reconstructed (row-normalized) genotype genotype genotype
Cluster
Hist1h2ap
Hist1h2ad
Hist1h2ad
Hist1h2ad
Hist1h2ad
Hist1h2ad
Hist1h2ad
Hist1h2ad
Global
Global
Malat1
Unc5b
Co118a1
Glul
Glad
Glul
Glad
Glad
Sema3g
Epas1
Fo62d1
Fo62d1 Control cluster Mutant 2 0 ⁻²cluster 2 3 4 5 6 7 Hist1h2ah Rrm2 Tk1 Aurkb Hist1h2ab HIST IN 280
HIST I