Dual Internal Standards with Metals and Molecules for MALDI Imaging of Kidney

Lipids

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Supporting Information

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Direct injection and laser ablation ICP-MS analysis

Nitrocellulose membranes (NC) (Whatman[®] Protran[®] nitrocellulose membranes, Dassel, Germany) of approximately 10x10 mm were fixed to glass slides next to the tissue sections and printed simultaneously. The printed membranes were collected and used for direct infusion ICP-MS as well as LA-ICP-MS.

Direct infusion ICP-MS

The collected NC membranes were weighed and digested in 150 μ L 69% HNO₃ (HNO₃, ultraquality, Carl Roth, Germany) for direct injection ICP-MS analysis. Digested samples were diluted in a ratio of 1:10 in 3.5% HNO₃ including 10 μ L ¹⁴¹Pr (100 μ g L⁻¹) as internal standard to a final volume of 500 μ L. For external calibration, a solution series from 20 ng L⁻¹ to 10 μ g L⁻¹ using ¹⁶⁹Tm standard solution was prepared. The Element XR sector field ICP-MS (Thermo Fisher Scientifc, Bremen, Germany) was connected to a Meinhard-type nebulizer with a cyclonic spray chamber (MicroMist, Glass Expansion, Melbourne, Australia) using a nebulizer gas flow rate of 1.0 L min⁻¹ Ar. The instrument was operated at plasma gas flow of 15 L min⁻¹, auxiliary gas flow of 1.00 L min⁻¹, and RF plasma power of 1250 W.

LA-ICP-MS

LA-ICP-MS analysis was performed on a commercial LA system (UP-213, ESI, Portland, USA) coupled to a sector field ICP-MS (Element XR, Thermo Fisher Scientific, Bremen, Germany). The ICP-MS was synchronized using the LA unit in an external trigger mode. The operating conditions are shown in Table S1. The ICP-MS operating conditions were tuned daily for maximum intensity and low oxide ratio ((ThO/Th) < 0.7%) using a reference glass slide (SRM 612, NIST, Maryland USA). The printed nitrocellulose membranes were ablated continuously in line scans. Overlapping line scans and high laser shot repetition rates were applied to ensure complete ablation of the membrane sample.

Mass traces of ¹⁶⁹Tm were recorded at low resolution (R=300). The analysis of a nitrocellulose membrane (with 2×2 mm average dimensions) required about 20 min. Data was exported to Origin 2016 (OriginLab Corporation, Northampton, MA) where data normalization and generation of color coded images were performed.

S2

ICP-MS		LA system			
RF plasma power	1257 W	Wavelength	213 nm		
Plasma gas flow (Ar)	15 L min ⁻¹	Helium gas flow	0.8 L min ⁻¹		
Sample gas flow (Ar)	1.3 L min ⁻¹	Laser energy	100% (5.16 J cm- ²)		
Auxiliary gas flow (Ar)	0.99 L min ⁻¹	Laser spot size	100 µm		
Mass resolution ($m/\Delta m$)	300 (LR)	Scan speed	100 μm s ⁻¹		
Sample time	2 ms	Repetition rate	20 Hz		
Scanning mode	E scan	Line overlapping	20%		
Detected isotope	¹⁶⁹ Tm				

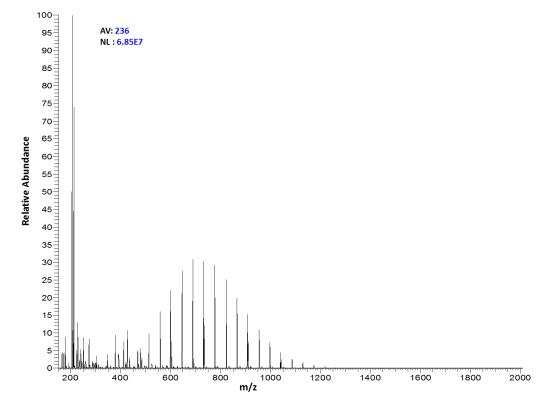


Figure S1. Mass spectrum showing background effect produced when colorless and yellow ink were used to dissolve DHB and the internal standards.

