

# GO\_Biological\_Process\_2021

Term

positive regulation of microtubule  
nucleation (GO:0090063)

positive regulation of nucleotide-binding  
oligomerization domain containing signaling  
pathway (GO:0070426)

regulation of microtubule nucleation  
(GO:0010968)

regulation of nucleotide-binding  
oligomerization domain containing 2  
signaling pathway (GO:0070432)

heat acclimation (GO:0010286)

cellular heat acclimation (GO:0070370)

regulation of transforming growth factor  
beta activation (GO:1901388)

G-quadruplex DNA unwinding (GO:0044806)

positive regulation of chromosome  
segregation (GO:0051984)

regulation of spindle assembly (GO:0090169)

0

2

4

6

Enrichment log(combined score)

# GO\_Cellular\_Component\_2021

Term

glial cell projection (GO:0097386)

meiotic spindle (GO:0072687)

endoplasmic reticulum tubular network  
membrane (GO:0098826)

polysomal ribosome (GO:0042788)

platelet dense granule membrane (GO:0031088)

nuclear stress granule (GO:0097165)

azurophil granule membrane (GO:0035577)

cytolytic granule (GO:0044194)

astral microtubule (GO:0000235)

astrocyte projection (GO:0097449)

RISC-loading complex (GO:0070578)

0

2

4

Enrichment log(combined score)

# GO\_Molecular\_Function\_2021

