

Supplementary Materials

Metabolic Impact of Anticancer Drugs Pd₂Spermine and Cisplatin on the Brain of Healthy Mice

Tatiana J. Carneiro ¹, Martin Vojtek ², Salomé Gonçalves-Monteiro ², João R. Neves ¹, Ana L.M. Batista de Carvalho ³, Maria Paula M. Marques ^{3,4}, Carmen Diniz ² and Ana M. Gil ^{1,*}

¹ Department of Chemistry and CICECO-Aveiro Institute of Materials, University of Aveiro, 3810-193 Aveiro, Portugal; tatiana.joao@ua.pt (T.J.C.), joaodrneves@ua.pt (J.R.N.)

² LAQV/REQUIMTE, Laboratory of Pharmacology, Department of Drug Sciences, Faculty of Pharmacy, University of Porto, 4150-755 Porto, Portugal; matovoj@gmail.com (M.V.), salomemonteiro8180@gmail.com (S.G.-M.), cdiniz@ff.up.pt (C.D.)

³ Molecular Physical-Chemistry R&D Unit, Department of Chemistry, University of Coimbra, 3004-535 Coimbra, Portugal; almbc@uc.pt (A.L.M.B.d.C.), pmc@ci.uc.pt (M.P.M.M.)

⁴ Department of Life Sciences, Faculty of Science and Technology, University of Coimbra, 3000-456 Coimbra, Portugal

* Correspondence: agil@ua.pt; Tel.: +351-234370707

Content of Supplementary Materials:

Figure S1. PCA and PLS-DA scores scatter plots for the ¹H NMR spectra of polar extracts from brain of healthy BALB/c mice after exposure to cDDP at a) 1 h, b) 12 h, and c) 48 h post-injections times, compared to controls.

Figure S2. PCA and PLS-DA scores scatter plots for the ¹H NMR spectra of polar extracts from brain of healthy BALB/c mice after exposure to Pd₂Spm at a) 1 h, b) 12 h, and c) 48 h post-injections times, compared to controls.

Figure S3. Examples of ¹H NMR spectral regions illustrating differences in a) polar and b) nonpolar metabolites between cDDP-exposed (blue trace) and Pd₂Spm-exposed (red trace), after 48 h of exposure.

Figure S4. PCA and PLS-DA scores scatter plots for the ¹H NMR spectra of nonpolar extracts from brain of healthy BALB/c mice after exposure to cDDP at a) 1 h, b) 12 h, and c) 48 h post-injections times, compared to controls.

Figure S5. PCA and PLS-DA scores scatter plots for the ¹H NMR spectra of nonpolar extracts from brain of healthy BALB/c mice after exposure to Pd₂Spm at a) 1 h, b) 12 h, and c) 48 h post-injections times, compared to controls.

Table S1. List of metabolites and corresponding spin systems identified in the 500MHz ¹H NMR spectra of polar and nonpolar extracts of brain from healthy BALB/c mice, at 1 h post-injection time with phosphate buffer saline.

Table S2. Significant metabolite variations (expressed in effect size, ES) in the polar (top) and nonpolar (bottom) metabolomes of mice brain exposed to Pd₂Spm, compared to cDDP, at 1, 12 and 48 h post-injection times.

