## **Supplementary Information**

### The Neuromelanin-related T<sub>2</sub>\* Contrast in Postmortem Human Substantia Nigra with 7T MRI

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**Supplementary Figure S1.** The schematic diagram of the histology-MRI image registration. The binary Perl image (B-I) (from the Perl stain image (B)) was registered to the binary KB image (A-I) (from the KB stain image (A)) to yield an aligned binary Perl image (B-II). Then, the resulting transformation information was used to warp the RGB channel information from the original Perl stain image (B) to generate the aligned Perl stain image (B-III). The original T<sub>1</sub>WI (C-I) was registered to the KB stain image (A) using red channel information, to yield the aligned T<sub>1</sub>WI (C-II). The same transformation information was used to warp the original T<sub>2</sub>\*WI (D-I) to the registered T<sub>2</sub>\*WI (D-II).



**Supplementary Figure S2.** Neuron-occupied areas (A-II and B-II) were extracted from 2D KB (A-I) and TH (B-I) stained images. Iron pigments (C-III) and unstained pigmented NM (C-II) were also extracted from 2D Perl stained images (C-I, the same two figures) to obtain iron and NM density maps. Then, the number of pixels occupied by neurons or pigments per 10×10 block was recorded to generate a density map (% occupied by neurons) as shown in D-II (only TH case shown). Then, the TH-positive neuron density image (D-II) ranging from 0 to 100% was registered to the closest KB stain image to yield the aligned density image (E-II) using the same transformation information to warp from the original TH stain image (D-I) to the aligned image (E-I).



**Supplementary Figure S3.** The polygonal regions of interest (ROIs). ROI-whole SN (A) based on the KB stain corresponding to the most of SN hypointensity on the  $T_2*WI$  (B), ROI-SNc (the substantia nigra pars compacta) based on the TH stain within the ROI-whole SN including the A9 cell group (C), and ROI-SNr (the substantia nigra pars reticulata) obtained by subtracting ROI-SNc from ROI-whole SN (D).



**Supplementary Figure S4.** The ex vivo substantia nigra MRI protocol (70-year-old female subject, at the rostral level the exiting third cranial nerve fibers). For the  $T_2*WI$ , an echo time of TE = 15.4 ms yielded the best visual MRI contrast among the ten different TEs, ranging from 3.1 to 40 ms (A, B, C). For the color-coded  $T_2*$  map (D), the color bar represents  $T_2*$  values (ms). The  $T_1WI$  (E) presents arch-shaped boundaries between the SN and the crus cerebri more distinctly than  $T_2*WI$ . The magnetization transfer  $T_1$ -weighted image (F) is unable to accurately depict NM-related contrasts in the substantia nigra pars compacta.

### **Supplementary Table S1**

Multiple R	Subject I	Subject II
T <sub>2</sub> * with Iron	0.49	0.31
T <sub>2</sub> * with Iron, NM	0.56	0.70
T <sub>2</sub> * with Iron, NM, Nissl	0.56	0.70
$T_2^*$ with Iron, NM, Nissl, TH	0.58	0.71

**Supplementary Table S1.** Coefficients of multiple correlation (multiple R) for  $T_2^*$  with histologic measures such as iron, NM, Nissl, and TH. These show that Nissl and TH are almost co-linear with NM and Iron since adding Nissl and TH to linear models did not increase multiple R much.

### **Supplementary Table S2**

Pearson's Correlation (Spearman's Correlation)	Nissl	ТН	NM
Subject I (six slices)			
TH	0.59* (0.68*)		
NM	0.82* (0.82*)	0.62* (0.66*)	
Iron	0.07 (0.30*)	0.01 (0.19*)	0.05 (0.30*)
Subject II (six slices)			
TH	0.69* (0.77*)		
NM	0.85* (0.86*)	0.69* (0.75*)	
Iron	-0.27* (-0.33*)	-0.17* (-0.12**)	-0.25* (-0.27*)

TH: tyrosine hydroxylase, NM: neuromelanin. \* P < 0.0001, \*\* P < 0.005

**Supplementary Table S2.** Pearson's correlations (Spearman's correlations in parentheses) between the densities of the histological components.