

Supporting Information

LC-MS-based metabolomics reveals metabolic signatures related to glioma stem-like cell self-renewal and differentiation

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Fig. S1. Flowchart of the LC-MS-based metabolomics approach.

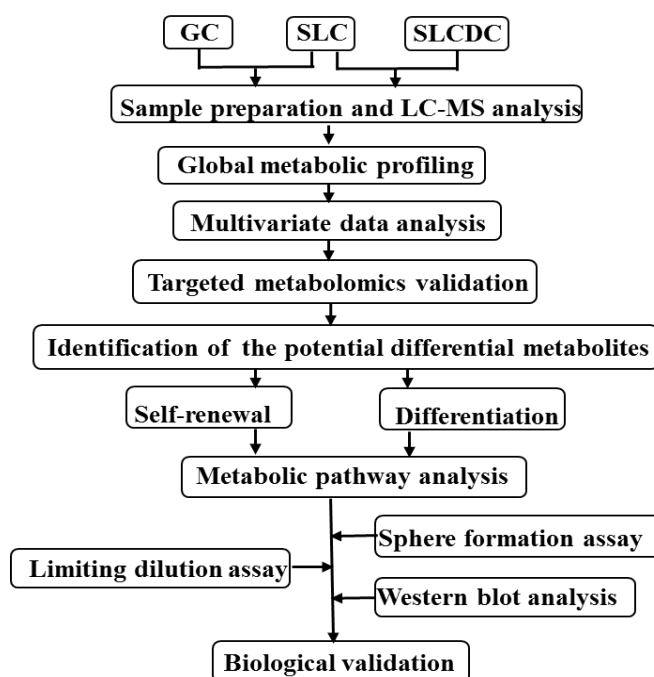
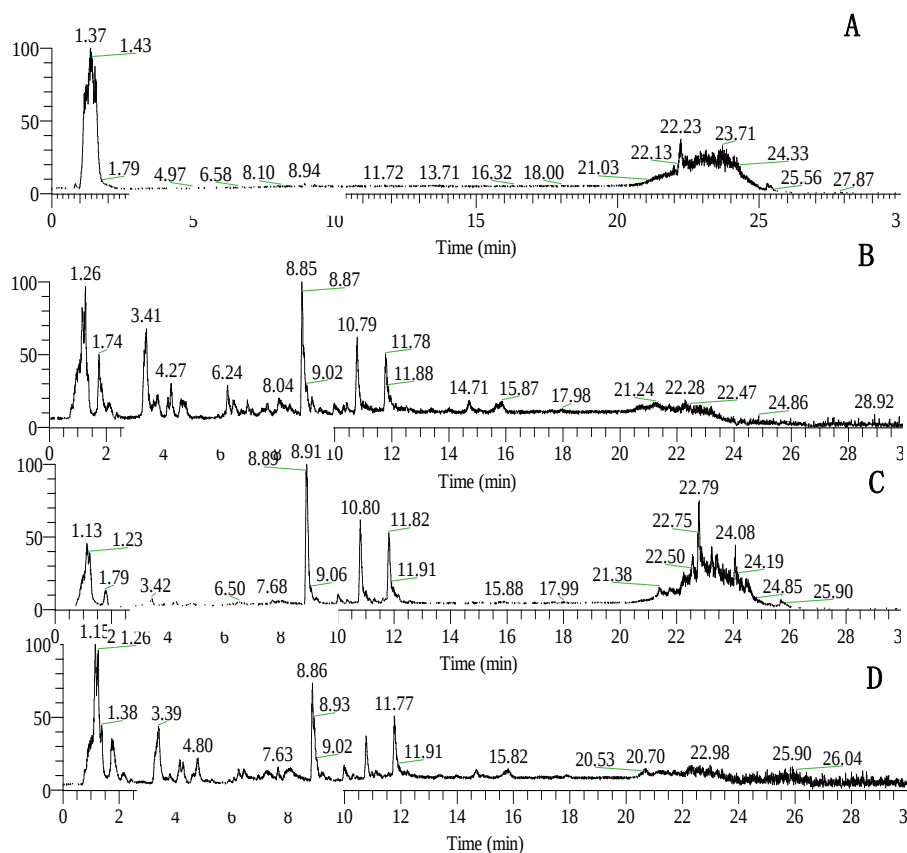


Fig. S2 The total ion chromatograms (TICs) of different cell samples. (A) Blank samples, (B) GC, (C) SLC, (D) SLDC.