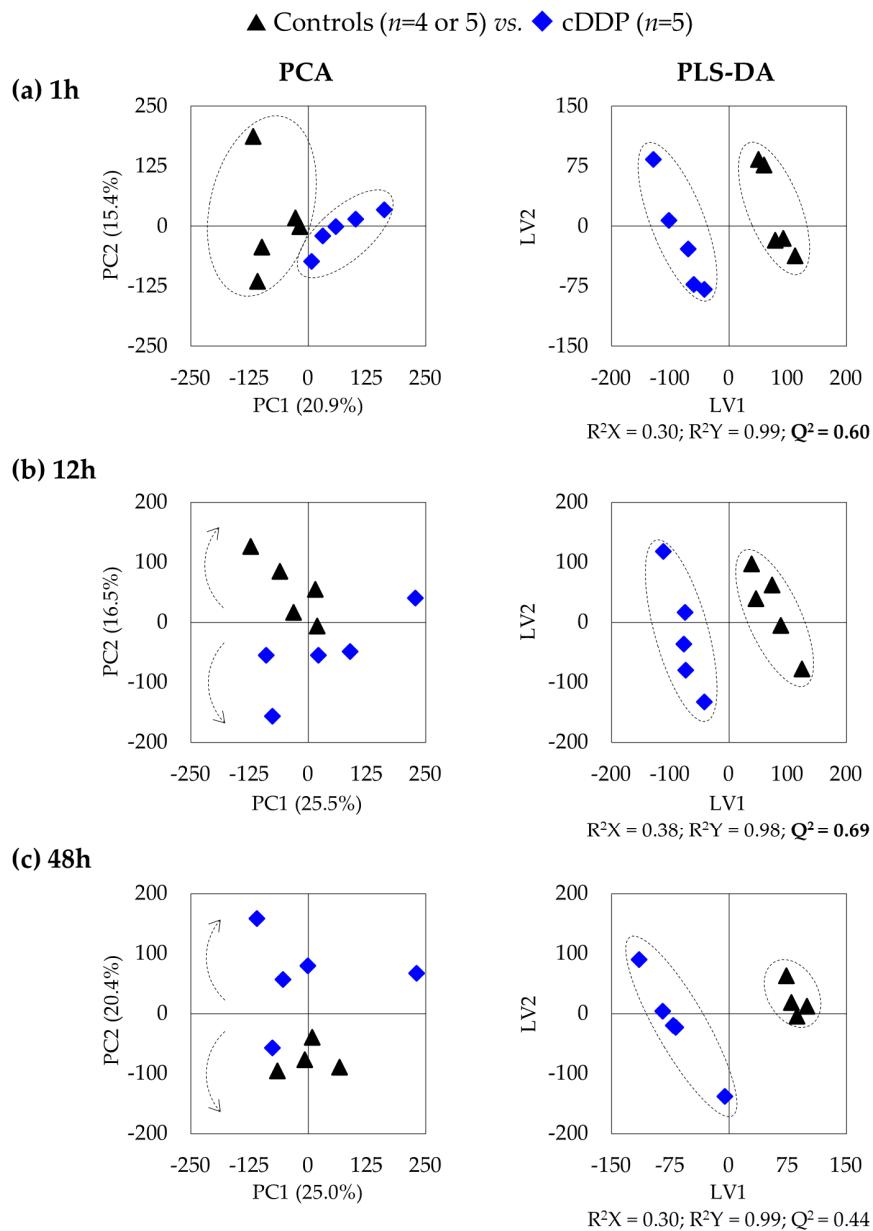
Figure S1

Figure S1. PCA and PLS-DA scores scatter plots for the ^1H NMR spectra of polar extracts from brain of healthy BALB/c mice after exposure to cDDP at a) 1 h, b) 12 h, and c) 48 h post-injections times, compared to controls. Validation parameters (R^2 and Q^2) are indicated for each PLS-DA model; Q^2 values > 0.5 are highlighted in bold.

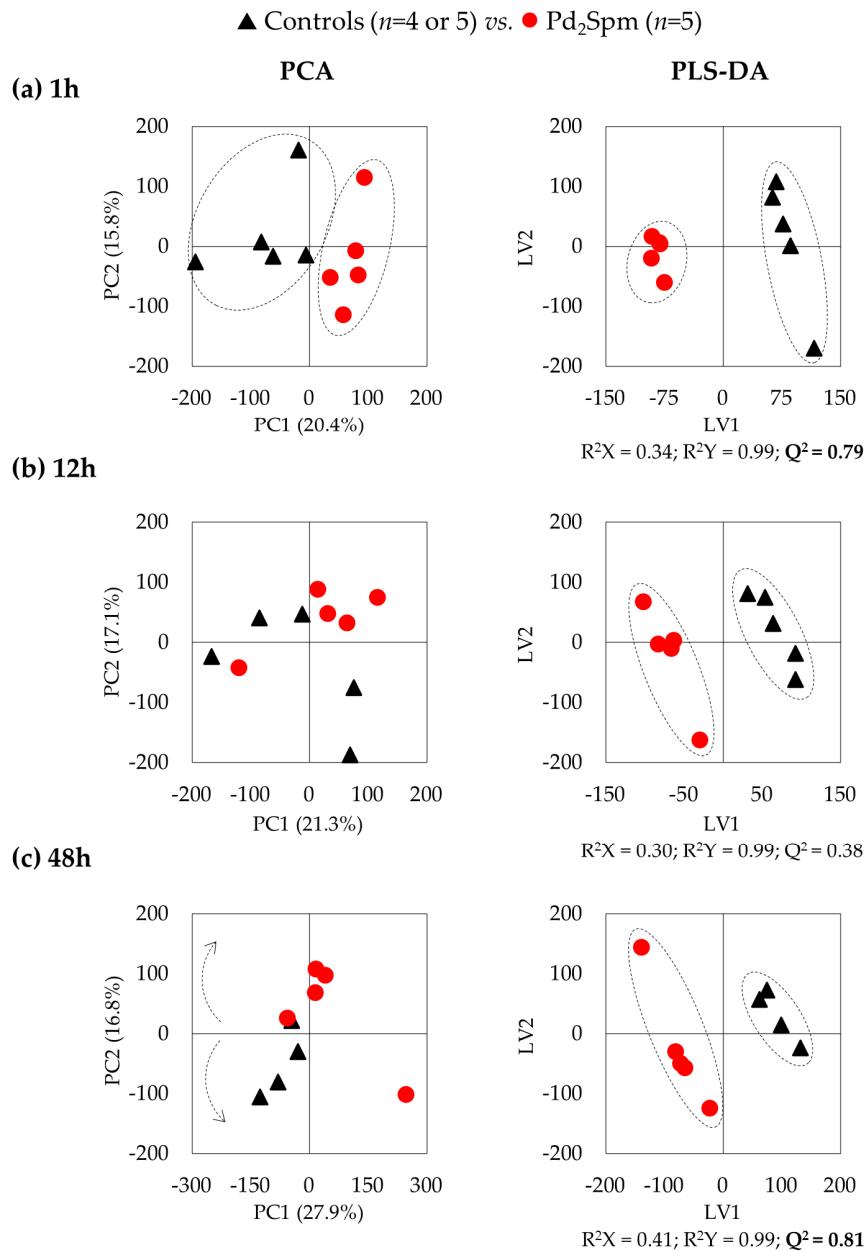
Figure S2

Figure S2. PCA and PLS-DA scores scatter plots for the ^1H NMR spectra of polar extracts from brain of healthy BALB/c mice after exposure to Pd_2Spm at a) 1 h, b) 12 h, and c) 48 h post-injections times, compared to controls. In PLS-DA models, Q^2 values > 0.5 (in bold) represent robust models.

