

Pathway Heat Map This is the heatmap associated with the statistical analysis of the data
Indicates ratios, *p*- and *q*-values for each comparison

0.55	Green: indicates significant difference ($p \leq 0.05$) between the groups shown; GREEN indicates a metabolite ratio of < 1
1.71	Red: indicates significant difference ($p \leq 0.05$) between the groups shown; RED indicates a metabolite ratio of > 1
1.43	Bold blue: narrowly missed statistical cutoff for significance; $0.05 < p < 0.10$
1.20	Non-colored text and cell: mean values are not significantly different for that comparison

PATHWAY SORT	SUPER PATHWAY	SUB PATHWAY	BIOCHEMICAL NAME	PLATFORM	COMP ID	Kegg	HMDB	Fold of Change				
								Day 0	Day 5		Day 10	
									(- CO)		(+CO)	
									(-CO)	(+CO)	(-CO)	(+CO)
1	Amino acid	Glycine, serine and threonine metabolism	glycine	GC/MS	11777	C00037	HMDB00123	1.06	0.65	1.23	0.71	1.01
3			dimethylglycine	GC/MS	5086	C01026	HMDB00092	0.90	1.11	1.16	1.13	1.90
6			beta-hydroxypyruvate	GC/MS	15686	C00168	HMDB01352	1.07	0.77	1.71	1.01	1.58
7			serine	GC/MS	1648	C00065	HMDB03406	0.79	0.91	0.64	0.81	0.38
11			homoserine	GC/MS	23642	C00263,C02926	HMDB00719	1.14	0.82	0.74	0.82	0.38
16			threonine	GC/MS	1284	C00188	HMDB00167	0.61	1.04	0.76	0.72	0.52
23		Alanine and aspartate metabolism	alanine	GC/MS	1126	C00041	HMDB00161	0.85	0.85	0.93	0.68	0.63
27			N-acetyl-beta-alanine	LC/MS pos	37432	C01073		1.09	0.93	1.14	1.14	1.48
29			aspartate	GC/MS	15996	C00049	HMDB00191	0.92	0.92	0.88	0.65	0.44
38		Glutamate metabolism	glutamate	LC/MS pos	57	C00025	HMDB0339	1.13	0.90	1.17	0.91	1.28
41			glutamate, gamma-methyl ester	LC/MS pos	33487			1.17	0.92	1.37	0.93	1.84
42			glutamine	GC/MS	1647	C00064	HMDB00641	0.95	0.79	0.76	1.89	0.33
44			gamma-aminobutyrate (GABA)	GC/MS	1416	C00334	HMDB00112	1.57	1.47	1.88	1.01	1.19
45			N-acetylglutamate	LC/MS pos	15720	C00624	HMDB01138	1.10	0.89	0.94	1.07	0.43
48			histidine	GC/MS	59	C00135	HMDB00177	0.94	0.99	1.10	0.64	1.13
66		Lysine metabolism	lysine	LC/MS pos	1301	C00047	HMDB00182	0.79	1.15	0.84	0.97	0.98
67			2-amino adipate	GC/MS	6146	C00956	HMDB00510	1.00	1.52	1.27	0.44	0.84
69			pipecolate	LC/MS pos	1444	C00408	HMDB00070	0.90	0.95	1.51	0.58	1.66
74		Phenylalanine & tyrosine metabolism	N6-acetylysine	LC/MS pos	36752	C02727	HMDB00206	0.81	1.01	1.23	0.60	1.47
78			phenylalanine	LC/MS pos	64	C00079	HMDB00159	0.99	1.06	1.07	0.78	1.05
90			tyrosine	LC/MS pos	1299	C00082	HMDB00158	1.00	0.96	0.97	0.65	0.70
126		Tryptophan metabolism	tryptophan	LC/MS pos	54	C00078	HMDB00929	0.92	1.02	0.99	0.74	0.77
157		Valine, leucine and isoleucine metabolism	isoleucine	LC/MS pos	1125	C00407	HMDB00172	0.88	1.00	0.81	0.84	0.85
158			leucine	LC/MS pos	60	C00123	HMDB00687	0.93	0.88	0.81	0.89	0.83