LC-(+) ESI/MS



LC-(-) ESI/MS

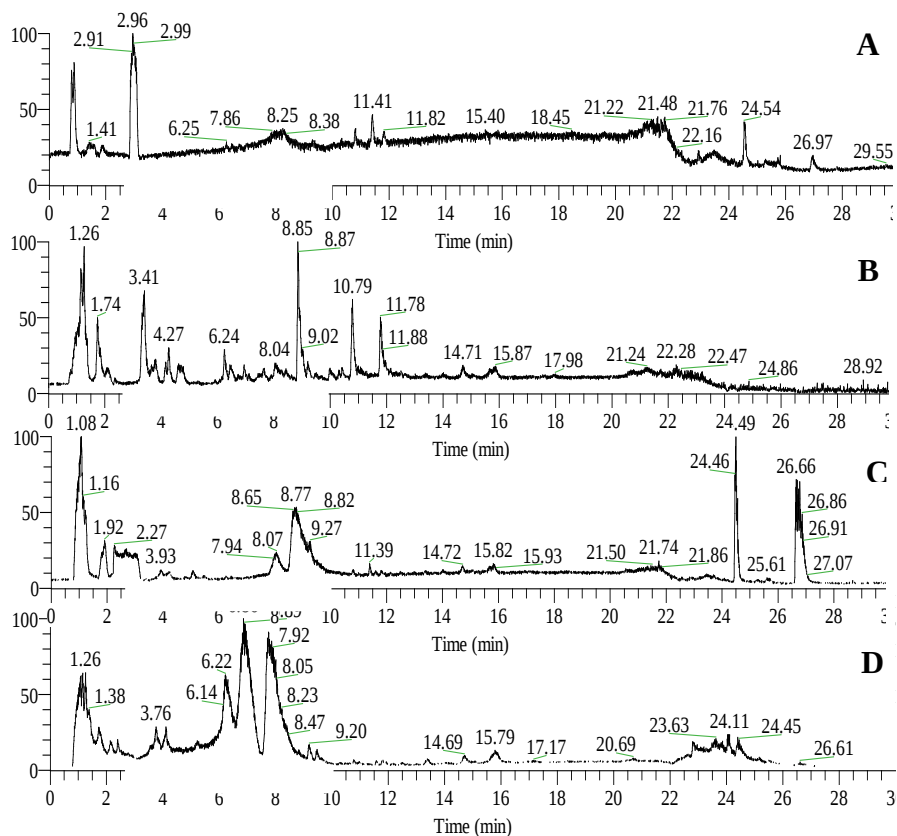


Fig. S3 The PCA score plots (95% confidence interval) of QC samples deriving from LC-(±) ESI-MS data.

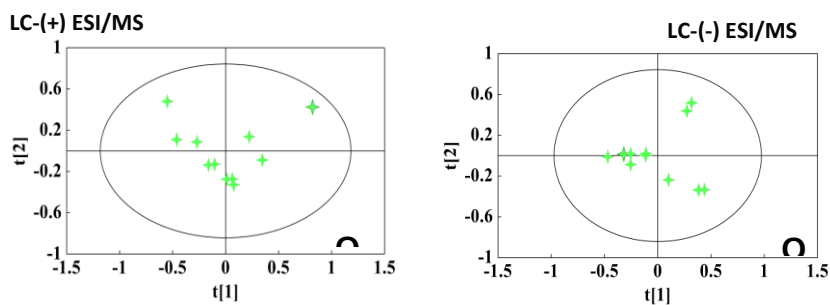


Fig. S4 Line plots of QC samples generated by PCA using component 1 and 2.

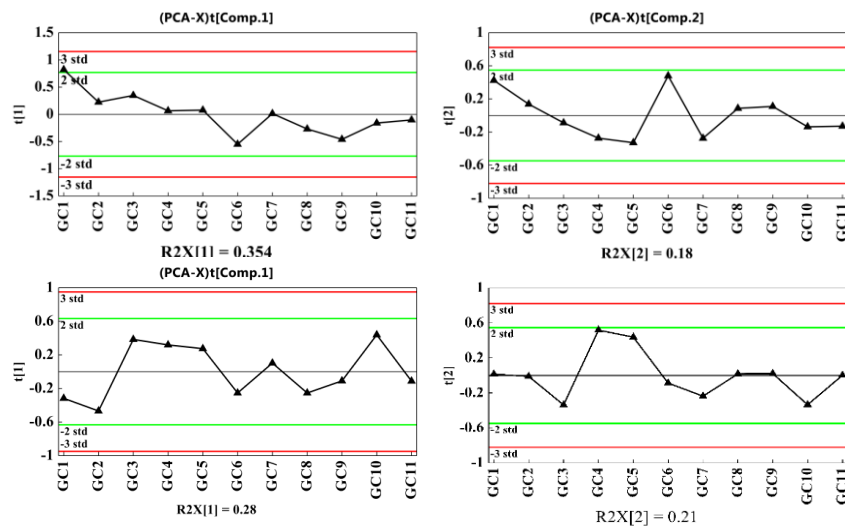


Fig. S5 The retention time deviation profiles deriving from LC-(±) ESI-MS.

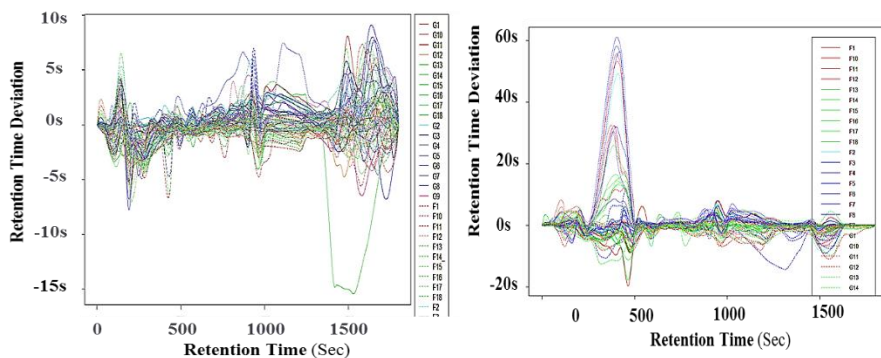


Fig. S6 Score plots of the OPLS-DA models (A, B) and plots of the permutation tests of the PLS-DA models (C,D) based on the LC-(-) ESI-MS data sets. (A, C) GCs vs SLCs, (B, D) SLCDs vs SLCs, (green ●) GCs, (red ▲) SLCs, (purple ■) SLCDs.

