

## Supplementary Information

A Holistic View into the Gallic Acid-induced Attenuation in Colitis

Based on the Microbiome-Metabolomics Analysis

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**Figure 1. LDA score plot**

**Figure 2. PCA score plot based on PICRUSt analysis**

**Figure 3. PCA scores plots of <sup>1</sup>H NMR data of feces and urine**

**Table 1. Relative abundance of major bacterial taxa among different groups**

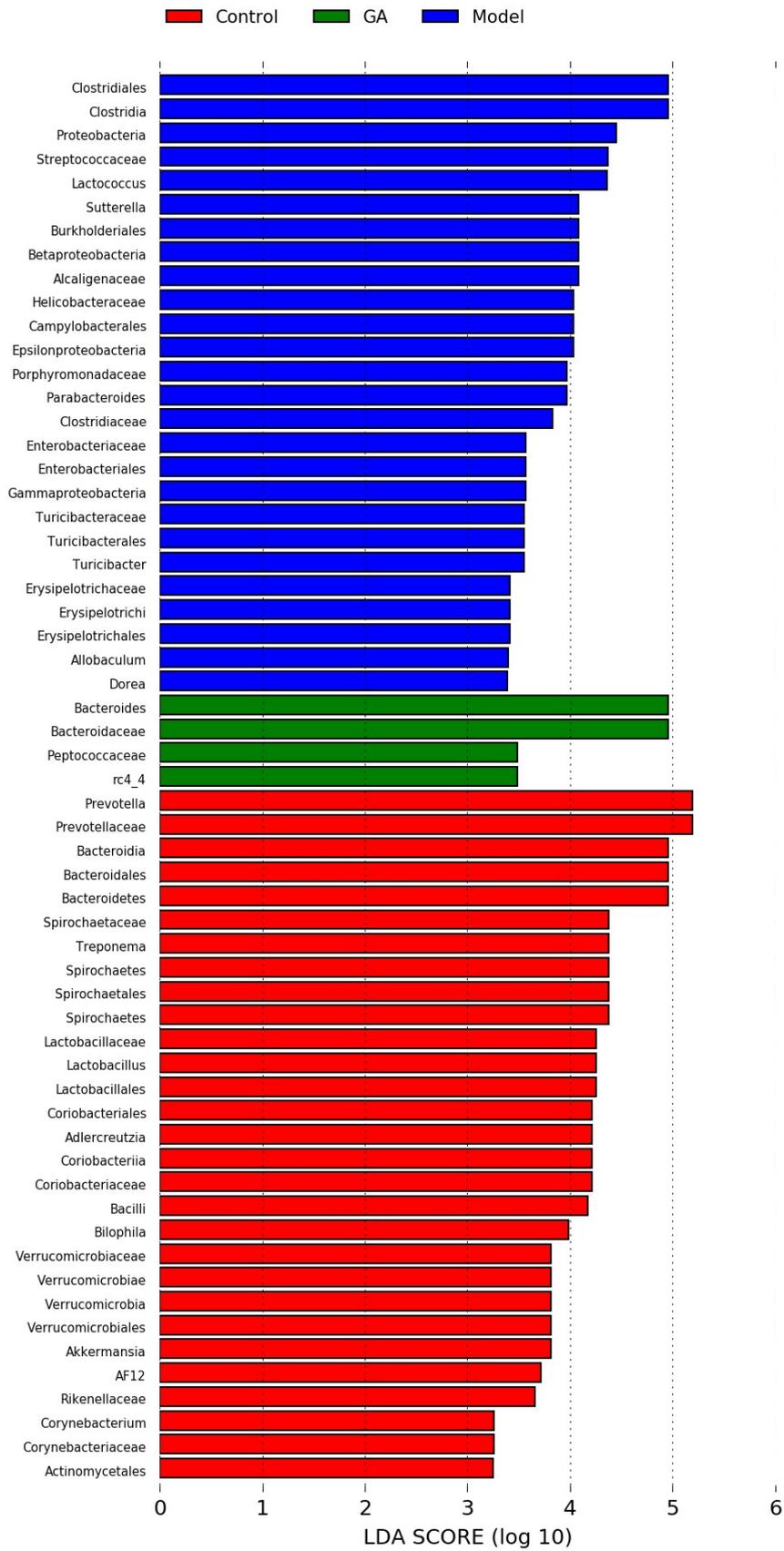
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**Table 3. <sup>1</sup>H NMR data of metabolites in rat urine**

