

GO id	Term	Annot. genes	Sign. genes	Expected	P-value	Adj. P-value
GO:0004653	polypeptide N-acetylgalactosaminyltransf...	9	3	0.18	0.0006	0.0407
GO:0097322	7SK snRNA binding	3	2	0.06	0.0012	0.0407
GO:0004594	pantothenate kinase activity	4	2	0.08	0.0023	0.0407
GO:0042577	lipid phosphatase activity	6	2	0.12	0.0056	0.0407