**Manuscript Title Page**

**1. Manuscript Title: Identification of novel BDNF-specific corticostriatal circuitries**

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**Abstract**

BDNF is released from axon terminals originating in the cerebral cortex onto striatal neurons. Here, we characterized BDNF neurons in the corticostriatal circuitry. First, we utilized *BDNF*-Cre and Ribotag transgenic mouse lines to label BDNF-positive neurons in the cortex and detected *BDNF* expression in all the subregions of the prefrontal cortex (PFC). Next, we used a retrograde viral tracing strategy, in combination with *BDNF*-Cre knockin mice, to map the cortical outputs of BDNF neurons in the dorsomedial and dorsolateral striatum (DMS and DLS). We found that *BDNF*-expressing neurons located in the medial prefrontal cortex (mPFC) project mainly to the DMS, and those located in the primary and secondary motor cortices (M1 and M2) and agranular insular cortex (AI) project mainly to the DLS. In contrast, *BDNF*-expressing orbitofrontal cortical (OFC) neurons differentially target the dorsal striatum (DS) depending on their mediolateral and rostrocaudal location. Specifically, the DMS is mainly innervated by the medial and ventral part of the orbitofrontal cortex (MO and VO) whereas the DLS receives projections specifically from the lateral part of the OFC (LO). Together, our study uncovers previously unknown BDNF corticostriatal circuitries. These findings could have important implications for the role of BDNF signaling in corticostriatal pathways.

**Significance Statement**

BDNF is released in axons upon neuronal depolarization. Surprisingly, careful mapping

of BDNF projecting neurons in the central nervous system (CNS) has not been conducted. Using

retrograde viral strategies in combination with transgenic mice, we mapped out corticostriatal

BDNF circuits. We found that, mPFC BDNF neurons project mainly to the DMS whereas the

motor cortex and AI project to the DLS. BDNF neurons in the OFC are anatomically segregated.

Whereas the DMS receives BDNF-positive projections from the VO, the DLS mainly receives

BDNF-positive projections from the LO. Our findings could be important to the study of BDNF

in corticostriatal circuitries.

**Introduction**

Brain Derived Neurotrophic Factor (BDNF) is a member of the nerve growth factor family of neurotrophic factors, which is highly expressed in the CNS (Hofer et al., 1990; Yan et al., 1997; Kowianski et al., 2018). The majority of BDNF in neurons is stored in presynaptic dense core vesicles and is released upon neuronal depolarization (Dieni et al., 2012; Song et al., 2017). Once BDNF is released at axon terminals, it binds the receptor tyrosine kinase, tropomyosin-related kinase B (TrkB). Activation of TrkB stimulates extracellular regulated kinase 1/2 (ERK1/2), Protein kinase C (PKC) and/or phosphoinositide 3 kinase (PI3K) signaling cascades resulting in the initiation of transcriptional and translational machineries (Huang and Reichardt, 2003; Leal et al., 2014; Zagrebelsky et al., 2020). In the adult brain, BDNF plays a crucial role in synaptic and structural plasticity (Panja and Bramham, 2014; De Vincenti et al., 2019), as well as in learning and memory (Bekinschtein et al., 2014; Miranda et al., 2019).

BDNF is highly expressed in the cerebral cortex of both rodents and humans (Hofer et al., 1990; Timmusk et al., 1993; Conner et al., 1997). Studies in rodents suggest that BDNF in the cortex contributes to learning and memory paradigms (Miranda et al., 2019). For example, absence of BDNF in the prelimbic cortex (PrL) alters fear expression in mice indicating a role for consolidation and expression of learned fear (Choi et al., 2010). In the OFC, BDNF is critical for goal-directed decision making and in selecting actions based on their consequences (Gourley et al., 2013). In the motor cortex, BDNF contributes to motor learning (Andreska et al., 2020).

The cerebral cortex is also the major source of BDNF in the striatum (Altar et al., 1997; Baquet et al., 2004; Strand et al., 2007). BDNF released from cortical terminals, binds to, and activates its receptor, TrkB, in the striatum (Altar et al., 1997; Baydyuk and Xu, 2014). BDNF/TrkB signaling in the striatum has important cellular and behavioral roles (Besusso et al.,

2013; Engeln et al., 2020). For example, TrkB signaling is required to control inhibition of

locomotor behavior in enkephalin (ENK) positive medium spiny neurons (MSN) (Besusso et al.,

2013), and Lobo and colleagues provided data to suggest that BDNF/TrkB signaling in dopamine

D1 receptor expressing (D1) MSN plays a role in stereotypy behaviors (Engeln et al., 2020).

Finally, rodent studies have suggested that BDNF in corticostriatal circuitries is linked to

addiction. For example, numerous studies investigated the role of BDNF in the PFC in relation

with cocaine use (review (Pitts et al., 2016)). Specifically, cocaine exposure regulates BDNF

signaling in the PFC and in turn BDNF influences the development and maintenance of cocaine-

related behaviors (Lu et al., 2010; Gourley et al., 2013; Zhang et al., 2015; Pitts et al., 2018). In

addition, activation of BDNF signaling in the DLS keeps alcohol intake in moderation

(Jeanblanc et al., 2009; Jeanblanc et al., 2013), whereas malfunctioning of BDNF/TrkB signaling

in the corticostriatal regions promotes compulsive heavy use of alcohol and other alcohol-

mediated behaviors (Logrip et al., 2009; Darcq et al., 2015; Darcq et al., 2016; Warnault et al.,

2016; Moffat et al., 2023).