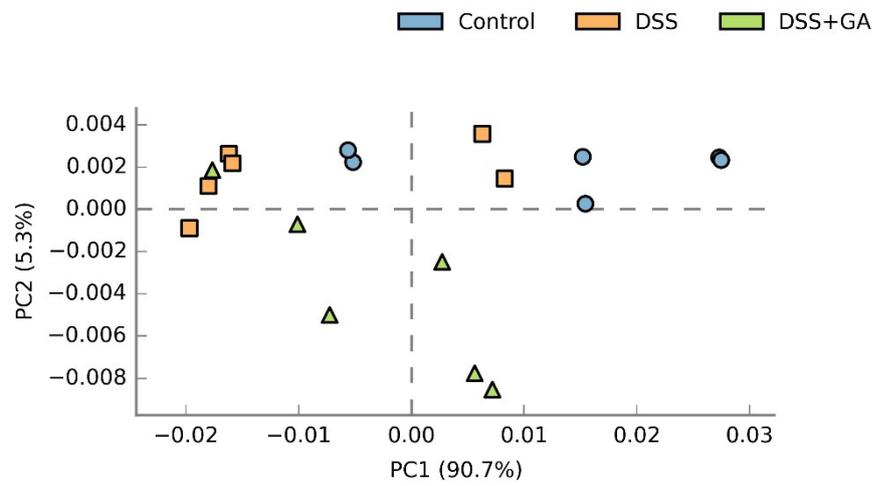
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**Table 5. Screened metabolites with significant difference in fold change (FC) value in urine samples**

