

Novel insights into mice multi-organ metabolism upon exposure to a potential anticancer Pd(II)-agent

Tatiana J. Carneiro¹, Rita Araújo¹, Martin Vojtek², Salomé Gonçalves-Monteiro², Carmen Diniz², Ana L.M. Batista de Carvalho³, M. Paula M. Marques^{3,4}, Ana M. Gil^{1,*}

¹ Department of Chemistry and CICECO – Aveiro Institute of Materials, University of Aveiro, 3810-193, Aveiro, Portugal;

² LAQV/REQUIMTE, Laboratory of Pharmacology, Department of Drug Sciences, Faculty of Pharmacy, University of Porto, 4050-313 Porto, Portugal;

³ “Química-Física Molecular”, University of Coimbra, 3004-535 Coimbra, Portugal;

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

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

Figure S1

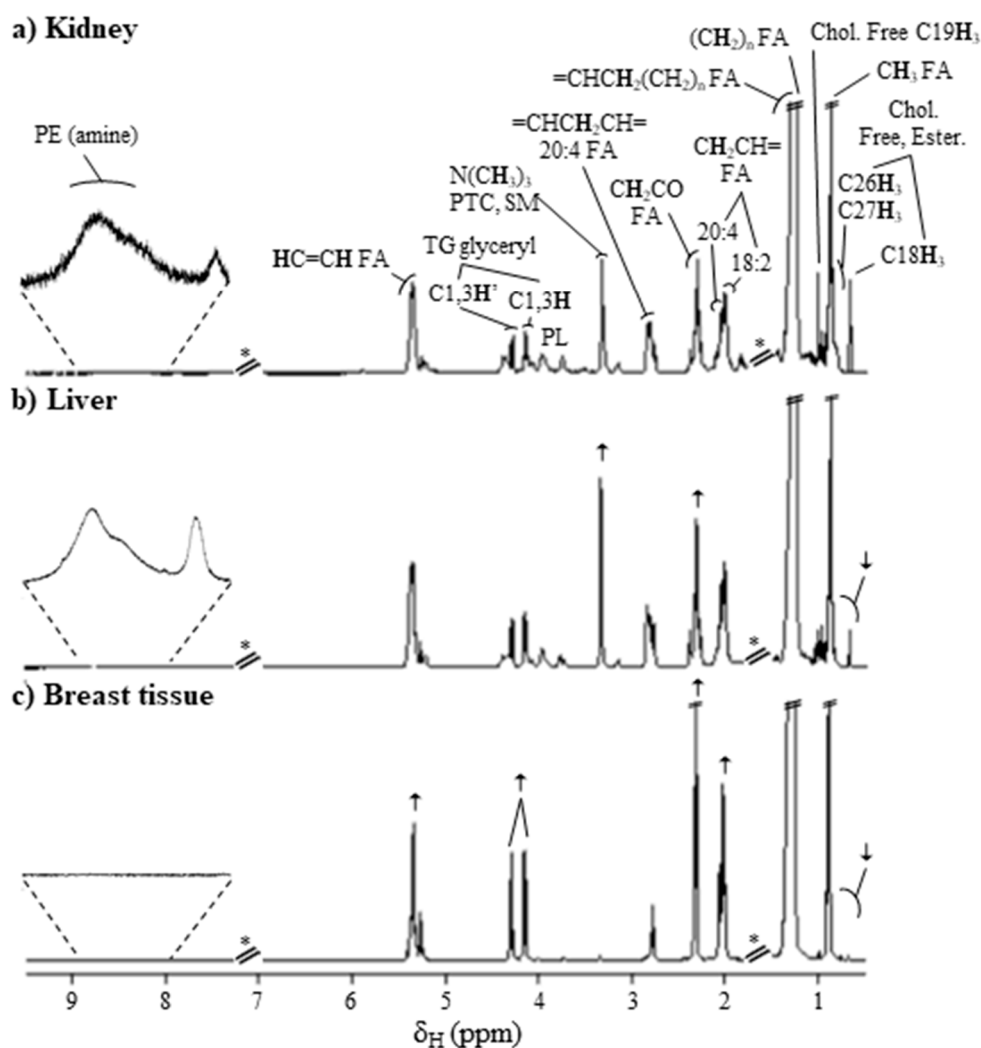


Figure S1. Average 500 MHz ^1H NMR spectra of lipophilic extracts of kidney, liver and breast tissue. Spectra obtained for (a) kidney, (b) liver and (c) breast tissue of BALB/c mice after 1h of post-injection with phosphate buffer saline (controls). * residual CDCl_3 (δ 7.15-7.37, and corresponding satellites at δ 7.04-7.06 and δ 7.45- δ 7.47) and water (δ 1.45-1.70) regions, removed from matrix used for multivariate analysis. Arrows identify noticeable differences in some metabolites between tissues. Abbreviations: Chol., cholesterol; Ester., esterified; FA, fatty acid; PL, phospholipids glyceryl moieties; PTC, phosphatidylcholine; PE, phosphatidylethanolamine; SM, sphingomyelin; TG, triacylglyceride.

