

# Dichotomous parvalbumin interneuron populations in dorsolateral and dorsomedial striatum

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## Author contribution:

Patricia Monteiro: Conception or design of the work; Acquisition or analysis or interpretation of data for the work; Drafting the work or revising it critically for important intellectual content; Final approval of the version to be published; Agreement to be accountable for all aspects of the work; Boaz Barak: Acquisition or analysis or interpretation of data for the work; Final approval of the version to be published; Agreement to be accountable for all aspects of the work; Yang Zhou: Acquisition or analysis or interpretation of data for the work; Final approval of the version to be published; Agreement to be accountable for all aspects of the work; Rebecca McRae: Acquisition or analysis or interpretation of data for the work; Final approval of the version to be published; Agreement to be accountable for all aspects of the work; Ian R. Wickersham: Acquisition or analysis or interpretation of data for the work; Final approval of the version to be published; Agreement to be accountable for all aspects of the work; Guoping Feng: Conception or design of the work; Acquisition or analysis or interpretation of data for the work; Drafting the work or revising it critically for important intellectual content; Final approval of the version to be published; Agreement to be accountable for all aspects of the work.

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# **Dichotomous parvalbumin interneuron populations in dorsolateral and dorsomedial striatum**

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**Running title:** Parvalbumin interneurons in dorsal striatum

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## **Key points summary:**

- Existence of two electrophysiological dichotomous populations of parvalbumin (PV) interneurons located in the dorsal striatum.
- Striatal PV interneurons in medial and lateral regions significantly differ in their intrinsic excitability.
- Parvalbumin interneurons in dorsomedial striatum, but not dorsolateral striatum, receive afferent glutamatergic input from cingulate cortex.

# **Dichotomous parvalbumin interneuron populations in dorsolateral and dorsomedial striatum**

## **ABSTRACT**

Dorsomedial striatum circuitry is involved in goal-directed actions or movements that upon repetition become habits, encoded by dorsolateral striatum. An inability to shift from habits can compromise action-control and prevent behavioral adaptation. But although these regions seem to be clearly behaviorally distinct, little is known about their distinct physiology.

Parvalbumin (PV) interneurons are a major source of striatal inhibition and are usually considered as a homogeneous population in the entire dorsal striatum. Here, we recorded PV interneurons in dorsal striatum slices from WT male mice and suggest the existence of two electrophysiological dichotomous populations. We found that PV interneurons located at the dorsomedial striatum region have increased intrinsic excitability as compared to PV interneurons in dorsolateral region. We also found that PV interneurons in dorsomedial region, but not dorsolateral striatum region, receive short-latency excitatory inputs from cingulate cortex. Therefore, our results demonstrate the importance of considering region specific parvalbumin interneuron populations when studying dorsal striatal function.

## INTRODUCTION

Daily goal-directed actions often become habitual automated responses after consecutive repetition (Yin and Knowlton, 2004; Yin et al., 2004, 2005, 2006; Hilário and Costa, 2008; Baldan Ramsey et al., 2011; Hilario et al., 2012). Striatum function is crucial for this habit-formation and for proper psychomotor behavior such as motor control, procedural learning and behavioral switching (Yin and Knowlton, 2006; Hilário and Costa, 2008; Steiner and Tseng, 2010; Parent, 2012). Adult striatum dysfunction results in loss of action-control (Graybiel, 2008) and its dysfunction has recently been linked to OCD and ASD (Ahmari et al., 2013; Burguière et al., 2014; Monteiro and Feng, 2015, 2017; Ahmari, 2016), thickening the list of previously known classic striatum-related disorders such as Parkinson and Huntington's disease (Kreitzer and Malenka, 2008; Plotkin and Surmeier, 2015).