**Figure 1. LDA score plot**

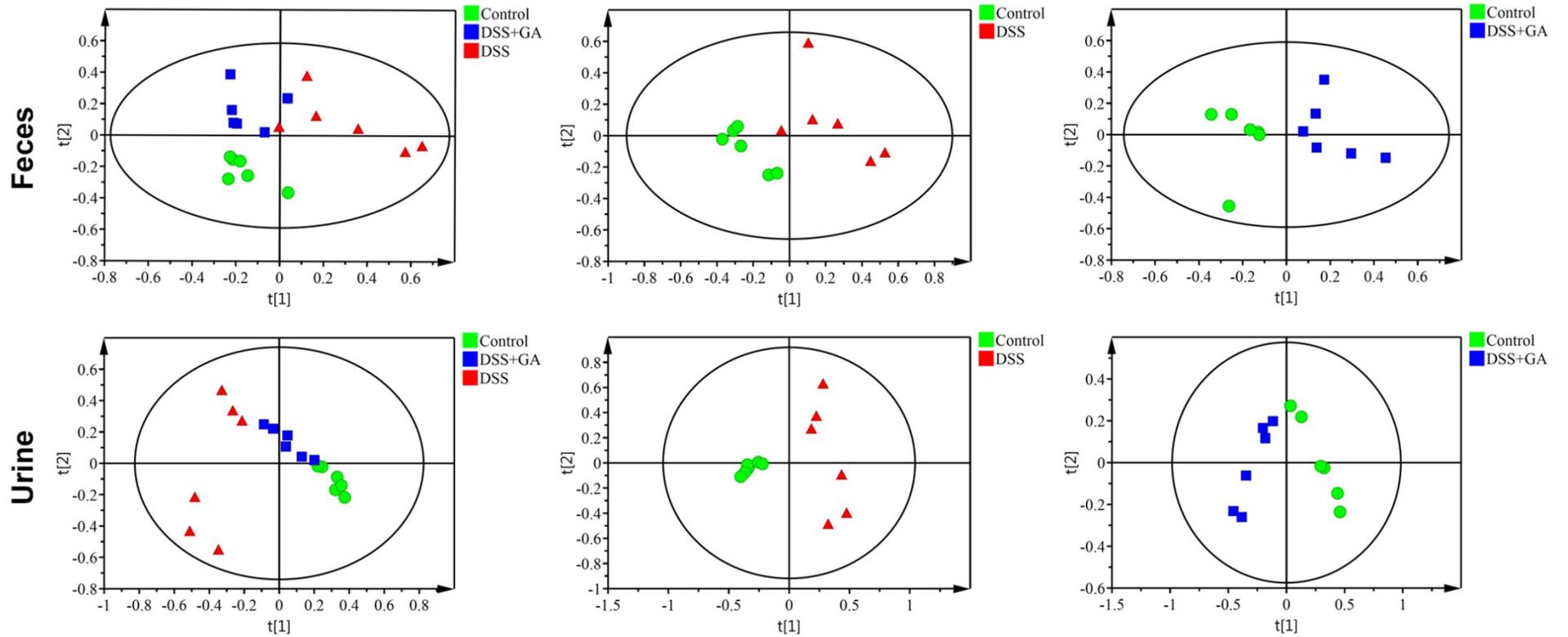


**Figure 2. PCA score plot based on PICRUSt analysis**



PICRUSt analysis shows the relative abundance of predicted microbial genes related to metabolism among groups based on Welch's t test.

**Figure 3. PCA scores plots of 1H NMR data of feces and urine**



**Table 1. Relative abundance of major bacterial taxa among different groups**

Phylum	Family	Genus	Relative abundance(%)
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			Control	DSS	DSS±GA
Actinobacteria	Corynebacteriaceae	Corynebacterium	0.19±0.17	0.00±0.00 <sup>###</sup>	0.04±0.05
	Micrococcaceae	Rothia	0.01±0.00	0.01±0.01	0.01±0.01
	Bifidobacteriaceae	Bifidobacterium	0.03±0.06	0.02±0.05	0.03±0.06
	Coriobacteriaceae	Adlercreutzia	0.01±0.00	0.00±0.00 <sup>###</sup>	0.01±0.01
Bacteroidetes	Bacteroidaceae	Bacteroides	0.77±0.36	5.54±4.57 <sup>###</sup>	19.70±9.77 <sup>*###</sup>
	Porphyromonadaceae	Parabacteroides	0.19±0.07	2.04±2.54 <sup>###</sup>	1.33±1.39 <sup>#</sup>
	Prevotellaceae	Prevotella	31.82±9.06	3.33±5.58 <sup>###</sup>	0.00±0.00 <sup>*###</sup>
	Rikenellaceae	AF12	0.05±0.04	0.00±0.00 <sup>###</sup>	0.00±0.00 <sup>###</sup>
	S24-7	g_	25.84±3.54	21.2±3.13	25.31±13.00
	[Odoribacteraceae]	Butyricimonas	0.03±0.03	0.04±0.04	0.00±0.00 <sup>*###</sup>
		Odoribacter	0.01±0.01	0.01±0.01	0.00±0.00
	[Paraprevotellaceae]	Paraprevotella	4.84±5.27	5.39±5.38	3.03±1.42
	[Paraprevotellaceae]	[Prevotella]	2.23±2.12	7.66±2.46 <sup>###</sup>	9.68±8.20
	Deferribacteres	Deferribacteraceae	Mucispirillum	0.01±0.01	0.01±0.01
Firmicutes	Lactobacillaceae	Lactobacillus	4.40±2.02	0.61±0.49 <sup>###</sup>	2.10±1.78
	Streptococcaceae	Lactococcus	0.00±0.00	0.02±0.02 <sup>###</sup>	0.00±0.00 <sup>**</sup>
	Turicibacteraceae	Turicibacter	0.06±0.10	0.69±0.63 <sup>###</sup>	0.13±0.13 <sup>*</sup>
	Clostridiaceae	Candidatus Arthromitus	0.00±0.00	0.00±0.00	0.01±0.01
		Clostridium	0.08±0.07	0.26±0.31	0.04±0.06
	Dehalobacteriaceae	Dehalobacterium	0.04±0.02	0.02±0.02	0.10±0.08
	Lachnospiraceae	Blautia	0.05±0.11	0.26±.32	0.10±0.14
		Coprococcus	0.20±0.32	0.35±0.16	0.57±0.44
		Dorea	0.04±0.02	0.20±0.19 <sup>###</sup>	0.10±0.13
		Lachnospira	0.01±0.01	0.00±0.00	0.04±0.07