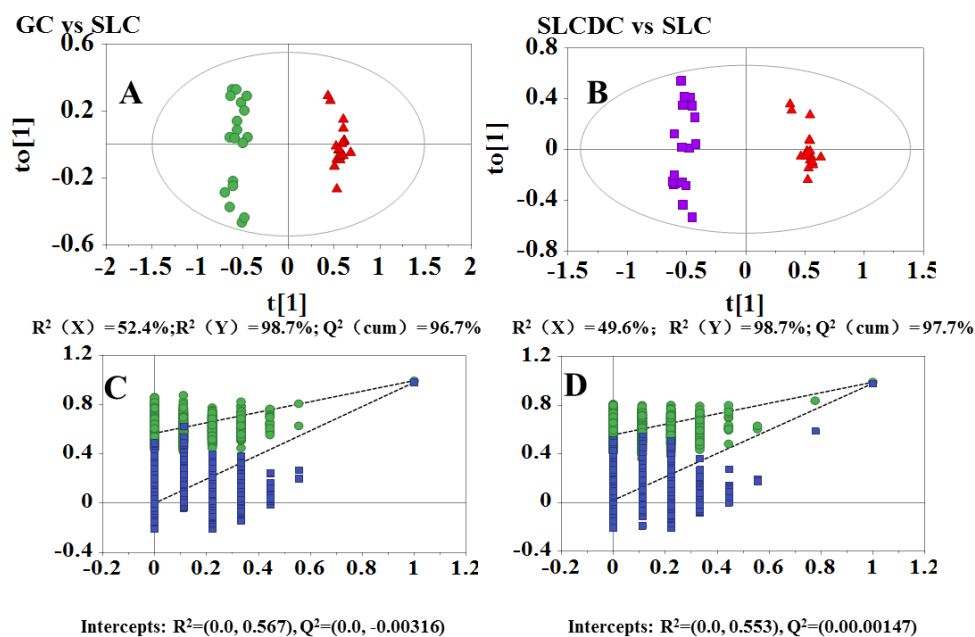


Fig. S7 The XICs of differential variables of the LC-MS/MS MRM analysis in positive ion mode (A) and negative ion mode (B).

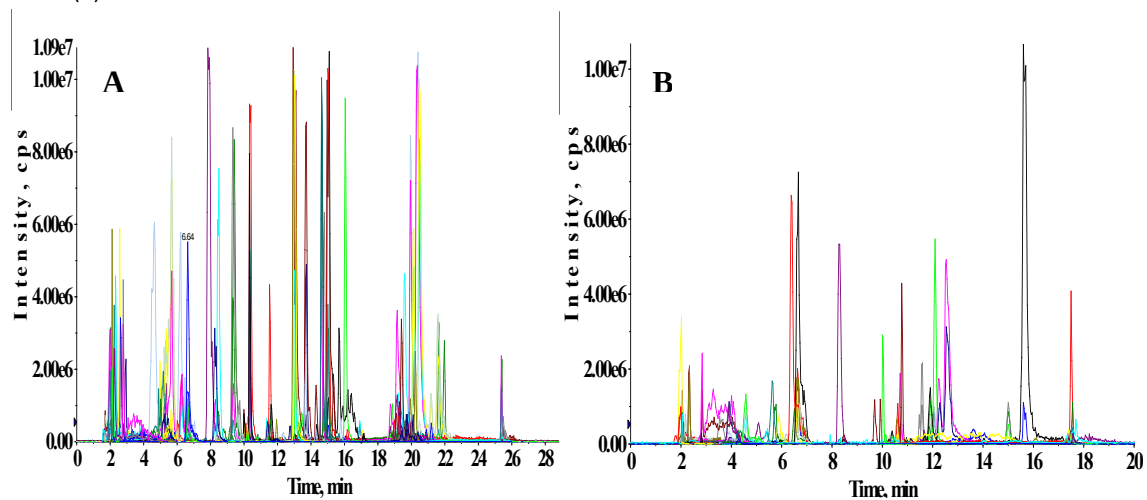
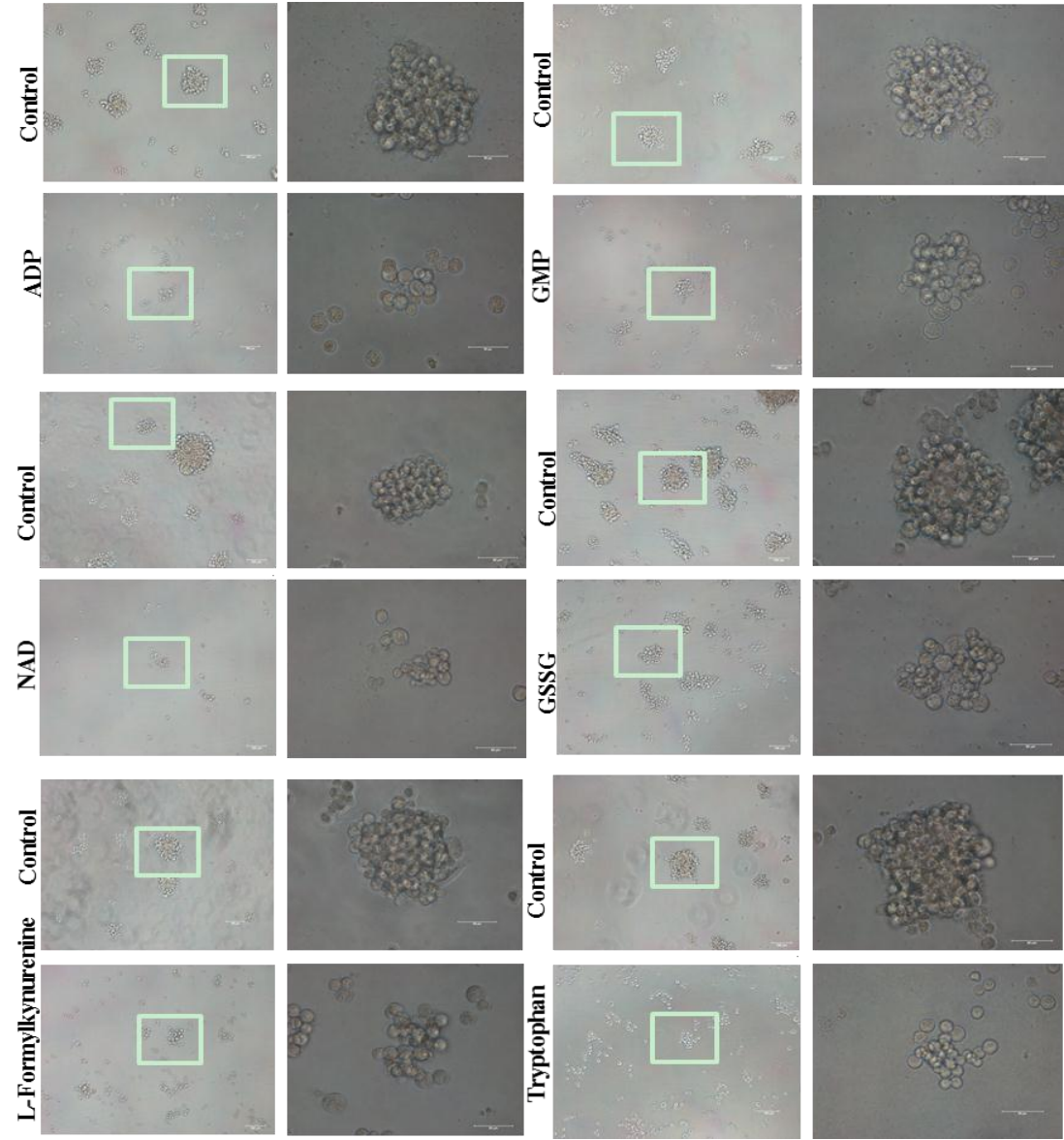


