

Figure S1. Heat map of r (Pearson's correlation coefficient) value among different samples including replicates. The color represented the r value.

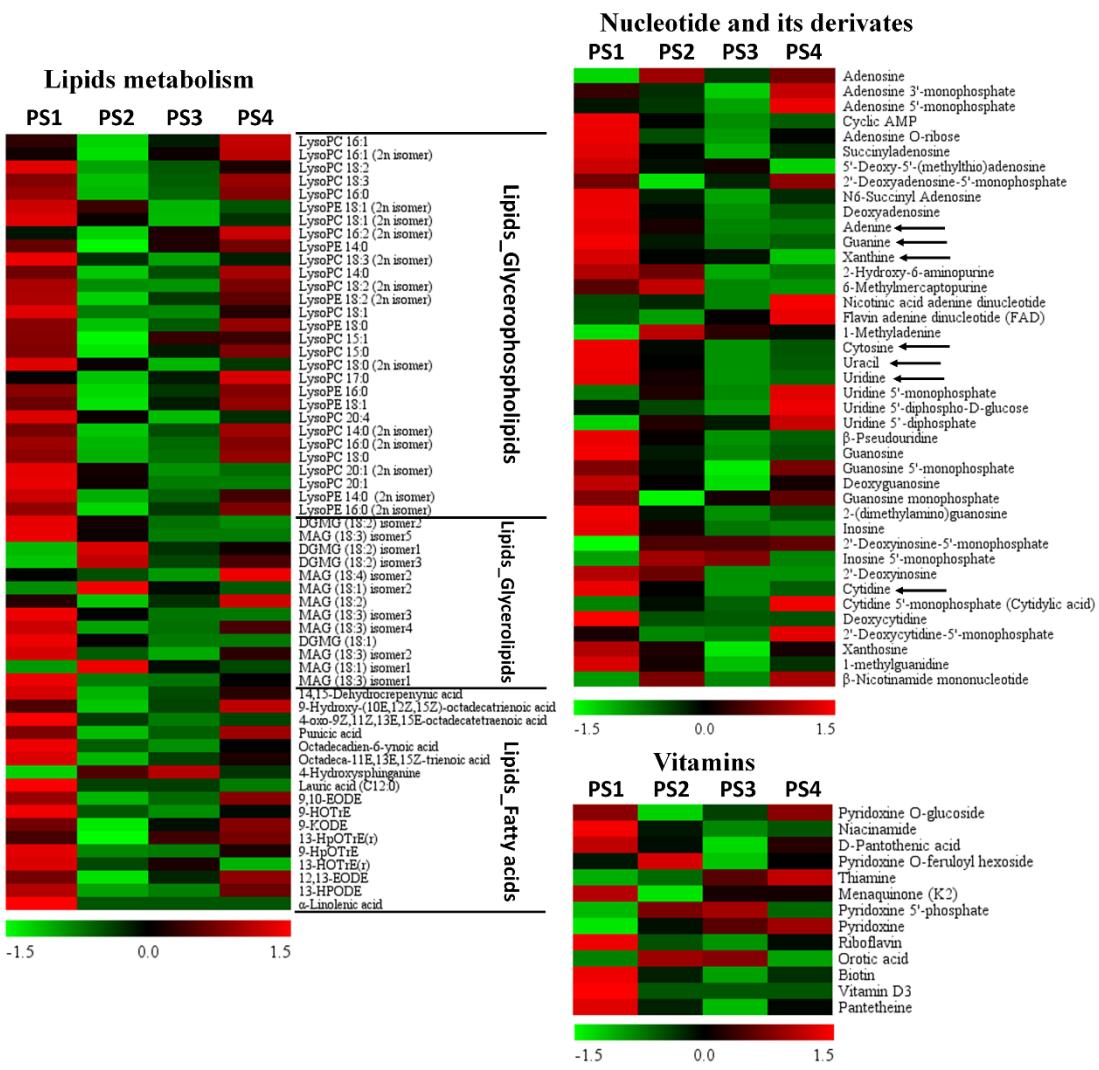


Figure S2. Distributions of accumulation profiles of lipids, nucleotides, and vitamins detected by widely-targeted UPLC-MC during four fruit developmental stages. The colors indicate the proportional content of each identified metabolites as determined by the average peak response area with R scale normalization. PS1, 2, 3, and 4 represents fruit samples collected at 27, 84, 125, 165 Days After Anthesis (DAA), respectively. Three independent replicates were performed for each stages.

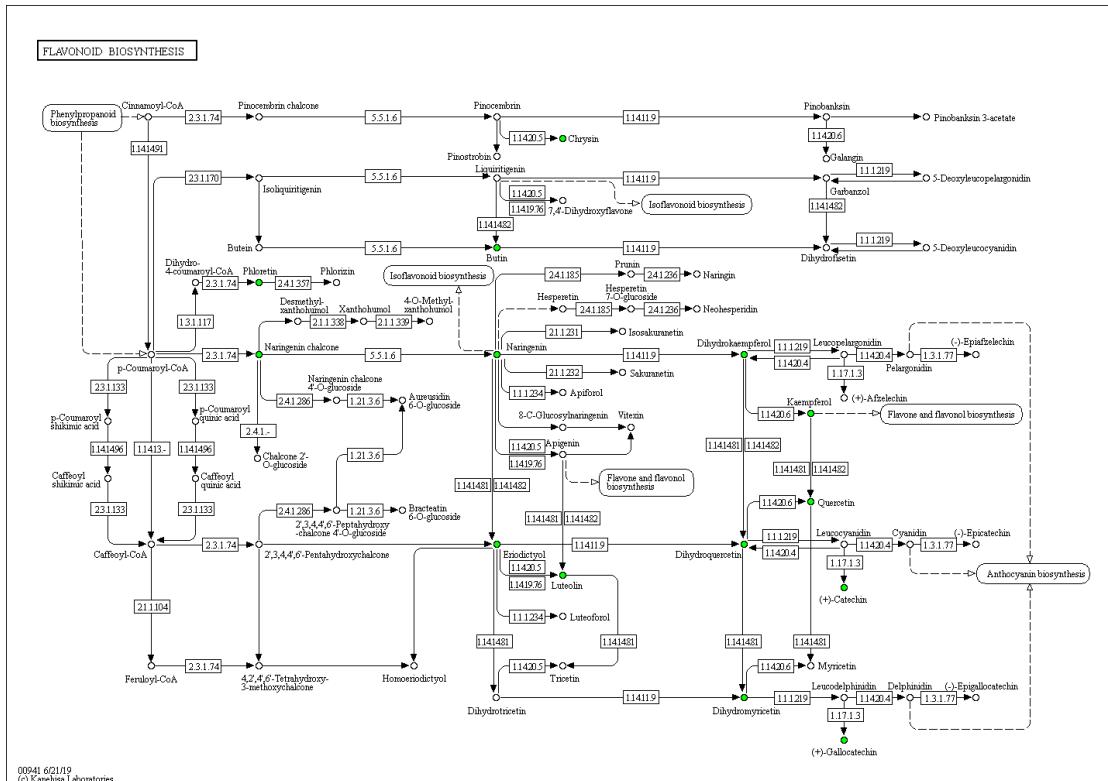


Figure S3. Differential metabolites of PS2 vs PS1 group in flavonoid biosynthesis pathway.

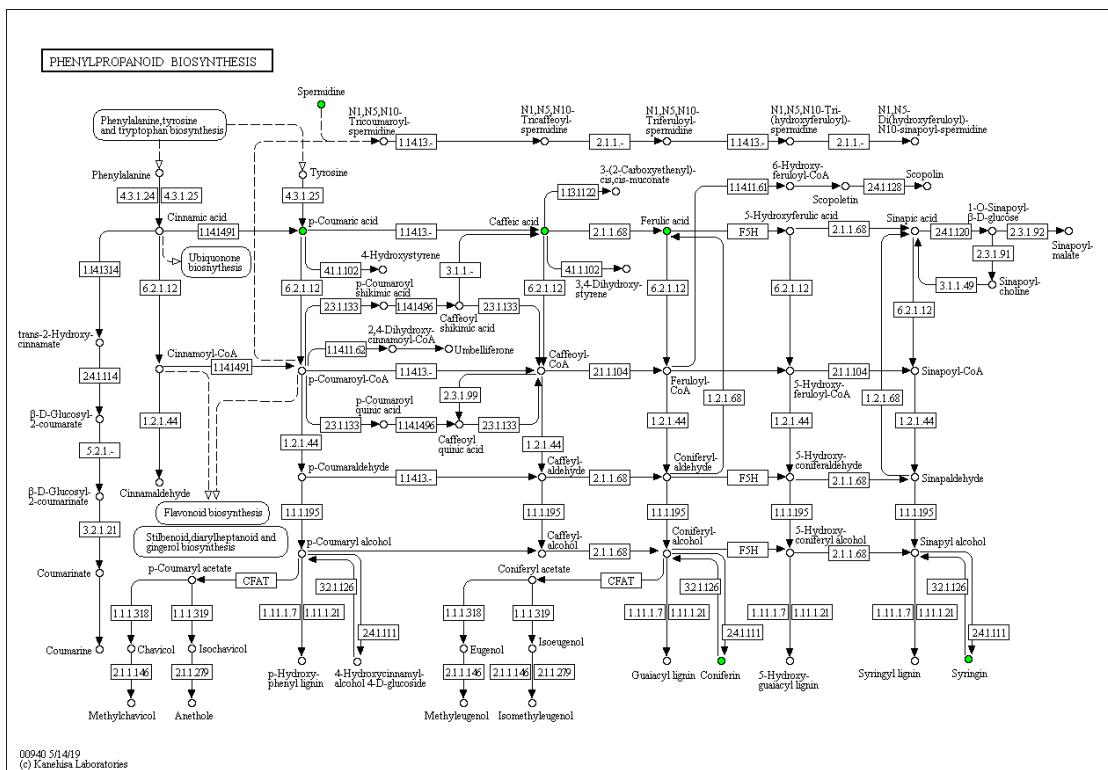


Figure S4. Differential metabolites of PS2 vs PS1 group in phenylpropanoid biosynthesis pathway.

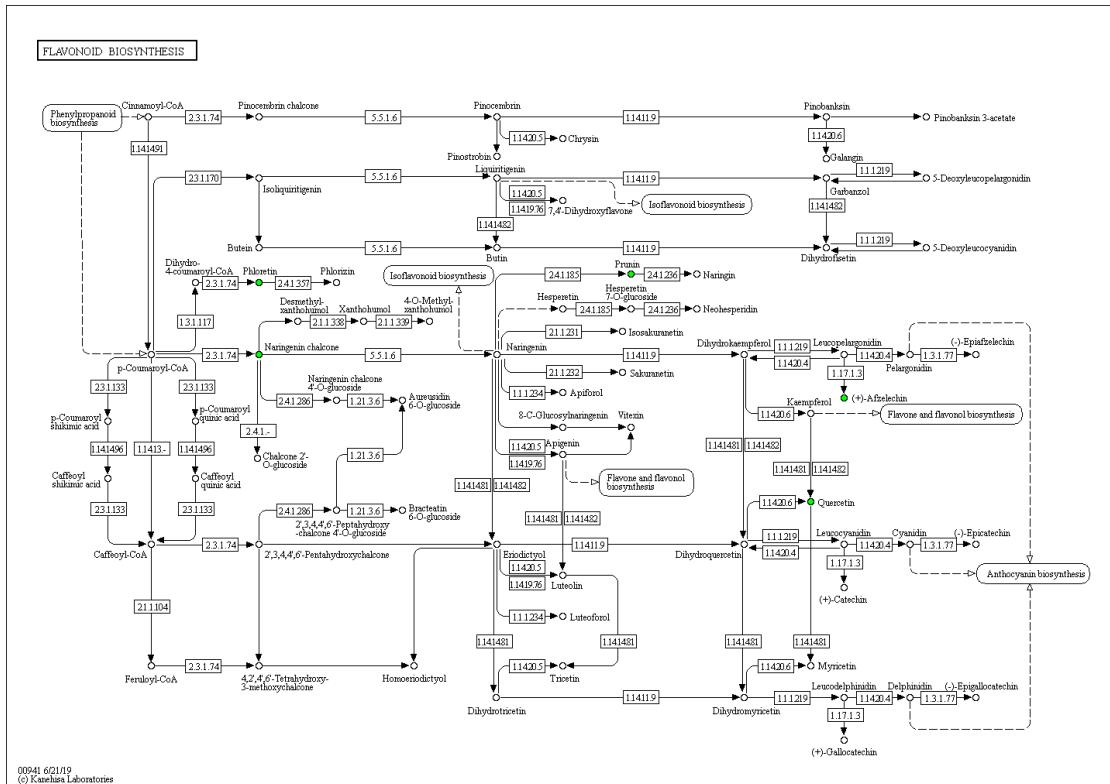


Figure S5. Differential metabolites of PS3 vs PS2 group in flavonoid biosynthesis pathway.

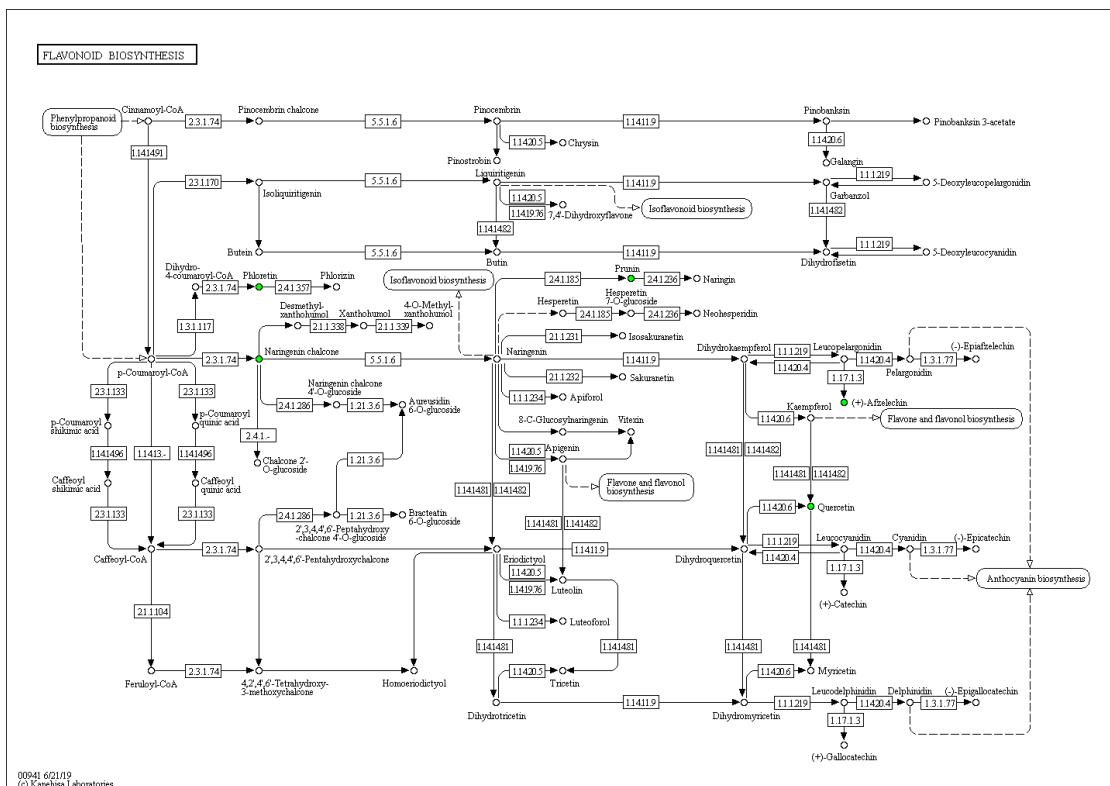


Figure S6. Differential metabolites of PS3 vs PS2 group in phenylpropanoid biosynthesis pathway.

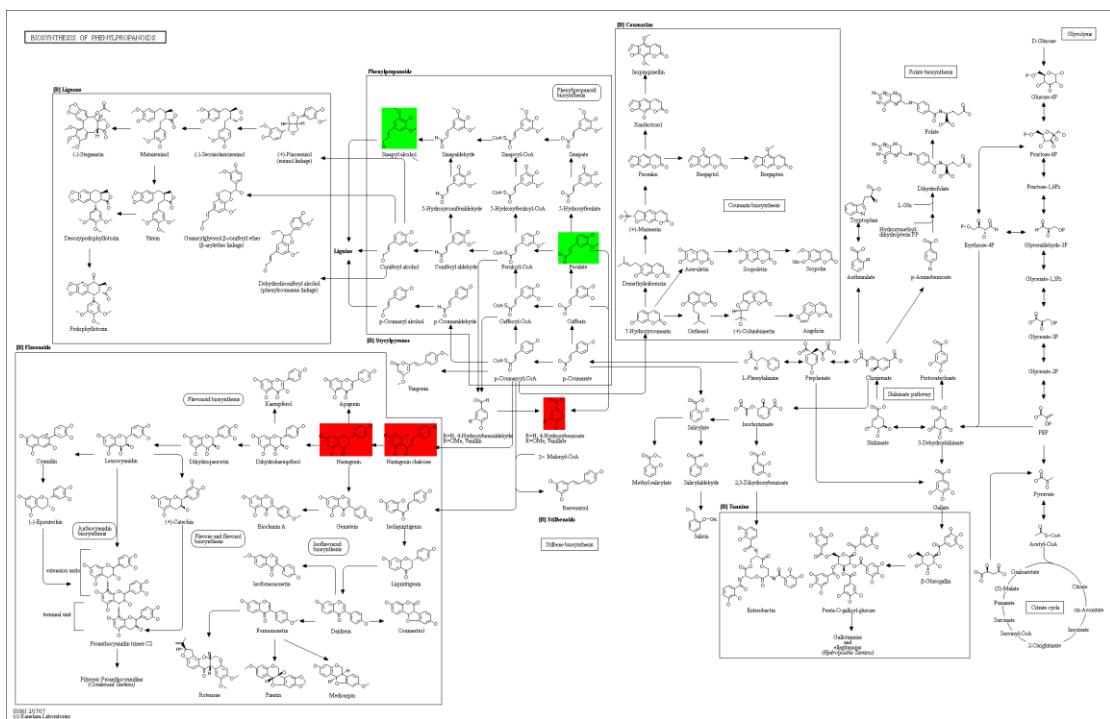


Figure S7. Differential metabolites of PS4 vs PS3 group in biosynthesis of phenylpropanoids pathway.

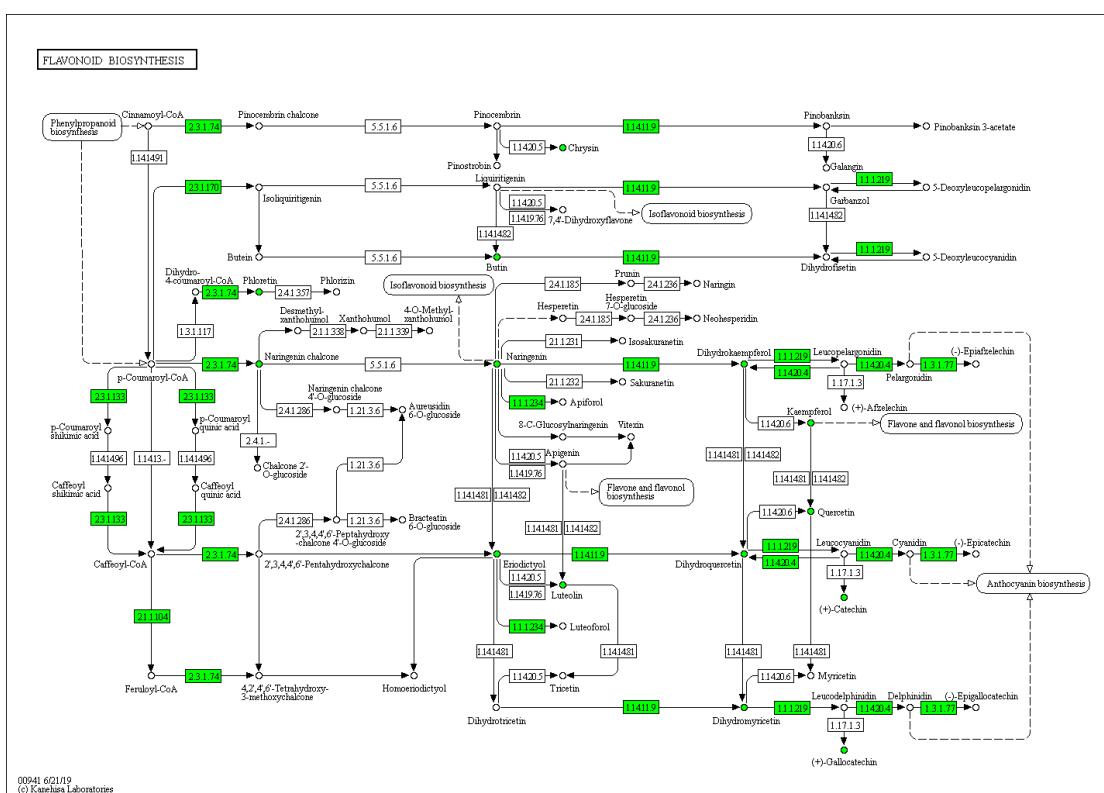
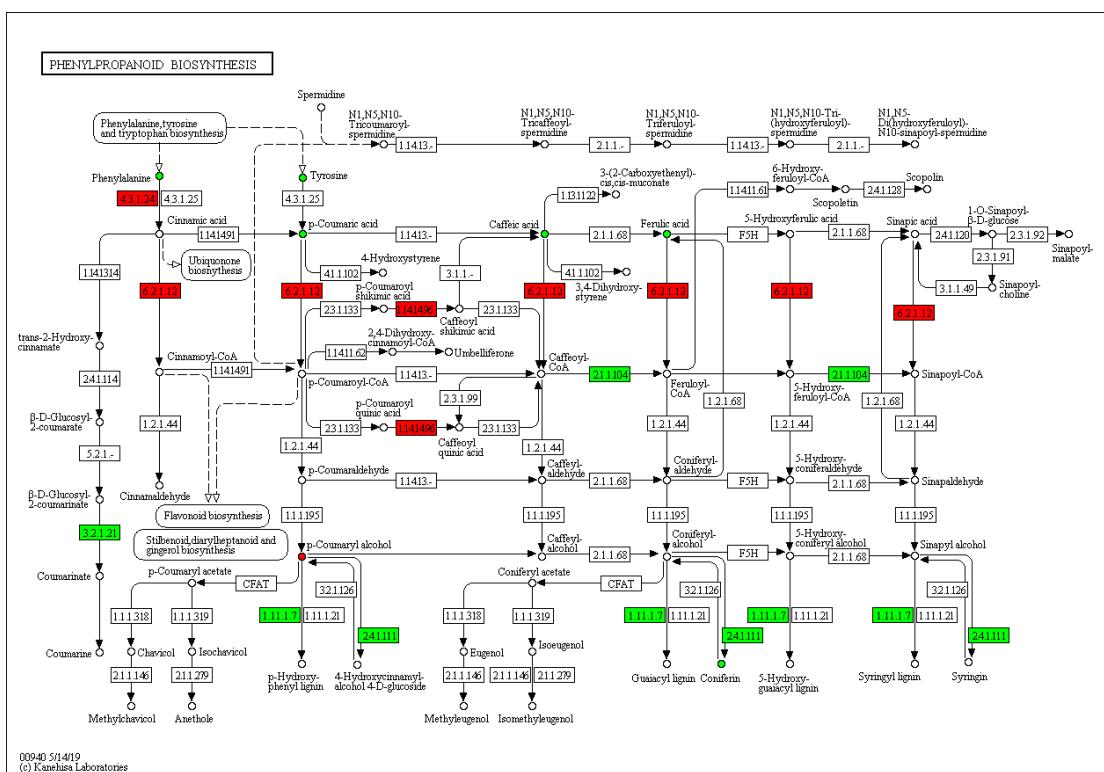


Figure S8. Differential metabolites of PS2 vs PS1 group in flavonoid biosynthesis pathway and phenylpropanoid biosynthesis pathway combined with RNA-seq results.

