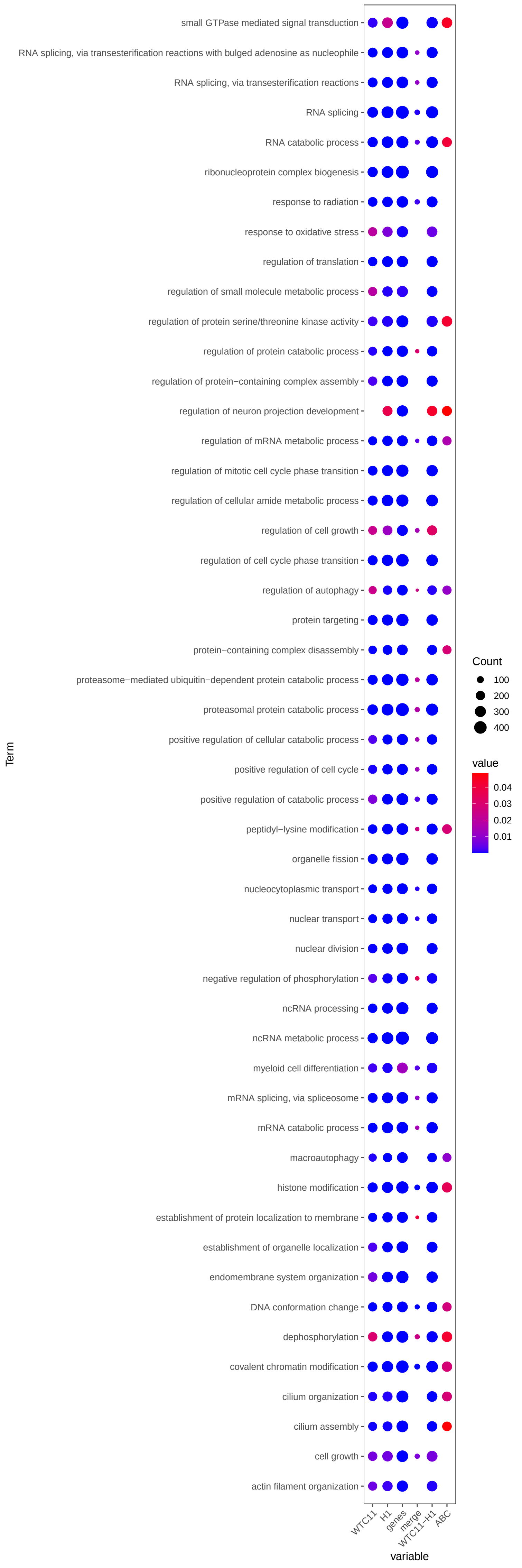
small GTPase mediated signal transduction
RNA splicing, via transesterification reactions with bulged adenosine as nucleophile
RNA splicing, via transesterification reactions
RNA splicing
RNA catabolic process
ribonucleoprotein complex biogenesis
response to radiation
response to oxidative stress
regulation of translation
regulation of small molecule metabolic process
regulation of protein serine/threonine kinase activity
regulation of protein catabolic process
regulation of protein−containing complex assembly
regulation of neuron projection development
regulation of mRNA metabolic process
regulation of mitotic cell cycle phase transition
regulation of cellular amide metabolic process
regulation of cell growth
regulation of cell cycle phase transition
regulation of autophagy
protein targeting
protein−containing complex disassembly
proteasome−mediated ubiquitin−dependent protein catabolic process
proteasomal protein catabolic process
positive regulation of cellular catabolic process
positive regulation of cell cycle
positive regulation of catabolic process
peptidyl−lysine modification
organelle fission
nucleocytoplasmic transport
nuclear transport
nuclear division
negative regulation of phosphorylation
ncRNA processing
ncRNA metabolic process
myeloid cell differentiation
mRNA splicing, via spliceosome
mRNA catabolic process
macroautophagy
histone modification
establishment of protein localization to membrane
establishment of organelle localization
endomembrane system organization
DNA conformation change
dephosphorylation
covalent chromatin modification
cilium organization
cilium assembly
cell growth
actin filament organization
Term
WTC11
H1
genes
merge
WTC11−H1
ABC
variable
Count
100
200
300
400
value
0.04
0.03
0.02
0.01