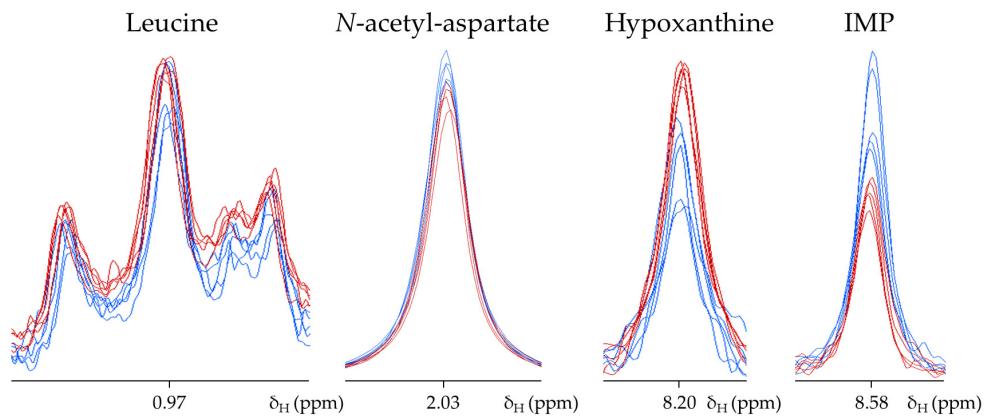
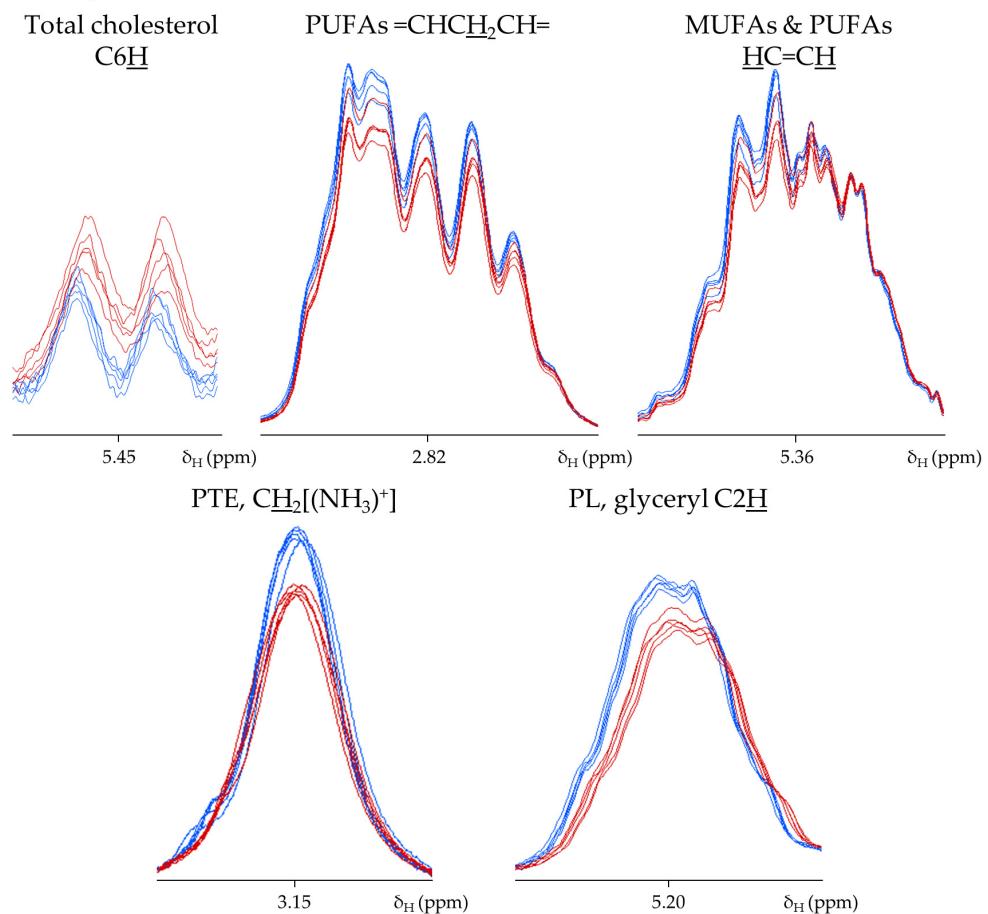
Figure S3**a) Polar metabolites****b) Nonpolar metabolites**

Figure S3. Examples of ¹H NMR spectral regions illustrating differences in a) polar and b) nonpolar metabolites between cDDP-exposed (blue trace) and Pd₂Spm-exposed (red trace), after 48 h of exposure.