Figure S2

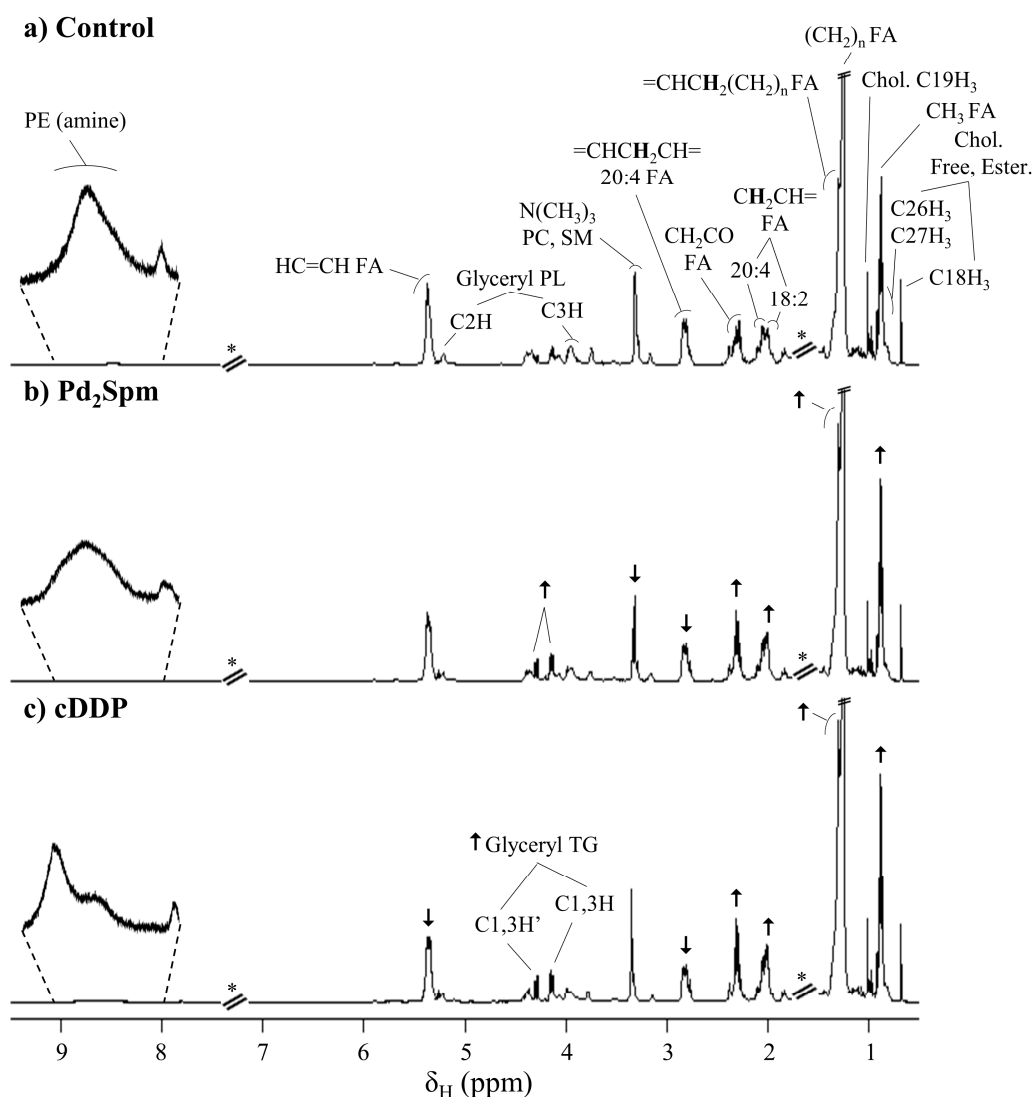


Figure S2. Average 500 MHz ^1H NMR spectra of lipophilic kidney extracts of mice exposed to cDDP and Pd₂Spm. Extracts correspond to BALB/c mice after 48 h of post-injection with (a) phosphate buffer saline (controls), (b) Pd₂Spm and (c) cDDP. * residual CDCl₃ (δ 7.15-7.37, and corresponding satellites at δ 7.04-7.06 and δ 7.45- δ 7.47) and water (δ 1.50-1.90) regions, removed from matrix used for multivariate analysis. Abbreviations as defined in Figure S1.

