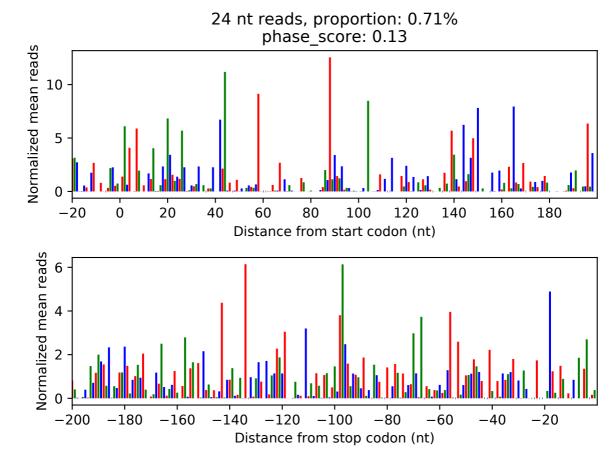


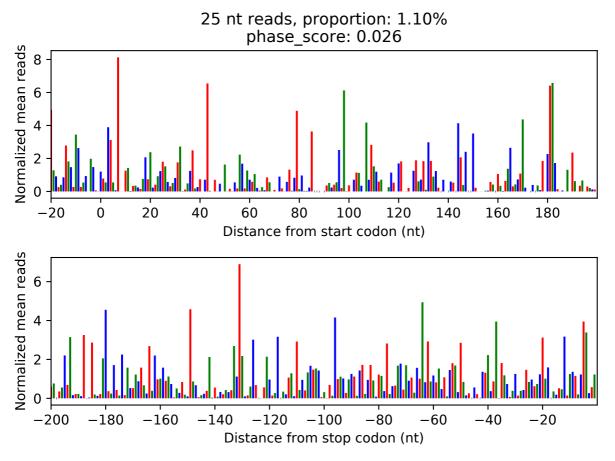
Normalized mean reads

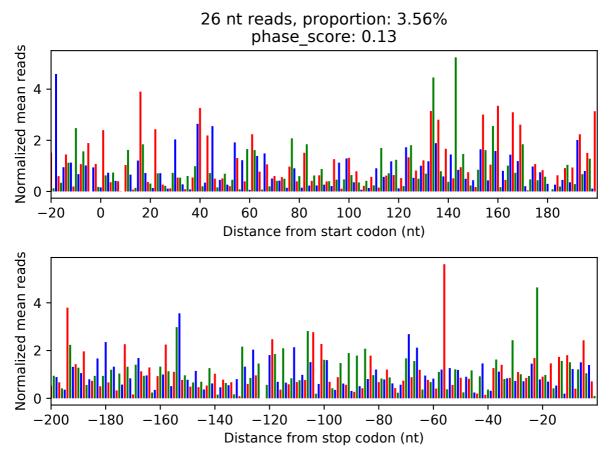
6

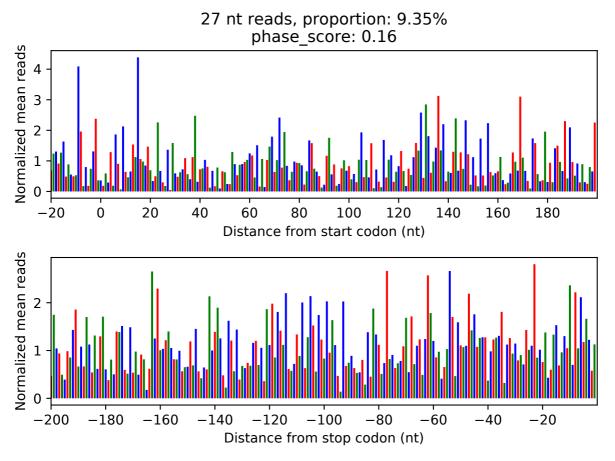
2

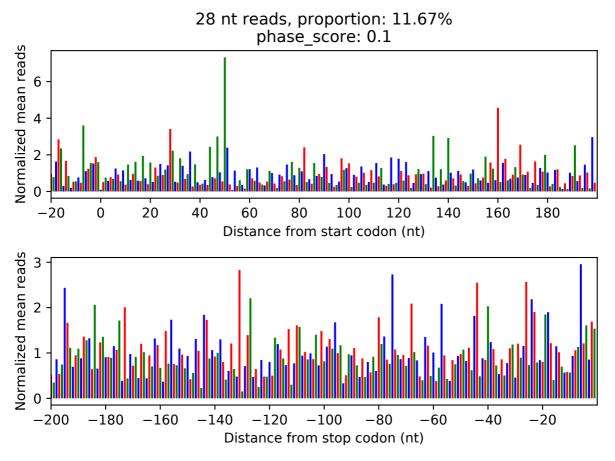
-20

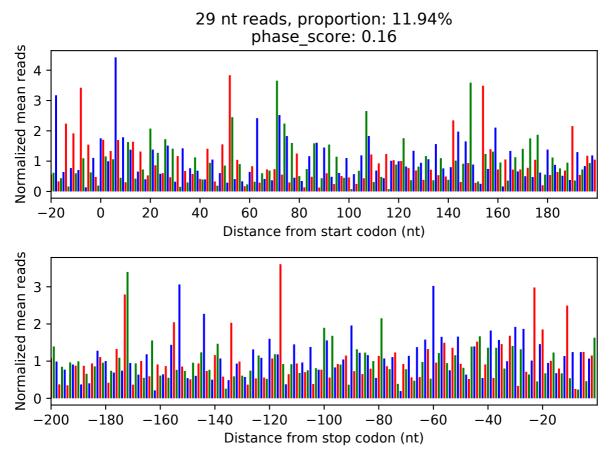






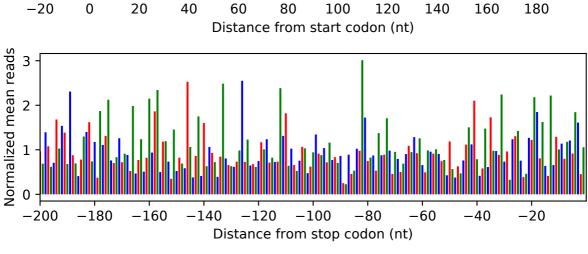


