Supplementary Table 9:94 KEGG Pathway in GLXBBX decoction.

KEGG Pathway ID	KEGG Pathway	P value
hsa04080	Neuroactive ligand-receptor interaction	2.51E-98
hsa01100	Metabolic pathways	6.96E-52
hsa04724	Glutamatergic synapse	8.25E-32
hsa04723	Retrograde endocannabinoid signaling	2.24E-31
hsa04727	GABAergic synapse	8.96E-23
hsa04713	Circadian entrainment	6.48E-22
hsa04726	Serotonergic synapse	4.25E-21
hsa03050	Proteasome	7.00E-20
hsa00250	Alanine, aspartate and glutamate metabolism	1.35E-19
hsa01130	Biosynthesis of antibiotics	2.95E-19
hsa04020	Calcium signaling pathway	2.01E-17
hsa05032	Morphine addiction	2.28E-17
hsa00591	Linoleic acid metabolism	4.04E-17
hsa04024	cAMP signaling pathway	4.03E-17
hsa04062	Chemokine signaling pathway	2.68E-16
hsa05033	Nicotine addiction	3.36E-16
hsa04725	Cholinergic synapse	1.88E-14
hsa00620	Pyruvate metabolism	7.64E-14
hsa00590	Arachidonic acid metabolism	5.59E-13
hsa00565	Ether lipid metabolism	1.13E-12
hsa04750	Inflammatory mediator regulation of TRP channels	1.21E-12
hsa04923	Regulation of lipolysis in adipocytes	2.16E-12
hsa04270	Vascular smooth muscle contraction	2.96E-12
hsa01200	Carbon metabolism	3.48E-12
hsa00564	Glycerophospholipid metabolism	5.16E-12
hsa04742	Taste transduction	8.44E-12
hsa00592	alpha-Linolenic acid metabolism	1.35E-11
hsa04915	Estrogen signaling pathway	3.27E-11
hsa03320	PPAR signaling pathway	6.94E-11
hsa04728	Dopaminergic synapse	9.36E-11
hsa04912	GnRH signaling pathway	2.70E-10
hsa00010	Glycolysis / Gluconeogenesis	3.54E-10
hsa00220	Arginine biosynthesis	1.19E-09
hsa04913	Ovarian steroidogenesis	1.17E-09
hsa00052	Galactose metabolism	4.66E-09
hsa00071	Fatty acid degradation	1.06E-08
hsa00260	Glycine, serine and threonine metabolism	1.41E-08
hsa04730	Long-term depression	2.45E-08
hsa00270	Cysteine and methionine metabolism	4.66E-08

hsa00500	Starch and sucrose metabolism	5.79E-08
hsa04611	Platelet activation	5.93E-08
hsa04921	Oxytocin signaling pathway	1.00E-07
hsa04924	Renin secretion	1.59E-07
hsa00350	Tyrosine metabolism	2.38E-07
hsa00330	Arginine and proline metabolism	3.28E-07
hsa04975	Fat digestion and absorption	5.07E-07
hsa01230	Biosynthesis of amino acids	6.47E-07
hsa04922	Glucagon signaling pathway	8.10E-07
hsa00380	Tryptophan metabolism	8.79E-07
hsa04720	Long-term potentiation	1.38E-06
hsa04971	Gastric acid secretion	1.51E-06
hsa00650	Butanoate metabolism	1.55E-06
hsa04540	Gap junction	2.15E-06
hsa04973	Carbohydrate digestion and absorption	2.50E-06
hsa00640	Propanoate metabolism	3.10E-06
hsa04066	HIF-1 signaling pathway	4.91E-06
hsa00280	Valine, leucine and isoleucine degradation	6.25E-06
hsa04972	Pancreatic secretion	9.58E-06
hsa00630	Glyoxylate and dicarboxylate metabolism	9.90E-06
hsa00561	Glycerolipid metabolism	1.01E-05
hsa00020	Citrate cycle (TCA cycle)	1.08E-05
hsa05030	Cocaine addiction	1.43E-05
hsa05031	Amphetamine addiction	1.51E-05
hsa00410	beta-Alanine metabolism	1.90E-05
hsa04911	Insulin secretion	2.11E-05
hsa00340	Histidine metabolism	2.54E-05
hsa04022	cGMP-PKG signaling pathway	3.76E-05
hsa04071	Sphingolipid signaling pathway	6.38E-05
hsa04970	Salivary secretion	7.75E-05
hsa04919	Thyroid hormone signaling pathway	9.63E-05
hsa04976	Bile secretion	1.18E-04
hsa04916	Melanogenesis	1.41E-04
hsa05142	Chagas disease (American trypanosomiasis)	1.42E-04
hsa04931	Insulin resistance	1.41E-04
hsa05200	Pathways in cancer	1.44E-04
hsa04920	Adipocytokine signaling pathway	1.50E-04
hsa00061	Fatty acid biosynthesis	2.63E-04
hsa04925	Aldosterone synthesis and secretion	3.76E-04
hsa05230	Central carbon metabolism in cancer	6.40E-04
hsa00360	Phenylalanine metabolism	7.21E-04
hsa00460	Cyanoamino acid metabolism	7.20E-04
hsa00670	One carbon pool by folate	7.55E-04

hsa01212	Fatty acid metabolism	0.001275309
hsa00860	Porphyrin and chlorophyll metabolism	0.001872446
hsa05211	Renal cell carcinoma	0.002129168
hsa00760	Nicotinate and nicotinamide metabolism	0.002599116
hsa04918	Thyroid hormone synthesis	0.002693559
hsa04930	Type II diabetes mellitus	0.003625251
hsa01210	2-Oxocarboxylic acid metabolism	0.003847084
hsa04014	Ras signaling pathway	0.007472606
hsa05231	Choline metabolism in cancer	0.008580881
hsa04977	Vitamin digestion and absorption	0.008670163
hsa05204	Chemical carcinogenesis	0.008706112
hsa05034	Alcoholism	0.008754223