Biondetti et al. Supplementary Material

Supplementary Table I Magnetic resonance imaging protocols

The state of a second second	TI-weighted	T2*-weighted iron- sensitive MRI	
0	neuromelanin-sensitive		
МКІ	MRI		
3D MP2RAGE	2D TSE	3D FLASH	
	Transverse, perpendicular to		
Sagittal	the longitudinal axis of the	Transverse	
	brainstem		
Anterior-posterior	Anterior-posterior	Right-left	
2.00	12	4, 7, 10, 13, 16, 19, 22, 25, 28,	
2 70	13	31, 34 and 37	
5000	890	40	
n/a	3	12	
700 and 2500	n/a	n/a	
4 and 5	90 and 180 (refocus)	20	
240	160	1000	
256 x 231 x 176	200 × 220 × 48	192 x 174 x 160	
x x		x x 2 reconstructed as	
	0 43 X 0 43 X 3	isotropic	
<i>n</i> /a	Cline coloct divertion	First-echo only, all encoding	
n/a	Slice-select direction	directions	
08:12	06:55	09:18	
	Sagittal Anterior-posterior 2 .98 5000 n/a 700 and 2500 4 and 5 240 256 x 231 x 176 1 x 1 x 1 n/a	T1-weighted anatomical MRIneuromelanin-sensitive MRI3D MP2RAGE2D TSE Transverse, perpendicular to the longitudinal axis of the brainstemSagittalthe longitudinal axis of the brainstemAnterior-posteriorAnterior-posterior2.98135000890 n/an/a3700 and 2500n/a4 and 590 and 180 (refocus)240160256 x 231 x 176200 x 220 x 481 x 1 x 10.43 x 0.43 x 3n/aSlice-select direction	

Parameter settings for all the MRI sequences

FLASH = fast low angle shot; FoV = field of view; MP2RAGE = magnetization prepared two rapid acquisition gradient echoes; n/a = not applicable; TE = echo time; TI = inversion time; TR = repetition time; TSE = turbo spin echo

Supplementary results

Following MRI quality control, we excluded neuromelanin-sensitive images acquired in one HC, two iRBD patients and five Parkinson's disease patients at V1, and two PDs at V2, mostly because of incorrect field of view placement excluding part of the substantia nigra pars compacta (SNc) from the image. Iron-sensitive images were unavailable in ten HCs, five iRBD patients and twenty-two Parkinson's disease patients at V1, and one HC, seven iRBD patients and twenty-two Parkinson's disease patients at V2.

The unavailability of some iron-sensitive MRI data sets was caused by errors in raw data transmission from the MRI system to the image reconstruction software. Raw data reconstruction was required for quantitative susceptibility mapping (QSM), which exploits the MRI signal phase, because the Siemens default multi-channel combination procedure

calculates the MRI phase incorrectly. This is a well-known issue in the QSM community.¹ Several solutions have been developed to overcome this issue, such as the ad-hoc reconstruction method implemented in our imaging center.^{1, 2} However, raw data retrieval is more susceptible to data communication issues from the MRI system. For future studies, we aim to try and retrieve these missing data.

For group difference analysis, all participants were included, as each contributed at least one measurement. For correlation analysis, only participants with corresponding complete pairs of measurements were included. The number of subjects retained for each analysis is listed in Supplementary Table 2.

Supplementary Table 2 Subjects retained for the analyses following MRI quality control	

	HCs		iRBDs		PDs	
	VI	V 2	VI	V 2	VI	V 2
Group difference / temporal	fitting analyses:					
Neuromelanin	54	28	41	21	130	81
Iron	45	27	38	14	113	61
DaT	35	18	32	15	48	40
Correlation analyses:						
DaT-neuromelanin	n/a	n/a	30	n/a	46	n/a
DaT-iron	n/a	n/a	28	n/a	43	n/a
Neuromelanin-iron	n/a	n/a	35	n/a	110	n/a

