

Abstract

 PV^{Cre} mice combined with AAV-FLEX vectors allowed efficient and specific targeting of PV⁺ interneurons in the striatum. However, diffusion of viral particles to the globus pallidus 28 caused massive transduction of PV⁺ projection neurons and subsequent anterograde transport of the transgene product to the subthalamic nucleus and the substantia nigra pars reticulata. Different AAV serotypes (1 and 9) and promoters (CBA and human synapsin) were evaluated. The combination of AAV1, a moderate expression level (human synapsin promoter) and a precise adjustment of the stereotaxic coordinates in the anterior and 34 dorsolateral part of the striatum were necessary to avoid transduction of PV^+ GP projection neurons. 36 Even in the absence of direct transduction due to diffusion of viral particles, GP PV^+ projection neurons could be retrogradely transduced via their terminals present in the dorsal 38 striatum. However, in the absence of diffusion, GP-Str PV+ projection neurons were poorly or not transduced suggesting that retrograde transduction did not significantly impair the 40 selective targeting of striatal PV^+ neurons. 41 Finally, a prominent reduction of the number of striatal PV⁺ interneurons (about 50%) was evidenced in the presence of the Cre recombinase suggesting that functional effects of AAV-

43 mediated transgene expression in PV⁺ striatal interneurons in PV^{Cre} mice should be analyzed with caution.

Introduction

 AAV vectors have gained increasing interest for gene therapy of neurological diseases (Kantor, McCown et al. 2014) (Hocquemiller, Giersch et al. 2016) as well as for functional studies in neuroscience research (Bedbrook, Deverman et al. 2018). Targeting specific neuronal subpopulations is a rapidly growing field allowing the manipulation of neuronal circuits using optogenetics (Hunnicutt, Jongbloets et al. 2016) or chemogenetics

 (Woloszynowska-Fraser, Wulff et al. 2017), the modeling of disease using human disease- causing transgenes (Grames, Dayton et al. 2018) as well as next-generation targeted gene therapy (Chtarto, Bockstael et al. 2013) (Dalkara, Byrne et al. 2013) (Vormstein-Schneider, Lin et al. 2020). Thanks to their ability to be axonally transported (Castle, Gershenson et al. 2014), AAV vectors are also useful for anatomical tracing (Zingg, Chou et al. 2017).

 The cell-type specificity of AAV-mediated transgene expression is dependent on the viral capsid and the regulatory elements driving transcription. Numerous vectors selectively targeting neurons or glial cells have been described. Selective gene expression into neurons can be achieved using pan-neuronal promoters such as the neuron-specific enolase (Klein, Meyer et al. 1998, Klein, Hamby et al. 2002) or the synapsin (Kugler, Lingor et al. 2003, Shevtsova, Malik et al. 2005) (Dashkoff, Lerner et al. 2016) (Nieuwenhuis, Haenzi et al. 2020) promoters. Specific targeting of excitatory neurons has been described using the calmodulin kinase II (CaMKIIα) promoter (Kim, Kim et al. 2015, Watakabe, Ohtsuka et al. 2015). Oligodendrocytes have been targeted using the myelin-basic protein promoter (Chen, McCarty et al. 1998, Chen, McCarty et al. 1999), astrocytes using assembled fragments of the glial fibrillary protein promoter (GFA) (Drinkut, Tereshchenko et al. 2012) (Meunier, Merienne et al. 2016) (Pignataro, Sucunza et al. 2017) (Dashkoff, Lerner et al. 2016) (Dashkoff, Lerner et al. 2016) and microglia using the F4/80 or the CD68 promoter (Rosario, Cruz et al. 2016). In some cases, the cellular specificity is obtained independently of the chosen capsid. For example, the GFA promoter drives expression mainly in astrocytes, when combined with AAV5 (Drinkut, Tereshchenko et al. 2012), AAV6 (Dirren, Towne et al. 2014) AAV8 (Pignataro, Sucunza et al. 2017), AAV9 (Dashkoff, Lerner et al. 2016) or AAV- D/J capsids (Jolle, Deglon et al. 2019). In contrast, the combination of a cell type-specific promoter and a capsid variant was necessary to transduce microglial cells (Rosario, Cruz et al. 2016). Finally, brain endothelial cell-specific targeting was obtained using a novel capsid 77 variant combined with a non-specific promoter (chicken β -actin promoter fused to cytomegalovirus enhancer sites) (Korbelin, Dogbevia et al. 2016).

 The development of vectors selectively targeting neuronal subpopulations is coming of age. Some examples are: GluA4-AAV a capsid variant designed to attach to glutamate receptor 4 (GluA4), selectively expressed by parvalbumin-positive (PV+) interneurons (Geiger, Melcher et al. 1995), combined with the SFFV (spleen focus-forming virus) promoter (Hartmann, Thalheimer et al. 2019); AAV-mDlx a vector targeting interneurons using mDlx transcriptional regulatory elements combined with AAV9 (Dimidschstein, Chen et al. 2016) or AAV5 (Lee, Vogt et al. 2014) capsids; AAV-TH which is targeting dopaminergic neurons thanks to the use of a fragment of the tyrosine hydroxylase promoter (Stauffer, Lak et al. 2016) and AAV PHP.eB with hybrid promoters containing specific enhancers targeting 88 PV- and VIP- interneurons (Vormstein-Schneider, Lin et al. 2020).

