Overlapping correction

Overlapping percentage between subnetworks

10
30
50
70
100

Regulation of nuclear beta catenin signaling and target gene transcription (p = 0.011)
Regulation of retinoblastoma protein (p = 0.031)
Regulation of CDC42 activity (p = 0.047)
Reelin signaling pathway (p = 0.033)
LPA receptor mediated events (p = 0.002)
PAR1–mediated thrombin signaling events (p = 0.006)
Hedgehog signaling events mediated by Gli proteins (p = 0.008)
ErbB receptor signaling network (p = 0.032)
ErbB2/ErbB3 signaling events (p = 0.047)
ErbB4 signaling events (p = 0.021)
Posttranslational regulation of adherens junction stability and disassembly (p = 0.023)
E–cadherin signaling in the nascent adherens junction (p = 0.014)
Regulation of RAC1 activity (p = 0.001)
CXCR4–mediated signaling events (p = 0.006)
PDGFR–beta signaling pathway (p = 0.001)
Ephrin B reverse signaling (p = 0.015)
IL8– and CXCR2–mediated signaling events (p = 0.043)
CD40/CD40L signaling (p = 0.045)
Signaling events regulated by Ret tyrosine kinase (p = 0.021)
Plasma membrane estrogen receptor signaling (p = 0.008)
Neurotrophic factor–mediated Trk receptor signaling (p = 0.005)
TCR signaling in naïve CD4+ T cells (p = 0.003)
TCR signaling in naïve CD8+ T cells (p = 0.004)
BCR signaling pathway (p = 0.022)
SHP2 signaling (p = 0.003)
Arf6 trafficking events (p = 0.012)
ALK2 signaling events (p = 0.028)
Endogenous TLR signaling (p = 0.033)
amb2 Integrin signaling (p = 0.003)
Beta3 integrin cell surface interactions (p = 0.008)