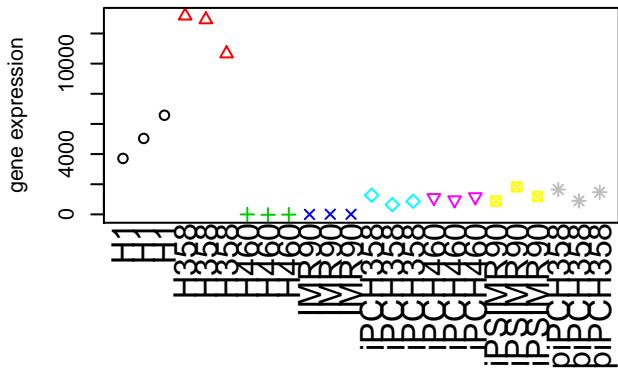
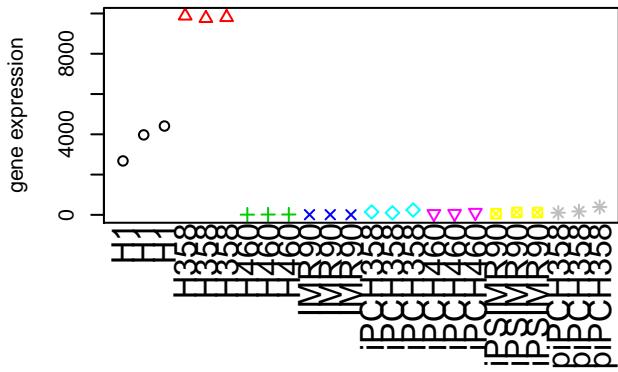


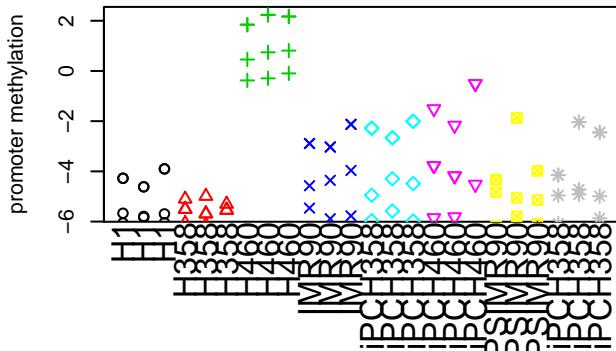
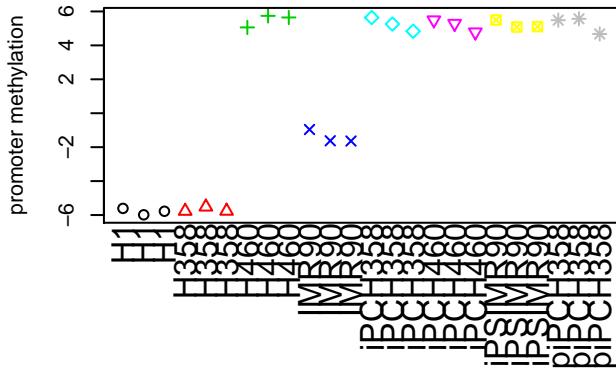
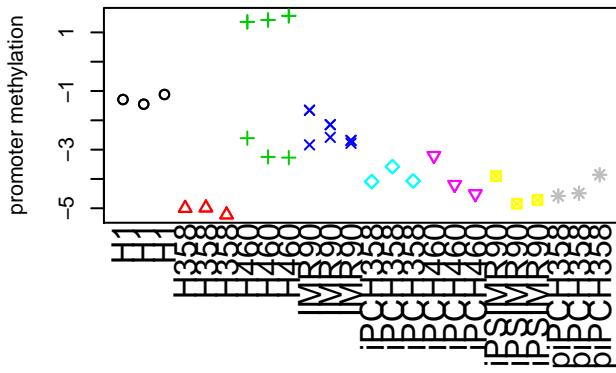
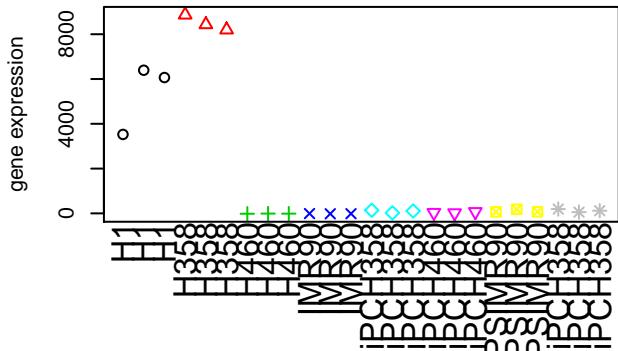
NM_021102.2 SPINT2
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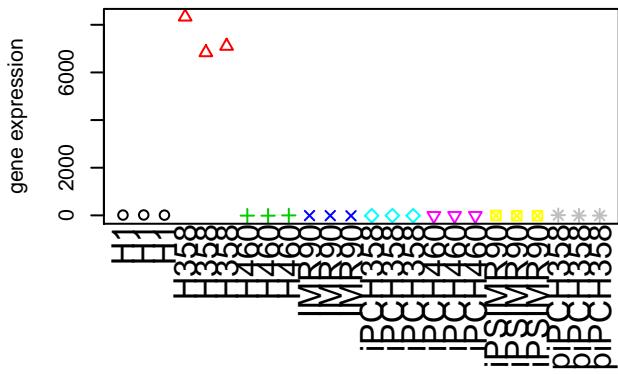
NM_002354.1 TACSTD1
COR= -7.79e-01 P= 3.71e-06



