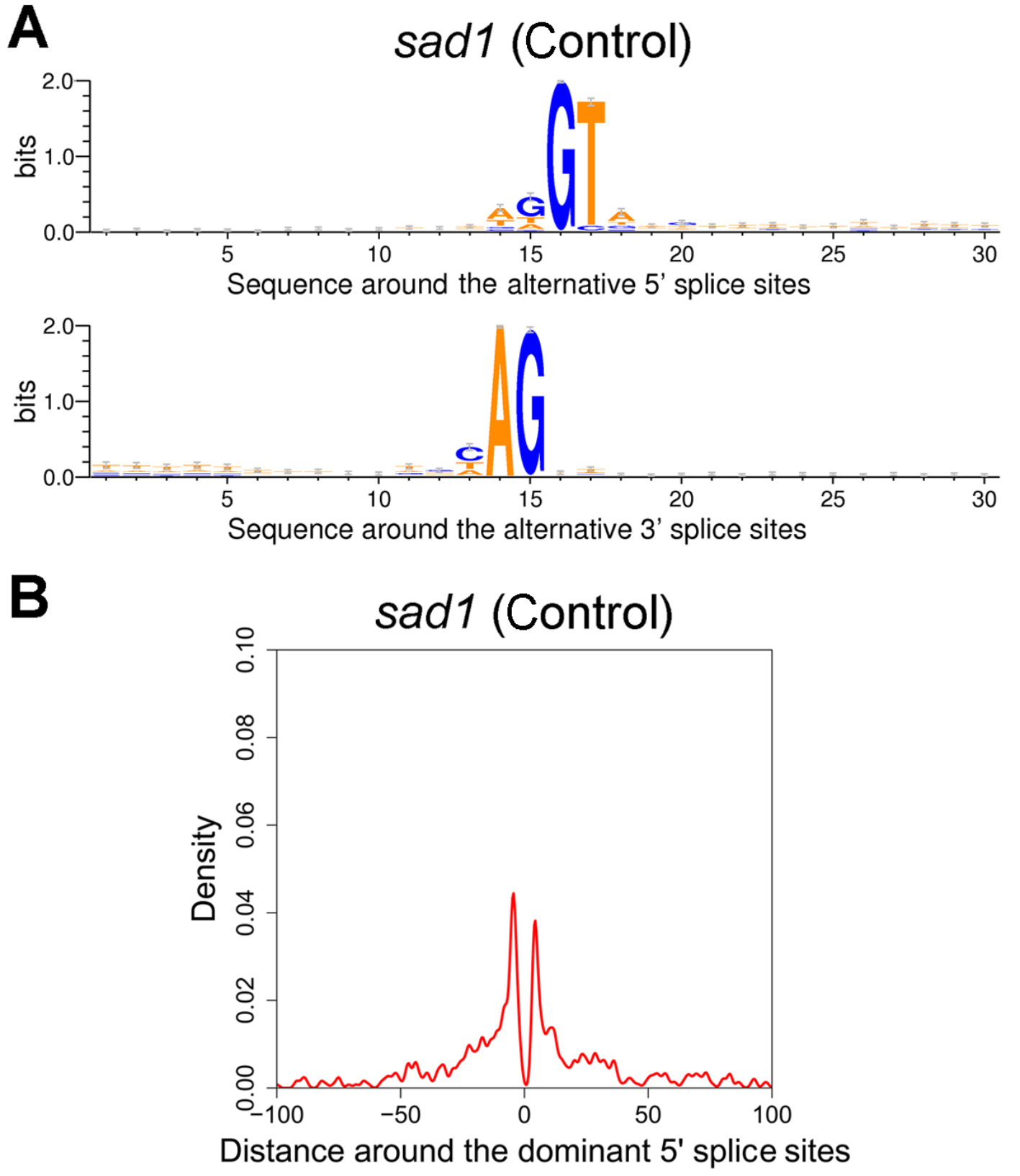
A
sad1 (Control)
bits
2.0
1.0
0.0
5
10
15
20
25
30
Sequence around the alternative 5' splice sites
bits
2.0
1.0
0.0
5
10
15
20
25
30
Sequence around the alternative 3' splice sites
B
sad1 (Control)
Density
0.00
0.02
0.04
0.06
0.08
0.10
−100
−50
0
50
100
Distance around the dominant 5' splice sites