 With the advent of single-cell RNA sequencing (Munoz-Manchado, Bengtsson Gonzales et al. 2018) (Gokce, Stanley et al. 2016) and methods for mapping chromatin accessibility (Buenrostro, Wu et al. 2015), new synthetic promoters will eventually be identified, which will further refine these molecular tools (Juttner, Szabo et al. 2019).

 Targeting specific neuronal subpopulations can also be achieved using a combination of Cre-driver mice (Hippenmeyer, Vrieseling et al. 2005) or rats (Liu, Brown et al. 2016) expressing the Cre recombinase under the control of a cell-type specific gene and an AAV vector harboring an inverted ORF flanked by 2 pairs of Cre recognition sites positioned so that expression occurs only when the Cre protein is present (Saunders, Johnson et al. 2012, Saunders and Sabatini 2015).

 In the striatum, projection neurons, also called medium-sized spiny neurons, relay motor output, expressing D1R- or D2R-type of dopamine receptors which, in response to dopamine respectively activate (D1R) or inhibit (D2R) efferent structures of the motor loop (Surmeier, Ding et al. 2007). The striatum also contains several classes of interneurons among which the (PV+) fast spiking and the cholinergic neurons which are thought to coordinate the activity of the projection neurons (Gritton, Howe et al. 2019).

 In the present study, we have focused on the targeting of PV+ interneurons of the dorso-lateral striatum which are key to the control of the sensorimotor striatum (Lee, Holley et al.

107) using PV^{Cre} driver mice (Hippenmeyer, Vrieseling et al. 2005).

Limitations of the AAV-FLEX/Cre driver mice targeting system have been previously

described: i) off-target expression in cells not expressing Cre (Fischer, Collins et al. 2019), ii)

expression of Cre in only a part of the targeted neuronal population (Saunders, Huang et al.

2016), iii) Cre recombinase-induced cellular abnormalities (He, Marioutina et al. 2014).

We show here that, despite the efficient and specific cellular targeting offered by the Cre-

lox system, the AAV delivery parameters have to be precisely adjusted to selectively target

the striatum with exclusion of the adjacent globus pallidus containing a high density of PV+

115 projection neurons. Furthermore, $PV⁺$ striatal interneurons were decreased by the Cre-

recombinase. These data suggest that functional effects of AAV-FLEX transgene expression

117 in PV⁺ striatal interneurons in PV^{Cre} mice should be analyzed with caution.

Material and Methods

Animals

121 Pvalbm1(*Cre*)Arbr (PV^{Cre}) mice [\(www.jax.org:](http://www.jax.org/) 008069) (Hippenmeyer, Vrieseling et al. 2005) in which a IRES-Cre-polyA cassette was introduced in the 3'UTR region of exon 5 of the PV gene, were genotyped using the following primers: Cre-forward, 5' GCG GTC TGG CAG TAA AAA CTA TC 3'; Cre-reverse, 5' GTG AAA CAG CAT TGC TGT CAC TT 3'; PVexon5 forward, 5' CAG AGC AGG CAT GGT GAC TA 3'; PVexon5 reverse, 5' CCA TTC GCC ATT AGT CTG GT 3. PCR conditions were: 4 min at 94°C followed by 25 cycles of 94°C for 30 127 sec, 60°C for 1 min, 72°C for 1 min. 128 Eleven weeks-old homozygous PV^{Cre} mice and wild-type (WT) C57Bl6-Ola-Hsd mice (En

Vigo) of both sexes were used for all experiments.

Plasmids

- AAV pCAG-FLEX-eGFP-WPRE (Plasmid #51502) and AAV phSyn1(S)-FLEX-eGFP-WPRE
- (Plasmid #51504) (Oh, Harris et al. 2014) were obtained from AddGene
- [\(http://www.addgene.org\)](http://www.addgene.org/).

pAAV2/1 and pAAV2/9 were provided by the Penn Vector Core (Philadelphia,

- Pennsylvania). pAd-helper was purchased from Stratagene (La Jolla, California),
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AAV production

- The 3 viruses used in this study were produced by triple transfection of HEK-293T cells (30
- 140 (10cm) plates; 5.0 X 10 6 cells per plate). The AAV helper plasmids; pAAV2/1 or pAAV2/9,
- expressing the AAV viral genes, were co-transfected with an adeno-helper plasmid (pAd-
- helper), expressing the adenoviral genes required for AAV replication and encapsidation
- together with the vector plasmids; AAV-phSyn1- FLEX -eGFP-WPRE or AAV-pCAG-FLEX-
- eGFP-WPRE in a 2:3:5 molar ratio. Fifty hours post-transfection, cells were harvested by
- low-speed centrifugation, medium was discarded, and cells were resuspended in Tris 50 mM
- 146 pH 8.5, NaCl 0.1M, EDTA 1mM and kept at -20°C. After five freezing/thawing cycles at -
- 20°C/37°C, the cell lysate was centrifuged 20 min at 11,000 rpm. The supernatant was
- recovered and treated with benzonase (50 units/ml, Sigma) for 30 min at 37°C and
- centrifuged again 20 min at 11,000 rpm to eliminate residual debris. The viruses were further
- purified by iodixanol gradient and microconcentrated as described previously (Zolotukhin,
- Potter et al. 2002).
- Viral genomes (vg) were titrated by quantitative polymerase chain reaction using universal primers located in the viral ITR sequence as previously described (Aurnhammer, Haase et 154 al. 2012). Titers were 1.72 x 10¹⁴ vg/ml for AAV1-hsyn- FLEX -eGFP, 1.03 x 10¹⁴ vg/ml for 155 AAV9-hsyn- FLEX -eGFP and 5.24 x 10¹³ vg/ml for AAV9-CBA- FLEX -eGFP.