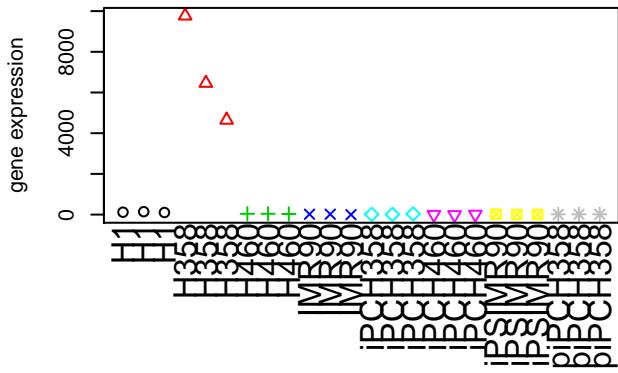
NM_004360.2 CDH1
COR= -4.19e-01 P= 2.07e-02



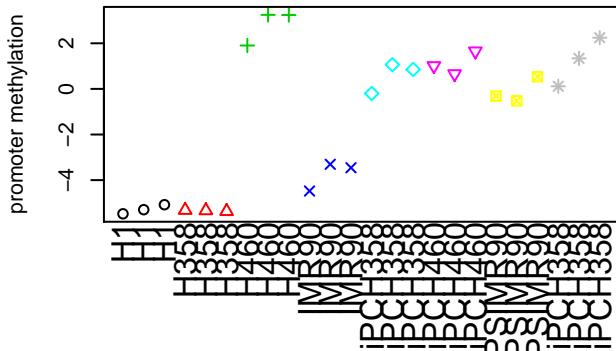
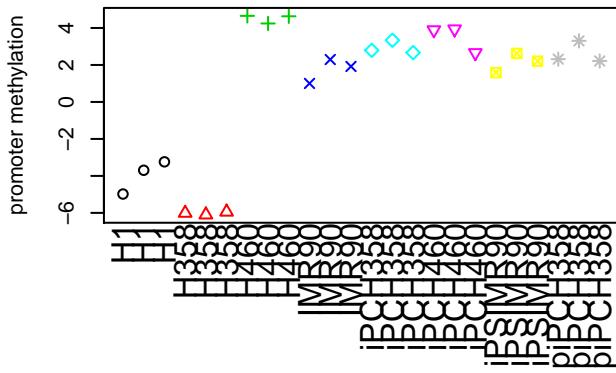
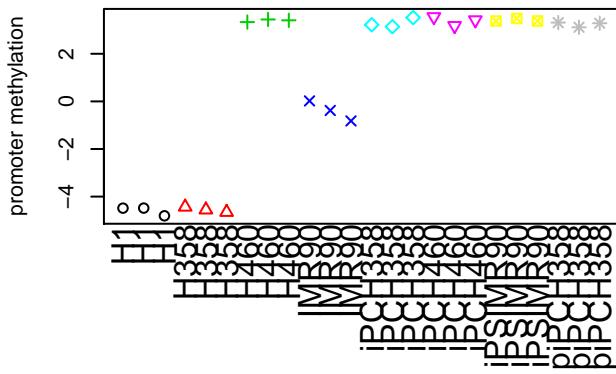
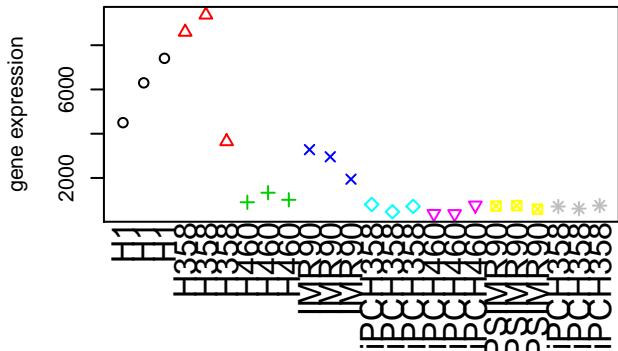
NM_002353.1 TACSTD2
COR= -6.07e-01 P= 8.31e-04



