

SUPPLEMENTARY TABLES

Supplementary Table 2. Amount of the three polymethoxyflavones—nobiletin, sinensetin, and tangeretin—in 1-gram water extracts of kososan (KS) and nobiletin-rich kososan (NKS).

	Nobiletin	Sinensetin	Tangeretin
KS	0.09 mg	N.D.	0.02 mg
NKS	2.44 mg	0.142 mg	0.516 mg

For this analysis, water extracts of KS and NKS was analyzed by high performance liquid chromatography under the following conditions: For determination of nobiletin, sinensetin, and tangeretin contents, an ODS column (Mightysil RP-18 GP S, 4.6 × 150 mm, 5 µm; Kanto Chemical Co., INC., Japan), equipped with a detector (an ultraviolet absorption photometer, wavelength: 338 nm) and operated at a column temperature of 40°C, mobile phase of 40% acetonitrile in H₂O, and a flow rate of 0.8 ml/min was used. An injection volume of 10 µL was used for all standard solutions and sample solutions tested. Abbreviation: N.D.: not detected.

Supplementary Table 3. Unannotated genes that showed significant recovery after Kampo formula treatment and a significant positive or negative correlation with immobility in the TST.

Gene_ID	Gene_name	Average read count (n = 5)					P-value (vs. P8/water)			Correlation (vs. TST)	
		R1/water	P8/water	P8/KS	P8/NKS	P8/HJG	R1/water	P8/KS	P8/NKS	P8/HJG	r
Saline/V-shaped recovered genes											
ENSMUSG00000082964	Rpl13-ps5	3.21	0.20	4.01	4.45	1.20	0.002	0.001	0.000	0.086	-0.419
ENSMUSG00000073752	Gm10570	24.84	9.82	18.44	18.56	13.34	0.001	0.032	0.030	0.310	-0.592
ENSMUSG00000077637	Gm22771	138.37	105.13	122.84	131.52	117.13	0.010	0.149	0.037	0.320	-0.410
ENSMUSG00000081431	Gm15483	22.83	13.29	21.87	22.55	21.38	0.040	0.060	0.045	0.073	-0.581
ENSMUSG00000102863	Gm37639	12.02	3.22	7.22	9.87	5.47	0.002	0.060	0.008	0.231	-0.590
ENSMUSG00000052865	Gm13619	7.21	2.01	6.63	8.08	3.84	0.018	0.028	0.009	0.250	-0.577
ENSMUSG00000097430	Gm10544	22.64	7.84	16.04	17.74	8.49	0.000	0.006	0.001	0.779	-0.441
ENSMUSG00000097648	9330185C12Rik	9.61	3.03	5.21	10.27	4.65	0.020	0.289	0.014	0.405	-0.419
ENSMUSG00000100455	Gm29170	398.86	334.44	380.46	394.50	364.66	0.013	0.070	0.020	0.226	-0.612
ENSMUSG00000106565	Gm43582	14.83	4.23	7.02	12.71	6.46	0.001	0.195	0.003	0.284	-0.455
ENSMUSG00000103283	ENSMUSG00000103283	87.70	68.78	82.17	91.76	75.62	0.042	0.139	0.016	0.434	-0.503
ENSMUSG00000056418	BC043934	33.25	16.43	27.45	27.86	14.98	0.005	0.045	0.039	0.731	-0.415
Saline/reverse V-shaped recovered genes											
ENSMUSG00000105003	Gm40055	3.21	9.64	6.02	3.63	6.88	0.019	0.281	0.034	0.436	0.552
ENSMUSG00000083914	Rps18-ps1	0.40	14.04	9.22	6.47	12.08	0.000	0.139	0.009	0.585	0.525
ENSMUSG00000047509	Gm6776	7.62	16.33	9.83	6.86	12.09	0.036	0.154	0.018	0.390	0.476
LPS/V-shaped recovered genes											
ENSMUSG00000100636	Gm3551	5.55	0.40	3.60	4.79	1.99	0.002	0.013	0.004	0.080	-0.591
ENSMUSG00000065694	Gm25411	6.97	1.78	4.61	5.99	3.58	0.011	0.084	0.024	0.215	-0.413
ENSMUSG00000097330	Gm26672	17.08	7.91	10.97	15.56	10.93	0.019	0.334	0.041	0.340	-0.477
ENSMUSG00000073371	Gm6594	34.74	21.75	33.51	33.68	33.02	0.030	0.046	0.043	0.054	-0.399
ENSMUSG00000087063	Gm15857	8.96	2.39	6.79	9.37	6.77	0.003	0.020	0.002	0.020	-0.413
ENSMUSG00000098164	Gm5493	9.18	2.97	5.19	9.17	3.59	0.032	0.303	0.032	0.733	-0.530
ENSMUSG00000069011	Gm10254	117.32	76.23	99.04	132.70	88.94	0.039	0.211	0.008	0.462	-0.472
ENSMUSG00000084828	Gm12367	17.11	9.28	13.17	16.56	11.33	0.030	0.224	0.040	0.494	-0.546
ENSMUSG00000102728	Gm37934	3.98	0.00	1.60	2.19	0.99	0.000	0.008	0.003	0.024	-0.465
ENSMUSG00000044081	Zfp85os	87.82	48.61	65.45	71.40	59.12	0.000	0.073	0.020	0.240	-0.594
ENSMUSG00000102193	Gm7299	5.57	0.40	3.39	5.80	2.98	0.015	0.049	0.014	0.065	-0.565
LPS/reverse V-shaped recovered genes											
ENSMUSG00000097301	AW121686	54.35	74.35	50.89	45.87	51.94	0.032	0.010	0.001	0.015	0.558
ENSMUSG00000085896	5330429C05Rik	7.37	20.00	14.19	7.17	15.70	0.034	0.449	0.030	0.590	0.490
ENSMUSG00000077316	n-R5s171	0.20	4.36	3.40	1.20	5.98	0.000	0.651	0.040	0.551	0.590
ENSMUSG00000080773	Gm12955	1.19	6.94	2.99	2.00	3.18	0.007	0.153	0.043	0.183	0.554
ENSMUSG00000086326	Gm13200	6.18	13.63	9.98	4.99	12.13	0.022	0.342	0.005	0.719	0.651

Correlation analysis was performed using Pearson's correlation coefficient.