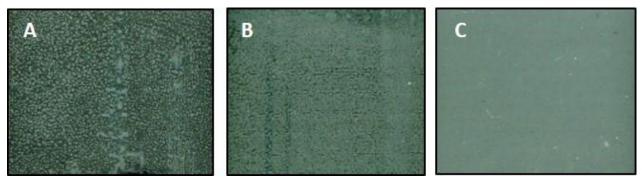
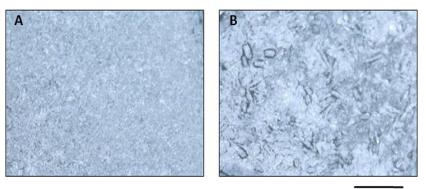


Figure S2. Effect of increasing organic solvent composition on the homogenity of the resulting printing output, where A) 20% MeOH, B) 50% MeOH, and C) 70% MeOH.



1.00 mm

Figure S3. Digital microscope camera images of matrix crystals after 5 printing cycles of ink solution (DHB = 120 mg mL^{-1} in 70% MeOH, 0.1% TFA). A) Pre-heating of the glass slide during 5s on a hot plate at 50 °C before each printing cycle, and B) at room temperature.

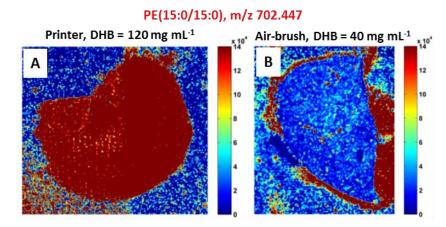


Figure S4. Effect of increasing DHB concentration on the ionization of the internal standard. Extracted ion images of $[M+K]^+$ of the PE(15:0/15:0) internal standard at m/z 702.447 obtained for (A): inkjet printer at 120 mg mL⁻¹ DHB, and (B): air-brush nebulizer at 40 mg mL⁻¹ DHB. Both tissues present 30 printing cycles and the same internal standard concentration.

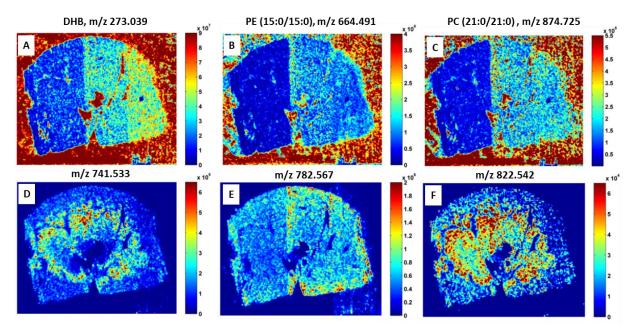


Figure S5. The effect of the number of printing cycles on the extracted ion images of a kidney tissue section with 20 printing cycles on the left side and 30 on the right side. (A) DHB, $([2M-2H_2O]^+ \text{ at } m/z \ 273.039)$. (B) $[M+H]^+$ adduct of the PE (15:0/15:0) internal standard at $m/z \ 664.491$. (C) $[M+H]^+$ adduct of the PC (21:0/21:0) internal standard at $m/z \ 741.533$, 782.567, and 822.542, respectively.

