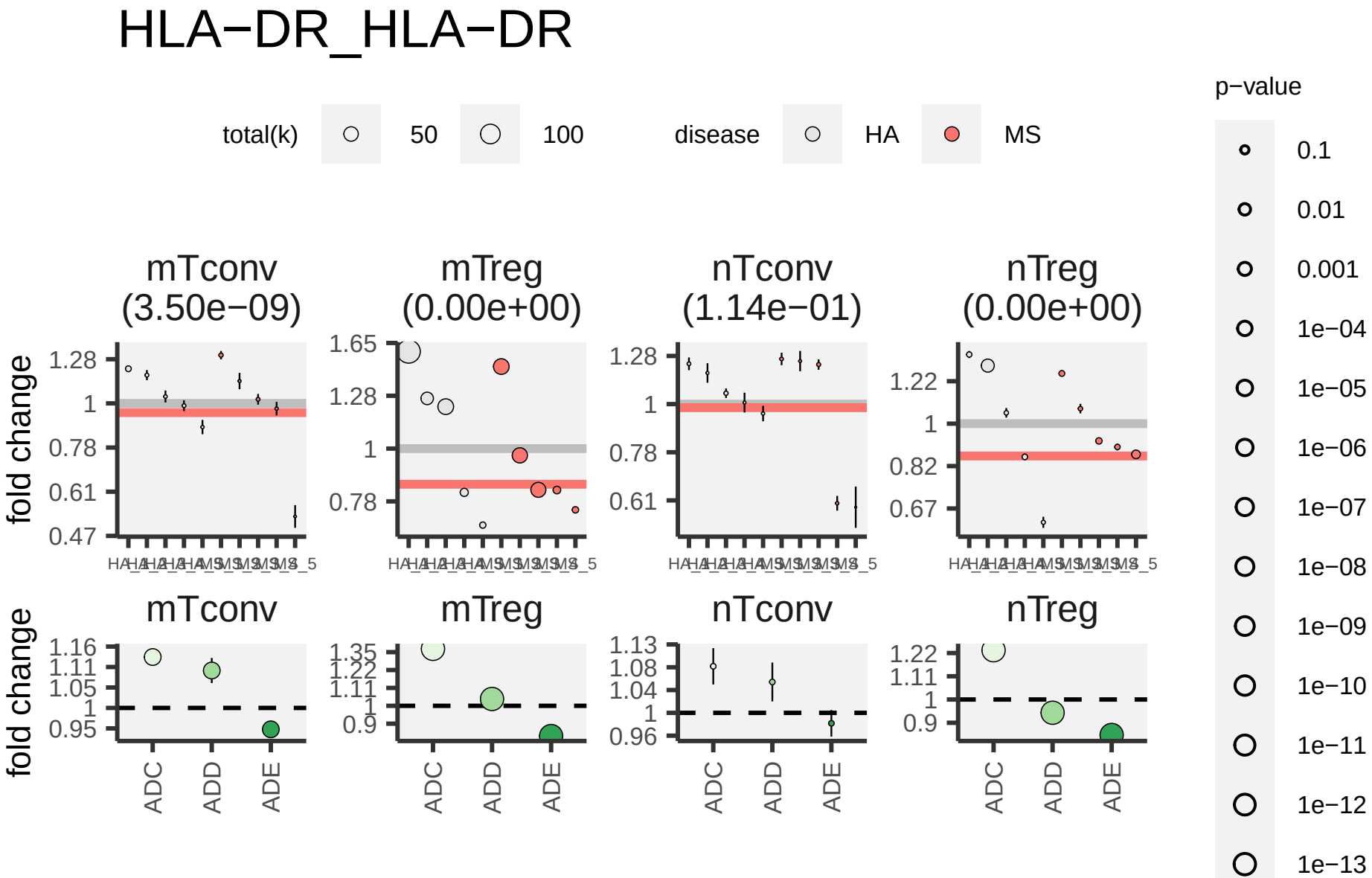
HLA−DR_HLA−DR
total(k)
50
100
disease
HA
MS
p−value
0.1
0.01
0.001
1e−04
1e−05
1e−06
1e−07
1e−08
1e−09
1e−10
1e−11
1e−12
1e−13
mTconv
(3.50e−09)
mTreg
(0.00e+00)
nTconv
(1.14e−01)
nTreg
(0.00e+00)
fold change
1.28
1
0.78
0.61
0.47
1.65
1.28
1
0.78
1.28
1
0.78
0.61
1.22
1
0.82
0.67
HA_1
HA_2
HA_3
HA_4
MS_1
MS_2
MS_3
MS_4
MS_5
mTconv
mTreg
nTconv
nTreg
fold change
1.16
1.11
1.05
1
0.95
1.35
1.22
1.11
1
0.9
1.13
1.08
1.04
1
0.96
1.22
1.11
1
0.9
ADC
ADD
ADE