Figure S3

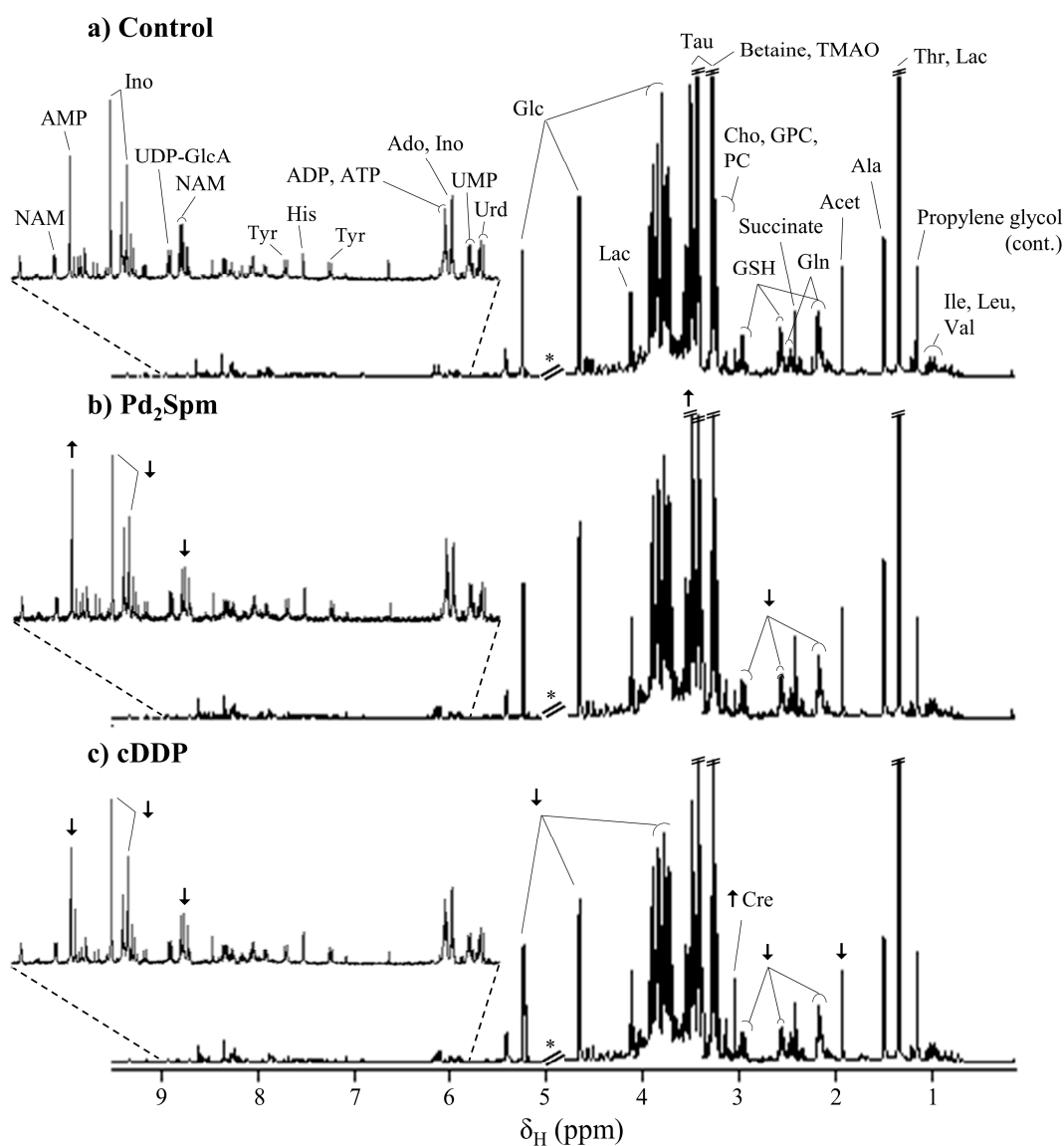


Figure S3. Average 500 MHz ^1H NMR spectra of aqueous liver extracts of mice exposed to Pd_2Spm and cDDP. Extracts were obtained for BALB/c mice after 12 h of post-injection with (a) phosphate buffer saline (controls), (b) Pd_2Spm and (c) cDDP. *Cut-off spectral region due to water suppression (δ_{H} 4.69-5.10). Abbreviations: 3-letter code used for amino acids; Acet, acetate; Ado, adenosine; ADP, adenosine diphosphate; AMP, adenosine monophosphate; ATP, adenosine triphosphate; Cho, choline; Cre, creatine; Glc, glucose; GPC, glycerophosphocholine; GSH, reduced glutathione; Ino, inosine; Lac, lactate; NAM, niacinamide; PC, phosphocholine; Tau, taurine; TMAO, trimethylamine-*N*-oxide; UDP-GlcA: uridine diphosphate-glucuronate; UMP, uridine monophosphate; Urd, uridine.

Supplementary tables S1, S2, S3 and S4

Table S1. List of metabolite variations in aqueous kidney extracts of mice exposed to cDDP and Pd₂Spm at 1, 12 and 48 h post-injection, compared to controls. Only variations with 95% significance level (p-value < 0.05) are shown; ES, effect size; ES error values in brackets correspond to > 75% of ES value; ^a variations remaining significant after False Discovery Rate correction; ^b metabolic variations specific of exposure to Pd₂Spm (these comprise both new variations and variations in the same metabolite but in opposite directions). 3-letter code used for amino acids; 3-HIBA, 3-hydroxyisobutyrate; ADP, adenosine diphosphate; AMP, adenosine monophosphate; Cho, choline; Cre, creatine; DMA, dimethylamine; DMSO₂, dimethyl sulfone (tentative assignment); Glc, glucose; PC, phosphocholine; Tau, taurine; TMA, trimethylamine; NAD⁺, nicotinamide adenine dinucleotide; UDP-GlcA, uridine-diphosphate-glucuronate; UMP, uridine monophosphate; Ui, unassigned i. [†]Tentative assignment. s: singlet, d: doublet, m: multiplet, t: triplet, dd: doublet of doublets.

Metabolite assignment	δ_H ppm multiplicity	cDDP vs. Controls (according to reference [15])						Pd ₂ Spm vs. Controls					
		1 h		12 h		48 h		1 h		12 h		48 h	
		ES ± Error	p-value	ES ± Error	p-value	ES ± Error	p-value	ES ± Error	p-value	ES ± Error	p-value	ES ± Error	p-value
3-HIBA	1.09 (d)	4.7 ± 2.5	1.9E-4 ^a	-2.2 ± 1.6	9.5E-3	- - -	-	2.7 ± 1.8	4.7E-3 ^a	- - -	-	- - -	-
ADP	8.28 (s)	1.8 ± (1.6)	2.9E-2	1.8 ± (1.5)	1.6E-2	- - -	-	2.3 ± 1.7	8.8E-3 ^a	- - -	-	- - -	-
Ala	1.48 (d)	-3.0 ± 1.9	9.5E-3 ^a	- - -	-	-2.2 ± 1.7	1.1E-2	-1.9 ± (1.6)	3.1E-2 ^a	- - -	-	- - -	-
Allantoin	5.39 (s)	- - -	-	-2.3 ± 1.6	9.3E-3	- - -	-	- - -	-	- - -	-	- - -	-
AMP	8.61 (s)	1.8 ± (1.5)	3.3E-2	1.9 ± (1.5)	2.8E-2	- - -	-	1.8 ± (1.6)	3.3E-2 ^a	- - -	-	- - -	-
Asn	2.96 (m)	- - -	-	-1.9 ± (1.5)	2.3E-2	- - -	-	- - -	-	- - -	-	- - -	-
Betaine	3.90 (s)	- - -	-	1.8 ± (1.5)	3.8E-2	- - -	-	- - -	-	- - -	-	- - -	-
Cho	3.21 (s)	-2.1 ± 1.6	1.6E-2	-3.2 ± 1.9	2.7E-3	- - -	-	-3.6 ± 2.1	8.2E-4 ^a	-2.7 ± 1.7	5.8E-3	- - -	-
Cre	3.93 (s)	- - -	-	-2.5 ± 1.6	1.1E-2	- - -	-	- - -	-	-2.3 ± 1.6	1.0E-2	-3.2 ± 2.0	1.1E-2
DMA	2.73 (s)	- - -	-	- - -	-	- - -	-	3.2 ± 2.0	5.5E-3 ^{a,b}	- - -	-	- - -	-
DMSO ₂ [†]	3.15 (s)	- - -	-	-1.9 ± (1.5)	3.2E-2	- - -	-	9.8 ± 4.7	2.1E-5 ^{a,b}	- - -	-	- - -	-
Fumarate	6.52 (s)	- - -	-	- - -	-	-3.5 ± 2.1	2.7E-3 ^a	2.9 ± 1.9	2.1E-2 ^{a,b}	2.0 ± 1.5	1.4E-2 ^b	2.2 ± 1.7	1.5E-2
Glc	5.23 (d)	- - -	-	1.9 ± (1.5)	2.7E-2	- - -	-	- - -	-	- - -	-	- - -	-
Hypoxanthine	8.21 (s)	- - -	-	- - -	-	-2.6 ± 1.8	3.3E-2	- - -	-	- - -	-	- - -	-
Ile	1.01 (d)	-3.1 ± 1.9	4.1E-3 ^a	-1.9 ± (1.5)	3.3E-2	-2.2 ± 1.7	1.5E-2	-2.2 ± 1.7	1.2E-2 ^a	- - -	-	- - -	-
Leu	0.96 (t)	-3.5 ± 2.1	1.2E-3 ^a	-1.8 ± (1.5)	3.0E-2	-1.8 ± (1.6)	2.5E-2	-3.5 ± 2.1	1.5E-3 ^a	- - -	-	- - -	-
<i>m</i> -Inositol	4.06 (t)	- - -	-	- - -	-	- - -	-	-2.2 ± 1.7	1.3E-2 ^a	-1.7 ± (1.4)	3.1E-2 ^b	- - -	-