Stereotaxic injections

 Adult mice (11 weeks-old) were used for unilateral intrastriatal injections. Briefly, the animals were anesthetized with a mixture of ketamine (100 mg/kg, Ketasol, Graeub AG) and xylasine

 (10 mg/kg, Rompun, Bayer). Injections were made according to coordinates defined by "The 161 Mouse Brain in stereotaxic coordinates, 3rd edition, Franklin, K.B.J. and Paxinos, G. AcademicPress, 2007" using a Kopf stereotaxic apparatus (David Kopf, Tujunga, California). 163 Viral particles diluted in 1 μ l of D-PBS (Biowhittaker, Lonza) were infused in the striatum, using a 34G needle at different coordinates (see Table 1). After injection, the needle was left in place for 5 min in order to allow diffusion of the viral suspension in the parenchyma. The needle was then slowly removed. Animals were maintained in a 12:12 hrs light-dark cycle with free access to food and water.

 Experimental procedures were approved by the "Affaires vétérinaires" of the Canton de Vaud" (Authorization n°VD3400).

Brain collection and immunohistochemistry

 Two weeks after viral injection, mice were euthanized with an overdose of pentobarbital (30 mg/kg in 0.9% NaCl). A 4% paraformaldehyde (PFA) solution at pH 7.4 in phosphate buffer saline (PBS, Bichsel AG) was freshly prepared before use. The mice were transcardially perfused consecutively with a PBS solution at pH 7.4 and with the ice-cold 4% PFA solution. Brains were collected and post-fixed in 4% PFA overnight at 4°C. Consecutive incubations of 24h in 20% and 30% sucrose solutions were performed to cryoprotect the brains which were then slowly frozen by consecutive immersions in 2-methyl-butane at -10°C and -20°C and finally stored at -80°C. A cryostat (Leica Biosystems, CM1850) was used to collect 25μm-thick coronal sections which were stored in an anti-freeze solution (glycerol 25%, ethylene glycol 30% and Na-phosphate buffer 50mM) at -20°C. The following antibodies were used to stain PV+ cells: guinea pig anti-PV (1:1000, cat. #195004, Synaptic Systems, Göttingen, Germany), biotinylated goat anti-guinea pig (1:200, cat. #BA-7000, Vector Laboratories, Burlingame. USA) and Cy3‐conjugated streptavidin (1:300, cat. #016-160-084, Jackson ImmunoResearch Laboratories, West Grove – USA). Free floating sections were stained as follows. Sections were washed 3 times for 10 min in Tris-buffered saline (TBS, 10mM Tris pH 7.6 and 0.9% NaCl) at room temperature (RT). Then, they were incubated 1h

 at RT in a blocking solution composed of 5% bovine serum albumin (BSA) in THST buffer (50mM Tris pH 7.6, 0.5M NaCl and Triton X-RT100 0.5%). Afterwards, they were incubated overnight at 4°C with the primary antibody in a THST solution containing 1% BSA. The second day, sections were first washed 3 times for 10 min in TBS at RT and then incubated for 1h at RT with the secondary antibody in THST. Sections were then washed 3 times for 10 min in TBS and incubated in the dark with the streptavidin conjugate for 1h at RT in THST. Finally, the sections were washed 3 times in PBS for 10 min at RT. The sections were mounted on microscope slides and covered with Vectashield mounting medium (Vector Laboratories, Burlingame, USA).

Image acquisition and quantifications

 Whole slide images were taken with a Zeiss Axioscan Z.1 slide-scanner (Carl Zeiss Microscopy, Germany) using a Plan-Apochromat 10x/0.45 or a Plan-Apochromat 20x/0.8 objective. All the images were taken with an Orca-Flash 4.0 V2 digital CMOS camera. 16 bits images were obtained. Cy3 was excited at 553 nm with a 555/30 nm LED at 50% power. A beam splitter at 568nm was used. The detection range was 578-640nm. eGFP was excited at 493nm with a 469/38 nm LED at 20% of power. A beam splitter at 498 was used. Detection range was 507-546nm. Confocal images were taken with a confocal microscope Zeiss LSM 800 (Carl Zeiss Microscopy, Germany) equipped with a 3x GaAsp detector. All images were collected with a Plan Apochromat 20x / 0.8 DIC II objective with a pinhole set at 36 µm. 8 bits resolution images have been obtained by bidirectional scanning and 4x averaging. Cy3 was excited with a 561 nm laser at 0.03% of power. 739V of master gain was applied and the detection range was 566-628 nm. eGFP was excited with a 488nm laser at 0.03% of power. 750V of master gain was applied and the detection range was 410- 546 nm. Controls with single-fluorescence were performed. No interference between red and green fluorescence was observed.

Images were processed using Zen Blue 2.3 (Carl Zeiss Microscopy, Germany) and

215 ImageJ/Fiji. Z-stacks were acquired on a 25 μ m thickness. Shown confocal images in Fig. 1,

2,5 and 6, are 2D maximal intensity projections of 35 images. .