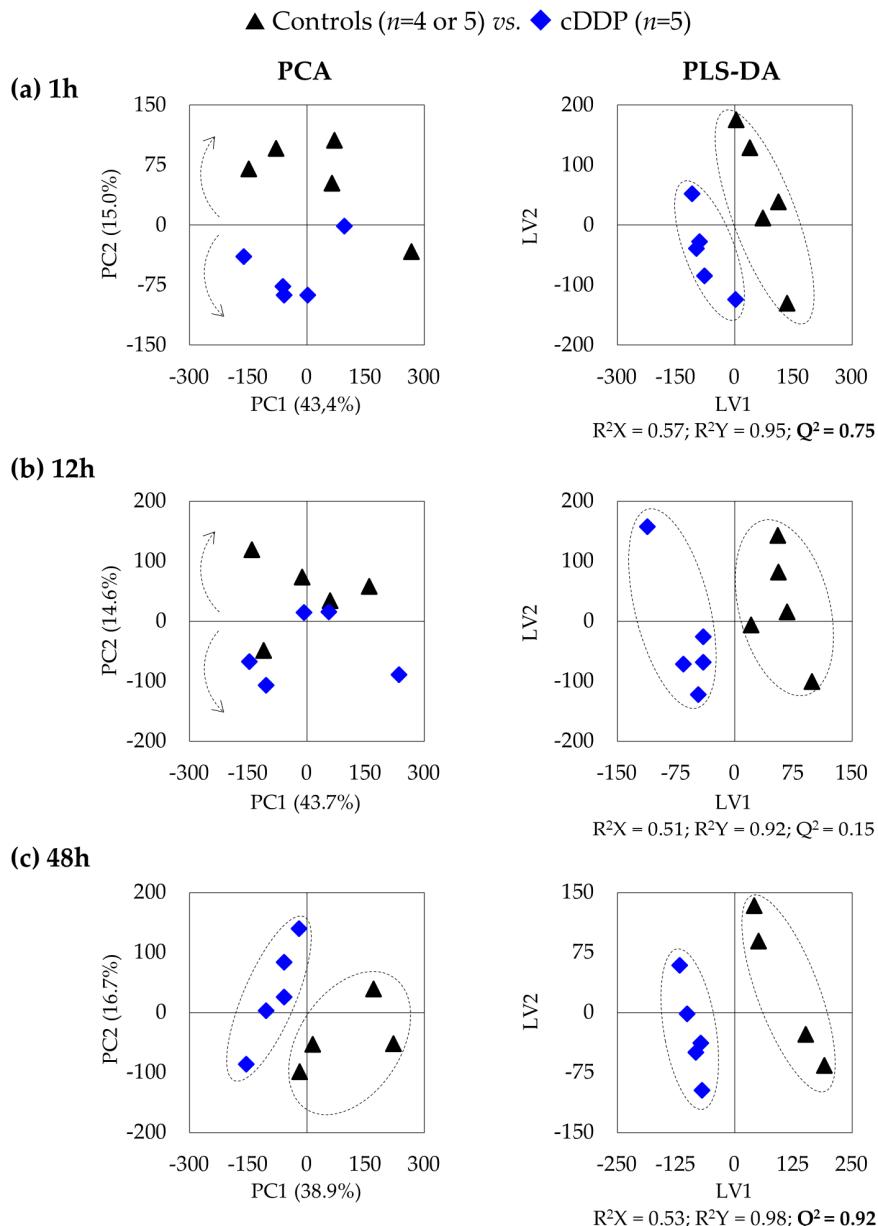
Figure S4

Figure S4. PCA and PLS-DA scores scatter plots for the ^1H NMR spectra of nonpolar extracts from brain of healthy BALB/c mice after exposure to cDDP at a) 1 h, b) 12 h, and c) 48 h post-injections times, compared to controls. In PLS-DA models, Q^2 values > 0.5 (in bold) represent robust models.