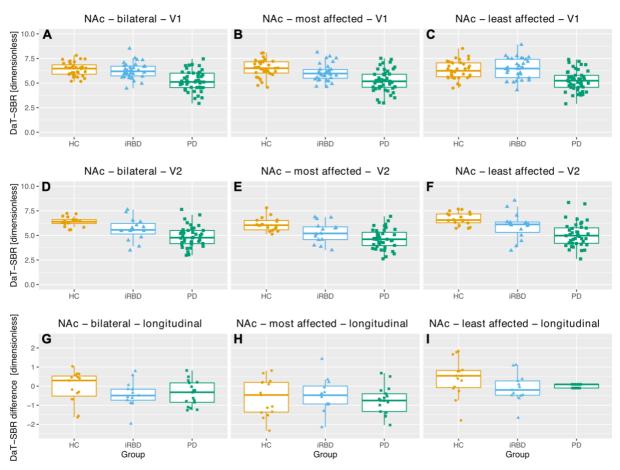
For each subject group and visit, the table shows the number of subjects retained for each analysis. Group difference and temporal fitting analyses only required one measurement in each subject. Correlation analyses required pairs of measurements in each subject. Notably, correlation analyses were only performed on subjects at VI

DaT = dopamine transporter; HC = healthy control subject; iRBD = idiopathic REM sleep behaviour disorder; n/a = not applicable; PD = Parkinson's disease; V1/V2 = visit one or two

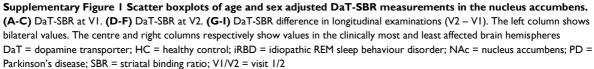
	Most affected hemisphere			Least affected h	Least affected hemisphere		
	Estimated	Estimated	Estimated	Estimated DaT-	Estimated	Estimated iRBD	
	DaT-SBR loss	beginning of the	iRBD time	SBR loss at PD	beginning of the	time placement	
	at PD onset	prodromal phase	placement	onset [%]	prodromal phase	[years]	
	[%]	in PD [years]	[years]		in PD [years]		
NAc	8.0	-1.1	0.1	9.0	-1.8	-1.3	
Limbic Cd	12.7	-1.9	-0.4	12.5	-3.2	-2.1	
Associative Cd	23.6	-4.4	-2.6	13.9	-2.3	-1.4	
Sensorimotor	27.1	-5.6	-3.8	17.0	-3.0	-1.7	
Cd							
Limbic Pu	29.7	-6.1	-4.4	19.6	-3.4	-2.9	
Associative Pu	48.0	-12.7	-10.3	34.8	-5.6	-4.6	
Sensorimotor	54.5	-20.7	-16.5	42.2	-7.8	-6.0	
Pu							

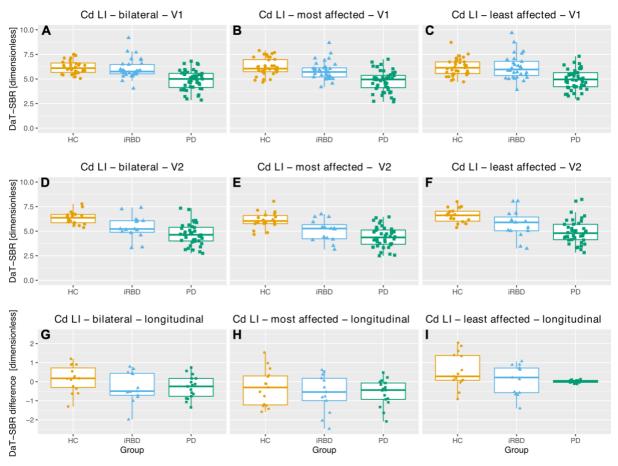
Supplementary Table 3 Estimated loss in DaT SBR over disease duration

Cd = caudate nucleus; DaT = dopamine transporter; iRBD = idiopathic REM sleep behaviour disorder; NAc = nucleus accumbens; PD = Parkinson's disease; Pu = putamen; SBR = striatal binding ratio

