## ENSG00000107738\_VSIR

total(k)

0.2

0.3

disease

HA

MS

01 02 03 (1.81e-01)(6.76e-04)(9.54e-02)1.22 fold change 1.22 0.82 0.82 0.82 0.67 02 03 fold change p-value 0.01 0.95