Sugars alcohols	Mad860	Negative	259	96.9	0.99	260.03	[M-H]-	D-Glucose 6-phosphate	Carbohydrates	1.60E+06	2.15E+06	9.30E+05	8.93E+05
	Mad55	Positive	183	59.3	0.77	182.079	[M+H]+	D-Sorbitol	Alcohols and polyols	5.70E+06	4.58E+06	5.69E+06	4.48E+06
	Mad56	Positive	183.08	111	0.77	182.079	[M+H]+	D-Mannitol	Alcohols and polyols	1.43E+07	1.27E+07	1.56E+07	1.19E+07
	Mad805	Negative	181.08	70.8	0.76	182.079	[M-H]-	Dulcitol	Alcohols and polyols	3.59E+06	4.22E+06	5.35E+06	2.96E+06
	Mad791	Negative	179.157	87	0.74	180.0634	[M-H]-	Inositol	Others	2.58E+05	2.89E+05	2.76E+05	4.98E+05
Nucleotides	Mad97	Positive	153	136.4	1.17	152	[M+H]+	Xanthine	Nucleotide and its derivates	8.60E+04	3.44E+03	2.22E+03	9.00E+00
	Mad157	Positive	245.1	113	1.42	244.0695	[M+H]+	Uridine	Nucleotide and its derivates	1.29E+07	3.41E+06	1.13E+06	1.51E+06
	Mad29	Positive	112	95	0.73	111.0433	[M+H]+	Cytosine	Nucleotide and its derivates	1.38E+06	5.48E+05	3.11E+05	3.91E+05
	Mad120	Positive	136.1	136.1	1.21	135.0545	[M+H]+	Adenine	Nucleotide and its derivates	7.45E+05	1.77E+05	5.76E+04	6.39E+04
	Mad107	Positive	113.027	95.9	1.19	112.027	[M+H]+	Uracil	Nucleotide and its derivates	1.50E+06	2.84E+05	1.03E+05	1.58E+05
	Mad109	Positive	152	135.1	1.19	151.049	[M+H]+	Guanine	Nucleotide and its derivates	2.99E+05	3.64E+04	1.68E+04	2.15E+04
Amino acids	Mad74	Positive	162.1	98.3	0.81	161.1	[M+H]+	2-Aminoadipic acid (L-Homoglutamic acid)	Amino acids	1.16E+05	5.51E+04	3.86E+04	7.40E+04
	Mad65	Positive	116.1	70	0.79	115.0633	[M+H]+		Amino acids	8.60E+06	6.56E+06	2.30E+06	2.30E+06
	Mad39	Positive	176.1	98.9	0.76	175.0957	[M+H]+	L-Citrulline	Amino acids	3.49E+05	2.87E+05	2.38E+05	2.78E+05
	Mad33	Positive	106	59.9	0.75	105.0426	[M+H]+	L-Serine	Amino acids	7.74E+05	6.65E+05	6.01E+05	5.39E+05
	Mad812	Negative	132	99.9	0.77	133.0375	[M-H]-	L-Aspartic acid	Amino acids	1.35E+06	5.99E+06	4.11E+06	1.05E+07
	Mad813	Negative	146.1	102.1	0.77	147.0532	[M-H]-	L-Glutamic acid	Amino acids	4.00E+06	7.34E+06	3.81E+06	8.75E+06
	Mad141	Positive	132.1	86	1.27	131.0946	[M+H]+	L-Leucine	Amino acids	1.26E+07	6.62E+06	5.17E+06	9.00E+00
	Mad196	Positive	166	120.1	1.96	165.079	[M+H]+	L-Phenylalanine	Amino acids	3.12E+07	3.81E+07	1.50E+07	1.39E+07
	Mad793	Negative	118	100.9	0.75	119.0582	[M-H]-	L-Threonine	Amino acids	6.97E+04	1.40E+05	1.35E+05	2.13E+05
	Mad15	Positive	147	83.2	0.67	146.1055	[M+H]+	L-(+)-Lysine	Amino acids	2.62E+05	2.86E+05	2.45E+05	4.01E+05
	Mad781	Negative	154.1	92.9	0.67	155.0695	[M-H]-	L-Histidine	Amino acids	7.71E+03	3.48E+04	2.19E+04	5.18E+04
	Mad78	Positive	118.1	71.8	0.81	117.079	[M+H]+	L-Valine	Amino acids	2.18E+07	1.00E+07	6.24E+06	1.03E+07
	Mad131	Positive	132	85.9	1.23	131.0946	[M+H]+	L-Isoleucine	Amino acids	1.60E+07	8.57E+06	1.49E+07	2.33E+07
	Mad784	Negative	173.1	131.1	0.69	174.1117	[M-H]-	L-(+)-Arginine	Amino acids	4.68E+04	1.89E+05	7.26E+04	9.42E+04

Mad796	Negative	118.1	101	0.75	119.0582	[M-H]-	L-Homoserine	Amino acids	1.47E+04	2.32E+04	1.68E+04	2.17E+04		
Mad797	Negative	131.054	71.8	0.75	132.054	[M-H]-	L-Asparagine	Amino acids	7.45E+06	1.55E+07	1.57E+07	2.38E+07		
Mad61	Positive	122.02	80.1	0.78	121.02	[M+H]+	L-Cysteine	Amino acids	7.12E+03	5.72E+03	1.02E+04	1.39E+04		
Mad136	Positive	138.084	103	1.25	137.084	[M+H]+	L-Tyramine	Amino acids	3.65E+04	6.42E+03	4.78E+03	5.20E+03		
Mad894	Negative	180.1	163	1.21	181.074	[M-H]-	L(-)-Tyrosine	Amino acids	7.29E+04	1.37E+05	1.41E+04	2.10E+04		
Mad105	Positive	150	72.8	1.18	149.051	[M+H]+	L-Methionine	Amino acids	4.68E+05	4.54E+06	2.36E+06	1.59E+06		
Mad798	Negative	145.069	127.1	0.75	146.069	[M-H]-	L-Glutamine	Amino acids	1.32E+04	6.35E+04	4.28E+04	9.48E+04		
Mad42	Positive	90.048	71.8	0.76	89.048	[M+H]+	L-Alanine	Amino acids	1.91E+06	7.88E+05	4.31E+05	4.15E+05		
Mad233	Positive	205.1	146.1	2.36	204.09	[M+H]+	L-Tryptophan	Amino acids	7.20E+06	4.59E+06	1.46E+06	2.59E+06		
Mad12	Positive	133	70	0.65	132.16	[M+H]+	L(+) -Ornithine	Amino acids	7.40E+06	5.77E+06	4.60E+06	6.96E+06		
Mad88	Positive	118	88	0.85	117.079	[M+H]+	Dl-Norvaline	Amino acids	2.07E+07	9.47E+06	6.03E+06	9.56E+06		
Mad82	Positive	269	134	0.81	268.055	[M+H]+	L-Homocystine	Amino acids	7.44E+05	6.25E+04	1.19E+04	7.31E+03		
Fatty acids	Mad746	Positive	496.3	478.3	8.65	495.3	[M+H]+	LysoPC 16:0	Lipids_Glycerophospholipids	6.94E+06	1.43E+06	2.24E+06	6.26E+06	
	Mad1344	Negative	452	255.3	8.57	453	[M-H]-	LysoPE 16:0	Lipids_Glycerophospholipids	2.42E+06	6.06E+05	1.20E+06	2.37E+06	
	Mad741	Positive	496.33	478.3	8.49	495.33	[M+H]+	LysoPC 16:0 (2n isomer)	Lipids_Glycerophospholipids	7.59E+06	1.48E+06	2.25E+06	6.63E+06	
	Mad764	Positive	524.36	507.4	9.57	523.36	[M+H]+	LysoPC 18:0	Lipids_Glycerophospholipids	1.37E+06	2.01E+05	3.09E+05	1.32E+06	
	Mad1343	Negative	452.3	255.3	8.4	453.3	[M-H]-	LysoPE 16:0 (2n isomer)	Lipids_Glycerophospholipids	3.92E+06	9.37E+05	1.81E+06	3.57E+06	
	Mad759	Positive	295.2	179.5	9.17	294.2	[M+H]+	9-Hydroxy-(10E,12Z,15Z)-octadecatrienoic acid	Lipids_Fatty acids	1.05E+05	2.19E+04	4.54E+04	1.88E+05	
	Mad751	Positive	279.3	201.5	8.81	278.3	[M+H]+	Punicic acid	Lipids_Fatty acids	2.24E+05	4.24E+04	6.87E+04	2.55E+05	
	Mad762	Positive	279.2	149.4	9.55	278.2	[M+H]+	Octadeca-11E,13E,15Z-trienoic acid	Lipids_Fatty acids	9.28E+06	4.46E+06	5.58E+06	6.61E+06	
	Mad682	Positive	318.3	300.9	6.63	317.3	[M+H]+	4-Hydroxysphinganine	Lipids_Fatty acids	9.13E+06	1.12E+07	1.19E+07	1.03E+07	
Organic acids	Mad1350	Negative	199	181.3	9.52	200	[M-H]-	Lauric acid (C12:0)	Lipids_Fatty acids	3.71E+03	3.24E+03	3.24E+03	3.16E+03	
	Mad1357	Negative	277.1	233.7	10.47	278.1	[M-H]-	α -Linolenic acid	Lipids_Fatty acids	7.76E+03	9.00E+00	9.00E+00	9.00E+00	
	Mad994						L(-)-Malic acid	Organic acids	2.23E+07	2.11E+07	1.75E+07	1.32E+07		
	Mad861	Negative	147	85	1	148	[M-H]-	Citramalate	Organic acids	1.07E+07	2.17E+06	9.66E+06	2.45E+07	

Mad370	Positive	627.1	447.3	3.33	626.1	[M+H]+	6-C-hexosyl-hesperetin O-hexoside	Flavone C-glycosides	9.04E+03	9.40E+04	1.27E+05	1.58E+05		
Mad1156	Negative	739.2	740	3.85	740.2	[M-H]-	Chrysoeriol 8-C-pentosyl-O-rutinoside	Flavone C-glycosides	6.58E+03	1.98E+04	2.34E+04	2.13E+04		
Mad1088	Negative	449.1	329.3	3.21	450.1	[M-H]-	Eriodictyol C-hexoside	Flavone C-glycosides	2.47E+06	6.55E+05	4.20E+05	4.43E+05		
Mad1097	Negative	447.1	327.3	3.32	448.1	[M-H]-	Luteolin C-hexoside	Flavone C-glycosides	2.48E+04	4.98E+04	2.30E+04	2.91E+04		
Flavonol	Mad386	Positive	713.1	303.6	3.41	712.1	[M+H]+	Quercetin 7-O-malonylhexosyl-hexoside	Flavonol	1.89E+04	4.69E+03	1.96E+03	3.86E+04	
	Mad443	Positive	611.2	303.7	3.73	610.2	[M+H]+	Quercetin 7-O-rutinoside	Flavonol	2.62E+06	1.62E+06	1.32E+06	5.61E+06	
	Mad1115	Negative	609.153	609.1	3.51	610.153	[M-H]-	Quercetin 3-O-rutinoside (Rutin)	Flavonol	3.01E+06	1.66E+06	1.33E+06	6.64E+06	
	Mad1251	Negative	301.043	151.1	4.92	302.043	[M-H]-	Quercetin	Flavonol	1.11E+07	1.59E+05	3.51E+04	5.35E+04	
	Mad629	Positive	287.048	287	5.52	286.048	[M+H]+	Kaempferol	Flavonol	4.14E+04	9.00E+00	9.00E+00	9.00E+00	
	Mad1147	Negative	593.159	285.1	3.77	594.1585	[M-H]-	Kaempferol 3-O-rutinoside (Nicotiflorin)	Flavonol	2.30E+05	7.14E+04	5.00E+04	1.76E+05	
	Mad1218	Negative	317	179	4.39	318.038	[M-H]-	Myricetin	Flavonol	2.09E+04	6.51E+03	5.64E+03	4.60E+03	
	Mad1184	Negative	447.101	287.1	4.03	448.101	[M-H]-	Kaempferol 3-O-glucoside (Astragalin)	Flavonol	4.27E+05	2.27E+05	1.38E+05	2.27E+05	
	Mad1220	Negative	287.06	125.1	4.43	288.063	[M-H]-	Aromadedrin (Dihydrokaempferol)	Flavonol	2.74E+05	9.00E+00	9.00E+00	9.00E+00	
	Mad1185	Negative	463	301	4.03	464.096	[M-H]-	Quercetin 4'-O-glucoside (Spiraeoside)	Flavonol	3.91E+07	3.57E+07	2.60E+07	2.91E+07	
	Mad432	Positive	465	302.8	3.68	464.096	[M+H]+	Quercetin 3-O-glucoside (Isotrifolin)	Flavonol	4.94E+07	4.64E+07	4.17E+07	4.24E+07	
	Mad465	Positive	449	286.9	3.86	448.101	[M+H]+	Kaempferol 3-O-galactoside (Trifolin)	Flavonol	5.56E+06	1.26E+06	5.55E+05	1.20E+06	
	Mad1208	Negative	431	285.1	4.29	432.106	[M-H]-	Kaempferol 3-O-rhamnoside (Kaempferin)	Flavonol	7.22E+06	1.60E+06	7.86E+05	1.15E+06	
	Mad363	Positive	741.22	433.2	3.28	740.216	[M+H]+	Kaempferol-3-O-robinoside-7-O-rhamnoside (Robinin)	Flavonol	9.00E+00	4.94E+04	6.00E+04	7.22E+04	
Flavanone	Mad508	Positive	611.2	303.1	4.09	610.19	[M+H]+	Hesperetin 7-O-neohesperidoside (Neohesperidin)	Flavanone	3.96E+05	1.24E+05	8.95E+04	1.45E+05	
	Mad481	Positive	581.179	57.1	3.92	580.1792	[M+H]+	Naringenin 7-O-neohesperidoside (Naringin)	Flavanone	6.61E+04	2.89E+04	2.01E+04	2.26E+04	
	Mad1187	Negative	433.121	122.9	4.05	434.1213	[M-H]-	Naringenin 7-O-glucoside (Prunin)	Flavanone	9.04E+06	3.64E+06	1.53E+06	2.04E+06	
	Mad624	Positive	273.069	153.1	5.41	272.0685	[M+H]+	Naringenin	Flavanone	7.00E+05	9.00E+00	9.00E+00	4.27E+03	
	Mad623	Positive	275.084	107	5.34	274.0841	[M+H]+	Phloretin	Flavanone	1.40E+06	1.99E+04	7.82E+03	9.40E+03	
	Mad595	Positive	289.063	153.1	4.85	288.063	[M+H]+	Eriodictyol	Flavanone	1.37E+05	9.00E+00	9.00E+00	9.00E+00	
	Mad1165	Negative	463.132	301.1	3.88	464.132	[M-H]-	Hesperetin 5-O-glucoside	Flavanone	7.60E+07	6.51E+07	5.49E+07	6.02E+07	

Proanthocyanidins & Anthocyanins	Mad490	Positive	611.19	303	3.99	610.19	[M+H]+	Hesperetin 7-rutinoside (Hesperidin)	Flavanone	4.66E+05	1.48E+05	9.18E+04	1.72E+05
	Mad1268	Negative	271.07	151.1	5.3	272.069	[M-H]-	Naringenin chalcone	Flavanone	7.30E+05	3.02E+03	9.00E+00	7.92E+03
	Mad1246	Negative	593	285.2	4.85	594.195	[M-H]-	Isosakuranetin-7-neohesperidoside (Poncirin)	Flavanone	3.94E+06	1.86E+05	2.05E+04	3.20E+04
	Mad1133	Negative	575	285.3	3.69	576.1268	[M-H]-	Procyanidin A1	Proanthocyanidins	2.75E+05	9.00E+00	9.00E+00	9.00E+00
	Mad1175	Negative	575	285.3	3.94	576.1268	[M-H]-	Procyanidin A2	Proanthocyanidins	9.00E+00	7.99E+04	5.22E+04	7.82E+04
	Mad1076	Negative	577	407.3	3.08	578.1424	[M-H]-	Procyanidin B2	Proanthocyanidins	4.63E+07	4.11E+07	3.44E+07	3.47E+07
	Mad1058	Negative	577.1	407.3	2.87	578.1424	[M-H]-	Procyanidin B3	Proanthocyanidins	4.35E+07	3.89E+07	3.20E+07	3.25E+07
	Mad356	Positive	477.1	315.6	3.22	477.1	Protonated	Rosinidin O-hexoside	Anthocyanins	3.77E+04	4.36E+03	9.00E+00	9.00E+00
	Mad248	Positive	449.1	287.3	2.45	449.1	Protonated	Cyanidin 3-O-glucoside (Kuromarin)	Anthocyanins	2.96E+07	1.86E+05	7.15E+05	3.72E+07
	Mad1017	Negative	465.1	285.3	2.41	466.1	[M-H]-	Cyanidin O-syringic acid	Anthocyanins	3.73E+06	1.85E+04	1.35E+05	1.16E+07
Catechin derivatives	Mad1111	Negative	865.1	407.2	3.46	866.1	[M-H]-	Catechin-catechin-catechin	Catechin derivatives	8.68E+06	9.27E+05	3.08E+05	4.15E+05
	Mad1116	Negative	561.1	271.3	3.55	562.1	[M-H]-	Epicatechin-epiafzelechin	Catechin derivatives	1.13E+06	1.23E+06	8.25E+05	9.32E+05
	Mad1054	Negative	289.079	125	2.81	290.079	[M-H]-	Catechin	Catechin derivatives	5.07E+07	8.51E+06	4.37E+06	5.11E+06
	Mad1082	Negative	289	78.8	3.17	290.3	[M-H]-	L-Epicatechin	Catechin derivatives	6.15E+07	4.24E+07	3.06E+07	3.01E+07
	Mad277	Positive	307	139.1	2.64	306	[M+H]+	Epigallocatechin (EGC)	Catechin derivatives	7.46E+05	4.50E+05	2.40E+05	5.06E+05
	Mad1015	Negative	305	125	2.36	306.074	[M-H]-	(+)-Gallocatechin (GC)	Catechin derivatives	1.13E+05	1.25E+04	2.06E+04	2.57E+04
	Mad1034	Negative	153.1	109.1	2.55	154.027	[M-H]-	Protocatechuic acid	Catechin derivatives	2.23E+06	1.97E+05	1.40E+05	2.10E+05
	Mad1071	Negative	137.1	137	2.99	138.032	[M-H]-	Protocatechuic aldehyde	Catechin derivatives	1.02E+05	1.98E+04	1.69E+04	2.10E+04

Table 3. Differentially accumulated metabolites among PS1 vs PS2, PS2 vs PS3, and PS3 vs PS4, respectively.

