ERG(SX,TH)
methylation
1.0
0.8
0.6
0.4
0.2
0
AD AO BL EG FT GA LG LI LV OV PA PO RA RV SB SG SX TH
P1 P2 P3 P4 P5 P6

ERG,P3
CpGnumber
140
120
100
80
60
40
20
0
p1 p2 p3 p4 p5 p6

FAM65B(TH)
methylation
1.0
0.8
0.6
0.4
0.2
0
AD AO BL EG FT GA LG LI LV OV PA PO RA RV SB SG SX TH
P1 P2 P3 P4 P5 P6

FAM65B,P6
CpGnumber
80
70
60
50
40
30
20
10
0
p1 p2 p3 p4 p5 p6

TGFB3(AO)
methylation
1.0
0.8
0.6
0.4
0.2
0
AD AO BL EG FT GA LG LI LV OV PA PO RA RV SB SG SX TH
P1 P2 P3 P4 P5

TGFB3,P4
CpGnumber
60
50
40
30
20
10
0
P1 P2 P3 P4 P5