



Figure S1 The results of western blot. The result showed that FMO3 expression in the liver tissue of mice treated with ranitidine and finasteride was significantly lower than in the saline-control and Andrioe-treated groups. ** $p < 0.01$ vs Saline group; t test.

Table S1 The results of sample sequencing data processing statistics

Sample ID	PE Reads	Raw Tags	Clean Tags	Effective Tags	AvgLen(bp)	GC(%)	Q20(%)	Q30(%)	Effective(%)
A1	79882	65646	57536	55953	415	54.34	96.61	93.42	70.04
A2	79873	64205	55290	54009	417	55.15	96.67	93.44	67.62
A3	80158	67473	59480	58151	415	54.27	96.68	93.53	72.55
A4	79944	67108	59113	57672	415	53.86	96.48	93.22	72.14
A5	79778	65248	56964	55605	417	54.44	96.58	93.31	69.7
A6	80127	65231	56760	55491	416	54.84	96.64	93.41	69.25
A7	79802	65495	57147	55230	415	54.83	96.59	93.33	69.21
A8	79817	65274	56990	55560	415	54.52	96.63	93.41	69.61
FEN1	80088	67152	59822	56824	415	54.46	96	92.14	70.95
FEN2	79947	67468	59875	57420	418	54.32	96.07	92.29	71.82
FEN3	80259	64784	56786	55039	415	54.7	96.06	92.29	68.58
FEN4	80168	67506	60133	57973	415	54.11	96.11	92.39	72.31
FEN5	7978	6475	5670	54806	415	53.87	96.02	92.21	68.69

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FEN6	7975 7	6667 1	5944 6	57728	415	54.34	96.17	92.43	72.38
FEN7	8003 6	6764 2	6024 7	56580	416	54.26	96.06	92.31	70.69
FEN8	8036 3	6647 6	5862 3	56923	416	54.37	96.02	92.17	70.83
R1	7988 0	6672 0	5907 4	58359	413	54.24	96.59	93.43	73.06
R2	8020 9	6680 1	5892 9	58311	414	54.29	96.65	93.47	72.7
R3	8012 5	6914 6	6184 8	60935	412	53.73	96.59	93.45	76.05
R4	7997 4	6725 6	5966 2	58542	412	53.82	96.45	93.16	73.2
R5	7968 1	6980 2	6287 0	61949	411	53.44	96.58	93.42	77.75
R6	8030 2	6903 0	6163 8	60710	412	54.25	96.56	93.37	75.6
R7	7978 6	6518 6	5675 8	55797	416	54.41	96.52	93.22	69.93
R8	7978 1	6752 4	6000 9	58603	413	54.27	96.57	93.39	73.45
W1	7997 3	6824 1	6103 7	58773	418	51.61	96.6	93.43	73.49
W2	8031 7	7046 6	6338 1	62267	418	52.17	96.62	93.44	77.53
W3	7982 4	7004 5	6317 3	62099	416	52.08	96.63	93.46	77.79

W4	7977 9	6885 9	6154 7	60488	418	51.68	96.45	93.19	75.82
W5	7992 1	7044 5	6356 3	62460	418	51.59	96.67	93.56	78.15
W6	7976 0	6869 1	6172 5	60874	420	51.06	96.7	93.63	76.32
W7	7972 1	6941 1	6243 5	61246	418	51.79	96.62	93.45	76.83
W8	8036 0	6978 7	6251 1	61299	419	51.51	96.58	93.39	76.28

