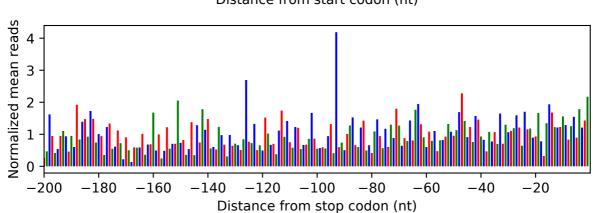
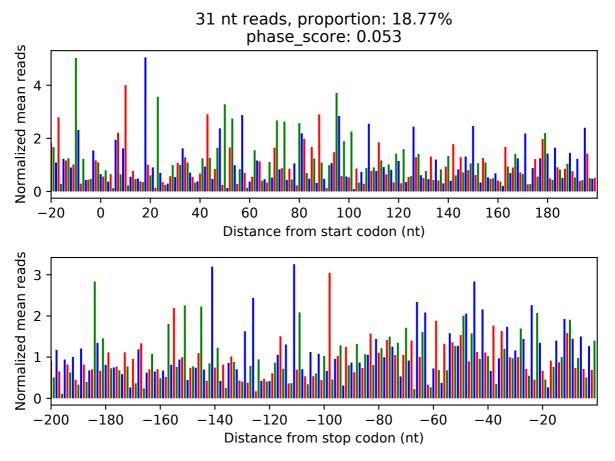


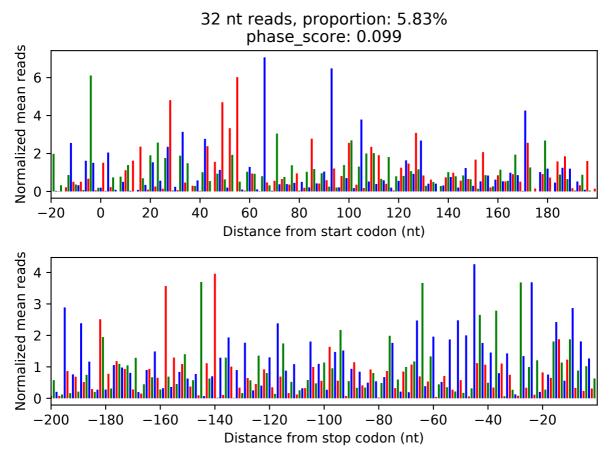
30 nt reads, proportion: 35.53% phase score: 0.066 Distance from start codon (nt)

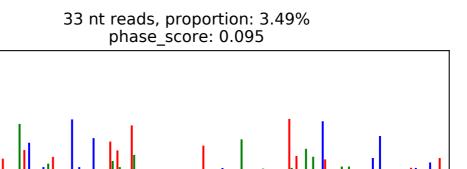


Normalized mean reads

-20







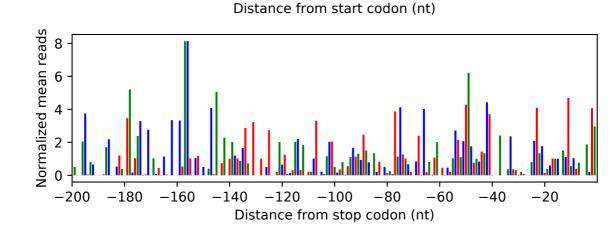
120

100

140

160

180



80

Normalized mean reads

10.0

7.5

5.0

2.5

0.0

-20

0

20

40

60

