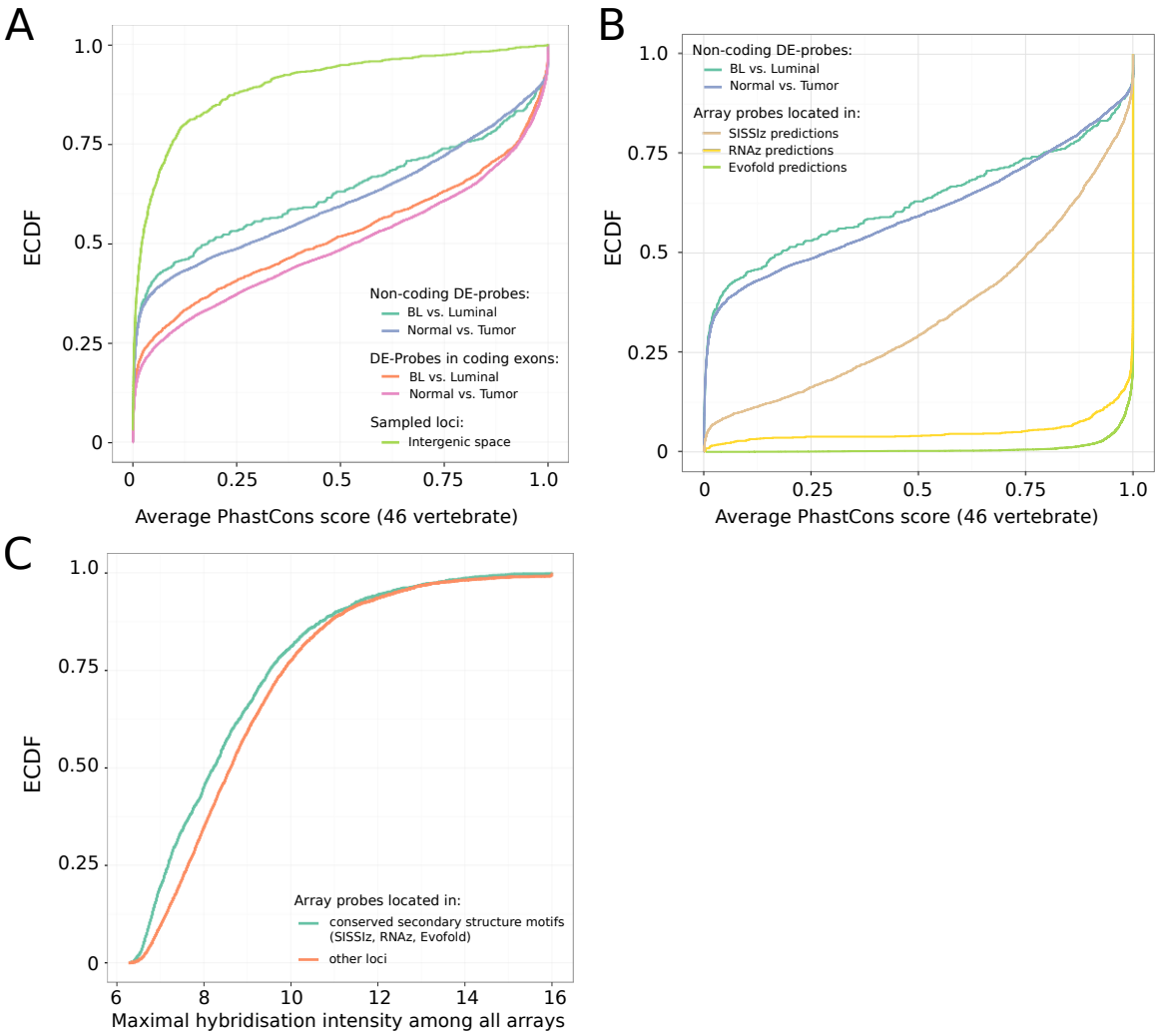
A
ECDF
1.0
0.75
0.5
0.25
0
0
0.25
0.5
0.75
1.0
Average PhastCons score (46 vertebrate)
Non-coding DE-probes:
BL vs. Luminal
Normal vs. Tumor
DE-Probes in coding exons:
BL vs. Luminal
Normal vs. Tumor
Sampled loci:
Intergenic space
B
ECDF
Non-coding DE-probes:
BL vs. Luminal
Normal vs. Tumor
Array probes located in:
SISSIz predictions
RNAz predictions
Evofold predictions
Average PhastCons score (46 vertebrate)
C
ECDF
1.0
0.75
0.50
0.25
0
6
8
10
12
14
16
Maximal hybridisation intensity among all arrays
Array probes located in:
conserved secondary structure motifs
(SISSIz, RNAz, Evofold)
other loci