Table S2 Alpha diversity index statistics

Sample ID	OTU	ACE	Chao1	Simpson	Shannon	Coverage
A1	349	356.2348	357.5714	0.0246	4.4439	0.9996
A2	346	352.2343	356.2	0.0271	4.387	0.9996
A3	359	363.8258	365.5	0.0288	4.471	0.9997
A4	355	361.7728	364.2308	0.0273	4.473	0.9997
R5	354	361.0806	362.5714	0.0339	4.4825	0.9997
R6	338	350.8598	349	0.0419	4.1996	0.9996
R7	349	355.8748	359	0.0195	4.5906	0.9997
R8	349	352.3699	352	0.0508	4.1888	0.9998
W1	54	192.1155	124	0.2482	1.8598	0.9996
W2	40	204.223	79	0.1964	2.0594	0.9998
W3	38	119.3596	71	0.2036	1.9729	0.9998
W4	52	123.5124	103	0.2025	2.1266	0.9997
W5	40	107.9454	58.3333	0.2038	2.0666	0.9998
W6	40	72.5537	49.3333	0.2591	1.8963	0.9999
W7	35	60.8773	45.5	0.2107	1.9424	0.9999
W8	43	111.0345	65.5	0.2583	1.8203	0.9998
A5	359	363.9935	363.1053	0.0228	4.4921	0.9997
A6	363	368.4088	367.5882	0.0208	4.5909	0.9997
A7	353	358.7041	363.9091	0.0225	4.467	0.9996
A8	364	368.4355	370.5	0.024	4.5061	0.9997
FEN1	317	331.6806	332.4762	0.0483	4.0035	0.9992
FEN2	309	327.5397	335.64	0.153	3.0112	0.9991
FEN3	311	320.7279	320.5	0.0384	4.2046	0.9994
FEN4	316	328.1112	328.0476	0.0425	4.0977	0.9994
FEN5	322	329.4535	330.0526	0.0347	4.1957	0.9995
FEN6	322	332.3943	333	0.0571	3.9718	0.9993
FEN7	312	327.3241	329.6522	0.0589	3.7382	0.9992

FEN8	321	330.7174	328.9167	0.0592	3.993	0.9995
R1	351	357.5436	358.5	0.0194	4.6551	0.9997
R2	355	367.3563	367.3529	0.0205	4.612	0.9996
R3	349	356.7963	354.7143	0.0344	4.3975	0.9997
R4	350	354.1829	353.2353	0.0278	4.5348	0.9998

Table S3 Species statistics of different grades of samples

Sample	Kindom	Phylum	Class	Order	Family	Genus	Species
A1	1	10	16	18	32	85	87
A2	1	10	16	17	32	85	85
A3	1	10	16	17	31	83	86
A4	1	10	16	18	33	85	87
A5	1	10	16	17	32	85	87
A6	1	10	16	17	30	84	86
A7	1	10	16	17	30	82	83
A8	1	10	16	17	31	85	87
FEN1	1	10	16	16	29	78	78
FEN2	1	10	16	16	30	77	78
FEN3	1	10	15	15	28	76	77
FEN4	1	10	15	15	26	75	76
FEN5	1	10	16	16	30	80	81
FEN6	1	10	16	16	29	80	81
FEN7	1	10	15	15	29	77	78
FEN8	1	10	16	17	31	80	81
R1	1	10	16	17	31	83	84
R2	1	10	16	17	31	82	83
R3	1	10	16	17	32	82	83
R4	1	10	16	17	30	81	82
R5	1	10	16	17	32	82	83
R6	1	10	16	16	31	80	81
R7	1	10	15	16	31	82	83
R8	1	10	16	17	30	82	84
W1	1	7	11	12	20	39	39
W2	1	6	8	9	16	29	29
W3	1	6	9	10	17	27	27

W4	1	6	8	9	17	39	39
W5	1	7	9	10	16	29	29
W6	1	6	8	9	15	28	28
W7	1	6	8	9	16	28	28
W8	1	8	11	12	19	36	37
Total	1	10	16	19	34	91	94

Table S4 Analysis on the difference of metabolic pathway of KEGG (FEN vs W)

