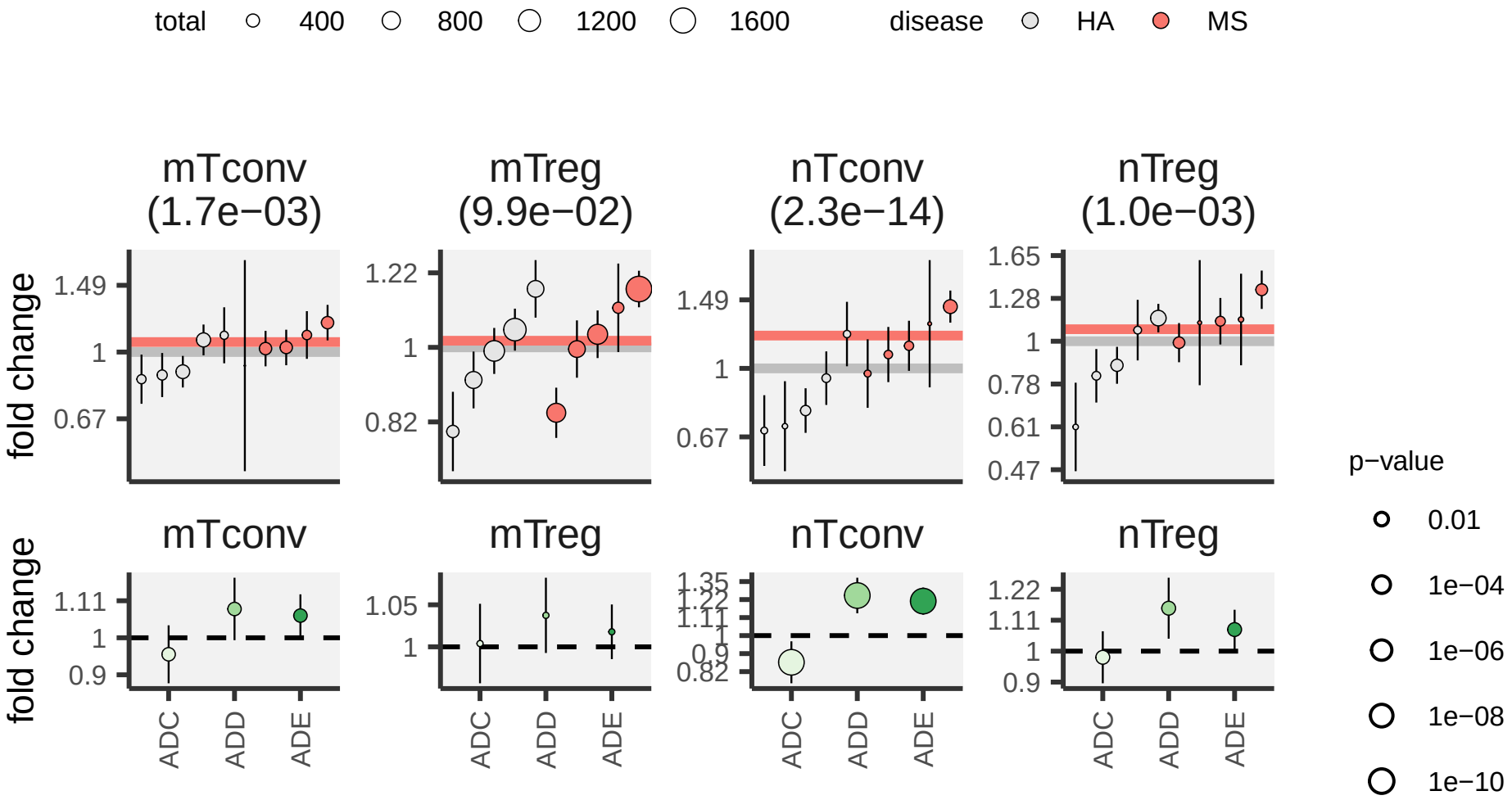
HMGB2


total
400
800
1200
1600
disease
HA
MS
mTconv
(1.7e−03)
mTreg
(9.9e−02)
nTconv
(2.3e−14)
nTreg
(1.0e−03)
fold change
1.49
1
0.67
1.22
1
0.82
1.49
1
0.67
1.65
1.28
1
0.78
0.61
0.47
mTconv
mTreg
nTconv
nTreg
fold change
1.11
1
0.9
1.05
1
1.35
1.22
1.11
1
0.9
0.82
1.22
1.11
1
0.9
ADC
ADD
ADE
p−value
0.01
1e−04
1e−06
1e−08
1e−10