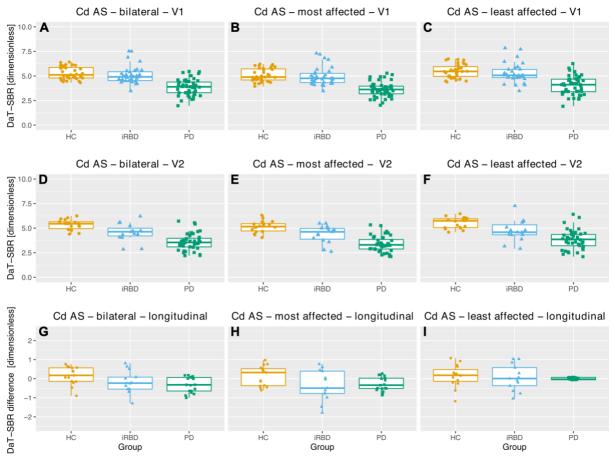


Supplementary figures

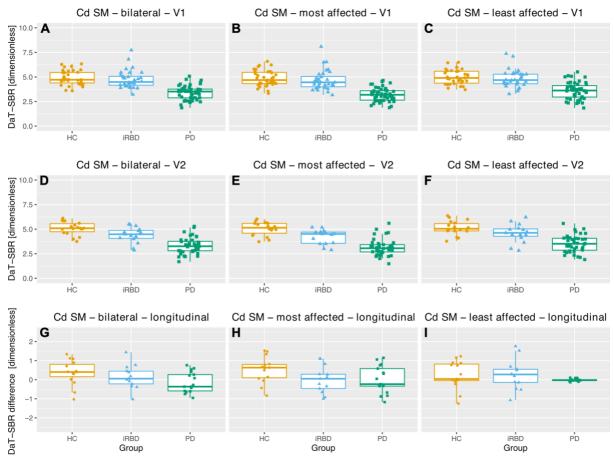




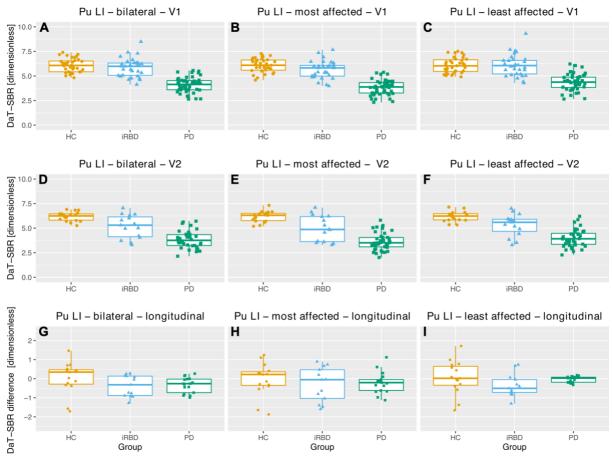
Supplementary Figure 2 Scatter boxplots of age and sex adjusted DaT-SBR measurements in the limbic caudate nucleus. (A-C) DaT-SBR at VI. (D-F) DaT-SBR at V2. (G-I) DaT-SBR difference in longitudinal examinations (V2 – VI). The left column shows bilateral values. The centre and right columns respectively show values in the clinically most and least affected brain hemispheres Cd = caudate nucleus; DaT = dopamine transporter; HC = healthy control; iRBD = idiopathic REM sleep behaviour disorder; LI = limbic; PD = Parkinson's disease; SBR = striatal binding ratio; VI/V2 = visit 1/2



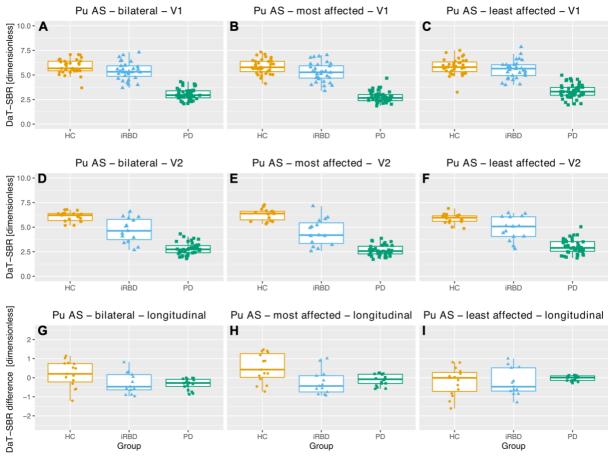
Supplementary Figure 3 Scatter boxplots of age and sex adjusted DaT-SBR measurements in the associative caudate nucleus. (A-C) DaT-SBR at V1. (D-F) DaT-SBR at V2. (G-I) DaT-SBR difference in longitudinal examinations (V2 – V1). The left column shows bilateral values. The centre and right columns respectively show values in the clinically most and least affected brain hemispheres AS = associative; Cd = caudate nucleus; DaT = dopamine transporter; HC = healthy control; iRBD = idiopathic REM sleep behaviour disorder; PD = Parkinson's disease; SBR = striatal binding ratio; V1/V2 = visit 1/2



