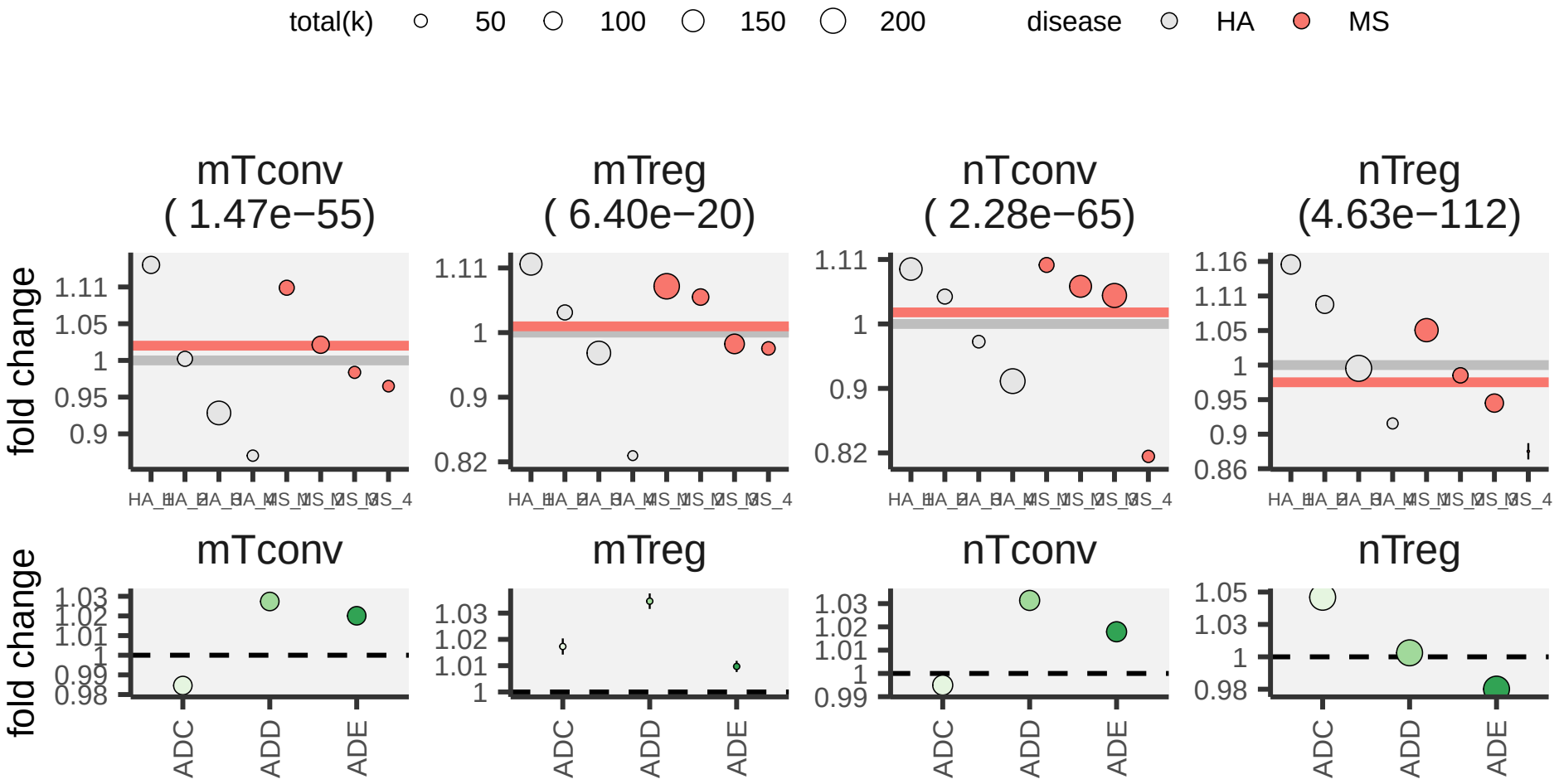
TCRab_TCRab


total(k)
50
100
150
200
disease
HA
MS
mTconv
(1.47e−55)
mTreg
(6.40e−20)
nTconv
(2.28e−65)
nTreg
(4.63e−112)
fold change
ADC
ADD
ADE