

A multimodal desorption electrospray ionisation workflow enabling visualisation of lipids and biologically relevant elements in a single tissue section

Catia Costa ¹, Janella De Jesus ^{2,3}, Chelsea Nikula ³, Teresa Murta ³, Geoffrey W. Grime ¹, Vladimir Palitsin ¹, Véronique Dartois ⁴, Kaya Firat ⁴, Roger Webb ¹, Josephine Bunch ³ and Melanie J. Bailey ^{1,2,*}

¹ University of Surrey Ion Beam Centre, Guildford GU2 7XH, UK; c.d.costa@surrey.ac.uk (C.C.); g.grime@surrey.ac.uk (G.W.G.); v.palitsin@surrey.ac.uk (V.P.); r.webb@surrey.ac.uk (R.W.)

² Department of Chemistry, University of Surrey, Guildford GU2 7XH, UK; janella.marie.de.jesus@npl.co.uk

³ The National Physical Laboratory, Teddington TW11 0LW, UK; c.jnikula@gmail.com (C.N.); tigoncalvesmurta@gmail.com (T.M.); josephine.bunch@npl.co.uk (J.B.)

⁴ Center for Discovery and Innovation, Hackensack Meridian School of Medicine, Nutley, NJ 07110, USA; veronique.dartois@hnh-cdi.org (V.D.); firat.kaya@hnh-cdi.org (K.F.)

* Correspondence: m.bailey@surrey.ac.uk; Tel.: +44-(0)1483682593

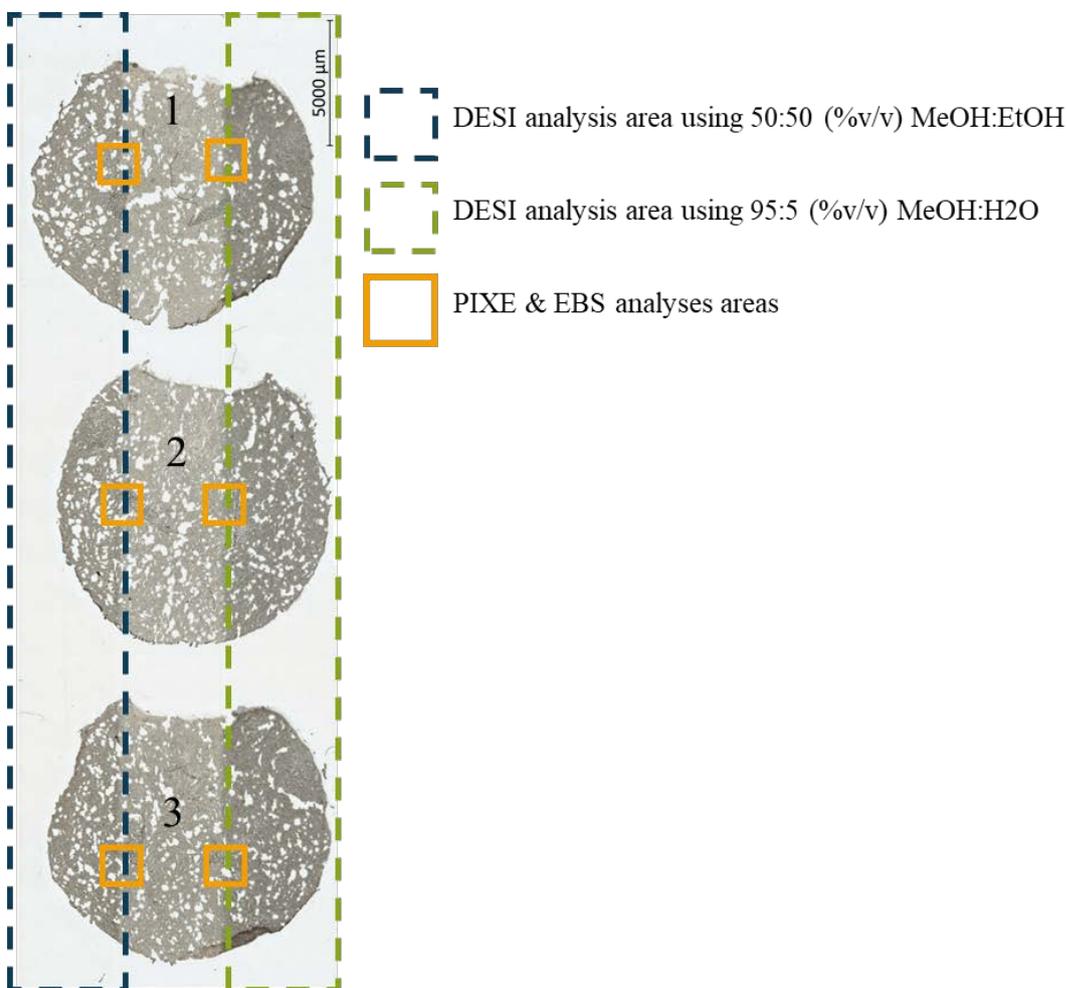
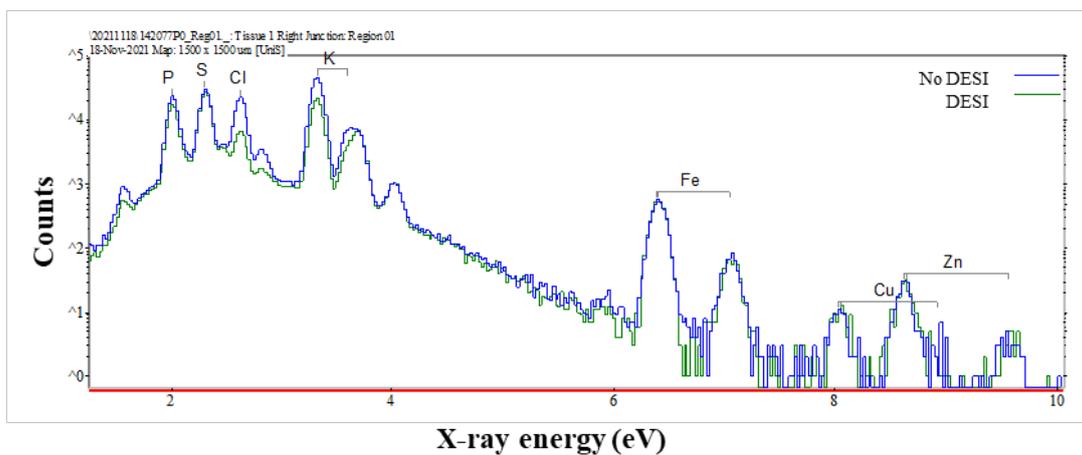


Figure S1. Areas analysed by DESI, PIXE & EBS on liver tissue homogenate sections.