		Roseburia	0.03±0.03	0.09±0.22	0.10±0.17
		[Ruminococcus]	0.19±0.15	0.35±0.47	0.13±0.08
	Peptococcaceae	rc4-4	0.02±0.02	0.13±0.03 <sup>###</sup>	0.37±0.25 <sup>*##</sup>
	Ruminococcaceae	Oscillospira	2.33±1.10	5.67±4.32	5.01±2.89
		Ruminococcus	4.59±1.34	2.75±0.85 <sup>###</sup>	3.50±1.63
	Veillonellaceae	Phascolarctobacterium	0.30±0.72	1.35±0.80 <sup>#</sup>	0.98±1.39
	Erysipelotrichaceae	Allobaculum	0.01±0.00	0.44±0.35 <sup>###</sup>	0.22±0.44 <sup>###</sup>
		p-75-a5	0.01±0.01	0.01±0.01	0.00±0.00
	Alcaligenaceae	Sutterella	0.42±0.17	2.70±1.49 <sup>###</sup>	0.78±0.59 <sup>*</sup>
Proteobacteria	Desulfovibrionaceae	Bilophila	0.02±0.02	0.00±0.00 <sup>#</sup>	0.00±0.00 <sup>###</sup>
		Desulfovibrio	0.09±0.12	0.10±0.07	0.03±0.05 <sup>*</sup>
	Helicobacteraceae	Flexispira	0.24±0.24	1.09±2.03	0.50±0.47
	Enterobacteriaceae	Proteus	0.00±0.00	0.01±0.01	0.01±0.01
Spirochaetes	Spirochaetaceae	Treponema	4.18±5.25	0.26±0.42 <sup>#</sup>	0.00±0.00 <sup>###</sup>
Tenericutes	Mycoplasmataceae	Mycoplasma	0.02±0.02	0.04±0.03	0.06±0.05
Verrucomicrobia	Verrucomicrobiaceae	Akkermansia	1.22±2.68	0.00±0.00 <sup>#</sup>	0.00±0.00 <sup>#</sup>

The data are expressed as the mean ± SD (n=6). p vales are calculated with Mann-Whitney U test. (vs. DSS :<sup>\*</sup> P<0.05,<sup>\*\*</sup>P<0.01;vs. control :<sup>#</sup> P<0.05,<sup>##</sup>P<0.01)