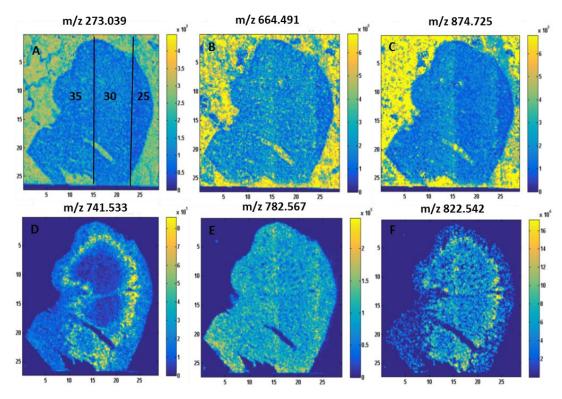


Figure S6. Kidney tissue section with 35 printing cycles on the left side, 30 on the middle part, and 25 on the right side. No enhancement of matrix and internal standard signal upon printing more than 30 cycles (A to C). (A) DHB, $[2M-2H_2O]^+$ (m/z 273.039). (B) $[M+H]^+$ adduct of the PE (15:0/15:0) internal standard (m/z 664.487). (C) $[M+H]^+$ adduct of the PC (21:0/21:0) internal standard (m/z 874.725). (D to F) Tissue lipids showing stable ionization at m/z 741.528, m/z 782.565, and m/z 822.542, respectively.

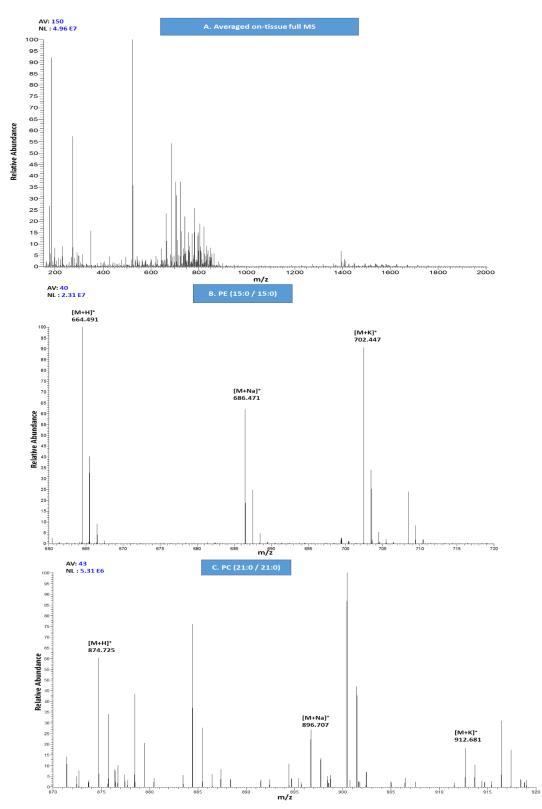


Figure S7. (A) Mass spectrum illustrates the spectral quality of a kidney tissue section printed with 30 cycles of matrix-internal standards mixture, tissue lipids are observed in the m/z range from 700-900. (B) H^+ , Na^+ and K^+ adducts of the lipidic internal standard PE (15:0/15:0) at m/z 664.491, m/z 686.470 and m/z 702.447, respectively. (C) H^+ (m/z 874.725), Na^+ (m/z 896.707), and K^+ (m/z 912.680) adducts of the lipidic internal standard PC (21:0/21:0).

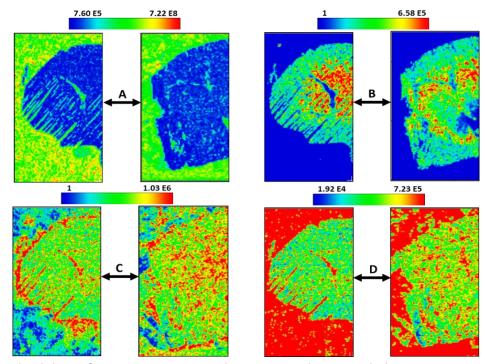


Figure S8. Reproducibility of MALDI images is demonstrated in two kidney tissue sections printed according to optimized parameters. (A) DHB signal at m/z 273.039. (B) Endogenous lipid at m/z 741.533. (C) $[M+K]^+$ of the PE (15:0/2115:0) internal standard at m/z 702.447. (D) $[M+Na]^+$ of the PC (21:0/21:0) internal standard at m/z 896.707.