Cells were manually counted on 2D maximal intensity projections of confocal images

transformed into the tiff format. Cells were counted by a blind observer using the Multi-point

219 tool in ImageJ. For evaluating the specificity, GFP⁺, PV⁺, and double-labeled cells were

counted. For each animal, five brain sections were selected, and three pictures were taken

for each section. To evaluate the efficacy in the dorsal striatum, the striatum was divided in

222 two by a horizontal line, and GFP⁺, PV⁺, and double-labeled cells were counted in the upper

part.

For the quantification of native GFP fluorescence intensity of individual cells (Fig. 6),

confocal images were used and the cells were delineated using the "free-hand" selection tool

of the ImageJ software. The mean fluorescence intensity was recorded for each individual

cell. Area with a similar surface in non-transduced area of the GP or striatum were measured

and the obtained values substracted.-

229 In order to evaluate Cre recombinase effect on PV^+ cells, WT and PV^C^+ mice were

230 compared. For each AAV-FLEX virus, three WT and three PVCre successfully injected

animals were available. The number of PV⁺ cells in the non-injected hemisphere was

counted on a section from the same region for each animal and divided by the surface in

233 mm². The images were transformed in .jpeg format at a resolution of 300dpi and countings

were performed by a blind observer using the Multi-point tool in ImageJ.

235 To determine viral toxicity the number of PV^+ cells of the injected hemisphere was compared to the corresponding area in the non-injected hemisphere. In order to count cells in the

whole striatum, images were acquired with the slide scanner. The images were transformed

in .jpeg at a resolution of 300dpi format and cells were counted by a blind observer using the

Multi-point tool in ImageJ.

Statistical analysis

 Data analyses and the creation of graphs were performed using GraphPad Prism 8 software (San Diego, CA) for Windows. Comparisons were performed by One-way ANOVA followed by Tukey's multiple comparisons test (Table 2), two-way ANOVA followed by Sidak post hoc test (for more than 2 groups) or by Student t-test (2 groups). Results were expressed as 246 mean \pm SD and statistical significance was established for a p value \leq 0.05.

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Results

Efficiency and cellular specificity of transgene expression in striatal PV⁺ interneurons 251 mediated by AAV2/9-FLEX vectors in PV^{Cre} mice.

252 AAV-FLEX vectors and PV^{Cre} mice were used to target transgene expression into striatal 253 PV⁺ interneurons. Since PV expression was weaker in the striatum than in other brain regions (for example in the cerebral cortex; see Suppl. Fig. 1), we used AAV-FLEX vectors 255 (Oh, Harris et al. 2014) with strong promoters; the non-specific CMV/chicken β -actin (CBA) promoter (Burger, Gorbatyuk et al. 2004) (Klein, Hamby et al. 2002) or the neuron-specific human synapsin promoter (hsyn) (Shevtsova, Malik et al. 2005) combined with the "Woodchuck hepatitis virus post-transcriptional regulatory element" (WPRE) which enhances mRNA stability and protein synthesis (Klein, Hamby et al. 2002) (Oh, Harris et al. 2014). In order to further maximize transduction efficiency, the vectors were encapsidated into AAV9 serotype (Cearley and Wolfe 2006) (Klein, Dayton et al. 2008).

262 The vectors (7.9 X 10⁹ vg in 1 μ I) were injected in the striatum of homozygous PV^{Cre} mice

using various coordinates (see Table 1). As a negative control, the vectors were also

injected in WT C57/Bl6 mice.

Table 1: Summary of the coordinates sets and of the number of animals used for each AAV-FLEX virus.

269 GFP⁺ cells were observed in the striatum with a distribution corresponding to the expected

AP: anterio-posterior, ML: mediol-lateral, DV: dorso-ventral, GP: globus pallidus. *These 4 mice were not included in the analysis of the efficiency and specificity of AAV2/1-hsyn-FLEXeGFP vector due to a drastic loss of PV-expressing striatal cells in this littermate precluding statistical analysis (see below and Suppl. Fig.2).

285 the Cre recombinase. The GFP⁺ cells which were not labeled by anti-PV antibodies could be

286 due to a low, undetectable PV expression in these neurons or to a potential toxicity of the

287 Cre recombinase (see below).

288

Figure 1. Efficiency and specificity of AAV2/9-FLEX-mediated PV⁺ 289 **striatal interneurons transduction in PVCre** 290 **(+/+) mice**

PVCre (+/+) mice were injected with AAV2/9-CBA-FLEX-eGFP (A,B,C,D) or AAV2/9-hsyn-FLEX-eGFP (E,F,G,H) vectors. A & E, Distribution of GFP⁺cells in the striatum (Axioscan 20-fold). Confocal pictures showing co-localization (D,H) of native GFP fluorescence (B,F) with parvalbumin staining (C,G). Arrows show cells expressing GFP but without detectable PV staining. Panel A surrounded area: transduced area in the cortex. Control wild-type mice injected with the same vectors did not harbor GFP⁺ cells (data not shown).

