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disease HA MS total(k) 0.2 0.4 01 02 03 (1.79e-13)(5.61e-02)(1)1.65 1.35 fold change 1.65 0.61 0.74 0.37 0.61 0.55 0.22 p-value HA_1HA_2HA_3HA_4MS_1 0.01 02 03 01 fold change 1.05 1.03 1e-04 1.22 1e-06 0.98 0.95 0.82 1e-08 ADD ADE ADE ADD 1e-10