Class1	Class2	FEN	W	FEN: rel. freq. (%)	W: rel. freq. (%)	p-val ues (correc ted)	Effect size	95.0% lower CI	95.0% upper CI
Metabolism	Carbohydrate metabolism	3101 849	5511 771	14.975 20522	15.926 48857	0	-0.9512 83352	-0.9709 06411	-0.9316 60293
Metabolism	Lipid metabolism	7656 78	1434 282	3.6965 64592	4.1444 16719	0	-0.4478 52127	-0.4583 53432	-0.4373 50821
Metabolism	Metabolism of cofactors and vitamins	1343 061	2226 544	6.4840 72597	6.4336 90292	1.90E- 13	0.05038 2305	0.03698 5362	0.06377 9248
Metabolism	Energy metabolism	1429 491	2447 315	6.9013 42099	7.0716 17159	0	-0.1702 75061	-0.1841 42747	-0.1564 07375
Metabolism	Nucleotide metabolism	1117 547	1731 120	5.3953 28938	5.0021 42306	0	0.39318 6632	0.38103 7844	0.40533 5419
Metabolism	Biosynthesis of other secondary metabolites	2809 70	5327 46	1.3564 75899	1.5393 91437	0	-0.1829 15538	-0.1893 76017	-0.1764 55059
Metabolism	Amino acid	2281	3942	11.017	11.392	0	-0.3749	-0.3921	-0.3578

ism	metabolism	995	521	08801	07628		88263	38135	38392
Metabolism	Metabolism of terpenoids and polyketides	388096	696824	1.873662208	2.013501554	0	-0.139839346	-0.147330068	-0.132348624
Metabolism	Xenobiotics biodegradation and metabolism	418000	716849	2.01803369	2.071364614	0	-0.053330924	-0.061031796	-0.045630051
Metabolism	Metabolism of other amino acids	423180	755284	2.043041858	2.182424124	0	-0.139382265	-0.147188053	-0.131576478
Metabolism	Glycan biosynthesis and metabolism	605924	1220926	2.925299152	3.527915798	0	-0.602616647	-0.612134406	-0.593098888
Genetic Information Processing	Translation	1060066	1538206	5.117820338	4.444709383	0	0.673110955	0.66139012	0.68483179
Metabolism	Global and	2853	4643	13.777	13.416	0	0.36081	0.34211	0.37950

ism	overview maps	768	194	51188	69967		2207	6202	8213
Human Diseases	Drug resistance	1676 13	3084 26	0.8092 0737	0.8912 09588	0	-0.0820 02218	-0.0869 78721	-0.0770 25715
Environmental Information Processing	Membrane transport	9533 44	1468 634	4.6025 84474	4.2436 78233	0	0.35890 624	0.34764 9834	0.37016 2647
Environmental Information Processing	Signal transduction	7294 07	1117 801	3.5214 54305	3.2299 31877	0	0.29152 2428	0.28163 0372	0.30141 4483
Cellular Processes	Cell motility	3962 39	2579 84	1.9129 75242	0.7454 55359	0	1.16751 9883	1.16095 3959	1.17408 5808
Genetic Information Processing	Folding, sorting and degradation	4464 97	7575 27	2.1556 12412	2.1889 05364	2.81E- 16	-0.0332 92952	-0.0412 3028	-0.0253 55624
Genetic Information Processing	Transcription	4622 5	6943 9	0.2231 66525	0.2006 46841	0	0.02251 9684	0.01999 1741	0.02504 7626

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Genetic Information Processing	Replication and repair	902255	1423866	4.355935375	4.114319259	0	0.241616116	0.230606005	0.252626227
Organismal Systems	Endocrine system	99947	199848	0.482527304	0.577469	0	-0.094941696	-0.09885805	-0.091025341
Environmental Information Processing	Signaling molecules and interaction	0	0	0	0	1.075	0	-7.56E-06	7.56E-06
Cellular Processes	Cell growth and death	149540	236691	0.721953966	0.683928361	0	0.038025605	0.033453795	0.042597416
Cellular Processes	Transport and catabolism	122564	266873	0.591718376	0.771140489	0	-0.179422113	-0.183834528	-0.175009699
Organismal Systems	Circulatory system	3708	225	0.0179016	0.000650147	0	0.017251454	0.016661517	0.01784139
Organismal System	Development	0	0	0	0	1.048780488	0	-7.56E-06	7.56E-06