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Table 2: Percentage of PV/GFP double-labeled cells among total GFP⁺ cells (specificity) or total PV⁺ 294 **cells (efficiency) in the dorsal striatum**

Data were analyzed using one-way ANOVA followed by Tukey's multiple comparisons test. Difference 296 between vectors were not significant for specificity. For efficiency, AAV2/1-hsyn-FLEX-eGFP was

297 significantly lower than AAV2/9-hsyn-FLEX-eGFP (p=0.0380)

298

299 *Transduction of contiguous and axonally-connected brain regions*

- 300 Examination of the whole brains revealed GFP⁺ cells in the globus pallidus (GP) (Fig. 2E)
- and in the cerebral cortex (CTX) (Fig. 2A) in a large proportion of the animals. These data
- suggest that viral particles diffused to these neighboring regions which contain a higher
- 303 density of PV⁺ neurons as compared to the striatum (Saunders, Huang et al. 2016).
- The localized cortical transduction was probably due to a reflux of the viral suspension along
- the needle and could be avoid using anti-reflux needles (Vazquez, Hagel et al. 2012,
- Casanova, Carney et al. 2014, Lueshen, Tangen et al. 2017).
- 307 The majority of GFP⁺ cells also expressed PV in the cerebral cortex (Fig. 2 B-D) as well as in the GP (Fig.2 F-H).
- Preferential transduction of the GP after injection of AAV vectors in the striatum has been
- previously reported (Tenenbaum, Jurysta et al. 2000) and was suggested to reflect a
- preferential diffusion of the viral particles along the vessels driven by the perivascular pump
- (Hadaczek, Yamashita et al. 2006).
- Several stereotaxic coordinates were tested in order to reduce diffusion to extra-striatal
- areas (see Table 1). The anteroposterior coordinate varied between +1.0 and +1.2, the
- dorsoventral coordinate varied between -2.75 and -3.0 and mediolateral coordinate was set
- at -1.8. For both promoters, the majority of the mice harbored a widespread transduction of
- the GP (see Table 1). With none of the coordinates was the GP transduction avoid in all

120 • Figure 2: Efficient transduction of PV⁺ neurons in globus pallidus and cerebral cortex.

PV^{Cre} (+/+) mice were injected with AAV2/9-CBA-FLEX-eGFP (A-D) or AAV2/9-hsyn-FLEX-eGFP (E-H) into the right striatum. GFP⁺ cells were present in the GP (E) and cortex (A). Co-localization (D,H) of native GFP fluorescence (B,F) with parvalbumin staining (C,G). (A, E) have been acquired with a Zeiss Axioscan Z.1 (Carl Zeiss Microscopy, Germany) using a 20x magnification and (B-D & F-H) have been acquired using Zeiss LSM 800 (Carl Zeiss Microscopy, Germany) with a 20x magnification.

321

- 322 Anterograde transport of GFP from GP PV⁺ neurons projecting to the SNr and STN
- 323 GP contains 40% of PV⁺ neurons which project to the subthalamic nucleus (STN), to the
- 324 substantia nigra pars reticulata (SNr) or to the striatum (Saunders, Huang et al. 2016).
- 325 Consistently, in animals with GP transduction, GFP⁺ fibers were evidenced in the STN (Fig.
- 326 3 A & B) and in the SNr (Fig. 3 C & D).

Figure 3. Anterograde transport of GFP in fibers of GP PV⁺ neurons projecting to the 329 **subthalamic nucleus and to the substantia nigra pars reticulata.**

PVCre (+/+) mice were injected with AAV2/9-hsyn-FLEX-eGFP into the right striatum. In mice with a GP transduction, GFP⁺ fibers were detected in the subthalamic nucleus (STN) (A & B) and in the substantia nigra reticulata (SNr) (C & D). B and D, enlargement of the GFP+ area in the STN (A) and SNr (C), respectively.

- 331 In the striatum, it was not possible to distinguish GFP⁺ fibers originating from striatal
- 332 interneurons and pallido-striatal projection neurons. A typical pattern of pallido-striatal fiber
- 333 tracts was evidenced in WT mice by staining with an anti-PV antibody (Fig.4 A & B). In PV^{Cre}
- 334 mice in which striatal AAV2/9 injection resulted in transduction of GP PV⁺ neurons, fiber
- 335 tracts with the same pattern were $GFP⁺(Fig.4 C & D)$, suggesting that, as expected they
- 336 originate from the pallido-striatal projection neurons.
- 337

139 Figure 4 Anterograde transport of GFP in fibers of GP PV⁺ neurons projecting to the striatum

(A) PV staining (in red) of a wild-type non-injected mouse. (B) Dorsal striatum PV⁺ interneurons and fibers from pallidal PV+ projection neurons. C) Mice with a massive GP transduction show pallido-striatal GFP+ fibers in the dorsal striatum (D). (B) Enlargement of the rectangle delineated in (A); (D) enlargement of the rectangle delineated in (C).

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- *Efficiency and cellular specificity of transgene expression in striatal PV⁺ interneurons*
- 342 *mediated by AAV2/1-FLEX vectors in PV^{Cre} mice.*
- Since AAV9 vectors diffusion to the GP and to the CTX hindered the selective targeting of
- the striatum, we used the AAV1 serotype which has previously been shown to allow region-
- targeted striatal transduction in rats (Burger, Gorbatyuk et al. 2004) (Bockstael, Chtarto et al.
- 346 2008). In mice, however, a moderate number of GFP⁺ cells in the GP were reported after
- intrastriatal injection of AAV2/1 vectors (Taymans, Vandenberghe et al. 2007).
- Furthermore, since the CBA promoter appeared stronger than the hsyn promoter (Fig.1), a
- higher level of GFP expression was obtained in cells distant from the injection site (e.g. note
- the prominent cortical transduction in Fig.1 A).