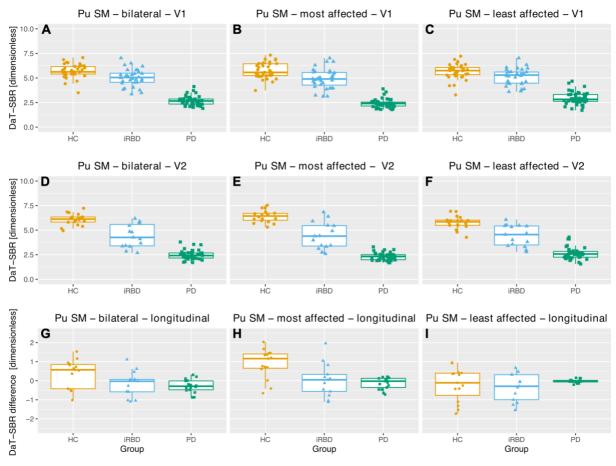
Supplementary Figure 4 Scatter boxplots of age and sex adjusted DaT-SBR measurements in the sensorimotor caudate nucleus. (A-C) DaT-SBR at V1. (D-F) DaT-SBR at V2. (G-I) DaT-SBR difference in longitudinal examinations (V2 – V1). The left column shows bilateral values. The centre and right columns respectively show values in the clinically most and least affected brain hemispheres Cd = caudate nucleus; DaT = dopamine transporter; HC = healthy control; iRBD = idiopathic REM sleep behaviour disorder; PD = Parkinson's disease; SBR = striatal binding ratio; SM = sensorimotor; V1/V2 = visit 1/2



Supplementary Figure 5 Scatter boxplots of age and sex adjusted DaT-SBR measurements in the limbic putamen. (A-C) DaT-SBR at VI. (D-F) DaT-SBR at V2. (G-I) DaT-SBR difference in longitudinal examinations (V2 – VI). The left column shows bilateral values. The centre and right columns respectively show values in the clinically most and least affected brain hemispheres DaT = dopamine transporter; HC = healthy control; iRBD = idiopathic REM sleep behaviour disorder; LI = limbic; PD = Parkinson's disease; Pu = putamen; SBR = striatal binding ratio; VI/V2 = visit I/2

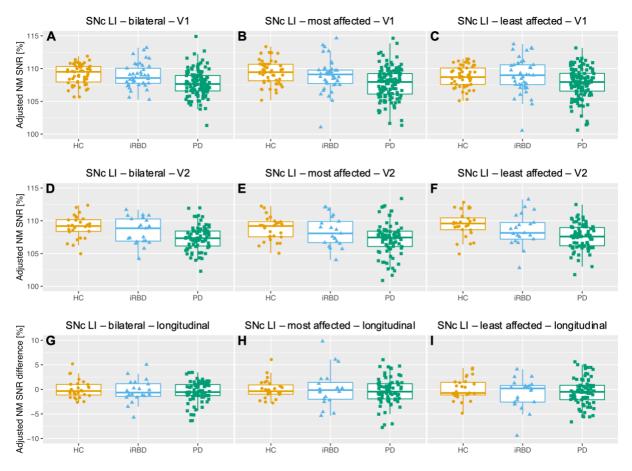


Supplementary Figure 6 Scatter boxplots of age and sex adjusted DaT-SBR measurements in the associative putamen. (A-C) DaT-SBR at V1. (D-F) DaT-SBR at V2. (G-I) DaT-SBR difference in longitudinal examinations (V2 – V1). The left column shows bilateral values. The centre and right columns respectively show values in the clinically most and least affected brain hemispheres AS = associative; DaT = dopamine transporter; HC = healthy control; iRBD = idiopathic REM sleep behaviour disorder; PD = Parkinson's disease; Pu = putamen; SBR = striatal binding ratio; V1/V2 = visit 1/2