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Cellular Processes	Cellular community	0	0	0	0	1.023809524	0	-7.56E-06	7.56E-06
Organismal Systems	Immune system	25437	46780	0.122805557	0.13517273	0	-0.012367173	-0.014317193	-0.010417153
Organismal Systems	Environmental adaptation	43076	63132	0.207963682	0.182422506	0	0.025541177	0.023110793	0.027971561
Organismal Systems	Nervous system	68232	135929	0.329412619	0.392772426	0	-0.063359807	-0.0665972	-0.060122413
Organismal Systems	Sensory system	0	0	0	0	1	0	-7.56E-06	7.56E-06
Human Diseases	Endocrine and metabolic diseases	31188	50408	0.150570418	0.145655985	4.70E-06	0.004914433	0.00280861	0.007020256
Organismal Systems	Excretory system	12094	32024	0.058387798	0.092534663	0	-0.034146865	-0.035606465	-0.032687265
Organis	Digestive	4973	1734	0.0240	0.0501	0	-0.0261	-0.0271	-0.0250

mal System s	system		2	08808	10421		01613	0973	93497
Human Disease s	Neurodege nerative diseases	5869 6	6645 7	0.2833 74415	0.1920 3023	0	0.09134 4185	0.08862 2217	0.09406 6154
Human Disease s	Substance dependenc e	9900	2375 1	0.0477 95535	0.0686 2949	0	-0.0208 33955	-0.0221 24973	-0.0195 42937
Human Disease s	Infectious diseases: Bacterial	1667 45	2722 20	0.8050 16812	0.7865 90865	9.96E- 14	0.01842 5947	0.01357 3582	0.02327 8312
Human Disease s	Infectious diseases: Parasitic	1144 5	2672 1	0.0552 54535	0.0772 11426	0	-0.0219 56892	-0.0233 35793	-0.0205 7799
Human Disease s	Infectious diseases: Viral	20	135	9.66E- 05	0.0003 90088	9.43E- 12	-0.0002 93531	-0.0003 79329	-0.0002 07734
Human Disease s	Cancers: Overview	1644 39	3067 42	0.7938 83832	0.8863 43601	0	-0.0924 59769	-0.0974 02682	-0.0875 16856
Human Disease s	Cancers: Specific types	1851 3	3779 8	0.0893 7765	0.1092 18873	0	-0.0198 41223	-0.0215 42043	-0.0181 40403
Human Disease s	Immune diseases	1150 4	2319 2	0.0555 39377	0.0670 14236	0	-0.0114 74859	-0.0128 13895	-0.0101 35823
Human Disease	Cardiovas cular	6	45	2.90E- 05	0.0001 30029	7.13E- 05	-0.0001 01062	-0.0001 53129	-4.90E- 05

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Table S5 Analysis on the difference of metabolic pathway of KEGG (R vs W)

Class1	Class2	R	W	R: rel. freq. (%)	W: rel. freq. (%)	p-values (corrected)	Effect size	95.0% lower CI	95.0% upper CI
Metabolism	Carbohydrate metabolism	4869 911	5511 771	15.717 15.663	15.926 48857	0	-0.2093 31947	-0.2270 25874	-0.1916 3802
Metabolism	Lipid metabolism	1070 975	1434 282	3.4564 6.6005	4.1444 16719	0	-0.6879 50714	-0.6972 01626	-0.6786 99802
Metabolism	Metabolism of cofactors and vitamins	1933 746	2226 544	6.2409 7.4169	6.4336 90292	0	-0.1927 16123	-0.2045 27452	-0.1809 04794
Metabolism	Energy metabolism	2012 550	2447 315	6.4953 06294	7.0716 17159	0	-0.5763 10866	-0.5884 92336	-0.5641 29395
Metabolism	Nucleotide metabolism	1742 909	1731 120	5.6250 6.6605	5.0021 42306	0	0.62292 43	0.61202 9644	0.63381 8956
Metabolism	Biosynthesis of other secondary metabolites	4273 51	5327 46	1.3792 3.3132	1.5393 91437	0	-0.1601 58305	-0.1659 68443	-0.1543 48168
Metabolism	Amino acid	3330	3942	10.750	11.392	0	-0.6419	-0.6571	-0.6267

