

Technical Note

## Enhanced binding of metabotropic glutamate receptor type 5 (mGluR5) PET tracers in the brain of parkinsonian primates

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Received 22 January 2008; revised 31 March 2008; accepted 5 April 2008  
Available online 20 April 2008

**The interplay between dopamine and glutamate in the basal ganglia regulates critical aspects of motor learning and behavior. Metabotropic glutamate receptors (mGluR) are increasingly regarded as key modulators of neuroadaptation in these circuits, in normal and disease conditions. Using PET, we demonstrate a significant upregulation of mGluR type 5 in the striatum of MPTP-lesioned, parkinsonian primates, providing the basis for therapeutic exploration of mGluR5 antagonists in Parkinson disease.**

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Increasing understanding of receptor function is revealing complex and dynamic interactions between neurotransmitter systems (Zhang and Sulzer, 2003), including heterosynaptic transmission, alternative second messengers and integration of intracellular cascades at different levels. In the basal ganglia, dopamine (DA) regulation of glutamate (Glu) neurotransmission is complex and the loss of DA-mediated inhibition in the striatum in Parkinson disease (PD) results in an imbalance in other neurotransmitters, mostly excitatory. Accordingly, Glu antagonists at ionotropic glutamate receptors (iGluR) (as well as anticholinergic drugs) have demonstrated antiparkinsonian effects, although their side effects are intolerable (Gubellini et al., 2004). In contrast, metabotropic (m)GluRs are activated when there is excess Glu in the synaptic cleft, that spills over to activate perisynaptic receptors — and therefore act as sensors and modulators when or where Glu transmission is enhanced (Gubellini et al., 2004; Konradi et al., 2004). This functional specificity makes them attractive pharmacological targets, although it cannot be excluded that drugs will also interact with mGluRs in other regions, masking or altering the effects (Gubellini et al., 2004). In contrast to iGluR antagonists, such

side effects may be subtle and not discernible in animal models (Gubellini et al., 2004). The prevalent distribution of mGluR5 in the striatum and limbic system supports their role modulating DA and Glu-dependent signaling and synaptic plasticity within the basal ganglia cortico-subcortical loops (Gubellini et al., 2004). While the role of mGluR5 in cocaine addiction has been firmly established (Chiamulera et al., 2001) using mGluR5 knockout mice, data from Parkinson models are equivocal (Armentero et al., 2006; Breyse et al., 2003; Mela et al., 2007; Oueslati et al., 2005; Samadi et al., 2007) except for a likely involvement in the pathophysiology of dyskinesias (Mela et al., 2007), a frequent complication of long-term L-DOPA replacement therapy. Taking advantage of novel PET tracers (Pellegrino et al., 2007; Wang et al., 2007) we examined the distribution of mGluR5 in the primate brain and the effect of DA denervation in MPTP-lesioned parkinsonian primates to determine whether mGluR5 drugs may be therapeutically relevant for PD.

### Methods

Eight young adult male primates (*Macaca fascicularis*) were included in the study. Animals were individually housed at the New England Regional Primate Center. Studies were conducted following NIH guidelines for animal use and care and were approved by the Internal Animal Care and use Committee at Harvard Medical Area and the Massachusetts General Hospital.

PET imaging studies (MicroPET P4, Concord Microsystems) were conducted in 4 control (naïve) and 4 MPTP-lesioned, parkinsonian primates. Systemic administration of MPTP to induce a stable parkinsonism and rating of parkinsonian severity was performed as previously reported by our group in detail (Jenkins et al., 2004). Imaging studies were performed under propofol anesthesia (0.3 mg/kg/min iv). In six animals mGluR5 and DA transporter binding were investigated in the same imaging session, using first a carbon-11 labeled pyridine analog, 2-(2-(5-[<sup>11</sup>C]methoxy)pyridin-3-yl)ethynyl)pyridine ([<sup>11</sup>C]MPEPy), which has fast binding kinetics.

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Available online on ScienceDirect (www.sciencedirect.com).

