
Figure 6 - Figure Supplement 1: Network characteristics for all gene set enrichment analyses

The detailed results for each network are presented in Figure 6 - Figure Supplement 2 (for 60 most abundant proteins) and Figure 6 - Figure Supplement 3 (for significantly differing proteins, divided by comparison).

Proteins used	N nodes	N edges	Avg. local clustering coefficient	Avg. node degree:
60 most abundant	41	32	0.361	1.56
Significant in 2/3, in any comparison	44	44	0.312	2
Significant in 2/3, Nurse vs forager	19	6	0.158	0.632
Significant in 2/3, Young vs mature	10	4	0.5	0.8
Significant in 2/3, Field vs lab	20	11	0.583	1.1