As detailed above, BDNF is crucial for cortical and striatal functions, yet careful

characterization of BDNF neurons in corticostriatal circuitry is lacking. Here, using a

combination of transgenic mouse lines together with viral-mediated gene delivery approaches,

we mapped out BDNF-expressing cortical neurons that project to the DMS and DLS.

**Methods**

**Reagents**

Mouse anti-NeuN antibody (MAB377) was obtained from Millipore (Billerica, MA). Rabbit

anti-VGLUT1 antibody (VGT1-3) was purchased from Mab Technologies (Stone Mountain,

GA). Chicken anti-GFP (A10262), donkey anti-mouse IgG AlexaFluor 594, anti-chicken AlexaFluor 488 and anti-rabbit IgG AlexaFluor 594 antibodies were purchased from Life Technologies (Grand Island, NY). Other common reagents were from Sigma Aldrich (St. Louis, MO) or Fisher Scientific (Pittsburgh, PA).

**Animals and Breeding**

Male C57BL/6J mice (6-8 weeks old at time of purchase) were obtained from The Jackson Laboratory. Male *BDNF*-Cre knockin mice, which express Cre recombinase at the endogenous *BDNF* locus, were obtained from Zach Knight, UCSF (Tan et al., 2016). Ribotag mice (ROSA26CAGGFP-L10a), which express the ribosomal subunit RPL10a fused to EGFP (EGFP-L10a) in Cre-expressing cells (Zhou et al., 2013), were purchased from The Jackson Laboratory (B6;129S4-Gt (ROSA)26Sortm9(EGFP/Rpl10a)Amc/J). Ribotag mice were crossed with *BDNF*-Cre mice allowing EGFP-L10a expression in *BDNF*-expressing cells. Mouse genotype was determined by poly-chain reaction (PCR) analysis of tail DNA.

Mice were individually housed on paper-chip bedding (Teklad #7084), under a reverse 12-hour light-dark cycle (lights on 1000 to 2200 h). Temperature and humidity were kept constant at $22 \pm 2°C$, and relative humidity was maintained at $50 \pm 5\%$. Mice were allowed access to food (Teklad Global Diet #2918) and tap water *ad libitum*. All animal procedures were approved by the university's Institutional Animal Care and Use Committee and were conducted in agreement with the Association for Assessment and Accreditation of Laboratory Animal Care.

**Virus information**

Recombinant adeno-associated virus (rAAV) retrograde EF1a Nuc-flox(mCherry)-EGFP (1 x

$10^{12}$ vg/ml), which expresses nuclear-localized mCherry by default but switches to nuclear-

localized EGFP expression in the presence of Cre (Back et al., 2019) (Addgene viral prep #

112677-AAVrg), and AAV1-pCAG-FLEX-EGFP, expressing EGFP in the presence of Cre

(Addgene viral prep # 51502-AAV1), were purchased from Addgene.

**Stereotaxic surgery and viral infection**

C57BL/6J (WT) or *BDNF*-Cre mice underwent stereotaxic surgery as described in Ehinger et al.

(Ehinger et al., 2020). Specifically, mice were anesthetized by vaporized isoflurane and were

placed in a digital stereotaxic frame (David Kopf Instruments, Tujunga, CA). A hole was drilled

above the site of viral injection. The injector (stainless tubing, 33 gauges; Small Parts

Incorporated, Logansport, IN) was slowly lowered into the target region. The injector was

connected to Hamilton syringes (10 µl; 1701, Harvard Apparatus, Holliston, MA), and the

infusion was controlled by an automatic pump at a rate of 0.1 µl/minute (Harvard Apparatus,

Holliston, MA). The injector remained in place for an additional 10 minutes to allow the virus to

diffuse and was then slowly removed. For experiments investigating BDNF-positive efferent

projections to the DS, *BDNF*-Cre animals were unilaterally infused with 0.5 µl of retro-AAV-

EF1aNuc-flox(mCherry)-EGFP targeting the DLS (anterior posterior (AP): +1.1, ML: ± 2.3, DV:

-2.85, infusion at -2.8 from bregma) or the DMS (AP: +1.1, ML: ± 1.2, DV: -3, infusion at -2.95

from bregma). For experiments investigating BDNF positive efferent projections from the OFC,

*BDNF*-Cre animals were unilaterally infused with 0.5 µl of AAV1-pCAG-FLEX-EGFP targeting

the OFC (AP: +2.58, ML: ± 1.2, DV: -2.85, infusion at -2.8 from bregma).

