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disease HA MS total(k) 0.25 0.50 0.75 1.00 mTconv mTreg nTconv nTreg (4.31e-16)(1.94e-01)(3.33e-18)(6.21e-01)1.49 fold change 1.11 1.22 1.22 1.22 400 0.82 0.9 0.82 0.82 0.67 0.82 p-value 0 0.01 mTconv nTconv mTreg nTreg fold change 1e-04 1.04 0.98 1e-06 0.96 0.9 1e-08 ADD ADD ADE 1e-10