

Table S11. Fraction of hubs annotated with GO terms in each network.

network	BP	CC	MF	none
Human-hq	97	96	74	3
Yeast-hq	92	96	71	2
Fly-hq	68	63	66	21
Athal	71	58	69	13
Ecoli	48	39	49	35
Human-all	85	86	61	10
Yeast-all	94	98	79	0

For each network and for each ontology (Biological process, Cellular component, Molecular function), we show the percentage of hubs annotated with at least some terms other than the root (the most general) term in this ontology (BP, CC, MF), and the percentage of hubs not annotated in any of the three ontologies (none).