

TableS5: Kyoto Encyclopedia of Genes and Genomes (KEGG) pathways analysis of the possible downstream target genes.

Id	Term	P value	Count
hsa04066	Hippo signaling pathway	0.00019568	8
hsa00590	Progesterone-mediated oocyte maturation	0.00356439	5
hsa04216	Propanoate metabolism	0.00497883	3
hsa01230	Cushing syndrome	0.00507869	6
hsa05140	Gastric acid secretion	0.0075956	4
hsa01210	Tight junction	0.00768132	6
hsa05206	Ovarian steroidogenesis	0.01529508	3
hsa05167	Melanogenesis	0.01994433	4
hsa00270	Protein digestion and absorption	0.0212688	4
hsa00220	Wnt signaling pathway	0.02414112	5