Fig. S10 Neurosphere formation assay in SLCs. Phase contrast photomicrographs of SLC treated with metabolites (Bar = 100 μ m, left; Bar = 50 μ m, right)



Sphere formation assay of SLC treated with metabolites, neurospheres with diameters larger than 50 μm were counted. The values shown are the means $\pm$ SD of at least 3 independent experiments.

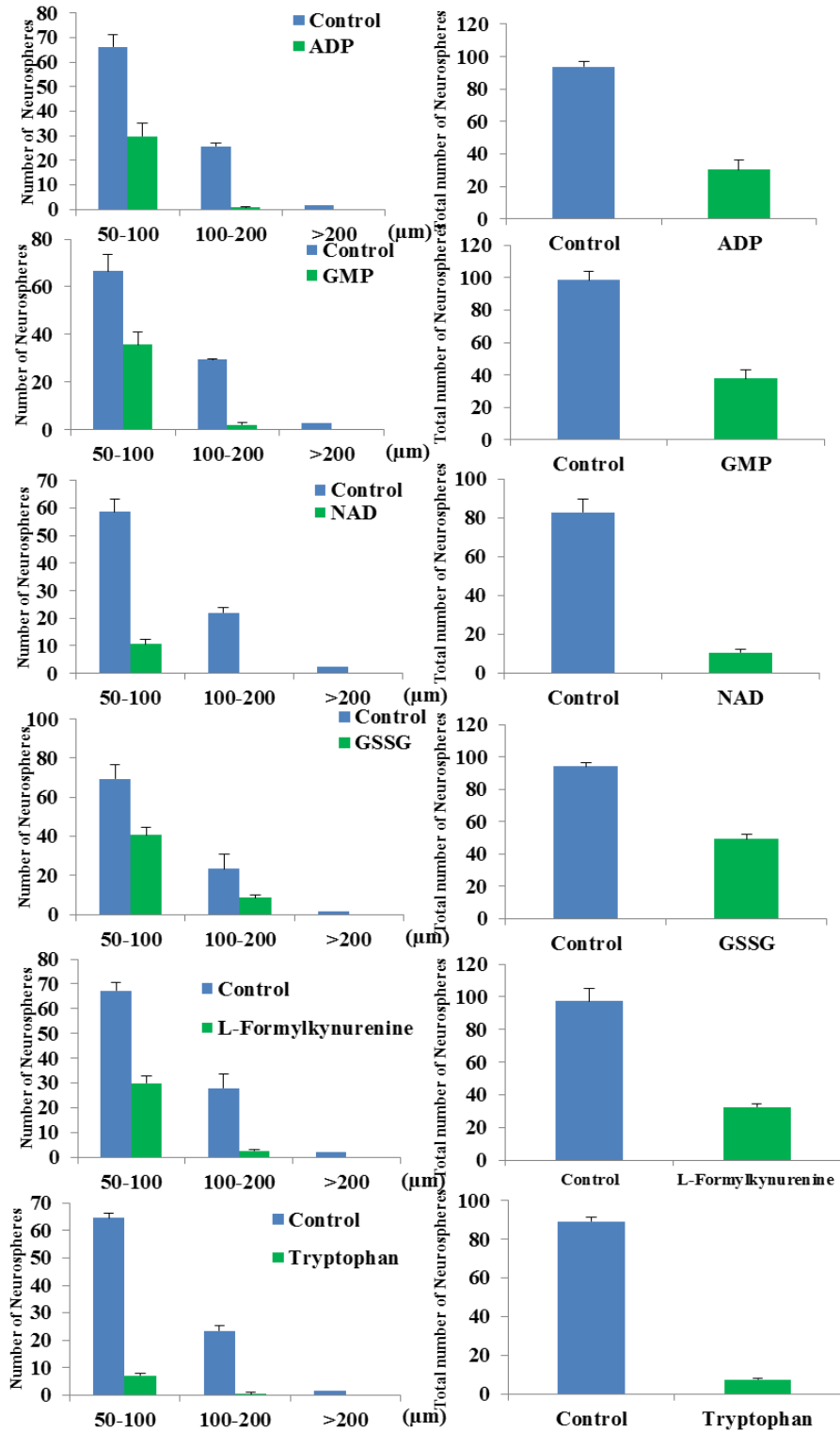


Fig. S11 Limiting dilution neurosphere assay in SLCs.