**Immunohistochemistry, imaging and quantification**

Following intraperitoneal (i.p.) administration of euthasol (200 mg/kg), mice were transcardially perfused with phosphate buffered saline, followed by 4% paraformaldehyde (PFA), pH 7.4. Brains were quickly removed post-perfusion and fixed for 24 hours in 4% PFA prior to cryoprotection in 30% sucrose solution for 3 days at 4°C. Brains were then sectioned to 30 µm by cryostat (CM3050, Leica, Buffalo Grove, IL), collected serially and stored at -80°C. PFA-fixed sections were permeabilized and blocked in PBS containing 0.3% Triton and 5% donkey serum for 4 hours at 4°C. Sections were then incubated for 18 hours at 4°C on an orbital shaker with the primary antibodies anti-NeuN (1/500), anti-GFP (1/1000) or anti-VGlut1 (1/1000) diluted in 3% bovine serum albumin (BSA) in PBS. Next, sections were washed in PBS and incubated for 4 hours at 4°C with Alexa Fluor 488-labeled donkey (1/500), Alexa Fluor 594-labeled donkey (1/500) antibodies in 3% BSA in PBS. After staining, sections were rinsed in PBS and were cover slipped using Prolong Gold mounting medium. Sections from rostral, rostrocaudal and caudal PFC were imaged on an Olympus Fluoview 3000 Confocal microscope (Olympus America, Center Valley PA) using manufacture recommended filter configurations, using the same parameters across mice and images. Captured images were used to quantify the number of fluorescent cells in subregions of rostral, rostrocaudal, and caudal PFC using FIJI ImageJ (NIH)(Schindelin et al., 2012). Specifically, the image of the PFC was precisely aligned to the anteroposterior (AP)-corresponding figure in the Paxinos atlas (Paxinos, 2004), using the BigWarp interface included in FIJI (J. A. Bogovic, 2016). By placing corresponding landmarks on the image and atlas, the image and the AP figure were overlayed with precision. Next, regions of interest (ROI) were traced on the overlayed image following the atlas delimitations using the polygon tool in FIJI. After adjusting the threshold (the same for each image), the number of

173  positive neurons within an ROI was automatically quantified using the counter plugin in FIJI. To

174  calculate the density of labeled neurons, the total number of labeled neurons in a region was

175  divided by the surface of the region in mm$^2$ using Fiji(Schindelin et al., 2012).

**Data Analysis**

Graphpad Prism 9 was used for statistical analysis. D'Agostino–Pearson normality test was used

to verify the normal distribution of variables. Data were analyzed using one- or two-way

ANOVA where appropriate. One-way ANOVA was followed by Tukey's multiple comparisons

test when appropriate. For two-way ANOVAs, significant main effects or interactions were

calculated, followed by Sidak's multiple comparisons test. p value cutoff for statistical

significance was set to 0.05.

**Results**

**Experimental strategy for evaluating *BDNF*-expressing neurons distribution in the PFC**

The cerebral cortex expresses high levels of *BDNF* message (Hofer et al., 1990; Timmusk et al., 1993), however, a careful analysis of *BDNF* containing neurons in the PFC has not been conducted. Therefore, we first assessed the distribution of *BDNF* expression in cortical neurons. To do so, we used a *BDNF*-Cre transgenic mouse line allowing Cre-recombinase expression only in *BDNF*-expressing cells (Tan et al., 2016), which was crossed with a Ribotag mouse line expressing GFP-fused ribosomal subunit RPL10 in the presence of Cre-recombinase (Zhou et al., 2013) (**Figure 1a**). The presence of GFP fused ribosomal subunit RPL10 (EGFP-L10a) enabled the visualization of *BDNF*-expressing cells.

**BDNF-positive neurons are detected in all prefrontal regions of the cortex**

A large number of *BDNF*-expressing neurons were detected in the PFC, in M2 and the PrL as well as in the ventrolateral OFC (VLO) (**Figure 1b-c**). In contrast, we did not detect green fluorescence in the dorsal or ventral striatum (**Figure 1d-e**), which confirms previous data indicating that striatal neurons do not express *BDNF* (Baydyuk and Xu, 2014). We then quantified the number of *BDNF*-expressing neurons in the different PFC subregions and extended frontal regions (M1 and AI) along the rostrocaudal axis (**Figure 1f**). Rostral and rostrocaudal quantification revealed that all PFC subregions exhibit the same density of BDNF-positive neurons (rostral: $735.7\pm7.5$ neurons/mm$^2$, rostrocaudal: $749.3\pm11.3$ neurons/mm$^2$). We did not find significant differences in the number of BDNF-positive neurons between rostral subregions (one-way ANOVA, Df = 3, F (3, 16) = 0.7636, p = 0.5309, n = 5) and rostrocaudal (one-way ANOVA, Df = 4, F (4, 20) = 1.888, p = 0.152, n = 5). However, caudal quantification

208     revealed a significantly lower density in BDNF-positive neurons in the MO compared to the

209     PrLCingulate area 1(Cg1) (**Figure 1f**) (one-way ANOVA, Df = 4, F (4, 20) = 3.667, p = 0.0214;

210     MO vs PrL/Cg1: p = 0.0122, n = 5).

212     **Experimental strategy for mapping of prefrontal BDNF-positive neurons projecting to the**

213     **DS**

214            The cerebral cortex is the major source of BDNF in the striatum (Altar, Cai et al. 1997,

Baquet, Gorski et al. 2004). The striatum is divided into the ventral and dorsal striatum

(Hunnicutt et al., 2016). We focused on the DS which is further divided into the lateral (DLS)

and medial (DMS) regions (Hunnicutt et al., 2016), and assessed whether *BDNF*-expressing

neurons in the PFC send projections to these two parts of the DS. To do so, we used a retrograde

viral strategy and infected the anterior part of the DLS or the DMS of *BDNF*-Cre mice with a

retrograde AAV-Nuc-flox-(mCherry)-EGFP viral construct (Back et al., 2019) (**Figure 2a**),

which enables mCherry expression in BDNF-negative neurons and EGFP expression in BDNF-

positive neurons (Back et al., 2019).

