

Pathway Name	Match Status	s	p	FDR	Impact
Aminoacyl-tRNA biosynthesis	16/48		4.85E-07	4.08E-05	0
Phenylalanine, tyrosine and tryptophan biosynthesis	4/4		4.60E-05	0.001930	1
Valine, leucine and isoleucine biosynthesis	4/8		0.002464	0.067298	0
Glyoxylate and dicarboxylate metabolism	8/32		0.003518	0.067298	0.1613
Arginine biosynthesis	5/14		0.004005	0.067298	0.3959
Alanine, aspartate and glutamate metabolism	7/28		0.006313	0.088386	0.5344
Histidine metabolism	5/16		0.007627	0.091526	0.3114
Nitrogen metabolism	3/6		0.009347	0.098146	0
Arginine and proline metabolism	8/38		0.01067	0.099587	0.2677
Pantothenate and CoA biosynthesis	5/19		0.016569	0.13918	0.0214
Taurine and hypotaurine metabolism	3/8		0.023114	0.16373	0.7142
Glutathione metabolism	6/28		0.024166	0.16373	0.1829
beta-Alanine metabolism	5/21		0.025339	0.16373	0.4552
Nicotinate and nicotinamide metabolism	4/15		0.030349	0.18209	0.1943
Phenylalanine metabolism	3/10		0.043788	0.24521	0.6190

带格式表格[Charles Wu]

Supplemental Table 3: Results of the Pathway Analysis of Metabolic Biomarkers.

FDR: false discovery rate.

设置格式[Charles Wu]: 字体: (默认) Times New Roman,
(中文)微软雅黑, 11 磅, 图案: 清除(白色), 字体颜色: 黑色