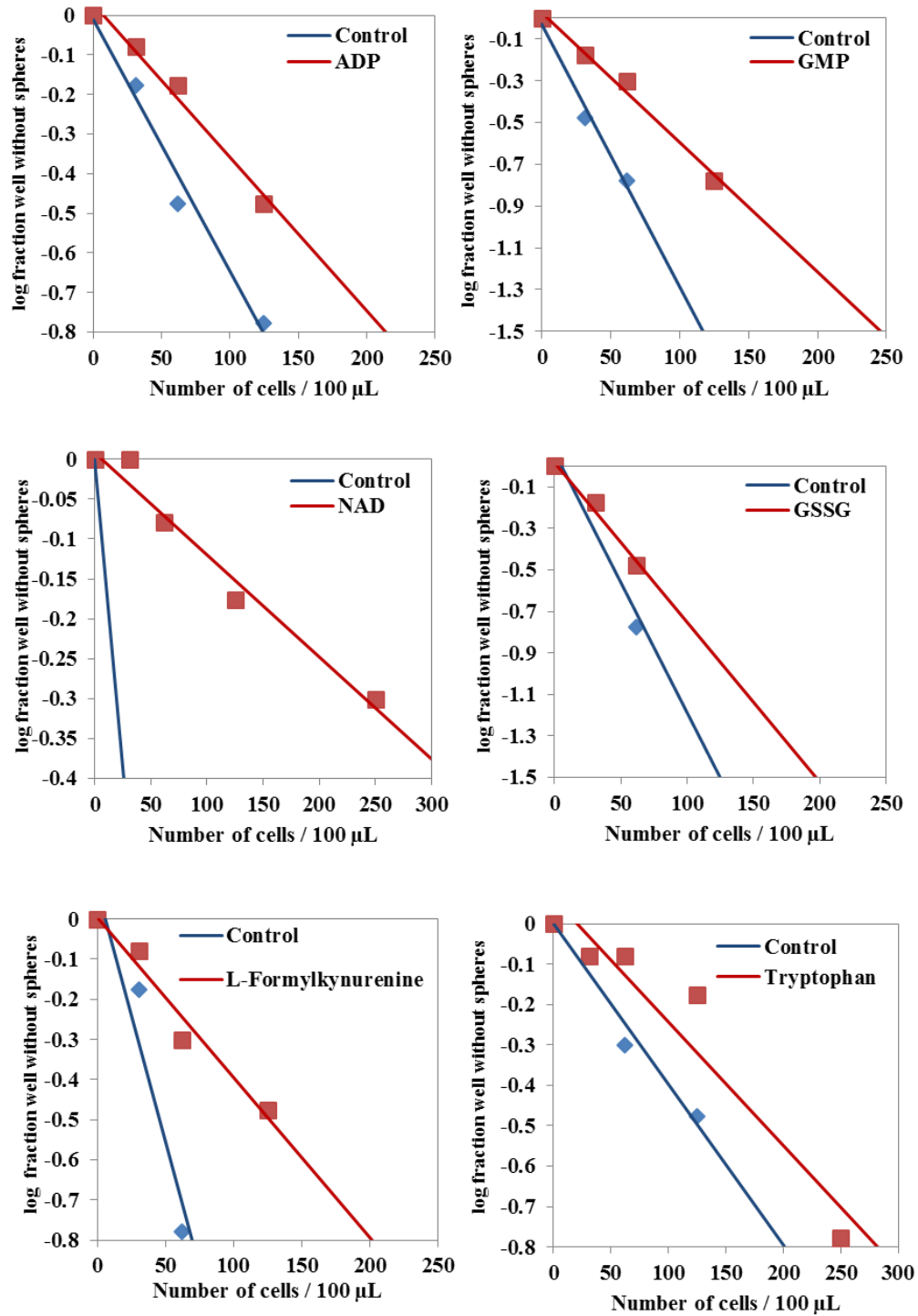


Table S1. The differential metabolites between GC and SLC group obtained by LC-(±) MS/MS analysis.