Index	Compounds	Class	Sample 1			Sample 2			VIP	Fold_Change	LogFC	type
PS1 vs PS2:			PS1a	PS1b	PS1c	PS2a	PS2b	PS2c				
Mad456	Syringetin 5-O-hexoside	Flavone	9.00E+00 0	9.00E+00 0	9.00E+00 0	1.64E+00 4	1.26E+00 4	3.21E+00 4	2.07758 4	2262.963	11.144	up
Mad493	Syringetin 7-O-hexoside	Flavone	9.00E+00 0	9.00E+00 0	4.32E+00 3	1.72E+00 4	1.00E+00 4	1.88E+00 4	1.54566 5	10.60396	3.40653 2	up
Mad370	6-C-hexosyl-hesperetin O-hexoside	Flavone C-glycosides	1.18E+00 4	9.51E+00 3	5.82E+00 3	1.45E+00 5	7.71E+00 4	5.99E+00 4	1.12152 1	10.3944	3.37773 4	up

Mad424	Luteolin 8-C-hexosyl-O-hexoside	Flavone C-glycosides	2.09E+0 4 0	9.00E+0 4 4	1.44E+0 4 4	5.62E+0 4 4	4.04E+0 4 4	6.24E+0 4 3	1.09370 3	4.503101 2	2.17091 9	up
Mad331	O-Feruloyl 2-hydroxylcoumarin	Coumarins	1.20E+0 4 4	1.42E+0 4 4	1.37E+0 4 4	1.18E+0 5 5	1.04E+0 5 5	1.03E+0 5 5	1.08883 2	8.145363 2	3.02597 9	up
Mad788	4-Hydroxy-3-methoxymandelate	Organic acids	9.00E+0 0 3	4.94E+0 0 0	9.00E+0 4 4	1.23E+0 4 4	1.58E+0 4 4	1.70E+0 4 4	1.53425 1	9.09641 1	3.18529 7	up
Mad115 1	Acetyl tryptophan	Amino acid derivatives	5.53E+0 5 5	3.70E+0 5 5	3.35E+0 5 6	5.94E+0 6 7	1.02E+0 7 7	8.27E+0 6 6	1.28930 2	19.40382 2	4.27826 8	up
Mad825	Anthranilate O-hexosyl-O-hexoside	Benzoic acid derivatives	1.68E+0 4 4	1.48E+0 4 4	1.50E+0 4 4	7.50E+0 4 4	1.42E+0 5 5	1.22E+0 5 5	1.04089 6	7.274678 6	2.86288 3	up
Mad116 8	Tricin 5-O-hexoside	Flavone	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	4.20E+0 4 4	2.78E+0 4 4	9.54E+0 4 4	2.20062 4	6118.519 4	12.5789 7	up
Mad28	2'-Deoxyinosine-5'-monophosphate	Nucleotide and its derivates	1.44E+0 5 5	1.77E+0 5 5	2.05E+0 5 7	1.18E+0 6 6	9.82E+0 6 6	6.99E+0 6 6	1.4988 4	54.39163 3	5.76531 3	up
Mad567	Ethyl 3,4-Dihydroxybenzoate (Ethyl protocatechuic)	Organic acids	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	8.65E+0 4 5	1.23E+0 5 5	1.45E+0 5 5	2.31398 7	13129.63 7	13.6805 4	up
Mad117 5	Procyanidin A2	Proanthocyanidins	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	9.98E+0 4 4	6.26E+0 4 4	7.74E+0 4 4	2.26627 8	8881.481 8	13.1165 8	up
Mad828	D-(+)-Sucrose	Carbohydrates	1.15E+0 6 5	7.58E+0 5 6	1.03E+0 6 6	8.94E+0 6 7	1.08E+0 7 7	8.06E+0 6 6	1.12527 7	9.462219 7	3.24217 9	up
Mad51	D(+)-Melezitose	Carbohydrates	2.62E+0 4 4	2.08E+0 4 4	1.74E+0 4 5	2.72E+0 5 5	2.00E+0 5 5	1.69E+0 5 5	1.13255 3	9.953416 3	3.31519 2	up
Mad105	L-Methionine	Amino acids	4.96E+0 5 5	6.17E+0 5 5	2.91E+0 5 6	3.88E+0 6 6	4.14E+0 6 6	5.60E+0 6 6	1.12976 1	9.700855 2	3.27811 up	up
Mad137	Pyridoxine 5'-phosphate	Vitamins	7.78E+0 3 0	9.00E+0 3 0	3.32E+0 3 4	1.28E+0 4 4	1.93E+0 4 4	1.71E+0 4 4	1.06268 9	4.428841 9	2.14692 up	up
Mad102 9	N-Acetyl-L-tyrosine	Amino acid derivatives	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	4.71E+0 4 4	4.68E+0 4 4	5.77E+0 4 4	2.21032 6	5614.815 2	12.4550 up	up
Mad850	1,5-Anhydro-D-glucitol	Alcohols and polyols	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	7.06E+0 4 4	7.99E+0 4 4	7.22E+0 4 4	2.25965 3	8248.148 5	13.0098 up	up
Mad45	1-Methyladenine	Nucleotide and its derivates	2.14E+0 4 4	1.85E+0 3 3	8.95E+0 4 4	7.38E+0 5 5	1.12E+0 5 5	1.36E+0 5 5	1.01529	6.587513 4	2.71973 4	up
Mad778	trans,trans-Muconic acid	Organic acids	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	3.99E+0 5 5	4.06E+0 5 5	4.55E+0 5 5	2.46720 6	46666.67 up	15.5101 up	up
Mad110 3	Glycitin	Isoflavone	2.25E+0 4 4	3.28E+0 0 0	9.00E+0 0 0	2.62E+0 5 5	3.29E+0 5 5	2.50E+0 5 5	1.39221 3	15.20548 up	3.92652 up	up
Mad347	Esculetin (6,7-dihydroxycoumarin)	Coumarins	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	1.00E+0 5 4	9.21E+0 4 4	7.04E+0 4 4	2.27877 2	9722.222 7	13.2470 up	up
Mad388	6-Methoxy-7,8-DihydroxyCoumarin	Coumarins	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	3.11E+0 4 4	5.03E+0 4 4	2.81E+0 4 4	2.16381 1	4055.556 8	11.9856 up	up
Mad363	Kaempferol-3-O-robinoside-7-O-rhamnoside (Robinin)	Flavonol	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	5.73E+0 4 4	3.93E+0 4 4	5.17E+0 4 4	2.20629 6	5492.593 7	12.4232 up	up
Mad110 4	Myricetin 3-O-galactoside	Flavonol	5.64E+0 4 4	9.10E+0 0 0	9.00E+0 0 0	5.00E+0 4 4	1.94E+0 5 5	3.65E+0 5 5	1.05604 6	4.131362 8	2.04661 up	up
Mad739	14,15-Dehydrocrepeninic acid	Lipids_Fatty acids	1.19E+0 5 5	1.63E+0 5 5	7.56E+0 4 4	1.27E+0 4 4	1.11E+0 4 4	1.12E+0 4 4	1.12460 9	0.097875 -	- 3.35292	dow n
Mad390	1-O-Feruloyl quinic acid	Quinate and its derivatives	8.33E+0 5 6	1.32E+0 5 6	9.18E+0 5 5	6.12E+0 4 4	3.25E+0 4 4	7.47E+0 4 4	1.27904 2	0.054836 - 4.18874	- 4.18874	dow n
Mad108 4	5-O-p-Coumaroylquinic acid	Quinate and its derivatives	8.12E+0 6 7	1.03E+0 7 7	1.13E+0 7 7	8.61E+0 5 5	7.74E+0 5 5	8.66E+0 5 5	1.18010 7	0.084152 - 3.57086	- 3.57086	dow n

Mad122 5	E-3,4,5'-Trihydroxy-3'-glucopyranosylstilbene	Others	6.87E+0 4 4	1.52E+0 5 4	8.21E+0 4 4	5.19E+0 3 3	9.00E+0 0 3	9.00E+0 0 0	1.86407 3 3	0.017199 -	- 5.86149	dow n
Mad97	Xanthine	Nucleotide and its derivates	8.34E+0 4 4	8.93E+0 4 4	8.54E+0 4 4	5.12E+0 3 3	5.20E+0 3 3	9.00E+0 0 0	1.45357 0 0	0.040019 -	- 4.64316	dow n
Mad2	Spermidine	Phenolamides	2.52E+0 6 6	2.43E+0 6 6	2.79E+0 5 5	1.52E+0 5 5	2.48E+0 5 5	2.97E+0 5 5	1.16839 4 4	0.090052 -	-3.4731	dow n
Mad240	N',N'',N''''-p-coumaroyl-cinnamoyl-caffeoxyl spermidine	Phenolamides	9.11E+0 3 4	1.44E+0 3 4	6.14E+0 3 3	2.52E+0 3 3	9.00E+0 0 0	9.00E+0 0 0	1.51497 8 8	0.085599 -	- 3.54627	dow n
Mad228	N'-p-Coumaroyl putrescine	Phenolamides	4.86E+0 4 4	5.24E+0 4 4	6.51E+0 4 4	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	2.22166 3 3	0.000163 -	- 12.5868	dow n
Mad356	Rosinidin O-hexoside	Anthocyanins	5.28E+0 4 4	2.84E+0 4 4	3.20E+0 3 3	7.17E+0 3 3	9.00E+0 0 0	5.89E+0 3 3	1.21537 8 8	0.115451 -	- 3.11465	dow n
Mad248	Cyanidin 3-O-glucoside (Kuromarin)	Anthocyanins	2.90E+0 7 7	3.25E+0 7 7	2.72E+0 5 5	3.77E+0 4 4	9.62E+0 4 4	8.40E+0 4 4	1.72089 8 8	0.006282 -	-7.3146	dow n
Mad430	5-O-p-Coumaroyl shikimic acid	Quinate and its derivatives	2.14E+0 5 5	2.34E+0 5 5	1.43E+0 5 4	2.86E+0 4 3	7.97E+0 3 3	2.19E+0 4 4	1.13923 7 7	0.098934 -	- 3.33739	dow n
Mad163	Niacinamide	Vitamins	3.94E+0 5 5	3.25E+0 5 5	4.31E+0 5 4	7.06E+0 4 4	5.03E+0 4 4	6.72E+0 4 4	1.00837 7 7	0.163565 -	- 2.61206	dow n
Mad715	LysoPE 14:0	Lipids_Glycerophospholipids	7.70E+0 3 3	8.98E+0 3 3	7.20E+0 3 3	9.00E+0 0 0	1.91E+0 3 3	9.00E+0 0 0	1.51509 7 7	0.080737 -	- 3.63063	dow n
Mad420	Phellodensin F	Others	1.87E+0 4 4	3.32E+0 4 4	1.85E+0 4 0	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	2.10411 0 0	0.000384 -	- 11.3484	dow n
Mad46	O-Phosphocholine	Cholines	1.79E+0 6 6	1.90E+0 6 6	2.29E+0 5 5	1.21E+0 5 5	1.01E+0 5 5	1.49E+0 5 6	1.25369 6 6	0.06204 -	- 4.01065	dow n
Mad718	LysoPC 15:1	Lipids_Glycerophospholipids	1.28E+0 4 4	1.48E+0 4 4	1.31E+0 4 3	4.20E+0 3 3	3.45E+0 3 3	9.00E+0 0 0	1.08178 0 0	0.188182 -	-2.4098	dow n
Mad768	MAG (18:3) isomer1	Lipids_Glycerolipids	1.99E+0 6 6	2.18E+0 6 6	1.06E+0 5 5	1.57E+0 5 5	1.78E+0 5 5	2.25E+0 5 6	1.10160 6 6	0.107075 -	- 3.22331	dow n
Mad101 3	Syringaldehyde O-glucoside	Others	2.99E+0 4 4	3.87E+0 4 4	2.71E+0 4 0	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	2.14953 4 4	0.000282 -	- 11.7913	dow n
Mad127 9	3,4-Dimethoxycinnamic acid	Hydroxycinnamoyl derivatives	7.17E+0 4 5	1.19E+0 4 4	4.72E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	2.25715 1 1	0.000113 -	- 13.1051	dow n
Mad134 7	9,10-EODE	Lipids_Fatty acids	3.86E+0 5 5	5.38E+0 5 5	2.89E+0 5 4	6.02E+0 4 4	6.14E+0 4 4	6.40E+0 4 8	1.01266 8 8	0.153009 -	- 2.70831	dow n
Mad134 9	9-KODE	Lipids_Fatty acids	1.11E+0 4 4	1.16E+0 4 3	7.18E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	1.98811 1 1	0.000904 -	-10.112	dow n
Mad133 9	13-HpOTrE(r)	Lipids_Fatty acids	4.80E+0 3 3	4.49E+0 3 3	1.68E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	1.82268 3 3	0.002461 -	- 8.66639	dow n
Mad127 2	4-Methoxycinnamic acid	Hydroxycinnamoyl derivatives	3.58E+0 5 5	6.46E+0 5 5	2.86E+0 5 0	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	2.46184 8 8	2.09E-05 -	- 15.5441	dow n
Mad135 2	12,13-EODE	Lipids_Fatty acids	5.21E+0 3 3	5.42E+0 3 3	3.39E+0 3 0	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	1.87728 4 4	0.001926 -	- 9.02031	dow n
Mad133 8	13-HPODE	Lipids_Fatty acids	1.42E+0 4 4	2.71E+0 4 4	1.36E+0 4 3	3.06E+0 3 3	2.71E+0 3 3	2.44E+0 3 2	1.00858 2 2	0.149545 -	- 2.74135	dow n
Mad111 1	Catechin-catechin-catechin	Catechin derivatives	8.45E+0 6 6	9.28E+0 6 6	8.30E+0 6 6	1.37E+0 5 5	6.87E+0 5 5	7.24E+0 5 9	1.12743 9 9	0.106838 -	-3.2265	dow n
Mad101 7	Cyanidin O-syringic acid	Anthocyanins	3.78E+0 6 6	4.19E+0 6 6	3.23E+0 6 4	5.54E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	2.21533 8 8	0.004948 -	- 7.65893	dow n
Mad116 1	Quercetin O-acetylhexoside	Flavonol	1.97E+0 7 7	2.21E+0 7 7	2.23E+0 7 6	4.47E+0 6 6	3.31E+0 6 6	2.93E+0 6 6	1.00540 9 9	0.167083 -	- 2.58137	dow n

Mad110 5	O-p-Coumaroyl quinic acid O-rutinoside derivative	Quinate and its derivatives	1.11E+0 4 4	1.65E+0 3 3	9.55E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	2.01816 8 8	0.000727 1.07991 1.07991	- 10.4262 -	dow n dow
Mad110 6	3-O-p-Coumaroyl quinic acid	Quinate and its derivatives	3.34E+0 7 7	4.12E+0 7 7	3.83E+0 6 6	6.94E+0 6 6	4.86E+0 6 6	2.55E+0 6 4	1.07991 4 4	0.127104 -	2.97592 -	n dow
Mad116 9	3-O-p-Coumaroyl shikimic acid	Quinate and its derivatives	3.09E+0 4 4	5.77E+0 4 4	3.75E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	2.18194 2 2	0.000214 -	12.1893 -	n dow
Mad135 7	α -Linolenic acid	Lipids_Fatty acids	8.62E+0 3 3	9.80E+0 3 3	4.86E+0 3 3	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	1.94812 9 9	0.00116 -	9.75192 -	dow n
Mad764	LysoPC 18:0	Lipids_Glycerophospholipids	1.05E+0 6 6	2.07E+0 6 5	9.78E+0 5 5	2.34E+0 5 5	1.95E+0 5 5	1.74E+0 5 2	1.00900 2 2	0.147145 -	2.76469 -	dow n
Mad124 9	Luteolin	Flavone	6.58E+0 4 4	8.09E+0 4 4	6.17E+0 4 4	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	2.25047 7 7	0.00013 -	12.9141 -	dow n
Mad125 1	Quercetin	Flavonol	1.10E+0 7 7	1.06E+0 7 7	1.17E+0 5 5	2.07E+0 5 5	1.42E+0 5 5	1.27E+0 5 5	1.55302 9 9	0.014294 -	6.12842 -	n dow
Mad629	Kaempferol	Flavonol	3.73E+0 4 4	4.62E+0 4 4	4.07E+0 4 4	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	2.18487 8 8	0.000217 -	12.1674 -	dow n
Mad105 4	Catechin	Catechin derivatives	5.12E+0 7 7	5.00E+0 7 7	5.09E+0 7 7	1.04E+0 7 7	7.06E+0 6 6	8.08E+0 6 6	1.00495 3 3	0.167916 -	2.57419 -	dow n
Mad969	Glutaric acid	Organic acids	1.70E+0 6 6	1.44E+0 6 6	1.34E+0 6 6	9.00E+0 0 0	9.00E+0 0 0	1.14E+0 6 6	1.81806 6 6	0.254468 -	1.97444 -	dow n
Mad786	2,6-Diaminooimelic acid	Amino acid derivatives	4.34E+0 6 6	6.48E+0 6 6	4.86E+0 6 6	4.58E+0 5 5	3.02E+0 5 5	2.63E+0 5 5	1.23972 2 2	0.065242 -	3.93805 -	dow n
Mad108 0	Caffeic acid	Hydroxycinnamoyl derivatives	3.08E+0 6 6	2.92E+0 6 6	3.03E+0 6 6	1.27E+0 5 5	9.20E+0 4 4	7.92E+0 4 4	1.39135 8 8	0.033023 -	4.92037 -	dow n
Mad117 0	Ferulic acid	Hydroxycinnamoyl derivatives	1.25E+0 5 5	1.26E+0 5 5	1.69E+0 5 5	2.99E+0 4 4	1.69E+0 4 4	2.30E+0 4 4	1.00200 7 7	0.16619 -	2.58909 -	dow n
Mad688	Chrysin	Flavone	5.59E+0 3 4	1.48E+0 3 3	8.38E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	1.97237 7 7	0.000938 -	10.0574 -	dow n
Mad444	Apigenin 5-O-glucoside	Flavone	1.13E+0 6 6	1.29E+0 6 6	1.35E+0 6 6	2.42E+0 5 5	1.70E+0 5 5	1.79E+0 5 5	1.02283 3 3	0.156764 -	2.67333 -	dow n
Mad631	Chrysoeriol	Flavone	1.07E+0 4 3	7.15E+0 3 3	9.65E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	1.97781 1 1	0.000982 -	9.99226 -	dow n
Mad624	Naringenin	Flavanone	6.11E+0 5 5	9.39E+0 5 5	5.51E+0 5 5	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	2.52183 0 0	1.29E-05 -	16.2478 -	dow n
Mad401	Chlorogenic acid methyl ester	Quinate and its derivatives	4.77E+0 6 6	5.11E+0 6 6	5.44E+0 6 6	1.80E+0 5 5	1.80E+0 5 5	1.81E+0 5 5	1.37563 2 2	0.035313 -	4.82364 -	dow n
Mad623	Phloretin	Flavanone	1.41E+0 6 6	1.31E+0 6 6	1.48E+0 6 6	2.74E+0 4 4	1.88E+0 4 4	1.35E+0 4 4	1.55567 4 4	0.014214 -	6.13651 -	dow n
Mad113 3	Procyanidin A1	Proanthocyanidins	2.65E+0 5 5	2.62E+0 5 5	2.99E+0 5 5	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	2.41829 6 6	3.27E-05 -	14.9009 -	dow n
Mad4	Gluconic acid	Carbohydrates	4.18E+0 4 4	2.98E+0 4 4	2.90E+0 4 4	9.00E+0 0 0	9.00E+0 0 0	1.66E+0 4 4	1.55885 9 9	0.165189 -	2.59781 -	dow n
Mad109	Guanine	Nucleotide and its derivates	2.51E+0 5 5	2.70E+0 5 5	3.76E+0 5 5	4.08E+0 4 4	3.27E+0 4 4	3.56E+0 4 4	1.08468 8 8	0.121628 -	3.03946 -	dow n
Mad962	Gallic acid	Benzoic acid derivatives	1.42E+0 5 5	1.30E+0 5 5	1.46E+0 5 5	2.28E+0 4 4	9.00E+0 0 0	9.00E+0 0 0	1.78811 6 6	0.054589 -	4.19526 -	dow n
Mad180	Deoxyguanosine	Nucleotide and its derivates	7.17E+0 4 4	5.78E+0 4 4	5.36E+0 4 4	1.55E+0 4 4	1.06E+0 4 4	9.00E+0 0 0	1.19197 7 7	0.142594 -	2.81001 -	dow n
Mad101	Deoxycytidine	Nucleotide and its derivates	6.28E+0 4 4	5.87E+0 4 4	4.62E+0 4 4	9.00E+0 0 0	1.39E+0 4 4	9.00E+0 0 0	1.67227 2 2	0.082993 -	3.59086 -	dow n

Mad105		Syringin	Hydroxycinnamoyl derivatives	9.22E+0 4	1.10E+0 5	7.23E+0 4	9.00E+0 0	9.00E+0 0	9.00E+0 0	2.28365 4	9.84E-05 -	-	dow
3		p-Coumaric acid	Hydroxycinnamoyl derivatives	3.52E+0 6	3.74E+0 6	4.70E+0 6	1.56E+0 5	1.13E+0 5	6.63E+0 4	1.42530 2	0.028035 -	-	dow
Mad113		Dihydroquercetin (Taxifolin)	Flavonol	7.46E+0 5	7.92E+0 5	5.19E+0 5	1.03E+0 4	9.97E+0 3	9.36E+0 3	1.54499 2	0.014404 -	-	dow
7		(+)-Gallocatechin (GC)	Catechin derivatives	1.11E+0 5	1.42E+0 5	8.67E+0 4	7.90E+0 3	1.32E+0 4	1.65E+0 4	1.10789 4	0.110686 -	-	dow
Mad117		Orobol (5,7,3',4'-tetrahydroxyisoflavone)	Isoflavone	1.23E+0 5	5.08E+0 4	2.21E+0 5	3.33E+0 3	9.00E+0 0	9.00E+0 0	1.92653 8	0.00848 -	-	dow
6		Eriodictyol	Flavanone	1.08E+0 5	2.10E+0 4	9.44E+0 0	9.00E+0 0	9.00E+0 0	9.00E+0 0	2.32642 3	6.55E-05 -	-	dow
Mad101		(+)-Jasmonic acid (JA)	Phytohormones	6.09E+0 5	6.02E+0 5	4.49E+0 5	3.77E+0 4	2.22E+0 4	2.29E+0 4	1.30193 7	0.04988 -	-	dow
5		Protocatechuic acid	Catechin derivatives	2.25E+0 6	2.38E+0 6	2.05E+0 5	2.90E+0 5	1.18E+0 5	1.83E+0 5	1.17466 6	0.088473 -	-	dow
Mad103		N-[-(Jasmonoyl)-(L)-Isoleucine (JA-L-Ile)]	Phytohormones	2.17E+0 5	2.06E+0 5	1.62E+0 5	5.29E+0 3	2.57E+0 3	6.44E+0 3	1.45354 6	0.024444 -	-	dow
4		L-Carnitine	Others	4.20E+0 4	3.87E+0 4	4.06E+0 4	3.38E+0 4	9.00E+0 0	9.00E+0 0	1.51812 1	0.278796 -	-	dow
Mad43		Fumaric acid	Organic acids	1.52E+0 6	1.08E+0 6	2.33E+0 6	9.00E+0 0	9.00E+0 0	9.00E+0 0	2.61316 6	5.48E-06 -	-	dow
Mad919		A-Ketoglutaric acid	Organic acids	2.36E+0 6	3.41E+0 6	2.52E+0 6	4.47E+0 5	4.02E+0 5	3.68E+0 5	1.03518 2	0.146803 -	-	dow
Mad903		Luteolin 7-O-glucoside (Cynaroside)	Flavone	2.70E+0 6	2.51E+0 6	2.91E+0 6	3.77E+0 5	2.71E+0 5	3.95E+0 5	1.07734 9	0.128448 -	-	dow
Mad113		Vitamin D3	Vitamins	4.21E+0 4	6.20E+0 4	4.92E+0 4	9.00E+0 0	9.00E+0 0	9.00E+0 0	2.21062 8	0.000176 -	-	dow
6		2'-Deoxyinosine	Nucleotide and its derivates	6.35E+0 4	8.04E+0 4	6.89E+0 4	1.19E+0 4	7.19E+0 3	1.11E+0 4	1.04904 8	0.14187 -	-	dow
Mad127		L-Homocystine	Amino acids	4.32E+0 5	7.70E+0 5	1.03E+0 6	1.16E+0 5	3.17E+0 4	3.97E+0 4	1.17330 2	0.083961 -	-	dow
1		Dihydromyricetin	Flavonol	5.23E+0 5	5.75E+0 5	2.81E+0 5	4.18E+0 4	9.00E+0 0	9.00E+0 0	1.92051 5	0.030325 -	-	dow
Mad376		4-Hydroxybenzoic acid	Organic acids	4.41E+0 5	2.91E+0 5	3.44E+0 5	1.88E+0 5	1.53E+0 5	9.00E+0 0	1.08637 1	0.316923 -	-	dow
Mad106		3-Hydroxy-3-methylpentane-1,5-dioic acid	Amino acid derivatives	5.37E+0 6	8.85E+0 6	5.35E+0 6	2.26E+0 5	2.92E+0 5	2.09E+0 5	1.35678 8	0.037149 -	-	dow
8		Naringenin chalcone	Flavanone	6.32E+0 5	1.01E+0 6	5.49E+0 5	9.03E+0 3	9.00E+0 0	9.00E+0 0	2.12425 4	0.00413 -	-	dow
Mad126		Aromadetrin (Dihydrokaempferol)	Flavonol	2.48E+0 5	2.94E+0 5	2.81E+0 5	9.00E+0 0	9.00E+0 0	9.00E+0 0	2.41777 8	3.28E-05 -	-	dow
0		Isosakuranetin-7-neohesperidoside (Poncirin)	Flavanone	3.61E+0 6	4.34E+0 6	3.88E+0 6	3.15E+0 5	1.15E+0 5	1.27E+0 5	1.32469 2	0.047084 -	-	dow
Mad124		5-oxoproline	Amino acid derivatives	6.45E+0 6	1.35E+0 7	8.53E+0 6	7.29E+0 5	1.07E+0 6	1.48E+0 6	1.08525 7	0.115133 -	-	dow
6		2-(Formylamino)benzoic acid	Organic acids	3.13E+0 5	3.07E+0 5	3.75E+0 5	9.00E+0 0	9.00E+0 0	9.00E+0 0	2.43968 8	2.71E-05 -	-	dow
Mad104		Coniferin	Hydroxycinnamoyl derivatives	1.45E+0 7	1.72E+0 7	1.31E+0 7	2.57E+0 6	1.42E+0 6	1.52E+0 6	1.08746 8	0.122991 -	-	dow
1												3.02337	n

