R-HSA-6	R-HSA.	R-HSA 147 ₄₂₉₀	R-HSA *3>3>55	immu *422475	innnune only	
	385 ₄₉₁	47 ₄ 290	3/3/55	*22 _{4>5}	'e only	
Pairwise path length in reference tree	0.012	0.029	0.014	0.161	0.028	
Jaccard distance over extended gene set	0.185	0.398	0.238	2.246	0.31	
Dijkstra BMA PPI	107.92	943.024	302.419	61498.08	4836.394	
WM distance over summary W2V	24.817	401.254	25.768	8212.347	NaN	
WM distance over gene symbols W2V	0.814	16.67	2.325	328.283	9500.508	
Cosine distance over over summary W2V	0.886	6.35	1.258	215.654	NaN	
Cosine distance over over NCBI summary W2V	3.384	54.313	7.583	737.154	101.634	
Cosine distance over gene symbols W2V	0.202	0.761	0.285	17.467	1.483	
Cosine distance GO MF description W2V	9.172	27.654	24.075	2348.945	206.04	
Cosine distance GO CC description W2V	5.542	52.808	28.758	2191.575	135.862	
Cosine distance GO BP description W2V	11.603	154.268	124.624	20266.078	361.471	
GO-distance (go_type=MF, measure=Wang, combine=BMA)	11.884	2588.84	301.245	2601.024	1954.441	
GO-distance (go_type=CC, measure=Wang, combine=BMA)	5.406	72.14	502.832	3835.697	5870.326	
GO-distance (go_type=BP, measure=Wang, combine=BMA)	24.084	826.678	1598.533	77040.866	48340.111	
Overlap distance over genes	0.239	3.488	0.7	281.89	11.426	
Overlap distance over gene traits	0.22	6.589	1.77	326.871	6.641	
Minkowski distance (p=2) over genes	0.001	0.003	0.001	0.221	0.016	
Minkowski distance $(p=2)$ over gene traits	0.001	0.007	0.003	0.276	0.013	
Minkowski distance (p=2) over gene trait frequency	0.001	0.018	0.006	0.327	0.192	
Minkowski distance (p=1) over genes	0.0	0.003	0.001	0.161	0.016	
Minkowski distance ($p=1$) over gene traits	0.0	0.007	0.003	0.204	0.014	
Minkowski distance (p=1) over gene trait frequency	0.001	0.016	0.005	0.279	0.192	
Kappa distance over genes	0.326	1.407	0.485	38.138	0.97	
Kappa distance over gene traits	0.524	1.664	0.536	41.809	0.638	
Jaccard distance over genes	0.0	0.001	0.001	0.053	0.008	
Jaccard distance over gene traits	0.0	0.004	0.002	0.066	0.009	
Cosine distance over gene trait frequency	0.001	0.013	0.004	0.115	0.197	
Random (uniform, (0,1))	0.012	0.031	0.016	0.129	0.014	