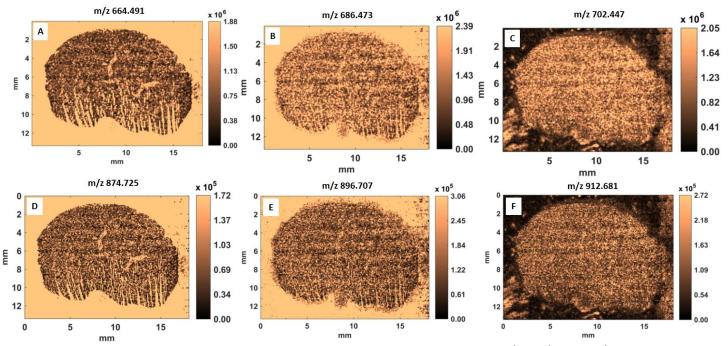


Figure S9. (A-C) Ion extracted images showing on-tissue distribution of H^+ , Na^+ , and K^+ adducts of IS PE (15:0/15:0) at m/z 664.491, 686.473, and 702.447, respectively. (D-F) Ion images showing on-tissue distribution of H^+ , Na^+ and K^+ adducts of IS PC (21:0/21:0) at m/z 874.725, m/z 896.707 and m/z 912.681, respectively.

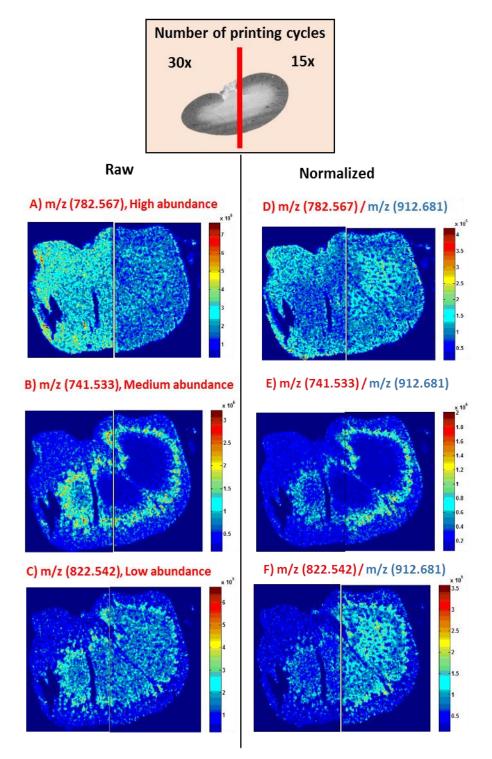
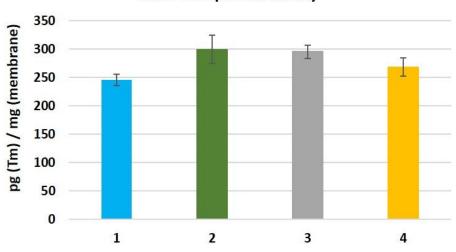


Figure S10. Color codes illustrate the effect of a proportional variation of matrix and internal standards amounts. Endogenous lipids with high, medium and low abundance are shown in panels A, B and C, respectively. Panels D, E and F show the same m/z images corrected by $M+K^+$ of PC internal standard, m/z 912.681.



