

# RELATIONSHIP BETWEEN NEUROMELANIN AND DOPAMINE TERMINALS WITHIN THE PARKINSON'S NIGROSTRIATAL SYSTEM

## Running Title: Neuromelanin and dopamine in Parkinson's

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## ABSTRACT

Parkinson's disease is characterized by the progressive loss of pigmented dopaminergic neurons in the substantia nigra and associated striatal deafferentation. Neuromelanin content is thought to reflect the loss of pigmented neurons, but available data characterising its relationship with striatal dopaminergic integrity are not comprehensive or consistent, and predominantly involve heterogeneous samples. In this cross-sectional study, we utilised neuromelanin-sensitive magnetic resonance imaging and the highly specific dopamine transporter positron emission tomography radioligand,  $^{11}\text{C}$ -PE2I, to assess the association between neuromelanin-containing cell levels in the substantia nigra pars compacta and nigrostriatal terminal density *in vivo*, in 30 patients with bilateral Parkinson's disease. Fifteen healthy controls also underwent neuromelanin-sensitive imaging. We used a novel approach taking into account the anatomical and functional subdivision of substantia nigra into dorsal/ventral tiers and striatal nuclei into pre/post-commissural sub-regions, in accordance with previous animal and post-mortem studies, and consider the clinically asymmetric disease presentation. *In vivo*, Parkinson's disease subjects displayed reduced neuromelanin levels in the ventral ( $-30\pm 28\%$ ) and dorsal tiers ( $-21\pm 24\%$ ) as compared to the control group ( $F_{1,43} = 11.95$ ,  $P = 0.001$ ). Within the Parkinson's disease group, nigral pigmentation was lower in the ventral tier as compared to the dorsal tier ( $F_{1,29} = 36.19$ ,  $P < 0.001$ ) and lower in the clinically-defined most affected side ( $F_{1,29} = 4.85$ ,  $P = 0.036$ ). Similarly, lower dopamine transporter density was observed in the ventral tier ( $F_{1,29} = 76.39$ ,  $P < 0.001$ ) and clinically-defined most affected side ( $F_{1,29} = 4.21$ ,  $P = 0.049$ ). Despite similar patterns, regression analysis showed no significant association between nigral pigmentation and nigral dopamine transporter density. However, for the clinically-defined most affected side, significant relationships were observed between pigmentation of the ventral nigral tier with striatal dopamine transporter binding in pre-commissural and post-commissural striatal sub-regions known to receive nigrostriatal projections from this tier, while the dorsal tier correlated with striatal projection sites in the pre-commissural striatum ( $P < 0.05$ , Benjamini-Hochberg corrected). In contrast, there were no statistically significant relationships between these two measures in the clinically-defined least affected side. These findings provide important insights into the topography of nigrostriatal neurodegeneration in Parkinson's disease, indicating that the characteristics of disease progression may fundamentally differ across hemispheres and support post-mortem data showing asynchrony in the loss of neuromelanin-containing versus tyrosine-hydroxylase positive nigral cells.

*Keywords: Parkinson's disease, magnetic resonance imaging, positron emission tomography, neuromelanin, dopamine transporter*

*List of abbreviated terms: AC = anterior commissure; ADD = signal-averaged images; AI = asymmetry index;  $^{11}\text{C-PE2I}$  = ( $^{11}\text{C}$ )N-(3-iodopro-2E-enyl)-2 $\beta$ -carbomethoxy-3 $\beta$ -(4'-methylphenyl)nortropine); CP = cerebral peduncle; CR = contrast ratio; DA = dopamine; DARTEL = diffeomorphic anatomical registration through exponentiated Lie algebra; DAT = dopamine transporter; HC = healthy control; FDR = false discovery rate; FSL = FMRIB software library; FWHM = full width at half maximum; GRAPPA = Generalized Autocalibrating Partial Parallel Acquisition;  $^{123}\text{I-}\beta\text{-CIT}$  = 2 $\beta$ -carbomethoxy-3 $\beta$ -(4[ $^{123}\text{I}$ ]iodophenyl)tropine;  $^{123}\text{I-FP-CIT}$  = N-(3-Fluoropropyl)-2 $\beta$ -carbomethoxy-3 $\beta$ -(4-[ $^{123}\text{I}$ ]iodophenyl)nortropine; LEDD = levodopa equivalent daily dose; MIAKAT = Molecular Imaging And Kinetic Analysis Toolbox; MNI = Montreal neurological institute; MPRAGE = Three-dimensional T1-weighted magnetization prepared rapid acquisition gradient echo; MPTP = 1-methyl-4-phenyl-1,2,3,6-tetrahydropyridine; 6-OHDA = 6-hydroxydopamine; PD = Parkinson's disease; PC = posterior commissure; ROI = region of interest; SN = substantia nigra; SNc = substantia nigra pars compacta; SNc<sup>dor</sup> = substantia nigra pars compacta dorsal tier; SNc<sup>ven</sup> = substantia nigra pars compacta ventral tier; SNr = substantia nigra pars reticulata; SPECT = single-photon emission computed tomography; SPM = statistical parametric mapping; SRTM = simplified reference tissue model; TAC = time-activity curve; TSE = turbo spin echo; UPDRS-III = Unified Parkinson's disease rating scale part-III*

## INTRODUCTION

The main pathological hallmarks of Parkinson's disease consist of the progressive loss of pigmented dopaminergic (DA) neurons in the substantia nigra pars compacta (SNc) and a reduction of dopaminergic striatal afferents (Fearnley and Lees, 1991). An important factor in the disease aetiology concerns the function of neuromelanin that is contained within pigmented SNc neurons. Neuromelanin is a dark insoluble complex compound that is synthesised as a by-product of DA oxidation during cytosolic DA homeostasis. Physiologically, neuromelanin is proposed to have a dual role. It sequesters potentially toxic organic chemical, exogenous and endogenous metals such as iron in redox-inactive state while its synthesis confers neuroprotection from cytosolic reactive/toxic quinones. However, neuromelanin released into the extracellular space upon neuronal death, as is the case in Parkinson's disease, further exacerbates neurodegeneration by releasing toxic metals and catecholaminergic products and inducing oxidative stress, local microglial activation and chronic inflammation (Zucca *et al.*, 2006; Zucca *et al.*, 2014; Zucca *et al.*, 2017).