**PFC-to-DMS BDNF-positive circuits**

225            First, we assessed whether *BDNF*-expressing cortical neurons project to the DMS by

infecting the DMS of *BDNF*-Cre mice with the retrograde AAV-Nuc-flox-(mCherry)-EGFP

(**Figure 2a-b**). As shown in **Figure 2c,** the DMS contains only cell nuclei labeled in red,

corresponding to intrastriatal connections. The prefrontal regions show a high density of

retrogradely EGFP labeled neurons (**Figure 2c and g**). Specifically, the mPFC including the PrL

and Cg1 represents the major BDNF-positive output to the DMS, accounting for 42.9% of PFC-

to-DMS projecting neurons (**Figure 2d-e**). In addition, 26.3% of the BDNF-positive projections

to the DMS are coming from M2 (**Figure 2d-e**). The MO and VO represent respectively 10.6%

and 15.4% of the BDNF-positive PFC-to-DMS projecting neurons (**Figure 2d-e**). Only 4.8% of

total BDNF-positive neurons project from the LO, M1 and AI to the DMS (**Figure 2**).

**Density of *BDNF*-expressing neurons in PFC subregions that project to the DMS**

To evaluate the density of the BDNF-positive PFC neurons projecting to the DMS within

each PFC subregion, we analyzed the number of retrogradely labeled neurons per mm$^2$ (**Figure**

**2f**). We found a significantly higher density of labeled BDNF-positive neurons in the Cg1

compared to the other regions, and in the PrL compared to LO, M1 and AI (**Figure 2f**) (one-way

ANOVA, Df = 7, F (7, 32) = 22.72, p <0.0001; Cg1 vs other regions : p < 0.0001, PrL vs LO : p

= 0.0206, PrL vs AI : p = 0.0159, PrL vs M1 : p = 0.0204, n = 5). In addition, the BDNF-positive

PFC neurons project bilaterally, but predominantly in an ipsilateral manner, and the topographic

pattern of PFC *BDNF*-expressing neurons is similar in the contralateral hemisphere (**Extended**

**Data Figure 2-1**).

***BDNF*-expressing PFC neurons project to the DLS**

Next, we assessed whether *BDNF*-expressing cortical neurons project to the DLS by

infecting *BDNF*-Cre mice with the retrograde AAV-Nuc-flox-(mCherry)-EGFP (**Figure 3a-b**).

Similar to the DMS, the DLS contained only red cell nuclei, corresponding to intrastriatal

connections (**Figure 3c**). Absence of EGFP labeled neurons indicated once more a lack of *BDNF*

message in the DLS. However, we found a high density of retrogradely EGFP labeled neurons in

the prefrontal regions (**Figure 3c and g**). Specifically, the motor cortex (M1/M2), and the AI are

the main regions that send BDNF-positive projection to the DLS, representing 80.9% and 12.6%, respectively, of the overall BDNF-positive projecting neurons (**Figure 3d-e**). Fewer BDNF-positive neurons in the OFC (MO, VO and LO) and mPFC (PrL and Cg1) project to the DLS, representing together only 6.4% of the overall BDNF-positive projecting neurons (**Figure 3d-e**).

**Density of PFC-to-DLS BDNF-positive projecting neurons in the PFC subregions**

We next analyzed the density of retrogradely labeled BDNF-positive neurons in the PFC (**Figure 3f**). We observed that a higher number of labeled BDNF-positive neurons are located in M2 compared to the MO, VO, LO, PrL and Cg1 (one-way ANOVA, $Df = 7$, $F (7, 32) = 6.412$, $p < 0.0001$; M2 vs MO : $p = 0.0004$, M2 vs PrL : $p = 0.0026$, M2 vs VO : $p = 0.0004$, M2 vs LO : $p = 0.0008$, M2 vs Cg1 : $p = 0.0033$, $n = 5$) (**Figure 3f**).

**Comparison of topographical distribution of BDNF-positive PFC to DLS and DMS projecting neurons along the rostrocaudal axis**

We performed a more detailed comparison of PFC topographical patterns of DMS and DLS projecting neurons along the rostrocaudal axis (**Figure 4**). There was no significant difference between the PrL, VO, LO and MO retrogradely labeled BDNF-positive neurons at the rostral position (**Figure 4a**). Interestingly, rostrocaudal and caudal analysis revealed a specific AI-to-DLS circuit (**Figure 4b-c**). In contrast with the rostral position, retrogradely labeled BDNF-positive neurons in the PrL/Cg1 projecting to the DMS or DLS exhibit an opposite distribution, with fewer DLS projecting neurons and an increased number of DMS projecting neurons along the rostrocaudal axis (**Figure 4b-c**). We then focused on the OFC and compared the mediolateral distribution of BDNF-positive projecting neurons along the rostrocaudal axis

(**Figure 4d**). We found that labeled *BDNF*-expressing neurons in the caudal LO are projecting

significantly more to the DLS than the DMS, whereas labeled *BDNF*-expressing neurons in the

rostral and rostrocaudal part of the VO are projecting significantly more to the DMS than the

DLS (**Figure 4d**) (two-way ANOVA; interaction effect, $F_{(8, 66)} = 6.398$, $p < 0.0001$, main

effect of rostrocaudal axis, $F_{(8, 66)} = 3.177$, $p = 0.0042$, main effect of OFC subregion, $F_{(1.66)}$

$= 3.441$, $p = 0.0681$, rostral VO : $p = 0.0007$, rostrocaudal VO : $p = 0.011$, caudal LO : $p =$

$0.0028$; n = 5).