Inter-sample variability

Figure S11. Metal amounts in mineralized membranes printed simultaneously with the tissues. Nitrocellulose membrane from each printing experiment weighed approx. 2-4 mg. Metal content in the different samples showed a mean value of 277 pg (Tm) / mg (membrane) with a RSD value of 11%. Error bars indicate the s.e.m

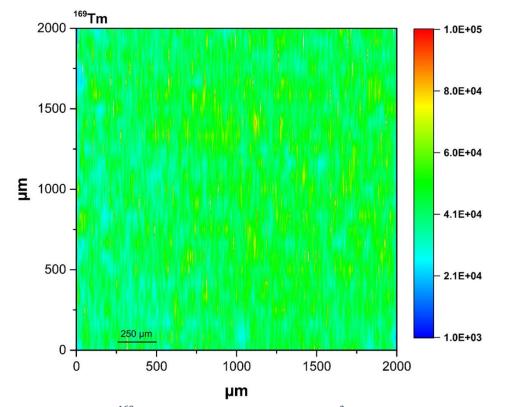


Figure S12. Intensity profile of ¹⁶⁹Tm using LA-ICP-MS of a 2 mm² area of a NC membrane, which received 30 printing cycles of the optimized matrix/internal standards mixture. RSD value of 2-4% was found between the different ablated lines within the membrane.

where the mean value in each sample is calculated using 3 replicates.

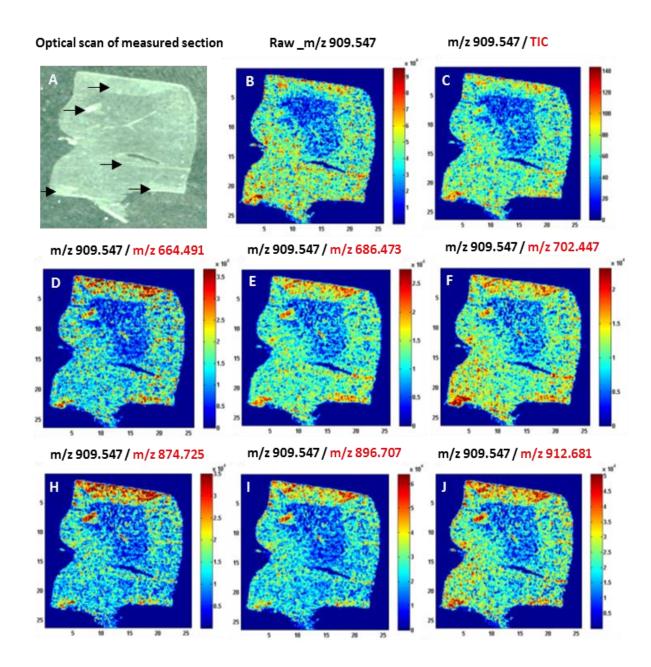
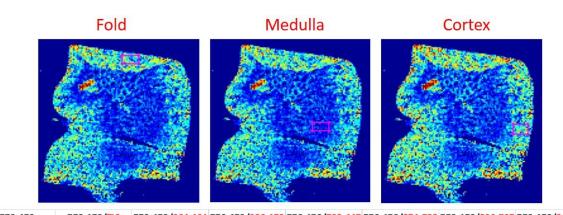


Figure S13. Improved reflection of tissue irregularities observed in internal standard normalized MALDI images for m/z 909.547 (PI(38:4), $[M+Na]^{\dagger}$). (A) Optical scan of the measured tissue. (B) Raw MALDI image of m/z 909.547. (C) TIC-normalized image. (D to F) Normalized images of m/z 909.547 to $[M+H]^{\dagger}$, $[M+Na]^{\dagger}$ and $[M+K]^{\dagger}$ of PE (15:0/15:0) at m/z 664.491, m/z 686.473, and m/z 702.447, respectively. (G to I) Normalized images of m/z 909.547 to $[M+H]^{\dagger}$, $[M+Na]^{\dagger}$ and $[M+K]^{\dagger}$ of PC(21:0/21:0) at m/z 874.725, m/z 896.707 and m/z 912.681, respectively.