167			valine	LC/MS pos	1649	C00183	HMDB00883	0.93	0.91	0.74	0.83	0.48
188		Cysteine, methionine, SAM, taurine metabolism	cysteine	GC/MS	31453	C00097	HMDB00574	0.97	1.01	1.11	1.14	0.74
198		S-adenosylhomocysteine (SAH) metabolism	S-adenosylhomocysteine (SAH)	LC/MS pos	15948	C00021	HMDB00939	1.22	0.98	0.87	0.83	0.68
213			arginine	GC/MS	1638	C00062	HMDB00517	0.90	0.80	0.96	0.53	0.60
218			proline	LC/MS pos	1898	C00148	HMDB00162	0.97	0.72	0.83	0.72	0.46
220		Urea cycle; arginine-, proline-, citrulline metabolism	citrulline	LC/MS pos	2132	C00327	HMDB00904	1.14	0.84	0.79	0.68	0.63
221			N-acetylornithine	LC/MS pos	15630	C00437	HMDB03357	0.75	1.10	1.17	0.49	0.84
238			5-methylthioadenosine (MTA)	LC/MS pos	1419	C00170	HMDB01173	1.22	0.81	0.91	0.81	0.32
253		Glutathione metabolism	5-oxoproline	LC/MS neg	1494	C01879	HMDB00267	1.01	0.94	0.84	1.55	0.67
321	Gamma-glutamyltranspeptidase (GGT) metabolism		gamma-glutamylvaline	LC/MS pos	32393		HMDB11172	1.06	0.86	0.76	0.72	0.24
322			gamma-glutamyl-2-aminobutyrate	LC/MS pos	37092			0.94	1.07	0.81	1.50	0.50
323			gamma-glutamylleucine	LC/MS pos	18369		HMDB11171	1.07	0.78	0.76	0.58	0.19
324			gamma-glutamylsoleucine	LC/MS pos	34456		HMDB11170	1.06	0.83	0.81	0.53	0.32
327			gamma-glutamylmethionine	LC/MS pos	37539			1.05	1.05	0.69	0.65	0.21

328	Peptide	gamma-glutamyl	gamma-glutamylglutamate	LC/MS pos	36738			0.96	0.93	1.08	0.81	0.78
329			gamma-glutamylglutamine	LC/MS pos	2730		HMDB11738	1.03	0.85	0.65	1.26	0.23
331			gamma-glutamylphenylalanine	LC/MS pos	33422		HMDB00594	1.10	1.00	0.80	0.58	0.41
332			gamma-glutamyltyrosine	LC/MS pos	2734			0.96	1.03	0.85	0.73	0.73
333			gamma-glutamylthreonine*	LC/MS pos	33364			1.02	0.94	0.69	0.54	0.31
336			gamma-glutamylalanine	LC/MS pos	37063			0.95	0.96	0.93	0.94	0.31
368	Carbohydrate	Aminosugars metabolism	N-acetylglucosamine	GC/MS	15095	C00140	HMDB00215	1.00	0.89	1.00	0.80	0.50
372			erythronate*	GC/MS	33477		HMDB00613	1.12	0.81	1.52	1.21	2.08
388		Fructose, mannose, galactose, starch, and sucrose metabolism	erythrose	GC/MS	27722	C01796	HMDB02649	1.58	0.64	1.37	0.83	1.91
390			galactitol (dulcitol)	GC/MS	1117	C01697	HMDB00107	0.88	1.44	1.02	0.71	0.57
391			galactose	GC/MS	12055	C01582	HMDB00143	1.18	1.00	1.00	0.36	1.00
397			maltose	GC/MS	15806	C00208	HMDB00163	0.97	0.70	1.01	0.67	1.12
402			mannose	GC/MS	584	C00159	HMDB00169	0.95	0.73	1.02	0.72	0.93
404			mannose-6-phosphate	GC/MS	1470	C00275	HMDB01078	1.00	1.29	1.02	0.93	1.18
410			sorbitol	GC/MS	15053	C00794	HMDB00247	0.87	1.11	1.44	1.27	1.62
416			trehalose	GC/MS	15573	C01083	HMDB00975	0.97	0.88	1.02	0.59	0.69
432		Glycolysis, gluconeogenesis, pyruvate metabolism	glycerate	GC/MS	1572	C00258	HMDB00139	1.18	0.78	1.33	0.84	1.70