Neuromelanin bound to metals such as iron and copper is highly paramagnetic, leading to T1-shortening and hyperintense signal on T1-weighted turbo spin-echo magnetic resonance imaging (MRI) sequences (Sulzer *et al.*, 2018). Hyperintensity in such images demonstrate direct associations with post-mortem neuromelanin-containing dopaminergic cell counts (Kitao *et al.*, 2013). In recent years several studies evaluating nigral depigmentation have demonstrated the ability for neuromelanin-sensitive MR to distinguish between Parkinson's disease and healthy individuals with high sensitivity and specificity, representing a potential *in vivo* index of neuromelanin content, disease progression and nigral cell death (Sasaki *et al.*, 2006; Kashihara *et al.*, 2011; Schwarz *et al.*, 2011; Matsuura *et al.*, 2013; Ogisu *et al.*, 2013; Ohtsuka *et al.*, 2013; Castellanos *et al.*, 2015; Reimao *et al.*, 2015; Fabbri *et al.*, 2017).

Overall estimations suggest that ~30% of neuromelanin-containing cells in the SNc are lost prior to the appearance of classical Parkinson's disease motor symptoms (Fearnley and Lees, 1991; Cheng *et al.*, 2010). In contrast, depletion of dopaminergic terminal markers in the principal projection sites of the striatum is in most cases much more pronounced at symptomatic threshold. Typically deficits of striatal DA integrity are estimated between 50-80% as compared to controls, although lower values of ~20% have been reported (Bernheimer *et al.*, 1973; Nyberg *et al.*, 1983; Kish *et al.*, 1988; Scherman *et al.*, 1989; Lee *et al.*, 2000; Cheng *et al.*,

2010). The difference between these estimates raises questions regarding the relationship between neuromelanin and the integrity of nigrostriatal terminals.

To date, three studies have examined pigmented cell density in the SNc at post-mortem, alongside striatal dopamine transporter (DAT) binding measured *in vivo* with  $^{123}\text{I}$ -FP-CIT single-photon emission computed tomography at 2.5-5 years ante-mortem, but with conflicting results (Colloby *et al.*, 2012; Kraemmer *et al.*, 2014; Saari *et al.*, 2017). Nigral cell counts were related to  $^{123}\text{I}$ -FP-CIT striatal specific binding ratios in Parkinson's disease dementia (Colloby *et al.*, 2012) and heterogeneous neurologic (Kraemmer *et al.*, 2014) samples. This correlation was not present in a small group of 11 Parkinson's disease patients (Saari *et al.*, 2017), leading Saari and colleagues to hypothesise that the relationship between surviving nigral cells and striatal dopamine is lost in later stages of the disease. To our knowledge, only two recent studies have assessed this relationship in a multimodal setting. Using neuromelanin-sensitive MRI, strong positive correlations were demonstrated between striatal DAT binding and both SN volume, as well as the SN-to-cerebral peduncle contrast ratios, in cohorts with idiopathic Parkinson's disease (Isaias *et al.*, 2016) and mixed parkinsonism (Kuya *et al.*, 2016). Despite these findings, the topography of the relationship between imaging biomarkers as it pertains to nigrostriatal damage has not yet been described in detail.

In descriptive anatomy, neuronal tracing and staining studies define two cellular tiers of the SNc; a calbindin-positive dorsal tier which encompasses the ventral tegmental area (VTA) and a calbindin-negative ventral tier which spreads laterally toward the SN pars reticulata (SNr) (Lynd-Balta and Haber, 1994; Haber *et al.*, 2000; Haber, 2014). In Parkinson's disease, the pattern of nigral neuronal loss differs to normal ageing, exhibiting an exponential decay with approximately 45% reduction over the first decade of illness, beginning and most prominently affecting the lateral ventral tier before spreading to the medial ventral and dorsal tiers. This is at variance to the linear fallout observed in healthy ageing that is estimated at 4.7-6% per decade and the relative sparing of the ventral compared to the dorsal tiers (Fearnley and Lees, 1991; Gibb and Lees, 1991). Meanwhile, the striatum is characterised by a rostrocaudal gradient of dopaminergic denervation with relative indemnity of caudate nucleus (Bernheimer *et al.*, 1973; Nyberg *et al.*, 1983; Kish *et al.*, 1988; Scherman *et al.*, 1989; Lee *et al.*, 2000; Cheng *et al.*, 2010; Oh *et al.*, 2012; Han *et al.*, 2016). This Parkinson's disease –related gradient is different to equivalent age-related declines of ~5-8% per decade across the putamen and caudate in healthy individuals (Ishibashi *et al.*, 2009; Shingai *et al.*, 2014). Evidence from animal work on

nigrostriatal topography indicates that neurons originating in the SNc dorsal tier project predominantly to the head of the caudate and anterior putamen, while those of the ventral tier project to the posterior putamen and posterior caudate nuclei (Carpenter and Peter, 1972; Szabo, 1980; Lynd-Balta and Haber, 1994; Haber *et al.*, 2000; Haber, 2014).

In this context, the current exploratory multimodal imaging study uses neuromelanin-sensitive MRI and PET with  $^{11}\text{C}$ -PE2I [a radioligand with high striatal binding (Jucaite *et al.*, 2006) and substantially greater DAT specificity than  $^{123}\text{I}$ -FP-CIT or  $^{123}\text{I}$ - $\beta$ -CIT (Abi-Dargham *et al.*, 1996; Emond *et al.*, 1997; Guilloteau *et al.*, 1998) that is highly correlated with Parkinson's disease symptom severity (Li *et al.*, 2018)] to examine the association between deficits in neuromelanin content in the SNc and reductions in DAT density in a Parkinson's disease cohort with bilateral disease. Importantly, we take into account the distinct anatomical sub-regions of both the SNc and the striatum, the reported topography of their neuronal connections and the lateralisation of clinical presentation, aiming to describe more comprehensively the association between the two imaging markers *in vivo* using an exploratory approach. While we expected there to be a relationship between neuromelanin and DAT density, based on available data, no specific hypotheses regarding its topography or extent were made.

## **MATERIALS AND METHODS**

### **Subjects**

A total of thirty non-demented mild-moderate stage Parkinson's disease patients and fifteen healthy controls were recruited from specialist movement disorder clinics and local advertisement respectively, under the TRANSEURO (<http://www.transeuro.org.uk/>) and PaMIR (Parkinson MRI Imaging Repository) research projects. Diagnosis of Parkinson's disease was performed by movement disorder specialists in accordance with the Parkinson's UK Brain Bank Criteria (Hughes *et al.*, 1992), excluding atypical parkinsonism, concomitant vascular load, history of cognitive impairment, psychiatric disorders and factors that would preclude MRI scanning.

Patients were instructed to withdraw from all standard release anti-Parkinson's disease medications 24 hours prior to motor and imaging assessments and 48 hours for prolonged release medications. This included levodopa, dopamine agonists, catechol-o-methyltransferase

and monoamine oxidase-B inhibitors. Caffeine in any form was not permitted within 12 hours prior to scan. Levodopa equivalent daily dosage (LEDD) for each participant was calculated (Tomlinson *et al.*, 2010).