Mad124 1	2'-Hydroxygenistein	Isoflavone	1.95E+0 4	2.85E+0 4	1.86E+0 4	9.00E+0 0	9.00E+0 0	9.00E+0 0	2.10000 5	0.000405	-	dow
Mad113 2	Quercetin 7-O- β -D-Glucuronide	Flavonol	2.74E+0 4	2.35E+0 4	2.57E+0 4	9.00E+0 0	9.00E+0 0	3.11E+0 3	1.68967 5	0.040836	-	dow
Mad528	p-Coumaraldehyde	Hydroxycinnamoyl derivatives	3.31E+0 5	5.58E+0 5	4.05E+0 5	9.00E+0 0	9.00E+0 0	9.00E+0 0	2.46742 2	2.09E-05	-	dow
Mad626	Butin	Flavone	5.96E+0 5	9.25E+0 5	6.15E+0 5	9.00E+0 0	9.00E+0 0	9.00E+0 0	2.52461 2	1.26E-05	-	dow
Mad105 1	Pantetheine	Vitamins	2.05E+0 4	2.56E+0 4	4.17E+0 3	9.00E+0 0	5.08E+0 3	3.18E+0 3	1.04645 1	0.164492	-	dow
Mad125 4	Morin	Flavonol	1.14E+0 7	1.12E+0 7	1.26E+0 7	2.15E+0 5	1.53E+0 5	1.31E+0 5	1.55432 3	0.014176	-	dow
Mad198	Deoxyadenosine	Nucleotide and its derivates	8.36E+0 5	8.97E+0 5	7.02E+0 5	1.36E+0 4	8.86E+0 4	9.66E+0 4	1.06857 8	0.13191	-	dow
Mad219	2-(dimethylamino)guanosine	Nucleotide and its derivates	2.58E+0 5	3.53E+0 5	2.65E+0 5	3.70E+0 4	2.73E+0 4	3.21E+0 4	1.11334 5	0.110046	-	dow
PS2 vs PS3:			PS2a	PS2b	PS2c	PS3a	PS3b	PS3c				

Mad759	9-Hydroxy-(10E,12Z,15Z)-octadecatrienoic acid	Lipids_Fatty acids	2.05E+0 4	2.36E+0 4	2.17E+0 4	4.66E+0 4	5.18E+0 4	3.78E+0 4	1.00813	2.069909	1.04956 7	up
Mad359	di-C,C-hexosyl-luteolin	Flavone C-glycosides	9.00E+0 0	2.22E+0 3	9.00E+0 0	6.66E+0 3	9.00E+0 0	2.72E+0 3	1.12605 7	4.195264	2.06876 1	up
Mad228	N'-p-Coumaroyl putrescine	Phenolamides	9.00E+0 0	9.00E+0 0	9.00E+0 0	9.00E+0 0	9.78E+0 3	9.00E+0 0	1.23457 1	362.8889	8.50338 4	up
Mad248	Cyanidin 3-O-glucoside (Kuromarin)	Anthocyanins	3.77E+0 5	9.62E+0 4	8.40E+0 4	2.54E+0 5	1.49E+0 6	4.00E+0 5	1.13274 6	3.84781	1.94403 8	up
Mad690	Azoxystrobin acid	Others	4.45E+0 4	8.10E+0 4	4.97E+0 4	1.18E+0 5	1.35E+0 5	1.78E+0 5	1.10328 6	2.460046	1.29868 5	up
Mad100	Piperidine	Alkaloids	2.45E+0 5	4.19E+0 5	5.20E+0 5	1.29E+0 6	8.42E+0 5	8.38E+0 5	1.09859 1	2.508446	1.32679 4	up
Mad722	LysoPC 16:1 (2n isomer)	Lipids_Glycerophospholipids	5.29E+0 4	6.19E+0 4	6.27E+0 4	1.05E+0 5	1.50E+0 5	1.25E+0 5	1.02669 6	2.140845	1.09818	up
Mad704	LysoPC 16:2 (2n isomer)	Lipids_Glycerophospholipids	9.18E+0 3	9.37E+0 3	9.07E+0 3	2.92E+0 4	4.13E+0 4	3.86E+0 4	1.40430 3	3.950036	1.98186 6	up
Mad715	LysoPE 14:0	Lipids_Glycerophospholipids	9.00E+0 0	1.91E+0 3	9.00E+0 0	2.63E+0 3	2.57E+0 3	3.89E+0 3	2.09661 5	4.71473	2.23717 5	up
Mad771	MAG (18:2)	Lipids_Glycerolipids	6.90E+0 4	9.88E+0 4	7.99E+0 4	1.98E+0 5	2.70E+0 5	1.59E+0 5	1.11501 9	2.531288	1.33987 2	up
Mad420	Phellodensin F	Others	9.00E+0 0	9.00E+0 0	9.00E+0 0	4.11E+0 3	9.00E+0 0	9.00E+0 0	1.15663 7	152.8889	7.25634	up
Mad718	LysoPC 15:1	Lipids_Glycerophospholipids	4.20E+0 3	3.45E+0 3	9.00E+0 0	5.89E+0 3	6.50E+0 3	8.22E+0 3	1.43644 7	2.690952	1.42811 7	up
Mad727	LysoPC 15:0	Lipids_Glycerophospholipids	1.08E+0 4	1.71E+0 4	1.42E+0 4	2.32E+0 4	2.61E+0 4	5.24E+0 4	1.00090 8	2.415677	1.27242 8	up
Mad761	LysoPC 17:0	Lipids_Glycerophospholipids	9.12E+0 3	1.03E+0 4	9.76E+0 3	2.73E+0 4	2.40E+0 4	5.91E+0 4	1.28998 4	3.783413	1.91968 8	up
Mad134 9	9-KODE	Lipids_Fatty acids	9.00E+0 0	9.00E+0 0	9.00E+0 0	6.38E+0 3	6.37E+0 3	9.00E+0 0	2.12820 8	472.5556	8.88434	up
Mad133 9	13-HpOTrE(r)	Lipids_Fatty acids	9.00E+0 0	9.00E+0 0	9.00E+0 0	2.35E+0 3	3.48E+0 3	2.72E+0 3	2.89936 6	316.6667	8.30682 1	up
Mad135 2	12,13-EODE	Lipids_Fatty acids	9.00E+0 0	9.00E+0 0	9.00E+0 0	1.99E+0 3	1.23E+0 3	9.00E+0 0	1.88524 3	119.5926	6.90198 4	up

Mad861	Citramalate	Organic acids	2.23E+0 6 6	2.15E+0 6 6	2.14E+0 6 7	1.29E+0 7	9.44E+0 6 6	6.65E+0 6 1	1.43605 1 1	4.446319 1 1	2.15261 1 up
Mad101 7	Cyanidin O-syringic acid	Anthocyanins	5.54E+0 4 0	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 5	3.43E+0 4 4	6.33E+0 3 3	1.40113 3 3	7.331715 1 1	2.87415 1 up
Mad102 0	Esculetin O-quinacyl esculetin O-quinic acid	Coumarins	9.00E+0 0 0	9.00E+0 6 6	3.22E+0 6 6	2.67E+0 6 6	2.07E+0 6 6	2.66E+0 6 7	2.89569 7 7	2.298124 7 7	1.20045 7 up
Mad102 5	2,3-dimethylsuccinic acid	Amino acid derivatives	3.37E+0 4 4	2.84E+0 4 4	2.59E+0 4 5	1.02E+0 5	5.14E+0 4 4	1.40E+0 5	1.20638 5 5	3.334091 3 3	1.73729 8 up
Mad969	Glutaric acid	Organic acids	9.00E+0 0 0	9.00E+0 6 6	1.14E+0 5 5	9.98E+0 5 5	9.39E+0 5 5	7.46E+0 7 7	2.77580 7 7	2.353472 8 up	1.23479 8 up
Mad102 3	Adipic acid	Organic acids	2.71E+0 4 4	2.59E+0 4 4	1.01E+0 5 5	1.16E+0 4 4	5.12E+0 5 5	1.26E+0 5 5	1.41091 5 5	4.646593 3 3	2.21617 3 up
Mad93	6-Aminocaproic acid	Organic acids	9.20E+0 5 6	2.03E+0 6 6	2.55E+0 6 6	5.50E+0 6 6	4.08E+0 6 6	4.15E+0 6 6	1.10218 6 6	2.496364 8 8	1.31982 8 up
Mad909	L-(+)-Tartaric acid	Organic acids	3.52E+0 4 4	5.18E+0 4 4	3.83E+0 5 5	3.05E+0 5 5	2.19E+0 5 5	1.35E+0 5 5	1.50221 5 5	5.259377 2 2	2.39489 2 up
Mad125 2	(+)-cis,trans-Abscisic acid (ABA)	Phytohormones	1.90E+0 5 5	2.12E+0 5 5	2.05E+0 5 5	4.20E+0 5 5	6.63E+0 5 5	4.47E+0 5 5	1.12236 6 6	2.520593 3 3	1.33376 3 up
Mad43	L-Carnitine	Others	3.38E+0 4 0	9.00E+0 0 0	9.00E+0 0 4	4.77E+0 4 4	5.07E+0 4 4	4.70E+0 4 4	2.49884 5 5	4.299485 4 4	2.10416 4 up
Mad778	trans,trans-Muconic acid	Organic acids	3.99E+0 5 5	4.06E+0 5 5	4.55E+0 5 5	1.25E+0 6 5	9.25E+0 5 5	7.81E+0 5 5	1.07543 4 4	2.346032 3 3	1.23022 3 up
Mad111 4	p-Coumaryl alcohol	Hydroxycinnamoyl derivatives	3.54E+0 4 4	3.13E+0 3 3	9.91E+0 5 5	4.10E+0 5 5	3.03E+0 5 5	2.94E+0 5 5	1.94520 3 3	13.1445 7 7	3.71638 7 up
Mad390	1-O-Feruloyl quinic acid	Quinate and its derivatives	6.12E+0 4 4	3.25E+0 4 4	7.47E+0 4 4	1.71E+0 4 3	9.56E+0 3 4	2.47E+0 4 4	1.22322 4 4	0.304988 n n	- dow
Mad577	Chrysin O-hexoside	Flavone	1.05E+0 5 4	7.90E+0 4 4	9.64E+0 4 4	4.82E+0 4 4	2.98E+0 4 4	5.31E+0 4 4	1.01210 7 7	0.467546 n n	- dow
Mad108 4	5-O-p-Coumaroylquinic acid	Quinate and its derivatives	8.61E+0 5 5	7.74E+0 5 5	8.66E+0 5 5	4.07E+0 5 5	2.62E+0 5 5	3.91E+0 5 5	1.10341 2 2	0.42383 n n	- dow
Mad750	MAG (18:1) isomer2	Lipids_Glycerolipids	4.66E+0 4 4	5.44E+0 4 4	5.37E+0 4 4	3.03E+0 4 4	2.68E+0 4 4	1.60E+0 4 4	1.00971 6 6	0.472527 n n	- dow
Mad334	Hydroxy-methoxycinnamate	Hydroxycinnamoyl derivatives	2.18E+0 6 6	1.26E+0 6 6	1.48E+0 5 5	2.48E+0 5 5	3.05E+0 5 5	3.18E+0 5 5	1.56559 1 1	0.177033 n n	- dow
Mad240	N,N'',N''''-p-coumaroyl-cinnamoyl-caffeoyle spermidine	Phenolamides	2.52E+0 3 0	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	1.10916 2 2	0.010638 n n	- dow
Mad11	Agmatine	Phenolamides	6.58E+0 3 3	4.46E+0 3 3	4.72E+0 3 3	5.05E+0 3 3	9.00E+0 0 0	9.00E+0 0 0	2.09834 2 2	0.321574 n n	1.63678 n dow
Mad356	Rosinidin O-hexoside	Anthocyanins	7.17E+0 3 0	9.00E+0 3 3	5.89E+0 0 3	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	2.13390 1 1	0.002066 n n	- dow
Mad386	Quercetin 7-O-malonylhexosyl-hexoside	Flavonol	4.67E+0 3 3	5.35E+0 3 3	4.06E+0 0 3	9.00E+0 0 0	5.86E+0 3 0	9.00E+0 0 0	2.03001 6 6	0.417472 n n	- dow
Mad430	5-O-p-Coumaroyl shikimic acid	Quinate and its derivatives	2.86E+0 4 3	7.97E+0 3 4	2.19E+0 4 0	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	3.31661 2 2	0.000462 n n	- dow
Mad585	N-Feruloyl tyramine	Phenolamides	9.00E+0 0 3	8.22E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	1.21724 2 2	0.003277 n n	- dow
Mad163	Niacinamide	Vitamins	7.06E+0 4 4	5.03E+0 4 4	6.72E+0 4 4	3.14E+0 4 4	3.54E+0 4 4	2.13E+0 4 3	1.00806 3 3	0.468368 n n	- dow
Mad323	Pyridoxine O-feruloyl hexoside	Vitamins	2.24E+0 4 4	1.19E+0 4 3	9.88E+0 3 4	1.88E+0 4 0	9.00E+0 0 0	9.00E+0 0 0	2.18798 8 8	0.425939 n n	- dow
											1.23128 n n

Mad235	Methoxyindoleacetic acid	Indole derivatives	3.18E+0 4	7.65E+0 4	1.39E+0 5	1.69E+0 4	2.90E+0 4	2.24E+0 4	1.14416 1.14416	0.276183 0.276183	-1.8563 -1.8563	dow n
Mad747	LysoPE 18:1 (2n isomer)	Lipids_Glycerophospholipids	5.27E+0 5	3.81E+0 5	5.31E+0 5	1.66E+0 5	1.19E+0 5	9.16E+0 4	1.38081 8	0.26171 0.26171	- 1.93396	dow n
Mad749	LysoPC 18:1 (2n isomer)	Lipids_Glycerophospholipids	8.22E+0 6	5.74E+0 6	5.90E+0 6	3.63E+0 6	2.57E+0 6	2.71E+0 6	1.04376 5	0.44864 0.44864	- 1.15637	dow n
Mad241	Indole	Indole derivatives	5.47E+0 3	1.61E+0 4	1.97E+0 4	9.00E+0 0	9.00E+0 0	3.40E+0 3	2.45861 9	0.08282 0.08282	- 3.59387	dow n
Mad46	O-Phosphocholine	Cholines	1.21E+0 5	1.01E+0 5	1.49E+0 5	1.47E+0 4	1.30E+0 4	8.88E+0 3	1.83401 3	0.098598 0.098598	- 3.34229	dow n
Mad755	LysoPC 18:0 (2n isomer)	Lipids_Glycerophospholipids	7.60E+0 4	5.30E+0 4	6.07E+0 4	3.29E+0 4	2.29E+0 4	2.46E+0 4	1.08942 4	0.423827 0.423827	- 1.23845	dow n
Mad963	Cyclic AMP	Nucleotide and its derivates	6.93E+0 4	4.82E+0 4	4.57E+0 4	2.41E+0 4	2.64E+0 4	2.39E+0 4	1.03084 2	0.455882 0.455882	- 1.13327	dow n
Mad107 9	1-O-beta-D-Glucopyranosyl sinapate	Hydroxycinnamoyl derivates	8.10E+0 5	4.32E+0 5	4.22E+0 5	1.49E+0 5	2.38E+0 5	8.61E+0 4	1.28500 1	0.284315 0.284315	- 1.81444	dow n
Mad111 1	Catechin-catechin-catechin	Catechin derivatives	1.37E+0 6	6.87E+0 5	7.24E+0 5	3.53E+0 5	2.32E+0 5	3.38E+0 5	1.18622 6	0.331895 0.331895	-1.5912 -1.5912	dow n
Mad118 8	Apigenin 7-O-glucoside (Cosmosin)	Flavone	2.66E+0 5	1.95E+0 5	2.09E+0 5	7.89E+0 4	8.42E+0 4	8.55E+0 4	1.18844 5	0.371045 0.371045	- 1.43033	dow n
Mad122 8	Luteolin O-eudesmic acid-O-hexoside	Flavone	3.27E+0 5	2.45E+0 5	1.80E+0 5	9.38E+0 4	3.14E+0 4	4.43E+0 4	1.44231 6	0.225399 0.225399	- 2.14945	dow n
Mad116 1	Quercetin O-acetylhexoside	Flavonol	4.47E+0 6	3.31E+0 6	2.93E+0 6	7.97E+0 5	2.33E+0 6	1.31E+0 6	1.07425 3	0.414286 0.414286	-1.2713 -1.2713	dow n
Mad119 8	Tricin 7-O-hexoside	Flavone	5.32E+0 5	3.05E+0 5	1.59E+0 5	2.22E+0 5	2.88E+0 4	5.29E+0 4	1.23257 4	0.30492 0.30492	-1.7135 -1.7135	dow n
Mad848	Trehalose 6-phosphate	Carbohydrates	9.43E+0 4	6.11E+0 4	5.65E+0 4	9.00E+0 0	6.47E+0 4	3.61E+0 4	1.51724 3	0.475739 0.475739	- 1.07176	dow n
Mad881	Glutathione oxidized	Amino acid derivatives	8.21E+0 4	1.01E+0 5	1.15E+0 5	4.65E+0 4	3.18E+0 4	1.06E+0 4	1.29303 4	0.298222 0.298222	- 1.74554	dow n
Mad195	6-Methylmercaptopurine	Nucleotide and its derivates	1.46E+0 7	1.22E+0 7	1.51E+0 7	1.06E+0 6	9.71E+0 5	1.16E+0 6	1.93936 7	0.076158 0.076158	- 3.71487	dow n
Mad208	Succinyladenosine	Nucleotide and its derivates	3.39E+0 5	2.95E+0 5	3.03E+0 5	1.47E+0 5	1.12E+0 5	1.80E+0 5	1.03138 4	0.468517 0.468517	- 1.09383	dow n
Mad754	LysoPC 20:4	Lipids_Glycerophospholipids	5.82E+0 5	4.22E+0 5	5.09E+0 5	2.80E+0 5	1.82E+0 5	2.08E+0 5	1.06061 9	0.442829 0.442829	- 1.17518	dow n
Mad769	LysoPC 20:1	Lipids_Glycerophospholipids	1.18E+0 5	7.26E+0 4	5.75E+0 4	2.83E+0 4	3.09E+0 4	2.38E+0 4	1.19172 8	0.334543 0.334543	- 1.57974	dow n
Mad65	L-Proline	Amino acids	3.57E+0 6	6.77E+0 6	9.35E+0 6	1.60E+0 6	3.17E+0 6	2.12E+0 6	1.10894 3	0.349924 0.349924	- 1.51489	dow n
Mad196	L-Phenylalanine	Amino acids	3.87E+0 7	3.79E+0 7	3.76E+0 7	1.50E+0 7	1.41E+0 7	1.59E+0 7	1.16683 3	0.394046 0.394046	- 1.34357	dow n
Mad157	Uridine	Nucleotide and its derivates	4.37E+0 6	2.57E+0 6	3.29E+0 6	1.18E+0 6	1.15E+0 6	1.05E+0 7	1.23567 7	0.330401 0.330401	- 1.59771	dow n
Mad120	Adenine	Nucleotide and its derivates	1.45E+0 5	1.72E+0 5	2.14E+0 5	5.19E+0 4	6.38E+0 4	5.72E+0 4	1.26065 8	0.325612 0.325612	- 1.61877	dow n
Mad132	N-Acetyl-L-glutamic acid	Amino acid derivatives	2.84E+0 5	2.51E+0 5	1.68E+0 5	5.19E+0 4	1.17E+0 5	3.72E+0 4	1.29839 1	0.293172 0.293172	- 1.77018	dow n
Mad35	5-Aminovaleric acid	Amino acid derivatives	1.14E+0 4	1.40E+0 4	1.66E+0 4	6.71E+0 3	9.00E+0 0	5.14E+0 3	1.58891 7	0.282357 0.282357	- 1.82441	dow n