434			glucose-6-phosphate (G6P)	GC/MS	31260	C00668	HMDB01401	1.06	1.19	0.87	0.92	1.38
436			glucose	GC/MS	20488	C00293	HMDB00122	1.17	0.93	1.25	0.99	1.85
439			fructose-6-phosphate	GC/MS	12021	C05345	HMDB00124	1.04	1.22	1.14	1.14	1.30
447			1,3-dihydroxyacetone	GC/MS	35963	C00184	HMDB01882	1.07	0.78	1.36	1.03	3.05
457	Energy	Glyoxylate and dicarboxylate metabolism	oxalate (ethanedioate)	GC/MS	20694	C00209	HMDB02329	1.41	0.83	1.59	0.94	1.17
465		Nucleotide sugars, pentose metabolism	sedoheptulose-7-phosphate	GC/MS	35649	C05382	HMDB01068	1.33	1.23	1.18	0.90	3.55
466			gluconate	GC/MS	587	C00257	HMDB00625	0.78	1.16	1.28	0.78	2.01
468			ribose	GC/MS	12083	C00121	HMDB00283	1.49	1.25	0.99	0.75	0.58
475			Isobor: ribulose 5-phosphate, xylulose 5-phosphate	GC/MS	37288			0.73	0.95	1.00	0.78	0.80
476			UDP-glucose	GC/MS	32344	C00029	HMDB00286	1.02	0.88	1.11	0.77	0.86
480			arabinose	GC/MS	575	C00181	HMDB00646	1.10	1.09	1.03	1.23	1.00
491		Krebs cycle	citrate	GC/MS	1564	C00158	HMDB00094	1.36	0.82	1.35	0.80	2.22
493			cis-aconitate	LC/MS neg	12025	C00417	HMDB00072	0.94	0.84	1.11	0.91	0.94
498			isocitrate	LC/MS pos	12110	C00311	HMDB00193, HMDB01874	0.98	0.94	1.50	1.06	1.82
500			alpha-ketoglutarate	GC/MS	33453	C00026	HMDB00208	1.08	1.09	1.49	0.92	1.80
501			succinate	LC/MS neg	1437	C00042	HMDB00254	1.00	1.24	0.70	0.93	0.34
504			fumarate	GC/MS	1643	C00122	HMDB00134	1.05	0.88	1.44	0.88	1.86
507		Oxidative phosphorylation	malate	GC/MS	1303	C00149	HMDB00156	0.80	1.27	1.58	0.75	2.66
511			phosphate	LC/MS neg	34425	C00009	HMDB01429	1.00	0.97	1.00	1.01	1.09
512			pyrophosphate (PPi)	GC/MS	2078	C00013	HMDB00250	1.14	0.62	0.94	0.44	0.68
512.5	Long chain fatty acid	Essential fatty acid	linoleate (18:2n6)	LC/MS neg	1105	C01595	HMDB00673	0.96	1.14	1.45	1.22	10.15
516			linolenate [alpha or gamma; (18:3n3 or 6)]	LC/MS neg	34025	C06427	HMDB01388	1.08	1.00	1.61	1.20	6.15
532		Medium chain fatty acid	laurate (12:0)	LC/MS neg	1645	C02679	HMDB00638	0.80	1.00	1.02	1.15	1.24
538		Long chain fatty acid	palmitate (16:0)	GC/MS	1336	C00249	HMDB00220	0.97	1.05	1.51	0.87	2.07
539			palmitoleate (16:1n7)	GC/MS	1507	C08362	HMDB03229	1.27	0.54	1.69	1.32	4.69
541			margarate (17:0)	LC/MS neg	1121		HMDB02259	0.89	0.91	1.22	1.07	1.54
542			10-heptadecenoate (17:1n7)	LC/MS neg	33971			0.95	0.92	1.42	1.14	4.03
543			stearate (18:0)	GC/MS	1358	C01530	HMDB00827	0.87	1.11	1.17	1.00	1.49
544			tuberculostearate	GC/MS	34163		HMDB04085	1.05	0.98	1.16	1.18	0.86
545			oleate (18:1n9)	LC/MS neg	1359	C00712	HMDB00207	1.03	1.21	1.30	1.18	6.42

554		10-nonadecenoate (19:1n9)	LC/MS neg	33972			1.23	1.00	1.07	0.99	8.63		
573		lignocerate (24:0)	GC/MS	1364	C08320	HMDB02003	1.00	1.00	1.23	1.00	3.49		