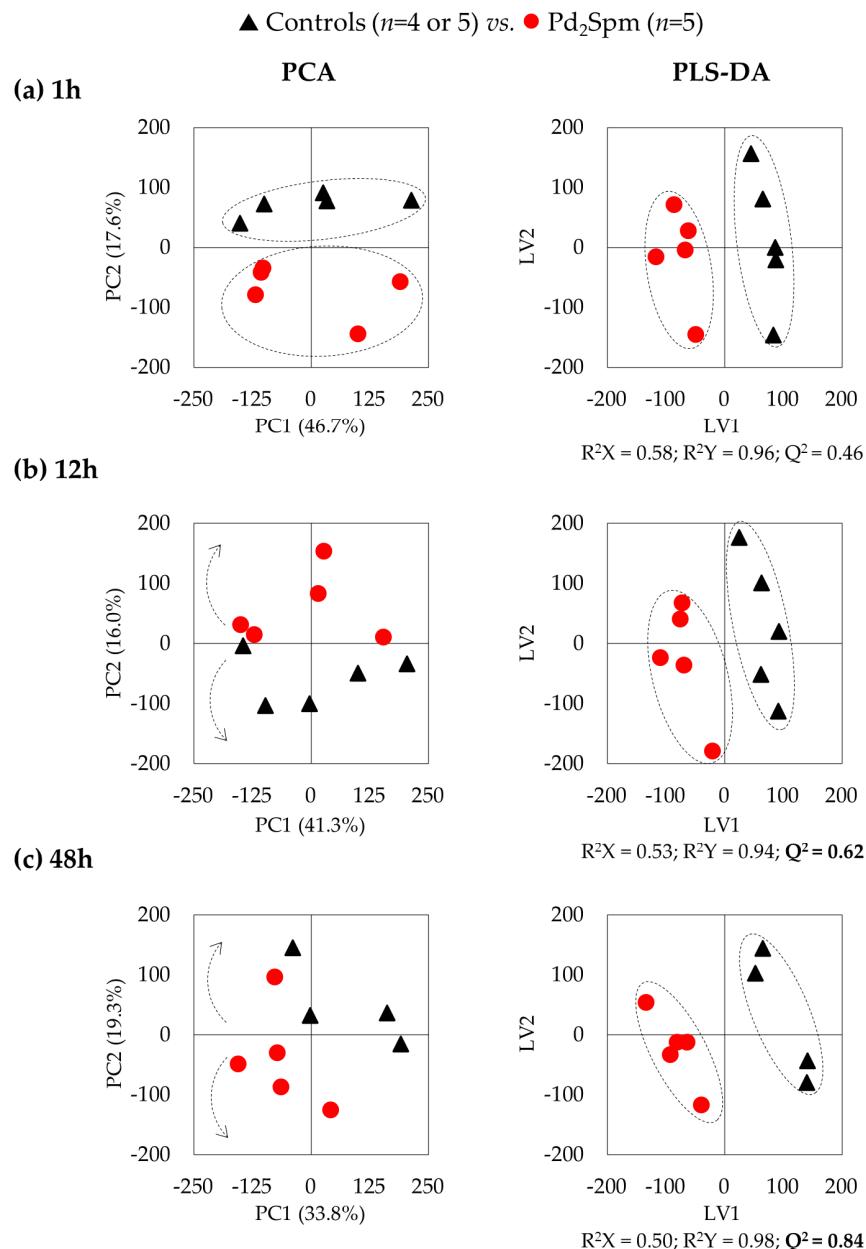
Figure S5

Figure S5. PCA and PLS-DA scores scatter plots for the ¹H NMR spectra of nonpolar extracts from brain of healthy BALB/c mice after exposure to Pd₂Spm at a) 1 h, b) 12 h, and c) 48 h post-injections times, compared to controls. In PLS-DA models, Q^2 values > 0.5 (in bold) represent robust models.

Table S1. List of metabolites and corresponding spin systems identified in the 500MHz ¹H NMR spectra of polar and nonpolar extracts of brain from healthy BALB/c mice, at 1 h post-injection time with phosphate buffer saline. ^a Predominant metabolites; ^b Metabolites identified, for the first time to our knowledge, by ¹H NMR in brain extracts of roedent models; * Possible contamination due to the extraction procedure; [†] Tentative assignment. Human Metabolome Database (HMDB ID) metabolite identification codes are indicated whenever possible. Abbreviations: 3-HBA, 3-hydroxybutyrate; ADP, adenosine diphosphate; AMP, adenosine monophosphate; ATP, adenosine triphosphate; DMA, dimethylamine; DMSO₂, dimethyl sulfone; GABA, γ-aminobutyrate; GPC, glycerophosphocholine; GSH, glutathione (reduced); HX, hypoxanthine; IMP, inosine monophosphate; MUFA, monounsaturated fatty acids; NAD⁺, nicotinamide adenine dinucleotide; PC, phosphocholine; PE, phosphoethanolamine; PL, phospholipid; PTC, phosphatidylcholine; PTE, phosphatidylethanolamine; PUFA, polyunsaturated fatty acids; SM, sphingomyelin; TMAO, Trimethylamine N-oxide; s, singlet; d, doublet; dd, double doublet; t, triplet; q, quartet; m, multiplet; br, broad signal.

Metabolite	δ_{H} ppm (multiplicity, assignment)	HMDB ID [61]
Polar extracts		
Amino acids and derivatives		
Alanine	1.48 (d, βCH ₃); 3.78 (q, αCH)	HMDB0000161
Asparagine [†]	2.83 (m, βCH ₂)	HMDB0000168
Aspartate	2.68 (dd, βCH); 2.81 (dd, β'CH); 3.90 (dd, αCH)	HMDB0000191
Creatine ^a	3.04 (s, N-CH ₃); 3.93 (s, N-CH ₂)	HMDB0000064
GABA ^a	1.90 (m, βCH ₂); 2.30 (t, γCH ₂); 3.01 (t, αCH ₂)	HMDB0000112
Glutamate ^a	2.04 (m, βCH); 2.11 (m, β'CH); 2.36 (m, γCH ₂); 3.76 (dd, αCH)	HMDB0000148
Glutamine	2.14 (m, βCH ₂); 2.45 (m, γCH ₂); 3.77 (t, αCH)	HMDB0000641
Glycine	3.55 (s, αCH ₂)	HMDB0000123
Histidine	7.06 (s, C4H ring); 8.00 (s, C2H ring)	HMDB0000177
Isoleucine	0.94 (t, δCH ₃); 1.01 (d, β'CH ₃); 1.99 (m, βCH)	HMDB0000177
Leucine	0.96 (t, δCH ₃ /δ'CH ₃); 1.71 (m, βCH ₂ /γCH)	HMDB0000687
Lysine	1.45 (m, γCH ₂); 1.73 (m, δCH ₂); 1.92 (m, βCH ₂)	HMDB0000182
N-acetylaspartate ^a	2.02 (s, CH); 2.50, 2.70 (m, CH ₂); 4.40 (m, CH ₃)	HMDB0000812
Phenylalanine	7.33 (d, C2H/C6H ring); 7.38 (m, C4H ring); 7.42 (t, C3H/C5H ring)	HMDB0000159
Taurine ^a	3.27 (t, S-CH ₂); 3.43 (t, N-CH ₂)	HMDB0000251
Tyrosine	6.90 (d, C3H/C5H ring); 7.20 (d, C2H/C6H ring)	HMDB0000158
Valine	0.99 (d, γCH ₃); 1.05 (d, γ'CH ₃); 2.27 (m, βCH); 3.61 (d, αCH)	HMDB0000883
Choline derivatives		
Choline ^a	3.21 (s, N(CH ₃) ₃)	HMDB0000097
GPC	3.23 (s, N(CH ₃) ₃); 3.92 (m, αCH ₂); 4.33 (m, PO ₃ -αCH ₂)	HMDB0000086
PC ^a	3.22 (s, N(CH ₃) ₃); 4.17 (m, PO ₃ -CH ₂)	HMDB0001565
Nucleotides and derivatives		
Adenine ^{b, †}	8.23 (s, C2H ring)	HMDB0000034
Adenosine	4.29 (q, C4'H ribose); 4.44 (dd, C3'H ribose); 6.10 (d, C1'H ribose); 8.12 (s, C8H ring); 8.27 (s, C2H ring)	HMDB0000050
ADP	6.15 (d, C1'H ribose); 8.27 (s, C2H ring); 8.54 (s, C8H ring)	HMDB0001341
AMP	4.51 (dd, C2'H ribose); 6.14 (d, C1'H ribose); 8.27 (s, C2H ring); 8.61 (s, C8H ring)	HMDB0000045
ATP	6.15 (d, C1'H ribose); 8.28 (s, C2H ring); 8.545 (s, C8H ring)	HMDB0000538
Organic		
Hypoxanthine ^b	8.18 (s, C2H); 8.20 (s, C8H)	HMDB0000157
IMP	8.24 (s, C8H ring); 8.58 (s, C8H ring)	HMDB0000175
Inosine	6.10 (d, C1'H); 8.24 (s, C8H ring); 8.35 (s, C2H ring)	HMDB0000195
NAD ⁺ [†]	8.43 (s, A8)	HMDB0000902
Uridine	5.90 (d, C5H ring); 5.92 (d, C1'H ring); 7.88 (d, C6H ring)	HMDB0000285
3-HBA	1.20 (d, CH ₃); 2.31 (m, CH ₂)	HMDB0000357
Acetate	1.92 (s, CH ₃)	HMDB0000042