(A)



(B)

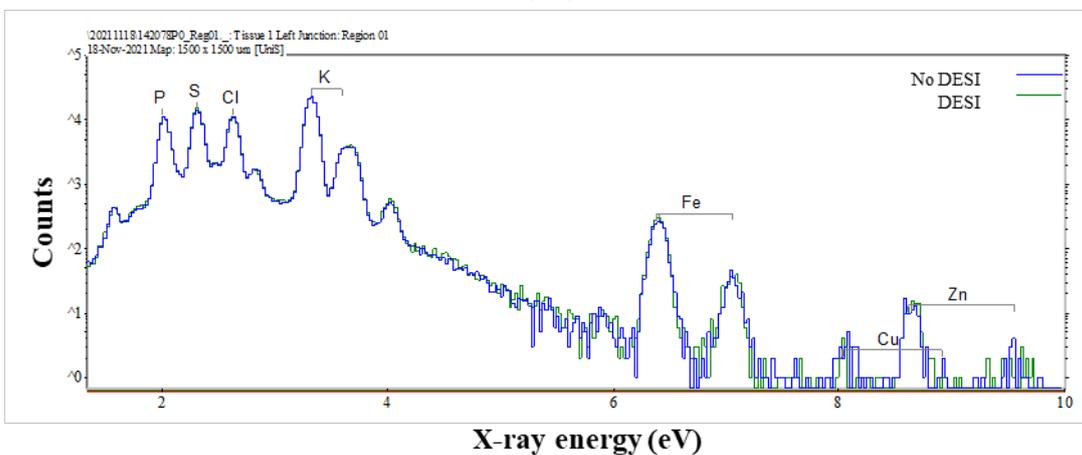


Figure S2. Overlay of X-ray spectra taken from regions of interest on areas not analysed (Reg01; blue) and analysed by DESI (Reg02; green) using (A) 95:5 (% v/v) MeOH:H₂O and (B) 50:50 (% v/v) MeOH:EtOH.

Table S1. T-test results between the average (n=3) of the ratios of the elemental peak area measured by PIXE after DESI analysis with two different solvents – MeOH/H₂O and MeOH/EtOH.

Table Analyzed	P	Table Analyzed	S	Table Analyzed	Cl
Column B	MeOH/EtOH	Column B	MeOH/EtOH	Column B	MeOH/EtOH
vs.	vs.	vs.	vs.	vs.	vs.
Column A	MeOH/H ₂ O	Column A	MeOH/H ₂ O	Column A	MeOH/H ₂ O
Unpaired t test		Unpaired t test		Unpaired t test	
P value	0.0004	P value	0.0016	P value	<0.0001
P value summary	***	P value summary	**	P value summary	****
Significantly different (P < 0.05)?	Yes	Significantly different (P < 0.05)?	Yes	Significantly different (P < 0.05)?	Yes
One- or two-tailed P value?	Two-tailed	One- or two-tailed P value?	Two-tailed	One- or two-tailed P value?	Two-tailed
t, df	t=11.28, df=4	t, df	t=7.566, df=4	t, df	t=25.22, df=4
How big is the difference?		How big is the difference?		How big is the difference?	
Mean of column A	0.6233	Mean of column A	0.7667	Mean of column A	0.2567
Mean of column B	1.007	Mean of column B	1.087	Mean of column B	0.96
Difference between means (B - A) ± SEM	0.3833 ± 0.03399	Difference between means (B - A) ± SEM	0.3200 ± 0.04230	Difference between means (B - A) ± SEM	0.7033 ± 0.02789
95% confidence interval	0.2890 to 0.4777	95% confidence interval	0.2026 to 0.4374	95% confidence interval	0.6259 to 0.7808
R squared (eta squared)	0.9695	R squared (eta squared)	0.9347	R squared (eta squared)	0.9938
F test to compare variances		F test to compare variances		F test to compare variances	
F, DFn, Dfd	13.86, 2, 2	F, DFn, Dfd	39.25, 2, 2	F, DFn, Dfd	9.000, 2, 2
P value	0.1346	P value	0.0497	P value	0.2
P value summary	ns	P value summary	*	P value summary	ns
Significantly different (P < 0.05)?	No	Significantly different (P < 0.05)?	Yes	Significantly different (P < 0.05)?	No
Data analyzed		Data analyzed		Data analyzed	
Sample size, column A	3	Sample size, column A	3	Sample size, column A	3
Sample size, column B	3	Sample size, column B	3	Sample size, column B	3
Table Analyzed	K	Table Analyzed	Fe	Table Analyzed	Zn
Column B	MeOH/EtOH	Column B	MeOH/EtOH	Column B	MeOH/EtOH
vs.	vs.	vs.	vs.	vs.	vs.
Column A	MeOH/H ₂ O	Column A	MeOH/H ₂ O	Column A	MeOH/H ₂ O
Unpaired t test		Unpaired t test		Unpaired t test	
P value	<0.0001	P value	0.0067	P value	0.0232
P value summary	****	P value summary	**	P value summary	*
Significantly different (P < 0.05)?	Yes	Significantly different (P < 0.05)?	Yes	Significantly different (P < 0.05)?	Yes
One- or two-tailed P value?	Two-tailed	One- or two-tailed P value?	Two-tailed	One- or two-tailed P value?	Two-tailed
t, df	t=21.57, df=4	t, df	t=5.163, df=4	t, df	t=3.577, df=4
How big is the difference?		How big is the difference?		How big is the difference?	
Mean of column A	0.4	Mean of column A	0.8067	Mean of column A	0.87
Mean of column B	1.027	Mean of column B	1.09	Mean of column B	1.107
Difference between means (B - A) ± SEM	0.6267 ± 0.02906	Difference between means (B - A) ± SEM	0.2833 ± 0.05487	Difference between means (B - A) ± SEM	0.2367 ± 0.06616
95% confidence interval	0.5460 to 0.7073	95% confidence interval	0.1310 to 0.4357	95% confidence interval	0.05296 to 0.4204
R squared (eta squared)	0.9915	R squared (eta squared)	0.8695	R squared (eta squared)	0.7618
F test to compare variances		F test to compare variances		F test to compare variances	
F, DFn, Dfd	7.444, 2, 2	F, DFn, Dfd	2.430, 2, 2	F, DFn, Dfd	5.254, 2, 2
P value	0.2368	P value	0.583	P value	0.3198
P value summary	ns	P value summary	ns	P value summary	ns
Significantly different (P < 0.05)?	No	Significantly different (P < 0.05)?	No	Significantly different (P < 0.05)?	No
Data analyzed		Data analyzed		Data analyzed	
Sample size, column A	3	Sample size, column A	3	Sample size, column A	3
Sample size, column B	3	Sample size, column B	3	Sample size, column B	3

