

ENSG00000075624_ACTB

total(k)



20



40



60



80

disease



HA



MS

p-value



0.01



0.001



1e-04



1e-05



1e-06



1e-07



1e-08



1e-09



1e-10



1e-11



1e-12

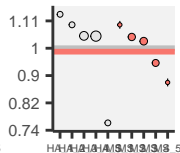
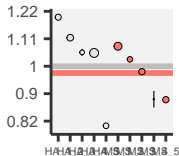
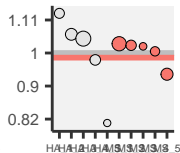
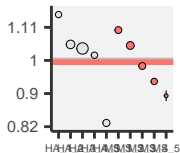
mTconv
(2.67e-02)

mTreg
(1.35e-19)

nTconv
(1.84e-17)

nTreg
(1.34e-06)

fold change



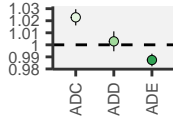
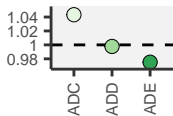
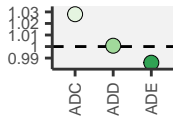
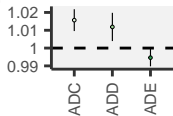
fold change

mTconv

mTreg

nTconv

nTreg



ADC

ADD

ADE

ADC

ADD

ADE

ADC

ADD

ADE

ADC

ADD

ADE