Niacinamide	7.60 (dd)	- - - -	- - - -	-1.7 ± (1.5) 3.6E-2	- - - -	- - - -	- - - -	- - - -	- - - -
PC	3.22 (s)	- - - -	1.9 ± (1.5) 1.7E-2	- - - -	- - - -	- - - -	- - - -	- - - -	- - - -
Phe	7.33 (d)	-3.2 ± 2.0 1.6E-2	- - - -	- - - -	-3.3 ± 2.0 1.6E-2 ^a	-1.6 ± (1.4) 3.6E-2	- - - -	- - - -	- - - -
Succinate	2.41 (s)	- - - -	- - - -	- - - -	2.0 ± (1.6) 2.0E-2 ^{a,b}	-1.5 ± (1.4) 4.3E-2 ^b	-1.6 ± (1.5) 3.2E-2 ^b	- - - -	- - - -
Tau	3.42 (t)	3.0 ± 1.9 3.2E-3 ^a	- - - -	- - - -	- - - -	- - - -	- - - -	- - - -	- - - -
TMA	2.89 (s)	- - - -	- - - -	-4.3 ± 2.4 1.1E-3 ^a	- - - -	- - - -	- - - -	- - - -	- - - -
Tyr	6.90 (d)	-2.1 ± 1.6 1.6E-2	- - - -	-3.1 ± 1.9 2.5E-3 ^a	-2.5 ± 1.8 7.0E-3 ^a	- - - -	- - - -	- - - -	- - - -
NAD ⁺	8.43 (s)	- - - -	1.7 ± (1.4) 1.6E-2	- - - -	2.4 ± 1.7 1.6E-2 ^a	1.7 ± (1.5) 3.2E-2	- - - -	- - - -	- - - -
UDP-GlcA	7.95 (d)	- - - -	1.7 ± (1.4) 3.0E-2	- - - -	2.9 ± 1.9 2.7E-3 ^a	- - - -	-2.1 ± 1.6 2.8E-2 ^b	- - - -	- - - -
UMP	5.99 (m)	1.8 ± (1.6) 2.8E-2	- - - -	- - - -	2.2 ± 1.7 1.2E-2 ^a	- - - -	- - - -	- - - -	- - - -
Val	1.05 (d)	-2.5 ± 1.7 8.9E-3 ^a	-2.1 ± 1.5 2.0E-2	-2.0 ± (1.6) 2.1E-2	-2.2 ± 1.7 1.0E-2 ^a	- - - -	- - - -	- - - -	- - - -
U1	0.89 (t)	- - - -	- - - -	-2.5 ± 1.8 1.6E-2	- - - -	1.9 ± (1.5) 1.9E-2 ^b	- - - -	- - - -	- - - -
U2	0.93 (s)	- - - -	-2.1 ± 1.6 1.2E-2	- - - -	- - - -	- - - -	- - - -	- - - -	- - - -
U3	1.62 (d)	- - - -	-2.8 ± 1.7 5.0E-3	- - - -	- - - -	- - - -	- - - -	- - - -	- - - -
U4	2.92 (s)	- - - -	-1.7 ± (1.4) 3.5E-2	- - - -	3.5 ± 2.1 1.1E-3 ^{a,b}	- - - -	2.1 ± 1.6 1.6E-2 ^b	- - - -	- - - -
U5	3.35 (s)	2.0 ± (1.6) 1.8E-2	1.8 ± (1.5) 2.2E-2	- - - -	- - - -	- - - -	- - - -	- - - -	- - - -

Table S2. List of metabolite variations in aqueous liver extracts of mice exposed to cDDP and Pd₂Spm at 1, 12 and 48 h post-injection, compared to controls. Only variations with 95% significance level (p-value < 0.05) are shown; ES, effect size; ES error values in brackets correspond to > 75% of ES value; ^a variations remaining significant after False Discovery Rate correction.; ^b metabolite variations specific of exposure to Pd₂Spm (these comprise both new variations and variations in the same metabolite but in opposite directions). [†] Tentative assignment. 3-HBA, 3-hydroxybutyrate; Ado, adenosine; ATP, adenosine triphosphate; GPC, glycerophosphocholine; GSH, glutathione (reduced); IMP, inosine monophosphate; Ino, inosine; other abbreviations as defined in Table S1.

