ENSG00000143106_PSMA5

total(k)

0.2

0.3

disease

HA

MS

01 02 03 (4.35e-02)(3.36e-04)(8.81e-01)1.22 fold change 1.22 1.22 0.9 0.82 0.82 0.82 0.67 0.67 0.74 02 03 01 fold change p-value 0.01 0.95