

GeDiPNet Cluster 599 -- Enriched GO Terms (Biological Process)

GO term enrichment: Biological Process terms (gene2go, human/tax_id=9606 only), upper-tail hypergeometric test, fold enrichment = (overlap / cluster gene count) / (term gene count / universe size), Benjamini-Hochberg FDR-corrected. Top 15 terms per cluster by q-value. See scripts/build_cluster_go_enrichment.php on gedipnet.bicnirrh.res.in. Data as of: 2026-10-20 03:23:14. · Rows: 7 · Generated: 2026-10-20 00:01:15

GO Term	GO ID	Overlap Genes	Overlap Gene Names	Term Gene Count	Cluster Gene Count	Fold Enrichment	P-value	FDR Q-value
fatty acid derivative metabolic process	GO:1901568	1	FAR1	1	1	18687.00	5.351e-5	1.943e-3
wax biosynthetic process	GO:0010025	1	FAR1	4	1	4671.75	2.141e-4	5.314e-3
long-chain fatty-acyl-CoA metabolic process	GO:0035336	1	FAR1	7	1	2669.57	3.746e-4	7.813e-3
glycerophospholipid biosynthetic process	GO:0046474	1	FAR1	7	1	2669.57	3.746e-4	7.813e-3
ether lipid biosynthetic process	GO:0008611	1	FAR1	8	1	2335.88	4.281e-4	8.531e-3
lipid biosynthetic process	GO:0008610	1	FAR1	33	1	566.27	1.766e-3	1.990e-2
lipid metabolic process	GO:0006629	1	FAR1	840	1	22.25	4.495e-2	1.007e-1