Metabolite assignment	δ_H ppm multiplicity	cDDP vs. Controls (according to reference [15])						Pd ₂ Spm vs. Controls					
		1 h		12 h		48 h		1 h		12 h		48 h	
		ES ± Error	p-value	ES ± Error	p-value	ES ± Error	p-value	ES ± Error	p-value	ES ± Error	p-value	ES ± Error	p-value
2-aminobutyrate [†]	0.80 (t)	- - - -	-	-1.6 ± (1.4)	4.4E-2	- - - -	-	-1.5 ± (1.4)	3.7E-2 ^a	- - - -	-	- - - -	-
3-HBA	1.20 (d)	- - - -	-	-1.8 ± (1.5)	3.1E-2	- - - -	-	-1.5 ± (1.4)	4.8E-2	- - - -	-	- - - -	-
Acetate	1.92 (s)	- - - -	-	- - - -	-	-2.5 ± 1.7	2.0E-2	- - - -	-	- - - -	-	-1.7 ± (1.5)	4.1E-2
Acetone	2.24 (s)	- - - -	-	-5.1 ± 2.6	5.2E-5 ^a	- - - -	-	- - - -	-	-1.9 ± (1.6)	3.7E-5 ^a	- - - -	-
Ado/ Ino	4.44 (dd)	- - - -	-	- - - -	-	- - - -	-	-1.8 ± (1.5)	2.0E-2 ^b	- - - -	-	- - - -	-
ADP	8.54 (s)	- - - -	-	- - - -	-	2.6 ± 1.8	1.6E-2	- - - -	-	- - - -	-	- - - -	-
AMP	4.51 (dd)	- - - -	-	- - - -	-	2.8 ± 1.8	3.8E-3	- - - -	-	- - - -	-	- - - -	-
Asp	2.67 (dd)	- - - -	-	- - - -	-	- - - -	-	- - - -	-	- - - -	-	-2.0 ± (1.6)	3.6E-2 ^b
ATP	8.52 (s)	- - - -	-	- - - -	-	2.3 ± 1.7	1.6E-2	- - - -	-	- - - -	-	- - - -	-
DMA	2.73 (s)	- - - -	-	-2.4 ± 1.6	6.0E-3 ^a	- - - -	-	3.2 ± 1.9	6.9E-3 ^{a,b}	- - - -	-	- - - -	-
Formate	8.46 (s)	- - - -	-	-1.9 ± (1.5)	7.9E-3 ^a	- - - -	-	- - - -	-	- - - -	-	- - - -	-
GPC	3.23 (s)	- - - -	-	- - - -	-	- - - -	-	- - - -	-	1.6 ± (1.5)	2.1E-2 ^b	- - - -	-
GSH	2.55 (m)	- - - -	-	-4.0 ± 2.1	2.7E-4 ^a	- - - -	-	- - - -	-	- - - -	-	- - - -	-
His	7.08 (s)	- - - -	-	2.2 ± 1.6	9.1E-3	- - - -	-	- - - -	-	- - - -	-	1.6 ± (1.5)	4.3E-2
Ile	1.01 (d)	- - - -	-	- - - -	-	- - - -	-	-2.3 ± 1.6	5.8E-3 ^{a,b}	- - - -	-	- - - -	-
IMP	8.58 (s)	- - - -	-	- - - -	-	1.7 ± (1.5)	3.2E-2	- - - -	-	- - - -	-	- - - -	-
Ino	8.35 (s)	- - - -	-	- - - -	-	- - - -	-	-1.7 ± (1.5)	2.5E-2 ^{a,b}	- - - -	-	- - - -	-
Lactate	4.10 (q)	-2.7 ± 1.7	6.9E-3	1.7 ± (1.5)	3.0E-2	- - - -	-	- - - -	-	- - - -	-	- - - -	-
Leu	0.96 (t)	- - - -	-	- - - -	-	- - - -	-	-2.0 ± 1.5	8.7E-3 ^{a,b}	- - - -	-	- - - -	-
Lys	1.71 (m)	- - - -	-	- - - -	-	- - - -	-	-1.8 ± (1.5)	1.3E-2 ^{a,b}	- - - -	-	2.6 ± 1.8	5.5E-3 ^b
NAD [†]	8.43 (s)	- - - -	-	- - - -	-	2.6 ± 1.8	9.9E-3	- - - -	-	- - - -	-	- - - -	-

Phe	7.33 (d)	- - - -	- - - -	- - - -	- - - -	-1.6 ±(1.4) 2.2E-2 ^{a,b}	- - - -	- - - -	- - - -
Thr, Lactate	1.33 (d)	-2.3 ± 1.6 3.2E-2	1.9 ±(1.5) 1.9E-2	- - - -	- - - -	- - - -	- - - -	- - - -	- - - -
Tyr	6.90 (d)	- - - -	- - - -	1.9 ±(1.6) 2.3E-2	-1.7 ±(1.4) 1.9E-2 ^{a,b}	- - - -	- - - -	- - - -	- - - -
UDP-GlcA	7.95 (d)	- - - -	-1.7 ±(1.4) 3.0E-2	- - - -	- - - -	- - - -	- - - -	- - - -	- - - -
Uridine	7.86 (d)	- - - -	- - - -	- - - -	-2.6 ± 1.7 2.4E-3 ^{a,b}	- - - -	- - - -	- - - -	- - - -
Val	1.05 (d)	- - - -	- - - -	- - - -	-1.9 ±(1.5) 1.1E-2 ^{a,b}	- - - -	- - - -	1.9 ±(1.6) 2.3E-2 ^b	- - - -
U6	0.73 (s)	-2.2 ± 1.6 1.6E-2	- - - -	2.6 ± 1.8 1.6E-2	- - - -	- - - -	- - - -	- - - -	- - - -
U7	0.85 (t)	- - - -	-1.6 ±(1.4) 1.6E-2	- - - -	- - - -	- - - -	- - - -	- - - -	- - - -
U8	3.10 (d)	- - - -	2.5 ± 1.7 9.8E-3 ^a	- - - -	2.5 ± 1.7 8.4E-3 ^a	- - - -	- - - -	- - - -	- - - -
U9	4.31 (d)	-1.6 ±(1.4) 4.7E-2	-1.8 ±(1.5) 2.1E-2	1.7 ±(1.5) 3.3E-2	- - - -	- - - -	- - - -	- - - -	- - - -
U10	6.18 (s)	-1.7 ±(1.5) 3.6E-2	-2.6 ± 1.7 7.8E-3 ^a	2.2 ± 1.7 2.3E-2	- - - -	- - - -	- - - -	- - - -	- - - -
U11	8.28 (br)	- - - -	-1.8 ±(1.5) 3.5E-2	1.8 ±(1.5) 4.2E-2	- - - -	- - - -	- - - -	- - - -	- - - -