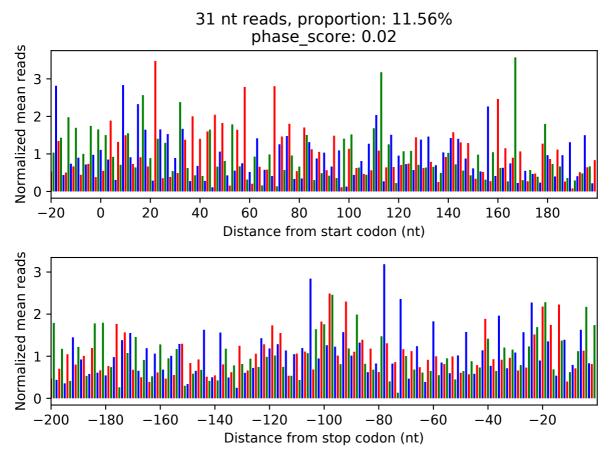


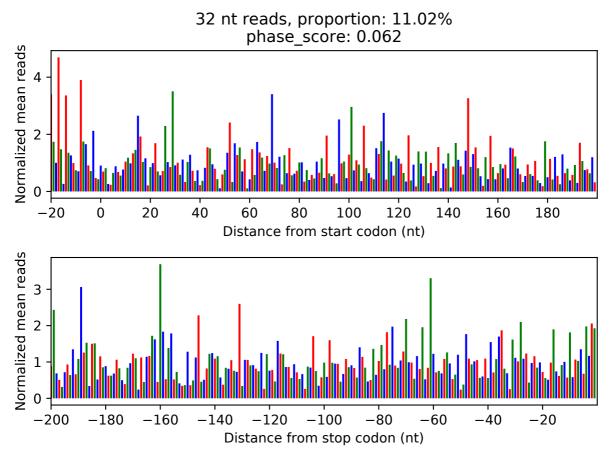


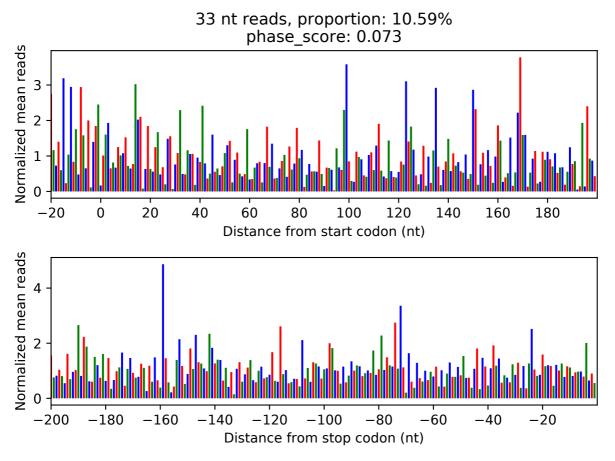
30 nt reads, proportion: 11.75% phase score: 0.11 Distance from start codon (nt)

Normalized mean reads









34 nt reads, proportion: 8.75% phase_score: 0.12