The Movement Disorders Society Unified Parkinson's Disease Rating Scale Part-III (UPDRS-III) (Goetz *et al.*, 2008) administered by two experienced raters was used to calculate total motor severity in the practically-defined OFF-medicated state and sub-divided into bradykinesia (items 4-8, 14), rigidity (item 3), tremor (items 15-18) and axial (items 1-2, 9-13) sub-scores. Clinical laterality was established based on clinical history recorded by the neurology unit at which patients were diagnosed. Asymmetry was also evaluated at research appointments during the study, derived from the sum of the UPDRS-III laterality items in the practically-defined OFF-medicated state.

Ethical approval was obtained from the local Research Ethics Committees for TRANSEURO (IRAS: 57821, 65071, 78574; EPN2013/758, IK2013/685) and PaMIR (IRAS: 124223). Participant consent was obtained in writing in accordance with the Declaration of Helsinki.

## **Image acquisition**

All scans were conducted at Invicro LLC (Hammersmith Hospital, London).

## **Magnetic resonance imaging**

MRI scans were acquired for all Parkinson's disease and healthy control participants on a 3T Siemens Magnetom Trio system with 32-channel head coil and consisted of neuromelanin-sensitive T1-weighted turbo spin-echo (TSE: TR=829ms; TE=12ms; flip angle=123°; echo train length=4; echo spacing=11.5ms; low specific absorption rate; FoV=256\*256mm; matrix size=320\*320) and high-resolution volumetric T1-weighted magnetization prepared rapid acquisition gradient echo (MPRAGE: TR=2300ms; TE=2.98ms; flip angle=9°; time to inversion=900ms; GRAPPA acceleration factor PE=2; FoV=240\*256mm; Matrix size=240\*256) sequences.

For the MPRAGE, one whole brain volume was acquired consisting of 160 contiguous slices of 1mm thickness. For the TSE, 12 slices of 2.5mm thickness and slice gap of 0.25mm were acquired parallel to the AC-PC line with coverage of the whole midbrain and upper pons. Scans

lasted 301 and 332 seconds respectively and patients were instructed to remain as still as possible for the duration.

### **Positron emission tomography**

Parkinson's disease participants (n=30) underwent  $^{11}\text{C}$ -PE2I ( $[^{11}\text{C}]\text{N}$ -(3-iodopro-2*E*-enyl)-2 $\beta$ -carbomethoxy-3 $\beta$ -(4'-methylphenyl)nortropane) PET scans, acquired on a Siemens Biograph TruePoint HI-REZ 6 PET/CT system. Patients were positioned supine such that the transaxial plane was parallel to the AC-PC plane and movement minimised using memory foam padding and video monitoring to aid repositioning.  $^{11}\text{C}$ -PE2I tracer volumes were prepared to 10ml using saline solution and administered intravenously as a single bolus injection followed immediately by 10ml saline flush (Injected dose= $325.85\pm 35.24$  MBq; Injected mass= $3.99\pm 1.89$   $\mu\text{g}$ ). Administration was at a rate of 1ml/s.

Dynamic emission data were acquired continuously while patients were at rest for 90 minutes post-injection. Data were binned into a dynamic series of 26 temporal frames (8x15s, 3x60s, 5x120s, 5x300s, 5x600s) and reconstructed with corrections for decay, scatter and attenuation using a filtered back-projection algorithm (direct inversion Fourier transform) with a matrix size of 128x128, zoom of 2.6 and 2mm isotropic pixel size and smoothed using a three-dimensional 5mm FWHM trans-axial Gaussian image filter. A low-dose CT transmission scan (0.36mSv) was acquired for attenuation and scatter correction

### **Image processing and region of interest analysis**

#### **Neuromelanin-sensitive magnetic resonance imaging**

For the quantitative evaluation of neuromelanin-sensitive MRI scans, study-specific templates were first created using grey and white matter MPAGE segmentations from both Parkinson's disease and healthy control participants and diffeomorphic anatomical registration through exponentiated Lie algebra (DARTEL) in SPM12. Software defaults were modified to enable template and MPAGE-to-template warp field generation in native voxel size (1mm<sup>3</sup>). Neuromelanin-sensitive T1-weighted images were co-registered to the corresponding MPAGE and DARTEL warp fields applied in one step to normalise all images to common space. The neuromelanin template could then be generated through voxel-wise averaging (mean) over the entire cohort (n=45).



Mesencephalic regions of interest (ROI) for the left/right substantia nigra pars compacta (SNc) were manually defined on five contiguous axial slices of the neuromelanin template on which the neuromelanin-related hyperintensity of the SNc was visible, at the level of the red nucleus and the inferior colliculus, in FSL (Jenkinson *et al.*, 2012). The volume of the SNc on each axial slice was divided into two sections representing the ventral (SNc<sup>ven</sup>) and dorsal (SNc<sup>dor</sup>) tiers, in accordance with post-mortem histological investigations on the spatial distribution of dopamine neurons relating to the striatal afferences of the SNc (Lynd-Balta and Haber, 1994; Haber *et al.*, 2000; Haber, 2014). The decussation of the superior cerebral peduncle (CP) was also defined on the same axial slices as a reference region (Fig. 1). Once all ROIs were delineated the inverse DARTEL warp fields for each participant were applied to generate individualised ROIs for sampling in native space. This was performed in order to reduce the effect of warping/normalisation on the absolute voxel intensities within the images. Registration was checked visually in FSL before extracting mean ROI intensity values using the `fslmeants` command. Neuromelanin contrast ratios (CR) were calculated in accordance with Sasaki and colleagues (Sasaki *et al.*, 2006) using the formula (SNc-CP)/CP for ventral and dorsal tiers separately. Contrast ratios were first calculated for the most/least affected sides for each ROI before averaging across sides to yield bilateral regional CRs. Contrast ratios of the whole SNc i.e. SNc<sup>ven</sup> and SNc<sup>dor</sup> combined (SNc<sup>com</sup>) were also calculated.

### **<sup>11</sup>C-PE2I positron emission tomography**

Pre-processing and kinetic modelling for MPRAGE and <sup>11</sup>C-PE2I PET was conducted using MIAKAT™ v3.4.2. (Molecular Imaging and Kinetic Analysis Toolbox, Imanova Imaging Centre, London, UK) (Gunn *et al.*, 2016), which utilises FSL (FMRIB Image Analysis Group, Oxford, UK) (Jenkinson *et al.*, 2012), SPM (Statistical Parametric Mapping, Wellcome Trust Centre for Neuroimaging, London, UK) and in-house pre-processing and kinetic modelling procedures within an integrated PET analytical framework and is implemented in MATLAB® (Mathworks, Natick, MA, USA).

