

relative transcript abundance

0.2
0.1
0.0

strain
atten
wt

genes

0.3 0.1 1.2 2.3 2.6 2.1 0.2 2.8 3.3 3.9 4.2 4.3 4.1 5.3 5.9 6.6 6.1 7.1 8.1 9.1 9.2 10.2 11.3 12.4 13.5 14.6 15.7 17.5 18.5 19.1 19.5