RT	m/z	Adduct ion	Elemental composition	Metabolite identification ^a	Product ions	P value	Fold Change	VIP
4.8	114.0916	[M+H-H ₂ O] ⁺	C ₆ H ₁₃ NO ₂	L-Leucine ^a	96.1, 105, 79.1, 69.1, 55.1	< 0.001	0.59	6.57
1.26	116.0709	[M+H] ⁺	C ₅ H ₉ NO ₂	L-Proline ^a	70.1, 68.05	< 0.001	1.28	3.35
1.25	118.0865	[M+H] ⁺	C ₅ H ₁₁ NO ₂	Valine ^a	72.1, 55.1	< 0.001	0.8	2.96
3.4	166.0861	[M+H] ⁺	C ₉ H ₁₁ NO ₂	Phenylalanine ^a	120.1, 103.1, 107.0, 131, 149.1	< 0.001	2.34	15.7
1.26	182.0812	[M+H] ⁺	C ₉ H ₁₁ NO ₃	L-Tyrosine ^a	136.1, 391.1, 119.0, 123.0, 147.0, 165.1	< 0.001	0.38	3.33
4.28	205.0971	[M+H] ⁺	C ₁₁ H ₁₂ N ₂ O ₂	L-Tryptophan ^a	188.1, 146.1, 159.1, 144.1, 118.1, 115.1	< 0.001	3.29	7.36
1.27	132.0769	[M+H] ⁺	C ₄ H ₉ N ₃ O ₂	Creatine ^a	90.1, 114.1, 115.1, 86.1, 87.1	< 0.001	6.74	7.76
4.16	220.1179	[M+H] ⁺	C ₉ H ₁₇ NO ₅	Pantothenic Acid ^a	90.1, 485.1, 124.1, 142.1, 160.1, 184.1, 202.1	< 0.001	1.63	3.4
2.38	130.0500	[M+H] ⁺	C ₅ H ₇ NO ₃	Pyrrolidonecarboxylic acid ^a	84.0, 56.1, 102.1	< 0.001	7.25	4.93
1.75	148.0426	[M+H-H ₂ O] ⁺	C ₅ H ₁₁ NO ₃ S	Methionine sulfoxide ^a	84.0, 102.1, 130.0, 56.1, 74.0	< 0.001	10.64	5.62
4.28	188.0706	[M+H] ⁺	C ₁₁ H ₉ NO ₂	Indoleacrylic acid ^a	146.1, 115.1, 170.1	< 0.001	3.2	8.44
3.4	131.0492	[M+H-H ₂ O] ⁺	C ₉ H ₈ O ₂	Trans-Cinnamic acid ^a	103.1, 77.0, 131.0	< 0.001	2.44	2.04
1.93	165.0546	[M+H] ⁺	C ₉ H ₈ O ₃	2-Hydroxycinnamic acid ^a	123.0, 91.1, 95.0, 103.1, 147.0	< 0.001	0.64	2.2
2.45	611.1441	[M-H] ⁻	C ₂₀ H ₃₂ N ₆ O ₁₂ S ₂	Oxidized glutathione ^a	306.1, 338.0, 272.1, 254.1, 160.0, 482.1, 593.1	< 0.001	14.65	2.99
0.86	188.1756	[M+H] ⁺	C ₉ H ₂₁ N ₃ O	N1-Acetylspermidine	72.1, 171.1, 117.1, 100.1, 58.1	< 0.001	0.02	13.89
0.85	245.2333	[M+H] ⁺	C ₁₂ H ₂₈ N ₄ O	N1-Acetylspermine ^a	100.1, 112.1, 129.1, 171.1, 84.1, 70.1	< 0.001	0.03	3.19
7.22	204.0865	[M+H-H ₂ O] ⁺	C ₈ H ₁₅ NO ₆	N-Acetyl-D-glucosamine ^a	96.0, 109.0, 126.1, 138.1, 144.1, 168.1, 186.1	< 0.001	5.18	5.65

Table S1. Continued.

RT	m/z	Adduct ion	Elemental composition	Metabolite identification ^a	Product ions	P value	Fold Change	VIP
4.1	246.1698	[M+H] ⁺	C ₁₂ H ₂₃ NO ₄	Pentanoylcarnitine	187.1, 144.1, 60.1, 85.0	< 0.001	20.07	3.94
11.02	398.3261	[M+H] ⁺	C ₂₃ H ₄₃ NO ₄	trans-2-Hexadecenoyl-carnitine	85.0, 339.3, 237.2, 144.1, 95.1, 60.1	< 0.001	9.51	2.5
11.95	400.3417	[M+H] ⁺	C ₂₃ H ₄₅ NO ₄	L-Palmitoylcarnitine	85.0, 341.3, 239.2, 144.1, 95.1, 60.1	< 0.001	13.59	4.18
12.37	426.3573	[M+H] ⁺	C ₂₅ H ₄₇ NO ₄	Oleoylcarnitine	85.0, 367.3, 265.3, 60.1, 144.1, 95.1	< 0.001	12.35	5.3

1.28	258.1099	[M+H] ⁺	C ₈ H ₂₀ NO ₆ P	Glycerophosphocholine	104.1, 60.1, 86.1, 125.0, 184.1	< 0.001	2.78	3.34
1.16	104.1074	[M+H] ⁺	C ₅ H ₁₃ NO	Choline ^a	60.1, 58.1, 59.1	< 0.001	4.14	10.31
11.91	468.3080	[M+H] ⁺	C ₂₂ H ₄₆ NO ₇ P	LysoPC(14:0)	184.1, 450.3, 125.0, 104.1	0.0158	2.9	4.04
13.03	482.3238	[M+H] ⁺	C ₂₃ H ₄₈ NO ₇ P	LPC(15:0) ^a	184.1, 464.3, 125.0, 104.1, 86.1	< 0.001	21.13	2.95
21.28	550.3863	[M+H] ⁺	C ₂₈ H ₅₆ NO ₇ P	LPC(20:1)	184.1, 532.4, 104.1, 86.1, 125.0	< 0.001	2.16	2.53
13.35	568.3391	[M+H] ⁺	C ₃₀ H ₅₀ NO ₇ P	LPC(22:6)	184.1, 550.4, 104.1, 125.0	< 0.001	35.53	2.54
15.67	480.3081	[M+H] ⁺	C ₂₃ H ₄₆ NO ₇ P	LPE(18:1) ^a	69.1, 95.1, 135.1, 265.3, 308.3, 339.3, 419.3	0.0144	1.63	4.7
3.33	152.0567	[M+H] ⁺	C ₅ H ₅ N ₅ O	Guanine ^a	135.0, 110.0, 107.0, 82.0	< 0.001	9.5	3.87
2.38	259.0219	[M-H] ⁻	C ₆ H ₁₃ O ₉ P	D-Glucose 6-phosphate ^a	78.96, 96.97, 138.98, 199.0, 222.8, 241.0	< 0.001	0.12	2.41
4.16	298.0966	[M+H] ⁺	C ₁₁ H ₁₅ N ₅ O ₃ S	5'-Methylthioadenosine ^a	136.1, 145.0, 97.0, 75.0, 61.0	< 0.001	2.63	6.58
22.09	326.3049	[M+H] ⁺	C ₂₀ H ₃₉ NO ₂	N-Oleylethanolamine	62.1, 309.3, 69.1, 95.1, 135.1, 163.1, 265.3	< 0.001	3.79	2.11
0.95	146.1651	[M+H] ⁺	C ₇ H ₁₉ N ₃	Spermidine ^a	72.1, 58.1, 75.1, 84.1, 112.1, 129.1	0.0019	3.97	6.71
7.12	606.0735	[M-H] ⁻	C ₁₇ H ₂₇ N ₃ O ₁₇ P ₂	UDP-N-acetylglucosamine	384.98, 402.99, 362.0, 282.0, 158.9, 176.9, 78.96	< 0.001	2.15	16.36
8.03	565.0471	[M-H] ⁻	C ₁₅ H ₂₄ N ₂ O ₁₇ P ₂	UDP-glucose ^a	323.0, 384.98, 241.0, 158.9, 402.99, 96.97, 78.96	< 0.001	0.47	2.62

Table S1. Continued.