**Table 2. <sup>1</sup>H NMR data of metabolites in rat feces**

Number	Metabolites	$\delta$ <sup>1</sup> H (ppm)
1	2-Hydroxyisobutyrate	1.35(s)
2	3-Methyl-2-oxovalerate	0.899(t), 1.093(d), 1.438(m), 1.662(m), 1.685, 2.928(m), 1.924(s)
3	Alanine	1.484(d), 3.780(q)
4	Acetate	1.924(s)
5	Asparagine	2.862(dd), 2.962(m), 3.993(dd)
6	Aspartate	2.666(dd), 2.805(dd), 3.898(dd)
7	Choline	3.191(s), 3.504(m), 4.085(m)
8	Creatine	3.045(s), 3.927(s)
9	Dimethylamine	2.719(s)
10	Formate	8.460(s)
11	Fumarate	6.525(s)
12	Glycerol	3.551(dd), 3.634(m), 3.767(m) 3.768
13	Glycine	3.567(s)
14	Histidines	3.144(dd), 3.249(dd), 4.003 (dd) , 7.093(s), 7.844(s)
15	Isoleucine	0.936(t), 0.993 (d), 1.249(m), 1.452(m), 1.971(m), 3.655(d)
16	Lactate	1.335(d), 4.125(q)
17	Leucine	0.925(d), 0.935(s), 1.703(m), 3.724 (d)
18	Methylamine	2.613(s)
19	Pyruvate	2.374(s)
20	Succinate	2.410(s)
21	Threonine	1.335(d), 3.578(d), 4.269(m)
22	Xanthine	7.909(s)
23	$\alpha$ -Glucose	3.418 (m), 3.547 (m), 3.726(m), 3.742(m), 3.772(m), 3.870 (m), 5.210(d)
24	1,3-dihydroxyacetone	3.573(s), 4.42(s)

25	2-Ketoisovalerate	1.127(d), 3.027(m)
26	3-Methyl-2-ketovalerate	0.888(t;7.4), 1.102(d;7.0), 1.461(m), 1.727(m), 2.939(m)
27	3-hydroxyphenylacetate	3.481(s), 6.757(m), 6.807(d), 6.906(dd), 7.245 (t)
28	Adenine	8.196(s), 8.213(s)
29	Butyrate	0.899(t), 1.553(m), 2.150(t)
30	Cadaverine	1.484(d), 1.727(m), 3.022(t)
31	DCA	6.859(d)
32	desaminotyrosine	3.817(d)
33	Ethanolamine	3.134(d), 7.850(d;7.6)
34	Glutarate	1.775(overlap), 2.181(t)
35	Hypoxanthine	8.196(s), 8.216(s)
36	Imidazole	7.327(s), 8.297(s)
37	Isocaproate	0.882(d), 1.449(m), 1.484 (d), 2.189 (t)
38	Malonate	3.112(s)
39	Methanol	3.366(s)
40	Methionine	2.142 (s), 2.169(m), 2.648(t), 3.853(m)
41	Methylsuccinate	1.104(d), 2.135 (d), 2.635 (m)
42	N-acetyl-glycoprotein	2.03(s)
43	NAG	2.042(s), 2.065(s)
44	Phenylacetate	3.542(s), 7.317(m), 7.375(m)
45	Propionate	1.060(t), 2.187(q), 2.374(s)
46	Sarcosine	2.705(s), 3.598(s)
47	Taurochenodeoxycholate	0.957, 0.672(s), 0.965(s)
48	TCA	0.67(s)
49	Trimethylamine	2.871(s)
50	Tyrosine	3.068(d), 3.140(dd), 3.941(d), 6.907(m), 7.196 (d/m)
51	Uracil	5.807(d), 7.542(d)
52	Urocanate	6.40(d), 7.318(d), 7.433(s), 7.885(s)
53	Valerate	0.899(t), 1.309(m), 2.178(t)

54	Valine	0.972(d), 1.013(d), 2.275(m), 3.617(d), 7.909(s)
55	$\alpha$ -arabinose	3.849(dd), 3.906(m), 3.993(t), 4.023(d), 5.242 (d)
56	$\alpha$ -galactose	3.64(dd), 4.08(m), 4.58(d), 5.274 (d)
57	$\alpha$ -ketoisocaproate	0.920 (s), 2.054(m), 2.616(d)
58	$\alpha$ -ketoisovalerate	1.127(d), 3.022(m)
59	$\alpha$ -xylose	3.545(dd), 3.630(m), 3.670 (t), 5.203(d)
60	$\beta$ -arabinose	3.525(dd), 3.670 (t), 3.856(dd), 3.945(m), 4.523(d)
61	$\beta$ -L-Fucose	1.251(d : 6.5), 3.452(m), 3.645 (m), 3.804 (m) , 4.561 (d;7.9)
62	$\beta$ -xylose	3.23(dd), 3.46(t), 3.63(m), 3.932(dd), 4.570(d)