Starting from the administration of [<sup>11</sup>C]MPEPy (10–13 mCi i.v., specific activity 900 mCi/μmol) dynamic volumetric data were acquired for 90 min. One hour after the data acquisition was

completed (i.e.150 min after injection) the carbon-11 labeled cocaine analog, 2β-[<sup>11</sup>C]carbomethoxy-3β-(4-fluorophenyl) tropane ([<sup>11</sup>C]CFT) was injected (8–10 mCi i.v., specific activity

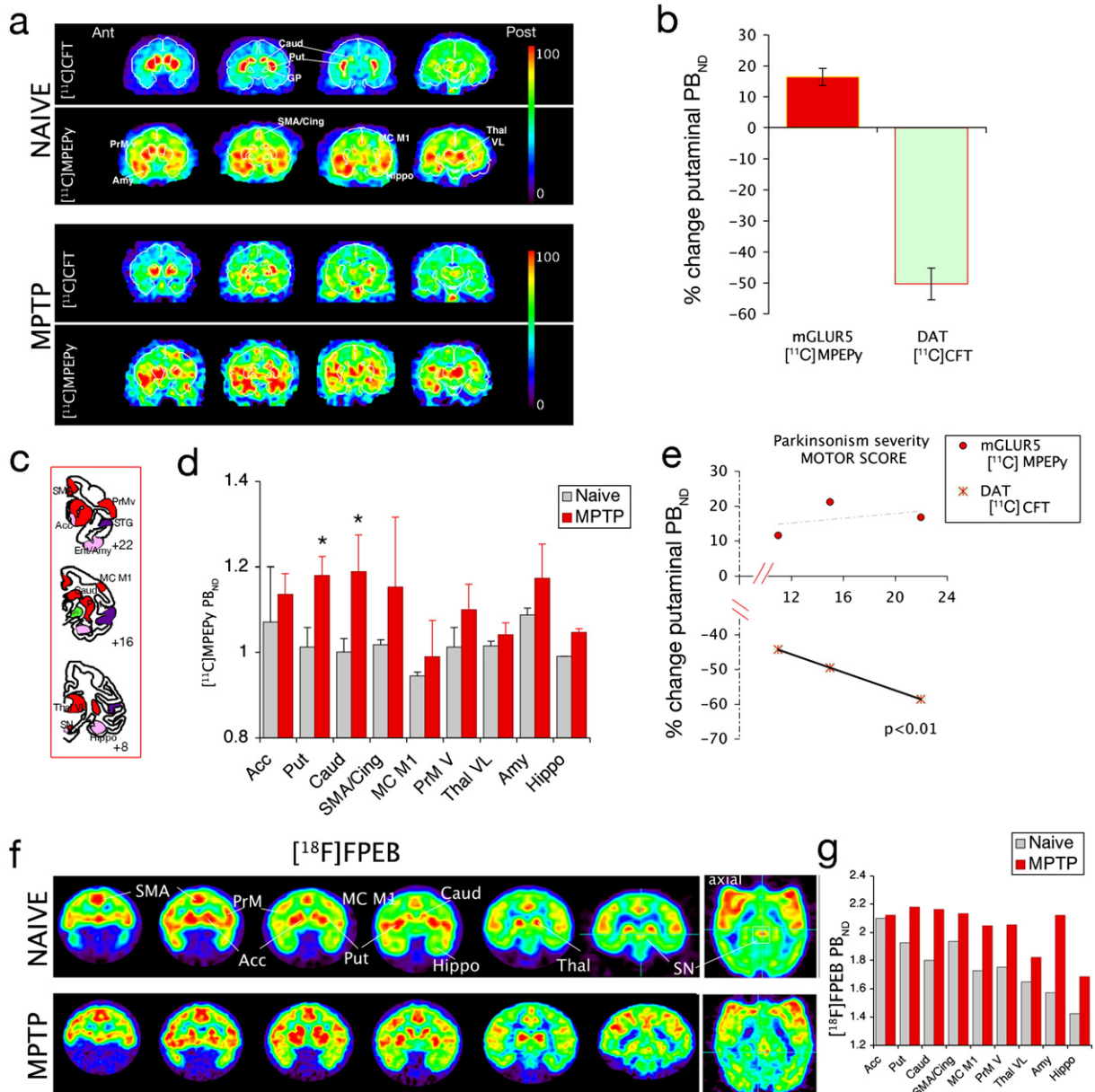


Fig. 1. a) Representative images of the distribution of the dopamine transporter (DAT) and mGluR5 in a naïve (top panel) and a parkinsonian primate. Identical coronal slices shown at 4 anteroposterior levels were acquired consecutively in the same imaging session, to facilitate delineation of the basal ganglia. Color scale is adjusted to maximal activity for each tracer. Distribution of [<sup>11</sup>C]CFT accumulation is illustrated at 40–45 min after administration of radioactivity (8–10 mCi i.v., specific activity 1400 mCi/μmol). [<sup>11</sup>C]MPEPy accumulation is illustrated at 10–25 min after administration of radioligand (10–13 mCi, iv., specific activity 900 mCi/μmol). b) Average change from naïve baseline in the putaminal binding in MPTP-lesioned animals. c) Schematic representation of anatomical regions integrated in the mesolimbic (pink) mesostriatal (red) and temporal (purple) loops that are affected in diseases in which the DA/Glu interaction appears to play a crucial pathogenic role, i.e. addiction, Parkinson's disease and schizophrenia, respectively. Globus pallidus pars interna is filled in green, to represent that no significant binding was observed in this region. d) ROI analysis of [<sup>11</sup>C]MPEPy binding demonstrated a significant increase in caudate and putamen. e) Putaminal change in [<sup>11</sup>C]MPEPy binding was not significantly correlated with the severity of parkinsonian signs (global score 0–24), unlike the change in [<sup>11</sup>C]CFT binding. f) Expression of mGluR5 in the brain of a naïve (top) and a parkinsonian primate, using the highly selective tracer [<sup>18</sup>F]FPEB delineated primary and downstream DA regions. SN/VTA are shown