Mad101	Asp-phe	Amino acid derivatives	5.84E+0 4	5.05E+0 4	5.39E+0 4	3.58E+0 4	1.89E+0 4	2.12E+0 4	1.02447 8	0.466216	-	1.10093	dow
Mad50	2-Hydroxy-6-aminopurine	Nucleotide and its derivates	1.44E+0 5	1.05E+0 5	1.28E+0 5	4.46E+0 4	7.30E+0 4	5.65E+0 4	1.02724 1	0.461804	-	1.11465	n
Mad125	Quercetin	Flavonol	2.07E+0 5	1.42E+0 5	1.27E+0 5	3.25E+0 4	3.98E+0 4	3.30E+0 4	1.46051 2	0.221218	-	2.17646	n
Mad104	Kynurenic acid	Organic acids	8.49E+0 4	1.31E+0 5	7.28E+0 4	2.70E+0 4	4.14E+0 4	3.19E+0 4	1.18640 7	0.347419	-	1.52525	dow
Mad107	Uracil	Nucleotide and its derivates	3.37E+0 5	2.15E+0 5	3.00E+0 5	8.90E+0 4	1.06E+0 5	1.15E+0 5	1.18269 2	0.36385	-	1.45859	dow
Mad786	2,6-Diaminoimelic acid	Amino acid derivatives	4.58E+0 5	3.02E+0 5	2.63E+0 5	8.81E+0 4	6.62E+0 4	1.32E+0 5	1.32049 2	0.279863	-	1.83721	n
Mad108	Caffeic acid	Hydroxycinnamoyl derivatives	1.27E+0 5	9.20E+0 4	7.92E+0 4	4.58E+0 4	4.21E+0 4	4.04E+0 4	1.06781 1	0.430248	-	1.21676	dow
Mad117	Ferulic acid	Hydroxycinnamoyl derivatives	2.99E+0 4	1.69E+0 4	2.30E+0 4	9.00E+0 0	2.18E+0 4	9.00E+0 0	2.33199 4	0.312579	-	1.67771	n
Mad444	Apigenin 5-O-glucoside	Flavone	2.42E+0 5	1.70E+0 5	1.79E+0 5	4.49E+0 4	8.40E+0 4	9.83E+0 4	1.13579 9	0.384433	-	1.3792	n
Mad118	Naringenin 7-O-glucoside (Prunin)	Flavanone	4.69E+0 6	3.18E+0 6	3.05E+0 6	1.52E+0 6	1.70E+0 6	1.37E+0 6	1.08405 6	0.42033	-	1.25041	dow
Mad362	Homovanillic acid	Hydroxycinnamoyl derivatives	5.18E+0 4	5.37E+0 4	3.54E+0 4	9.00E+0 0	2.84E+0 4	2.00E+0 4	1.59762 7	0.34357	-	1.54132	n
Mad863	Shikimic acid	Organic acids	9.44E+0 5	5.63E+0 5	5.31E+0 5	1.75E+0 5	1.91E+0 5	2.56E+0 5	1.26493 2	0.305201	-	1.71217	dow
Mad401	Chlorogenic acid methyl ester	Quinate and its derivatives	1.80E+0 5	1.80E+0 5	1.81E+0 5	1.04E+0 5	6.87E+0 4	7.87E+0 4	1.04518 4	0.464695	-	1.10564	n
Mad623	Phloretin	Flavanone	2.74E+0 4	1.88E+0 4	1.35E+0 4	9.00E+0 0	1.36E+0 4	9.85E+0 3	1.47705	0.392948	-	1.34759	n
Mad118	3-Hydroxy-4-methoxycinnamic acid	Hydroxycinnamoyl derivatives	2.43E+0 4	2.26E+0 4	1.86E+0 4	9.00E+0 0	9.00E+0 0	9.00E+0 0	3.37679 3	0.000412	-	11.2443	n
Mad947	Methylmalonic acid	Organic acids	9.38E+0 6	1.12E+0 7	1.06E+0 7	4.53E+0 6	7.15E+0 6	2.91E+0 6	1.00704 4	0.467928	-	1.09564	n
Mad108	Nicotinic acid	Nicotinic acid derivatives	2.34E+0 5	2.27E+0 5	2.38E+0 5	1.35E+0 4	3.04E+0 4	9.00E+0 0	2.26231	0.062817	-	3.9927	n
Mad4	Gluconic acid	Carbohydrates	9.00E+0 0	9.00E+0 0	1.66E+0 4	9.00E+0 0	9.00E+0 0	9.00E+0 0	1.28416 4	0.001625	-	9.26557	dow
Mad894	L-(-)-Tyrosine	Amino acids	1.19E+0 5	1.31E+0 5	1.61E+0 5	1.33E+0 4	1.58E+0 4	1.31E+0 4	1.82013 7	0.102676	-	3.28382	n
Mad109	Guanine	Nucleotide and its derivates	4.08E+0 4	3.27E+0 4	3.56E+0 4	1.64E+0 4	1.97E+0 4	1.42E+0 4	1.04641 9	0.461045	-	1.11702	n
Mad71	Inosine	Nucleotide and its derivates	1.97E+0 5	1.20E+0 5	1.07E+0 5	3.55E+0 4	2.93E+0 4	8.07E+0 4	1.18335	0.34316	-	1.54305	n
Mad962	Gallic acid	Benzoic acid derivatives	2.28E+0 4	9.00E+0 0	9.00E+0 0	9.00E+0 0	9.00E+0 0	9.00E+0 0	1.30809 5	0.001183	-	-9.723	dow
Mad180	Deoxyguanosine	Nucleotide and its derivates	1.55E+0 4	1.06E+0 4	9.00E+0 0	9.00E+0 0	9.00E+0 0	9.00E+0 0	2.23603 1	0.001034	-	9.91737	n
Mad927	S-(5'-Adenosy)-L-homocysteine	Amino acid derivatives	1.54E+0 4	1.72E+0 4	8.69E+0 3	6.59E+0 3	9.48E+0 3	9.00E+0 0	1.45454 4	0.389416	-	1.36061	n
Mad960	Xanthosine	Nucleotide and its derivates	2.31E+0 4	2.14E+0 4	2.05E+0 4	1.02E+0 4	1.57E+0 4	9.00E+0 0	1.50129	0.3986	-	1.32699	n

Mad113	p-Coumaric acid	Hydroxycinnamoyl derivatives	1.56E+0 5 5	1.13E+0 4 4	6.63E+0 4 4	6.54E+0 4 4	3.39E+0 0 0	9.00E+0 0 0	1.69734 7 7	0.29618 -	-	dow
Mad124	Oroborol (5,7,3',4'-tetrahydroxyisoflavone)	Isoflavone	3.33E+0 3 0	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	1.13626 2 2	0.008065 -	1.75546 n	dow
Mad129	(+)-Jasmonic acid (JA)	Phytohormones	3.77E+0 4 4	2.22E+0 4 4	2.29E+0 3 3	5.92E+0 3 3	2.18E+0 4 4	9.00E+0 0 0	1.62499 1 1	0.334891 -	1.57824 n	dow
Mad218	D-Pantothenic acid	Organic acids	5.51E+0 3 3	3.92E+0 3 3	6.10E+0 3 3	2.42E+0 3 3	9.00E+0 0 0	2.54E+0 3 3	1.46784 8 8	0.319961 -	1.64403 n	dow
Mad988	3-Hydroxy-3-methyl butyric acid	Organic acids	6.88E+0 4 4	9.37E+0 4 4	9.90E+0 4 4	1.57E+0 4 4	2.06E+0 4 4	9.00E+0 0 0	1.94289 6 6	0.138849 -	2.84841 n	dow
Mad928	Succinic acid	Organic acids	1.01E+0 7 7	1.18E+0 7 7	1.02E+0 7 7	4.45E+0 6 6	6.90E+0 6 6	2.80E+0 6 6	1.05294 6 6	0.44081 -	1.18177 n	dow
Mad221	Theobromine	Alkaloids	8.24E+0 3 3	7.67E+0 3 3	7.35E+0 3 3	5.07E+0 3 3	9.00E+0 0 0	6.54E+0 3 3	1.33078 1 1	0.499527 -	1.00137 n	dow
Mad72	Trigonelline	Alkaloids	1.02E+0 6 6	1.47E+0 6 6	1.71E+0 5 5	5.34E+0 5 5	3.58E+0 5 5	2.15E+0 5 5	1.35857 3 3	0.263571 -	1.92373 n	dow
Mad178	2'-Deoxyinosine	Nucleotide and its derivates	1.19E+0 4 4	7.19E+0 3 3	1.11E+0 4 4	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	3.19944 8 8	0.000894 -	10.1269 n	dow
Mad332	N-Acetyl-5-hydroxytryptamine	Tryptamine derivatives	5.78E+0 3 3	4.71E+0 3 3	7.06E+0 3 3	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	3.07581 9 9	0.001538 -	-9.3443 n	dow
Mad82	L-Homocystine	Amino acids	1.16E+0 5 5	3.17E+0 4 4	3.97E+0 4 4	1.33E+0 4 4	9.48E+0 3 3	1.30E+0 4 4	1.38454 6 6	0.190928 -	-2.3889 n	dow
Mad106	4-Hydroxybenzoic acid	Organic acids	1.88E+0 5 5	1.53E+0 5 5	9.00E+0 0 0	1.19E+0 5 5	9.00E+0 0 0	9.00E+0 0 0	1.31573 8 8	0.349017 -	1.51863 n	dow
Mad943	3-Hydroxy-3-methylpentane-1,5-dioic acid	Amino acid derivatives	2.26E+0 5 5	2.92E+0 5 5	2.09E+0 5 5	9.59E+0 4 4	1.06E+0 5 5	4.35E+0 4 4	1.22004 3 3	0.337552 -	-1.56682 n	dow
Mad126	Naringenin chalcone	Flavanone	9.03E+0 3 0	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	1.22837 0 0	0.002984 -	-8.3885 n	dow
Mad124	Isosakuranetin-7-neohesperidoside (Poncirin)	Flavanone	3.15E+0 5 5	1.15E+0 5 5	1.27E+0 5 5	2.21E+0 4 4	2.30E+0 4 4	1.65E+0 4 4	1.71237 9 9	0.110592 -	-3.17668 n	dow
Mad23	γ -aminobutyric acid	Organic acids	3.09E+0 5 5	3.18E+0 5 5	3.15E+0 5 5	1.26E+0 5 5	2.01E+0 5 5	1.26E+0 5 5	1.01289 2 2	0.480892 -	1.05622 n	dow
Mad63	N,N-Dimethylglycine	Amino acid derivatives	2.08E+0 5 5	1.68E+0 5 5	1.52E+0 5 5	5.94E+0 4 4	1.24E+0 5 5	6.28E+0 4 4	1.00544 9 9	0.466288 -	-1.10071 n	dow
Mad906	Aminomalonic acid	Organic acids	3.48E+0 5 5	3.95E+0 5 5	3.42E+0 5 5	1.55E+0 5 5	2.48E+0 5 5	8.70E+0 4 4	1.02936 3 3	0.451613 -	-1.14684 n	dow
Mad115	Genistein 7-O-Glucoside (Genistin)	Isoflavone	2.74E+0 4 4	1.12E+0 4 4	1.37E+0 4 4	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	3.30817 6 6	0.000516 -	-10.9196 n	dow
Mad111	Vitexin 2"-O-beta-L-rhamnoside	Flavone C-glycosides	5.84E+0 4 4	2.37E+0 4 4	2.95E+0 4 4	1.07E+0 4 4	1.03E+0 4 4	1.21E+0 4 4	1.22638 1 1	0.296595 -	-1.75343 n	dow
Mad104	Coniferin	Hydroxycinnamoyl derivatives	2.57E+0 6 6	1.42E+0 6 6	1.52E+0 6 6	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	4.22418 3 3	4.90E-06 -	-17.6387 n	dow
Mad465	Kaempferol 3-O-galactoside (Trifolin)	Flavonol	1.37E+0 6 6	1.35E+0 6 6	1.06E+0 6 6	7.30E+0 5 5	4.90E+0 5 5	4.46E+0 5 5	1.06379 4 4	0.440741 -	-1.182 n	dow
Mad365	Afzelechin (3,5,7,4'-Tetrahydroxyflavan)	Flavanone	1.12E+0 5 5	1.13E+0 5 5	9.89E+0 4 4	4.07E+0 4 4	7.14E+0 4 4	3.95E+0 4 4	1.02235 2 2	0.468046 -	-1.09528 n	dow
Mad125	Morin	Flavonol	2.15E+0 5 5	1.53E+0 5 5	1.31E+0 5 5	3.30E+0 4 4	4.77E+0 4 4	3.80E+0 4 4	1.42247 5 5	0.237876 -	-2.07172 n	dow
Mad198	Deoxyadenosine	Nucleotide and its derivates	1.36E+0 5 5	8.86E+0 4 4	9.66E+0 4 4	4.11E+0 4 4	4.58E+0 4 4	2.01E+0 4 4	1.22159 4 4	0.333126 -	-1.58586 n	dow

Mad219	2-(dimethylamino)guanosine	Nucleotide and its derivates	3.70E+0 4	2.73E+0 4	3.21E+0 4	1.72E+0 4	9.15E+0 3	8.61E+0 3	1.18902 7	0.362656	-	dow
Mad891	β -Pseudouridine	Nucleotide and its derivates	6.11E+0 4	4.80E+0 4	3.59E+0 4	1.95E+0 4	1.69E+0 4	1.54E+0 4	1.18484 9	0.357241	-	dow
PS3 vs PS4:												
			PS3a	PS3b	PS3c	PS4a	PS4b	PS4c				
Mad759	9-Hydroxy-(10E,12Z,15Z)-octadecatrienoic acid	Lipids_Fatty acids	4.66E+0 4	5.18E+0 4	3.78E+0 4	1.88E+0 5	1.99E+0 5	1.76E+0 5	1.60784 2	4.133627	2.04740 8	up
Mad70	sn-Glycero-3-phosphocholine	Cholines	2.95E+0 5	3.07E+0 5	2.38E+0 5	6.83E+0 5	6.64E+0 5	7.35E+0 5	1.28143 5	2.478571	1.30950 9	up
Mad359	di-C,C-hexosyl-luteolin	Flavone C-glycosides	6.66E+0 3	9.00E+0 0	2.72E+0 3	1.57E+0 4	1.22E+0 4	3.07E+0 3	1.64522 3	3.298541	1.72182 8	up
Mad699	MAG (18:4) isomer2	Lipids_Glycerolipids	4.51E+0 4	7.12E+0 4	2.48E+0 4	1.79E+0 5	1.30E+0 5	6.86E+0 4	1.18267 2	2.676116	1.42014 1	up
Mad11	Agmatine	Phenolamides	5.05E+0 3	9.00E+0 0	9.00E+0 0	9.61E+0 3	3.29E+0 4	1.58E+0 4	2.79814 8	11.50552	3.52425 5	up
Mad228	N'-p-Coumaroyl putrescine	Phenolamides	9.00E+0 0	9.78E+0 3	9.00E+0 0	8.09E+0 3	1.15E+0 4	9.00E+0 0	1.18123 2	2.000306	1.00022 1	up
Mad248	Cyanidin 3-O-glucoside (Kuromarin)	Anthocyanins	2.54E+0 5	1.49E+0 6	4.00E+0 5	8.75E+0 6	7.45E+0 7	2.83E+0 7	2.57015 8	52.02892	5.70124 2	up
Mad456	Syringetin 5-O-hexoside	Flavone	4.38E+0 4	9.26E+0 4	1.35E+0 4	3.10E+0 5	8.69E+0 4	7.55E+0 4	1.18468 6	3.151434	1.65600 9	up
Mad493	Syringetin 7-O-hexoside	Flavone	2.83E+0 4	7.66E+0 4	1.20E+0 4	2.66E+0 5	4.89E+0 4	6.59E+0 4	1.13830 9	3.257485	1.70375 9	up
Mad298	C-hexosyl-luteolin O-hexoside	Flavone C-glycosides	9.00E+0 0	9.00E+0 0	9.00E+0 0	1.16E+0 4	1.09E+0 4	9.00E+0 0	2.47747 4	833.6667	9.70332 7	up
Mad386	Quercetin 7-O-malonylhexasyl-hexoside	Flavonol	9.00E+0 0	5.86E+0 3	9.00E+0 0	9.00E+0 0	4.10E+0 4	7.48E+0 4	1.74309 1	19.70211	4.30027 8	up
Mad430	5-O-p-Coumaroyl shikimic acid	Quinate and its derivatives	9.00E+0 0	9.00E+0 0	9.00E+0 0	8.63E+0 3	9.00E+0 0	9.00E+0 0	1.35818 6	320.2963	8.32326 3	up
Mad585	N-Feruloyl tyramine	Phenolamides	9.00E+0 0	9.00E+0 0	9.00E+0 0	9.00E+0 0	9.00E+0 0	1.34E+0 4	1.41577 8	496.963	8.95699 5	up
Mad100	Piperidine	Alkaloids	1.29E+0 6	8.42E+0 5	8.38E+0 5	2.29E+0 6	4.48E+0 6	3.98E+0 6	1.47073 1	3.619529	1.85580 2	up
Mad235	Methoxyindoleacetic acid	Indole derivatives	1.69E+0 4	2.90E+0 4	2.24E+0 4	6.93E+0 4	3.99E+0 4	3.74E+0 4	1.06403 2	2.146413	1.10192 8	up
Mad745	LysoPC 18:3	Lipids_Glycerophospholipids	2.61E+0 5	3.32E+0 5	5.81E+0 5	1.40E+0 6	1.28E+0 6	9.16E+0 5	1.38912 6	3.063032	1.61496 1	up
Mad746	LysoPC 16:0	Lipids_Glycerophospholipids	1.47E+0 6	1.85E+0 6	3.41E+0 6	6.75E+0 6	6.89E+0 6	5.15E+0 6	1.33461 4	2.791976	1.48128 7	up
Mad704	LysoPC 16:2 (2n isomer)	Lipids_Glycerophospholipids	2.92E+0 4	4.13E+0 4	3.86E+0 4	1.02E+0 5	8.56E+0 4	9.19E+0 4	1.29808 9	2.56187	1.35719 7	up
Mad715	LysoPE 14:0	Lipids_Glycerophospholipids	2.63E+0 3	2.57E+0 3	3.89E+0 3	1.01E+0 4	1.18E+0 4	8.77E+0 3	1.47581 3	3.374037	1.75447 6	up
Mad719	LysoPC 14:0	Lipids_Glycerophospholipids	1.43E+0 4	1.29E+0 4	2.06E+0 4	5.84E+0 4	7.32E+0 4	4.60E+0 4	1.52047 6	3.715481	1.89354 9	up
Mad765	LysoPE 18:0	Lipids_Glycerophospholipids	6.88E+0 3	8.20E+0 3	1.29E+0 4	2.46E+0 4	3.35E+0 4	2.51E+0 4	1.38337 1	2.973553	1.57218 8	up
Mad751	Punicic acid	Lipids_Fatty acids	7.20E+0 4	7.26E+0 4	6.14E+0 4	2.47E+0 5	2.65E+0 5	2.54E+0 5	1.54900 2	3.718447	1.8947	up
Mad771	MAG (18:2)	Lipids_Glycerolipids	1.98E+0 5	2.70E+0 5	1.59E+0 5	8.37E+0 5	1.17E+0 6	9.95E+0 5	1.67946 1	4.787879	2.25938 7	up

Mad217	trans-zeatin N-glucoside	Phytohormones	3.40E+0 5	4.66E+0 5	3.73E+0 5	1.39E+0 6	7.70E+0 5	6.34E+0 5	1.12142 5	2.369805	1.24476 8	up
Mad241	Indole	Indole derivatives	9.00E+0 0	9.00E+0 0	3.40E+0 3	3.92E+0 3	6.69E+0 3	8.13E+0 3	2.49782 1	5.482738	2.45489 7	up
Mad420	Phellodensin F	Others	4.11E+0 3	9.00E+0 0	9.00E+0 0	6.84E+0 3	9.00E+0 0	7.38E+0 3	1.27026 3	3.446948	1.78531 9	up
Mad46	O-Phosphocholine	Cholines	1.47E+0 4	1.30E+0 4	8.88E+0 3	1.82E+0 4	2.91E+0 4	3.48E+0 4	1.11386 8	2.244396	1.16632 7	up
Mad730	MAG (18:3) isomer2	Lipids_Glycerolipids	5.51E+0 3	7.84E+0 3	7.99E+0 3	1.94E+0 4	2.10E+0 4	1.40E+0 4	1.26692 3	2.549203	1.35004 6	up
Mad761	LysoPC 17:0	Lipids_Glycerophospholipids	2.73E+0 4	2.40E+0 4	5.91E+0 5	1.66E+0 5	2.52E+0 5	1.89E+0 5	1.75347 1	5.498188	2.45895 6	up
Mad768	MAG (18:3) isomer1	Lipids_Glycerolipids	2.75E+0 5	1.80E+0 5	1.50E+0 5	5.04E+0 5	2.60E+0 5	4.96E+0 5	1.01932 4	2.082645	1.05841 7	up
Mad788	4-Hydroxy-3-methoxymandelate	Organic acids	1.72E+0 4	1.43E+0 4	7.01E+0 3	4.73E+0 4	3.86E+0 4	3.10E+0 4	1.37420 5	3.035575	1.60197	up
Mad109 6	4-hydroxycoumarin di-glucoside	Coumarins	9.00E+0 0	9.00E+0 0	9.00E+0 0	2.00E+0 4	1.47E+0 5	1.26E+0 5	4.03781 1	10851.85	13.4056 5	up
Mad134 7	9,10-EODE	Lipids_Fatty acids	9.99E+0 4	9.93E+0 4	8.53E+0 4	3.27E+0 5	3.87E+0 5	3.44E+0 5	1.54631 7	3.718805	1.89483 9	up
Mad134 9	9-KODE	Lipids_Fatty acids	6.38E+0 3	6.37E+0 3	9.00E+0 0	1.87E+0 4	1.60E+0 4	9.97E+0 3	1.75511 1	3.501058	1.80779 1	up
Mad133 9	13-HpOTrE(r)	Lipids_Fatty acids	2.35E+0 3	3.48E+0 3	2.72E+0 3	6.26E+0 3	1.08E+0 4	7.54E+0 3	1.33842	2.877193	1.52466 2	up
Mad135 2	12,13-EODE	Lipids_Fatty acids	1.99E+0 3	1.23E+0 3	9.00E+0 0	6.77E+0 3	8.22E+0 3	7.01E+0 3	2.02033 6	6.813255	2.76834 4	up
Mad133 8	13-HPODE	Lipids_Fatty acids	3.16E+0 3	3.66E+0 3	3.07E+0 3	1.04E+0 4	1.38E+0 4	1.12E+0 4	1.51639 8	3.579373	1.83970 7	up
Mad861	Citramalate	Organic acids	1.29E+0 7	9.44E+0 6	6.65E+0 6	1.84E+0 7	2.89E+0 7	2.61E+0 7	1.24688 6	2.531908	1.34022 5	up
Mad124 0	Disinapoyl hexoside	Hydroxycinnamoyl derivatives	9.00E+0 0	9.00E+0 0	9.00E+0 0	1.05E+0 4	3.02E+0 4	2.89E+0 4	3.76119 3	2577.778	11.3319 1	up
Mad101 7	Cyanidin O-syringic acid	Anthocyanins	9.00E+0 0	3.43E+0 5	6.33E+0 4	1.47E+0 6	2.74E+0 7	6.00E+0 6	3.03255 7	85.82138	6.42326 5	up
Mad113 0	Tricin O-saccharic acid	Flavone	1.42E+0 4	1.57E+0 4	9.00E+0 0	1.15E+0 5	1.35E+0 4	7.41E+0 3	1.58934 5	4.544117	2.184	up
Mad107 7	3-O-p-coumaroyl shikimic acid O-hexoside	Quinate and its derivatives	1.39E+0 4	1.37E+0 4	1.15E+0 4	2.50E+0 4	8.95E+0 4	2.99E+0 4	1.30760 3	3.693095	1.88483	up
Mad848	Trehalose 6-phosphate	Carbohydrates	9.00E+0 0	6.47E+0 4	3.61E+0 4	6.96E+0 4	7.20E+0 4	7.54E+0 4	1.72067 4	2.152586	1.10607 1	up
Mad104 8	2-Isopropylmalate	Organic acids	1.66E+0 5	2.07E+0 5	1.75E+0 5	3.15E+0 5	3.99E+0 5	4.47E+0 5	1.13891 5	2.118613	1.08312	up
Mad874	Guanosine 5'-monophosphate	Nucleotide and its derivates	2.58E+0 4	2.40E+0 4	2.78E+0 4	7.94E+0 4	5.20E+0 4	5.50E+0 4	1.22257 4	2.402062	1.26427 3	up
Mad713	LysoPC 14:0 (2n isomer)	Lipids_Glycerophospholipids	2.84E+0 4	2.35E+0 4	3.89E+0 4	1.04E+0 5	1.41E+0 5	8.96E+0 4	1.51510 9	3.685022	1.88167 3	up
Mad741	LysoPC 16:0 (2n isomer)	Lipids_Glycerophospholipids	1.58E+0 6	1.85E+0 6	3.32E+0 6	7.10E+0 6	7.10E+0 6	5.70E+0 6	1.38149 8	2.948148	1.55980 9	up
Mad764	LysoPC 18:0	Lipids_Glycerophospholipids	2.30E+0 5	2.24E+0 5	4.73E+0 5	1.27E+0 6	1.47E+0 6	1.22E+0 6	1.62252 4	4.271845	2.09485 9	up