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**Abbreviation:** TCA, Taurocholic acid; DCA, Deoxycholic acid; NAG, N-Acetyl group contained glycoprotein

**Table 3. <sup>1</sup>H NMR data of metabolites in rat urine**

1H NMR Data of Metabolites in Rat Urine		
Number	Metabolites	δ <sup>1</sup> H (ppm)
1	1-Methylnicotinamide	4.476(s),8.181(t),8.894(d),8.960(d),9.272(s)
2	2-Hydroxy-3-methylvalerate	0.8(t),0.93(d),1.15(m),1.35(m),1.75(m),3.87(d)
3	2PY	6.67(d), 7.97(dd), 8.34(d), 3.65(s)
4	3-Hydeoxyphenylacetate	3.490(s),6.794(m),6.872(m or ov),7.258(tr or ov)
5	3-Hydroxyisovalerate	1.275(s), 2.357(s)
6	4-Hydroxyhippurate	3.978(s),6.976(d),7.765(d)
7	4-hydroxyphenylacetate	6.87(d)
8	4PY	6.70(d),7.83(dd) ,8.56(d) ,3.90(s)
9	4-Pyridoxate	2.429(s),7.834(s)
10	Acetamide	1.99(s)
11	Acetoacetate	2.266(s),3.434(s)
12	Acetone	2.229(s)
13	Acetylcarnitine	2.159(s), 2.504(m), 2.631(m), 3.202(s), 3.621(d), 3.829(dd),
14	Allantoin	5.38(s), 6.02(s), 7.27(s), 8.01(s)
15	Betaine	3.273(s),3.927(s)
16	Carnitine	2.41(t),3.230(s),3.405(m),4.554(m)
17	cis-aconitate	3.123(s/d),5.711(t)
18	Creatinine	3.043(s),4.052(s)
19	Dimethyl sulfone	3.145(s)
20	Galactarate	3.95(s), 4.26(s)
21	Galactonate	4.26(s)
22	Glycerate 3-phosphate	4.22(m), 4.06(d), 3.84(d)

23	Glycolate	3.932(s)
24	Guanidoacetate	3.801(s)
25	Guanine	7.69(s)
26	Hippurate	3.975(d), 7.553(t), 7.642(t), 7.837(d), 8.508(s)
27	Hypotaurine	3.36(t)
28	Indole-3-acetate	3.681 (s), 7.161 (t), 7.216 (t), 7.507 (d), 7.635(d)
29	Isobutyrate	1.053 (d), 2.382(m)
30	Isopropanol	1.146 (d), 4.028(m)
31	Isovalerate	0.895(d), 1.929(m), 2.049(d)
32	Isovalerylglycine	0.925(d), 1.994(m), 2.166(d),3.746(d)
33	Ketoleucine	0.944 (d), 2.100 (m), 2.617(d)
34	Lactose	3.276 (t), 3.507 (m), 3.604 (m), 3.710 (m), 3.768 (m), 3.847 (m), 3.936 (m), 4.455 (d), 4.660 (d), 5.211(d)
35	Methylguanidine	2.828 (s), 3.356(s)
36	N, N-dimethylglycine	2.925 (s), 3.712(s)
37	N6, N6, N6-trimethyllysine	3.109(s)
38	N-acetylglutamate	4.13(t), 1.83(m), 2.27(t), 2.04(s)
39	N-Acetylglycine	2.044 (s), 3.763 (d), 7.997(s)
40	N-methylhydantoin	2.91(s), 4.08(s)
41	N-Nitrosodimethylamine	3.162 (s), 3.802(s)
42	N-Phenylacetylglycine	3.67(s), 3.74(d), 7.98(s)
43	O-Acetylcholine	2.143 (s), 3.202 (s), 3.711 (t/m), 4.529(m)
44	Oxypurinol	8.209(s)
45	Pantothenate	0.94(s), 0.90(s), 3.52(s), 3.40(s), 4.00(s), 3.45(t), 2.43(t)
46	p-cresol	2.253 (s), 6.817 (d), 7.130(d)
47	p-Cresol glucuronide	2.299 (s), 7.055 (m), 7.237(m)
48	p-cresol sulfate	2.347(s), 7.217 (d), 7.293(s,d,d)
49	Phenylacetylglycine	3.680 (s), 3.746 (d), 7.365 (t), 7.425(t)