576		hexacosanoate (26:0)	GC/MS	21143		HMDB02356	1.01	1.12	1.29	1.00	3.90		
589		Fatty acid, ester	n-Butyl Oleate	GC/MS	36802		0.99	1.04	0.91	0.80	1.28		
591			4-hydroxybutyrate (GHB)	GC/MS	34585	C00989	HMDB00710	1.00	1.04	0.90	1.08	1.80	
596	Lipid		8-hydroxyoctanoate	LC/MS neg	21239		HMDB00711	1.13	0.87	0.81	1.13	1.78	
599			2-hydroxymyristate	LC/MS neg	32413	C13790	HMDB02261	0.82	1.18	1.00	1.12	1.27	
602			2-hydroxystearate	LC/MS neg	17945	C03045		1.02	1.24	1.01	0.86	1.05	
604			2-hydroxypalmitate	LC/MS neg	35675			0.93	1.02	0.83	1.13	1.51	
608			13-HODE + 9-HODE	LC/MS neg	37752			2.81	3.31	1.91	0.77	12.18	
615			sebacate (decanedioate)	LC/MS neg	32398	C08277	HMDB00792	1.02	1.06	1.13	1.08	1.58	
616			azelate (nonanedioate)	LC/MS neg	18362	C08261	HMDB00784	0.81	1.24	1.00	1.21	1.37	
619			hexadecanedioate	LC/MS neg	35678		HMDB00672	0.86	0.77	1.15	1.24	1.88	
626		Fatty acid, amide	stearamide	LC/MS pos	37487	C13846		0.97	1.04	0.85	1.37	1.20	
632.6		Fatty acid, branched	15-methylpalmitate	GC/MS	38295			1.00	1.82	1.88	0.66	7.46	
679			palmitoyl ethanolamide	LC/MS neg	38165			0.95	0.86	1.15	1.13	1.07	
679.1		Endocannabinoid	stearoyl ethanolamide	GC/MS	38625			1.03	0.97	1.22	0.87	0.95	
744		Fatty alcohol, long chain	1-octadecanol	GC/MS	21128	D01924	HMDB02350	0.60	0.84	1.87	1.21	1.76	
746			ethanolamine	GC/MS	34285	C00189	HMDB00149	0.91	0.89	1.29	0.91	1.10	
749			diethanolamine	GC/MS	20691	C06772	HMDB04437	0.83	1.05	0.96	0.92	1.12	
750		Glycerolipid metabolism	glycerol	GC/MS	15122	C00116	HMDB00131	1.06	0.83	1.24	0.76	1.86	
752			glycerol 3-phosphate (G3P)	GC/MS	15365	C00093	HMDB00126	1.02	0.69	0.70	0.96	0.14	
760		Inositol metabolism	myo-inositol	GC/MS	19934	C00137	HMDB00211	0.98	1.00	0.69	0.88	0.32	
763			inositol 1-phosphate (I1P)	GC/MS	1481		HMDB00213	0.94	0.80	1.03	1.01	0.40	
776		Lysolipid	2-palmitoylglycerophosphoethanolamine*	LC/MS pos	35684			0.96	0.79	0.60	0.92	0.28	
813			1-docosahexaenyl glycerophosphocholine*	LC/MS pos	33822			2.13	1.00	1.89	1.14	0.46	
955			adenine	GC/MS	554	C00147	HMDB00034	1.22	0.61	0.71	0.82	0.25	
956			adenosine	LC/MS pos	555	C00212	HMDB00050	1.18	0.98	1.13	0.72	0.69	
964		Purine metabolism, adenine containing	adenosine 5'-monophosphate (AMP)	LC/MS pos	32342	C00020	HMDB00045	1.07	0.89	0.83	0.92	0.27	
966.5	Nucleotide		adenosine 5'-diphosphoglucose	LC/MS neg	35652	C06192,C00498	HMDB06369, HMDB06557	0.92	0.72	0.99	0.84	0.36	
970			adenosine 3',5'-cyclic monophosphate (cAMP)	LC/MS neg	2831	C00575	HMDB00058	0.83	0.72	0.91	0.52	1.60	
1016		Pyrimidine metabolism, thymine containing	thymine	GC/MS	604	C00178	HMDB00262	1.07	0.94	0.78	0.88	0.51	
1019			thymidine 5'-monophosphate	LC/MS neg	12023	C00364	HMDB01227	1.14	0.83	0.75	1.24	0.41	