First, MPRAGE images were brain extracted, segmented and rigid-body registered to the Montreal Neurological Institute (MNI) template (Mazziotta *et al.*, 1995). MPRAGE images in “pseudo-MNI” space (MPRAGE<sub>REG</sub>) were then used for manual subcortical ROI delineation in Analyze11.0. Afferent striatal dopaminergic connections from mesencephalic neurons show a

distinctive dorsal-ventral pattern. The nigral dorsal tier projects to the ventromedial striatum while the ventral tier projects to the central associative and dorsal striatum (Lynd-Balta and Haber, 1994; Haber *et al.*, 2000; Haber, 2014). Striatal sub-division was conducted in accordance with the anatomical-landmark-based methods derived from work on the *in vivo* distribution of D3 receptors using  $^{11}\text{C}$ -(+)-PHNO (Tziortzi *et al.*, 2011). In brief, the putamen was sub-divided into 3 sections on the axial plane; posterior to the AC (post-commissural putamen), anterior to the AC and dorsal to the posterior commissure (pre-commissural dorsal putamen), anterior to the AC and ventral to the posterior commissure (pre-commissural ventral putamen). The caudate was sub-divided into two sections on the axial plane; anterior to the AC (pre-commissural caudate) and posterior to the AC (post-commissural caudate) (Fig. 1A, Fig. 1B). ROIs of the whole putamen and caudate nucleus were formed by merging pre-commissural and post-commissural sub-divisions of the putamen and caudate respectively. Furthermore, the striatum was defined as the combination of the putamen and caudate nucleus. Cerebellar grey matter was automatically defined by applying the deformation fields estimated from the spatial normalisation (12-parameter affine transformation followed by non-linear warp) of the MNI template to the MPRAGE<sub>REG</sub>, to an MNI-based regional atlas (CIC Atlas v1.2.). Grey matter segmentation maps were then applied to isolate cerebellar grey matter. ROIs for the SNc<sup>ven</sup> and SNc<sup>dor</sup>, which were defined on the study-specific template as described in the previous section, were also brought into MPRAGE<sub>REG</sub> space by applying the inverse DARTEL warp fields and rigid-body registration parameters (native-to-pseudo-MNI) in one step.

Motion correction was conducted on the dynamic PET data to correct for intra-scan head movement using frame-to-frame rigid registration with the 16<sup>th</sup> frame (corresponding to 780-900 seconds post-injection) as a reference, chosen due to high signal-to-noise ratio. No image artefacts due to substantial head movement were observed for any of the patients included in this study. Realigned frames corresponding to 10-90 minutes were subsequently summed to obtain signal-averaged (ADD) images that were then co-registered to the subjects MPRAGE<sub>REG</sub> using normalised mutual information as the cost function. The derived registration parameters were applied to bring the dynamic PET data into alignment with the MPRAGE<sub>REG</sub> and corresponding ROI map. PET image processing was evaluated at each step before ROI maps were applied to the dynamic PET frames to generate regional time-activity curves (TACs) for all ROIs on the most and least affected sides separately as well as bilaterally. The simplified reference tissue model (SRTM) (Lammertsma and Hume, 1996) was used to calculate regional non-displaceable binding potential (BP<sub>ND</sub>). Cerebellar grey matter was used as a reference as

previous studies have demonstrated negligible tracer uptake in this region (Hall *et al.*, 1999; Halldin *et al.*, 2003; Jucaite *et al.*, 2006).

Registered frames between 10-90 minutes were subsequently summed to obtain signal-averaged (ADD) images that were co-registered to the subjects MPRAGE<sub>REG</sub> using normalised mutual information as a cost function. Approximated co-registration matrices were then applied to both ADD images and realigned dynamic frames so that all images were in register with the subject MPRAGE<sub>REG</sub> and the corresponding ROI map.

< *Fig. 1* >

### **Statistical analysis**

All statistical analyses were performed using R version 3.4.3 (R Core Team, 2017) and packages “afex” (Singmann *et al.*, 2017), “emmeans” (Lenth, 2018) and “Hmisc” (Harrell Jr, 2017).

Demographic data were compared between Parkinson’s disease and healthy control groups using independent *t*-tests and Fisher’s exact test.

To assess differences in neuromelanin CRs between groups and regions of the SNc, a mixed analysis of variance (ANOVA) was conducted with group (Parkinson’s disease, healthy control) as the between-subjects factor and region (bilateral SNc<sup>ven</sup>, bilateral SNc<sup>dor</sup>) as the within-subjects factor.

To assess lateralisation of SNc neuromelanin content, CR asymmetry indices (AI) (Seibyl *et al.*, 1995) were calculated for the SNc<sup>ven</sup> and SNc<sup>dor</sup> according to clinical laterality in the Parkinson’s disease group (most/least affected) and right/left in the healthy control group and compared using one-tailed independent *t*-tests. Within the Parkinson’s disease group, lateralisation was assessed for both SNc neuromelanin content and DAT density using 2-way repeated measures ANOVA, with neuromelanin CR or <sup>11</sup>C-PE2I BP<sub>ND</sub> as the dependent variable and region (SNc<sup>ven</sup>, SNc<sup>dor</sup>) and side (most/least affected) as within-subjects factors.

Linear relationships between neuromelanin CRs in the SNc and  $^{11}\text{C}$ -PE2I BP<sub>ND</sub> values in the striatum were tested using simple linear regression. Not all regional combinations between SNc and striatum were assessed; instead, planned tests were performed based on previous anatomical work mapping connectivity between sub-regions of the substantia nigra and striatum (Lynd-Balta and Haber, 1994; Haber *et al.*, 2000; Haber, 2014). As such, regression analyses were conducted between neuromelanin CR (independent variable) in the SNc<sup>ven</sup> (Fig. 1, C7) and  $^{11}\text{C}$ -PE2I BP<sub>ND</sub> (dependent variable) in the precommissural dorsal putamen and caudate (Fig. 1, A2-3) as well as the post-commissural putamen and caudate (Fig. 1, B4-5); and between neuromelanin CR in the SNc<sup>dor</sup> (Fig. 1, C6) and  $^{11}\text{C}$ -PE2I BP<sub>ND</sub> in the pre-commissural ventral putamen, dorsal putamen and caudate (Fig. 1, A1-3). Regression analyses were also conducted between neuromelanin CR and  $^{11}\text{C}$ -PE2I BP<sub>ND</sub> values within the SNc<sup>ven</sup> and SNc<sup>dor</sup>. Analyses were performed for the most and least affected sides separately.

