EmapPlot (Top 15) GO: MF – siRNA_01_vs_Control

anaphase−promoting complex binding
(p=2.3e−05, Size=6)

cytoskeletal motor activity
(p=2.3e−05, Size=18)

helicase activity
(p=5.3e−07, Size=25)

tubulin binding
(p=7.7e−09, Size=44)

ATP−dependent activity, acting on DNA
(p=2.6e−11, Size=27)

DNA helicase activity
(p=1.5e−10, Size=18)

ATP hydrolysis activity
(p=3.4e−07, Size=46)

microtubule motor activity
(p=7.3e−07, Size=16)

catalytic activity, acting on DNA
(p=2.7e−12, Size=40)

microtubule binding
(p=9.3e−11, Size=40)

single−stranded DNA helicase activity
(p=2.7e−12, Size=14)

number of genes
10
20
30
40

p.adjust
2.0e−05
1.5e−05
1.0e−05
5.0e−06

single−stranded DNA binding
(p=5.5e−10, Size=25)

cyclin−dependent protein serine/threonine kinase regulator activity
(p=1.1e−05, Size=12)

DNA secondary structure binding
(p=1.0e−06, Size=12)

damaged DNA binding
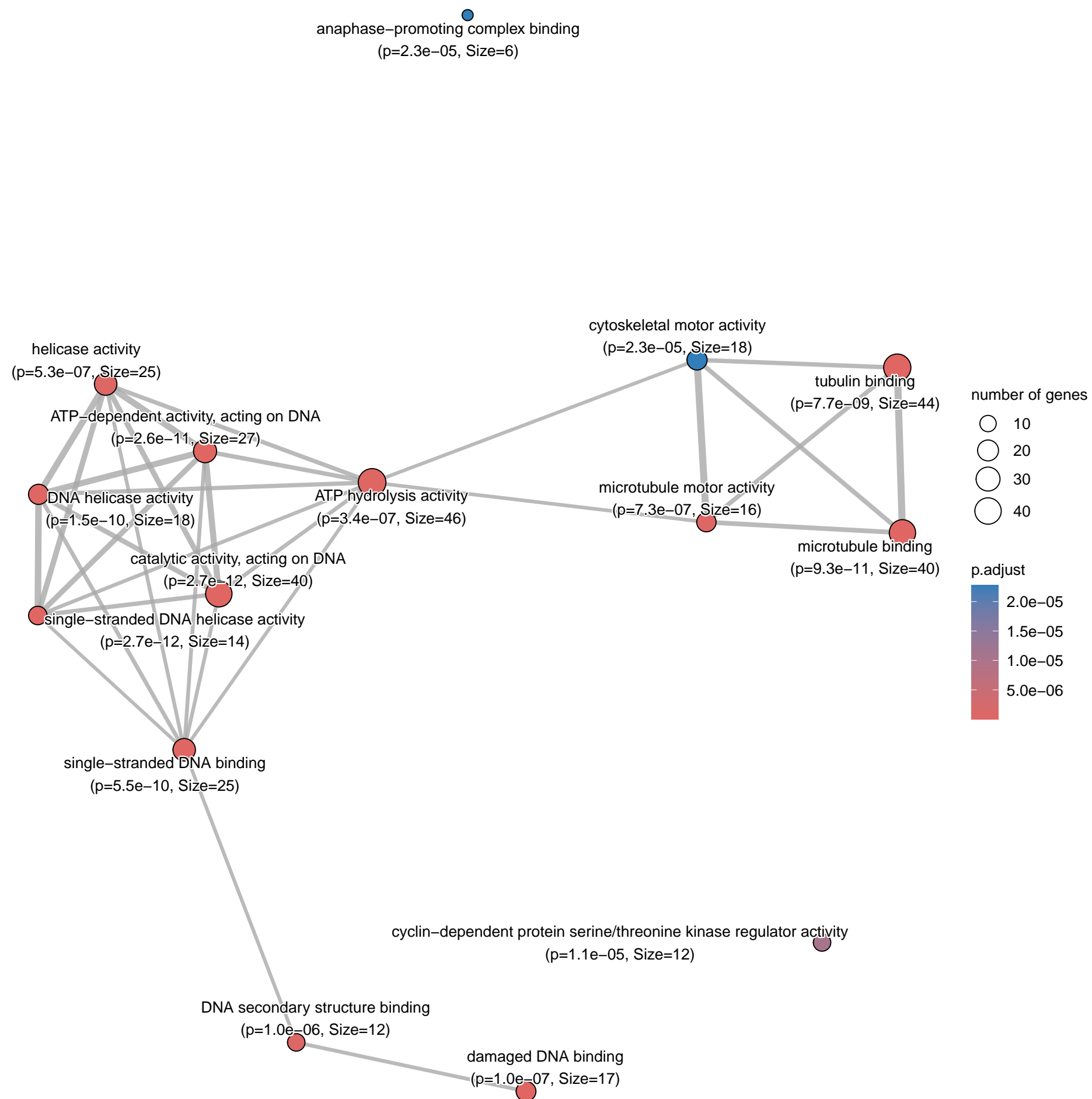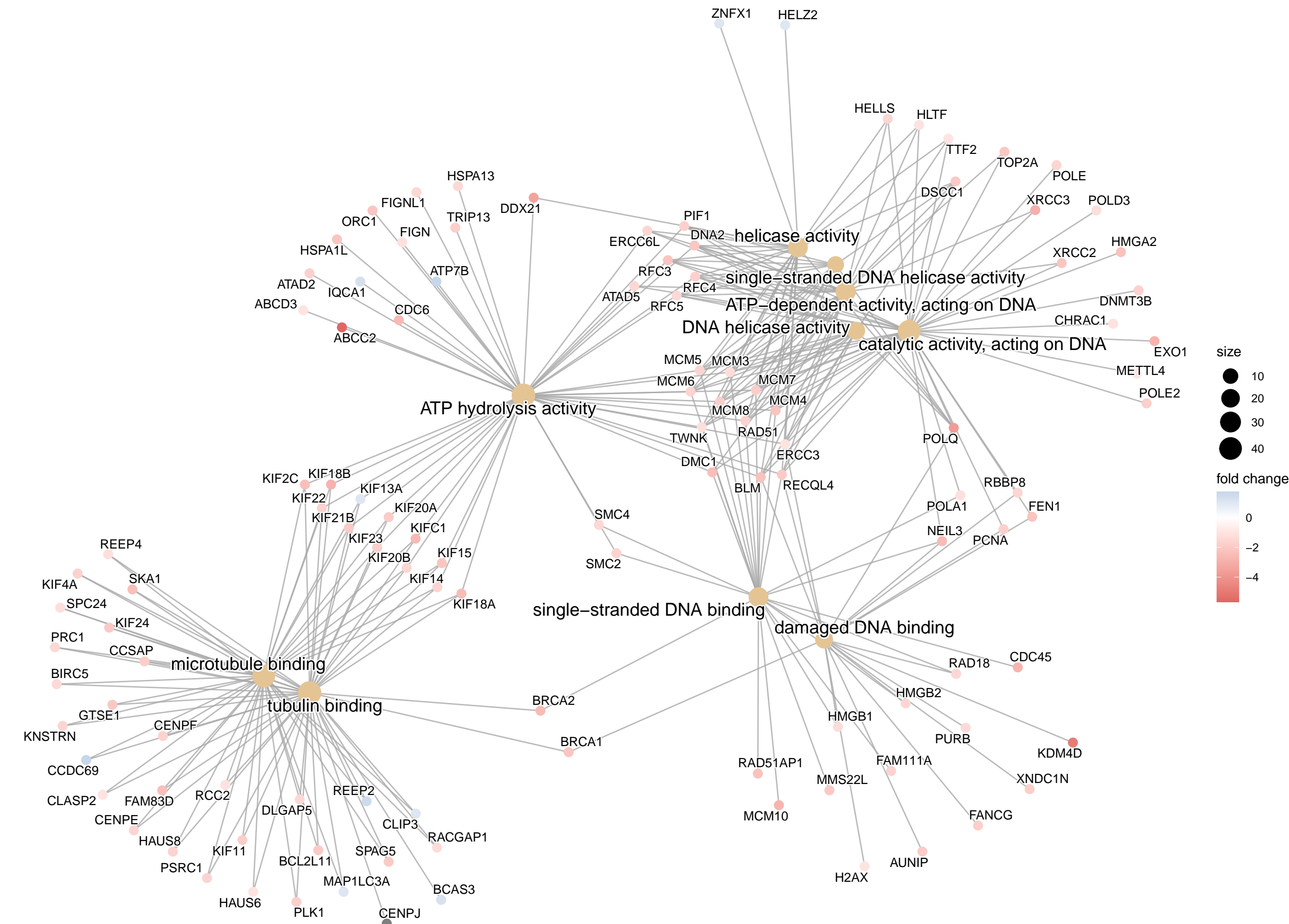(p=1.0e−07, Size=17)

Chord Chart / CnetPlot (Top 10) GO: MF – siRNA_01_vs_Control

DotPlot (Top 15) GO: MF – siRNA_01_vs_Control