

Estimated fold change in population-wide mutation rate

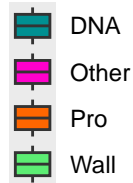
Kruskal-Wallis, $p = 0.02$

0.042

0.0055

0.93

mode_of_action



100
10
1
0.1

DNA

Other

Pro

Wall

mode_of_action