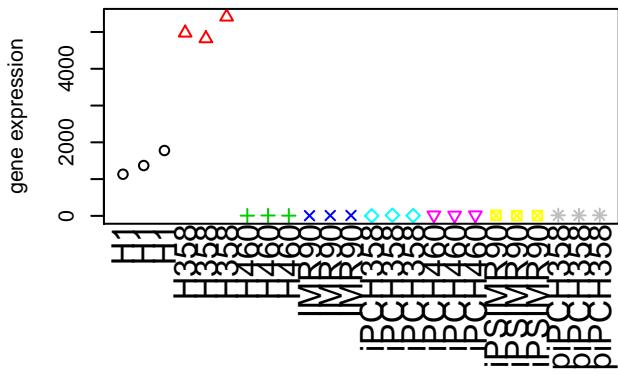
NM_005562.1 LAMC2
COR= -7.07e-01 P= 5.59e-05



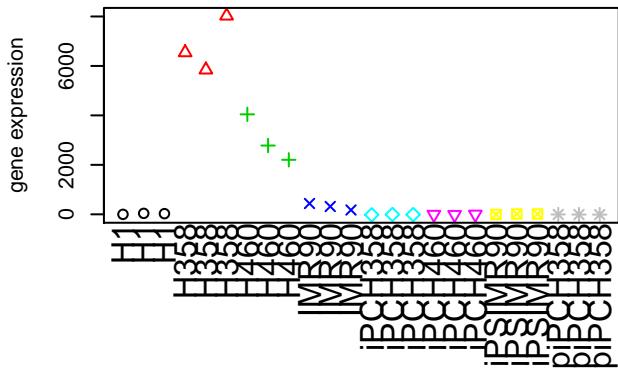
NM_145899.1 HMGA1
COR= -8.23e-01 P= 4.03e-07



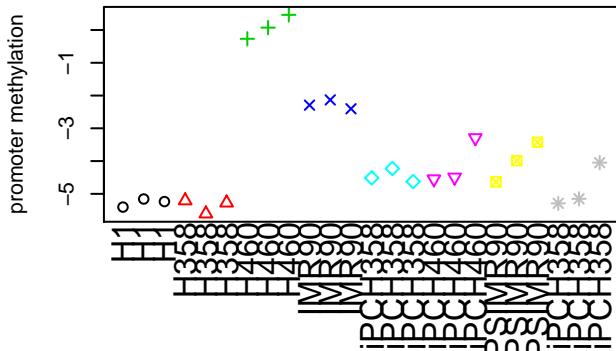
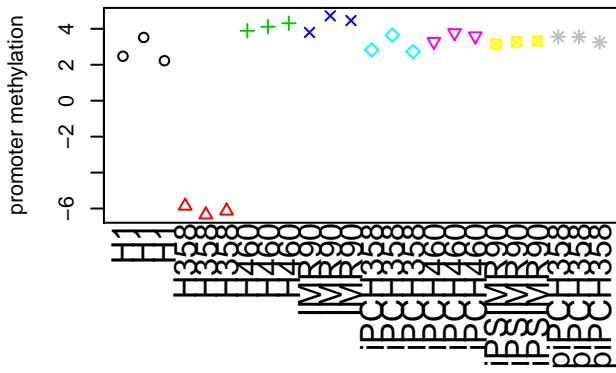
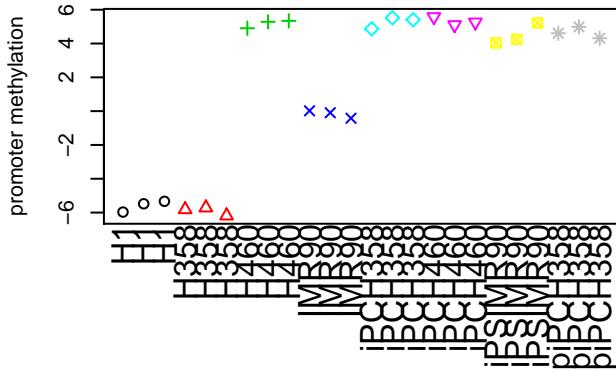
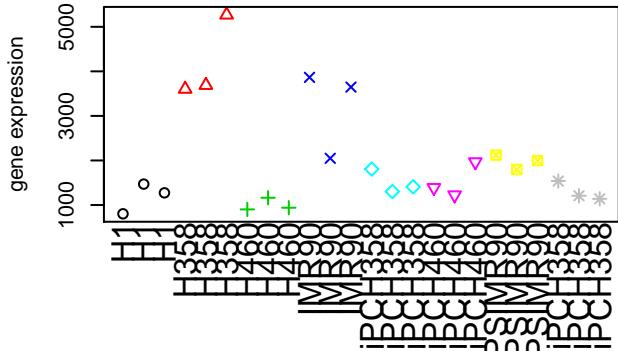
NM_005558.3 LAD1
COR= -7.85e-01 P= 2.76e-06



