ENSG00000164442_CITED2

total(k)

0.2

0.3

disease

HA

MS

01 02 03 (2.16e-02)(2.86e-02)(4.01e-04)fold change 1.22 1.22 1.22 0.82 0.82 0.82 0.74 0.67 02 03 01 fold change p-value 0.01