Pearson's product moment correlation coefficient (one-tailed) was used to explore relationships between each measure of disease severity (disease duration, UPDRS-III subscores; bradykinesia, rigidity, tremor, axial) and neuromelanin CRs for bilateral nigral regions (SNc<sup>ven</sup>, SNc<sup>dor</sup>) as well as  $^{11}\text{C}$ -PE2I BP<sub>ND</sub> for bilateral nigral and striatal sub-regions (SNc<sup>ven</sup>, SNc<sup>dor</sup>, pre-commissural dorsal putamen, pre-commissural ventral putamen, pre-commissural caudate, post-commissural putamen and post-commissural caudate).

The Benjamini-Hochberg procedure was used to control the false-discovery rate (FDR) across all planned correlation and regression analyses simultaneously, at a desired threshold of  $P < 0.05$ .

Normal distribution was assessed graphically using normal probability plots as appropriate, extreme values were evaluated using Z-scores, boxplots and studentized residuals, independence of observations using Durbin-Watson statistic and homoscedasticity assessed by residual plots and Levene's test.

## **Data Availability**

The authors confirm that data presented in this article is original. The data that support the findings of this study are available from the corresponding author, upon reasonable request.

## RESULTS

### Demographics

Clinical and demographic characteristics for the Parkinson's disease and healthy control groups are summarized in Table 1. There were no significant differences in age or gender between groups.

< Table 1 >

### Neuromelanin in the substantia nigra in Parkinson's disease compared to healthy controls

Mixed ANOVA showed a main effect of group ( $F_{1,43}=11.95, P=0.001$ ), in which neuromelanin CRs were significantly lower in Parkinson's disease (Mean<sub>EM</sub>=0.095, 95% CI [0.081, 0.109]) as compared to the healthy controls (Mean<sub>EM</sub>=0.129, 95% CI [0.115, 0.143]). There was also a main effect of region ( $F_{1,43}=41.35, P < 0.001$ ) whereby SNc<sup>ven</sup> (Mean<sub>EM</sub>=0.100, 95% CI [0.090, 0.110]) showed significantly lower neuromelanin CRs than SNc<sup>dor</sup> (Mean<sub>EM</sub>=0.124, 95% CI [0.114, 0.134]). However, no significant group by region interaction was detected ( $F_{1,43}=0.71, P=0.404$ ) (Fig. 2). Including age and gender as covariates did not alter results.

< Fig. 2 >

### Lateralisation of neuromelanin and dopamine transporters in Parkinson's disease

Asymmetry indices showed no significant differences between Parkinson's disease and healthy control groups for SNc<sup>ven</sup> ( $t_{43}=0.75, P=0.230$ ) or SNc<sup>dor</sup> ( $t_{43}=1.42, P=0.081$ ). Removal of one extreme outlier (Table 2) did not alter results.

Two-way repeated measures ANOVA conducted on the Parkinson's disease group revealed no significant interaction between region (SNc<sup>ven</sup>, SNc<sup>dor</sup>) and side (most/least affected) on either neuromelanin CR ( $F_{1,29}=0.17, P=0.685$ ) (Fig. 3, A) or <sup>11</sup>C-PE2I BP<sub>ND</sub> ( $F_{1,29}=0.34, P=0.563$ ) (Fig. 3, B). However, for neuromelanin CR, there was a marginally significant main effect of side ( $F_{1,29}=4.85, P=0.036$ ), whereby neuromelanin CR was lower in the most affected hemisphere (Mean=0.097, 95% CI [0.084, 0.109]) than in the least affected hemisphere (Mean=0.104, 95% CI [0.092, 0.117]), and a main effect of region ( $F_{1,29}=36.19, P<0.001$ ) in

which  $\text{SNc}^{\text{ven}}$  (Mean=0.087, 95% CI [0.074, 0.100]) was significantly lower than  $\text{SNc}^{\text{dor}}$  (Mean=0.114, 95% CI [0.101, 0.127]). Similarly for  $^{11}\text{C-PE2I BP}_{\text{ND}}$ , there was a marginally significant main effect of side ( $F_{1,29}=4.21$ ,  $P=0.049$ ), in which  $^{11}\text{C-PE2I BP}_{\text{ND}}$  was lower in the most affected hemisphere (Mean=0.633, 95% CI [0.584, 0.684]) than in the least affected hemisphere (Mean=0.667, 95% CI [0.618, 0.717]), and a main effect of region ( $F_{1,29}=76.39$ ,  $P<0.001$ ), whereby  $\text{SNc}^{\text{ven}}$  (Mean=0.607, 95% CI [0.558, 0.655]) was significantly lower than  $\text{SNc}^{\text{dor}}$  (Mean=0.694, 95% CI [0.646, 0.743]). Including age and gender as covariates did not alter results.

< Fig. 3 >

In terms of percentage of regional neuromelanin CR reduction, the Parkinson's disease group showed a ventro-dorsal pattern of neurodegeneration as compared to healthy controls as follows  $\text{SNc}^{\text{ven}}$  (-30.03%) >  $\text{SNc}^{\text{dor}}$  (-21.45%) (Table 2).

< Table 2 >

### **Relationship between neuromelanin and DAT density in the nigrostriatal system**

On the most affected side, significant positive relationships were found between neuromelanin CR in the  $\text{SNc}^{\text{ven}}$  and  $^{11}\text{C-PE2I BP}_{\text{ND}}$  in the pre-commissural dorsal putamen ( $F_{1,28}=13.82$ ,  $P=0.001$ ) and caudate ( $F_{1,28}=15.13$ ,  $P=0.001$ ) and the post-commissural putamen ( $F_{1,28}=13.05$ ,  $P=0.001$ ) (Fig. 4). Neuromelanin CR in the  $\text{SNc}^{\text{dor}}$  was positively associated with  $^{11}\text{C-PE2I BP}_{\text{ND}}$  in the pre-commissural dorsal putamen ( $F_{1,28}=7.99$ ,  $P=0.009$ ), and caudate ( $F_{1,28}=23.12$ ,  $P<0.001$ ) (Fig. 5). Trends were found between neuromelanin CR in the  $\text{SNc}^{\text{ven}}$  and  $^{11}\text{C-PE2I BP}_{\text{ND}}$  in the post-commissural caudate ( $F_{1,28}=5.13$ ,  $P=0.031$ ) and between neuromelanin CR in the  $\text{SNc}^{\text{dor}}$  and  $^{11}\text{C-PE2I BP}_{\text{ND}}$  in the pre-commissural ventral putamen ( $F_{1,28}=5.22$ ,  $P=0.030$ ), however, after Benjamini-Hochberg FDR correction across all correlational tests, these results did not remain significant. Inclusion of additional covariates age and gender in the above regression models had only minimal impact upon the beta coefficients associated with the main independent variable (difference of  $3.96\pm 2.92\%$ ) and did not result in changes to significance level. No significant relationship was found between neuromelanin CR and  $^{11}\text{C-PE2I BP}_{\text{ND}}$  within either  $\text{SNc}^{\text{ven}}$  or  $\text{SNc}^{\text{dor}}$  (Fig. 4, Fig. 5).

