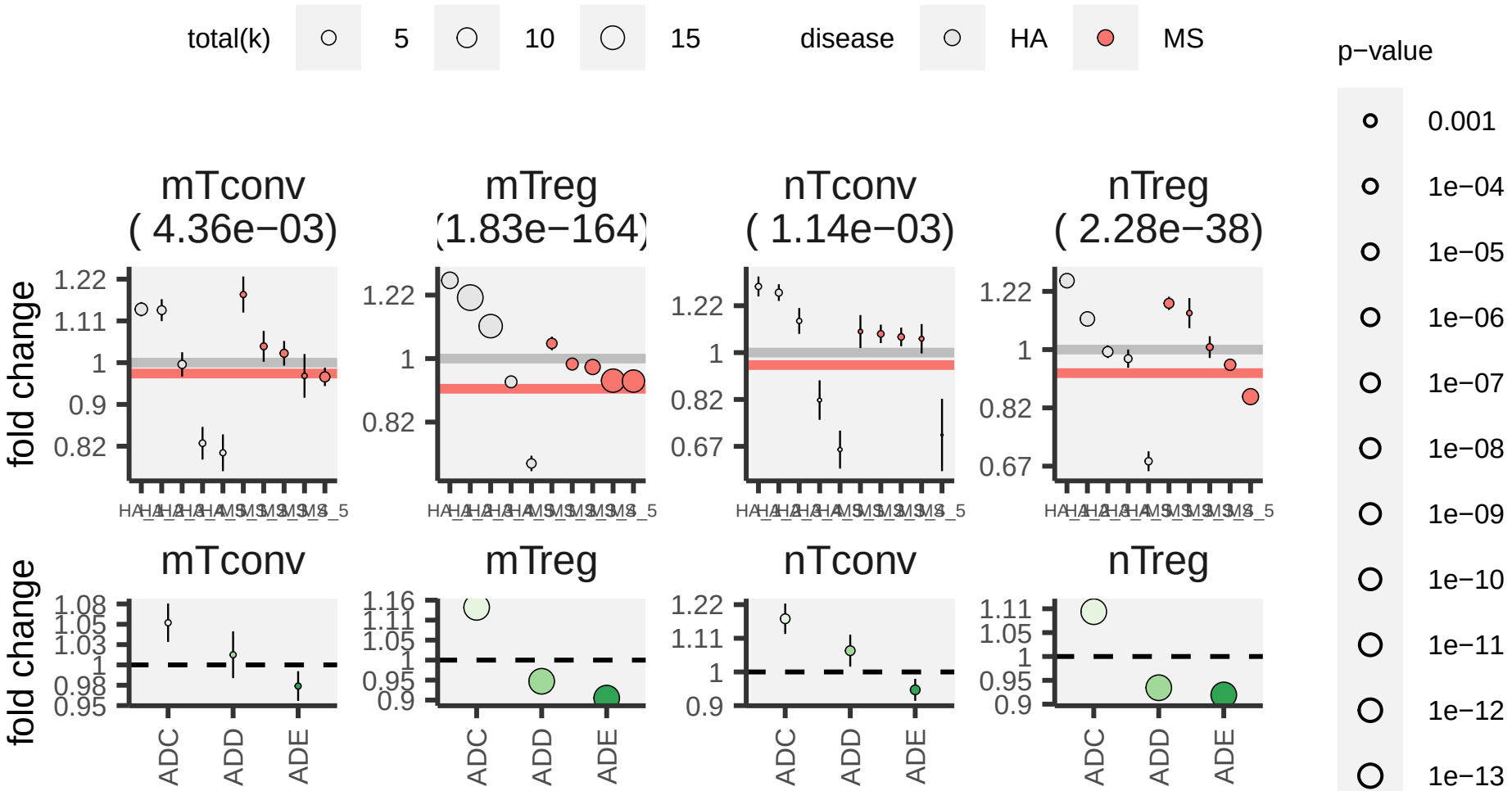
# TIGIT_TIGIT


total(k)
5
10
15
disease
HA
MS
p−value
0.001
1e−04
1e−05
1e−06
1e−07
1e−08
1e−09
1e−10
1e−11
1e−12
1e−13
1e−14
mTconv
( 4.36e−03)
mTreg
(1.83e−164)
nTconv
( 1.14e−03)
nTreg
( 2.28e−38)
fold change
ADC
ADD
ADE