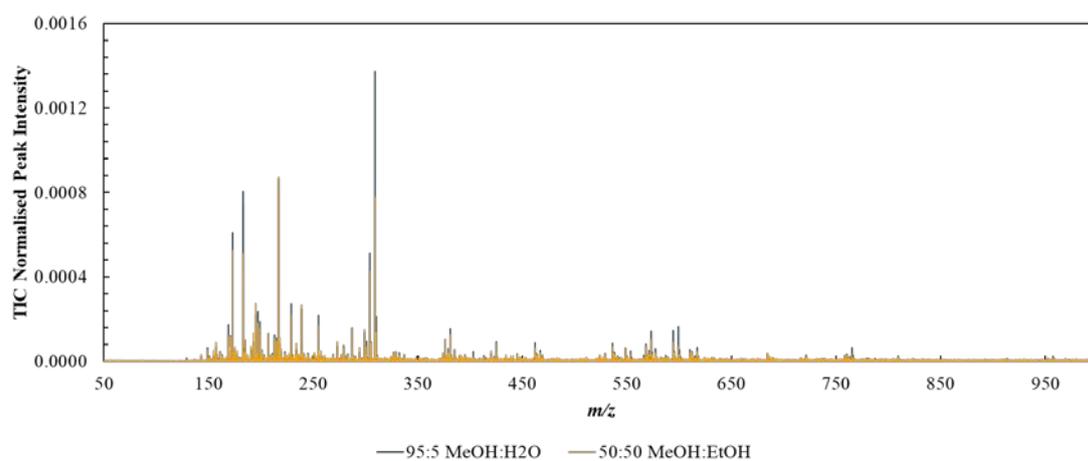


Figure S3. Spectra acquired with DESI using the two solvent systems 95:5 (%v/v) methanol/water (top) or 50:50 (%v/v) methanol/ethanol from the background (PET-only).

Table S2. Top 50 lipid features, adducts formed and measured *m/z* detected using each spray solvent system.

95:5 (%v/v) MeOH:H ₂ O			50:50 (%v/v) MeOH:EtOH		
Assignment	Adduct	Measured <i>m/z</i>	Assignment	Adduct	Measured <i>m/z</i>
PC 36:4	[M+K] ⁺	820.538	PC 36:4	[M+K] ⁺	820.538
PC 38:4	[M+K] ⁺	848.568	PC 38:4	[M ⁺ K] ⁺	848.568
PC 34:2	[M+K] ⁺	796.534	PC 38:7	[M ⁺ H] ⁺	804.562
PC 38:7	[M+H] ⁺	804.562	PC 40:7	[M ⁺ H] ⁺	832.596
PC 40:7	[M+H] ⁺	832.592	PC 34:2	[M+K] ⁺	796.54
PC 36:5	[M+H] ⁺	780.560	PC 36:4	[M ⁺ H] ⁺	782.58
PC 34:1	[M+K] ⁺	798.550	PA 42:3	[M ⁺ K] ⁺	821.542
PC 36:4	[M+H] ⁺	782.574	PC 34:1	[M ⁺ K] ⁺	798.552
PC 36:2	[M+K] ⁺	824.566	PI 32:0	[M ⁺ K] ⁺	849.576
PA 42:3	[M+K] ⁺	821.542	PC 36:2	[M+K] ⁺	824.572
PG 42:7	[M+H] ⁺	849.570	PC 38:5	[M ⁺ H] ⁺	808.596
PC 38:5	[M+H] ⁺	808.592	PC 36:3	[M ⁺ K] ⁺	822.55
PC 36:3	[M+K] ⁺	822.546	PA 40:1	[M ⁺ K] ⁺	797.544
PA 40:1	[M+K] ⁺	797.540	PC 38:5	[M+K] ⁺	846.554
PC 38:5	[M+K] ⁺	846.548	PC 38:6	[M ⁺ H] ⁺	806.578
FA 18:2	[M+K] ⁺	319.226	PC 38:4	[M ⁺ H] ⁺	810.612
PC 38:6	[M+H] ⁺	806.572	PC 38:5	[M ⁺ Na] ⁺	830.578
PC 38:6	[M+K] ⁺	844.536	PC 38:6	[M ⁺ K] ⁺	844.536
PC 38:4	[M+H] ⁺	810.606	PC 38:3	[M ⁺ K] ⁺	850.578
PA 42:1	[M+K] ⁺	825.572	PA 40:0	[M ⁺ Na] ⁺	783.582
PA 40:0	[M+Na] ⁺	783.582	PC 40:6	[M ⁺ H] ⁺	834.606
PC 38:3	[M+K] ⁺	850.574	PA 42:1	[M ⁺ K] ⁺	825.572
PG 38:4	[M+H] ⁺	799.554	PG 38:4	[M+H] ⁺	799.558
PC 38:5	[M+Na] ⁺	830.578	PG 40:5	[M+Na] ⁺	847.56
PC 38:6	[M+Na] ⁺	828.552	PC 36:1	[M ⁺ K] ⁺	826.578
LPC 16:0	[M+K] ⁺	534.302	FA 18:2	[M+K] ⁺	319.23
FA 20:5	[M+H ₃ O] ⁺	321.244	PG 40:6	[M ⁺ H] ⁺	823.55
PC 36:1	[M+K] ⁺	826.572	PG 40:6	[M ⁺ Na] ⁺	845.542
PC 34:2	[M+H] ⁺	758.578	PA 42:0	[M ⁺ Na] ⁺	811.616
PG 40:5	[M+Na] ⁺	847.556	LPC 18:3	[M ⁺ H] ⁺	518.33
PC 40:6	[M+H] ⁺	834.606	PC 36:3	[M ⁺ H] ⁺	784.592
PG 40:6	[M+H] ⁺	823.550	PC32:1	[M+K] ⁺	770.522
LPC 18:3	[M+H] ⁺	518.328	PC 36:2	[M ⁺ H] ⁺	786.612
PC 40:6	[M+K] ⁺	872.564	PE O-40:6	[M ⁺ Na] ⁺	800.556
PC 32:1	[M+K] ⁺	770.516	PC 34:1	[M ⁺ H] ⁺	760.592
LPC 20:4	[M+K] ⁺	582.302	FA 20:5	[M ⁺ H] ⁺	303.234
PG 40:6	[M+Na] ⁺	845.536	LPC 20:4	[M ⁺ Na] ⁺	566.33
PS 36:1	[M+Na] ⁺	812.542	LPC 20:3	[M ⁺ H] ⁺	546.362