Parvalbumin (PV) interneurons are critical circuit modulators and are thought to be malleable throughout life in the adult brain (Plotkin et al., 2005; Kepecs and Fishell, 2014; Dehorter et al., 2015). In rodent striatum, PV interneurons represent only a small neuronal percentage (~1%) but provide prominent feedforward inhibition to medium-spiny projection neurons (MSNs) (Tepper et al., 2008; Gittis et al., 2010). Decreased numbers of striatal PV interneurons, but not MSNs, have been reported in post-mortem caudate and putamen tissue from patients with Tourette syndrome, suggesting a link between action-control and interneuron pathology in specific striatum sub-regions (Kataoka et al., 2010; Xu et al., 2015, 2016). In rodents, DMS and DLS regions are part of an homogeneous structure, lacking the anatomical segregation between caudate and putamen regions seen in primates (Reep et al., 2003; Voorn et al., 2004). However, PV expression, is more abundant laterally than medially in rodents (Kita et al., 1990; Todtenkopf et al., 2004). Given this anatomical segregation of PV expression, we asked whether striatum PV interneurons could be electrophysiologically different in DLS

versus DMS regions. Our data provides the first evidence of dichotomous physiological properties between PV interneurons located in the two striatum regions.

## **MATERIALS AND METHODS**

### **Ethical Approval**

All animal procedures were reviewed and approved by the MIT Committee on Animal Care (CAC). Only male mice were used for experiments. Detailed methods are described below.

### **Experimental animals**

PV-*Cre* mice (*Pvalb*<sup>tm1(cre)Arbr</sup>) were purchased from the Jackson Laboratory (JAX Stock No:008069) and ROSA26-*stop*<sup>flox</sup>-tdTomato cKI mice were gently provided by Dr. Fan Wang (Duke University School of Medicine). Both mouse lines have been previously described (Hippenmeyer et al., 2005; Arenkiel et al., 2011). Specific primers were designed in this study to distinguish PV-cre heterozygous from homozygous mice:

Primer 5'-GCTCAGAGCCTCCATTCCCT-3'

Primer 5'-GCTCAGAGCCTCCATTCCCT-3'

Primer 5'-CAGCCTCTGTTCCACATACTTC-3'

ROSA26-*stopflox*-tdTomato<sup>+/+</sup> mice were crossed with PV-*Cre* mice to generate PV-*Cre*<sup>+/-</sup>:ROSA26-*stopflox*-tdTomato<sup>+/+</sup> mice for electrophysiology recordings. Animals were housed on controlled environment (12h light/dark cycles, 24h temperature and humidity monitoring), 2-5 mice *per* cage and *ad libitum* food and water. All experimental procedures were reviewed and approved by the MIT Committee on Animal Care (CAC).

### **Slice preparation**

Acute striatal slices were prepared from 6-8-week-old PV-*cre*<sup>+/-</sup>:ROSA26-*stopflox*-tdTomato<sup>+/+</sup>. Animals were anesthetized by avertin intraperitoneal injection (tribromoethanol,

20 mg/ml, 0.5 mg/g body weight) and transcardially perfused with cutting NMDG-based aCSF (mM): 92 N-methyl-D-glucamine (NMDG), 2.5 KCl, 1.20 NaH<sub>2</sub>PO<sub>4</sub>, 30 NaHCO<sub>3</sub>, 20 HEPES, 25 glucose, 2 thiourea, 5 Na-ascorbate, 3 Na-pyruvate, 0.5 CaCl<sub>2</sub>·2H<sub>2</sub>O, 10 MgSO<sub>4</sub>·7H<sub>2</sub>O (~300 mOsm, 7.2-7.4 pH) (Ting et al., 2014). Following decapitation, brain was removed and coronal brain slices (300 μm) were prepared using a Vibratome 1000 Plus (Leica Microsystems, USA). Slices were recovered in cutting solution at 32–34°C for 10-15 minutes and transferred to room-temperature carbogenated regular aCSF (mM): 119 NaCl, 2.5 KCl, 1.2 NaH<sub>2</sub>PO<sub>4</sub>, 24 NaHCO<sub>3</sub>, 12.5 glucose, 2 MgSO<sub>4</sub>·7H<sub>2</sub>O, 2 CaCl<sub>2</sub>·2H<sub>2</sub>O (~300 mOsm, 7.2-7.4 pH). All slices were allowed to recover at least ≥1h prior to whole-cell recordings.