in coronal and axial reconstruction. Distribution of [18F]FPEB accumulation is illustrated at 60–70 min after administration of radioligand (0.8–1.2 mCi i.v., specific activity 1900 mCi/μmol); g) Regional values in binding potential follow the pattern described above for [<sup>11</sup>C]MPEPy. Acc = Accumbens, Amy = Amygdala, Caud = Caudate, Cing = cingulate Cortex, Ent = Entorhinalis cortex, GP = Globus Pallidus, Hippo = Hippocampus, MC M1 = Primary Motor Cortex, PBN<sub>D</sub> = in vivo binding potential, PrM = Premotor Cortex, Put = Putamen, SMA = supplementary motor area, SN = substantia nigra, Thal = Thalamus, V = ventral, VL = ventrolateral.

1400 mCi/ $\mu$ mol) and imaging data were acquired for 90 min. Transmission imaging was done using a cobalt-57 source for processing maps for attenuation correction. Image processing was done using filtered backprojection and software provided by the manufacturer (Asipro 6.0, Concord Microsystems/Siemens). In two animals imaging studies of mGluR5 and DAT were done in different session to avoid background activity, since [ $^{18}$ F]FPEB has long retention time. After administration of [ $^{18}$ F]FPEB (3-[ $^{18}$ F]fluoro-5-(2-pyridinylethynyl)benzotrile) (0.8–1.2 mCi i.v., specific activity 1900 mCi/ $\mu$ mol) dynamic volumetric data were acquired for 120 min followed by transmission imaging and data reconstruction as above.

The synthesis and labeling of these tracers has been described by us (Brownell et al., 2003; Pellegrino et al., 2007; Wang et al., 2007). Regions of interest (ROIs) were delineated based on MRI high resolution anatomical images acquired on a Siemens 3T Trio system, as previously reported by our group (Jenkins et al., 2004; Sanchez-Pernaute et al., 2007) and anatomical atlas (Fig. 1c). *In vivo* binding potential ( $BP_{ND}$ ) (Innis et al., 2007) of these ligands was calculated using the cerebellum as reference tissue (Zhu et al., 2007).

#### Statistical analysis

Results are shown as mean  $\pm$  SD. Two-tailed unpaired t test was used for comparison between conditions and simple regression analysis to assess the correlation with motor signs.

