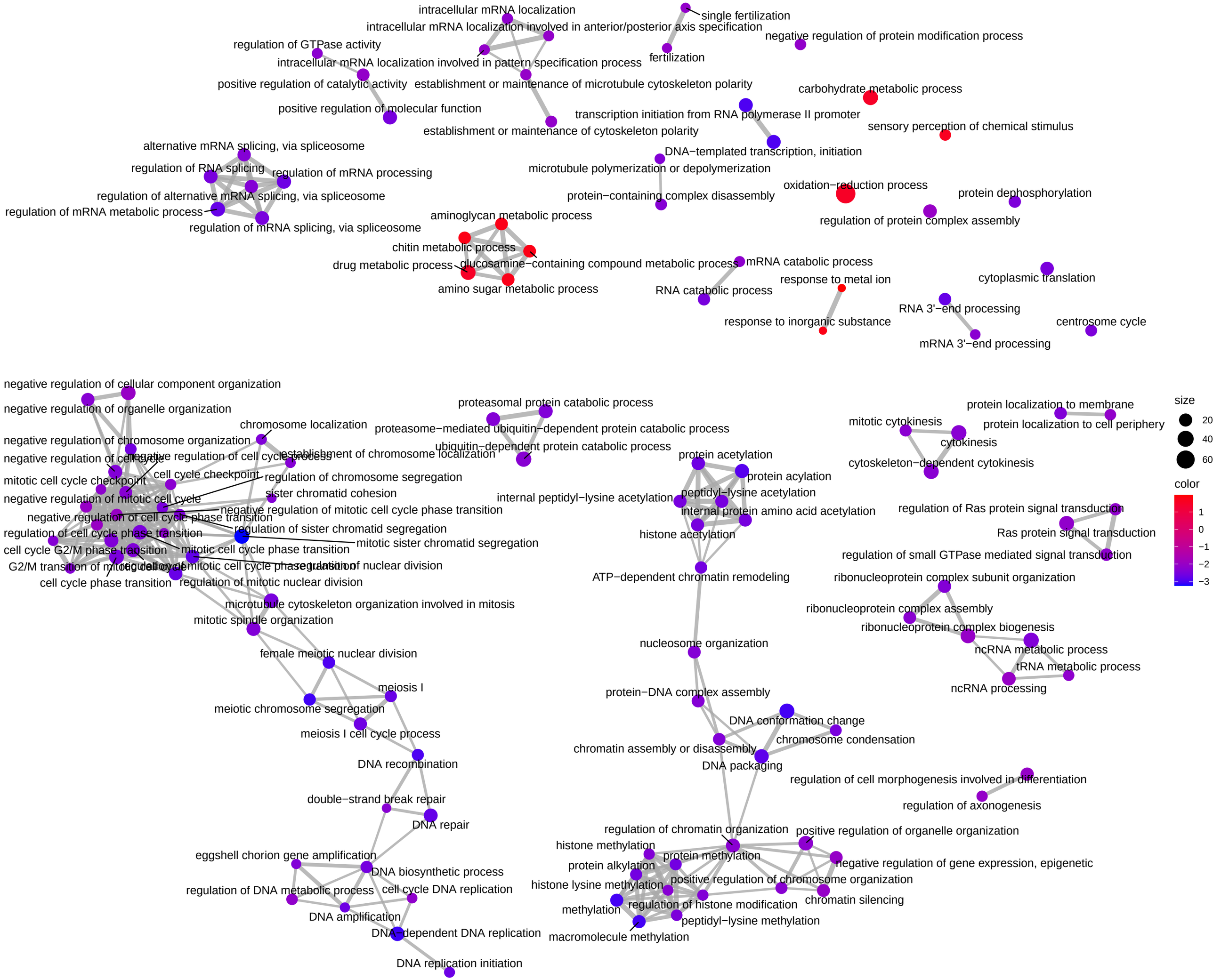
intracellular mRNA localization
intracellular mRNA localization involved in anterior/posterior axis specification
regulation of GTPase activity
intracellular mRNA localization involved in pattern specification process
positive regulation of catalytic activity
establishment or maintenance of microtubule cytoskeleton polarity
positive regulation of molecular function
establishment or maintenance of cytoskeleton polarity
single fertilization
fertilization
negative regulation of protein modification process
carbohydrate metabolic process
transcription initiation from RNA polymerase II promoter
sensory perception of chemical stimulus
DNA−templated transcription, initiation
alternative mRNA splicing, via spliceosome
regulation of RNA splicing
regulation of mRNA processing
microtubule polymerization or depolymerization
regulation of alternative mRNA splicing, via spliceosome
protein−containing complex disassembly
oxidation−reduction process
protein dephosphorylation
regulation of mRNA metabolic process
regulation of mRNA splicing, via spliceosome
regulation of protein complex assembly
aminoglycan metabolic process
chitin metabolic process
drug metabolic process
glucosamine−containing compound metabolic process
mRNA catabolic process
amino sugar metabolic process
RNA catabolic process
response to metal ion
cytoplasmic translation
RNA 3'−end processing
response to inorganic substance
centrosome cycle
mRNA 3'−end processing
negative regulation of cellular component organization
negative regulation of organelle organization
chromosome localization
negative regulation of chromosome organization
negative regulation of cell cycle process
establishment of chromosome localization
negative regulation of cell cycle
cell cycle checkpoint
regulation of chromosome segregation
mitotic cell cycle checkpoint
sister chromatid cohesion
negative regulation of mitotic cell cycle
negative regulation of mitotic cell cycle phase transition
negative regulation of cell cycle phase transition
regulation of sister chromatid segregation
regulation of cell cycle phase transition
mitotic sister chromatid segregation
cell cycle G2/M phase transition
mitotic cell cycle phase transition
G2/M transition of mitotic cell cycle
regulation of nuclear division
cell cycle phase transition
regulation of mitotic nuclear division
microtubule cytoskeleton organization involved in mitosis
mitotic spindle organization
female meiotic nuclear division
meiosis I
meiotic chromosome segregation
meiosis I cell cycle process
DNA recombination
double−strand break repair
DNA repair
eggshell chorion gene amplification
DNA biosynthetic process
regulation of DNA metabolic process
cell cycle DNA replication
DNA amplification
DNA−dependent DNA replication
DNA replication initiation
proteasomal protein catabolic process
proteasome−mediated ubiquitin−dependent protein catabolic process
ubiquitin−dependent protein catabolic process
protein acetylation
protein acylation
internal peptidyl−lysine acetylation
peptidyl−lysine acetylation
internal protein amino acid acetylation
histone acetylation
ATP−dependent chromatin remodeling
nucleosome organization
protein−DNA complex assembly
DNA conformation change
chromosome condensation
chromatin assembly or disassembly
DNA packaging
regulation of chromatin organization
positive regulation of organelle organization
histone methylation
protein methylation
protein alkylation
negative regulation of gene expression, epigenetic
histone lysine methylation
positive regulation of chromosome organization
chromatin silencing
methylation
regulation of histone modification
peptidyl−lysine methylation
macromolecule methylation
protein localization to membrane
mitotic cytokinesis
protein localization to cell periphery
cytokinesis
cytoskeleton−dependent cytokinesis
regulation of Ras protein signal transduction
Ras protein signal transduction
regulation of small GTPase mediated signal transduction
ribonucleoprotein complex subunit organization
ribonucleoprotein complex assembly
ribonucleoprotein complex biogenesis
ncRNA metabolic process
tRNA metabolic process
ncRNA processing
regulation of cell morphogenesis involved in differentiation
regulation of axonogenesis
size
20
40
60
color
1
0
−1
−2
−3