On the clinically least affected side, there were no significant relationships between neuromelanin CR and  $^{11}\text{C}$ -PE2I BP<sub>ND</sub> within the ventral or dorsal SNc or with  $^{11}\text{C}$ -PE2I BP<sub>ND</sub> in striatal sub-regions (Fig. 4, Fig. 5).

< *Fig. 4* >

< *Fig. 5* >

### **Relationship between clinical severity and neuromelanin and DAT density**

SNc neuromelanin CR was inversely related to disease duration, particularly within the ventral tier, while no association with UPDRS-III bradykinesia, rigidity or axial subscales was evident. In contrast, SNc  $^{11}\text{C}$ -PE2I BP<sub>ND</sub> appeared to show an opposing pattern whereby negative correlations were evident with UPDRS-III bradykinesia, rigidity or axial subscales but not disease duration. In the striatum,  $^{11}\text{C}$ -PE2I BP<sub>ND</sub> appeared to correlate relatively consistently with both disease duration and UPDRS-III bradykinesia, rigidity and axial subscales. No significant correlations were found between UPDRS-III tremor and any imaging measure (Fig. 6).

< *Fig. 6* >

## **DISCUSSION**

### **Neuromelanin loss in the substantia nigra**

The present study assessed the integrity of nigrostriatal pathways *in vivo* in a moderate-stage Parkinson's disease cohort by using neuromelanin-sensitive MR imaging and  $^{11}\text{C}$ -PE2I PET. As expected, Parkinson's disease subjects had reduced neuromelanin when compared to controls, in line with previous investigations. While Sasaki *et al.* (2006) in their seminal paper examining neuromelanin *in vivo* report contrast ratio values almost double to what is reported here, our data show a larger effect size in equivalent tests ( $d = 1.1$  and  $0.8$ ), with both studies having similar sensitivity to detect  $d > 0.8$  with 80% power at  $\alpha$  of 0.05. The disparity could be due to disease severity with the current cohort at 6.8 years of illness as compared to 2.5 years, or to the size and placement of regions-of-interest. The size of the circular cursors used for

sampling by Sasaki *et al.* (2006) was not detailed, though they state that pixels in the high signal intensity areas were measured. In contrast, the semi-automated sampling method used here was not as influenced by the local hyperintensity in individual subjects. In addition, our cohort was ~15 years younger.

### **Relationship between nigral neuromelanin and striatal dopamine transporters**

Our findings demonstrate that the relationship between nigral neuromelanin and striatal DAT in Parkinson's disease exhibits a remarkably lateralised pattern between the most and least affected brain hemispheres. Moderate to strong linear associations between the two measures were detected across the two regions but only in the most affected side. It is possible this lateralisation could be explained by ceiling effects and/or hemispheric lagging akin to that which has been observed in Parkinson's disease with striatal dopaminergic markers such as aromatic L-amino acid decarboxylase, vesicular monoamine transporter (type 2) and dopamine transporter (Lee *et al.*, 2000; Nandhagopal *et al.*, 2009). However, this alone seems unlikely given that significant depigmentation was found in both sides of the nigra, indicating that the neurodegenerative process in the least affected hemisphere had already begun in our cohort. If we assume that the neurodegenerative process occurs in a similar manner for both hemispheres, we might expect some degree of correlation between the two imaging measures on the least affected side, especially between regions with substantial SNc cell and striatal DAT loss. That this was not the case may be strong evidence to the contrary.

Alternatively, these measures might reflect different aspects of disease progression. In a comprehensive post-mortem investigation, Kordower and colleagues recently observed that loss of melanin-containing neurons in the SNc was consistently outweighed by loss of tyrosine hydroxylase-positive neurons in the first two decades of illness (Kordower *et al.*, 2013). Thus while DAT imaging may yield markers reflecting dopaminergic phenotype and neuronal dysfunction, neuromelanin markers may relate more closely to structure and neurodegeneration. It is possible this may explain why here, DAT tended to correlate with bradykinetic/rigid/axial severity while nigral neuromelanin correlated with disease duration, particularly within the nigra. Interestingly, these differential trajectories appear to converge and become less variable over time (Kordower *et al.*, 2013). It is feasible that this could account for the lateralisation demonstrated here, where associations become apparent only when the extent of nigral neuromelanin loss comes in line with that of tyrosine hydroxylase-positive cell density.



## **Relationship between neuromelanin and dopamine transporters within the substantia nigra**

In line with evidence from post-mortem data, Parkinson's disease subjects displayed a ventral to dorsal pattern of nigral depigmentation (Fearnley and Lees, 1991; Gibb and Lees, 1991; Kordower *et al.*, 2013) and showed a tendency for greater loss in the clinically-defined most affected side. Despite DAT density following a similar pattern of distribution we did not observe any relationship between the two imaging markers. Similar results have also been shown in a small cohort of young healthy males (Ito *et al.*, 2017). Recent nuclear imaging studies using  $^{11}\text{C}$ -FeCIT and  $^{18}\text{F}$ -FE-PE2I PET in early and *de novo* patients have shown that the loss of DAT in the striatum (-35-70%) exceeds the loss in the substantia nigra (-25-30%), as compared to healthy controls (Caminiti *et al.*, 2017; Fazio *et al.*, 2018). One explanation for this relates to evidence suggesting that distal axonal degeneration occurs initially before proceeding retrograde towards the cell body (Calo *et al.*, 2016; Kurowska *et al.*, 2016; Tagliaferro and Burke, 2016). If this is the case then the relatively modest DAT loss in the nigra may represent delayed, slower or variable progression rate to that in the striatum. While this is yet to be studied, it could explain why the relationship with DAT differs between the nigra and striatum.