NM_201525.1 GPR56
COR= -8.45e-01 P= 1.00e-07

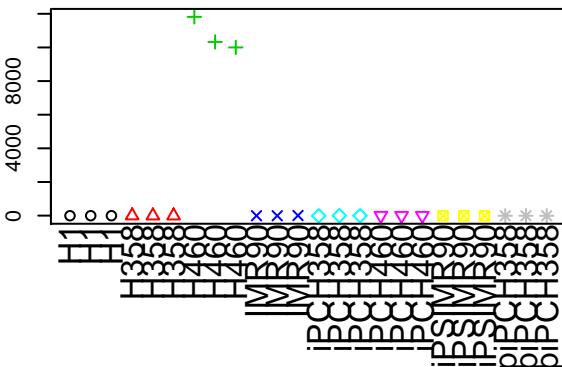


NM_004566.2 PFKFB3
COR= -1.89e-01 P= 1.88e-01

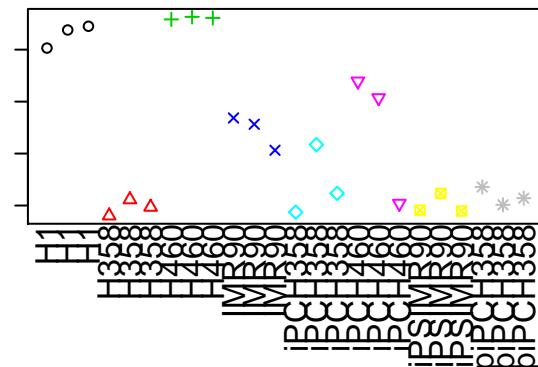


NM_005218.3 DEFB1
COR= 6.08e-01 P= 8.18e-04

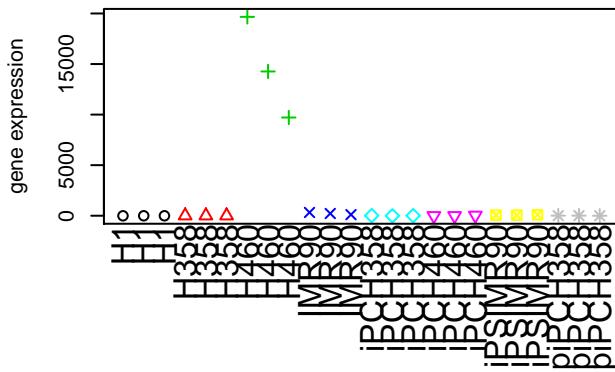
gene expression



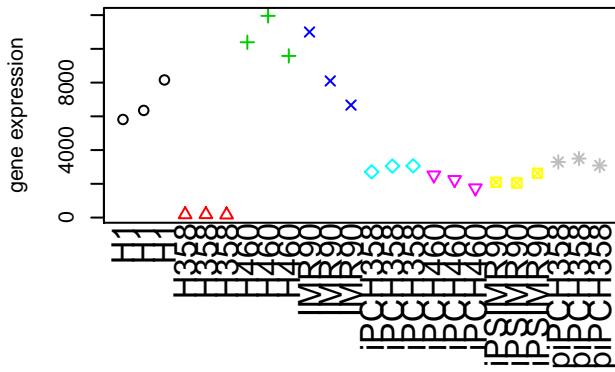
promoter methylation



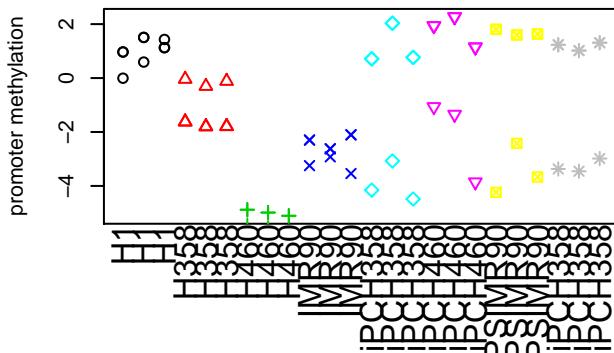
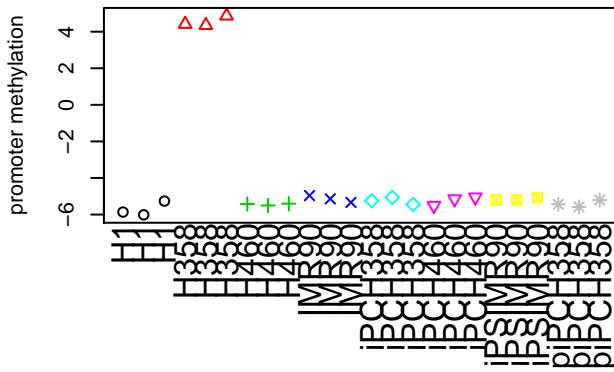