**The DS receives inputs from BDNF-positive OFC neurons**

Next, we used an anterograde viral strategy to confirm the presence of a BDNF-specific

circuit between the OFC and DS. To do so, we infected the OFC of *BDNF*-Cre mice with an

AAV-Flex-GFP virus allowing visualization of projections extended by BDNF-positive OFC

neurons (**Figure 5a-c**). As shown in **Figure 5d**, BDNF-positive projections were localized in

both the DLS and in the DMS.

Finally, to determine whether the BDNF-positive OFC neurons form synapses with DLS

neurons, we stained neurons with anti-vesicular glutamate transporter 1 (VGLUT1) antibodies

labeling the cortical glutamatergic presynaptic compartment. GFP-positive projections and

VGLUT1 were co-labeled in the DLS (**Figure 6**), suggesting that BDNF-positive glutamatergic

neurons from the OFC extend projections and form synapses with neurons in the DLS.

**Discussion**

In the present study we mapped out cortical *BDNF*-expressing neurons projecting to the DLS and DMS. We found that *BDNF*-expressing neurons located in the mPFC project mainly to the DMS, whereas the motor cortex and the AI mainly project to the DLS. Interestingly, we observed an anatomical segregation along the rostrocaudal axis of OFC BDNF-positive neurons that project to the DLS and DMS. Thus, our data define novel BDNF neural circuits connecting the PFC to the DS.

**Characterization of BDNF neurons in the PFC and the DS**

In 1990, Hofer et al. used *in situ* hybridization to characterize *BDNF* message in the brain and reported that *BDNF* mRNA is abundant in the mouse cerebral cortex (Hofer et al., 1990). In contrast, BDNF protein was detected in the striatum, even though *BDNF* mRNA was absent (Spires et al., 2004; Gharami et al., 2008). Altar et al. reported that BDNF protein found in the striatum is synthesized, anterogradely transported from the cell bodies located in the cerebral cortex and released presynaptically (Altar et al., 1997). Although more recent *BDNF* expression surveys have been conducted (Singer et al., 2018; Leschik et al., 2019; Wosnitzka et al., 2020), a careful mapping of *BDNF*-expressing neurons in cortical areas and their striatal projections has not been previously undertaken. Crossing a *BDNF*-Cre mouse line (Tan et al., 2016) with a Ribotag reporter line (Zhou et al., 2013) enabled labeling of *BDNF*-expressing neurons in the prefrontal cerebral cortex. Using this approach, which has superior accuracy compared to *in situ* hybridization and immunohistochemistry, we provide a definitive proof that a majority of neurons in prefrontal regions of the cortex express *BDNF*. Similar to what was previously

reported (Hofer et al., 1990), we did not detect *BDNF*-expressing neurons in the striatum.

Interestingly, although basal *BDNF* levels in the rodent striatum are negligible, a robust increase

in *BDNF* message in the striatum is detected in response to behaviors such as voluntary alcohol

intake (McGough et al., 2004; Jeanblanc et al., 2009) and cocaine administration (Liu et al.,

2006), as well as in response to exercise (Marais et al., 2009) and stress (Miyanishi et al., 2021).

Reconciling these potentially conflicting findings merits further investigation. One possibility is

that these stimuli increase *BDNF* expression in cortical regions projecting to the striatum.

Specifically, *BDNF* mRNA can be found in axon terminals (Lau et al., 2010), and recent studies

show that presynaptic protein translation occurs within neuronal projections (Costa et al., 2019;

Hafner et al., 2019). Thus, detectable *BDNF* in the striatum could be explained by the presence

of *BDNF* mRNA within cortical neuron projections in the striatum.

**Characterization of corticostriatal BDNF circuitries**

Our viral tracing strategy enabled us to explore BDNF-specific circuitries. For instance,

we uncovered a specific AI-to-DLS BDNF circuit. Interestingly, Haggerty et al. recently found

that stimulation of the AI-to-DLS circuit decreases alcohol binge drinking in male mice which in

turn reshapes glutamatergic synapses in the DLS from AI inputs (Haggerty et al., 2022). In

addition, we previously found that activation of BDNF/TrkB signaling in the DLS keeps alcohol

intake in moderation (Jeanblanc et al., 2009; Jeanblanc et al., 2013). Thus, it is tempting to

suggest a role for BDNF in AI-to-DLS synaptic adaptations that gate alcohol intake.

