ENSG00000107738_VSIR

total(k)

0.2

0.3

disease

HA

MS

01 02 03 (4.22e-04)(6.69e-02)(2.73e-01)1.22 1.28 fold change 1.22 0.9 0.82 0.78 0.82 0.67 02 03 fold change p-value 0.01 0.95