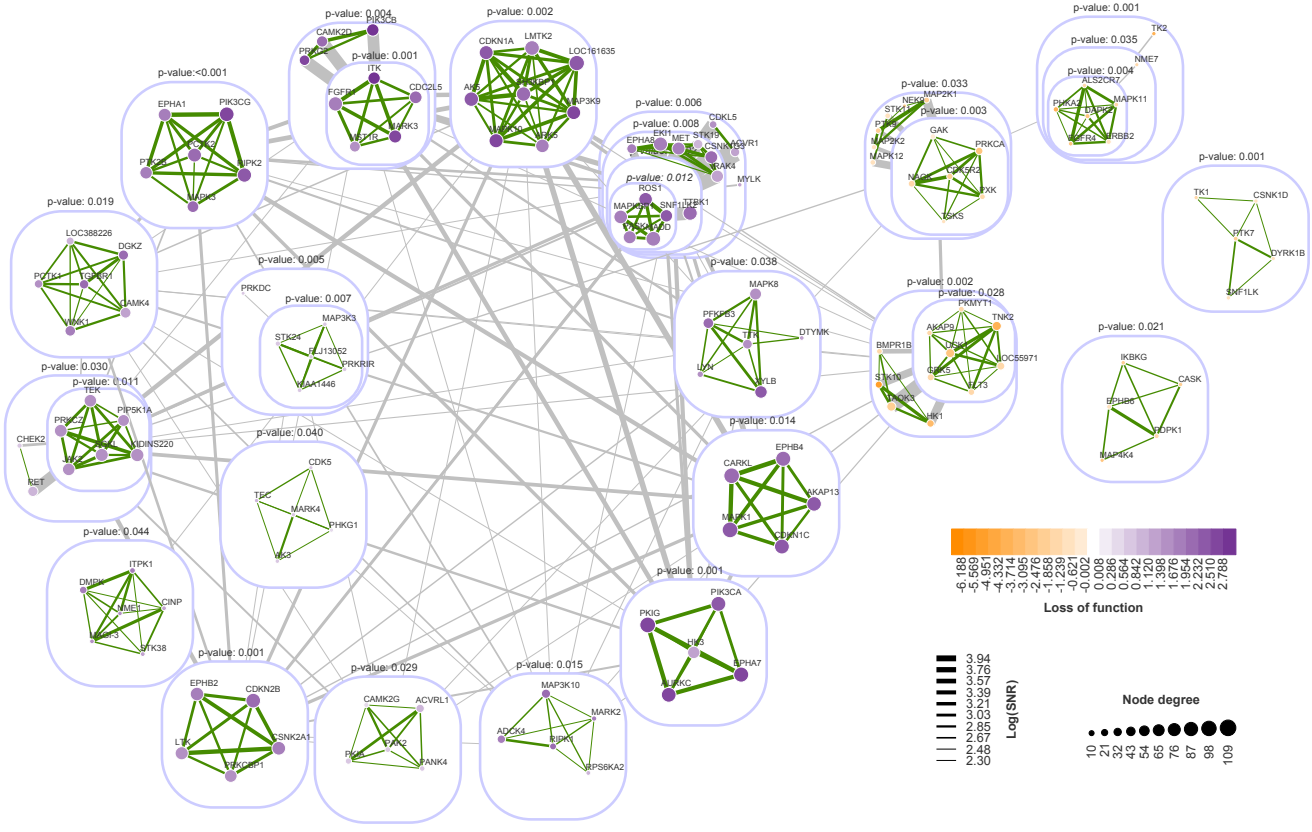
p-value: 0.004
PIK3CB
CAMK2D
PRKCE
p-value: 0.001
ITK
CD2L5
FGFR1
MAPK3
MAPK1
p-value:<0.001
EPHA1
PIK3CG
PIP5K1A
PIP4K2
MAPK3
p-value: 0.002
CDKN1A
LMTK2
LOC161635
AKT2
MAP3K9
MAPK1
MAPK3
p-value: 0.006
CDKL5
p-value: 0.008
EPHA2
EIF2AK3
MET
STK10
CSNK1G3
TRAK4
MYLK
p-value: 0.012
ROS1
MAPK10
SNF1LK
ITPKB
TTBK1
PTK2B
p-value: 0.019
LOC388226
DGKZ
PDPK1
TGFBR2
CAMK4
MAK1
p-value: 0.005
PRKDC
p-value: 0.007
MAP3K3
STK24
RIPK3
PRKRIR
MAP3K11
p-value: 0.030
p-value: 0.011
TEK
PRKCB
PIP5K1A
CHEK2
MIDINS220
MAP2K2
RET
p-value: 0.040
CDK5
TEC
MARK4
PHKG1
MAP3K3
p-value: 0.044
ITPK1
DMPK
CINP
MINK1
MAP3K3
STK38
p-value: 0.001
EPHB2
CDKN2B
LTK
CSNK2A1
PRKCBP1
p-value: 0.029
CAMK2G
ACVRL1
PRKCE
TRAK2
PANK4
p-value: 0.015
MAP3K10
MARK2
ADCK4
RIPK1
RPS6KA2
p-value: 0.038
MAPK8
PFKFB3
TTK
DTYMK
UCK2
MYLB
p-value: 0.014
EPHB4
CARKL
AKAP13
MAPK1
CDKN1C
p-value: 0.001
PIK3CA
PKIG
HIS3
ALPHA7
MAPKKC
p-value: 0.033
NEK9
MAP2K1
STK11
PTK2
MAP2K2
MAPK12
p-value: 0.003
GAK
PRKCA
MAPK8
CDK5R2
PXK
IGKS
p-value: 0.001
TK2
p-value: 0.035
NME7
p-value: 0.004
ALS2CR7
PHKA2
MAPK11
DAPP1
MAPK3
MAPK1
TRRAP
TRRBB2
p-value: 0.001
TK1
CSNK1D
PTK7
DYRK1B
SNF1LK
p-value: 0.002
p-value: 0.028
PKMYT1
AKAP9
TNK2
DGKA
LOC55971
CDK5
MLT3
BMPR1B
STK10
MAPK3
HIK1
p-value: 0.021
IKBKG
CASK
EPHB6
PDPK1
MAP4K4
-6.188
-5.569
-4.951
-4.332
-3.714
-3.095
-2.476
-1.858
-1.239
-0.621
-0.002
0.008
0.286
0.564
0.842
1.120
1.398
1.676
1.954
2.232
2.510
2.788
Loss of function
3.94
3.76
3.57
3.39
3.21
3.03
2.85
2.67
2.48
2.30
Log(SNR)
Node degree
10
21
32
43
54
65
76
87
98
109