### **Electrophysiology recordings**

Slices were transferred to the recording chamber (RC-27L, Warner Instruments) and constantly perfused with carbogenated regular aCSF at 30±2°C, ~2 ml/minute rate. Borosilicate glass microelectrodes (King Precision Glass) were pulled on a P-97 horizontal puller (Sutter Instruments) and backfilled either with KGlu, CsCl or CsGlu internal (KGlu mM: 145 K-Gluconate, 10 HEPES, 1 EGTA, 2 MgATP, 0.3 NaGTP and 2 MgCl<sub>2</sub>; pH adjusted to 7.3 with KOH and osmolarity adjusted to ~300 mOsm with sucrose. CsGlu mM: 110 CsOH, 110 D-Gluconic acid, 15 KCl, 4 NaCl, 5 TEA-Cl, 20 HEPES, 0.2 EGTA, 5 Lidocaine N-ethyl chloride, 4 MgATP, 0.3 NaGTP; pH adjusted to 7.3 with KOH and osmolarity adjusted to ~300 mOsm with K<sub>2</sub>SO<sub>4</sub>. CsCl mM: 103 CsCl, 12 CsOH, 12 Methanesulfonic acid, 5 TEA-Cl, 10 HEPES, 0.5 EGTA, 5 Lidocaine N-ethyl chloride, 4 MgATP, 0.3 NaGTP, 10 Phosphocreatine, 4 NaCl; pH adjusted to 7.3 with KOH and osmolarity adjusted to ~300 mOsm with K<sub>2</sub>SO<sub>4</sub>), presenting a typical resistance around 2-4 MΩ.

Slices were visualized under IR-DIC (infrared-differential interference contrast) using a BX-51WI microscope (Olympus) and recordings were obtained after seal rupture and internal equilibrium from visually identified *tdTomato*-positive cells. All recordings were always

obtained in parallel for both cell populations (DLS-PV and DMS-PV) along the lateral-medial striatal axis within each brain slice. Only the DLS region close to the corpus callosum and DMS close to the lateral ventricles, were targeted for recordings to avoid any potential “intermediate” region. Input resistance was obtained through extrapolation from the I-V plot as well as direct measurement through -150pA hyperpolarizing current step.

Picrotoxin 100  $\mu$ M and TTX 1  $\mu$ M were added for mEPSC experiments and AP-V 50  $\mu$ M, NBQX 10  $\mu$ M and TTX 1  $\mu$ M were added for mIPSC experiments (all drugs from Tocris).

Striatal EPSCs and EPSPs were evoked by a 0.1ms stimulation step (Isoflex, AMPI) delivered at 0.05Hz frequency by a platinum iridium concentric bipolar electrode (CBAPC75, FHC) placed in cingulate cortex layers 4/5 (stimulation electrode 1-1.5mm away; stimulation intensity ranged from 0.1 to 3.0mA). Latency was measured as the time between the stimulation (electrode artifact) and the onset of the resulting EPSC. EPSPs and action potential (AP) graphically displayed traces were obtained at -50 mV by gradual increasing stimulation intensity until AP firing. Stimulation artifacts were partly removed for clarity of the EPSP figure. Summary EPSP data (onset time, peak, rise and decay time) were extracted from the largest evoked EPSP at -70 mV from pairs of PV-MSN adjacently located (same stimulation electrode location and intensity for each PV-MSN pair). Analysis of EPSCs and EPSPs was performed on averaged responses from 3-5 sweeps. PV cells were identified based on *tdTomato* signal and all presented hyperpolarized resting membrane potential (~ -85mV) and typical AP-shape upon current-injection. Putative MSNs were identified based on soma size (12-20  $\mu$ m), absence of *tdTomato* signal, typical hyperpolarized resting membrane potential (~ -85mV) and typical AP-shape upon current-injection. EPSP rise time was measured from 10% to 90% of the peak amplitude of the synaptic response. Decay time was measured by a single exponential fit. Onset of Cg-evoked DMS-EPSPs was measured as the delay between the stimulus delivery and the EPSP foot. Data was acquired using a MultiClamp 700B amplifier and a Digidata 1440A. Signals were filtered at 1-2 KHz and digitized at 10 KHz. For current-clamp recordings,

bridge balance was adjusted and pipette capacitance neutralized. Series resistance compensation ( $<20\text{ M}\Omega$ ) was performed in voltage-clamping mode. Theoretical liquid junction potential was estimated to be around  $-11\text{ mV}$  and not corrected-for.