#### Results and discussion

The distribution of [ $^{11}$ C]MPEPy in the brain of naïve ( $n=3$ ) and MPTP-lesioned, parkinsonian primates ( $n=3$ ) was compared to that of [ $^{11}$ C]CFT, a cocaine analog that binds to the DA transporter (DAT) as described (Brownell et al., 2003) (Fig. 1a). In naïve animals [ $^{11}$ C]MPEPy rapidly accumulated in discrete cortical and subcortical regions encompassing the premotor and cingulate cortices, superior temporal gyrus and limbic (paraentorhinal/amygdala/hippocampal) cortex, the nucleus accumbens, caudate and putamen (predominantly at rostral levels), the ventral thalamus and the midbrain. This distribution corresponds to areas that have been shown to display high mGluR5 mRNA expression in the rodent brain (Messinger et al., 2002). Of interest is the lack of binding in the globus pallidus, which agrees with mRNA data in rodent (but not with published immunohistochemistry (Smith et al., 2000)).

MPTP-lesioned animals had a significant loss of [ $^{11}$ C]CFT binding in the putamen ( $t_{1,3}=8.27$ ;  $p<0.05$ ) with typical preservation of DA innervation of the nucleus accumbens (Fig. 1b, (Jenkins et al., 2004)). Regional analysis of [ $^{11}$ C]MPEPy was performed in cortical and subcortical areas to examine the motor and limbic DA loops (color coded in Fig. 1c, at 3 coronal levels of the macaque brain). We found a significant enhancement of binding in the motor regions of the striatum (putamen  $t_{1,4}=4.56$ ;  $p=0.01$ ; caudate  $t_{1,4}=3.57$ ;  $p=0.02$ ) (Fig. 1d). The average increase in the motor striatum,  $18.6 \pm 8.1\%$  was moderate (16% in the putamen, Fig. 1b) and not significantly correlated with the loss of [ $^{11}$ C]CFT binding, (Fig. 1e) or with the severity of the parkinsonian score – although the slope of the regression was positive (0.34). We acknowledge that the small volumes of ROIs are vulnerable for partial volume effects and the recorded activity might be less than the “real” activity. However, in this case it means that enhancement of mGluR5 accumulation is even more than in the presented data. The loss of [ $^{11}$ C]CFT binding was directly correlated with the severity of the parkinsonian signs

( $p<0.005$ ) measured by the global motor score in a rating scale based on the motor subscale of the UPDRS (Fig. 1e) as we have previously described in this model (Jenkins et al., 2004). To confirm that the change in [ $^{11}$ C]MPEPy binding reflected changes in mGluR5, we examined in 2 other primates, the distribution of the novel compound [ $^{18}$ F]FPEB (Hamill et al., 2005; Wang et al., 2007), which has exceptionally high affinity to mGluR5. The reported *in vitro*  $B_{max}/K_d$  value based on the saturation binding studies in rhesus caudate-putamen tissue is 210 (Patel et al., 2007) and *in vivo* studies have shown that it is subgroup specific (Wang et al., 2007). These imaging studies confirmed mGluR5 binding to DA target regions (Fig. 1f) matching cortical and subcortical areas with identical distribution to that we described previously using fMRI and amphetamine (Jenkins et al., 2004) (i.e. the areas in which DA release induces an increase in regional cerebral blood volume). Interestingly, there was no significant binding in the pallidal complex, either in the naïve or DA denervated condition. Given the high affinity of this tracer, it was feasible to resolve the VTA/SN region (Fig. 1f, see axial reconstruction in the far right panel) although more studies are necessary for quantification of MPTP-induced changes using this tracer (Fig. 1g).

In conclusion, here we identified a significant enhancement in [ $^{11}$ C]MPEPy mGluR5 binding in the striatum of MPTP-lesioned parkinsonian primates. Because, in principle, DA denervation in PD enhances Glu transmission, further amplification through upregulation of striatal mGluR5 appears to be a possible pathogenic mechanism involved in aspects of disease progression, as well as in the development of long-term complications. Thus, these novel selective mGluR5 PET tracers are valuable tools for *in vivo* mechanistic studies in patients and in the development of novel therapeutic approaches for PD.

#### Acknowledgments

Supported by NIH-1R01 EB001850 and NIH-1P50 NS39793. We are grateful to Jack McDowell for the technical support and Dr BG Jenkins for the MRI studies.

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