	Ascorbate	4.52 (d, C4H ring)	HMDB0000044
	Formate	8.46 (s, CH)	HMDB0000142
	Fumarate	6.52 (s, CH)	HMDB0000134
	Lactate ^a	1.33 (d, CH ₃); 4.10 (q, CH)	HMDB0000190
	Succinate	2.41 (s, CH ₂)	HMDB0000254
Other compounds	Acetone ^b	2.24 (s, CH ₃)	HMDB0001659
	DMA ^b	2.73 (s, (CH ₃) ₂)	HMDB0000087
	DMSO ₂ ^{b, †}	3.15 (s, CH ₃)	HMDB0004983
	Ethanol *	1.19 (t, CH ₃); 3.65 (q, CH ₂)	HMDB0000108
	<i>m</i> -Inositol ^a	3.28 (t, C5H); 3.54 (dd, C1H/ C3H) 3.62 (t, C4H/ C6H); 4.06 (t, C2H)	HMDB0000211
	Niacinamide	7.60 (m, C5H); 8.72 (m, C4H, C6H); 8.94 (m, C2H)	HMDB0001406
	PE ^{b, †}	3.99 (m, PO ₃ -CH ₂)	HMDB0000224
	Propylene glycol *	1.15 (d, CH ₃); 3.87 (m, CH)	HMDB0001881
	TMAO ^b	3.27 (s, CH ₃)	HMDB0000925
Nonpolar extracts			
Cholesterol	Free ^a	0.68 (s, C18H ₃); 0.86 (d, C26H ₃); 0.87 (d, C27H ₃); 0.91 (s, C27H ₃); 0.95 (m, C9H); 0.99 (m, C14H); 1.01 (s, C19H ₃); 1.07 (m, C1H); 1.10 (m, C17H); 1.12 (m, C23H/ C24H); 1.13 (m, C22H); 1.49 (m, C11H); 1.51 (m, C2H); 1.84 (m, C4H); 1.87 (t, C16H); 1.95 (t, C7H); 1.99 (t, C7H'); 2.26 (t, C4H'); 3.53 (m, C3H); 5.45 (d, C6H)	HMDB0000067
	Esterified	0.68 (s, C18H ₃); 0.86 (d, C26H ₃); 0.87 (d, C27H ₃); 0.91 (s, C27H ₃); 0.95 (m, C9H); 0.99 (m, C14H); 1.01 (s, C19H ₃); 1.10 (m, C17H); 1.12 (m, C23H/ C24H); 1.13 (m, C22H); 1.15 (m, C1H); 1.84 (t, C4H); 1.87 (t, C15H'); 1.95 (t, C7H); 1.99 (t, C7H'); 2.31 (t, C4H'); 5.45 (d, C6H)	Undefined
Fatty acids (FA)	Saturated	0.89 (br, CH ₃); 1.25 (br, (CH ₂) _n); 2.30 (m, CH ₂ CO)	Undefined
	Monounsaturated	2.02 (ω 7/ ω 9 CH ₂ CH ₂ CH=); 2.30 (m, ω 7/ ω 9 CH ₂ CO); 5.34 (m, HC=CH)	16:1 (Δ^9 ; ω 7), HMDB0031053 / 18:1 (Δ^9 ; ω 9), HMDB0000207
	Polyunsaturated ^a	0.93 (t, ω 6 CH ₃); 0.98 (t, ω 3 CH ₃); 1.30 (m, (CH ₂) _n); 1.33 (m, ω 3/ ω 6 =CHCH ₂ (CH ₂) _n); 2.05 (m, CH ₂ CH=); 2.30 (m, CH ₂ CO); 2.77 (t, 18:2 =CHCH ₂ CH=); 2.81 (m, ω 3/ ω 6 =CHCH ₂ CH=); 2.82 (m, 22:6 =CHCH ₂ CH=); 2.84 (m, 22:6 CH ₂ CO); 5.37 (m, HC=CH)	18:2 ($\Delta^{9,12}$; ω 6), HMDB0000673 22:6 ($\Delta^{4,7,10,13,16,19}$; ω 3), HMDB0002183 and other undefined ω 3/ ω 6 FAs
	PL resonances	3.75 (br, CH ₂ N(CH ₃) ₃); 3.94 (br, Glyceryl C3H ₂); 4.38 (br, POCH ₂); 4.40 (br, Glyceryl C1H ₂); 5.20 (br, Glyceryl C2H)	Undefined
Phospholipids	PTC ^a	3.31 (s, N(CH ₃) ₃); 4.12 (m, Glyceryl, C1H ₂)	Undefined
	PTE ^a	3.15 (br, CH ₂ [(NH ₃) ⁺])); 8.60 (br, [(NH ₃) ⁺])	Undefined
	SM	3.29 (s, N(CH ₃) ₃)	Undefined

