GO id	Term	Annot. genes	Sign. genes	Expected	P-value	Adj. P-value
GO:0003823	antigen binding	60	6	1.03	0.00054	0.0382
GO:0003976	UDP-N-acetylglucosamine-lysosomal-enzyme	3	2	0.05	0.00086	0.0382
GO:0016780	phosphotransferase activity, for other s	18	3	0.31	0.00333	0.0382
GO:0005126	cytokine receptor binding	149	8	2.55	0.00408	0.0382
GO:0034987	immunoglobulin receptor binding	6	2	0.10	0.00418	0.0382
GO:0035064	methylated histone binding	40	4	0.68	0.00470	0.0382
GO:0042393	histone binding	124	7	2.12	0.00541	0.0382
GO:0005164	tumor necrosis factor receptor binding	25	3	0.43	0.00861	0.0382
GO:0042887	amide transmembrane transporter activity	9	2	0.15	0.00969	0.0382