LPC 18:0	[M+K] ⁺	562.334
PC 42:11	[M+H] ⁺	852.538
PA 42:0	[M+Na] ⁺	811.610
PC 40:6	[M+Na] ⁺	856.584
PS 38:3	[M+Na] ⁺	836.536
LPC 20:4	[M+Na] ⁺	566.330
PC 36:3	[M+H] ⁺	784.590
PC 42:8	[M+K] ⁺	896.558
PE O-40:6	[M+Na] ⁺	800.554
PC 34:1	[M+H] ⁺	760.592
LPC 20:3	[M+H] ⁺	546.360
PC 36:2	[M+H] ⁺	786.610

PC 34:5	[M+K] ⁺	790.548
PG 42:6	[M ⁺ H] ⁺	851.582
LPC 20:4	[M ⁺ K] ⁺	582.304
FA 18:1	[M+K] ⁺	321.244
LPC 18:0	[M ⁺ K] ⁺	562.334
PI 34:2	[M+K] ⁺	873.574
PE 38:2	[M ⁺ H] ⁺	772.538
PC 40:5	[M ⁺ K] ⁺	874.58
FA 22:6	[M ⁺ K] ⁺	367.204
FA 20:4	[M ⁺ Na] ⁺	327.234
FA 18:2	[M ⁺ K] ⁺	319.208
PE 38:1	[M ⁺ K] ⁺	812.548

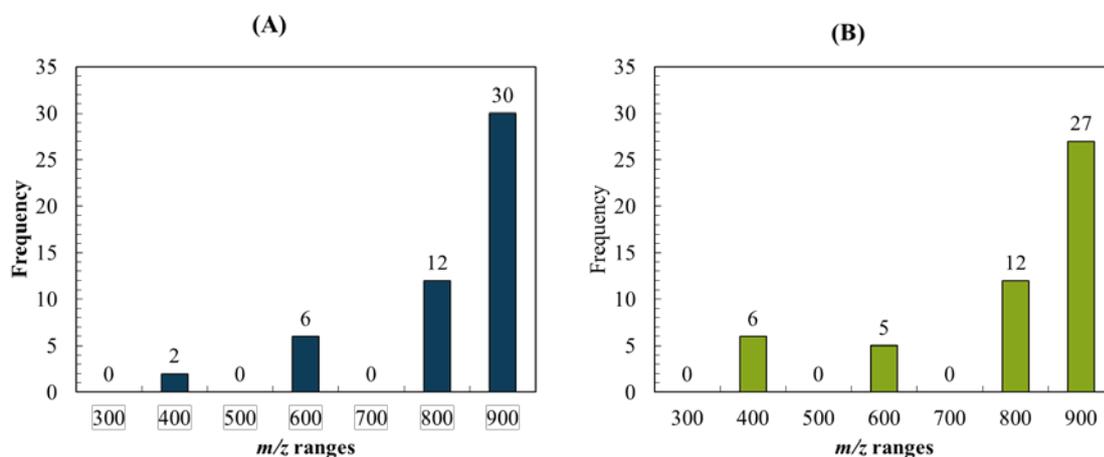


Figure S4. Histogram of the number of lipid features in the top 50 detected per m/z range (bin size 100) for each spray solvent – (A) MeOH/H₂O and (B) MeOH/EtOH.

Table S3. T-test results comparing the measured normalised (to TIC) peak intensities for the top 10 lipids measured using the two solvent systems.

	Discovery?	P value	Mean of 95:5 (%v/v) MeOH:H ₂ O	Mean of 50:50 (%v/v/ MeOH:EtOH	Difference	SE of difference	t ratio	df	q value
PC 36:4 [M+K] ⁺	Yes	<0.000001	0.01117	0.01398	-0.002816	0.0003404	8.27	16	<0.000001
PC 38:4 [M+K] ⁺	Yes	<0.000001	0.01014	0.01349	-0.003351	0.0003131	10.7	16	<0.000001
PC 34:2 [M+K] ⁺	Yes	0.000786	0.009648	0.01054	-0.0008878	0.000215	4.13	16	0.000893
PC 38:7 [M+H] ⁺	Yes	<0.000001	0.007512	0.01219	-0.004677	0.0002612	17.91	16	<0.000001
PC 40:7 [M+H] ⁺	Yes	<0.000001	0.006606	0.0115	-0.004899	0.0002348	20.86	16	<0.000001
PC 34:1 [M+K] ⁺	Yes	0.006748	0.007	0.007458	-0.0004578	0.0001472	3.109	16	0.006816
PC 36:4 [M+H] ⁺	Yes	<0.000001	0.006516	0.008541	-0.002026	0.0001099	18.44	16	<0.000001
PC 36:2 [M+K] ⁺	Yes	0.000004	0.006268	0.007224	-0.0009567	0.000139	6.884	16	0.000005
PA 42:3 [M+K] ⁺	Yes	0.000001	0.00551	0.006727	-0.001217	0.0001635	7.439	16	0.000002