Table S2. Significant metabolite variations (expressed in effect size, ES) in the polar (top) and nonpolar (bottom) metabolomes of mice brain exposed to Pd₂Spm, compared to cDDP, at 1, 12 and 48 h post-injection times. Only variations with |ES| > ES Error and *p*-value < 0.05 are shown. [†] Tentative assignment. [‡] Partial integration of resonance peak. ^a Metabolic variation statistically significant after False Discovery Rate (FDR) correction; ^b Metabolite observed to vary only in the direct comparison of the two drugs (and not in comparison with controls). Abbreviations as defined in Tables 1 and S1.

		$\delta_{\text{H}}/\text{ppm}$ (multiplicity)	Pd ₂ Spm vs. cDDP							
	Metabolite Family/ assignment		1h		12h		48h			
		ES	\pm Error	<i>p</i> -value	ES	\pm Error	<i>p</i> -value	ES	\pm Error	<i>p</i> -value
Amino acids and derivatives	Alanine	1.48 (d)	-3.2 \pm 1.9	2.4×10^{-3} ^a	—	—	—	1.5 \pm 1.4	4.9×10^{-2}	
	Leucine	0.96 (t)	-1.8 \pm 1.5	2.4×10^{-2}	—	—	—	3.1 \pm 1.8	5.1×10^{-3} ^a	
	N-acetylaspartate	2.03 (s)	—	—	—	—	—	-3.0 \pm 1.8	1.7×10^{-3} ^a	
	Taurine ^b	3.43 (t)	—	—	—	—	—	-2.4 \pm 1.6	1.3×10^{-2}	
	Valine	1.05 (d)	-3.0 \pm 1.8	1.7×10^{-3} ^a	—	—	—	—	—	
Choline derivatives	Glycerophosphocholine	3.23 (s)	2.5 \pm 1.7	1.4×10^{-2}	—	—	—	—	—	
	Phosphocholine ^b	3.21 (s)	—	—	—	—	—	-2.1 \pm 1.5	1.2×10^{-2}	
	Adenine ^{†, b}	8.23 (s)	—	—	—	—	—	-2.2 \pm 1.6	2.0×10^{-2}	
Nucleotides and derivatives	Adenosine	4.29 (q)	—	—	—	—	—	1.5 \pm 1.4	1.6×10^{-2}	
	ADP	8.54 (s)	1.9 \pm 1.5	1.8×10^{-2}	—	—	—	-1.6 \pm 1.4	3.9×10^{-2}	
	HX	8.20 (s)	—	—	—	—	—	2.9 \pm 1.8	6.2×10^{-3} ^a	
	IMP	8.58 (s)	—	—	—	—	—	-3.0 \pm 1.8	4.8×10^{-3} ^a	
	Inosine	8.35 (s)	-1.8 \pm 1.5	2.8×10^{-2}	—	—	—	—	—	
Organic acids	Formate	8.46 (s)	—	—	-1.8 \pm 1.5	3.4×10^{-2}	—	—	—	
Other compounds	Acetone	2.24 (s)	-5.7 \pm 2.8	1.7×10^{-3} ^a	—	—	—	—	—	
	DMA	2.73 (s)	4.4 \pm 2.3	2.3×10^{-3} ^a	—	—	—	2.2 \pm 1.6	1.7×10^{-2}	
	DMSO ₂ [†]	3.15 (s)	14.5 \pm 6.5	7.9×10^{-3}	3.1 \pm 1.8	7.9×10^{-3}	—	—	—	
Unassigned resonances	U1	0.80 (t)	-2.2 \pm 1.6	2.1×10^{-2}	—	—	—	—	—	
	U4 ^b	2.06 (st)	—	—	—	—	—	2.4 \pm 1.6	3.2×10^{-2}	
	U5 ^b	6.08 (s)	—	—	—	—	—	-1.8 \pm 1.5	4.5×10^{-2}	
Cholesterol	Free, C3H	3.53 (m)	—	—	—	—	—	3.1 \pm 1.8	7.9×10^{-3} ^a	
	Total, C6H	5.45 (d)	—	—	—	—	—	3.2 \pm 1.9	2.7×10^{-3} ^a	
	Total, C7H & C7H'	1.95-1.97	—	—	—	—	—	1.9 \pm 1.5	2.5×10^{-2}	
	Total, C26H ₃	0.86 (d)	—	—	—	—	—	1.8 \pm 1.5	2.8×10^{-2}	
	Total, C27H ₃	0.87 (d)	—	—	—	—	—	4.3 \pm 2.3	1.6×10^{-4} ^a	
Fatty acids	CH ₃	0.89 (br)	—	—	—	—	—	2.8 \pm 1.7	7.9×10^{-3} ^a	
	Saturated, (CH ₂) _n	1.25 (br)	2.8 \pm 1.7	7.9×10^{-3} ^a	—	—	—	2.3 \pm 1.6	7.2×10^{-3} ^a	
	MUFAs (Δ^9 ; $\omega 7/\omega 9$), CH ₂ CH ₂ CH=	2.02 (m)	—	—	—	—	—	1.9 \pm 1.5	2.1×10^{-2}	
	MUFAs, HC=CH	5.34 (m)	—	—	—	—	—	2.5 \pm 1.6	7.9×10^{-3} ^a	
	PUFAs ($\omega 3$), CH ₃	0.98 (t)	—	—	—	—	—	-1.6 \pm 1.4	3.6×10^{-2}	
	PUFAs, (CH ₂) _n	1.30 (m)	—	—	—	—	—	2.3 \pm 1.6	1.1×10^{-2} ^a	
	PUFAs ($\omega 3/\omega 6$), =CHCH ₂ (CH ₂) _n	1.33 (m)	—	—	—	—	—	3.4 \pm 1.9	9.9×10^{-4} ^a	
	PUFAs, CH ₂ CH=	2.05 (m)	—	—	—	—	—	-1.9 \pm 1.5	1.6×10^{-2} ^a	
	PUFAs ($\omega 3/\omega 6$), =CHCH ₂ CH=	2.81 (m)	—	—	—	—	—	-2.4 \pm 1.6	6.5×10^{-3} ^a	
	PUFAs, HC=CH	5.37 (m)	—	—	—	—	—	-2.7 \pm 1.7	3.0×10^{-3} ^a	
	18:2 ($\Delta^{9,12}$; $\omega 6$), =CHCH ₂ CH=	2.77 (t)	-2.5 \pm 1.7	4.29×10^{-3} ^a	-1.3 \pm 1.4	1.6×10^{-2} ^a	—	-2.0 \pm 1.5	1.5×10^{-2} ^a	
	22:6 ($\Delta^{4,7,10,13,16,19}$; $\omega 3$), CH ₂ CO	2.38 (m)	—	—	—	—	—	-3.8 \pm 2.1	3.3×10^{-4} ^a	
	22:6 ($\Delta^{4,7,10,13,16,19}$; $\omega 3$), =CHCH ₂ CH=	2.82 (m)	—	—	—	—	—	-3.1 \pm 1.8	1.4×10^{-3} ^a	
Phospholipids	CH ₂ N(CH ₃) ₃	3.75 (br)	2.3 \pm 1.6	8.9×10^{-3} ^a	1.7 \pm 1.4	3.4×10^{-2}	—	—	—	
	POCH ₂	4.38 (br)	—	—	—	—	—	-5.9 \pm 2.9	1.2×10^{-4} ^a	
	Glyceryl C3H ₂	3.94 (br)	—	—	—	—	—	-5.9 \pm 2.9	5.8×10^{-5} ^a	
	Glyceryl C1H ₂	4.40 (br)	—	—	—	—	—	-4.1 \pm 2.2	2.0×10^{-4} ^a	
	Glyceryl C2H	5.20 (br)	—	—	—	—	—	-4.9 \pm 2.5	8.9×10^{-5} ^a	
	PTC & SM, N(CH ₃) ₃	3.29 - 3.31	—	—	—	—	—	-5.6 \pm 2.7	5.7×10^{-5} ^a	
	PTC, Glyceryl C1H ₂	4.12 (m)	—	—	—	—	—	-3.4 \pm 2.0	7.0×10^{-4} ^a	

