

Supplementary Table8: 97 KEGG Pathway in XFZY decoction.

KEGG Pathway ID	KEGG Pathway	P-value
hsa04080	Neuroactive ligand-receptor interaction	1.48E-81
hsa01100	Metabolic pathways	9.22E-47
hsa04724	Glutamatergic synapse	5.32E-29
hsa04723	Retrograde endocannabinoid signaling	1.41E-20
hsa04727	GABAergic synapse	8.25E-20
hsa03050	Proteasome	8.79E-18
hsa00250	Alanine, aspartate and glutamate metabolism	1.33E-16
hsa04024	cAMP signaling pathway	3.15E-16
hsa04726	Serotonergic synapse	3.66E-15
hsa04713	Circadian entrainment	6.55E-15
hsa04062	Chemokine signaling pathway	1.19E-14
hsa04913	Ovarian steroidogenesis	6.51E-14
hsa00260	Glycine, serine and threonine metabolism	6.76E-14
hsa05033	Nicotine addiction	2.44E-13
hsa00591	Linoleic acid metabolism	3.21E-13
hsa04020	Calcium signaling pathway	7.30E-13
hsa05032	Morphine addiction	1.52E-12
hsa04750	Inflammatory mediator regulation of TRP channels	4.13E-12
hsa04742	Taste transduction	5.67E-12
hsa04270	Vascular smooth muscle contraction	1.03E-11
hsa01130	Biosynthesis of antibiotics	1.30E-11
hsa04923	Regulation of lipolysis in adipocytes	3.09E-11
hsa04912	GnRH signaling pathway	5.08E-10
hsa00562	Inositol phosphate metabolism	1.04E-09
hsa04725	Cholinergic synapse	2.11E-09
hsa00220	Arginine biosynthesis	2.23E-09
hsa00565	Ether lipid metabolism	2.24E-09
hsa01200	Carbon metabolism	4.47E-09
hsa00500	Starch and sucrose metabolism	8.79E-09
hsa00340	Histidine metabolism	1.24E-08
hsa04728	Dopaminergic synapse	1.39E-08
hsa00330	Arginine and proline metabolism	1.41E-08
hsa04611	Platelet activation	2.72E-08
hsa00564	Glycerophospholipid metabolism	3.53E-08
hsa04730	Long-term depression	5.81E-08
hsa01230	Biosynthesis of amino acids	7.28E-08
hsa00360	Phenylalanine metabolism	1.05E-07
hsa00592	alpha-Linolenic acid metabolism	1.23E-07
hsa00052	Galactose metabolism	2.30E-07
hsa00590	Arachidonic acid metabolism	6.70E-07

hsa04540	Gap junction	7.29E-07
hsa04975	Fat digestion and absorption	9.60E-07
hsa00970	Aminoacyl-tRNA biosynthesis	1.14E-06
hsa04720	Long-term potentiation	1.14E-06
hsa05031	Amphetamine addiction	1.14E-06
hsa04071	Sphingolipid signaling pathway	1.14E-06
hsa04915	Estrogen signaling pathway	1.42E-06
hsa00380	Tryptophan metabolism	1.73E-06
hsa04921	Oxytocin signaling pathway	2.22E-06
hsa00410	beta-Alanine metabolism	3.16E-06
hsa03320	PPAR signaling pathway	5.79E-06
hsa04920	Adipocytokine signaling pathway	5.89E-06
hsa04931	Insulin resistance	9.12E-06
hsa04922	Glucagon signaling pathway	1.04E-05
hsa04976	Bile secretion	1.24E-05
hsa05030	Cocaine addiction	1.25E-05
hsa02010	ABC transporters	1.58E-05
hsa04924	Renin secretion	1.70E-05
hsa00010	Glycolysis / Gluconeogenesis	1.70E-05
hsa04070	Phosphatidylinositol signaling system	1.87E-05
hsa04973	Carbohydrate digestion and absorption	2.17E-05
hsa00350	Tyrosine metabolism	3.80E-05
hsa04066	HIF-1 signaling pathway	4.76E-05
hsa04925	Aldosterone synthesis and secretion	9.00E-05
hsa05231	Choline metabolism in cancer	1.10E-04
hsa04911	Insulin secretion	1.13E-04
hsa04971	Gastric acid secretion	1.37E-04
hsa00270	Cysteine and methionine metabolism	1.78E-04
hsa00280	Valine, leucine and isoleucine degradation	2.14E-04
hsa04919	Thyroid hormone signaling pathway	2.26E-04
hsa04022	cGMP-PKG signaling pathway	4.58E-04
hsa05142	Chagas disease (American trypanosomiasis)	5.20E-04
hsa05211	Renal cell carcinoma	5.62E-04
hsa00650	Butanoate metabolism	6.87E-04
hsa00630	Glyoxylate and dicarboxylate metabolism	6.87E-04
hsa00670	One carbon pool by folate	0.001174613
hsa03013	RNA transport	0.001261309
hsa05200	Pathways in cancer	0.001310766
hsa00061	Fatty acid biosynthesis	0.001495345
hsa04910	Insulin signaling pathway	0.001764061
hsa00071	Fatty acid degradation	0.001957722
hsa00460	Cyanoamino acid metabolism	0.002534462
hsa04930	Type II diabetes mellitus	0.002613734

hsa00020	Citrate cycle (TCA cycle)	0.002743375
hsa04014	Ras signaling pathway	0.002719308
hsa00790	Folate biosynthesis	0.0030754
hsa04972	Pancreatic secretion	0.003942301
hsa04916	Melanogenesis	0.003905856
hsa05230	Central carbon metabolism in cancer	0.005853769
hsa00051	Fructose and mannose metabolism	0.005903938
hsa04370	VEGF signaling pathway	0.006259346
hsa01212	Fatty acid metabolism	0.006464457
hsa04970	Salivary secretion	0.007716165
hsa05143	African trypanosomiasis	0.008241077
hsa00983	Drug metabolism - other enzymes	0.00905385
hsa04914	Progesterone-mediated oocyte maturation	0.009175644
hsa04015	Rap1 signaling pathway	0.009615289