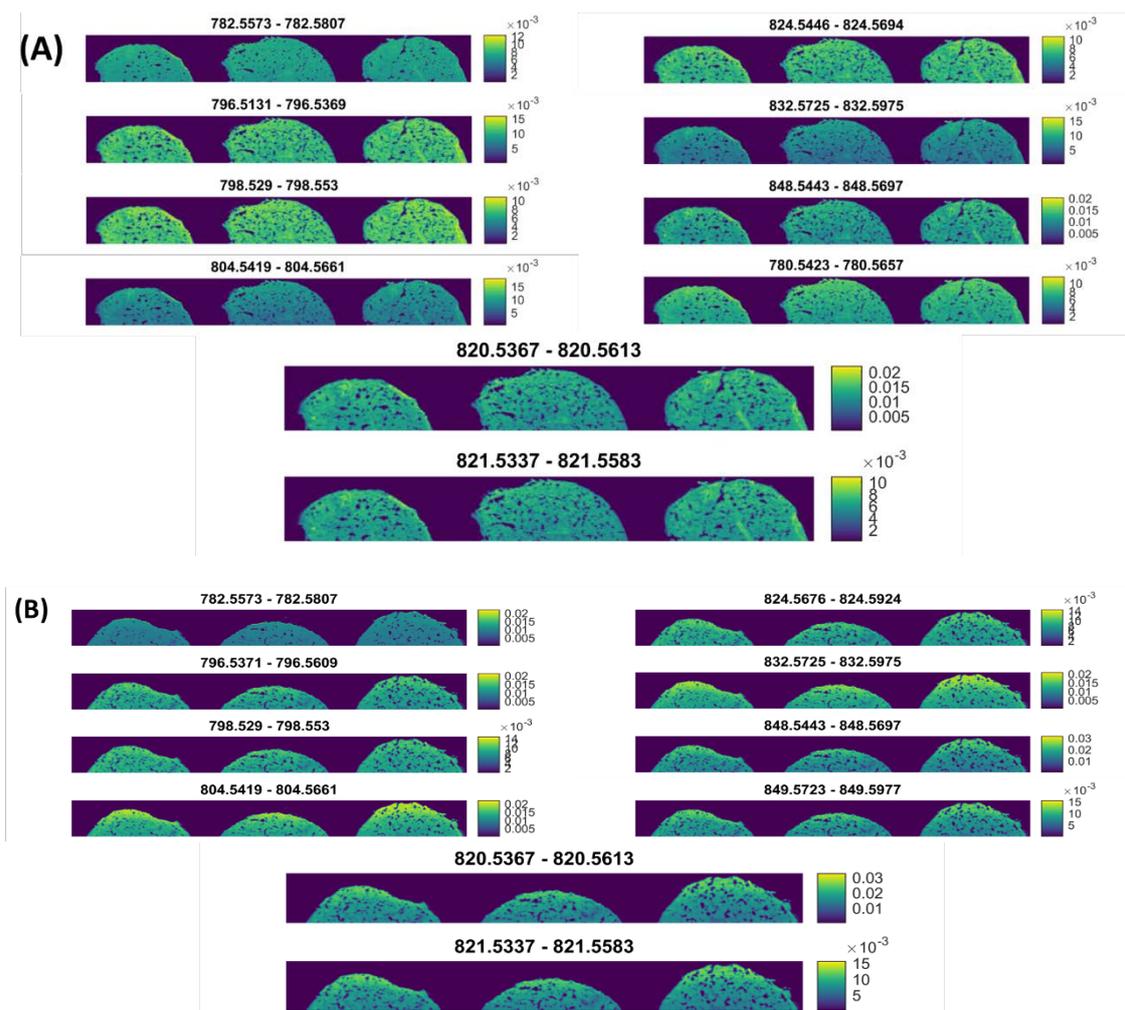


Figure S5. Resulting DESI ion maps for the top 10 most abundant lipids imaged using (A) 95:5 (%v/v) MeOH:H₂O or (B) 50:50 (%v/v) MeOH:EtOH spray solvents on liver tissue homogenates.

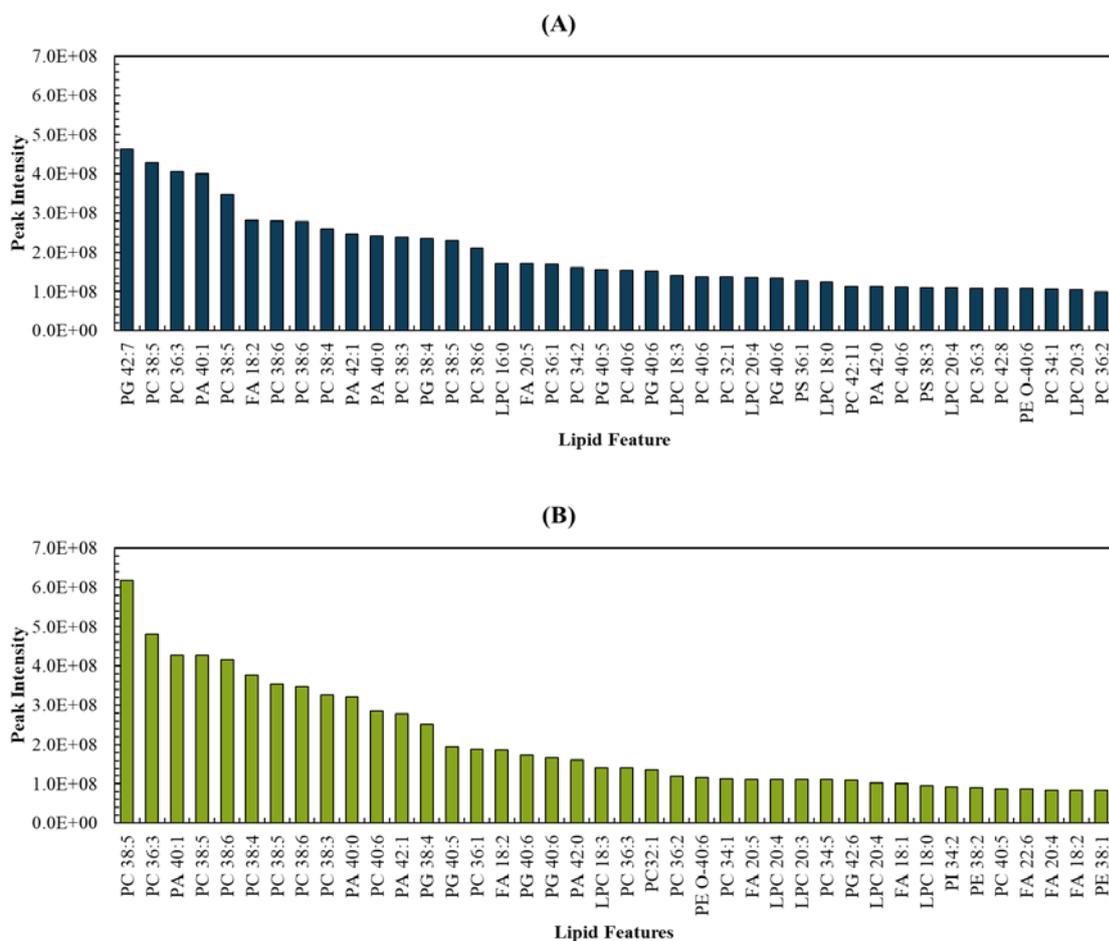


Figure S6. Remaining 40 features in the top 50 most intense lipid features and their respective intensities for (A) MeOH:H₂O and (B) MeOH:EtOH.

