A
Repeat-anti-repeat hybrid
Length of the longest helix
30
20
10
0.000
0.100
0.200
0.300
0.400
0.500
Prop. non-pairing nucleotides
B
SVM Cross-validation: AUC = 0.9901
Sensitivity
1.000
0.750
0.500
0.250
0.000
0.000
0.250
0.500
0.750
1.000
1-Specificity
C
SVM true-positive probability
Frequency
20
10
0
0.965
0.970
0.975
0.980
0.985
0.990
Pr(true positive)