ism	metabolism	889	521	11517	07628		61108	65739	56476
Metabolism	Metabolism of terpenoids and polyketides	5237 53	6968 24	1.6903 61063	2.0135 01554	0	-0.3231 40491	-0.3296 65894	-0.3166 15088
Metabolism	Xenobiotics biodegradation and metabolism	6016 89	7168 49	1.9418 91803	2.0713 64614	0	-0.1294 7281	-0.1362 70264	-0.1226 75356
Metabolism	Metabolism of other amino acids	6428 97	7552 84	2.0748 86552	2.1824 24124	0	-0.1075 37572	-0.1145 35481	-0.1005 39662
Metabolism	Glycan biosynthesis and metabolism	8434 20	1220 926	2.7220 54724	3.5279 15798	0	-0.8058 61075	-0.8142 69909	-0.7974 5224
Genetic Information Processing	Translation	1610 639	1538 206	5.1981 78248	4.4447 09383	0	0.75346 8865	0.74305 9014	0.76387 8716
Metabolism	Global and	4303	4643	13.888	13.416	0	0.47208	0.45542	0.48874

ism	overview maps	396	194	78543	69967		5758	9782	1734
Human Diseases	Drug resistance	2183 33	3084 26	0.7046 48187	0.8912 09588	0	-0.1865 61401	-0.1908 66108	-0.1822 56694
Environmental Information Processing	Membrane transport	1509 030	1468 634	4.8702 45239	4.2436 78233	0	0.62656 7006	0.61643 4496	0.63669 9516
Environmental Information Processing	Signal transduction	1066 990	1117 801	3.4436 04811	3.2299 31877	0	0.21367 2934	0.20495 3858	0.22239 2009
Cellular Processes	Cell motility	8081 52	2579 84	2.6082 30738	0.7454 55359	0	1.86277 5379	1.85646 8103	1.86908 2656
Genetic Information Processing	Folding, sorting and degradation	6149 23	7575 27	1.9846 03231	2.1889 05364	0	-0.2043 02134	-0.2112 27795	-0.1973 76473
Genetic Information Processing	Transcription	6808 3	6943 9	0.2197 31156	0.2006 46841	0	0.01908 4315	0.01685 5496	0.02131 3134

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Genetic Information Processing	Replication and repair	1374190	1423866	4.435062461	4.114319259	0	0.320743201	0.310922059	0.330564344
Organismal Systems	Endocrine system	143039	199848	0.461644241	0.577469	0	-0.115824759	-0.119304934	-0.112344584
Environmental Information Processing	Signaling molecules and interaction	0	0	0	0	1.075	0	-5.99E-06	5.99E-06
Cellular Processes	Cell growth and death	219593	236691	0.708714713	0.683928361	0	0.024786352	0.020747487	0.028825216
Cellular Processes	Transport and catabolism	174803	266873	0.564159413	0.771140489	0	-0.206981076	-0.210917547	-0.203044606
Organismal Systems	Circulatory system	5082	225	0.016401653	0.000650147	0	0.015751506	0.015286676	0.016216336
Organismal System	Development	0	0	0	0	1.048780488	0	-5.99E-06	5.99E-06

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Cellular Processes	Cellular community	0	0	0	0	1.023809524	0	-5.99E-06	5.99E-06
Organismal Systems	Immune system	37758	46780	0.121860215	0.13517273	0	-0.013312516	-0.015052687	-0.011572344
Organismal Systems	Environmental adaptation	69906	63132	0.225614709	0.182422506	0	0.043192203	0.04099257	0.045391836
Organismal Systems	Nervous system	92265	135929	0.297776172	0.392772426	0	-0.094996254	-0.097834824	-0.092157683
Organismal Systems	Sensory system	0	0	0	0	1	0	-5.99E-06	5.99E-06
Human Diseases	Endocrine and metabolic diseases	45984	50408	0.148408817	0.145655985	0.004266266	0.002752832	0.000888971	0.004616692
Organismal Systems	Excretory system	7686	32024	0.024805806	0.092534663	0	-0.067728857	-0.068889691	-0.066568023
Organis	Digestive	7238	1734	0.0233	0.0501	0	-0.0267	-0.0276	-0.0258