### **Antibodies**

Mouse and rabbit anti-parvalbumin antibodies (PV235 and PV27, 1:5000 dilution) from Swant; mouse anti-DARPP32 (611520, 1:1000) from BD Biosciences; rabbit anti-Somatostatin (AB5494, 1:200) from Millipore; rabbit anti-Neuropeptide Y (T-4070, 1:250) from Peninsula Laboratories International; goat anti-ChAT antibody (AB144P, 1:500 dilution) from Millipore; chicken anti-GFP (ab13970, 1:5000) from Abcam.

### **Morphological experiments and quantification of PV immunoreactivity**

For morphological experiments Biocytin (0.3-0.5% w/v) was included in KGlu internal solution during patch clamp recordings. Recorded slices were fixed in 4 % PFA (paraformaldehyde), kept in 4 % PFA overnight at  $4\text{ }^{\circ}\text{C}$ , and then transferred to PBS. Next day slices were incubated for 2h in AlexaFluor-488 streptavidin conjugate (1:1000, Invitrogen), washed 3x in PBS and mounted for imaging. For TSA-immunohistochemistry, rabbit anti-parvalbumin antibody (Swant, PV27) was used together with Tyramide signal amplification (TSA) (Invitrogen, T-20922) according to TSA kit instructions. Quantification of PV-immunoreactivity was performed using cellSens imaging software (Olympus).

### **IHC and microscope imaging**

Mice were anesthetized by isoflurane inhalation and transcardially perfused with PBS solution followed by 4 % PFA (paraformaldehyde) fixative solution. Brains were kept in 4 % PFA overnight at  $4\text{ }^{\circ}\text{C}$ , then transferred to PBS and sectioned at 50 microns. Sections were washed 3x in PBS and permeabilized for 5 minutes using a PBS solution containing 10% MeOH, 3%



H<sub>2</sub>O<sub>2</sub>, and 0.5% NaBH<sub>2</sub>, except for ChAT immunostaining and virus experiments. Slices were again washed 3x PBS and incubated in 1.2 % Triton-X 100 for 15 minutes, followed by another 3x PBS wash. Blocking was then performed for 1h in PBS containing 2% BSA and 0.2% Triton-X. Primary antibodies were incubated overnight at 4°C for detection of the respective antigens. Next day sections were washed 3x PBS, followed by 2-4h incubation with secondary antibody (species-specific Alexa-conjugated antibody, 1:1500, Invitrogen) at room temperature. Lastly, sections were washed 3x PBS, stained for DAPI, mounted in ProlongGold (Invitrogen) or Fluoro-Gel (Electron Microscopy Sciences) and imaged using Olympus Fluoview FV1000 confocal and Olympus BX61 microscopes. Serial sections were reconstructed as 3D models using BioVis3D software (<http://www.biovis3d.com/>, Montevideo, Uruguay).

### **Surgeries and viruses**

6-8 weeks old mice were anesthetized with isoflurane, placed on a stereotactic frame and viral injections performed using a Nanoject device (Drummond scientific). For monosynaptic tracing experiments we injected ~0.2 µl of undiluted AAV2/8-synP-DIO-sT-P2A-EGFP-P2A-B19G (2.29e12 gc/ml [gc = genome copies]) at DLS or DMS (bregma coordinates for guide cannula implantation: DLS ML+2.75, AP+0.5, 2.0; DMS ML+2.0, AP+0.5, 2.3; please note that the injector/infusion cannula further extends 1mm DV). 2-3 weeks later we injected ~0.2µl of undiluted RV-ΔG-RFP(EnvA) (titer 2.75E9 i.u./ml [i.u. = infectious units]) at the same bregma location. All stereotaxic coordinates are from bregma/skull. Mice were sacrificed 7days later to observe expression pattern. AAV-synP-DIO-sTpEpG and RVΔG-RFP(EnvA) were generated by I.R.W. from MIT Genetic Neuroengineering Group (Massachusetts Institute of Technology) (Kohara et al., 2014).

### **Experimental Design and Statistical Analysis**

All statistical analyses were performed using Prism (GraphPad Software) and MATLAB Software (MathWorks). Non-normal distributions were assumed for all the datasets regardless of variance and sample size. Pairwise comparisons were performed using Mann Whitney test for unpaired data and Wilcoxon Signed Rank test for paired data comparisons, with a threshold of  $p < 0.05$  for significance. All datasets are presented as means  $\pm$  s.e.m. Further details on particular analyses can also be found on the respective figures/results section and Table 1.