Table S3. List of metabolite variations found in aqueous breast tissue extracts of mice exposed to cDDP and Pd₂Spm at 1, 12 and 48 h post-injection, compared to controls. Only variations with 95% significance level (p-value < 0.05) are shown; ES, effect size; ES error values in brackets correspond to > 75% of ES value; ^a variations remaining significant after False Discovery Rate correction, ^b metabolite variations specific of exposure to Pd₂Spm (these comprise both new variations and variations in the same metabolite but in opposite directions). [†]Tentative assignment. Abbreviations as defined in Tables S1 and S2.

Metabolite assignment	δ_H ppm multiplicity	cDDP <i>vs.</i> Controls (according to reference [15])						Pd ₂ Spm <i>vs.</i> Controls					
		1 h		12 h		48 h		1 h		12 h		48 h	
		ES \pm Error	p-value	ES \pm Error	p-value	ES \pm Error	p-value	ES \pm Error	p-value	ES \pm Error	p-value	ES \pm Error	p-value
3-HBA	1.20 (d)	1.8 \pm (1.5)	4.3E-2	- - -	-	- - -	-	- - -	-	- - -	-	- - -	-
Ado/Ino	6.10 (d)	- - -	-	-2.4 \pm 1.6	6.1E-3 ^a	- - -	-	- - -	-	-1.9 \pm (1.5)	1.5E-2	- - -	-
ADP	8.54 (s)	- - -	-	1.7 \pm (1.4)	3.1E-2 ^a	- - -	-	1.9 \pm (1.5)	1.6E-2	4.5 \pm 2.3	1.8E-4 ^a	- - -	-
Creatinine	3.04 (s)	- - -	-	- - -	-	- - -	-	- - -	-	- - -	-	1.6 \pm (1.5)	3.2E-2 ^b
DMA	2.73 (s)	- - -	-	- - -	-	- - -	-	3.6 \pm 2.0	4.9E-3 ^{a,b}	- - -	-	- - -	-
DMSO ₂ [†]	3.15 (s)	- - -	-	- - -	-	- - -	-	8.8 \pm 4.0	1.8E-6 ^{a,b}	- - -	-	- - -	-
Formate	8.46 (s)	- - -	-	- - -	-	- - -	-	- - -	-	-1.6 \pm (1.4)	4.7E-2 ^b	- - -	-
Gln	2.45 (m)	- - -	-	- - -	-	-2.2 \pm 1.6	1.2E-2	- - -	-	- - -	-	- - -	-
Gly	3.56 (s)	- - -	-	- - -	-	- - -	-	-1.6 \pm (1.4)	4.6E-2 ^b	- - -	-	- - -	-
GSH	2.55 (m)	- - -	-	- - -	-	- - -	-	1.8 \pm (1.5)	2.0E-2 ^b	- - -	-	- - -	-
Guanine [†]	7.68 (s)	- - -	-	- - -	-	-1.9 \pm (1.6)	3.1E-2	- - -	-	-1.9 \pm (1.5)	2.5E-2	-1.8 \pm (1.6)	3.2E-2
Ino	8.35 (s)	- - -	-	-2.8 \pm 1.7	2.4E-3 ^a	- - -	-	- - -	-	-2.3 \pm 1.6	6.8E-3 ^a	- - -	-
Tyr	6.90 (d)	- - -	-	- - -	-	- - -	-	- - -	-	-1.6 \pm (1.4)	3.7E-2 ^b	- - -	-
U12	1.25 (s)	- - -	-	2.0 \pm 1.5	2.6E-2 ^a	- - -	-	2.1 \pm 1.6	1.7E-2	2.0 \pm 1.5	2.4E-2	- - -	-

Table S4. List of metabolite variations found in lipophilic extracts of kidney, liver and breast tissue of mice exposed to cDDP and Pd₂Spm at 1, 12 and 48 h post-injection, compared to controls. Only variations with 95% significance level (p-value < 0.05) are shown; ES, effect size; ES error values in brackets correspond to > 75% of ES values; ^a variations remaining significant after False Discovery Rate correction, ^b metabolite variations specific of exposure to Pd₂Spm (these comprise both new variations and variations in the same metabolite but in opposite directions).