352 coordinates were evaluated to avoid GP transduction. Anteroposterior coordinates varied 353 between +1.0 and +1.2 and dorsoventral coordinates between -2.75 and -3.0. The 354 mediolateral coordinates were set at -1.8. A total of 5 mice were tested. GP transduction 355 could be avoided only with the most anterior and dorsal coordinates (AP=+1.2; ML=-1.8; 356 DV=-2.75) (n=3) (Fig. 5A). 357 The efficiency of PV⁺ interneurons transduction, evaluated as the number of double-labeled 358 PV+/GFP+ cells (Fig. 5 D), relative to the total number of PV+ cells (Fig.5 C) was approx.. 359 93%) (Table 2). The specificity of the targeting, evaluated as the number of double-labeled 360 PV+/GFP+ cells relative to the total number of GFP+ cells (Fig. 5 B), was higher than approx. 361 89% (Table 2). As for AAV9, injection into wild-type mice did not result in GFP⁺ cells. 362 In order to confirm on a larger number of animals that the established stereotaxic 363 coordinates (AP=+1.2; ML=-1.8; DV=-2.75) allow to avoid AAV1 viral particles diffusion to 364 the GP, 4 additional mice, taken from a different littermate of $PV^{Cre}(+/+)$ mice, were injected 365 with AAV2/1-hsyn-FLEX-eGFP. However, in this littermate, the loss of PV⁺ cells in the 366 striatum was aggravated and only few PV^+ and GFP $^+$ cells were observed, precluding a 367 relevant statistical analysis (Suppl.Fig. 2A). However, despite an apparently normal amount 368 of PV⁺ cells in the GP, no GFP⁺ cells were observed (Suppl.Fig. 2B).

351 Therefore, the hsyn promoter was selected for further experiments. Several stereotaxic

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Figure 5 Efficiency and specificity of AAV2/1-FLEX-hsyn-mediated PV⁺ 370 **striatal interneurons transduction in PVCre** 371 **(+/+) mice**

PV^{Cre} (+/+) mice were injected with AAV2/1-hsyn-FLEX-eGFP vectors. (A) Distribution of GFP⁺ cells in the striatum (Axioscan 20-fold). Confocal pictures showing co-localization (D) of native GFP fluorescence (B) with parvalbumin staining (C). Control wild-type mice injected with the same vectors did not harbor GFP⁺ cells (data not shown)

373 *Retrograde transduction of pallido-striatal neurons by AAV-FLEX vectors?*

374 Since vectors were injected in the dorsolateral striatum in which pallido-striatal PV⁺ neurons 375 project, in the absence of diffusion, GP PV⁺ neurons could nevertheless be retrogradely 376 transduced.

377 Therefore, we examined the GP of 4 AAV2/9-hsyn and 3 AAV2/1-hsyn-injected mice

378 apparently devoided of direct GP transduction as shown in Fig. 2 E-H.

379 In one AAV2/9-injected mice, few GFP⁺ cells were observed in the GP (Fig. 6 D). As

380 expected these native GFP fluorescent cells were also PV^+ (Fig. 6 E & F). However, the

381 level of fluorescence intensity of GFP⁺ cells in the GP was lower than the fluorescence of

382 directly transduced striatal cells (Fig.6 A), also labeled by anti-PV immunofluorescence

383 (Fig.6 B & C). Whether these GFP⁺ cells resulted from retrotransduction of pallido-striatal

384 projection neurons or residual diffusion of viral particles remains to be determined. The

385 mean fluorescence intensity of the GFP⁺ cells in the GP (n=4) was drastically lower than

386 striatal GFP⁺ cells in the same animal (n=4), respectively 5.90±3.69 and 76.98±7.78 A.U.

387 In 2 other AAV2/9-injected mice, no GFP⁺ cells were detected in the GP but some fibers

388 were evidenced (Fig.6 G-I). The origin of these GFP+/PV+ fibers in the GP remains to be

389 determined. Finally, in the last AAV2/9-injected mice, no GFP⁺cells of fibers were observed.

390 These data suggest that if AAV2/9 retrograde transduction of pallido-striatal neurons

391 occurred, it was very inefficient.

392 In AAV2/1-injected mice, devoided of direct GP transduction, no GFP+ cells were detected in 393 the GP (data not shown).

395 Figure 6. Retrograde transduction of GP PV⁺ neurons mediated by AAV2/9-hsyn-FLEX-eGFP 396 **injected in the dorsal striatum?**

PV^{Cre} (+/+) mice were injected with AAV2/9-hsyn-FLEX-eGFP vector. Animals showing no efficient and widespread transduction (as shown in Fig.2 E-H) were further examined. (A-C) Transduced cells in the dorsal striatum. (D-F) Mice showing 4 GFP⁺ cells in the GP- (G-I) 1 out 2 mice showing GFP⁺ fibers in the GP.

- 398 *Decrease of PV-expressing cells in the striatum of PV^{Cre} mice*
- 399 Continuous postnatal Cre expression has previously been reported to cause a decrease of
- 400 cell numbers of some cell types such as immune cells (Schmidt-Supprian and Rajewsky
- 401 2007) (Zeitrag, Alterauge et al. 2020) or retinal pigmented epithelium cells (He, Marioutina et
- 402 al. 2014).