## RESULTS

To study dorsal striatum PV interneurons, we started by crossing PV-*Cre* mice with ROSA26-stop<sup>fllox</sup>-*tdTomato* reporter mice, to fluorescently label PV interneurons for cell-targeted patch clamp recordings. Immunostaining of PV-*Cre:tdTomato*<sup>+</sup> striatal brain sections, showed faithful colocalization between *tdTomato* signal and PV immunostaining, indicating successful *tdTomato* expression under the control of endogenous PV promoter (Fig. 1). To further confirm correct PV interneurons labeling by this strategy, we also performed separate immunostaining protocols with other well-known striatal markers. Colocalization was not observed between *tdTomato* native signal and immunostaining for other striatal markers such as DARPP32, SOM, NPY, or ChAT, indicating a correct reporting strategy (Fig. 1).

We next examined passive and active membrane properties by performing whole-cell intracellular electrophysiology. Recordings from PV interneurons located in DMS and DLS regions showed that these populations markedly differed in intrinsic excitability (Fig. 2A-E and Table1). PV located in DMS showed higher intrinsic excitability, as represented by their leftshifted I-V curve (Fig. 2A). Input resistance obtained through extrapolation from the I-V plot as well as direct measurement through a -150 pA hyperpolarizing current step, showed significantly higher values for DMS-PV than DLS-PV (Fig. 2B) (DMS-PV mean= 43 M $\Omega$ ; DLS-PV mean= 28 M $\Omega$ ). In addition, significantly higher membrane resistance was observed for the DMS population (Fig. 2B) (DMS-PV mean= 118 M $\Omega$ ; DLS-PV mean= 85 M $\Omega$ ). Active membrane properties also showed that DMS-PV interneurons are more excitable, displaying a left-shifted I-F curve (Fig. 2C). Action potential triggering (rheobase) also required less current injection compared to DLS-PV population (DMS-PV mean= 424 pA; DLS-PV mean= 584 pA) (Fig. 2D). Resting membrane potential (DMS-PV= -86 mV; DLS-PV= -85 mV) and spike threshold (DMS-PV mean= -39 mV; DLS-PV mean= -41 mV; Fig. 2E) were not different between the two populations, and thus do not seem to contribute for the observed excitability differences.

But because differences in excitability can also arise from differences in membrane surface area, we measured whole-cell capacitance (roughly proportional to membrane surface area). Capacitance recordings confirmed smaller values for DMS-PV than DLS-PV (Fig. 2F), suggesting that the differences in excitability could be attributable, at least in part, to different membrane areas.

To explore if these differences in electrophysiology could accompany differences in morphology, we acquired morphological data from DMS-PV and DLS-PV by using TSA-enhanced parvalbumin immunostaining (Fig. 3) and by using biocytin-filled patch pipettes during electrophysiology recordings. We found that DLS-PV consistently displayed more extensive arborization and complexity than DMS-PV interneurons. This morphological observation supports whole-cell capacitance measurements, which indicated larger capacitance values for DLS-PV than DMS-PV. Quantification of PV-immunoreactivity also showed stronger detection in the more lateral regions of the striatum compared to DMS (Fig.3).

We next investigated striatal synaptic connectivity for both PV populations. Significantly higher mEPSC frequency was observed in DMS-PV interneurons as compared to DLS-PV interneurons, with no significant differences in amplitude (Fig. 2G and Table1). In addition, DMS-PV received significantly fewer inhibitory synapses than DLS-PV, with no differences in amplitude (Fig. 2H and Table1). These results could suggest the existence of two discrete PV populations with functional differences in synaptic connectivity. Given that local synapses within the striatum are inhibitory, the mEPSC frequency data suggested that either DMS-PV received more excitatory inputs from a shared afferent source with DLS-PV, or that DMS-PV received inputs from an afferent source distinct from DLS-PV.

To resolve this question at the cellular level, we decided to generate modified pseudotyped rabies-virus and performed for the first time *in vivo Cre*-dependent monosynaptic retrograde tracing specifically from striatal PV interneurons.