	778.478	778.478/TIC	778.478/664.491	778.478/686.473	778.478/702.447	778.478/874.725	778.478/ <mark>896.707</mark>	778.478/912.68
				Fold				
avg	78951.38	122.81	39887.16	27795.82	22967.90	42253.25	64662.98	55277.57
stdev	21476.53	38.53	12911.01	7129.54	5775.75	13522.21	18286.00	14202.42
rsd	27.20	31.37	32.37	25.65	25.15	32.00	28.28	25.69
				Medulla				
avg	41594.96	57.03	13294.88	10811.85	9380.80	12110.12	20591.45	19991.35
stdev	9525.90	14.52	3356.07	2042.76	1606.40	2538.37	3675.16	3455.46
rsd	22.90	25.46	25.24	18.89	17.12	20.96	17.85	17.28
				Cortex				
avg	86864.65	131.96	34150.45	25822.16	24469.28	29607.06	53337.28	52443.89
stdev	18544.02	29.69	8884.57	4391.21	3884.72	7630.15	11291.18	9789.22
rsd	21.35	22.50	26.02	17.01	15.88	25.77	21.17	18.67
	754.536	754.536/TIC	754.536/664.491	754.536/686.473	754.536/702.447	754.536/874.725	754.536/896.707	754.536/912.68

		754.550	7 34.3 30/110	734.330/004.431	134.330/000.473	134.330/102.441	134.330/014.123	134.330/030.707	1 34.330/ 512.001
					Fold				
a]	avg	45717.28	92.27	25176.95	19579.69	13582.36	26041.86	38597.79805	32665.08
	stdev	11034.78	26.92	7553.80	4089.61	3174.74	7236.63	8409.485431	7509.60
	rsd	24.14	29.18	30.00	20.89	23.37	27.79	21.79	22.99
	Medulla								
32:1)+	avg	41224.18	52.12	10717.90	10754.05	8474.25	9319.13	19419.51	17340.52
	stdev	8819.66	13.42	2936.73	1683.67	1521.63	2201.89	3237.85	3144.14
	rsd	21.39	25.75	27.40	15.66	17.96	23.63	16.67	18.13
Ĵ					Cortex				
	avg	43038.59	63.75	13740.25	12085.05	12047.11	14476.04	26131.57	26007.62
	stdev	10909.54	14.15	3815.47	2589.03	3021.71	3659.00	5769.83	6230.95
	rsd	25.35	22.20	27.77	21.42	25.08	25.28	22.08	23.96

Figure S14. Mean relative intensity calculated in different ROIs (regions of interest) in the tissue section for m/z 778.478 and m/z 754.536 (marked in magenta: Upper, Middle and Right, each ROI contains approx. 100 pixels). The tables show average, standard deviation and relative standard deviation for non-normalized, TIC-normalized and different internal standards adducts–normalized intensity at m/z 664.491, m/z 686.473, m/z 702.447, m/z 874.725, m/z 896.707 and m/z 912.681, corresponding to $[M+H]^+$, $[M+Na]^+$, $[M+K]^+$ for PE(15:0/15:0) and PC(21:0/21:0), respectively.

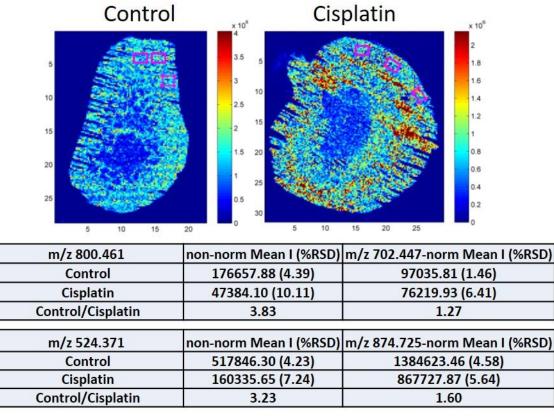


Figure S15. Raw and IS-normalized mean intensity and %RSD over 3 ROIs (highlighted in the MALDI images in magenta, each ROI has 75 pixel) in the cortex of control and cisplatin-treated tissues for m/z 524.371 and m/z 800.461.