We observed that the mPFC exhibits a large number of BDNF-positive neurons

projecting to the DMS. These results are in accordance with previous behavioral studies showing

that the DMS receives projections from associative cortices such as the PrL (Friedman et al.,

2015; Hart et al., 2018; Shipman et al., 2019; Vicente et al., 2020). We also found that a high

number of BDNF-positive neurons in the motor cortex extends projections to the DLS. The

motor cortex is known for innervating the DLS (Yin and Knowlton, 2006), and Andreska et al.

recently reported that BDNF in this corticostriatal circuit is essential for motor learning

(Andreska et al., 2020).

We also show herein that OFC BDNF-positive neurons project to the DS. Interestingly

these projections are organized in a mediolateral manner. Specifically, the DMS receives

projections from BDNF-positive neurons located in the MO and VO whereas BDNF-positive

neurons projecting to the DLS are located specifically in the LO. It is known that the OFC

projects to the DS (Gremel and Costa, 2013; Gremel et al., 2016; Zimmermann et al., 2017), and

previous studies showed that the DMS receives input from the MO (Gourley et al., 2016; Green

et al., 2020). The DMS plays an important role for goal directed behavior (Vandaele et al., 2019)

and BDNF in the MO is essential to sustain goal-sensitive action in mice (Gourley et al., 2016).

Specifically, Gourley et al. showed that *BDNF* knockdown in the MO decreases behavioral

sensitivity to reinforcer devaluation. Thus, it is likely that this BDNF circuitry and perhaps

BDNF released by MO neurons within the DMS regulate goal-directed action control.

We found that the majority of LO BDNF-positive neurons project to the DLS. The

functional implication of this circuitry is unknow. Gourley and colleagues reported that BDNF in

both VLO and MO participates in goal-directed behavior (Gourley et al., 2013; Gourley et al.,

2016; Zimmermann et al., 2017), and Pitts et al. reported that inhibiting TrkB signaling in the

DLS blocks habit formation (Pitts et al., 2018). In addition, mechanistic target of rapamycin

complex 1 (mTORC1) in the VLO is involved in habitual alcohol seeking (Morisot et al., 2019).

Putting these studies together, it is plausible that BDNF in the LO neurons projecting to the DLS

plays a role in inflexible behavior.

370            Another possible role for BDNF in this corticostriatal circuit is to shape and modulate the

synaptic output of striatal neurons. Using a two-neuron microcircuit approach in primary cortico-

striatal neurons, Paraskevopoulou and colleagues showed that BDNF and glutamate co-released

from cortical projections are required to modulate inhibitory synaptic transmission of striatal

neurons (Paraskevopoulou et al., 2019). Therefore, co-release of BDNF with glutamate in the

DLS and DMS by PFC neurons may modulate inhibitory sensitivity and dendritic morphology of

striatal neurons in these PFC-DS circuits.

**BDNF-TrkB signaling in the DS**

379            Most neurons in the striatum are MSN which are divided into two subpopulations, D1

MSN and dopamine D2 receptor (D2) expressing MSN (Gerfen et al., 1990). In the adult brain,

both D1 and D2 MSN express TrkB receptors (Lobo et al., 2010). Engeln and colleagues showed

that a subset of mice congenitally lacking TrkB receptor in D1-MSN in the DS exhibit repetitive

circling behavior, suggesting a role of BDNF signaling in D1-MSN in this type of behavior

(Engeln et al., 2020). Thus, it would be of great interest to map which dorsal striatal neuronal

subtype receives inputs from cortical neuron. In addition, Nestler and colleagues provided data to

suggest that activation of BDNF-TrkB signaling in D1 versus D2 MSN triggers opposite effects

on cocaine and morphine-dependent rewarding behaviors in the nucleus accumbens (NAc) (Lobo

et al., 2010; Koo et al., 2014). Additional studies are warranted to shed a light on the nature of

cortical neurons projecting to the NAc. Furthermore, the DS is also organized into patch and

matrix compartments which have distinct connectivity and genetic signature (Martin et al.,

2019). As both of patch and matrix neurons express TrkB (Costantini et al., 1999), more studies

are required to decipher the contribution of BDNF/TrkB signaling in subpopulation of DS

neurons.

**Comparison between our findings and the literature**

Using a retrograde viral tracing, we analyzed the spatial profiles of BDNF-positive

projecting neurons in the PFC. We were able to confirm that these BDNF-positive circuits

exhibit the same medial–lateral gradient of corticostriatal projections first reported in rats and

primates (Selemon and Goldman-Rakic, 1988; Haber, 2003; Schilman et al., 2008). Although

our data are generally consistent with the excitatory corticostriatal circuits described in the

literature, several points might contribute to some discrepancies. For example, Balsters et al.

compared corticostriatal circuits between human, non-human primates and mice and found

significant differences in cortical projections to the anterior putamen and caudate body (Balsters

et al., 2020). Many previous studies  have been done in other animal models such as rat and

primate (Berendse et al., 1992; Haber et al., 2006; Schilman et al., 2008; Hoover and Vertes,

2011; Mailly et al., 2013). Neuroanatomical differences exist across species and detailed

explorations of similarities and differences between mice and other species are of importance.