50	Phosphorylcholine	3.203 (s), 3.671 (t), 4.190 (dd)
51	Propylene glycol	1.14(d), 3.44(dd), 3.54(dd), 3.88(m)
52	Succinimide	2.78(s)
53	Thiamine	9.42(s), 8.04(s), 5.45(s), 3.89(t), 3.18(t), 2.56(s), 2.49(s)
54	Trans-Aconitate	3.435 (s), 6.571 (s)
55	Trigonelline	4.438 (s), 8.084 (t), 8.837 (t/d), 9.122 (s)
56	TMAO	3.266(s)
57	$\beta$ -Glucose	3.243 (t), 3.478 (m), 3.492 (m), 3.864 (m), 3.927 (s), 4.660(d)
58	4-hydroxyphenylacetate	3.45(s), 6.87(d), 7.16(d)
59	3-hydroxyisobutyrate	1.06(d), 2.48(m), 3.52(m), 3.71(m)
60	GPC	3.24(s), 3.69(m), 4.33(m)
61	NAG	2.04(s)
62	PC	3.22(s), 3.60(m), 4.17(m)
63	Dimethylglycine	2.92(s), 3.72(s)
64	2-Hydroxyisobutyrate	1.35(s)
65	3-Methyl-2-oxovalerate	0.899(t), 1.093(d), 1.438(m), 1.662(m), 2.928(m)
66	Alanine	1.489(d), 3.780(q)
67	Acetate	1.927(s)
68	Asparagine	2.862(dd), 2.962(m), 3.995(dd)
69	Choline	3.191(s), 3.504(t/m), 4.085(m)
70	Citrulline	1.558(m), 1.865(m), 3.124(dd), 3.743(t)
71	Creatine	3.028/3.047(s), 3.926(s)
72	DMA	2.719(s)
73	Formate	8.467(s)
74	Fumarate	6.524(s)
75	Glycine	3.566(s)
76	Isoleucine	0.936(t), 0.995(d), 1.249(m), 1.452(m), 1.971(m), 3.655(d)
77	Lactate	1.331(d), 4.115(q)

78	Leucine	0.925(d), 0.936(s/d), 1.703(m), 3.725(d)
79	Pyruvate	2.373 (s)
80	p-Hydroxyphenylacetate	3.445(s), 6.865 (d), 7.169(d)
81	Succinate	2.413(s)
82	Taurine	3.243(t), 3.435(t)
83	Threonine	1.330(d), 3.579(d), 4.269(m)
84	Xanthine	7.909(s)

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**Abbreviation:** 2PY, N<sup>1</sup>-methyl-2-pyridone-5-carboxamide; 4PY, N<sup>1</sup>-methyl-4-pyridone-3-carboxamide; TMAO, Trimethylamine-N-oxide; GPC, Glycerophosphocholine; NAG, N-Acetyl group contained glycoprotein; PC, Phosphorylcholine; DMA, Dimethylamine;