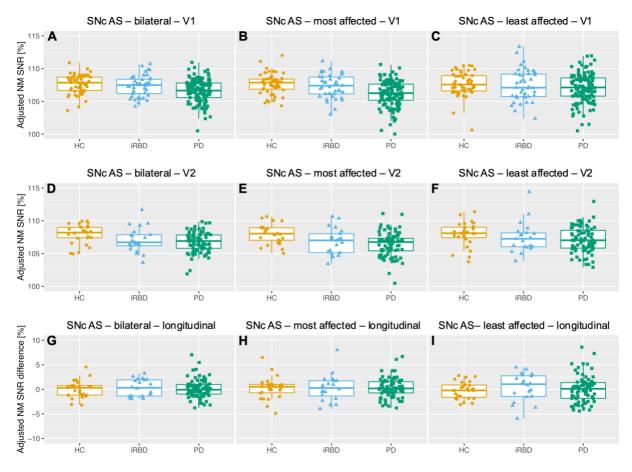


Supplementary Figure 7 Scatter boxplots of age and sex adjusted DaT-SBR measurements in the sensorimotor putamen. (A-C) DaT-SBR at VI. (D-F) DaT-SBR at V2. (G-I) DaT-SBR difference in longitudinal examinations (V2 – VI). The left column shows bilateral values. The centre and right columns respectively show values in the clinically most and least affected brain hemispheres

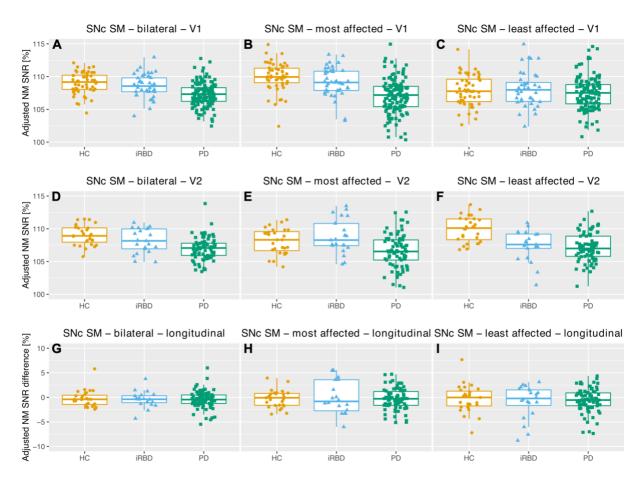
DaT = dopamine transporter; HC = healthy control; iRBD = idiopathic REM sleep behaviour disorder; PD = Parkinson's disease; Pu = putamen; SBR = striatal binding ratio; SM = sensorimotor; VI/V2 = visit 1/2



Supplementary Figure 8 Scatter boxplots of age and sex adjusted neuromelanin SNR measurements in the limbic SNc. (A-C) NM SNR at VI. (D-F) NM SNR at V2. (G-I) NM SNR difference in longitudinal examinations (V2 - VI). The left column shows bilateral values. The centre and right columns respectively show values in the clinically most and least affected brain hemispheres HC = healthy control; IRBD = idiopathic REM sleep behaviour disorder; LI = limbic; NM = neuromelanin; PD = Parkinson's disease; SNc = substantia nigra pars compacta; SNR = signal-to-noise ratio; VI/V2 = visit I/2

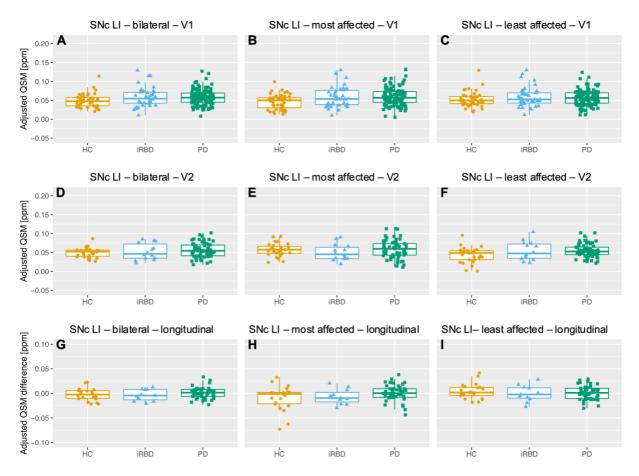


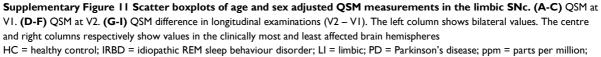
Supplementary Figure 9 Scatter boxplots of age and sex adjusted neuromelanin SNR measurements in the associative SNc. (A-C) NM SNR at VI. (D-F) NM SNR at V2. (G-I) NM SNR difference in longitudinal examinations (V2 - VI). The left column shows bilateral values. The centre and right columns respectively show values in the clinically most and least affected brain hemispheres AS = associative; HC = healthy control; IRBD = idiopathic REM sleep behaviour disorder; NM = neuromelanin; PD = Parkinson's disease; SNc = substantia nigra pars compacta; SNR = signal-to-noise ratio; VI/V2 = visit I/2