1022		Pyrimidine metabolism, uracil containing	uracil	GC/MS	605	C00106	HMDB00300	0.91	0.61	0.74	0.75	1.26	
1036		Purine and pyrimidine metabolism	methylphosphate	GC/MS	37070			0.99	0.84	1.02	1.08	1.15	
1047			Biotin metabolism	biotin	LC/MS pos	568	C00120	HMDB00030	1.24	0.91	1.36	0.97	2.01
1063			nicotinamide	LC/MS pos	594	C00153	HMDB01406	0.96	0.82	1.23	0.77	1.34	
1066		Nicotinate and nicotinamide metabolism	nicotinamide adenine dinucleotide (NAD+)	LC/MS pos	5278	C00003	HMDB00902	0.97	0.76	0.63	1.03	0.14	
1077	Cofactors and vitamins		nicotinate	LC/MS pos	1504	C00253	HMDB01488	1.26	0.83	1.21	0.92	0.60	
1079			nicotinic acid mononucleotide (NaMN)	LC/MS pos	32461		HMDB01132	1.27	1.09	0.73	1.11	0.77	
1080			nicotinate ribonucleoside*	LC/MS pos	33471	C05841	HMDB06809	1.17	1.08	0.90	0.98	0.83	

1085	Pantothenate and CoA metabolism	pantothenate	LC/MS pos	1508	C00864	HMDB00210	0.94	1.07	0.70	1.25	0.54	
1089		acetyl CoA	LC/MS neg	32484	C00024	HMDB01206	1.06	1.20	1.18	1.06	0.63	
1119	Vitamin B6 metabolism	pyridoxine (Vitamin B6)	LC/MS pos	608	C00314	HMDB02075	1.17	0.92	1.31	0.94	1.73	
1128	Benzene metabolism	2-(hydroxymethyl)phenol	GC/MS	37492	C02323		1.40	0.77	0.97	0.67	1.20	
1137		4-hydroxybenzoate	GC/MS	21133	C00156	HMDB00500	1.08	0.98	0.95	0.86	0.86	
1150	Chemical Xenobiotics	bicine	GC/MS	20708		HMDB11727	0.87	1.18	0.96	0.90	1.19	
1151		glycolate (hydroxyacetate)	GC/MS	15737	C00160	HMDB00115	1.08	0.90	1.29	1.06	1.61	
1157		glycerol 2-phosphate	GC/MS	27728	C02979	D01488	HMDB02520	1.08	0.66	0.80	0.84	0.55
1161		heptaethylene glycol	LC/MS pos	38154			1.21	0.98	1.39	0.96	2.08	
1162		hexaethylene glycol	LC/MS pos	38133			1.19	0.93	1.62	0.92	2.22	
1165		tetraethylene glycol	LC/MS pos	38101			0.96	0.65	1.65	1.02	1.94	
1165.5		diethylene glycol	GC/MS	38099			2.50	0.77	1.06	0.74	1.47	
1175		trizma acetate	GC/MS	20710	C07182			0.11	0.41	1.32	0.39	0.56
1196.5		cyclohexylamine*	GC/MS	33452	C00571			1.09	1.17	1.10	0.94	1.25
1206.6		dexpanthenol	LC/MS pos	38314	D03726	HMDB04231	0.86	0.80	0.98	0.68	0.29	
1207.9		3-(2-pyridinyl)pyridine	LC/MS pos	38122			1.08	1.08	1.21	1.04	1.31	
1212		salicylate	GC/MS	1515	C00805	HMDB01895	0.81	0.85	1.91	0.87	0.92	
1287	Food component/Plant	4-hydroxybenzyl alcohol	GC/MS	15127	C17467	HMDB11724	22.52	0.91	0.89	0.55	1.02	
1330		ergothioneine	LC/MS pos	37459	C05570	HMDB03045	1.02	0.81	0.49	0.98	0.11	
1334	Sugar, sugar substitute, starch	5-hydroxymethylfurfural	LC/MS pos	37429	C11101			1.77	0.79	1.19	1.41	2.25
1358		trehalose 6-phosphate	GC/MS	15802	C00689	HMDB01124	1.00	0.69	0.53	0.56	0.24	
1360		erythritol	GC/MS	20699	C00503	HMDB02994	1.20	0.81	1.43	0.82	1.47	
1380.1	Bacterial	mycothione (MSSM)*	LC/MS pos	38682			0.99	0.83	0.57	0.85	0.12	