,000) and mounted in a heated chamber for observation. D1R-QDs were detected using a mercury lamp and appropriate excitation/emission filters. Images were obtained with an acquisition time of 50 ms with up to 1,000 consecutive frames at an acquisition rate of 30 Hz. Signals were detected using an EMCCD camera (Quantum, Roper Scientific). QDs were followed on randomly selected dendritic regions for up to 30 minutes. QD recording sessions were processed using Metamorph software (Universal Imaging Corp.). The instantaneous diffusion coefficient D was calculated for each trajectory, from linear fits of the first 4 points of the mean square displacement versus time [MSD(t)] function, using MSD(t) = <r²>(t) = 4Dt. The 2-dimensional trajectories of single molecules in the plane of focus were constructed by correlation analysis between consecutive images using a Vogel algorithm.

**Immunocytochemistry and image analysis.** For immunostaining, surface D1R-CFP or GluN2B were specifically stained using a rabbit polyclonal antibody against GFP (1:200; Invitrogen) or a rabbit polyclonal antibody raised against GluN2B with extracellular epitope (1:1,000 for 15 minutes at 37°C in culture medium). Briefly, neurons were then fixed with 4% PFA for 15 minutes, washed, and incubated with secondary antibody anti-rabbit Alexa Fluor 568 (1:1,000 for 30 minutes; Molecular Probes). For PSD-95 and total D1R immunostaining, we used a monoclonal antibody raised in mouse anti-GFP (1:1,000 for 60 minutes at 37°C, 500 for 30 minutes at room temperature; UC Davis/NIH NeuroMab Facility) or a monoclonal antibody raised in mouse anti-D1R (1:500 for 60 minutes at 37°C, Invitrogen). Cultures were treated with a 37°C prewarmed mixture of 4% paraformaldehyde in 0.1 M PB (pH 7.4) for 20 minutes at 14°C, rinsed 3 times in PBS, and permeabilized with 0.3% Triton X-100 in PBS for 50 seconds at room temperature. The cultures were then incubated in PBS with 0.3% BSA (Sigma-Aldrich) for 60 minutes at room temperature to block nonspecific binding sites. Coverslips were placed cell-side-down onto a 200-μl drop of primary antibodies (in PBS-BSA). Cultures were washed 3 times in PBS-BSA and incubated at room temperature for 30–60 minutes with a secondary Alexa Fluor 568–conjugated goat anti-mouse IgG (1:400 in PBS for D1R, 1:1,000 for PSD-95; Invitrogen). For D1R immunostaining, coverslips were further incubated with Alexa Fluor 568–conjugated donkey anti-goat immunoglobulin G (1:400 in PBS; Invitrogen) to increase the fluorescence signal (88). The slides were then rinsed and mounted in Vectashield (Vector Laboratories), and preparations were kept at 4°C until quantification. For analysis, immunofluorescent stainings were imaged on an inverted confocal spinning-disk microscope (Leica). Fluorescence intensity was measured using Metamorph software (Universal Imaging) and corrected for background noise in the different conditions. Surface D1R was measured on n = 5 neurons (i.e., N = 21 dendritic fields) in the PSD-95 condition, and on n = 6 neurons (i.e., N = 20 dendritic fields) in the shPSD condition.

**Statistics**

Data are expressed as mean ± SEM unless otherwise specified. Analysis involving comparisons of 2 groups were performed using 2-tailed t test. Western blotting experiments involving several experimental groups were analyzed using 1-way ANOVA followed by Dunnett. Behavioral data (rating scores and locomotor activity) in rat and macaque models were analyzed using Kruskal-Wallis followed by Dunn’s multiple-comparisons test. QD experiments, reporting diffusion coefficient distribution, were analyzed using Mann-Whitney test. Analyses were completed using STATA (Intercooled Stata 6.0; Stata Corp.). A P value less than 0.05 was considered significant.

**Study approval**

All experiments were carried out in accordance with the European Communities Council Directive of November 24, 1986 (86/609/EEC). The Ethical Committee of CNRS, Région Aquitaine (Bordeaux, France), approved the rat experiments. Monkey experiments were performed in an AAALAC-accredited facility following acceptance of study design by the Institute of Lab Animal Science IACUC (Chinese Academy of Science, Beijing, China).

**Acknowledgments**

This work was supported by Agence Nationale de la Recherche grants ANR-07-MNP TRAFINLID and ANR-08-MNP-018 MCHPRIMAPARK (to B. Bloch and E. Bezard), by a Federation pour la Recherche sur le Cerveau grant (to E. Bezard), by Biothèque Primate — CNRS Life Sciences Department (to E. Bezard), by Conseil Régional d’Aquitaine (to A. Berthet, E. Bezard, L. Groc, and D. Choquet), by a postdoc fellowship from Fondation pour la Recherche Médicale (to B. Dehay), by ERC grant 232942 (to D. Choquet), and by FP7-PEOPLE-2009-RG (to B. Dehay). We thank the Bordeaux Imaging Center for technical assistance, B. Rousseau and L. Hao for animal care, M.-L. Thiolat for general technical assistance, and V. Guyonnet-Dupeyrat (Université Bordeaux Segalen Vectorology Core facility) and T. Leste-Lasserre, G. Drutel, and A. Brochard (Université Bordeaux Segalen qPCR Core facility). Université Bordeaux Segalen and CNRS provided infrastructural support.

Received for publication May 2, 2012, and accepted in revised form August 16, 2012.

Address correspondence to: Erwan Bezard, CNRS UMR 5293, IMN, Université Bordeaux Segalen, 146 rue Léo Saignat, 33076 Bordeaux cedex, France. Phone: 5575.71687; Fax: 5569.86182; E-mail: erwan.bezard@u-bordeaux2.fr.

2. Gerfen CR. The neostriatal mosaic: multiple levels of compartmental organization. J Neural Transm Suppl. 1992;36:43–59.