RT	m/z	Adduct ion	Elemental composition	Metabolite identification ^a	Product ions	P value	Fold Change	VIP
4.61	426.0218	[M-H] ⁻	C ₁₀ H ₁₅ N ₅ O ₁₀ P ₂	ADP ^a	78.96,134.0,158.9,272.96,328.0,408.0,346.1,96.97	0.0394	3.83	5.5
4.14	347.0392	[M-H] ⁻	C ₁₀ H ₁₃ N ₄ O ₈ P	IMP ^a	78.96,96.97,135.0,150.98,211.0	< 0.001	0.29	3.59
3.67	362.0503	[M-H] ⁻	C ₁₀ H ₁₄ N ₅ O ₈ P	GMP ^a	78.96,96.97,150.0,211.0	< 0.001	6.73	5.29
4.25	363.0344	[M-H] ⁻	C ₁₀ H ₁₃ N ₄ O ₉ P	XMP	211.0,78.96,96.7,108.0,151.0	< 0.001	33.98	3.39
3.68	283.0680	[M-H] ⁻	C ₁₀ H ₁₂ N ₄ O ₆	Xanthosine ^a	151.0,108.0	< 0.001	37.38	3.2
3.89	323.0280	[M-H] ⁻	C ₉ H ₁₃ N ₂ O ₉ P	UMP ^a	78.96,96.97,111.0,150.98,192.99,211.0,280.0	< 0.001	5.26	4.21
3.32	284.0986	[M+H] ⁺	C ₁₀ H ₁₃ N ₅ O ₅	Guanosine ^a	152.1,135.0,110.0	< 0.001	11.09	3.52

Table S2. The differential metabolites between SLCDC and SLC group obtained by LC-(±) MS/MS analysis.

RT	m/z	Adduct ion	Elemental composition	Metabolite identification ^a	Product ions	P value	Fold Change	VIP
2.38	130.0500	[M+H] ⁺	C ₅ H ₇ NO ₃	Pyrrolidonecarboxylic acid ^a	84.0,56.1,102.1	0.0014	4.41	3.96
1.27	132.0769	[M+H] ⁺	C ₄ H ₉ N ₃ O ₂	Creatine ^a	90.1,114.1,86.1,73.0,60.1	0.0017	3.53	5.57
3.4	166.0861	[M+H] ⁺	C ₉ H ₁₁ NO ₂	Phenylalanine ^a	120.1,103.1,107.0,131.0,149.1	0.0029	1.97	15.99
4.28	205.0971	[M+H] ⁺	C ₁₁ H ₁₂ N ₂ O ₂	L-Tryptophan ^a	188.1,146.1, 159.1,144.1,118.1,115.1	< 0.001	2.21	6.39
1.26	182.0811	[M+H] ⁺	C ₉ H ₁₁ NO ₃	L-Tyrosine ^a	136.1,91.1,119.05,123.0,147.0,165.1	0.0408	0.66	2.59
4.7	261.1444	[M+H] ⁺	C ₁₁ H ₂₀ N ₂ O ₅	Gamma-Glu-Leu	86.1,132.1,198.1,244.1	< 0.001	7.85	2.89
3.3	209.0919	[M+H] ⁺	C ₁₀ H ₁₂ N ₂ O ₃	Kynurenine ^a	146.1,94.1,136.1,192.1,174.1,150.1,120.0,99.0,	< 0.001	58.84	3.39

Table S2. Continued.

RT	m/z	Adduct ion	Elemental composition	Metabolite identification ^a	Product ions	P value	Fold Change	VIP
4.16	220.1179	[M+H] ⁺	C ₉ H ₁₇ NO ₅	Pantothenic Acid ^a	90.1,85.1,124.1,142.1,160.1,184.1,202.1	< 0.001	3.92	9.79
4.15	218.1023	[M-H] ⁻	C ₉ H ₁₇ NO ₅	Pantothenic Acid ^a	88.0,146.1,71.0,99.0,116.1	0.0179	2.9	6.49
3.72	237.0869	[M+H] ⁺	C ₁₁ H ₁₂ N ₂ O ₄	L-Formylkynurenine ^a	146.1,136.1,174.1,192.1,202.05,94.1	0.0097	24.55	2.34
4.28	188.0706	[M+H] ⁺	C ₁₁ H ₉ NO ₂	Indoleacrylic acid ^a	146.1,115.1,170.1	< 0.001	2.23	7.47