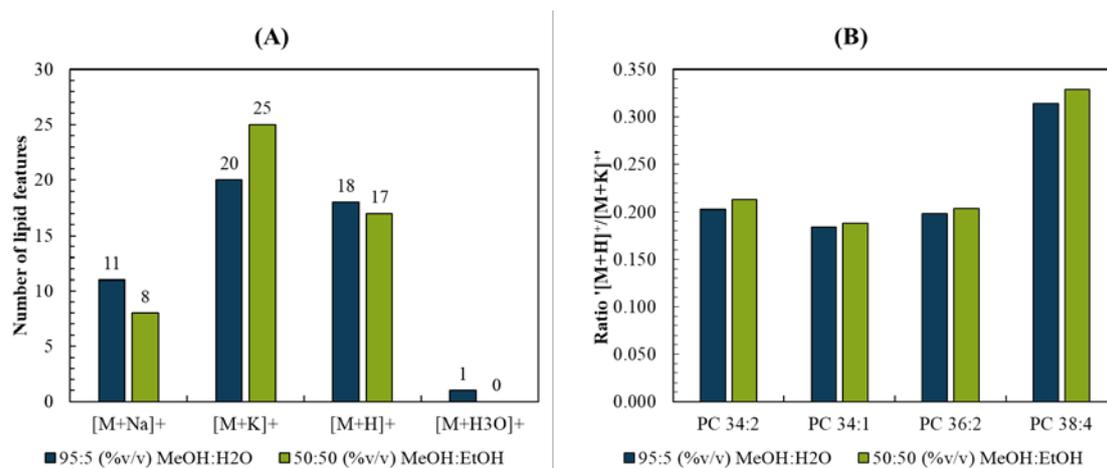


Figure S7. (A) Number of lipid features detected in the top 50 lipids with a specific adduct for each of the two solvent mixtures – 95:5 (%v/v) MeOH:H₂O (blue) and 50:50 (%v/v) MeOH:EtOH (green). (B) Ratio of average (n=9) peak intensities measured for [M+H]⁺ and [M+K]⁺ ions formed for a selection of PC lipids using the two solvent mixtures – 95:5 (%v/v) MeOH:H₂O (blue) and 50:50 (%v/v) MeOH:EtOH (green).

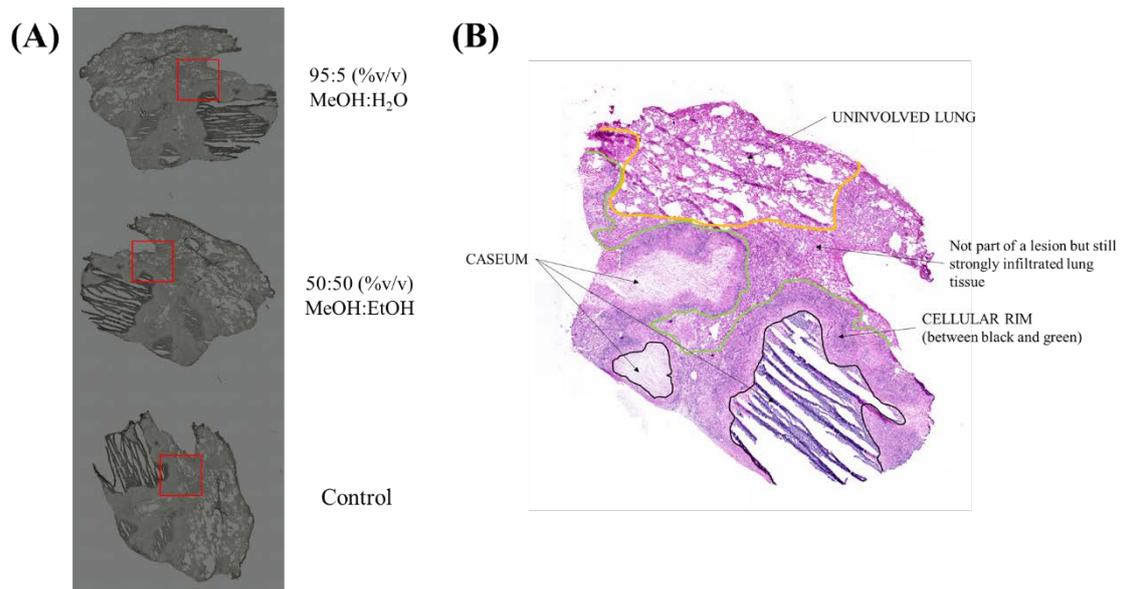


Figure S8. (A) Optical images of sequential fresh frozen rabbit lung tissues containing a caseous granuloma (a lesion caused by tuberculosis) showing the regions analysed using ion beam analysis; (B) Hematoxylin and eosin (H&E) stained section and the different regions of the granuloma.

Table S4. T-test results of the elemental peak area measured by PIXE from a lung tissue section after DESI analysis with two different solvents – MeOH/H₂O and MeOH/EtOH.

