

Table S1. Differentially expressed serum endogenous metabolites in Nor, Cis and MEAG groups.

Metabolite	RT, min	Measured m/z	Adduct	Formula	Cis vs. Nor				MEAG vs. Cis			
					FC	V	P-value	Trend	FC	V	P-value	Trend
L-phenylalanine	1.287 /	166.0860 /	(M+H)+ /	C9H11NO2	2.6645 /	1.0618 /	2.8132x10 <sup>-5</sup> /	↑/↑.	-1.8279 /	1.1342 /	7.9870x10 <sup>-4</sup> /	↓/↓.
	1.343	164.0715	(M-H)-		3.7194	1.3234	7.0602x10 <sup>-7</sup>		-2.1005	1.1563	9.3350x10 <sup>-4</sup>	
L-carnitine	0.835	162.1128	(M+H)+	C7H15NO3	3.3979	1.0839	1.4820x10 <sup>-5</sup>	↑	-2.5086	1.1840	4.0438x10 <sup>-4</sup>	↓
Creatine	0.858	132.0763	(M+H)+	C4H9N3O2	-3.9873	1.1408	1.5123x10 <sup>-8</sup>	↓	2.3159	1.2268	1.5702x10 <sup>-4</sup>	↑
Phosphocholine	11.821	184.0728	(M)+	C5H15NO4P	1.6580	1.0996	5.2402x10 <sup>-6</sup>	↑	-1.0745	1.1725	2.6454x10 <sup>-4</sup>	↓
LysoPE (16:0/0:0)	11.727	454.2934	(M+H)+	C21H44NO7P	-4.0524	1.1468	5.0280x10 <sup>-9</sup>	↓	2.0566	1.2379	1.4216x10 <sup>-4</sup>	↑
sStearoyl-L-carnitine	12.937	428.3734	(M+H)+	C25H49NO4	-4.6133	1.1197	3.3669x10 <sup>-7</sup>	↓	2.5077	1.1712	4.1441x10 <sup>-4</sup>	↑
L-tryptophan	4.118	203.0826	(M-H)-	C11H12N2O2	3.3316	1.2970	7.2559x10 <sup>-6</sup>	↑	-2.1848	1.1798	6.7676x10 <sup>-4</sup>	↓
Citric acid	0.922	191.0188	(M-H)-	C6H8O7	3.5848	1.1097	1.2978x10 <sup>-3</sup>	↑	-1.5454	1.1738	5.8196x10 <sup>-4</sup>	↓
Ascorbic acid	0.919	175.0243	(M-H)-	C6H8O6	-3.1022	1.0061	6.5670x10 <sup>-3</sup>	↓	2.6874	1.2327	4.2563x10 <sup>-4</sup>	↑
Hyochoic acid	6.792	407.2796	(M-H)-	C24H40O5	5.2868	1.0319	7.4059x10 <sup>-3</sup>	↑	-10.7310	1.2635	2.0890x10 <sup>-5</sup>	↓
Docosahexaenoic acid	11.173	327.2325	(M-H)-	C22H32O2	-3.4037	1.1619	4.2741x10 <sup>-4</sup>	↓	3.6107	1.2709	2.8350x10 <sup>-5</sup>	↑
L-tyrosine	1.216	180.0660	(M-H)-	C9H11NO3	5.2108	1.2661	1.4192x10 <sup>-5</sup>	↑	-3.2157	1.1713	5.2116x10 <sup>-4</sup>	↓
L-pyroglutamic acid	0.952	128.0350	(M-H)-	C5H7NO3	3.6354	1.3284	4.3286x10 <sup>-7</sup>	↑	-1.5794	1.0661	3.4123x10 <sup>-3</sup>	↓

LysoPE, lysophosphatidylethanolamine; RT, retention time; measured m/z, measured mass-to-charge ratio; FC, fold change; V, variable importance in projection; Nor, normal control group; Cis, cisplatin group; MEAG, methanolic extract of BaiYangJie. iYangJie. BaiYangJie. iYangJie. atin group; Mrease in the corresponding comparison.

Table SII. Topological properties of core targets.

No.	Gene name	Degree centrality	Betweenness centrality	Closeness centrality
1	AKT1	117	3287.6824	0.6923
2	EGFR	101	2511.0270	0.6592
3	ALB	103	2134.2258	0.6571
4	BCL2	98	2081.5960	0.6469
5	HIF1A	92	1869.2987	0.6449
6	HSP90AA1	95	1563.5806	0.6409
7	SRC	102	1432.9989	0.6350
8	ESR1	93	1420.6189	0.6311
9	PTGS2	82	1304.1351	0.6292
10	MAPK3	85	1108.3928	0.6106
11	GSK3B	73	965.2620	0.6000

For each centrality metric, genes are ranked independently. Thus, the three gene columns list the top-ranked genes for degree centrality, betweenness centrality and closeness centrality, respectively, and row-wise entries are not intended for direct one-to-one comparison.