



FAM115A 7.9e-02 CTAGE4 1.0e-02 ARHGEF35 2.8e-06 CTAGE8 5.1e-02 ARHGEF5 1.1e-01 TPK1 4.8e-03 CNTNAP2 7.9e-03 CUL1 3.4e-01 EZH2 2.8e-01 ZNF786 2.2e-01