Table Analyzed	Phosphorus	Table Analyzed	Sulphur	Table Analyzed	Chlorine	Table Analyzed	Potassium	Table Analyzed	Iron
Column B	MeOH:H2O	Column B	MeOH:H2O	Column B	MeOH:H2O	Column B	MeOH:H2O	Column B	MeOH:H2O
vs.	vs.	vs.	vs.	vs.	vs.	vs.	vs.	vs.	vs.
Column A	Control	Column A	Control	Column A	Control	Column A	Control	Column A	Control
Unpaired t test		Unpaired t test		Unpaired t test		Unpaired t test		Unpaired t test	
P value	0.0042	P value	0.0641	P value	<0.0001	P value	<0.0001	P value	0.354
P value summary	**	P value summary	ns	P value summary	****	P value summary	****	P value summary	ns
Significantly different (P < 0.05)?	Yes	Significantly different (P < 0.05)?	No	Significantly different (P < 0.05)?	Yes	Significantly different (P < 0.05)?	Yes	Significantly different (P < 0.05)?	No
One- or two-tailed P value?	Two-tailed	One- or two-tailed P value?	Two-tailed	One- or two-tailed P value?	Two-tailed	One- or two-tailed P value?	Two-tailed	One- or two-tailed P value?	Two-tailed
t, df	t=4.484, df=6	t, df	t=2.265, df=6	t, df	t=9.781, df=6	t, df	t=11.11, df=6	t, df	t=1.004, df=6
How big is the difference?		How big is the difference?		How big is the difference?		How big is the difference?		How big is the difference?	
Mean of column A	169825	Mean of column A	166966	Mean of column A	239292	Mean of column A	327173	Mean of column A	2756
Mean of column B	103629	Mean of column B	131636	Mean of column B	20466	Mean of column B	90625	Mean of column B	2233
Difference between means (B - A) ± SEM	-66197 ± 14763	Difference between means (B - A) ± SEM	-35330 ± 15598	Difference between means (B - A) ± SEM	-218826 ± 22373	Difference between means (B - A) ± SEM	-236548 ± 21288	Difference between means (B - A) ± SEM	-523.1 ± 520.8
95% confidence interval	-102321 to -30073	95% confidence interval	-73496 to 2836	95% confidence interval	-273572 to -164081	95% confidence interval	-288637 to -184459	95% confidence interval	-1797 to 751.2
R squared (eta squared)	0.7702	R squared (eta squared)	0.4609	R squared (eta squared)	0.941	R squared (eta squared)	0.9537	R squared (eta squared)	0.1439
F test to compare variances		F test to compare variances		F test to compare variances		F test to compare variances		F test to compare variances	
F, DFn, Dfd	1.472, 3, 3	F, DFn, Dfd	2.191, 3, 3	F, DFn, Dfd	737.9, 3, 3	F, DFn, Dfd	7.637, 3, 3	F, DFn, Dfd	13.91, 3, 3
P value	0.7585	P value	0.536	P value	0.0002	P value	0.129	P value	0.0578
P value summary	ns	P value summary	ns	P value summary	***	P value summary	ns	P value summary	ns
Significantly different (P < 0.05)?	No	Significantly different (P < 0.05)?	No	Significantly different (P < 0.05)?	Yes	Significantly different (P < 0.05)?	No	Significantly different (P < 0.05)?	No
Data analyzed		Data analyzed		Data analyzed		Data analyzed		Data analyzed	
Sample size, column A	4	Sample size, column A	4	Sample size, column A	4	Sample size, column A	4	Sample size, column A	4
Sample size, column B	4	Sample size, column B	4	Sample size, column B	4	Sample size, column B	4	Sample size, column B	4
Table Analyzed	Phosphorus	Table Analyzed	Sulphur	Table Analyzed	Chlorine	Table Analyzed	Potassium	Table Analyzed	Iron
Column C	MeOH:EtOH	Column C	MeOH:EtOH	Column C	MeOH:EtOH	Column C	MeOH:EtOH	Column C	MeOH:EtOH
vs.	vs.	vs.	vs.	vs.	vs.	vs.	vs.	vs.	vs.
Column A	Control	Column A	Control	Column A	Control	Column A	Control	Column A	Control
Unpaired t test		Unpaired t test		Unpaired t test		Unpaired t test		Unpaired t test	
P value	0.0556	P value	0.3721	P value	0.2547	P value	0.1484	P value	0.3264
P value summary	ns	P value summary	ns	P value summary	ns	P value summary	ns	P value summary	ns
Significantly different (P < 0.05)?	No	Significantly different (P < 0.05)?	No	Significantly different (P < 0.05)?	No	Significantly different (P < 0.05)?	No	Significantly different (P < 0.05)?	No
One- or two-tailed P value?	Two-tailed	One- or two-tailed P value?	Two-tailed	One- or two-tailed P value?	Two-tailed	One- or two-tailed P value?	Two-tailed	One- or two-tailed P value?	Two-tailed
t, df	t=2.369, df=6	t, df	t=0.9643, df=6	t, df	t=1.259, df=6	t, df	t=1.658, df=6	t, df	t=1.068, df=6
How big is the difference?		How big is the difference?		How big is the difference?		How big is the difference?		How big is the difference?	
Mean of column A	169825	Mean of column A	166966	Mean of column A	239292	Mean of column A	327173	Mean of column A	2756
Mean of column C	121313	Mean of column C	146604	Mean of column C	195866	Mean of column C	259388	Mean of column C	2127
Difference between means (C - A) ± SEM	-48512 ± 20475	Difference between means (C - A) ± SEM	-20362 ± 21115	Difference between means (C - A) ± SEM	-43427 ± 34487	Difference between means (C - A) ± SEM	-67785 ± 40881	Difference between means (C - A) ± SEM	-629.8 ± 589.4
95% confidence interval	-98613 to 1588	95% confidence interval	-72029 to 31306	95% confidence interval	-127814 to 40961	95% confidence interval	-167818 to 32248	95% confidence interval	-2072 to 812.5
R squared (eta squared)	0.4834	R squared (eta squared)	0.1342	R squared (eta squared)	0.209	R squared (eta squared)	0.3142	R squared (eta squared)	0.1599
F test to compare variances		F test to compare variances		F test to compare variances		F test to compare variances		F test to compare variances	
F, DFn, Dfd	2.231, 3, 3	F, DFn, Dfd	4.848, 3, 3	F, DFn, Dfd	1.379, 3, 3	F, DFn, Dfd	3.171, 3, 3	F, DFn, Dfd	2.680, 3, 3
P value	0.527	P value	0.2273	P value	0.7979	P value	0.3686	P value	0.4396
P value summary	ns	P value summary	ns	P value summary	ns	P value summary	ns	P value summary	ns
Significantly different (P < 0.05)?	No	Significantly different (P < 0.05)?	No	Significantly different (P < 0.05)?	No	Significantly different (P < 0.05)?	No	Significantly different (P < 0.05)?	No
Data analyzed		Data analyzed		Data analyzed		Data analyzed		Data analyzed	
Sample size, column A	4	Sample size, column A	4	Sample size, column A	4	Sample size, column A	4	Sample size, column A	4
Sample size, column C	4	Sample size, column C	4	Sample size, column C	4	Sample size, column C	4	Sample size, column C	4