Metabolite family/ assignment		δ_H ppm multiplicity	cDDP vs. Controls						Pd ₂ Spm vs. Controls					
			1 h		12 h		48 h		1 h		12 h		48 h	
			ES \pm Error	p-value	ES \pm Error	p-value	ES \pm Error	p-value	ES \pm Error	p-value	ES \pm Error	p-value	ES \pm Error	p-value
KIDNEY														
Cholesterol	Free/ Esterified, C18H ₃	0.68 (s)	1.9 \pm (1.5)	2.0E-2 ^a	- - -	-	- - -	-	- - -	-	- - -	-	- - -	-
	Free, C4H'	2.27 (t)	1.7 \pm (1.4)	3.2E-2 ^a	- - -	-	- - -	-	- - -	-	- - -	-	- - -	-
	Esterified, C4H'	2.31 (t)	1.7 \pm (1.5)	2.5E-2 ^a	- - -	-	-2.9 \pm 1.9	7.7E-3 ^a	- - -	-	- - -	-	- - -	-
	Free, C3H	3.53 (m)	- - -	-	- - -	-	- - -	-	- - -	-	- - -	-	-1.6 \pm (1.5)	4.2E-2 ^b
Fatty acids	CH ₃	0.89 (br)	- - -	-	- - -	-	1.7 \pm (1.5)	4.4E-2	- - -	-	- - -	-	2.1 \pm (1.6)	1.7E-2
	=CHCH ₂ (CH ₂) _n	1.30 (br)	-1.6 \pm (1.4)	3.2E-2 ^a	- - -	-	- - -	-	- - -	-	- - -	-	1.7 \pm (1.5)	4.3E-2 ^b
	CH ₂ CO	2.30 (m)	-1.6 \pm (1.4)	3.7E-2 ^a	- - -	-	- - -	-	- - -	-	- - -	-	1.6 \pm (1.5)	4.9E-2 ^b
	18:2 =CHCH ₂ CH=	2.77 (t)	-1.6 \pm (1.4)	3.2E-2 ^a	- - -	-	- - -	-	- - -	-	- - -	-	- - -	-
	18:2 HC=CH	5.34 (m)	- - -	-	- - -	-	- - -	-	- - -	-	- - -	-	1.7 \pm (1.5)	1.6E-2 ^b
	18:2/ 20:4 CH ₃	0.98 (t)	1.5 \pm (1.4)	4.4E-2	- - -	-	- - -	-	- - -	-	- - -	-	-1.8 \pm (1.6)	2.7E-2 ^b
	20:4 =CHCH ₂ CH=	2.81 (m)	- - -	-	- - -	-	- - -	-	- - -	-	- - -	-	-1.6 \pm (1.5)	4.9E-2 ^b
	20:4 HC=CH	5.37 (m)	1.6 \pm (1.4)	3.2E-2 ^a	- - -	-	- - -	-	- - -	-	- - -	-	- - -	-
Phospholipids	CH ₂ N(CH ₃) ₃	3.75 (br)	1.8 \pm (1.5)	2.4E-2 ^a	- - -	-	-2.6 \pm 1.8	7.9E-3 ^a	- - -	-	- - -	-	-1.6 \pm (1.5)	4.7E-2
	Glycerol C3H ₂	3.94 (br)	2.5 \pm 1.7	4.6E-3 ^a	- - -	-	-3.9 \pm 2.2	7.0E-4 ^a	- - -	-	- - -	-	- - -	-
	Glycerol C1H ₂	4.38 (br)	- - -	-	- - -	-	- - -	-	- - -	-	- - -	-	-1.6 \pm (1.5)	5.0E-2 ^b
	Glycerol C2H	5.20 (br)	2.0 \pm 1.5	3.2E-2 ^a	- - -	-	-2.1 \pm 1.6	2.0E-2	- - -	-	- - -	-	- - -	-
	PE CH ₂ NH ₃	3.16 (br)	- - -	-	- - -	-	-1.8 \pm (1.5)	1.6E-2	- - -	-	- - -	-	-1.8 \pm (1.6)	2.8E-2
	PTC/SM, N(CH ₃) ₃	3.29/3.31 (s)	1.7 \pm (1.4)	3.2E-2 ^a	- - -	-	-1.6 \pm (1.5)	5.0E-2	- - -	-	- - -	-	-1.7 \pm (1.5)	3.5E-2
Triglycerides	Glycerol C1, 3H	4.14 (dd)	- - -	-	- - -	-	- - -	-	- - -	-	- - -	-	1.8 \pm (1.6)	2.9E-2 ^b
	Glycerol C2H	5.26 (m)	-1.8 \pm (1.5)	2.3E-2 ^a	- - -	-	1.7 \pm (1.5)	4.3E-2	- - -	-	- - -	-	- - -	-
Lipid ratios	(CH ₂) _n /CH ₃	-	- - -	-	- - -	-	- - -	-	- - -	-	- - -	-	-1.6 \pm (1.5)	4.9E-2 ^b
	HC=CH/CH ₃	-	- - -	-	- - -	-	- - -	-	- - -	-	- - -	-	-1.7 \pm (1.5)	3.2E-2 ^b