Mad812	L-Aspartic acid	Amino acids	4.62E+0 6 6	4.84E+0 6 6	2.87E+0 6 6	7.70E+0 6 6	1.14E+0 7 7	1.23E+0 7 7	1.24904 4	2.546634	1.34859 2	up
Mad813	L-Glutamic acid	Amino acids	3.28E+0 6 6	5.22E+0 6 6	2.93E+0 6 6	6.68E+0 6 6	1.08E+0 7 7	8.77E+0 6 6	1.169	2.296588	1.19949 2	up
Mad794	2-Aminoethanesulfonic acid	Organic acids	1.07E+0 4 4	1.02E+0 4 4	2.91E+0 3 3	1.82E+0 4 4	2.29E+0 4 4	1.68E+0 4 4	1.19864 8	2.431751	1.28199 6	up
Mad781	L-Histidine	Amino acids	1.80E+0 4 4	2.54E+0 4 4	2.22E+0 4 4	3.28E+0 4 4	7.41E+0 4 4	4.85E+0 4 4	1.12860 9	2.368902	1.24421 9	up
Mad35	5-Aminovaleric acid	Amino acid derivatives	6.71E+0 3 0	9.00E+0 3 0	5.14E+0 3 3	1.76E+0 4 4	1.27E+0 4 4	1.65E+0 4 4	1.83560 4	3.94637	1.98052 6	up
Mad111 5	Quercetin 3-O-rutinoside (Rutin)	Flavonol	9.16E+0 5 6	1.82E+0 6 6	1.24E+0 5 5	7.26E+0 5 6	5.30E+0 6 6	1.39E+0 7 7	1.02098 7	5.011569	2.32526 2	up
Mad969	Glutaric acid	Organic acids	9.98E+0 5 5	9.39E+0 5 5	7.46E+0 5 5	1.85E+0 6 6	1.94E+0 6 6	2.17E+0 6 6	1.19448 5	2.221394	1.15146 5	up
Mad111 2	N-Acetyl-l-leucine	Amino acid derivatives	2.40E+0 4 4	2.32E+0 4 4	1.86E+0 4 4	5.55E+0 4 4	7.44E+0 4 4	9.59E+0 4 4	1.46192 4	3.431611	1.77888 6	up
Mad991	2-Methylsuccinic acid	Organic acids	1.26E+0 6 6	1.30E+0 6 6	1.15E+0 6 6	2.63E+0 6 6	2.83E+0 6 6	3.20E+0 6 6	1.23663 1	2.334232	1.22294 8	up
Mad93	6-Aminocaproic acid	Organic acids	5.50E+0 6 6	4.08E+0 6 6	4.15E+0 6 6	9.19E+0 6 6	1.45E+0 7 7	1.41E+0 7 7	1.31653 1	2.752367	1.46067 3	up
Mad114 7	Kaempferol 3-O-rutinoside (Nicotiflorin)	Flavonol	4.78E+0 4 4	3.29E+0 4 4	6.93E+0 4 4	4.82E+0 4 4	1.44E+0 5 5	3.37E+0 5 5	1.10517 6	3.528	1.81885 1	up
Mad624	Naringenin	Flavanone	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	1.28E+0 4 4	9.00E+0 0 0	1.41504 1	474.7407	8.89099 6	up
Mad362	Homovanillic acid	Hydroxycinnamoyl derivatives	9.00E+0 0 4	2.84E+0 4 4	2.00E+0 4 4	3.13E+0 4 4	6.02E+0 4 4	5.27E+0 4 4	1.78792 8	2.978785	1.57472 4	up
Mad118 2	3-Hydroxy-4-methoxycinnamic acid	Hydroxycinnamoyl derivatives	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	1.37E+0 4 4	9.00E+0 0 0	9.00E+0 0 0	1.40315 3	508.0741	8.98889 5	up
Mad4	Gluconic acid	Carbohydrates	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	1.66E+0 4 4	1.43637 8	615.4815	9.26557 2	up
Mad180	Deoxyguanosine	Nucleotide and its derivates	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	2.99E+0 4 4	1.48E+0 4 4	9.00E+0 0 0	2.58133 7	1655.889	10.6933 9	up
Mad960	Xanthosine	Nucleotide and its derivates	1.02E+0 4 4	1.57E+0 4 0	9.00E+0 0 0	3.04E+0 4 4	1.71E+0 4 4	1.56E+0 4 4	1.63209 2	2.435447	1.28418 7	up
Mad798	L-Glutamine	Amino acids	3.80E+0 4 4	7.40E+0 4 4	1.64E+0 4 4	7.64E+0 5 5	1.07E+0 5 5	1.01E+0 5 5	1.13345	2.214953	1.14727 6	up
Mad113 5	Kaempferol 3-O-robinobioside (Biorobin)	Flavonol	5.35E+0 4 4	2.71E+0 4 4	6.17E+0 4 4	5.00E+0 5 5	1.48E+0 5 5	3.72E+0 5 5	1.18841 1	4.005622	2.00202 6	up
Mad218	D-Pantothenic acid	Organic acids	2.42E+0 3 0	9.00E+0 3 3	2.54E+0 3 3	7.37E+0 3 3	3.31E+0 3 3	3.58E+0 3 3	1.56770 7	2.869793	1.52094 7	up
Mad988	3-Hydroxy-3-methyl butyric acid	Organic acids	1.57E+0 4 4	2.06E+0 4 0	9.00E+0 0 4	7.21E+0 4 4	2.73E+0 4 4	3.90E+0 4 4	1.84766 6	3.811727	1.93044 5	up
Mad80	Betaine	Alkaloids	9.63E+0 4 5	1.28E+0 5 5	1.10E+0 5 5	2.71E+0 5 5	2.99E+0 5 5	2.01E+0 5 5	1.19632 2	2.306312	1.20558 8	up
Mad909	L-(+)-Tartaric acid	Organic acids	3.05E+0 5 5	2.19E+0 5 5	1.35E+0 5 5	4.44E+0 5 5	5.18E+0 5 5	5.13E+0 5 5	1.16854 4	2.23824	1.16236 5	up
Mad221	Theobromine	Alkaloids	5.07E+0 3 0	9.00E+0 3 3	6.54E+0 4 4	1.38E+0 3 3	5.73E+0 3 3	4.46E+0 3 3	1.43619 9	2.064722	1.04594 7	up
Mad72	Trigonelline	Alkaloids	5.34E+0 5 5	3.58E+0 5 5	2.15E+0 5 5	2.55E+0 6 6	2.11E+0 6 6	1.26E+0 6 6	1.70023 3	5.347787	2.41894 2	up

Mad7	Putrescine	Phenolamides	2.58E+0 5 5	5.10E+0 9.00E+0 0	1.56E+0 9.00E+0 0	1.29E+0 9.00E+0 0	8.59E+0 8.57E+0 3	3.96E+0 9.00E+0 0	1.15869 5 3	2.754329 1.37538 318.0741	1.46170 1 8.31321	up	
Mad332	N-Acetyl-5-hydroxytryptamine	Tryptamine derivatives	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	8.57E+0 3 0	9.00E+0 0 0	1.37538 8 0	318.0741 9 8	8.31321 9 up		
Mad106 8	4-Hydroxybenzoic acid	Organic acids	1.19E+0 5 0	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	2.43E+0 5 4	2.00E+0 5 3	1.54676 5 3	3.722202 3.722202 3.722202	1.89615 6 up		
Mad126 8	Naringenin chalcone	Flavanone	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	1.70E+0 4 4	6.75E+0 3 3	2.4783 2.4783	879.963 879.963	9.78129 9 up		
Mad830	Uridine 5'-diphosphate	Nucleotide and its derivates	1.60E+0 5 5	1.21E+0 5 5	1.55E+0 5 5	3.42E+0 5 5	2.79E+0 5 5	2.60E+0 5 2	1.10387 2 2	2.020642 2.020642 2.020642	1.01481 4 up		
Mad987	ethylmalonate	Organic acids	9.15E+0 3 0	9.00E+0 4 0	1.38E+0 4 4	1.87E+0 4 4	1.81E+0 4 4	3.14E+0 4 2	1.73849 2 2	2.970513 2.970513 2.970513	1.57071 2 up		
Mad115 5	Genistein 7-O-Glucoside (Genistin)	Isoflavone	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	7.98E+0 3 0	9.00E+0 0 0	1.36821 8 0	296.2222 296.2222 296.2222	8.21053 6 up		
Mad104 1	Coniferin	Hydroxycinnamoyl derivatives	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	6.49E+0 5 7	1.75185 7 7	24037.7 24037.7 24037.7	14.5530 1 up		
Mad105 1	Pantetheine	Vitamins	5.66E+0 3 0	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	6.85E+0 3 3	5.95E+0 3 3	1.18863 4 4	2.2559 3 4	1.17370 3 up		
Mad123 0	Formononetin 7-O-glucoside (Ononin)	Isoflavone	9.06E+0 3 3	7.12E+0 3 3	8.62E+0 3 3	9.67E+0 4 4	3.66E+0 4 4	2.77E+0 4 4	1.69485 4 4	6.491935 9 9	2.69864 9 up		
Mad122 5	E-3,4,5'-Trihydroxy-3'-glucopyranosylstilbene	Others	8.14E+0 3 0	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	1.36405 1 1	0.00331 0.00331 0.00331	- -	dow	
Mad97	Xanthine	Nucleotide and its derivates	9.00E+0 0 0	3.07E+0 3 3	3.57E+0 3 3	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	2.25702 4 4	0.004061 0.004061 0.004061	- 7.94403 n	dow	
Mad690	Azoxystrobin acid	Others	1.18E+0 5 5	1.35E+0 5 5	1.78E+0 5 5	4.63E+0 4 4	4.35E+0 4 4	3.83E+0 4 4	1.46526 3 3	0.297216 0.297216 0.297216	- 1.75042 n	dow	
Mad141	L-Leucine	Amino acids	4.79E+0 6 6	4.74E+0 6 6	5.98E+0 6 6	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	4.92612 1 1	1.74E-06 19.1318 n	- -	dow	
Mad121	γ -Glu-Cys	Amino acid derivatives	6.67E+0 3 3	9.76E+0 3 0	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	4.09E+0 3 3	1.31632 8 8	0.249894 0.249894 0.249894	- 2.00061 n	dow	
Mad66	Glycyl-L-proline	Amino acid derivatives	3.90E+0 3 3	4.78E+0 3 3	3.68E+0 3 3	4.74E+0 3 3	9.00E+0 0 0	9.00E+0 0 0	2.27279 2 2	0.384951 0.384951 0.384951	- 1.37725 n	dow	
Mad121 5	Benzoic acid	Benzoic acid derivatives	2.58E+0 5 5	2.39E+0 5 5	3.41E+0 5 5	1.37E+0 5 5	1.10E+0 5 5	1.41E+0 5 5	1.14895 4 4	0.463007 0.463007 0.463007	- 1.11089 n	dow	
Mad102 3	Adipic acid	Organic acids	1.16E+0 5 4	5.12E+0 5 5	1.26E+0 5 4	5.40E+0 4 4	3.40E+0 4 4	3.86E+0 4 4	1.05681 6 6	0.431787 0.431787 0.431787	- 1.21161 n	dow	
Mad117 0	Ferulic acid	Hydroxycinnamoyl derivatives	9.00E+0 0 4	2.18E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	9.00E+0 0 0	1.46378 9 9	0.001238 0.001238 9.65834 n	- -	dow	
Mad108	Nicotinic acid	Nicotinic acid derivatives	1.35E+0 4 4	3.04E+0 4 4	9.00E+0 0 0	9.00E+0 0 0	1.69E+0 4 4	9.00E+0 0 0	1.28565 9 9	0.385297 0.385297 0.385297	- 1.37596 n	dow	
Mad915	S-(methyl)glutathione	Amino acid derivatives	5.34E+0 4 4	5.27E+0 4 4	4.66E+0 4 4	2.92E+0 4 4	1.72E+0 4 4	1.24E+0 4 4	1.29243 2 2	0.385069 0.385069 0.385069	- 1.37681 n	dow	
Mad137	Pyridoxine 5'-phosphate	Vitamins	2.94E+0 4 4	1.88E+0 4 4	2.83E+0 4 4	1.92E+0 4 4	9.00E+0 0 0	1.86E+0 4 4	1.59193 1 1	0.494235 0.494235 0.494235	- 1.01673 n	dow	
Mad53	L-Saccharopine	Amino acid derivatives	4.58E+0 5 5	7.55E+0 5 5	8.24E+0 5 5	2.56E+0 5 5	1.77E+0 5 5	3.25E+0 5 5	1.26805 5 5	0.372116 1.26805 1.26805	- 1.42618 n	dow	
Mad43	L-Carnitine	Others	4.77E+0 4 4	5.07E+0 4 4	4.70E+0 4 4	9.00E+0 0 0	3.41E+0 4 4	3.10E+0 4 4	1.67622 1 1	0.447792 0.447792 0.447792	-1.1591 n n	dow	
Mad139	Orotic acid	Vitamins	4.66E+0 4 4	5.95E+0 4 4	6.63E+0 4 4	5.10E+0 4 4	9.00E+0 0 0	3.48E+0 4 4	1.66929 2 2	0.497732 0.497732 0.497732	- 1.00656 n	dow n	

Mad9	trans-Citridic acid	Organic acids	4.44E+0 5	6.42E+0 5	6.40E+0 5	2.31E+0 5	2.01E+0 5	2.91E+0 5	1.2193	0.418888	-	dow
Mad114 2	Sinapyl alcohol	Hydroxycinnamoyl derivatives	9.19E+0 3	5.45E+0 3	9.00E+0 0	9.00E+0 0	9.00E+0 0	4.29E+0 3	1.26877 6	0.294082	-	dow
Mad778	trans,trans-Muconic acid	Organic acids	1.25E+0 6	9.25E+0 5	7.81E+0 5	2.48E+0 5	3.63E+0 5	6.08E+0 5	1.20189 5	0.412382	-	dow

Table 4. KEGG pathway assignments of differentially accumulated metabolites among PS1 vs PS2, PS2 vs PS3, and PS3 vs PS4, respectively.

#Kegg_pathway	ko_id	Cluter_frequency	Genome_frequency	P-value	Corrected_P-value
PS1 vs PS2_KEGG_stat					
Flavonoid biosynthesis	ko0094 1	13 out of 53 24.5283018867925%	21 out of 221 9.50226244343891%	0.00010348 5	0.0081753
Biosynthesis of phenylpropanoids	ko0106 1	15 out of 53 28.3018867924528%	28 out of 221 12.6696832579186%	0.00027082 9	0.0213955
Degradation of aromatic compounds	ko0122 0	28.3018867924528% 5 out of 53 9.43396226415094%	12.6696832579186% 9 out of 221 4.07239819004525%	0.03812386 7	1
Linoleic acid metabolism	ko0059 1	2 out of 53 3.77358490566038%	2 out of 221 0.904977375565611%	0.05668449 2	1
alpha-Linolenic acid metabolism	ko0059 2	2 out of 53 3.77358490566038%	2 out of 221 0.904977375565611%	0.05668449 2	1
Phenylpropanoid biosynthesis	ko0094 0	7 out of 53 13.2075471698113%	16 out of 221 7.23981900452489%	0.05833098 3	1
Isoflavonoid biosynthesis	ko0094 3	7 out of 53 13.2075471698113% 3 out of 53 5.66037735849057%	16 out of 221 5 out of 221 2.26244343891403%	0.09069238 4	1
Biosynthesis of secondary metabolites	ko0111 0	24 out of 53 45.2830188679245%	84 out of 221 38.0090497737557%	0.13829530 9	1
Benzoate degradation	ko0036 2	2 out of 53 3.77358490566038%	3 out of 221 1.35746606334842%	0.14365248 0.22012015	1
Tyrosine metabolism	ko0035 0	3 out of 53 5.66037735849057%	7 out of 221 3.16742081447964% 1 out of 221	0.23981900 1	1
Biosynthesis of unsaturated fatty acids	ko0104 0	1 out of 53 1.88679245283019%	1 out of 221 0.452488687782805%	0.23981900 5	1
Rheumatoid arthritis	ko0532 3	1 out of 53 1.88679245283019%	1 out of 221 0.452488687782805%	0.23981900 5	1
Benzoic acid family	ko0711 0	1 out of 53 1.88679245283019%	1 out of 221 0.452488687782805%	0.23981900 5	1
Fatty acid degradation	ko0007 1	1 out of 53 1.88679245283019%	1 out of 221 0.452488687782805%	0.23981900 5	1
Anthocyanin biosynthesis	ko0094 2	1 out of 53 1.88679245283019%	1 out of 221 0.452488687782805%	0.23981900 5	1
Styrene degradation	ko0064 3	1 out of 53 1.88679245283019%	1 out of 221 0.452488687782805%	0.23981900 5	1

	ko0521	1 out of 53 1.88679245283019%	1 out of 221	0.23981900		
Renal cell carcinoma	1	1 out of 53 1.88679245283019%	0.452488687782805%	5	1	
Ubiquinone and other terpenoid-quinone biosynthesis	ko0013	0	2 out of 53 3.77358490566038%	4 out of 221 1.80995475113122%	0.24358587	1
	ko0062	3	2 out of 53 3.77358490566038%	4 out of 221 1.80995475113122%	0.24358587	
Toluene degradation	ko0023	0	6 out of 53 11.3207547169811%	19 out of 221 8.5972850678733%	0.28828480	1
Purine metabolism	ko0036	0	3 out of 53 5.66037735849057%	9 out of 221 4.07239819004525%	0.37025254	
Phenylalanine metabolism	ko0079	0	1 out of 53 1.88679245283019%	2 out of 221	0.42295351	1
Folate biosynthesis	ko0010	0	1 out of 53 1.88679245283019%	0.904977375565611%	0.42295351	
Steroid biosynthesis	ko0036	0	1 out of 53 1.88679245283019%	0.904977375565611%	0.42295351	1
Bisphenol degradation	ko0421	3	1 out of 53 1.88679245283019%	0.904977375565611%	0.42295351	
Longevity regulating pathway - worm	ko0094	2	1 out of 53 1.88679245283019%	0.904977375565611%	0.44353927	1
Flavone and flavonol biosynthesis	ko0107	4	4 out of 53 7.54716981132075%	6.33484162895928%	0.50466255	
Biosynthesis of plant hormones	ko0003	0	4 out of 53 7.54716981132075%	15 out of 221	0.56260403	1
Pentose phosphate pathway	ko0062	0	1 out of 53 1.88679245283019%	6.78733031674208%	0.56260403	
Pyruvate metabolism	ko0023	0	1 out of 53 1.88679245283019%	3 out of 221 1.35746606334842%	0.56260403	1
Caffeine metabolism	ko0523	2	1 out of 53 1.88679245283019%	3 out of 221 1.35746606334842%	0.56260403	
Choline metabolism in cancer	ko0002	1	1 out of 53 1.88679245283019%	3 out of 221 1.35746606334842%	0.56260403	1
Citrate cycle (TCA cycle)	ko0062	0	1 out of 53 1.88679245283019%	3 out of 221 1.35746606334842%	0.56260403	
Polycyclic aromatic hydrocarbon degradation	ko0407	4	1 out of 53 1.88679245283019%	3 out of 221 1.35746606334842%	0.56260403	1
Plant hormone signal transduction	ko0520	5	1 out of 53 1.88679245283019%	3 out of 221 1.35746606334842%	0.56260403	
Pathways in cancer	ko0062	0	1 out of 53 1.88679245283019%	3 out of 221 1.35746606334842%	0.56260403	1
Aminobenzoate degradation	ko0095	7	3 out of 53 5.66037735849057%	12 out of 221 5.42986425339367%	0.58196195	
Isoquinoline alkaloid biosynthesis	0	0	1 out of 53 1.88679245283019%	4 out of 221 1.80995475113122%	0.66894342	1

	ko0072			0.66894342		
Carbon fixation pathways in prokaryotes	0	1 out of 53 1.88679245283019%	4 out of 221 1.80995475113122%		2	1
Vitamin B6 metabolism	ko0075	0	1 out of 53 1.88679245283019%	4 out of 221 1.80995475113122%	0.66894342	
	ko0033	2	1 out of 53 1.88679245283019%	4 out of 221 1.80995475113122%	2	1
Carbapenem biosynthesis	ko0065	0	1 out of 53 1.88679245283019%	4 out of 221 1.80995475113122%	0.66894342	
Butanoate metabolism	ko0106	6	1 out of 53 1.88679245283019%	4 out of 221 1.80995475113122%	0.66894342	
Biosynthesis of alkaloids derived from terpenoid and polyketide	ko0019	0	1 out of 53 1.88679245283019%	4 out of 221 1.80995475113122%	0.66894342	
Oxidative phosphorylation	ko0497	2	1 out of 53 1.88679245283019%	4 out of 221 1.80995475113122%	2	1
Bile secretion	ko0056	6	2 out of 53 3.77358490566038%	4.52488687782805%	0.73870410	
Glycerophospholipid metabolism	ko0492	4	1 out of 53 1.88679245283019%	5 out of 221 2.26244343891403%	0.74980055	
Glucagon signaling pathway	ko0415	2	1 out of 53 1.88679245283019%	5 out of 221 2.26244343891403%	0.74980055	
AMPK signaling pathway	ko0106	2	1 out of 53 1.88679245283019%	5 out of 221 2.26244343891403%	0.74980055	
Biosynthesis of terpenoids and steroids	ko0120	2	1 out of 53 1.88679245283019%	5 out of 221 2.26244343891403%	0.74980055	
Carbon metabolism	ko0076	0	2 out of 53 3.77358490566038%	4.97737556561086%	0.78823887	
Nicotinate and nicotinamide metabolism	ko0497	0	2 out of 53 3.77358490566038%	4.97737556561086%	2	1
Vitamin digestion and absorption	ko0027	7	2 out of 53 3.77358490566038%	4.97737556561086%	0.78823887	
Cysteine and methionine metabolism	ko0077	0	2 out of 53 3.77358490566038%	4.97737556561086%	2	1
Pantothenate and CoA biosynthesis	ko0152	0	1 out of 53 1.88679245283019%	6 out of 221 2.71493212669683%	0.81119208	
Antifolate resistance	ko0030	3	1 out of 53 1.88679245283019%	6 out of 221 2.71493212669683%	0.81119208	
Lysine biosynthesis	ko0048	0	1 out of 53 1.88679245283019%	6 out of 221 2.71493212669683%	0.81119208	
Glutathione metabolism	ko0096	0	2 out of 53 3.77358490566038%	5.42986425339367%	0.82949425	
Glucosinolate biosynthesis	ko0040	6	1 out of 53 1.88679245283019%	7 out of 221 3.16742081447964%	0.85773543	
Phenylalanine, tyrosine and tryptophan biosynthesis	ko0040	0	1 out of 53 1.88679245283019%	7 out of 221 3.16742081447964%	0.85773543	