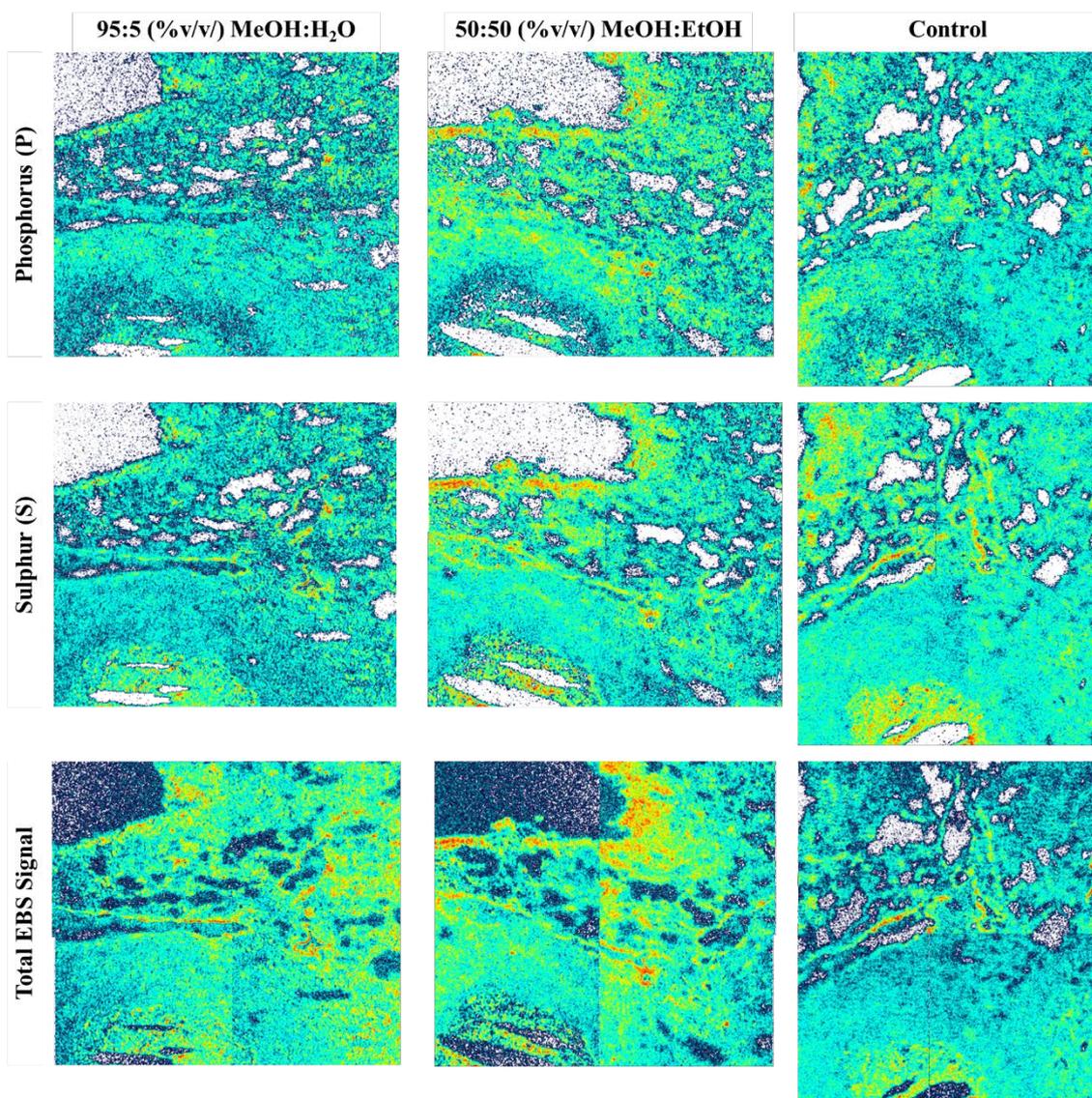


Figure S9. Phosphorus (P), sulphur (S) PIXE maps and total EBS maps taken from fresh frozen lung tissue sections after DESI analysis using 95:5 (%v/v) MeOH:H₂O or 50:50 (%v/v) MeOH:EtOH. A third section (Control) was also analysed – no DESI measurements were taken on this sample.

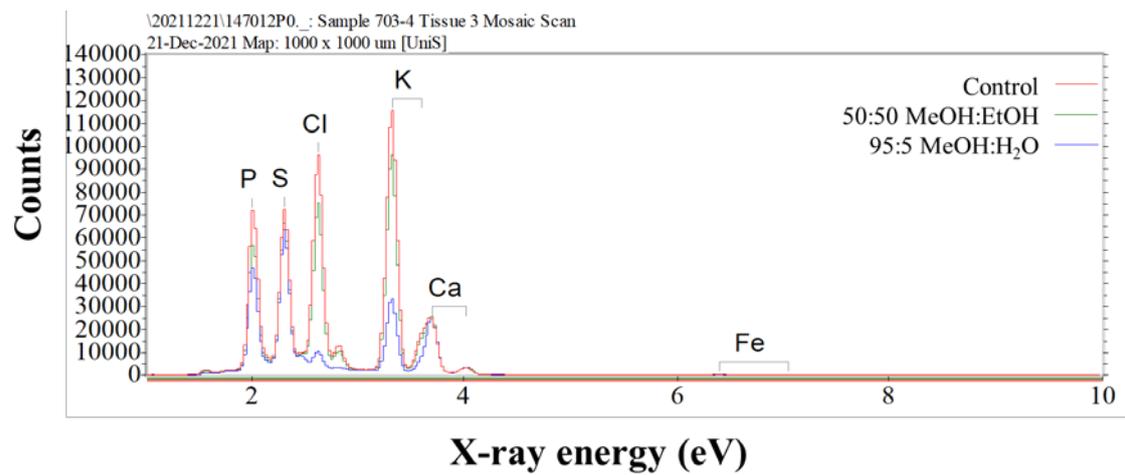


Figure S10. Overlay of the X-ray spectra taken from the squares containing the caseum in each of the lung tissue section samples.