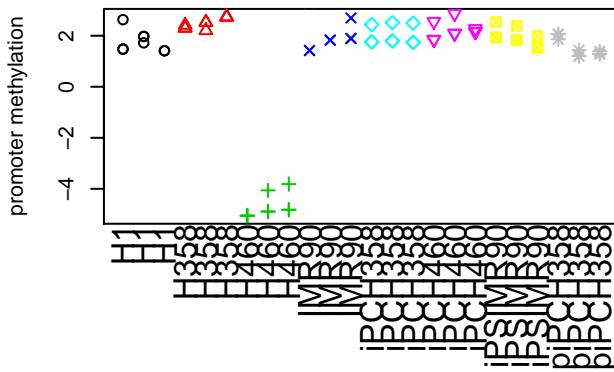
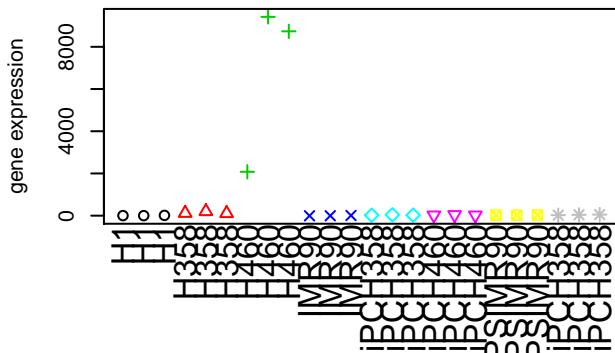
NM_002727.2 SRGN
COR= -9.49e-01 P= 7.88e-13



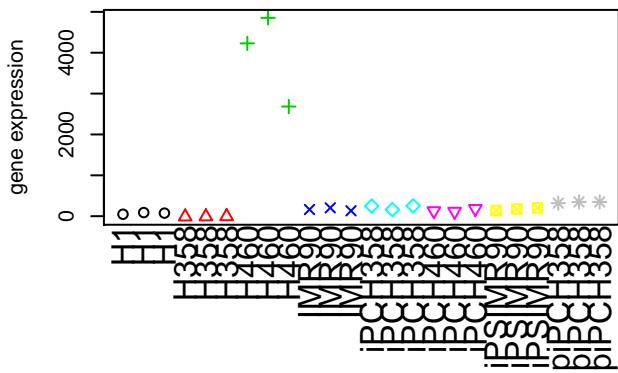
NM_004181.3 UCHL1
COR= -4.87e-01 P= 7.86e-03



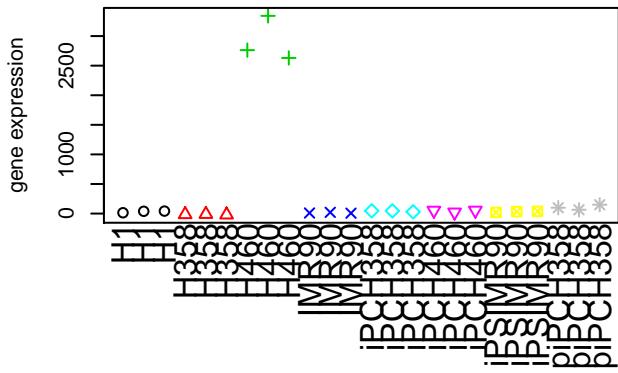
NM_000691.3 ALDH3A1
COR= -5.88e-01 P= 1.25e-03



NM_012307.2 EPB41L3
COR= -2.39e-01 P= 1.30e-01



NM_021136.2 RTN1
COR= -3.62e-01 P= 4.10e-02



NM_005559.2 LAMA1
COR= -1.92e-01 P= 1.84e-01

