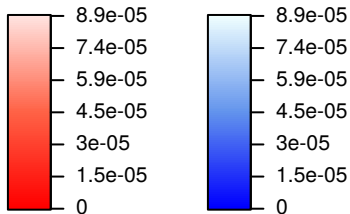


p-values up

p-values down

reporter, distinct-directional (both)



Max node size:
772 genes

Min node size:
12 genes

Max edge width:
321 genes

Min edge width:
3 genes

