## ENSG00000150593\_PDCD4

total(k)

0.2

0.3

disease

HA

MS

01 02 03 (8.41e-01)(2.49e-03)(1.86e-01)1.28 fold change 1.22 0.78 0.78 0.82 0.61 0.61 02 01 03 fold change p-value 0.01 0.9 0.95