Unknowns	U1	0.82 (br)	- - - -	- - - -	- - - -	-3.0 ± 1.8	4.4E-2 ^b	- - - -	- - - -		
	U2	2.38 (br)	- - - -	- - - -	- - - -	- - - -	- - - -	-1.8 ± (1.6)	3.1E-2 ^b		
	U3	4.07 (br)	1.9 ± (1.5)	1.9E-2 ^a	- - - -	- - - -	- - - -	- - - -	-1.6 ± (1.5)	3.9E-2 ^b	
	U4	4.20 (dd)	- - - -	- - - -	- - - -	- - - -	- - - -	11.1 ± 5.0	1.7E-7 ^{a,b}	2.7 ± 1.8	6.5E-3 ^b
	U5	4.69 (d)	- - - -	- - - -	- - - -	- - - -	- - - -	- - - -	- - - -	-2.7 ± 1.8	3.9E-3 ^b
	U6	5.90 (d)	- - - -	- - - -	- - - -	-1.9 ± (1.6)	2.6E-2	- - - -	- - - -	-1.9 ± (1.6)	2.3E-2
	U7	7.88 (br)	1.8 ± (1.5)	2.7E-2 ^a	- - - -	-3.1 ± 2.0	2.6E-3 ^a	- - - -	- - - -	-2.1 ± (1.6)	3.2E-2
LIVER											
Cholesterol	Free/ Esterified, C26H ₃	0.86 (d)	- - - -	- - - -	- - - -	1.6 ± (1.5)	4.3E-2	- - - -	- - - -	- - - -	
	Free, C19H ₃	1.01 (s)	- - - -	- - - -	- - - -	- - - -	- - - -	1.6 ± (1.4)	4.5E-2 ^b	- - - -	
Fatty acids	CH ₃	0.89 (br)	- - - -	- - - -	- - - -	- - - -	- - - -	2.0 ± 1.5	3.0E-2 ^b	- - - -	
	(CH ₂) _n	1.25 (br)	- - - -	- - - -	- - - -	- - - -	- - - -	2.1 ± 1.6	1.7E-2 ^b	- - - -	
	18:2/ 20:4 CH ₃	0.98(t)	- - - -	- - - -	- - - -	2.4 ± 1.7	6.9E-3 ^a	- - - -	- - - -	- - - -	
	20:4 CH ₂ CH=	2.07 (q)	-3.2 ± 2.0	1.9E-2 ^a	- - - -	- - - -	- - - -	-2.2 ± 1.7	1.6E-2 ^b	- - - -	
	20:4 =CHCH ₂ CH=	2.81 (m)	-2.2 ± 1.7	1.7E-2 ^a	- - - -	- - - -	- - - -	- - - -	- - - -	- - - -	
	20:4 HC=CH	5.37 (m)	-2.4 ± 1.7	1.3E-2 ^a	- - - -	- - - -	- - - -	-1.6 ± (1.5)	3.7E-2	- - - -	
	22:6 CH ₂ CO	2.37 (br)	-2.2 ± 1.7	4.6E-2	- - - -	1.9 ± (1.6)	2.9E-2	-1.9 ± (1.6)	4.1E-2	- - - -	
BREAST TISSUE											
Cholesterol	Free/ Esterified, C3H	1.02 (s)	- - - -	-1.7 ± (1.4)	4.0E-2 ^a	- - - -	- - - -	-1.8 ± (1.5)	3.2E-2	- - - -	
Fatty Acids	CH ₃	0.89 (br)	1.7 ± (1.4)	3.0E-2	- - - -	- - - -	- - - -	- - - -	- - - -	- - - -	
	(CH ₂) _n	1.25 (br)	- - - -	-2.7 ± 1.7	4.7E-3 ^a	- - - -	- - - -	-1.7 ± (1.5)	3.0E-2 ^a	- - - -	
	=CHCH ₂ (CH ₂) _n	1.30 (br)	- - - -	2.2 ± 1.6	1.2E-2 ^a	- - - -	- - - -	- - - -	- - - -	- - - -	
	CH ₂ CO	2.30 (m)	- - - -	1.8 ± (1.5)	3.2E-2 ^a	- - - -	- - - -	- - - -	- - - -	- - - -	
	18:2, HC=CH	5.34 (m)	- - - -	2.9 ± 1.8	2.9E-3 ^a	- - - -	- - - -	- - - -	1.8 ± (1.5)	2.2E-2 ^a	
	18:2, =CHCH ₂ CH=	2.77 (t)	- - - -	2.2 ± 1.6	8.1E-3 ^a	- - - -	- - - -	- - - -	1.6 ± (1.4)	1.6E-2 ^a	
	20:4, HC=CH	5.36 (m)	- - - -	2.3 ± 1.6	6.9E-3 ^a	- - - -	- - - -	- - - -	1.7 ± (1.4)	3.3E-2 ^a	
	20:4, CH ₂ CH=	2.07 (q)	- - - -	2.2 ± 1.6	8.8E-3 ^a	- - - -	- - - -	1.5 ± (1.4)	4.7E-2	- - - -	
Triglycerides	Glyceryl C1, 3H	4.14 (dd)	- - - -	1.7 ± (1.4)	3.9E-2 ^a	- - - -	- - - -	- - - -	- - - -	- - - -	
	Glyceryl C1, 3H'	4.29 (dd)	- - - -	1.8 ± (1.5)	2.9E-2 ^a	- - - -	- - - -	- - - -	- - - -	- - - -	
	Glyceryl C2H	5.26 (m)	- - - -	4.4 ± 2.3	1.3E-4 ^a	- - - -	- - - -	1.5 ± (1.4)	4.8E-2	2.6 ± 1.7	4.3E-3 ^a
Lipid ratios	(CH ₂) _n /CH ₃	-	-1.7 ± (1.4)	4.5E-2	-1.7 ± 1.4	2.9E-2 ^a	- - - -	- - - -	-2.1 ± 1.5	1.3E-2 ^a	
	HC=CH/CH ₃	-	-1.8 ± (1.5)	3.7E-2	- - - -	- - - -	- - - -	- - - -	-1.6 ± (1.4)	3.7E-2 ^a	
Unknowns	U4	4.20 (dd)	- - - -	-5.0 ± 2.5	3.4E-4 ^a	-3.2 ± 2.0	2.3E-2	-1.9 ± (1.5)	2.0E-2 ^a	-4.3 ± 2.3	8.1E-4 ^a

U8	2.17 (s)	3.1 ± 1.8	1.5E-3 ^a	- - -	-	- - -	-	- - -	-	- - -	-	- - -	-
U9	3.60 (dd)	3.0 ± 1.8	1.8E-3 ^a	-5.3 ± 2.6	3.0E-4 ^a	-6.6 ± 3.3	2.4E-3 ^a	-3.2 ± 1.9	2.3E-3 ^a	-5.1 ± 2.5	9.3E-4 ^a	-6.0 ± 3.1	1.3E-3 ^a
U10	3.64 (s)	- - -	-	-4.0 ± 2.1	4.3E-4 ^a	-4.0 ± 2.3	1.6E-2	-2.3 ± 1.6	1.5E-2 ^a	-4.0 ± 2.1	2.4E-3 ^a	- - -	-
U11	4.38 (br)	- - -	-	-1.7 ± (1.4)	3.8E-2 ^a	- - -	-	- - -	-	- - -	-	- - -	-