## ENSG00000107738\_VSIR

total(k)

0.2

disease

HA

MS

01 02 03 (4.50e-03) (8.20e-02)(4.63e - 02)1.22 -1.28 fold change 0.78 0.82 0.61 01 02 03 fold change p-value 0.01 4DE