0.86	188.1756	[M+H] ⁺	C ₉ H ₂₁ N ₃ O	N1-Acetylspermidine	72.1,171.1,117.1, 100.1, 58.1	< 0.001	0.57	10.24
1.16	245.2333	[M+H] ⁺	C ₁₂ H ₂₈ N ₄ O	N1-Acetylspermine ^a	100.1,112.1,129.1,171.1,84.1,70.1	0.0019	2.83	2.46
1.77	306.076	[M-H] ⁻	C ₁₀ H ₁₇ N ₃ O ₆ S	Glutathione ^a	143.0,128.0,160.0,210.1,99.1,254.1	< 0.001	1.92	5.43
7.22	204.0865	[M+H-H ₂ O] ⁺	C ₈ H ₁₅ NO ₆	N-Acetyl-D-glucosamine ^a	138.1,144.1,126.1,168.1,186.1,109.0,96.0,84.0	< 0.001	16.15	13.34
4.16	298.0966	[M+H] ⁺	C ₁₁ H ₁₅ N ₅ O ₃ S	5'-Methylthioadenosine ^a	136.1,145.0,97.0,75.0,61.0	< 0.001	1.79	5.38
1.12	162.1124	[M+H] ⁺	C ₇ H ₁₅ NO ₃	L-Carnitine ^a	103.0,85.0,60.1	< 0.001	6.41	3.98
1.27	204.1231	[M+H] ⁺	C ₉ H ₁₇ NO ₄	Acetylcarnitine ^a	85.0,145.0, 60.1,102.1	0.0018	6.97	2.18
1.26	218.1386	[M+H] ⁺	C ₁₀ H ₁₉ NO ₄	Propionylcarnitine	85.0,159.1,144.1,60.1	0.0049	13.95	3.42
4.1	246.1698	[M+H] ⁺	C ₁₂ H ₂₃ NO ₄	Pentanoylcarnitine	187.1,85.0,144.1,60.1	0.0026	16.47	4.15
10.71	372.3104	[M+H] ⁺	C ₂₁ H ₄₁ NO ₄	Tetradecanoylcarnitine	85.0,60.1,95.1, 144.1,313.2,211.2	0.0044	5.02	2.09
11.95	400.3417	[M+H] ⁺	C ₂₃ H ₄₅ NO ₄	L-Palmitoylcarnitine	85.0,341.3,239.2,144.1,95.1,60.1	< 0.001	7.25	3.63
12.37	426.3573	[M+H] ⁺	C ₂₅ H ₄₇ NO ₄	Oleoylcarnitine	85.0,367.3,265.3,60.1,144.1,95.1	0.0017	3.53	2.99
14.5	428.3730	[M+H] ⁺	C ₂₅ H ₄₉ NO ₄	Stearoylcarnitine	85.0,369.3,267.3,144.1,95.1,60.1	< 0.001	15.53	2.44
1.28	258.1099	[M+H] ⁺	C ₈ H ₂₀ NO ₆ P	Glycerophosphocholine	104.1,60.1,86.1,125.0,184.1	< 0.001	2.99	3.77

Table S2. Continued

RT	m/z	Adduct ion	Elemental composition	Metabolite identification ^a	Product ions	P value	Fold Change	VIP
1.16	104.1074	[M+H] ⁺	C ₅ H ₁₃ NO	Choline ^a	60.1,58.1,59.1	0.0466	1.61	5.01
20.69	524.3707	[M+H] ⁺	C ₂₆ H ₅₄ NO ₇ P	LPC(18:0)	184.1, 506.36, 125.0, 104.1, 86.1	< 0.001	3.36	7.53
13.35	568.3391	[M+H] ⁺	C ₃₀ H ₅₀ NO ₇ P	LPC(22:6)	184.1, 550.4, 104.1, 125.0	< 0.001	15.24	2.08
14.9	480.3081	[M+H] ⁺	C ₂₃ H ₄₆ NO ₇ P	LPE(18:1) ^a	339.3, 419.3, 462.3, 308.3, 265.3, 69.1,	< 0.001	0.59	2.36
20.54	482.3237	[M+H] ⁺	C ₂₃ H ₄₈ NO ₇ P	LPE(18:0)	341.3, 310.3, 421.3, 464.3, 267.3, 69.1	0.0016	2.04	4.21
2.4	664.1154	[M+H] ⁺	C ₂₁ H ₂₈ N ₇ O ₁₄ P ₂	NAD ^a	136.1, 232.1, 348.1, 428.0, 97.0	< 0.001	3.71	2.84
3.32	284.0986	[M+H] ⁺	C ₁₀ H ₁₃ N ₅ O ₅	Guanosine ^a	152.1, 135.0, 110.0	< 0.001	4.4	2.67
3.9	325.0428	[M+H] ⁺	C ₉ H ₁₃ N ₂ O ₉ P	UMP ^a	97.0, 113.0, 227.1, 69.0	0.0063	6.31	2.06
3.67	362.0503	[M-H] ⁻	C ₁₀ H ₁₄ N ₅ O ₈ P	GMP ^a	78.96, 96.97, 150.0, 211.0	< 0.001	2.4	2.67
8.03	565.0471	[M-H] ⁻	C ₁₅ H ₂₄ N ₂ O ₁₇ P ₂	UDP-glucose ^a	323.0, 384.98, 241.0, 158.9, 402.99,	< 0.001	1.73	2.67
7.12	606.0735	[M-H] ⁻	C ₁₇ H ₂₇ N ₃ O ₁₇ P ₂	UDP-N-acetylglucosamine	384.98,402.99,362.0,282.0,158.9,176.9	< 0.001	4.43	29.66
3.33	152.0567	[M+H] ⁺	C ₅ H ₅ N ₅ O	Guanine ^a	135.0, 110.0, 107.0, 82.0	0.0389	4.1	3.09
1.84	137.0458	[M+H] ⁺	C ₅ H ₄ N ₄ O	Hypoxanthine ^a	119.0, 110.0, 94.0, 82.0, 67.0, 55.0	0.0331	0.68	2.6
1.75	148.0426	[M+H-H ₂ O] ⁺	C ₅ H ₁₁ NO ₃ S	Methionine sulfoxide ^a	84.0, 102.1, 130.05, 56.1, 74.0	< 0.001	3.76	2.9