Furthermore, in contrast with previous literature (Berendse et al., 1992; Hoover and Vertes,

2011), in our comparison of the topographical distribution of BDNF-positive neurons, we

observed VO-to-DMS and mPFC to-DLS BDNF-positive projecting neurons. An explanation to

some of the discrepancies between corticostriatal circuit mapping studies is the experimental

strategy used by us and others. Specifically, unlike the majority of studies that used anterograde

strategies to map corticostriatal circuits (Berendse et al., 1992; Haber et al., 2006; Schilman et

al., 2008; Hoover and Vertes, 2011; Mailly et al., 2013), we utilized a retrograde strategy in

which a retrograde AAV was used (Tervo et al., 2016). Retrograde AAV is known to enter at the

postsynaptic compartment of neurons and then be retrogradely transported to the cell body

(Tervo et al., 2016). Thus, unlike an anterograde tracer which labels the entire fibers, we

specifically labeled neurons projecting and forming synapses at the site of infection. However,

Tervo et al. have provided good evidence to support the entry of AAV retrograde at axonal

terminals (Tervo et al., 2016), additional studies will be necessary to fully rule out its ability to

infect axons of passage. Although our anterograde tracing confirmed both the sparse and dense

*BDNF*-expressing projections, anterograde tracing is not appropriate for detection of sparse

projections.

Pan et al. reported tracing data consistent with our data using a fluorescent latex

microsphere retrograde injection in mice (Pan et al., 2010). The authors reported an intense

retrograde labeling in the PrL-IL and Cg1-M2 and no labeling in the LO-AI when injecting the

retrograde tracer in the anterior DMS (aDMS). Pan et al. further found an intense retrograde

labeling in the LO-AI and Cg1-M2 when injecting the fluorescent latex microsphere in the

anterior DLS (aDLS). In addition, using a retrograde viral strategy, Green et al. also reported

neurons located in the Cg1/PrL and IL/MO projecting to the aDMS and neurons located in the AI

and M1/M2 projecting to the aDLS (Green et al., 2020).

It is important to note that our study focused specifically on the anterior part of the DS.

Interestingly, Pan et al. found no retrograde labeling in the PrL, IL, MO, LO, AI, Cg1 and M2

when they injected the retrograde tracer in posterior part of the DS (pDS) (Pan et al., 2010).

Since the pattern of projection in the striatum differs along the AP axis (Mailly et al., 2013;

Hunnicutt et al., 2016), further work is needed to decipher PFC to pDS BDNF circuits in mice.

In addition, using our BDNF-positive retrograde tracing strategy, the carefully mapped

corticostrial circuities were restricted to the DMS and DLS and therefore lack dorsocentral

inputs. Further tracing work is needed to describe the complete mediolateral PFC-to-DS

circuitries.

In summary, in this study we mapped out *BDNF*-expressing neurons in the PFC and

deciphered BDNF-specific corticostriatal circuits. Furthermore, the discovery of LO-to-DLS

BDNF microcircuitry highlights the importance for deciphering the function of BDNF in the

context of microcircuits composed by very localized neuronal ensembles.

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**Figure legends**

**Figure 1: BDNF-positive neurons are detected in the PFC**

(**a**) Experimental strategy for detection of *BDNF*-expressing neuron in the cortex. *BDNF*-Cre

mice were crossed with Ribotag mice in which GFP-fused ribosomal subunit RPL10 is expressed

in the presence of Cre recombinase, enabling the detection of *BDNF*-expressing neurons. (**b-c**)

Representative images of cortical neurons. NeuN (red) and GFP (green) co-localization shows a

high number of *BDNF*-expressing neurons in the mPFC, MC and OFC. (**d-e**) Representative

images of the striatum. Neurons in the DMS and DLS do not express *BDNF*. (**f**) Topographical

quantification of BDNF-positive neurons in the PFC along the rostrocaudal axis. One-way

ANOVA followed by Tukey's post hoc test, $*p = 0.05$, $n = 5$ mice. Scale bar is indicated on each

panel. mPFC: medial prefrontal cortex, M1: primary motor cortex, M2: secondary motor cortex,

PrL; prelimbic cortex, Cg1: Cingulate area 1, OFC: orbitofrontal cortex, VLO: ventrolateral

OFC, MO: medial OFC, DMS: dorsomedial striatum, DLS: dorsolateral striatum, FrA: frontal

association cortex.

**Figure 2: BDNF-positive neurons in the PFC project to the DMS**

(**a**) Retrograde viral strategy to map *BDNF*-expressing cortical neurons projecting to the DMS. The DMS of *BDNF*-Cre mice was injected with a retrograde AAV EF1a Nuc-flox(mCherry)-EGFP viral construct enabling nuclear-localized mCherry to be expressed by default and to switch to nuclear-localized EGFP expression in the presence of Cre. (**b, h**) Retro AAV EF1a Nuc-flox(mCherry)-EGFP was infused in the DMS, and BDNF-positive neurons distribution was examined in the PFC. (**b**) Rostrocaudal distribution of the viral infection site. (**c**) *BDNF-*

expressing neurons were identified by EGFP labeled nuclei and BDNF-negative neurons by

mCherry nuclei. Lower panel shows BDNF-negative neurons (red) in the DMS. Upper panel

shows *BDNF*-expressing neurons (green) in the PFC. (**d**) Percentage of total BDNF-positive

projecting neurons per PFC subregion. (**e**) Number of BDNF-positive neurons projecting to the

DMS per PFC subregion. (**f**) Density of BDNF-positive neurons projecting to the DMS per PFC

subregion. (**g**) Representative images of PFC along the rostrocaudal axis. Contralateral BDNF-

positive PFC projections to the DS were also quantified (**Extended Data Figure 2-1**). One-way

ANOVA, followed by Tukey's post hoc test. $^{\#}p < 0.05$, $^{\#\#}p < 0.01$ (PrL compared to other

structures), $**p < 0.01$, $***p < 0.001$ (Cg1 compared to other structures). n = 5 mice, scale bar is

indicated on each panel. MO: medial OFC, VO: ventral OFC, LO: lateral OFC, PrL; prelimbic

cortex, Cg1: Cingulate area 1, M1: primary motor cortex, M2: secondary motor cortex, AI:

anterior insular cortex, DMS: dorsomedial striatum.