403 Therefore, we compared the number of PV^+ cells in WT and PV^{Cre} homozygous mice.

404 The number of PV⁺ cells was drastically (approx. 2-fold) reduced in the striatum (Fig. 7 G)

405 but not in the CTX (Fig. 7 H) of PV^{Cre} mice as compared to WT mice (compare Fig. 7 A-C to 406 Fig. 7 D-F).

407 In order to determine whether Cre effect could be aggravated by viral transduction, injected 408 and non-injected hemispheres of PV^{Cre} mice were compared for the 3 vectors (Fig. 7 I). No 409 difference was observed. As a control, the number of PV⁺ cells in the left and right 410 hemispheres of 3 non-injected WT mice were compared. No difference was observed $(40.61\pm3.88$ and 39.92 ± 3.19 cells/mm² for the right and left hemispheres respectively;

412 student t test: p=0.8240).

AAV does not affect the number of PV⁺cells in the striatum

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415 Figure 7. Decrease of striatal PV⁺ cells in PV^{Cre} mice

Wild-type (A-C) and PVCre (D-F) mice were injected with different AAV-FLEX vectors. *Cre decreases the number of PV⁺ cells in the striatum but not in the cortex.* In the striatum (B & E), the number of PV⁺ cells was significantly reduced in PV^{Cre} mice (two-way ANOVA with Sidak's multiple comparison test; p=0.0201 for AAV2/9-CBA-FLEX-eGFP; p=0.0064 for AAV2/9-hsyn-FLEX-eGFP; p=0.0004 for AAV2/1-hsyn-FLEX-eGFP) (G). Circles, AAV2/9-CBA-FLEX-eGFP; squares, AAV2/9 hsyn-FLEX-eGFP; triangles, AAV2/1-hsyn-FLEX-eGFP. In the cortex (C & F), the number of PV⁺ cells in PV^{Cre} mice was not significantly different from the number of PV⁺ cells in WT mice (Student t test) (H). Circles, wild-type mice; squares, PV^{Cre} mice. *Viral toxicity* The number of PV⁺ cells in the injected hemisphere of PVCre mice was compared to the contralateral non-injected hemisphere (I). Statistical analysis did not reveal a significant difference (twoway ANOVA with Sidak's multiple comparison test). Circles, AAV2/9-CBA-FLEX-eGFP; squares, AAV2/9-hsyn-FLEX-eGFP; triangles, AAV2/1-hsyn-FLEX-eGFP.

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- 417

418 **Discussion**

419 In order to study the role of different types of neurons in complex neuronal circuits, it

420 is crucial to have available molecular tools allowing to selectively deliver genetic information

421 in targeted neuronal subpopulations. Furthermore, it is key to avoid transducing neurons with

422 similar functions in other brain regions which could perturb the behavioral and functional

423 outcomes. Thus, both cell type-specific and region-specific targeting are required.

424 In the present study, we focused on the targeting of transgene expression into mice

425 PV⁺ interneurons of the dorsal striatum. These interneurons, also called "fast-spiking",

426 although representing a very small proportion of striatal neurons (approx. 0.5%), are key to

427 the coordination of striatal projections neurons which modulate the sensory-motor loop.

EXPLO PV^{Cre} transgenic mice combined with Cre-dependent AAV vectors have been widely used to target PV-expressing neurons in the cerebral cortex, hippocampus and thalamus (Daigle, Madisen et al. 2018) (Madisen, Zwingman et al. 2010). However, very few studies 431 aimed at targeting striatal PV⁺ interneurons (Abreu, Gama et al. 2018, Enterria-Morales, Lopez-Lopez et al. 2020).

In order to achieve efficient and selective transduction of striatal PV⁺ interneurons, 434 we compared 2 different AAV serotypes known to efficiently transduce the CNS, AAV2/1 435 (Burger, Gorbatyuk et al. 2004) (Bockstael, Chtarto et al. 2008) and AAV2/9 (Ciesielska,

436 Hadaczek et al. 2012) (Klein, Dayton et al. 2008), and 2 different transcription promoters, the 437 human synapsin (hsyn) and the hybrid chicken β -actin/CMV (CBA) promoters.

 We show that, due to the diffusion of viral particles in the brain parenchyma, it is difficult to target the dorsal striatum while avoiding diffusion to the GP. In contrast to the 440 sparse distribution of PV^+ striatal interneurons, the GP contains a high density of PV^+ neurons (40% of the total) which project to the STN, the SNr and the striatum (Saunders, 442 Huang et al. 2016). Transduction of PV⁺ GP projection neurons severely compromised the 443 targeting of PV⁺ striatal interneurons. Precise stereotaxic coordinates adjustment in the dorsolateral striatum, AAV1 capsids (diffusing less than AAV9), and the hsyn promoter (having a moderate transcriptional activity) were necessary to avoid direct transduction of 446 GP PV⁺ neurons. Such an extrastriatal PV⁺ neuronal transduction after striatal injections of AAV-FLEX vectors, has been previously described and authors *a posteriori* excluded from their behavioral analysis the animals showing a transduction in untargeted areas (Xu, Li et al. 2016).