First, a conditional helper virus (AAV-synP-DIO-sTpEpG) was injected either into DMS or DLS region of PV-*Cre* mice. After *Cre*-mediated recombination, this AAV helper virus expressed EGFP and two proteins: TVA (receptor for the avian EnvA, necessary for subsequent rabies virus infection) and B19G (the rabies glycoprotein necessary for monosynaptic retrograde spreading) (Wall et al., 2010). Then, two weeks post injection, modified EnvA-ΔG-RFP-rabies-virus (EnvA-pseudotyped, G-deleted, RFP-rabies) was injected into the same location, allowing monosynaptic retrograde tracing from striatum PV interneurons (Fig. 4A).

Histological analysis revealed that DMS-PV interneurons receive dense afferent innervation from cingulate cortex (Cg) (Movie 1 and Fig. 4B,C,F, Fig. 5A-C). This cortical projection was not observed for PV interneurons in DLS region (Movie 2 and Fig. 4D,E,G). Both populations of PV interneurons in DMS and DLS seemed to integrate histological inputs from thalamus and GP regions.

To functionally test this identified cortical-DMS-PV projection, we placed a stimulation electrode in deep-layer Cg and recorded post-synaptic evoked responses from DMS region. Given that PV interneurons in DLS do not received inputs from Cg, we compared DMS PV interneurons with neighboring MSN-*tdTomato* negative neurons.

Stimulation of cingulate cortex evoked a reliable glutamatergic excitatory response in both DMS-MSN neurons and DMS-PV interneurons (Fig. 5D-E and Table1). The average latency of cortical-evoked DMS-EPSCs was significantly shorter in PV interneurons than DMS-MSNs, indicating a rapid cortical recruitment of striatal inhibition (Fig. 5D-E and Table1). Cortical-evoked subthreshold DMS-EPSPs had a trend for faster onset time and trend for larger amplitude in DMS-PV interneurons compared to neighboring DMS-MSNs (Fig. 5F-G and Table1). Rise time was not significantly different between MSN and PV interneurons, although DMS-EPSPs decay time had a trend for slower in MSNs (Fig. 5H-I and Table1). These observations indicated that DMS-PV interneurons are strongly enervated by cingulate cortex and cortical inputs may affect PV interneurons more strongly than MSN neurons, potentially

allowing PV interneurons to control the recruitment and spread of second messengers more efficiently in the striatum. Moreover, these findings demonstrate for the first time that, similar to medium-spiny neurons, PV interneurons also receive glutamatergic inputs from distinct cortical regions along the medial-lateral axis of the striatum.

## DISCUSSION

Our findings demonstrate the existence of two dichotomous PV interneuron populations along the medial-lateral axis of the dorsal striatum. Whole-cell recordings revealed that DMS-PV interneurons have more excitable intrinsic membrane properties as compared to PV interneurons in dorsolateral region. Given the important role of the DMS region during initial behavioral learning (Yin et al., 2005; Ragozzino, 2007), having a more excitable PV population in this region could provide means for efficient behavioral inhibition during an initial ‘trial-and-error’ learning phase. In a scenario where an unexpected outcome occurs and behavioral flexibility is required (*e.g.* early learning phase), a more excitable PV population, such as the DMS-PV population, could therefore be more easily triggered and inhibit/shape local MSNs firing to promote behavioral adaptation. On the other hand, once a task has been extensively repeated and a specific behavioral strategy has proven to be effective, it should be preserved and not easily changed (habitual behavior; DLS-mediated). In this second habitual scenario, stronger inputs would thus be needed to activate the less excitable DLS-PV interneurons and stop an ongoing habitual response.

Moreover, the fact that associative Cg inputs specifically target DMS-PV interneurons, but not PV interneurons in dorsolateral striatum, further suggests that these discrete PV populations in lateral and medial territories are part of distinct corticostriatal networks and may in fact contribute to the hypothesized parallel/competing roles of DLS and DMS in controlling behavior (Hilario et al., 2012). Our data provide the first evidence of dichotomous physiological properties between PV interneurons along the medial-lateral axis of the dorsal striatum, and highlight the importance of considering region specific parvalbumin interneuron populations when studying the dorsal striatum.

