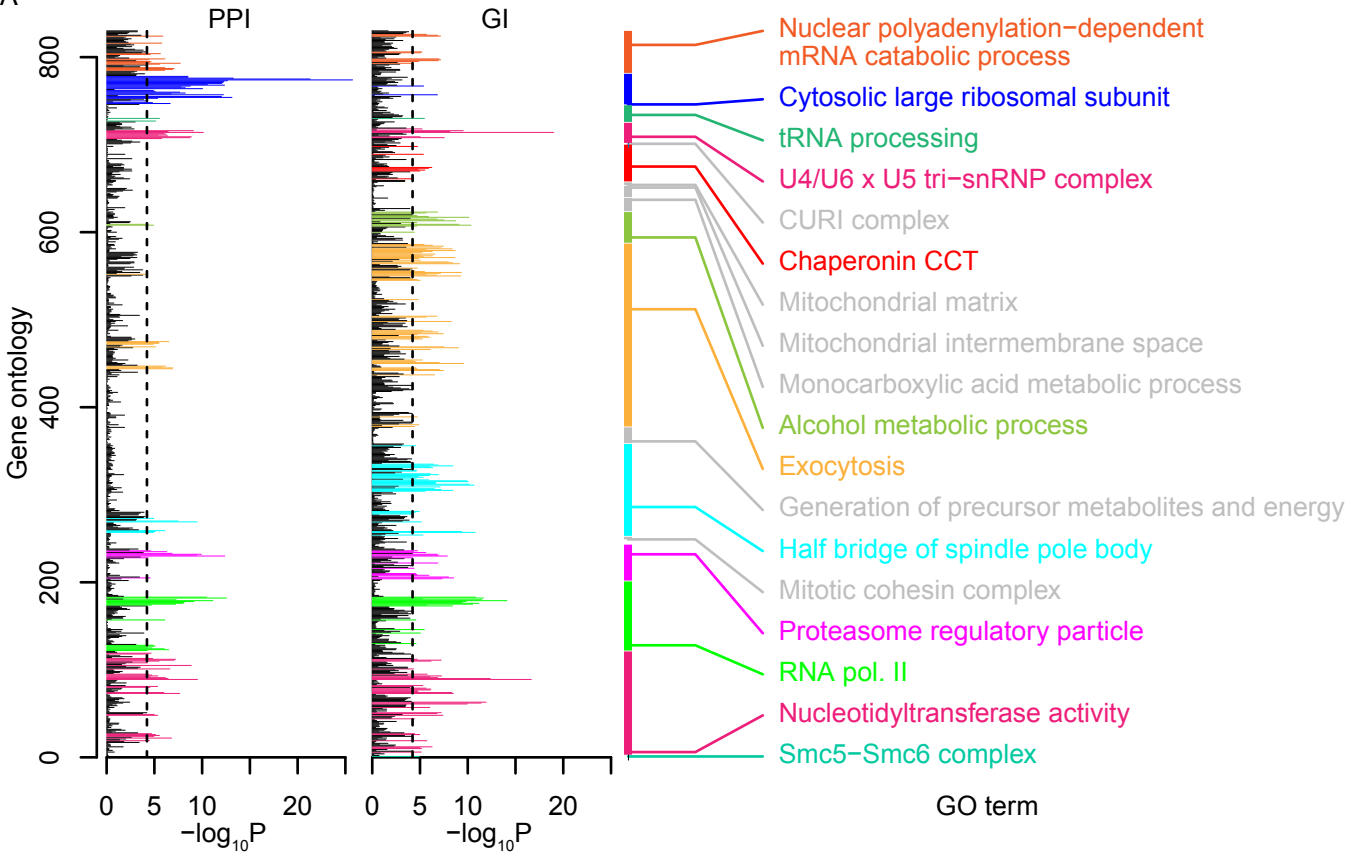
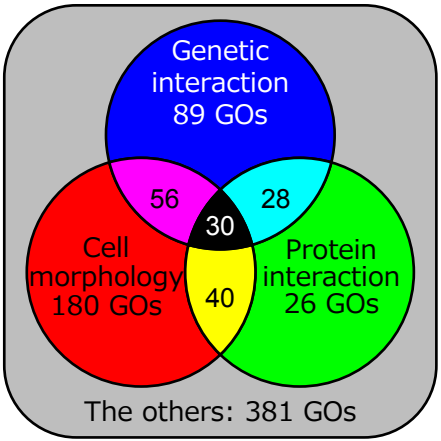
A


PPI
GI
Gene ontology
0
200
400
600
800
0 5 10 20
−log10P
0 5 10 20
−log10P
Nuclear polyadenylation−dependent mRNA catabolic process
Cytosolic large ribosomal subunit
tRNA processing
U4/U6 x U5 tri−snRNP complex
CURI complex
Chaperonin CCT
Mitochondrial matrix
Mitochondrial intermembrane space
Monocarboxylic acid metabolic process
Alcohol metabolic process
Exocytosis
Generation of precursor metabolites and energy
Half bridge of spindle pole body
Mitotic cohesin complex
Proteasome regulatory particle
RNA pol. II
Nucleotidyltransferase activity
Smc5−Smc6 complex
GO term

B


Genetic interaction 89 GOs
56
28
30
Cell morphology 180 GOs
40
Protein interaction 26 GOs
The others: 381 GOs