**Extended Data Figure 2-1: Contralateral BDNF-positive PFC projection to the DS**

The DMS (**a**) or the DLS (**b**) of *BDNF*-Cre mice were injected with a retrograde AAV EF1a

Nuc-flox(mCherry)-EGFP viral construct and retrogradely labeled neurons in the PFC were

quantified. One-way ANOVA ($F_{(7, 32)} = 20.37$; $p<0.0001$), followed by Tukey's post hoc test.

***$p < 0.001$, * M2 compared to other structures. n = 5 mice. MO: medial OFC, VO: ventral

OFC, LO: lateral OFC, PrL; prelimbic cortex, Cg1: Cingulate area 1, M1: primary motor cortex,

M2: secondary motor cortex, AI: anterior insular cortex.

**Figure 3: BDNF-positive neurons in the PFC project to the DLS**

(**a**) Retrograde viral strategy to map *BDNF*-expressing cortical neurons projecting to the DLS. The DLS of *BDNF*-Cre mice was injected with a retrograde AAV EF1a Nuc-flox(mCherry)-EGFP viral construct enabling nuclear-localized mCherry to be expressed by default and to switch to nuclear-localized EGFP expression in the presence of Cre. (**b, h**) Retro AAV EF1a Nuc-flox(mCherry)-EGFP was infused in the DLS, and BDNF-positive neurons distribution was examined in the PFC. (**b**) Rostrocaudal distribution of the viral infection site. (**c**) BDNF-positive

neurons were identified by EGFP labeled nuclei and BDNF-negative neurons by mCherry nuclei.

Lower panel shows BDNF-negative neurons (red) in the DLS. Upper panel shows BDNF-

positive neurons (green) in the PFC. (**d**) Percentage of total BDNF-positive projecting neurons

per PFC subregion. (**e**) Number of BDNF-positive neurons projecting to the DLS per PFC

subregion. (**f**) Density of BDNF-positive neurons projecting to the DLS per PFC subregion. (**g**)

Representative images of PFC along the rostrocaudal axis. One-way ANOVA by Tukey's post

hoc test. $*p<0.05$, $**p < 0.01$, $***p < 0.001$, M2 to other structures.  n = 5 mice, scale bar is

indicated on each panel. VO: ventral OFC, LO: lateral OFC, PrL; prelimbic cortex, Cg1:

Cingulate area 1, M1: primary motor cortex, M2: secondary motor cortex, AI: anterior insular

cortex, DLS: dorsolateral striatum.

**Figure 4: Comparison of topographical distribution of BDNF-positive PFC to DLS and DMS projecting neurons along the rostrocaudal axis**

(**a-c**) Comparison of BDNF-positive PFC neurons projecting to the DMS versus DLS along the rostrocaudal axis. (**d**) Comparison of the distribution of BDNF-positive projecting neurons in the MO, VO and LO. Two-way ANOVA, followed by Sidak's post hoc test. *p<0.05, **p $< 0.01$, ***p $< 0.001$, n = 5 mice. PFC: Prefrontal cortex, DLS: dorsolateral striatum, DMS: dorsomedial

726  striatum, OFC: orbitofrontal cortex, MO: medial orbitofrontal cortex, VO: ventral orbitofrontal

727  cortex, LO: lateral orbitofrontal cortex, PrL; prelimbic cortex, Cg1: Cingulate area 1, AI: anterior

728  insular cortex.

**Figure 5: The DS receives projections from BDNF-positive OFC neurons**

(**a-b**) Anterograde AAV1-Flex-GFP virus was injected in the OFC of *BDNF*-Cre mice allowing

visualization of projections extended by BDNF-positive OFC neurons. (c-d) Representative

735     images of the VO, DLS and DMS, showing GFP-positive projections localized in the DLS and in

736     the DMS. DAPI staining is used to label nuclei. n = 2, scale bar is indicated on each panel. OFC:

737     orbitofrontal cortex, DLS: dorsolateral striatum, DMS: dorsomedial striatum, DS: dorsal

738     striatum, VO: ventral orbitofrontal cortex.

**Figure 6: Glutamatergic BDNF-positive OFC neurons form synapses with DLS neurons**

AAV-Flex-GFP virus was injected in the OFC of *BDNF*-Cre mice allowing visualization of projections in the DLS. GFP-positive projections in the DLS were stained for VGLUT1, a specific marker of cortical glutamatergic presynaptic compartment. Representative images of the DLS labeled with GFP in green and VGLUT1 in red. n = 2, scale bar is indicated on each panel.

DLS: dorsolateral striatum, OFC: orbitofrontal cortex.