	ko0041			0.85773543		
beta-Alanine metabolism	0	1 out of 53 1.88679245283019%	7 out of 221 3.16742081447964%	4	1	
	ko0031			0.85773543		
Lysine degradation	0	1 out of 53 1.88679245283019%	7 out of 221 3.16742081447964%	4	1	
	ko0106		13 out of 221	0.86351783		
Biosynthesis of alkaloids derived from histidine and purine	5	2 out of 53 3.77358490566038%	5.88235294117647%	5	1	
	ko0112		38 out of 221	0.86365721		
Microbial metabolism in diverse environments	0	7 out of 53 13.2075471698113%	17.1945701357466%	4	1	
	ko0022			0.89296918		
Arginine biosynthesis	0	1 out of 53 1.88679245283019%	8 out of 221 3.61990950226244%	1	1	
	ko0202		10 out of 221	0.93970094		
Two-component system	0	1 out of 53 1.88679245283019%	4.52488687782805%	7	1	
	ko0025		10 out of 221	0.93970094		
Alanine, aspartate and glutamate metabolism	0	1 out of 53 1.88679245283019%	4.52488687782805%	7	1	
	ko0113		37 out of 221			
Biosynthesis of antibiotics	0	5 out of 53 9.43396226415094%	16.7420814479638%	0.97306332	1	
Biosynthesis of alkaloids derived from ornithine, lysine and nicotinic acid	ko0106		13 out of 221	0.97480329		
	4	1 out of 53 1.88679245283019%	5.88235294117647%	3	1	
Biosynthesis of alkaloids derived from shikimate pathway	ko0106		13 out of 221	0.97480329		
	3	1 out of 53 1.88679245283019%	5.88235294117647%	3	1	
Arginine and proline metabolism	ko0033		13 out of 221	0.97480329		
	0	1 out of 53 1.88679245283019%	5.88235294117647%	3	1	
Pyrimidine metabolism	ko0024		14 out of 221	0.98122360		
	0	1 out of 53 1.88679245283019%	6.33484162895928%	8	1	
Mineral absorption	ko0497		14 out of 221	0.98122360		
	8	1 out of 53 1.88679245283019%	6.33484162895928%	8	1	
Central carbon metabolism in cancer	ko0523		22 out of 221	0.98499784		
	0	2 out of 53 3.77358490566038%	9.95475113122172%	1	1	
Metabolic pathways	ko0110	31 out of 53	154 out of 221	0.98503892		
	0	58.4905660377358%	69.683257918552%	2	1	
Biosynthesis of plant secondary metabolites	ko0106		40 out of 221	0.98587726		
	0	5 out of 53 9.43396226415094%	18.0995475113122%	3	1	
2-Oxocarboxylic acid metabolism	ko0121		15 out of 221			
	0	1 out of 53 1.88679245283019%	6.78733031674208%	0.98603109	1	
Aminoacyl-tRNA biosynthesis	ko0097		20 out of 221	0.99689648		
	0	1 out of 53 1.88679245283019%	9.04977375565611%	8	1	
Biosynthesis of amino acids	ko0123		32 out of 221	0.99906761		
	0	2 out of 53 3.77358490566038%	14.4796380090498%	5	1	
Protein digestion and absorption	ko0497		24 out of 221	0.99909757		
	4	1 out of 53 1.88679245283019%	10.8597285067873%	9	1	
ABC transporters	ko0201		30 out of 221	0.99986623		
	0	1 out of 53 1.88679245283019%	13.5746606334842%	6	1	

PS2 vs PS3_KEGG_stat

Dopaminergic synapse	ko0472 8	3 out of 62 4.83870967741935%	3 out of 221 1.35746606334842%	0.02131148 5	1
Ubiquinone and other terpenoid-quinone biosynthesis	ko0013 0	3 out of 62 4.83870967741935%	4 out of 221 1.80995475113122%	0.06794257 7	1
Retrograde endocannabinoid signaling	ko0472 3	3 out of 62 4.83870967741935%	4 out of 221 1.80995475113122%	0.06794257 7	1
Phenylalanine metabolism	ko0036 0	5 out of 62 8.06451612903226%	9 out of 221 4.07239819004525%	0.07250532 7	1
Pertussis	ko0513 3	2 out of 62 3.2258064516129%	0.904977375565611%	0.07778691 9	1
Estrogen signaling pathway	ko0491 5	2 out of 62 3.2258064516129%	0.904977375565611%	0.07778691 9	1
Novobiocin biosynthesis	ko0040 1	2 out of 62 3.2258064516129%	0.904977375565611%	0.07778691 9	1
Propanoate metabolism	ko0064 0	2 out of 62 3.2258064516129%	0.904977375565611%	0.07778691 9	1
Melanogenesis	ko0491 6	2 out of 62 3.2258064516129%	0.904977375565611%	0.07778691 9	1
Tyrosine metabolism	ko0035 0	4 out of 62 6.45161290322581%	7 out of 221 3.16742081447964%	0.09873435 2	1
Phenylalanine, tyrosine and tryptophan biosynthesis	ko0040 0	4 out of 62 6.45161290322581%	7 out of 221 3.16742081447964%	0.09873435 2	1
Phenylpropanoid biosynthesis	ko0094 0	7 out of 62 11.2903225806452%	7.23981900452489%	0.12422984 4	1
Purine metabolism	ko0023 0	8 out of 62 12.9032258064516%	19 out of 221 8.5972850678733%	0.12473851 6	1
GABAergic synapse	ko0472 7	3 out of 62 4.83870967741935%	5 out of 221 2.26244343891403%	0.13584776 3	1
Nicotinate and nicotinamide metabolism	ko0076 0	5 out of 62 8.06451612903226%	4.97737556561086%	0.16399892 4	1
Cocaine addiction	ko0503 0	2 out of 62 3.2258064516129%	3 out of 221 1.35746606334842%	0.19073778 8	1
Plant hormone signal transduction	ko0407 5	2 out of 62 3.2258064516129%	3 out of 221 1.35746606334842%	0.19073778 8	1
Amphetamine addiction	ko0503 1	2 out of 62 3.2258064516129%	3 out of 221 1.35746606334842%	0.19073778 8	1
Caffeine metabolism	ko0023 2	2 out of 62 3.2258064516129%	3 out of 221 1.35746606334842%	0.19073778 8	1
cAMP signaling pathway	ko0402 4	3 out of 62 4.83870967741935%	6 out of 221 2.71493212669683%	0.21810944 6	1
Degradation of aromatic compounds	ko0122 0	4 out of 62 6.45161290322581%	9 out of 221 4.07239819004525%	0.22338982	1

	ko0106	10 out of 62	28 out of 221	0.22629652	
Biosynthesis of phenylpropanoids	1	16.1290322580645%	12.6696832579186%	1	1
	ko0492	1 out of 62	1 out of 221	0.28054298	
Oxytocin signaling pathway	1	1.61290322580645%	0.452488687782805%	6	1
	ko0426	1 out of 62	1 out of 221	0.28054298	
Adrenergic signaling in cardiomyocytes	1	1.61290322580645%	0.452488687782805%	6	1
	ko0491	1 out of 62	1 out of 221	0.28054298	
Thyroid hormone signaling pathway	9	1.61290322580645%	0.452488687782805%	6	1
	ko0421	1 out of 62	1 out of 221	0.28054298	
Longevity regulating pathway - multiple species	3	1.61290322580645%	0.452488687782805%	6	1
	ko0202	1 out of 62	1 out of 221	0.28054298	
Biofilm formation - <i>Pseudomonas aeruginosa</i>	5	1.61290322580645%	0.452488687782805%	6	1
	ko0541	1 out of 62	1 out of 221	0.28054298	
Dilated cardiomyopathy (DCM)	4	1.61290322580645%	0.452488687782805%	6	1
	ko0497	1 out of 62	1 out of 221	0.28054298	
Pancreatic secretion	2	1.61290322580645%	0.452488687782805%	6	1
	ko0401	1 out of 62	1 out of 221	0.28054298	
Ras signaling pathway	4	1.61290322580645%	0.452488687782805%	6	1
	ko0105	1 out of 62	1 out of 221	0.28054298	
Biosynthesis of enediyne antibiotics	9	1.61290322580645%	0.452488687782805%	6	1
	ko0402	1 out of 62	1 out of 221	0.28054298	
Calcium signaling pathway	0	1.61290322580645%	0.452488687782805%	6	1
	ko0096	1 out of 62	1 out of 221	0.28054298	
Betalain biosynthesis	5	1.61290322580645%	0.452488687782805%	6	1
	ko0516	1 out of 62	1 out of 221	0.28054298	
Human papillomavirus infection	5	1.61290322580645%	0.452488687782805%	6	1
	ko0511	1 out of 62	1 out of 221	0.28054298	
Vibrio cholerae infection	0	1.61290322580645%	0.452488687782805%	6	1
	ko0492	1 out of 62	1 out of 221	0.28054298	
Relaxin signaling pathway	6	1.61290322580645%	0.452488687782805%	6	1
	ko0497	1 out of 62	1 out of 221	0.28054298	
Salivary secretion	0	1.61290322580645%	0.452488687782805%	6	1
	ko0202	1 out of 62	1 out of 221	0.28054298	
Biofilm formation - <i>Escherichia coli</i>	6	1.61290322580645%	0.452488687782805%	6	1
	ko0152	1 out of 62	1 out of 221	0.28054298	
Endocrine resistance	2	1.61290322580645%	0.452488687782805%	6	1
	ko0411	1 out of 62	1 out of 221	0.28054298	
Oocyte meiosis	4	1.61290322580645%	0.452488687782805%	6	1
	ko0401	1 out of 62	1 out of 221	0.28054298	
MAPK signaling pathway	0	1.61290322580645%	0.452488687782805%	6	1
	ko0098	1 out of 62	1 out of 221	0.28054298	
Drug metabolism - other enzymes	3	1.61290322580645%	0.452488687782805%	6	1

	ko0496	1 out of 221	0.28054298		
Aldosterone-regulated sodium reabsorption	0	1 out of 62 1.61290322580645%	0.452488687782805%	6	1
	ko0491	1 out of 221	0.28054298		
GnRH signaling pathway	2	1 out of 62 1.61290322580645%	0.452488687782805%	6	1
	ko0090	1 out of 221	0.28054298		
Carotenoid biosynthesis	6	1 out of 62 1.61290322580645%	0.452488687782805%	6	1
	ko0401	1 out of 221	0.28054298		
Rap1 signaling pathway	5	1 out of 62 1.61290322580645%	0.452488687782805%	6	1
	ko0094	1 out of 221	0.28054298		
Anthocyanin biosynthesis	2	1 out of 62 1.61290322580645%	0.452488687782805%	6	1
	ko0105	1 out of 221	0.28054298		
Biosynthesis of vancomycin group antibiotics	5	1 out of 62 1.61290322580645%	0.452488687782805%	6	1
	ko0467	1 out of 221	0.28054298		
Leukocyte transendothelial migration	0	1 out of 62 1.61290322580645%	0.452488687782805%	6	1
	ko0502	1 out of 221	0.28054298		
Prion diseases	0	1 out of 62 1.61290322580645%	0.452488687782805%	6	1
	ko0491	1 out of 221	0.28054298		
Progesterone-mediated oocyte maturation	4	1 out of 62 1.61290322580645%	0.452488687782805%	6	1
	ko0437	1 out of 221	0.28054298		
Apelin signaling pathway	1	1 out of 62 1.61290322580645%	0.452488687782805%	6	1
	ko0496	1 out of 221	0.28054298		
Vasopressin-regulated water reabsorption	2	1 out of 62 1.61290322580645%	0.452488687782805%	6	1
	ko0711	1 out of 221	0.28054298		
Benzoic acid family	0	1 out of 62 1.61290322580645%	0.452488687782805%	6	1
	ko0516	1 out of 221	0.28054298		
HTLV-I infection	6	1 out of 62 1.61290322580645%	0.452488687782805%	6	1
	ko0406	1 out of 221	0.28054298		
Chemokine signaling pathway	2	1 out of 62 1.61290322580645%	0.452488687782805%	6	1
	ko0434	1 out of 221	0.28054298		
Hedgehog signaling pathway	0	1 out of 62 1.61290322580645%	0.452488687782805%	6	1
	ko0093	1 out of 221	0.28054298		
Caprolactam degradation	0	1 out of 62 1.61290322580645%	0.452488687782805%	6	1
	ko0036	1 out of 221	0.28054298		
Chlorocyclohexane and chlorobenzene degradation	1	1 out of 62 1.61290322580645%	0.452488687782805%	6	1
	ko0461	1 out of 221	0.28054298		
Platelet activation	1	1 out of 62 1.61290322580645%	0.452488687782805%	6	1
	ko0471	1 out of 221	0.28054298		
Circadian rhythm	0	1 out of 62 1.61290322580645%	0.452488687782805%	6	1
	ko0007	1 out of 221	0.28054298		
Fatty acid degradation	1	1 out of 62 1.61290322580645%	0.452488687782805%	6	1
	ko0496	1 out of 221	0.28054298		
Endocrine and other factor-regulated calcium reabsorption	1	1 out of 62 1.61290322580645%	0.452488687782805%	6	1

Puromycin biosynthesis	ko0023	1	1 out of 62 1.61290322580645%	0.452488687782805%	1 out of 221	0.28054298	6	1
Tryptophan metabolism	ko0038	0	4 out of 62 6.45161290322581%	4.52488687782805%	10 out of 221	0.29645966	9	1
	ko0096						0.30764076	
Tropane, piperidine and pyridine alkaloid biosynthesis	ko0095	0	3 out of 62 4.83870967741935%	7 out of 221 3.16742081447964%	6		1	
Isoquinoline alkaloid biosynthesis	ko0503	0	2 out of 62 3.2258064516129%	4 out of 221 1.80995475113122%	8		1	
Alcoholism	ko0503	4	2 out of 62 3.2258064516129%	4 out of 221 1.80995475113122%	8		1	
Morphine addiction	ko0065	2	2 out of 62 3.2258064516129%	4 out of 221 1.80995475113122%	8		1	
Butanoate metabolism	ko0107	0	2 out of 62 3.2258064516129%	4 out of 221 1.80995475113122%	8		1	
Biosynthesis of plant hormones	ko0492	0	5 out of 62 8.06451612903226%	6.78733031674208%	5		1	
Glucagon signaling pathway	ko0106	2	2 out of 62 3.2258064516129%	5 out of 221 2.26244343891403%	2		1	
Biosynthesis of terpenoids and steroids	ko0491	2	2 out of 62 3.2258064516129%	5 out of 221 2.26244343891403%	2		1	
Thyroid hormone synthesis	ko0501	8	2 out of 62 3.2258064516129%	5 out of 221 2.26244343891403%	2		1	
Parkinson's disease	ko0421	2	2 out of 62 3.2258064516129%	5 out of 221 2.26244343891403%	2		1	
Longevity regulating pathway - worm	ko0202	2	1 out of 62 1.61290322580645%	0.904977375565611%	4		1	
Quorum sensing	ko0472	4	1 out of 62 1.61290322580645%	0.904977375565611%	4		1	
Long-term potentiation	ko0059	0	1 out of 62 1.61290322580645%	0.904977375565611%	4		1	
alpha-Linolenic acid metabolism	ko0472	2	1 out of 62 1.61290322580645%	0.904977375565611%	4		1	
Cholinergic synapse	ko0059	5	1 out of 62 1.61290322580645%	0.904977375565611%	4		1	
Linoleic acid metabolism	ko0079	1	1 out of 62 1.61290322580645%	0.904977375565611%	4		1	
Folate biosynthesis	ko0514	0	1 out of 62 1.61290322580645%	0.904977375565611%	4		1	
Amoebiasis	ko0040	6	1 out of 62 1.61290322580645%	0.904977375565611%	4		1	
Benzoxazinoid biosynthesis		2	1 out of 62 1.61290322580645%	0.904977375565611%	4		1	

Ovarian steroidogenesis	ko0491 3	1 out of 62 1.61290322580645%	2 out of 221 0.904977375565611%	0.48329905 4	1
Bisphenol degradation	ko0036 3	1 out of 62 1.61290322580645%	2 out of 221 0.904977375565611%	0.48329905 4	1
Staurosporine biosynthesis	ko0040 4	1 out of 62 1.61290322580645%	2 out of 221 0.904977375565611%	0.48329905 4	1
Nicotine addiction	ko0503 3	1 out of 62 1.61290322580645%	2 out of 221 0.904977375565611%	0.48329905 4	1
Chagas disease (American trypanosomiasis)	ko0514 2	1 out of 62 1.61290322580645%	2 out of 221 0.904977375565611%	0.48329905 4	1
Cell cycle - yeast	ko0411 1	1 out of 62 1.61290322580645%	2 out of 221 0.904977375565611%	0.48329905 4	1
Phospholipase D signaling pathway	ko0497 2	1 out of 62 1.61290322580645%	2 out of 221 0.904977375565611%	0.48329905 4	1
Gastric acid secretion	ko0471 1	1 out of 62 1.61290322580645%	2 out of 221 0.904977375565611%	0.48329905 4	1
Circadian entrainment	ko0427 3	1 out of 62 1.61290322580645%	2 out of 221 0.904977375565611%	0.48329905 4	1
Vascular smooth muscle contraction	ko0033 0	1 out of 62 1.61290322580645%	2 out of 221 0.904977375565611%	0.48329905 4	1
Prodigiosin biosynthesis	ko0491 3	1 out of 62 1.61290322580645%	2 out of 221 0.904977375565611%	0.48329905 4	1
Insulin signaling pathway	ko0411 0	1 out of 62 1.61290322580645%	2 out of 221 0.904977375565611%	0.48329905 4	1
Meiosis - yeast	ko0033 3	1 out of 62 1.61290322580645%	2 out of 221 0.904977375565611%	0.52005100 4	1
Arginine and proline metabolism	ko0106 0	4 out of 62 6.45161290322581%	13 out of 221 5.88235294117647%	0.52005100 3	1
Biosynthesis of alkaloids derived from shikimate pathway	ko0024 3	4 out of 62 6.45161290322581%	13 out of 221 5.88235294117647%	0.58808356 3	1
Pyrimidine metabolism	ko0094 0	4 out of 62 6.45161290322581%	14 out of 221 6.33484162895928%	0.58808356 2	1
Flavone and flavonol biosynthesis	ko0036 4	4 out of 62 6.45161290322581%	14 out of 221 6.33484162895928%	0.62957968 2	1
Benzoate degradation	ko0472 2	1 out of 62 1.61290322580645%	3 out of 221 1.35746606334842%	0.62957968 7	1
Serotonergic synapse	ko0511 6	1 out of 62 1.61290322580645%	3 out of 221 1.35746606334842%	0.62957968 7	1
Biofilm formation - Vibrio cholerae	ko0520 1	1 out of 62 1.61290322580645%	3 out of 221 1.35746606334842%	0.62957968 7	1
Pathways in cancer	ko0491 0	1 out of 62 1.61290322580645%	3 out of 221 1.35746606334842%	0.62957968 7	1