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## FIGURES

### **Figure 1. Native *tdTomato* expression shows faithful colocalization with PV staining in striatal sections from PV-cre:*tdTomato* mice.**

*Pvalb<sup>tm1(cre)Arbr</sup>*: ROSA26-*stop<sup>flox</sup>-tdTomato* mice express *tdTomato* fluorescence in striatal parvalbumin interneurons. Fluorescence images of histological sections show colocalization of native *tdTomato* expression (red) with endogenous PV immunoreactivity, but not DARPP32, SOM, NPY or ChAT.

### **Figure 2. DMS-PV interneurons have distinct intrinsic excitability and synaptic physiology from DLS-PV interneurons.**

Higher intrinsic excitability for DMS-PV is shown by their left-shifted IV curve (A) and higher input and membrane resistance (B). (C) IF-plot with representative current injection steps demonstrates increased excitability for DMS-PV interneurons compared to DLS-PV. Please note that the graph only displays current injections up to 740 pA but many DLS-PV required much higher current injections for firing. Given that above 740 pA, many DMS-PV entered in “depolarization blockade” and become inexcitable, the graph is only displayed up to that value. (D) Lower rheobase current values indicate higher excitability of DMS-PV compared to DLS-PV. (E) No difference in firing threshold. (F) DMS-PV interneurons have smaller capacitance ( $p < 0.001$ , two-sample Kolmogorov-Smirnov test). (G) Summary bar graphs and cumulative probability curves (200 events *per* interneuron) show significantly increased mEPSC frequency in DMS-PV, with no difference in amplitude. (H) Summary bar graphs and cumulative probability curves (140 events *per* interneuron) show that DMS-PV interneurons have reduced mIPSC frequency; with no difference in amplitude. Example traces are shown at the bottom for G,H. \* $p < 0.05$ ; \*\* $p < 0.01$ ; \*\*\* $p < 0.001$ ; All statistical analyses are in Table 1.

### **Figure 3. DLS-PV interneurons have more abundant parvalbumin expression and display more complex morphology than DMS-PV.**

(A) Parvalbumin interneurons display significantly higher PV intensity *per cell* in dorsolateral striatum compared to dorsomedial striatum (\*\*\* $p < 0.001$ ;  $n = 80$  DLS-PV cells and  $n = 80$  DMS-PV cells; two-way ANOVA for the left cumulative frequency curves and Kolmogorov-Smirnov test for right histogram distribution). (B) Reconstructed morphology of PV interneurons in DLS (top) and DMS (bottom), after TSA-enhanced parvalbumin immunostaining.

### **Figure 4. Cre-dependent monosynaptic retrograde tracing from striatum parvalbumin interneurons.**

(A) Adult PV-*Cre* mice are injected in the DLS or DMS with AAV expressing TVA, EGFP and rabies glycoprotein (AAV-DIO-sTpEpG) in a *Cre*-dependent manner. Two weeks later, the same mice are injected with EnvA-pseudotyped monosynaptic rabies that can only infect PV interneurons expressing TVA, and can only retrogradely spread from PV interneurons expressing rabies glycoprotein (G). One week after rabies injection, direct inputs onto striatal PV interneurons become labeled (RFP). (B, C, F) DMS-PV interneurons receive dense inputs from cingulate cortex (Cg). RFP labelled Cg-axons can be observed in coronal section. (D, E, G) DLS-PV interneurons receive inputs from motor (M1, M2), somatosensory cortex and from other PV within DLS, but not cingulate cortex (Cg). LO - lateral orbital cortex, FrA - frontal association cortex, PrL - prelimbic cortex, Cg1 - cingulate cortex, area 1, Cg2 - cingulate cortex,

area 2, M1 - primary motor cortex, M2 - secondary motor cortex, S1 - primary somatosensory cortex, S2 - secondary somatosensory cortex, RSC - retrosplenial cortex.