	PTE CH ₂ [NH ₃] ⁺	3.15 (br)	—	—	—	—	—	-8.8 ± 4.1	9.9 × 10 ⁻⁷ ^a
	PTE [(NH ₃) ⁺]	8.60 (br)	—	—	—	—	—	-2.3 ± 1.6	7.4 × 10 ⁻³ ^a
	SM	5.70 (m)	—	—	—	—	—	2.8 ± 1.7	4.0 × 10 ⁻³ ^a
Unassigned resonances	U2	0.60 (d)	—	—	—	—	—	-2.3 ± 1.6	1.1 × 10 ⁻² ^a
	U5	2.61 (s)	5.2 ± 2.6	3.9 × 10 ⁻⁵ ^a	-2.2 ± 1.6	9.4 × 10 ⁻³ ^a	—	-1.6 ± 1.4	4.0 × 10 ⁻²
	U6	2.99 (s)	4.4 ± 2.3	1.3 × 10 ⁻³ ^a	—	—	—	-2.1 ± 1.5	1.6 × 10 ⁻² ^a
	U9	3.84 (d)	-1.8 ± 1.5	2.8 × 10 ⁻²	—	—	—	—	—
	U10	3.90 (br)	—	—	—	—	—	5.5 ± 2.7	3.4 × 10 ⁻⁵ ^a
	U11	5.29 (t)	—	—	—	-2.0 ± 1.5	1.6 × 10 ⁻² ^a	—	—
	U12	8.34 (br)	—	—	—	—	—	4.0 ± 2.2	2.2 × 10 ⁻⁴ ^a
	U13	0.89 (s)	—	—	—	—	—	1.8 ± 1.5	2.6 × 10 ⁻²
	U14	2.20 (q)	—	—	—	—	—	2.5 ± 1.7	4.0 × 10 ⁻³ ^a
	U15	4.22 (br)	—	—	—	—	—	3.5 ± 2.0	1.4 × 10 ⁻³ ^a
	U16	7.77 (br)	—	—	—	—	—	2.7 ± 1.7	4.4 × 10 ⁻³ ^a