mal System s	system		2	5993	10421		50491	75994	24988
Human Disease s	Neurodege nerative diseases	8377 2	6645 7	0.2703 65854	0.1920 3023	0	0.07833 5624	0.07599 0748	0.08068 05
Human Disease s	Substance dependenc e	3267	2375 1	0.0105 4392	0.0686 2949	0	-0.0580 8557	-0.0590 3601	-0.0571 3513
Human Disease s	Infectious diseases: Bacterial	2414 79	2722 20	0.7793 49615	0.7865 90865	0.0010 68603	-0.0072 4125	-0.0115 19192	-0.0029 63307
Human Disease s	Infectious diseases: Parasitic	8287	2672 1	0.0267 45474	0.0772 11426	0	-0.0504 65953	-0.0515 6185	-0.0493 70055
Human Disease s	Infectious diseases: Viral	87	135	0.0002 80784	0.0003 90088	0.0178 15056	-0.0001 09304	-0.0002 03679	-1.49E- 05
Human Disease s	Cancers: Overview	2302 24	3067 42	0.7430 25215	0.8863 43601	0	-0.1433 18386	-0.1476 71194	-0.1389 65579
Human Disease s	Cancers: Specific types	2305 0	3779 8	0.0743 91598	0.1092 18873	0	-0.0348 27275	-0.0362 93621	-0.0333 60929
Human Disease s	Immune diseases	1731 1	2319 2	0.0558 69542	0.0670 14236	0	-0.0111 44694	-0.0123 48871	-0.0099 40516
Human Disease	Cardiovas cular	25	45	8.07E- 05	0.0001 30029	0.0566 11348	-4.93E- 05	-0.0001 04772	6.08E-0 6

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Table S6 Analysis on the difference of metabolic pathway of COG (FEN vs W)

Class1	Class2	FEN	W	FEN: rel. freq. (%)	W: rel. freq. (%)	p-val ues (corre cted)	Effect size	95.0% lower CI	95.0% upper CI
INFORMA TION STORAG E AND PROCES SING	RNA processing and modification	795 0	177 02	0.0242 10278	0.0317 69065	0	-0.007 558787	-0.008 272124	-0.006 845451
INFORMA TION STORAG E AND PROCES SING	Chromatin structure and dynamics	794 3	174 45	0.0241 88961	0.0313 07838	0	-0.0071 18877	-0.007 829791	-0.006 407963
METABO LISM	Energy production and conversion	163 1110	268 534 7	4.9672 48671	4.8192 83963	0	0.1479 64708	0.1386 40832	0.1572 88584
CELLULA R PROCES SES AND SIGNALI NG	Cell cycle control, cell division, chromosome partitioning	385 875	544 647	1.1751 1209	0.9774 56006	0	0.1976 56084	0.1931 5042	0.2021 61748
METABO LISM	Amino acid transport and	243 776	398 004	7.4237 75766	7.1428 1907	0	0.2809 56696	0.2697 21392	0.2921 91999

	metabolism	7	1						
METABOLISM	Nucleotide transport and metabolism	834 995	125 441 3	2.5428 25318	2.2512 44422	0	0.2915 80896	0.2849 30729	0.2982 31063
METABOLISM	Carbohydrate transport and metabolism	281 815 2	450 218 4	8.5821 6906	8.0798 88055	0	0.5022 81005	0.4903 18641	0.5142 43369
METABOLISM	Coenzyme transport and metabolism	130 230 4	230 056 5	3.9659 29835	4.1287 31225	0	-0.162 801391	-0.171 282219	-0.154 320563
METABOLISM	Lipid transport and metabolism	800 689	145 445 9	2.4383 52638	2.6102 5891	0	-0.171 906272	-0.178 645637	-0.165 166908
INFORMATION STORAGE AND PROCESSING	Translation, ribosomal structure and biogenesis	194 265 6	291 143 3	5.9160 05317	5.2250 31389	0	0.6909 73928	0.6810 06575	0.7009 41282
INFORMATION STORAGE AND PROCESSING	Transcription	261 840 2	423 114 2	7.9738 66786	7.5934 59908	0	0.3804 06878	0.3688 16852	0.3919 96904
INFORMATION STORAGE AND	Replication, recombination and repair	201 646 3	335 857 5	6.1407 71104	6.0274 99104	0	0.1132 72001	0.1029 48548	0.1235 95454