## **Comparison with previous studies**

A few studies (Colloby *et al.*, 2012; Kraemmer *et al.*, 2014; Saari *et al.*, 2017) have evaluated the relationship between nigral neuromelanin and striatal DAT, using ante-mortem DAT-specific SPECT ( $^{123}\text{I}$ -FP-CIT or  $^{123}\text{I}$ - $\beta$ -CIT) with either post-mortem histochemistry of the SNc (conducted ~2.5-5 years after SPECT assessment) or neuromelanin-sensitive MRI. Positive correlations have been noted in a mixed dementia cohorts (Colloby *et al.*, 2012) and in general neurological samples (Kraemmer *et al.*, 2014; Kuya *et al.*, 2016) and were in line with those found in smaller Parkinson's disease samples (Isaias *et al.*, 2016) and in the current Parkinson's disease-only cohort.

In contrast, Saari and colleagues found no significant correlations between nigral neuromelanin and striatal DAT in a small group of 11 Parkinson's disease and 7 individuals with mixed parkinsonism (Saari *et al.*, 2017). This may have been due to lack of power and/or inclusion of mixed parkinsonism, as correlation coefficients appear to graduate from zero/weak to moderate

upon removal of non-Parkinson's disease patients from the analysis. The authors suggested that the relationship between nigral neuromelanin cell density and striatal DAT may dissipate as the disease progresses. Indeed, while results showing that nigral neuronal density in a mixed dementia cohort accounts for 58%, 40% and 20% of the variance in posterior and anterior putamen and caudate DAT respectively, they appear to be driven mostly by non-Parkinson's disease individuals (Colloby *et al.*, 2012), with the Parkinson's disease dementia data tending towards an asymptote. In the current Parkinson's disease cohort, although significant relationships were found between both tiers of the nigra and their striatal afferents in the most affected side, it appeared that this association was strongest between the dorsal tier and pre-commissural caudate, which retains the highest DAT expression across the striatum (Nandhagopal *et al.*, 2009; Oh *et al.*, 2012; Han *et al.*, 2016). However, normative data from a group of healthy young men show that nigral neuromelanin accumulation does not correlate with nigral DAT at baseline (Ito *et al.*, 2017). In addition, preclinical work demonstrates correlations in more pathologically advanced 6-OHDA mouse models but not in mild MPTP regimes (Alvarez-Fischer *et al.*, 2007). Thus, it is possible that the relationship between nigral neuromelanin content and striatal DAT is constrained by both ceiling and floor effects and may be evident only at some stages of Parkinson's disease progression.

Isaias and colleagues found no relationship between neuromelanin and DAT asymmetries, performing analyses on the whole nigra, putamen and caudate (Isaias *et al.*, 2016). Moreover, correlational analyses of absolute values involved collapsing across the most/least affected sides. Others have found positive results but using asymmetry indices calculated between left and right hemispheres (Kraemmer *et al.*, 2014; Kuya *et al.*, 2016), thus limiting pathological relevance and interpretability. The only study to perform correlations separately for most/least affected sides showed significant associations between asymmetry indices but no correlations of absolute values (Saari *et al.*, 2017). Importantly however, these authors distinguished most/least affected sides as those with higher/lower cell count and/or DAT density. In the present study, while we found greater demelanisation in the clinically most affected side we also noted that a significant proportion of our patients (~38%) displayed the opposite i.e. greater neuromelanin loss in the clinically least affected side, which was discordant to the side with greatest DAT loss. This finding has been discussed recently (Isaias *et al.*, 2016) and possibly stands as an important methodological factor to explain the discordance between results.

## Limitations and considerations

There are some limitations in the current report. Delineation of the SNc was based on the hyperintense area of the midbrain on neuromelanin MR images as it is not easily visible on standard structural scans, which could introduce bias towards overall greater values. We attempted to resolve this by employing an automated procedure in which the SNc was delineated on neuromelanin templates created via normalisation of structural scans from both Parkinson's disease and healthy control groups. In doing so, we were able to remain objective and consistent across individuals. Moreover, our data correspond well with percentage losses in a less advanced subset of 9 patients (1-14 year disease duration, Mean = 8 years) from Kordower and colleagues (Kordower *et al.*, 2013), whose values indicate ~22% loss in the dorsal and ~35% in the ventral tiers. Second, while striatal regions of interest were defined according to anatomical-landmark-based guidelines derived from a recent study on *in vivo* distribution of D3 receptors, parcellation could potentially be improved using connectivity-based methods such as probabilistic tractography (Chowdhury *et al.*, 2013). Age and gender have recently been shown to have significant effects over neuromelanin levels in healthy individuals (Xing *et al.*, 2018). While we attempted to account for this through addition of covariates in our analyses, this constitutes incomplete control and thus the influence of these variables should be considered here and in future work. In addition, although striatal DAT in Parkinson's disease is well characterised using mostly SPECT ligands such as <sup>123</sup>I-FP-CIT, we did not obtain <sup>11</sup>C-PE2I scans for healthy controls. This would have enabled parallel analysis from which we could ascertain the normative nigrostriatal state across both brain hemispheres using the two imaging markers. Lastly, it has been shown that chronic exposure to dopaminergic drugs including levodopa and dopamine agonists can down-regulate striatal DAT to varying degrees (-4-7.2%), depending on the exposure dose (Guttman *et al.*, 2001; Fahn *et al.*, 2004). As such, chronic exposure should be considered as a confounding factor in the current study.

## Conclusions

The current study provides important insights into the relationship between neuromelanin content in the SNc and striatal DAT density in moderate stage Parkinson's disease, as measured *in vivo* using neuromelanin-sensitive MRI and <sup>11</sup>C-PE2I PET. Reduction of nigral pigmentation in Parkinson's disease displays an uneven pattern of association with the loss of striatal dopaminergic function towards the clinically most affected side while no relationship was found with nigral DAT. These findings may be indicative of a lag in disease progression or differences

in the pathologic processes measured that could manifest with heterogeneous rate of decline, convergence and symmetry. However, further work including longitudinal imaging assessment on the demelanisation trajectories in the nigra, in both tiers and on the most and least affected side would provide a strong basis from which we could start to understand the relationship between these pathologic processes.

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## **COMPETING INTERESTS**

The authors report no competing interests.