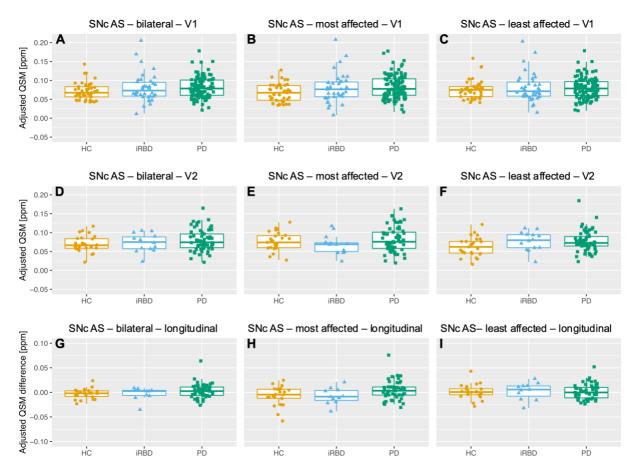
Supplementary Figure 10 Scatter boxplots of age and sex adjusted neuromelanin SNR measurements in the sensorimotor SNc. (A-C) NM SNR at VI. (D-F) NM SNR at V2. (G-I) NM SNR difference in longitudinal examinations (V2 – VI). The left column shows bilateral values. The centre and right columns respectively show values in the clinically most and least affected brain hemispheres

HC = healthy control; IRBD = idiopathic REM sleep behaviour disorder; NM = neuromelanin; PD = Parkinson's disease; SM = sensorimotor; SNc = substantia nigra pars compacta; SNR = signal-to-noise ratio; VI/V2 = visit $\frac{1}{2}$

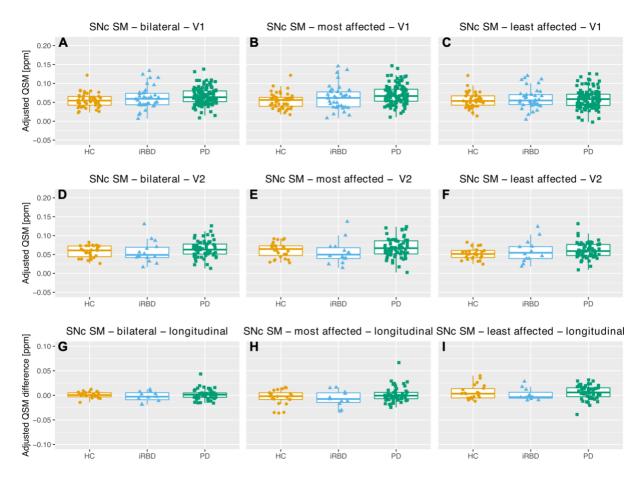




QSM = quantitative susceptibility mapping; SNc = substantia nigra pars compacta; V1/V2 = visit 1/2



Supplementary Figure 12 Scatter boxplots of age and sex adjusted QSM measurements in the associative SNc. (A-C) QSM at V1. (D-F) QSM at V2. (G-I) QSM difference in longitudinal examinations (V2 – V1). The left column shows bilateral values. The centre and right columns respectively show values in the clinically most and least affected brain hemispheres AS = associative; HC = healthy control; IRBD = idiopathic REM sleep behaviour disorder; PD = Parkinson's disease; ppm = parts per million; QSM = quantitative susceptibility mapping; SNc = substantia nigra pars compacta; V1/V2 = visit 1/2



Supplementary Figure 13 Scatter boxplots of age and sex adjusted QSM measurements in the sensorimotor SNc. (A-C) QSM at VI. (D-F) QSM at V2. (G-I) QSM difference in longitudinal examinations (V2 – VI). The left column shows bilateral values. The centre and right columns respectively show values in the clinically most and least affected brain hemispheres HC = healthy control; IRBD = idiopathic REM sleep behaviour disorder; PD = Parkinson's disease; ppm = parts per million; SM = sensorimotor; QSM = quantitative susceptibility mapping; SNc = substantia nigra pars compacta; VI/V2 = visit I/2

References

1. Robinson SD, Bredies K, Khabipova D, Dymerska B, Marques JP, Schweser F. An illustrated comparison of processing methods for MR phase imaging and QSM: combining array coil signals and phase unwrapping. *NMR Biomed*. Apr 2017;30(4)doi:10.1002/nbm.3601

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