PROCES SING									
CELLULAR PROCESSES AND SIGNALING	Cell wall/membrane/envelope biogenesis	238 202 3	453 292 4	7.2540 17559	8.1350 55893	0	-0.881 038335	-0.892 454763	-0.869 621906
CELLULAR PROCESSES AND SIGNALING	Cell motility	615 110	772 156	1.8732 05565	1.3857 57233	0	0.4874 48333	0.4818 82626	0.4930 14039
CELLULAR PROCESSES AND SIGNALING	Posttranslational modification, protein turnover, chaperones	105 847 8	181 908 6	3.2234 02124	3.2646 40282	0	-0.041 238158	-0.048 876077	-0.033 600239
METABOLISM	Inorganic ion transport and metabolism	155 587 0	317 871 6	4.7381 18943	5.7047 1341	0	-0.966 594467	-0.976 080125	-0.957 108808
METABOLISM	Secondary metabolites biosynthesis, transport and catabolism	383 771	744 720	1.1687 04741	1.3365 1895	0	-0.167 814209	-0.172 573237	-0.163 055181
POORLY	General	383	675	11.684	12.115	0	-0.431	-0.445	-0.417

CHARACTERIZED	function prediction only	6868	1015	48325	77436		291108	228777	353439
POORLY CHARACTERIZED	Function unknown	2457564	4518638	7.484063927	8.109417385	0	-0.625353458	-0.636863551	-0.613843365
CELLULAR PROCESSES AND SIGNALING	Signal transduction mechanisms	2038226	3242537	6.20704636	5.819250385	0	0.387795976	0.377500958	0.398090993
CELLULAR PROCESSES AND SIGNALING	Intracellular trafficking, secretion, and vesicular transport	791901	1487278	2.411590383	2.669157846	0	-0.257567463	-0.264313263	-0.250821662
CELLULAR PROCESSES AND SIGNALING	Defense mechanisms	907769	1404743	2.764445291	2.52103561	0	0.243409681	0.236448798	0.250370564
CELLULAR PROCESSES AND SIGNALING	Extracellular structures	16	735	4.87E-05	0.001319075	1.13E-122	-0.00127035	-0.001373397	-0.001167302

NG									
CELLULAR PROCESSES AND SIGNALING	Nuclear structure	0	0	0	0	1	0	-4.74E-06	4.74E-06
CELLULAR PROCESSES AND SIGNALING	Cytoskeleton	5391	10370	0.016417309	0.018610621	5.87E-14	-0.002193311	-0.00276401	-0.001622613

Table S7 Analysis on the difference of metabolic pathway of COG (R vs W)

Class1	Class2	R	W1	R: rel. freq. (%)	W: rel. freq. (%)	p-values (corrected)	Effect size	95.0% lower CI	95.0% upper CI
INFORMATION STORAGE AND PROCESSING	RNA processing and modification	6762	17702	0.013342264	0.031769065	0	-0.018426801	-0.018996238	-0.017857365
INFORMATION STORAGE AND PROCESSING	Chromatin structure and dynamics	4840	17445	0.00954992	0.031307838	0	-0.021757918	-0.022298407	-0.021217429
METABOLISM	Energy production and conversion	2455957	2685347	4.845907602	4.819283963	1.74E-10	0.026623639	0.018460659	0.034786619
CELLULAR PROCESSES AND SIGNALING	Cell cycle control, cell division, chromosome partitioning	661520	544647	1.305260962	0.977456006	0	0.327804956	0.32374699	0.331862923
METABOLISM	Amino acid transport and	374466	398004	7.388684288	7.14281907	0	0.245865218	0.235982664	0.255747771

