A
0
1
2
FPM
500
400
300
200
100
0
simulation
Expressed transcripts
ENST00000484745 >
ENST00000440132 >
txrevise promoters
ENST00000484745 >
ENST00000440132 >
txrevise 3′ ends
ENST00000484745 >
ENST00000440132 >
500
1000
1500
2000
2500
Distance from region start (bp)

B
ENST00000484745 promoter usage
1.00
0.75
0.50
0.25
0.00
0
1
2
genotype

C
ENST00000484745 3'end usage
1.00
0.75
0.50
0.25
0.00
0
1
2
genotype