**Table 4. Screened metabolites with significant difference in fold change (FC) value in feces samples.**

Number	Metabolites	DSS / Control			□	DSS+GA / Control			□	DSS+GA / DSS		
		FC	p value	Trend		□	FC	p value		Trend	□	FC
1	Valerate	2.618	0.007	↑**						0.450	0.012	↓*
2	Butyrate	1.987	0.014	↑*						0.532	0.013	↓*
3	Glutarate	1.740	0.01	↑*						0.578	0.012	↓*
4	Taurochenodeoxycholate	1.655	0.009	↑**								
5	Glycine	1.563	0.005	↑**		1.544	0.002	↑**				
6	Methionine	1.517	0.006	↑**						0.712	0.013	↓*
7	3-Methyl-2-ketovalerate	1.515	0.015	↑*						0.640	0.013	↓*
8	Propionate	1.509	0.014	↑*						0.635	0.007	↓**
9	Methylsuccinate	1.503	0.014	↑*						0.699	0.024	↓*
10	3-Methyl-2-oxovalerate	1.465	0.029	↑*						0.627	0.013	↓*
11	Acetate	1.465	0.027	↑*						0.688	0.030	↓*
12	Dimethylamine	0.809	0.002	↓**								
13	Adenine	0.800	0.046	↓*								
14	Uracil	0.776	0.025	↓*								
15	Fumarate	0.739	<0.001	↓***		0.733	0.003	↓**				
16	Deoxycholic acid	0.637	<0.001	↓***		0.703	0.002	↓**				
17	Asparagine					1.264	<0.001	↑***				
18	Desaminotyrosine					1.261	0.005	↑**		1.212	0.032	↓*
19	Alpha-arabinose					1.208	0.005	↑**				
20	Xanthine					0.750	<0.001	↓***				
21	α-galactose									1.234	<0.001	↑***
22	β-xylose									1.280	0.003	↑**
23	β-arabinose									1.249	0.004	↑**
24	α-xylose									1.240	0.008	↑**

25

Isocaproate

□

□

□

□

□

□

□

□

0.782

0.018

↓\*

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\*p<0.05; \*\*p<0.01, \*\*\*p<0.001

**Table 5. Screened metabolites with significant difference in fold change (FC) value in urine samples**

Number	Metabolites	DSS / Control			□	DSS+GA / Control			□	DSS+GA / DSS		
		FC	p value	Trend		FC	p value	Trend		FC	p value	Trend
1	Oxypurinol	0.337	<0.001	↓***					2.771	<0.001	↑***	
2	Xanthine	0.588	<0.001	↓***					1.691	<0.001	↑***	
3	4-Hydroxyhippurate	0.714	<0.001	↓***		0.648	<0.001	↓***				
4	Hippurate	0.719	<0.001	↓***		0.650	<0.001	↓***				
5	cis-aconitate	0.792	<0.001	↓***								
6	Galactarate	0.815	<0.001	↓***								
7	Isovalerate	1.421	<0.001	↑***		1.280	<0.001	↑***				
8	Acetoacetate	0.723	0.045	↓*								
9	Succinate	1.605	0.021	↑*								
10	Fumarate	0.806	0.020	↓*								
11	Isopropanol	0.707	0.016	↓*		0.742	0.033	↓**				
12	Acetate	2.154	0.011	↑*		1.316	0.002	↑**				
13	Taurine	0.646	0.008	↓**								
14	trans-Aconitate	0.506	0.008	↓**								
15	Trigonelline	0.830	0.006	↓**					1.219	0.002	↑**	
16	4-Pyridoxate	0.741	0.005	↓**		0.628	<0.001	↓***				
17	2-Hydroxyisobutyrate	1.542	0.003	↑**		1.338	<0.001	↑***				
18	Formate	1.659	0.001	↑**					0.661	0.010	↓**	
19	Indole-3-acetate					0.674	<0.001	↓***				
20	succinimide					1.239	<0.001	↑***				
21	p-Cresol glucuronide					0.753	<0.001	↓***				
22	Glycolate					0.783	<0.001	↓***				
23	Asparagine					0.800	<0.001	↓***				
24	Phenylacetyl glycine	□	□	□	□	0.797	<0.001	↓***	□	□	□	

\*p<0.05; \*\*p<0.01, \*\*\*p<0.001