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**Figure 1: Region of interest definition in the substantia nigra and striatum.** Bottom right is a sagittal midline view of a 3D T1-weighted MPRAGE template, derived using DARTEL with 30 Parkinson's disease and 15 healthy control participants. The red line annotations illustrate the position of the coronal and axial planes corresponding to the sections depicted in panels A, B and C. Panels **A and B** represent  $^{11}\text{C}$ -PE2I ADD images overlaid onto the structural MPRAGE<sub>REG</sub> from two levels of the coronal plane either side of the anterior commissure along the y-axis. Regions of the pre-commissural striatum (**Panel A**) consist of ventral putamen (1 - red), dorsal putamen (2 - green) and caudate (3 - purple). Regions of the post-commissural striatum (**Panel B**) include putamen (4 - orange) and caudate (5 - blue). **Panel C** shows an axial slice of the neuromelanin-sensitive T1-weighted template derived using DARTEL (n = 45) at the level of the substantia nigra pars compacta, including delineation of the dorsal (6 - red) (SNc<sup>dor</sup>) and ventral (7 - blue) (SNc<sup>ven</sup>) tiers, as well as cerebral peduncle (8 - yellow). Descriptive anatomy demonstrates projections from the dorsal tier (6) to striatal regions 1-3 while the ventral tier (7) projects to regions 2-5 (Lynd-Balta and Haber, 1994; Haber *et al.*, 2000; Haber, 2014). MPRAGE = Three-dimensional T1-weighted magnetization prepared rapid acquisition gradient echo; ROI = region of interest;  $^{11}\text{C}$ -PE2I = ( $^{11}\text{C}$ )N-(3-iodopro-2E-enyl)-2 $\beta$ -carbomethoxy-3 $\beta$ -(4'-methylphenyl)nortropane); ADD = signal-averaged images; SNc<sup>ven</sup> = substantia nigra pars compacta ventral tier; SNc<sup>dor</sup> = substantia nigra pars compacta dorsal tier.

**Figure 2: Neuromelanin loss in the ventral and dorsal tiers of the substantia nigra pars compacta in Parkinson's disease as compared to healthy controls.** Strip plot illustrating neuromelanin CR values for individuals of both Parkinson's disease and healthy control groups and for each sub-region of the substantia nigra pars compacta (ventral tier, dorsal tier). Solid black points represent individuals within the Parkinson's disease group and hollow grey points represent individuals within the healthy control group. Crossbars represent 95% confidence intervals about the estimated marginal mean derived from the mixed ANOVA analysis. PD =

Parkinson's disease; HC = Healthy control; SNc = substantia nigra pars compacta; CR = neuromelanin contrast ratio.

**Figure 3: Lateralisation of neuromelanin loss and DAT density in the ventral and dorsal tiers of the substantia nigra pars compacta for the clinically-defined most and least affected sides.** Strip plots showing neuromelanin CR (A) and  $^{11}\text{C}$ -PE2I  $\text{BP}_{\text{ND}}$  (B) values for individuals within the Parkinson's disease group as a function of the clinically-defined most/least affected side and sub-region of the substantia nigra pars compacta (ventral tier, dorsal tier). Solid blue points represent the ventral tier while hollow red points represent the dorsal tier. Points between the clinically-defined most and least affected sides are connected with either a solid blue line (ventral tier) or dashed red line (dorsal tier) to illustrate asymmetry within individuals. Crossbars represent 95% confidence intervals about the mean. SNc = substantia nigra pars compacta; DAT = dopamine transporter; CR = neuromelanin contrast ratio;  $\text{BP}_{\text{ND}}$  = non-displaceable binding potential;  $^{11}\text{C}$ -PE2I = ( $^{11}\text{C}$ ]N-(3-iodopro-2E-enyl)-2 $\beta$ -carbomethoxy-3 $\beta$ -(4'-methylphenyl)nortropane).

**Figure 4: Association between neuromelanin in the substantia nigra pars compacta ventral tier and nigrostriatal dopamine transporter binding.** Scatterplots and simple linear regression showing positive relationships between neuromelanin CR in the  $\text{SNc}^{\text{ven}}$  and  $^{11}\text{C}$ -PE2I  $\text{BP}_{\text{ND}}$  in the pre-commissural dorsal putamen (A), pre-commissural caudate (B), post-commissural putamen (C), post-commissural caudate (D) and  $\text{SNc}^{\text{ven}}$  (E) in the clinically-defined most affected (solid blue points, solid line) and least affected sides (hollow black points, dashed line). Grey shaded areas represent 95% confidence intervals. \*\*indicates significant result following Benjamini-Hochberg FDR correction for all correlational tests. \*indicates significance at  $P < 0.05$ . MA = most affected; LA = least affected;  $\text{SNc}^{\text{ven}}$  = substantia nigra pars compacta ventral tier; CR = neuromelanin contrast ratio;  $\text{BP}_{\text{ND}}$  = non-displaceable binding potential;  $^{11}\text{C}$ -PE2I = ( $^{11}\text{C}$ ]N-(3-iodopro-2E-enyl)-2 $\beta$ -carbomethoxy-3 $\beta$ -(4'-methylphenyl)nortropane).

**Figure 5: Association between neuromelanin in the substantia nigra pars compacta dorsal tier and nigrostriatal dopamine transporter binding.** Scatterplots and simple linear

regression showing positive relationships between neuromelanin CR in the SNc<sup>dor</sup> and <sup>11</sup>C-PE2I BP<sub>ND</sub> in the pre-commissural dorsal putamen (A), pre-commissural ventral putamen (B), pre-commissural caudate (C) and SNc<sup>dor</sup> (D) in the clinically-defined most affected (solid red points, solid line) and least affected sides (hollow black points, dashed line). Grey shaded areas represent 95% confidence intervals. \*\*indicates significant result following Benjamini-Hochberg FDR correction for all correlational tests. \*indicates significance at  $P < 0.05$ . MA = most affected; LA = least affected; SNc<sup>dor</sup> = substantia nigra pars compacta dorsal tier; CR = neuromelanin contrast ratio; BP<sub>ND</sub> = non-displaceable binding potential; <sup>11</sup>C-PE2I = ([<sup>11</sup>C]N-(3-iodopro-2E-enyl)-2 $\beta$ -carbomethoxy-3 $\beta$ -(4'-methylphenyl)nortropene).

**Figure 6: Relationship between measures of clinical severity and regional neuromelanin and dopamine transporter binding.** Pearson's correlation matrix with measures of clinical severity by row and neuromelanin CR in the SNc and <sup>11</sup>C-PE2I BP<sub>ND</sub> in the SNc and striatum by column. Tiles are colour coded and labelled with the value of the correlation coefficient. \*\*indicates significant result following Benjamini-Hochberg FDR correction for all correlational tests. \*indicates significance at  $P < 0.05$ . Pre-C = Pre-Commissural; Post-C = Post-Commissural; SNc<sup>dor</sup> = substantia nigra pars compacta dorsal tier; SNc<sup>ven</sup> = substantia nigra pars compacta ventral tier; UPDRS-III = Unified Parkinson's Disease Rating Scale Part III; BP<sub>ND</sub> = non-displaceable binding potential; <sup>11</sup>C-PE2I = ([<sup>11</sup>C]N-(3-iodopro-2E-enyl)-2 $\beta$ -carbomethoxy-3 $\beta$ -(4'-methylphenyl)nortropene).