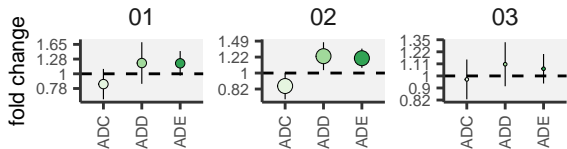
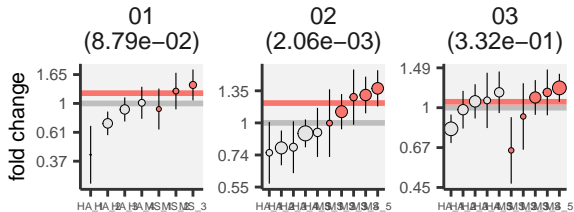


ENSG00000163599_CTLA4

HA
 MS
 total(k)
 0.01
 0.02
 0.03
 0.04
 0.05



p-value

0.01