 450 None of the vectors provided any GFP⁺ cell in WT mice demonstrating the tight Cre-451 dependency of transgene expression. With all tested vectors, the proportion of GFP+ cells 452 stained by anti-PV antibodies was higher than 85%. However, GFP⁺ cells not expressing PV 453 at a detectable level were observed. This could be due to an effect of the Cre recombinase 454 towards PV⁺ neurons. Indeed, we observed that the number of PV⁺ cells was drastically 455 reduced (\sim 2-fold) in the striatum of PV^{Cre} mice. This Cre-mediated effect was specific to 456 striatal PV⁺ neurons since no reduction of PV⁺ cells was observed in the cerebral cortex. A 457 similar phenomenon has been reported in retinal pigmented cells in which Cre expression 458 resulted in age- and dosage- dependent attenuation of β catenin and phalloidin stainings in 459 VDM2-Cre mice (He, Marioutina et al. 2014). It remains to be determined whereas Cre effect 460 on PV⁺ cells is also present in heterozygous Pvalbm1/Arbr mice. During the writing of this 461 manuscript, Enteria-Moralez and collaborators reported a similar effect of Cre towards PV⁺ 462 cells in the striatum of homozygous Pvalbm1/Arbr mice. In this study, heterozygous Cre (+/-)

 mice showed a non-significant decrease of PV mRNA. However, the small number of mice (n=2 for some groups) precluded valid statistical analysis.

 AAV vectors have previously been reported to have a toxic effect on specific classes of neurons in particular when expressing GFP. Indeed, nigral injection of AAV2/1 vectors in rats resulted in a decrease of the number of TH-expressing neurons, which was further aggravated in the presence of GFP (Albert, Voutilainen et al. 2019). AAV2/9 vectors expressing GFP were furthermore shown to elicit an inflammatory response in the rat striatum (Samaranch, San Sebastian et al. 2014).

 Another potential off-target effect could result from retrograde transport of AAV viral 472 particles in pallidostriatal fibers and transgene expression in GP PV⁺ projection neurons (Saunders, Huang et al. 2016). AAV retrograde transport depends on the capsid serotype and inconsistent data have been reported. For example, AAV1 retrograde transport has been repeatedly found to be negligible (Burger, Gorbatyuk et al. 2004) (Bockstael, Chtarto et al. 2008) (Oh, Harris et al. 2014) (Tervo, Hwang et al. 2016) although another report shows that it occurred with a moderate efficiency in nigrostriatal dopaminergic neurons (Taymans, Vandenberghe et al. 2007). Surprisingly, efficient retrograde infection was demonstrated with AAV2/1-FLEX vectors in several types of neurons (Rothermel, Brunert et al. 2013) whereas, in the same study, a constitutive AAV2/1 vector expressing mCherry from the CBA promoter but not one expressing eGFP from the hsyn promoter, was proficient for retrograde transduction. Similarly, inconsistent reports of AAV9 serotype retrograde transport efficiency have been published (Klein, Dayton et al. 2008) (Tervo, Hwang et al. 2016) (Cearley and Wolfe 2006) (Gallagher, Watson et al. 2008). In a comparative study, (Tervo, Hwang et al. 2016) AAV1 ability to undergo retrograde transport was found to be lower than that of AAV9. Different species (rat versus mice), different neuronal pathways, expression levels driven by 487 the chosen promoter and titers of the vector may explain these discrepancies. 488 To our knowledge, retrograde transduction of GP PV⁺ neurons following intrastriatal AAV injection has never been described. In our study, in the absence of apparent diffusion, only a

490 very low number of GFP⁺ cells were detected in 1 out of 4 AAV2/9-injected mice examined,

and their native green fluorescence was lower than at the target site in the striatum.

492 Furthermore, it cannot be excluded that the observed GFP⁺ cells in the GP resulted from a low level of diffusion rather than retrograde transport. In contrast, with AAV2/1 vector, no retrograde transduction of pallidostriatal neurons was observed in our conditions.

 In conclusion, caution should be taken when interpreting functional and behavioral 496 data obtained in the PV^{Cre} mice/AAV-FLEX system. First, it is key to avoid transducing PV neurons in other brain regions which will perturb the behavioral and functional outcomes. GP 498 PV⁺ neurons project to the STN, SNr and striatum (Saunders, Huang et al. 2016), nuclei which are all involved in the motor loop. Thus, transduction of these neurons might drastically modify the functional effects of transgene expression. Secondly, the herein 501 described Cre-mediated deleterious effect toward PV⁺ striatal interneurons could reflect neuronal cell death or loss of PV expression. In the latter case, these neurons are probably dysfunctional since PV, a calcium-binding protein is important for neuronal activity. Perturbation of the synaptic plasticity of these fast-spiking neurons which are thought to coordinate the activity of striatal projection neurons, might severely affect the motor loop. It should be noted that, in a model of AAV-mediated Cre expression, neuronal programmed cell death inducing behavioral perturbations were described (Rezai Amin, Gruszczynski et al. 2019)

 In order to achieve targeted gene delivery into fast-spiking striatal interneurons 510 without the use of PV^{Cre} mice and AAV-Flex vectors, a more selective transcriptional targeting could be designed. Viral vectors with specific enhancers allowing to target neuronal or glial cell populations in wild-type animals are coming to an era (Dimidschstein, Chen et al. 2016) (Meunier, Merienne et al. 2016) (Pignataro, Sucunza et al. 2017) (Vormstein-Schneider, Lin et al. 2020).

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- **Marcelo Duarte-Azevedo**: Conceptualization, Methodology, Validation, Formal analysis,
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- **Sibilla Sander:** Conceptualization, Methodology, Validation, Formal analysis, Investigation, Data
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- **Cheryl Jeanneret**: Validation, Formal analysis, Investigation, Data curation, Writing-Review and
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