	metabolism	3	1						
METABOLISM	Nucleotide transport and metabolism	137 962 9	125 441 3	2.7221 79036	2.2512 44422	0	0.4709 34615	0.4649 94384	0.4768 74845
METABOLISM	Carbohydrate transport and metabolism	538 966 0	450 218 4	10.634 46728	8.0798 88055	0	2.5545 79228	2.5434 74335	2.5656 84121
METABOLISM	Coenzyme transport and metabolism	187 910 8	230 056 5	3.7077 13019	4.1287 31225	0	-0.421 018207	-0.428 394145	-0.413 642269
METABOLISM	Lipid transport and metabolism	109 215 5	145 445 9	2.1549 57199	2.6102 5891	0	-0.455 301711	-0.461 093976	-0.449 509446
INFORMATION STORAGE AND PROCESSING	Translation, ribosomal structure and biogenesis	298 849 4	291 143 3	5.8966 69116	5.2250 31389	0	0.6716 37727	0.6629 04824	0.6803 7063
INFORMATION STORAGE AND PROCESSING	Transcription	458 051 7	423 114 2	9.0379 27843	7.5934 59908	0	1.4444 67934	1.4339 43399	1.4549 92469
INFORMATION STORAGE AND	Replication, recombination and repair	306 546 1	335 857 5	6.0485 34548	6.0274 99104	5.59E -06	0.0210 35444	0.0119 69606	0.0301 01282

PROCES SING									
CELLULAR PROCESSES AND SIGNALING	Cell wall/membrane/envelope biogenesis	326 981 6	453 292 4	6.4517 52295	8.1350 55893	0	-1.683 303598	-1.693 169778	-1.673 437419
CELLULAR PROCESSES AND SIGNALING	Cell motility	953 023	772 156	1.8804 32516	1.3857 57233	0	0.4946 75283	0.4898 33593	0.4995 16973
CELLULAR PROCESSES AND SIGNALING	Posttranslational modification, protein turnover, chaperones	147 229 2	181 908 6	2.9050 14622	3.2646 40282	0	-0.359 62566	-0.366 198328	-0.353 052993
METABOLISM	Inorganic ion transport and metabolism	218 121 2	317 871 6	4.3038 01659	5.7047 1341	0	-1.400 91175	-1.409 179992	-1.392 643508
METABOLISM	Secondary metabolites biosynthesis, transport and catabolism	482 003	744 720	0.9510 51668	1.3365 1895	0	-0.385 467282	-0.389 499754	-0.381 43481
POORLY	General	566	675	11.168	12.115	0	-0.947	-0.959	-0.935

CHARACTERIZED	function prediction only	0255	1015	38476	77436		389596	583689	195503
POORLY CHARACTERIZED	Function unknown	3409487	4518638	6.727340492	8.109417385	0	-1.382076894	-1.392027152	-1.372126635
CELLULAR PROCESSES AND SIGNALING	Signal transduction mechanisms	3396240	3242537	6.701202518	5.819250385	0	0.881952134	0.872719511	0.891184757
CELLULAR PROCESSES AND SIGNALING	Intracellular trafficking, secretion, and vesicular transport	978971	1487278	1.931631136	2.669157846	0	-0.737526709	-0.74321095	-0.731842469
CELLULAR PROCESSES AND SIGNALING	Defense mechanisms	1621842	1404743	3.20009531	2.52103561	0	0.6790597	0.672698208	0.685421192
CELLULAR PROCESSES AND SIGNALING	Extracellular structures	7	735	1.38E-05	0.001319075	5.98E-193	-0.001305263	-0.001404864	-0.001205662

NG									
CELLULAR PROCESSES AND SIGNALING	Nuclear structure	0	0	0	0	1	0	-3.69E-06	3.69E-06
CELLULAR PROCESSES AND SIGNALING	Cytoskeleton	7139	10370	0.014086132	0.018610621	0	-0.004524489	-0.005012983	-0.004035994