Gap junction	ko0454	0	1 out of 62 1.61290322580645%	3 out of 221 1.35746606334842%	0.62957968	7
	ko0491	1	1 out of 62 1.61290322580645%	3 out of 221 1.35746606334842%	0.62957968	7
Insulin secretion	ko0492	4	1 out of 62 1.61290322580645%	3 out of 221 1.35746606334842%	0.62957968	7
	ko0472	4	1 out of 62 1.61290322580645%	3 out of 221 1.35746606334842%	0.62957968	7
Renin secretion	ko0421	1	1 out of 62 1.61290322580645%	3 out of 221 1.35746606334842%	0.62957968	7
	ko0002	0	1 out of 62 1.61290322580645%	3 out of 221 1.35746606334842%	0.62957968	7
Glutamatergic synapse	ko0491	7	1 out of 62 1.61290322580645%	3 out of 221 1.35746606334842%	0.62957968	7
	ko0523	1	1 out of 62 1.61290322580645%	3 out of 221 1.35746606334842%	0.62957968	7
Longevity regulating pathway	ko0068	0	1 out of 62 1.61290322580645%	3 out of 221 1.35746606334842%	0.62957968	7
	ko0492	3	1 out of 62 1.61290322580645%	3 out of 221 1.35746606334842%	0.62957968	7
Citrate cycle (TCA cycle)	ko0062	0	1 out of 62 1.61290322580645%	3 out of 221 1.35746606334842%	0.62957968	7
	ko0003	0	1 out of 62 1.61290322580645%	3 out of 221 1.35746606334842%	0.62957968	7
Prolactin signaling pathway	ko0031	0	1 out of 62 1.61290322580645%	3 out of 221 1.35746606334842%	0.63144570	7
	ko0096	6	2 out of 62 3.2258064516129%	7 out of 221 3.16742081447964%	0.63144570	7
Choline metabolism in cancer	ko0041	0	2 out of 62 3.2258064516129%	7 out of 221 3.16742081447964%	0.63144570	7
	ko0475	0	2 out of 62 3.2258064516129%	7 out of 221 3.16742081447964%	0.70911854	3
Methane metabolism	ko0046	0	2 out of 62 3.2258064516129%	8 out of 221 3.61990950226244%	0.73492858	3
	ko0472	1	1 out of 62 1.61290322580645%	4 out of 221 1.80995475113122%	0.73492858	3
Regulation of lipolysis in adipocytes	ko0072	0	1 out of 62 1.61290322580645%	4 out of 221 1.80995475113122%	0.73492858	3
	ko0475	1	1 out of 62 1.61290322580645%	4 out of 221 1.80995475113122%	0.73492858	3
Pyruvate metabolism	ko0106	6	1 out of 62 1.61290322580645%	4 out of 221 1.80995475113122%	0.73492858	3
	ko0474	0	1 out of 62 1.61290322580645%	4 out of 221 1.80995475113122%	0.73492858	3
Pentose phosphate pathway						
Lysine degradation						
Glucosinolate biosynthesis						
beta-Alanine metabolism						
Cyanoamino acid metabolism						
Inflammatory mediator regulation of TRP channels						
Carbon fixation pathways in prokaryotes						
Synaptic vesicle cycle						
Biosynthesis of alkaloids derived from terpenoid and polyketide						
Olfactory transduction						

	ko0062			0.73492858		
Toluene degradation	3	1 out of 62 1.61290322580645%	4 out of 221 1.80995475113122%		3	1
	ko0050			0.73492858		
Starch and sucrose metabolism	0	1 out of 62 1.61290322580645%	4 out of 221 1.80995475113122%		3	1
	ko0033			0.73492858		
Carbapenem biosynthesis	2	1 out of 62 1.61290322580645%	4 out of 221 1.80995475113122%		3	1
	ko0019			0.73492858		
Oxidative phosphorylation	0	1 out of 62 1.61290322580645%	4 out of 221 1.80995475113122%		3	1
	ko0028			0.73492858		
Valine, leucine and isoleucine degradation	0	1 out of 62 1.61290322580645%	4 out of 221 1.80995475113122%		3	1
	ko0111	22 out of 62	84 out of 221	0.73680994		
Biosynthesis of secondary metabolites	0	35.4838709677419%	38.0090497737557%		9	1
Biosynthesis of alkaloids derived from ornithine, lysine and nicotinic acid	4	3 out of 62 4.83870967741935%	5.88235294117647%		2	1
	ko0106			0.75989460		
Biosynthesis of alkaloids derived from histidine and purine	5	3 out of 62 4.83870967741935%	5.88235294117647%		2	1
	ko0112			0.80280981		
Microbial metabolism in diverse environments	0	9 out of 62 14.5161290322581%	17.1945701357466%		8	1
	ko0421			0.81066327		
Ferroptosis	6	1 out of 62 1.61290322580645%	5 out of 221 2.26244343891403%		4	1
	ko0056			0.81066327		
Glycerophospholipid metabolism	4	1 out of 62 1.61290322580645%	5 out of 221 2.26244343891403%		4	1
	ko0402			0.81066327		
cGMP-PKG signaling pathway	2	1 out of 62 1.61290322580645%	5 out of 221 2.26244343891403%		4	1
	ko0073			0.81066327		
Thiamine metabolism	0	1 out of 62 1.61290322580645%	5 out of 221 2.26244343891403%		4	1
	ko0492			0.81066327		
Aldosterone synthesis and secretion	5	1 out of 62 1.61290322580645%	5 out of 221 2.26244343891403%		4	1
	ko0092			0.81066327		
Sulfur metabolism	0	1 out of 62 1.61290322580645%	5 out of 221 2.26244343891403%		4	1
	ko0415			0.81066327		
AMPK signaling pathway	2	1 out of 62 1.61290322580645%	5 out of 221 2.26244343891403%		4	1
	ko0063			0.81066327		
Glyoxylate and dicarboxylate metabolism	0	1 out of 62 1.61290322580645%	5 out of 221 2.26244343891403%		4	1
	ko0094			0.81066327		
Isoflavonoid biosynthesis	3	1 out of 62 1.61290322580645%	5 out of 221 2.26244343891403%		4	1
	ko0497			0.82431413		
Bile secretion	6	2 out of 62 3.2258064516129%	4.52488687782805%		3	1
	ko0025			0.82431413		
Alanine, aspartate and glutamate metabolism	0	2 out of 62 3.2258064516129%	4.52488687782805%		3	1
	ko0474			0.82431413		
Taste transduction	2	2 out of 62 3.2258064516129%	4.52488687782805%		3	1

2-Oxocarboxylic acid metabolism	ko0121 0 ko0123 0 ko0497 4 ko0026 1 ko0077 0 ko0030 0 ko0090 8 ko0027 0 ko0120 0 ko0113 0 ko0110 0 ko0094 1 ko0062 7 ko0206 0 ko0523 0 ko0106 0 ko0022 0 ko0497 8 ko0026 0 ko0408 0 ko0097 0	3 out of 62 4.83870967741935% 7 out of 62 11.2903225806452% 5 out of 62 8.06451612903226% 1 out of 62 1.61290322580645% 1 out of 62 1.61290322580645% 1 out of 62 1.61290322580645% 1 out of 62 1.61290322580645% 2 out of 62 3.2258064516129% 2 out of 62 3.2258064516129% 8 out of 62 12.9032258064516% 40 out of 62 64.5161290322581% 4 out of 62 6.45161290322581% 2 out of 62 3.2258064516129% 1 out of 62 1.61290322580645% 4 out of 62 6.45161290322581% 8 out of 62 12.9032258064516% 1 out of 62 1.61290322580645% 1 out of 62 1.61290322580645%	15 out of 221 32 out of 221 24 out of 221 10.8597285067873% 6 out of 221 2.71493212669683% 6 out of 221 2.71493212669683% 6 out of 221 2.71493212669683% 11 out of 221 11 out of 221 37 out of 221 154 out of 221 69.683257918552% 21 out of 221 12 out of 221 5.42986425339367% 7 out of 221 3.16742081447964% 22 out of 221 40 out of 221 18.0995475113122% 8 out of 221 3.61990950226244% 14 out of 221 6.33484162895928% 9 out of 221 4.07239819004525% 9 out of 221 4.07239819004525% 20 out of 221 9.04977375565611%	0.84583784 0.85482286 0.86000702 0.86500992 0.86500992 0.86500992 0.86500992 0.86521054 0.86521054 0.86521054 0.87761867 0.88551301 0.89272807 0.89734343 0.90393729 0.91432442 0.92958078 0.93176854 0.94159779 0.95162934 0.95162934 0.95547776 0.95547776	9 3 2 7 7 7 7 1 1 1 1 1 1 6 6 4 7 9 7 1 6 6 9	1 1
Biosynthesis of amino acids						
Protein digestion and absorption						
Monobactam biosynthesis						
Pantothenate and CoA biosynthesis						
Lysine biosynthesis						
Zeatin biosynthesis						
Cysteine and methionine metabolism						
Carbon metabolism						
Biosynthesis of antibiotics						
Metabolic pathways						
Flavonoid biosynthesis						
Aminobenzoate degradation						
Phosphotransferase system (PTS)						
Central carbon metabolism in cancer						
Biosynthesis of plant secondary metabolites						
Arginine biosynthesis						
Mineral absorption						
Glycine, serine and threonine metabolism						
Neuroactive ligand-receptor interaction						
Aminoacyl-tRNA biosynthesis						

	ko0202	10 out of 221				
Two-component system	0	1 out of 62 1.61290322580645%	4.52488687782805%	0.96577548	1	
	ko0497	11 out of 221		0.97583197		
Vitamin digestion and absorption	7	1 out of 62 1.61290322580645%	4.97737556561086%	4	1	
	ko0048	12 out of 221		0.98296729		
Glutathione metabolism	0	1 out of 62 1.61290322580645%	5.42986425339367%	6	1	
	ko0201	30 out of 221		0.99966783		
ABC transporters	0	2 out of 62 3.2258064516129%	13.5746606334842%	2	1	

PS3 vs PS4_KEGG_stat						
	ko0023			0.01104455		
Caffeine metabolism	2	3 out of 50 6%	3 out of 221 1.35746606334842%	6	1	
	ko0106		13 out of 221	0.04696710		
Biosynthesis of alkaloids derived from histidine and purine	5	6 out of 50 12%	5.88235294117647%	7	1	
	ko0059		2 out of 221	0.05039078		
Linoleic acid metabolism	1	2 out of 50 4%	0.904977375565611%	6	1	
	ko0047		2 out of 221	0.05039078		
D-Glutamine and D-glutamate metabolism	1	2 out of 50 4%	0.904977375565611%	6	1	
	ko0496		2 out of 221	0.05039078		
Proximal tubule bicarbonate reclamation	4	2 out of 50 4%	0.904977375565611%	6	1	
	ko0091		2 out of 221	0.05039078		
Nitrogen metabolism	0	2 out of 50 4%	0.904977375565611%	6	1	
	ko0094			0.07775665		
Isoflavonoid biosynthesis	3	3 out of 50 6%	5 out of 221 2.26244343891403%	2	1	
	ko0408			0.11977911		
Neuroactive ligand-receptor interaction	0	4 out of 50 8%	9 out of 221 4.07239819004525%	9	1	
	ko0472			0.12908324		
Glutamatergic synapse	4	2 out of 50 4%	3 out of 221 1.35746606334842%	6	1	
	ko0523			0.12908324		
Choline metabolism in cancer	1	2 out of 50 4%	3 out of 221 1.35746606334842%	6	1	
	ko0036			0.12908324		
Benzoate degradation	2	2 out of 50 4%	3 out of 221 1.35746606334842%	6	1	
	ko0034			0.13085854		
Histidine metabolism	0	3 out of 50 6%	6 out of 221 2.71493212669683%	6	1	
	ko0497		24 out of 221	0.14289466		
Protein digestion and absorption	4	8 out of 50 16%	10.8597285067873%	4	1	
	ko0096			0.19309890		
Tropane, piperidine and pyridine alkaloid biosynthesis	0	3 out of 50 6%	7 out of 221 3.16742081447964%	5	1	
	ko0031			0.19309890		
Lysine degradation	0	3 out of 50 6%	7 out of 221 3.16742081447964%	5	1	
	ko0472			0.22113176		
Retrograde endocannabinoid signaling	3	2 out of 50 4%	4 out of 221 1.80995475113122%	5	1	
	ko0033			0.22113176		
Carbapenem biosynthesis	2	2 out of 50 4%	4 out of 221 1.80995475113122%	5	1	

Toluene degradation	ko0062	3	2 out of 50 4%	4 out of 221 1.80995475113122%	0.22113176	5	1
Ether lipid metabolism	ko0056	5	1 out of 50 2%	1 out of 221 0.452488687782805%	0.22624434	4	1
Long-term depression	ko0473	0	1 out of 50 2%	1 out of 221 0.452488687782805%	0.22624434	4	1
Huntington's disease	ko0501	6	1 out of 50 2%	1 out of 221 0.452488687782805%	0.22624434	4	1
Dioxin degradation	ko0062	1	1 out of 50 2%	1 out of 221 0.452488687782805%	0.22624434	4	1
Bacterial chemotaxis	ko0203	0	1 out of 50 2%	1 out of 221 0.452488687782805%	0.22624434	4	1
Anthocyanin biosynthesis	ko0094	2	1 out of 50 2%	1 out of 221 0.452488687782805%	0.22624434	4	1
C5-Branched dibasic acid metabolism	ko0066	0	1 out of 50 2%	1 out of 221 0.452488687782805%	0.22624434	4	1
Caprolactam degradation	ko0093	0	1 out of 50 2%	1 out of 221 0.452488687782805%	0.22624434	4	1
Fatty acid degradation	ko0007	1	1 out of 50 2%	1 out of 221 0.452488687782805%	0.22624434	4	1
Benzoic acid family	ko0711	0	1 out of 50 2%	1 out of 221 0.452488687782805%	0.22624434	4	1
Arginine biosynthesis	ko0022	0	1 out of 50 2%	8 out of 221 3.61990950226244%	0.26109795	2	1
Metabolic pathways	ko0110	0	37 out of 50 74%	154 out of 221 69.683257918552%	0.28399863	6	1
Glycerophospholipid metabolism	ko0056	4	2 out of 50 4%	5 out of 221 2.26244343891403%	0.31671517	4	1
Valine, leucine and isoleucine biosynthesis	ko0029	0	2 out of 50 4%	5 out of 221 2.26244343891403%	0.31671517	4	1
GABAergic synapse	ko0472	7	2 out of 50 4%	5 out of 221 2.26244343891403%	0.31671517	4	1
Taurine and hypotaurine metabolism	ko0043	0	2 out of 50 4%	5 out of 221 2.26244343891403%	0.31671517	4	1
Glyoxylate and dicarboxylate metabolism	ko0063	0	2 out of 50 4%	5 out of 221 2.26244343891403%	0.31671517	4	1
Ferroptosis	ko0421	6	2 out of 50 4%	5 out of 221 2.26244343891403%	0.31671517	4	1
Biosynthesis of alkaloids derived from ornithine, lysine and nicotinic acid	ko0106	4	4 out of 50 8%	13 out of 221 5.88235294117647%	0.33427111	8	1
Arginine and proline metabolism	ko0033	0	4 out of 50 8%	13 out of 221 5.88235294117647%	0.33427111	8	1

Microbial metabolism in diverse environments	ko0112 0	10 out of 50 20%	38 out of 221 17.1945701357466%	0.34239703 1	1
ABC transporters	ko0201 0	8 out of 50 16%	30 out of 221 13.5746606334842%	0.35874073 6	1
mTOR signaling pathway	ko0415 0	1 out of 50 2%	2 out of 221 0.904977375565611%	0.40209790 2	1
Folate biosynthesis	ko0079 0	1 out of 50 2%	2 out of 221 0.904977375565611%	0.40209790 2	1
Porphyrin and chlorophyll metabolism	ko0086 0	1 out of 50 2%	2 out of 221 0.904977375565611%	0.40209790 2	1
Nicotine addiction	ko0503 3	1 out of 50 2%	2 out of 221 0.904977375565611%	0.40209790 2	1
Phototransduction	ko0474 4	1 out of 50 2%	2 out of 221 0.904977375565611%	0.40209790 2	1
Circadian entrainment	ko0471 3	1 out of 50 2%	2 out of 221 0.904977375565611%	0.40209790 2	1
Phospholipase D signaling pathway	ko0407 2	1 out of 50 2%	2 out of 221 0.904977375565611%	0.40209790 2	1
Long-term potentiation	ko0040 0	1 out of 50 2%	2 out of 221 0.904977375565611%	0.40209790 2	1
Benzoxazinoid biosynthesis	ko0071 2	1 out of 50 2%	2 out of 221 0.904977375565611%	0.40209790 2	1
Carbon fixation in photosynthetic organisms	ko0501 0	1 out of 50 2%	2 out of 221 0.904977375565611%	0.40209790 2	1
Amyotrophic lateral sclerosis (ALS)	ko0513 4	1 out of 50 2%	2 out of 221 0.904977375565611%	0.40209790 2	1
Pertussis	ko0036 3	1 out of 50 2%	2 out of 221 0.904977375565611%	0.40209790 2	1
Bisphenol degradation	ko0038 3	1 out of 50 2%	10 out of 221 0.904977375565611%	0.40246439 2	1
Tryptophan metabolism	ko0202 0	3 out of 50 6%	10 out of 221 4.52488687782805%	0.40246439 8	1
Two-component system	ko0025 0	3 out of 50 6%	10 out of 221 4.52488687782805%	0.40246439 8	1
Alanine, aspartate and glutamate metabolism	ko0030 0	3 out of 50 6%	10 out of 221 4.52488687782805%	0.40964348 8	1
Lysine biosynthesis	ko0077 0	2 out of 50 4%	6 out of 221 2.71493212669683%	0.40964348 8	1
Pantothenate and CoA biosynthesis	ko0106 0	2 out of 50 4%	6 out of 221 2.71493212669683%	0.41628772 8	1
Biosynthesis of plant secondary metabolites		10 out of 50 20%	40 out of 221 18.0995475113122%	1	1

Biosynthesis of secondary metabolites	ko0111 0 ko0023	20 out of 50 40% 5 out of 50 10%	84 out of 221 38.0090497737557% 19 out of 221 8.5972850678733%	0.43209234 7 0.43685733	
Purine metabolism	ko0121 0	4 out of 50 8%	15 out of 221 6.78733031674208%	0.45238205 4 5	1
2-Oxocarboxylic acid metabolism	ko0076 0	3 out of 50 6%	11 out of 221 4.97737556561086%	0.47116080 2 2	1
Nicotinate and nicotinamide metabolism	ko0120 0	3 out of 50 6%	11 out of 221 4.97737556561086%	0.47116080 2 2	1
Carbon metabolism	ko0097 0	5 out of 50 10%	20 out of 221 9.04977375565611%	0.48826943 7 0.49626132	1
Aminoacyl-tRNA biosynthesis	ko0035 0	2 out of 50 4%	7 out of 221 3.16742081447964%	0.49626132 2 0.49626132	1
Tyrosine metabolism	ko0041 0	2 out of 50 4%	7 out of 221 3.16742081447964%	0.49626132 2 0.53633151	1
beta-Alanine metabolism	ko0048 0	3 out of 50 6%	12 out of 221 5.42986425339367%	0.53633151 8 8	1
Glutathione metabolism	ko0503 1	1 out of 50 2%	3 out of 221 1.35746606334842%	0.53860523 1 0.53860523	1
Amphetamine addiction	ko0406 8	1 out of 50 2%	3 out of 221 1.35746606334842%	0.53860523 1 0.53860523	1
FoxO signaling pathway	ko0012 0	1 out of 50 2%	3 out of 221 1.35746606334842%	0.53860523 1 0.53860523	1
Primary bile acid biosynthesis	ko0062 0	1 out of 50 2%	3 out of 221 1.35746606334842%	0.53860523 1 0.53860523	1
Pyruvate metabolism	ko0454 0	1 out of 50 2%	3 out of 221 1.35746606334842%	0.53860523 1 0.53860523	1
Gap junction	ko0472 0	1 out of 50 2%	3 out of 221 1.35746606334842%	0.53860523 1 0.53860523	1
Dopaminergic synapse	ko0003 8	1 out of 50 2%	3 out of 221 1.35746606334842%	0.53860523 1 0.53860523	1
Pentose phosphate pathway	ko0503 0	1 out of 50 2%	3 out of 221 1.35746606334842%	0.53860523 1 0.53860523	1
Cocaine addiction	ko0523 0	1 out of 50 2%	3 out of 221 1.35746606334842% 22 out of 221	0.53860523 1 0.58589714	1
Central carbon metabolism in cancer	ko0123 0	5 out of 50 10%	9.95475113122172% 32 out of 221	0.62170716 4 0.64404675	1
Biosynthesis of amino acids	ko0026 0	7 out of 50 14%	14.4796380090498% 9 out of 221 4.07239819004525%	0.64404675 3 0.64404675	1
Glycine, serine and threonine metabolism	ko0036 0	2 out of 50 4%	9 out of 221 4.07239819004525%	0.64404675 3 0.64404675	1
Phenylalanine metabolism		2 out of 50 4%	9 out of 221 4.07239819004525%		1

Degradation of aromatic compounds	ko0122 0	2 out of 50 4%	9 out of 221 4.07239819004525%	0.64404675 3	1
Synaptic vesicle cycle	ko0472 1	1 out of 50 2%	4 out of 221 1.80995475113122%	0.64442971 9	1
Valine, leucine and isoleucine degradation	ko0028 0	1 out of 50 2%	4 out of 221 1.80995475113122%	0.64442971 9	1
Butanoate metabolism	ko0065 0	1 out of 50 2%	4 out of 221 1.80995475113122%	0.64442971 9	1
Olfactory transduction	ko0474 0	1 out of 50 2%	4 out of 221 1.80995475113122%	0.64442971 9	1
Neomycin, kanamycin and gentamicin biosynthesis	ko0052 4	1 out of 50 2%	4 out of 221 1.80995475113122%	0.64442971 9	1
Starch and sucrose metabolism	ko0503 0	1 out of 50 2%	4 out of 221 1.80995475113122%	0.64442971 9	1
Alcoholism	ko0050 4	1 out of 50 2%	4 out of 221 1.80995475113122%	0.64442971 9	1
Vitamin B6 metabolism	ko0013 0	1 out of 50 2%	4 out of 221 1.80995475113122%	0.64442971 9	1
Ubiquinone and other terpenoid-quinone biosynthesis	ko0024 0	1 out of 50 2%	4 out of 221 1.80995475113122%	0.65240144 9	1
Pyrimidine metabolism	ko0474 0	3 out of 50 6%	14 out of 221 6.33484162895928%	0.70444234 1	1
Taste transduction	ko0402 2	2 out of 50 4%	10 out of 221 4.52488687782805%	0.72635835 1	1
cGMP-PKG signaling pathway	ko0092 2	1 out of 50 2%	5 out of 221 2.26244343891403%	0.72635835 5	1
Sulfur metabolism	ko0094 0	1 out of 50 2%	5 out of 221 2.26244343891403%	0.74694010 5	1
Phenylpropanoid biosynthesis	ko0026 0	3 out of 50 6%	16 out of 221 7.23981900452489%	0.78970132 4	1
Monobactam biosynthesis	ko0152 1	1 out of 50 2%	6 out of 221 2.71493212669683%	0.78970132 8	1
Antifolate resistance	ko0090 3	1 out of 50 2%	6 out of 221 2.71493212669683%	0.78970132 8	1
Zeatin biosynthesis	ko0062 8	1 out of 50 2%	6 out of 221 2.71493212669683%	0.78970132 8	1
Aminobenzoate degradation	ko0106 7	2 out of 50 4%	12 out of 221 5.42986425339367%	0.80027291 6	1
Biosynthesis of phenylpropanoids	ko0096 1	5 out of 50 10%	28 out of 221 12.6696832579186%	0.81031811 4	1
Glucosinolate biosynthesis	ko0096 6	1 out of 50 2%	7 out of 221 3.16742081447964%	0.83860799 6	1

	ko0206			0.83860799		
Phosphotransferase system (PTS)	0	1 out of 50 2%	7 out of 221 3.16742081447964%	6	1	
	ko0040			0.83860799		
Phenylalanine, tyrosine and tryptophan biosynthesis	0	1 out of 50 2%	7 out of 221 3.16742081447964%	6	1	
	ko0497		14 out of 221	0.86804734		
Mineral absorption	8	2 out of 50 4%	6.33484162895928%	2	1	
	ko0046			0.87631640		
Cyanoamino acid metabolism	0	1 out of 50 2%	8 out of 221 3.61990950226244%	8	1	
	ko0113		37 out of 221	0.89510301		
Biosynthesis of antibiotics	0	6 out of 50 12%	16.7420814479638%	2	1	
	ko0497		10 out of 221	0.92767320		
Bile secretion	6	1 out of 50 2%	4.52488687782805%	1	1	
	ko0027		11 out of 221	0.94481225		
Cysteine and methionine metabolism	0	1 out of 50 2%	4.97737556561086%	3	1	
	ko0106		13 out of 221	0.96801147		
Biosynthesis of alkaloids derived from shikimate pathway	3	1 out of 50 2%	5.88235294117647%	7	1	
	ko0094		21 out of 221	0.97300535		
Flavonoid biosynthesis	1	2 out of 50 4%	9.50226244343891%	8	1	
	ko0094		14 out of 221	0.97570102		
Flavone and flavonol biosynthesis	4	1 out of 50 2%	6.33484162895928%	6	1	
	ko0107		15 out of 221	0.98157034		
Biosynthesis of plant hormones	0	1 out of 50 2%	6.78733031674208%	3	1	