## ENSG00000051108\_HERPUD1

total(k)

disease

HA

MS

mTconv mTreg nTconv nTreg (5.83e-19)(2.44e-24)(6.09e-08)(6.03e-10)1.49 fold change 1.22 -1.22 1.35 1.22 0.82 0.82 0.74 0.82 0.67 mTconv nTconv mTreg nTreg fold change p-value 1.11 1e-08 0.9 0.9 1e-10 ADD ADE ADD ADE ADD

0.25

0.50

0.75

1.00