**Figure 5. DMS-PV interneurons receive afferent glutamatergic input from Cingulate cortex (Cg).**

(A) Cg projection neurons (RFP+) revealed by *Cre*-dependent retrograde monosynaptic labeling from DMS-PV infected interneurons (GFP+). (B) Zoomed image from square area depicted in A; Cg axons (red) projecting to DMS-PV interneurons (green) can be seen in detail. (C) White arrows show DMS-PV interneurons co-infected by AAV-sTpEpG (green) and RV- $\Delta$ G-RFP(EnvA) (red); Blue arrow shows a DMS-PV interneuron expressing only RFP, due to rabies retrograde labeling from other synaptically connected PV interneurons. (D, E) Brain slice cartoon and representative traces showing that stimulation of Cg reliably evokes post-synaptic excitatory responses (NBQX-sensitive EPSC) in DMS-PV interneurons and neighboring MSNs; Cumulative distribution of EPSC latency after Cg stimulation shows significantly shorter latencies for DMS-PV interneurons compared to MSNs. (F-I) Summary data extracted from the largest evoked subthreshold EPSP in each pair of PV-interneuron and its respective neighboring-MSN. Representative subthreshold EPSP traces are represented on the left side for both cell types. Each trace results from a gradual increase in the stimulation intensity until action-potential triggering (insets show the full action-potential trace). Statistical analyses are in Table 1.

**Movie 1. Cre-dependent monosynaptic retrograde tracing from parvalbumin interneurons in DMS.**

3D model reconstructed from serial sections showing afferent cells that project onto dorsomedial striatal PV interneurons.

**Movie 2. Cre-dependent monosynaptic retrograde tracing from parvalbumin interneurons in DLS.**

3D model reconstructed from serial sections showing afferent cells that project onto dorsolateral striatal PV interneurons.

**Table 1. Experimental Design and Statistical Analysis**

Figure	Measurement	N	Mean±s.e.m.	Statistical test and P value
Fig.2B (top)	Input resistance (MΩ)	DMS-PV=25 cells	42.76± 2.01	Mann Whitney test, ***p<0.0001
		DLS-PV=25 cells	28.21 ± 1.255	
Fig.2B (bottom)	Membrane resistance (MΩ)	DMS-PV=25 cells	118.1 0± 8.68	Mann Whitney test, **p=0.0013
		DLS-PV=25 cells	84.70 ± 7.44	
Fig. 2D	Rheobase current (pA)	DMS-PV=25 cells	424.4 ± 26.05	Mann Whitney test, **p=0.0019
		DLS-PV=25 cells	583.6 ± 36.17	
Fig. 2E	Firing threshold (mV)	DMS-PV=15 cells	-39.40± 1.17	Mann Whitney test, p=0.1624
		DLS-PV=15 cells	-41.20 ± 1.34	
Fig. 2G	mEPSC Frequency (Hz)	DMS-PV=12 cells	22.31± 1.78	Mann Whitney test, *p=0.0229
		DLS-PV=10 cells	15.49± 1.26	
	mEPSC Amplitude (pA)		15.25 ± 0.83	Mann Whitney test, p=0.1062
		13.08 ± 0.80		
Fig. 2H	mIPSC Frequency (Hz)	DMS-PV=16 cells	5.68 ± 0.49	Mann Whitney test, **p=0.0098
		DLS-PV=16 cells	8.17± 0.79	
	mIPSC Amplitude (pA)		29.66± 2.01	Mann Whitney test, p=0.1011
		24.42 ± 1.43		
Fig. 5D, 5E	EPSC Latency (ms)	DMS-PV=8 cells	5.49± 0.44	Wilcoxon Signed Rank test, *p=0.0156
		DMS-MSN=8 cells	7.04 ± 0.73	
Fig. 5F	EPSP Onset time (ms)	DMS-PV=5 cells	5.42± 0.71	Wilcoxon Signed Rank test, p=0.1250
		DMS-MSN=5 cells	6.88 ± 1.32	
Fig. 5G	EPSP Peak (mV)	DMS-PV=5 cells	7.73± 1.96	Wilcoxon Signed Rank test, p=0.4375
		DMS-MSN=5 cells	6.31 ± 1.43	
Fig. 5H	EPSP Rise time (ms)	DMS-PV=5 cells	3.58± 0.41	Wilcoxon Signed Rank test, p=1.0000
		DMS-MSN=5 cells	3.67 ± 0.52	
Fig. 5I	EPSP Decay time (ms)	DMS-PV=5 cells	19.22± 1.64	Wilcoxon Signed Rank test, p=0.0625
		DMS-MSN=5 cells	27.45± 1.43	











