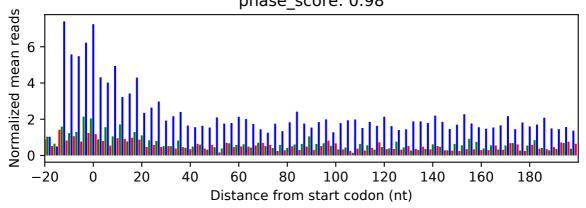
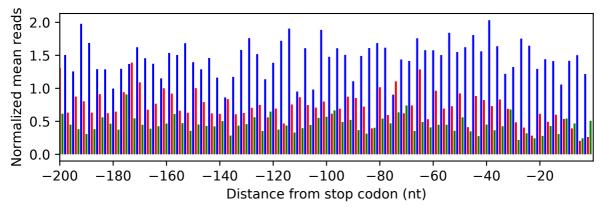
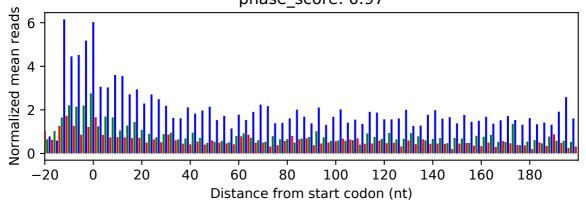
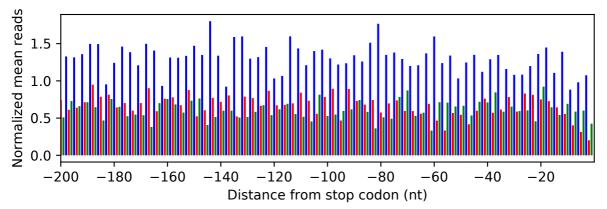
26 nt reads, proportion: 0.95% phase score: 0.98



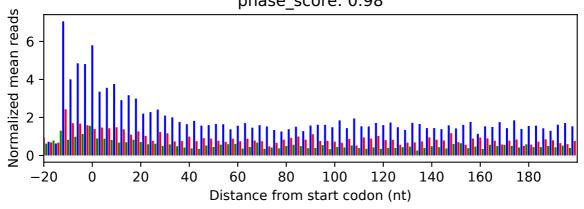


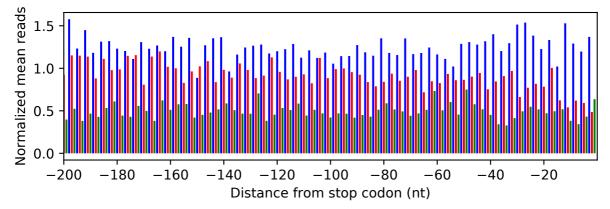
27 nt reads, proportion: 2.29% phase score: 0.97



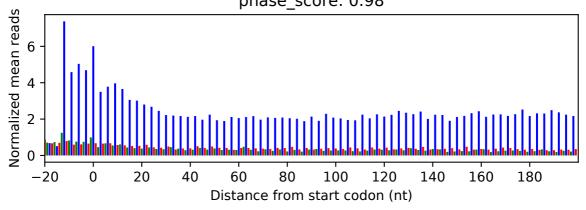


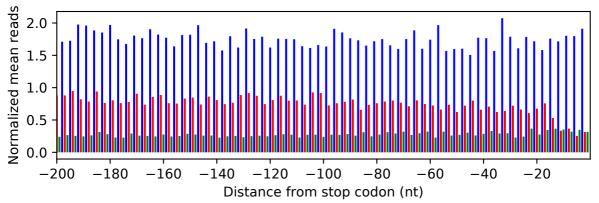
28 nt reads, proportion: 4.80% phase score: 0.98



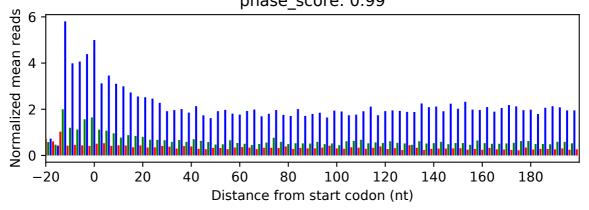


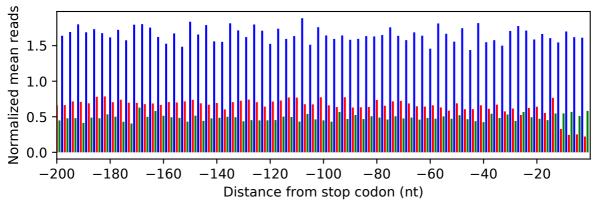
29 nt reads, proportion: 20.39% phase score: 0.98



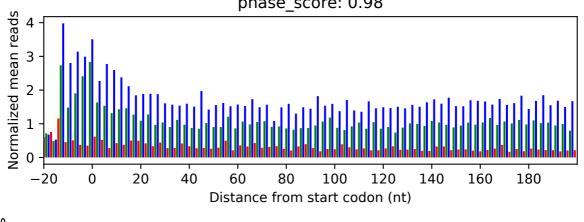


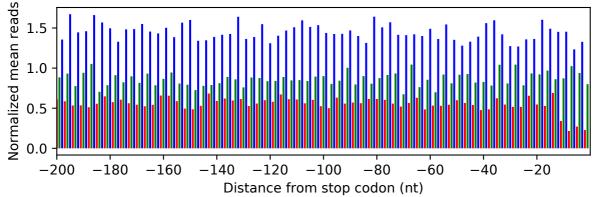
30 nt reads, proportion: 24.66% phase score: 0.99



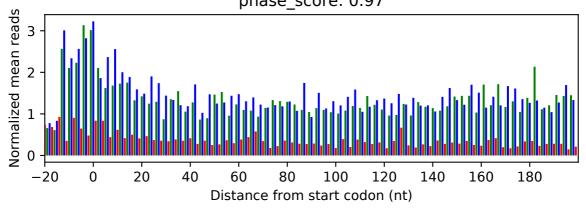


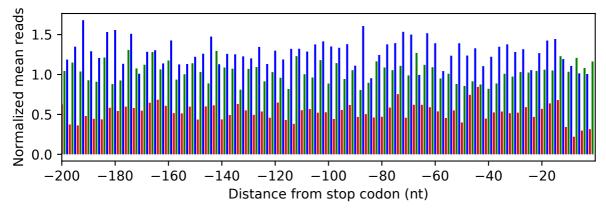
31 nt reads, proportion: 11.20% phase score: 0.98



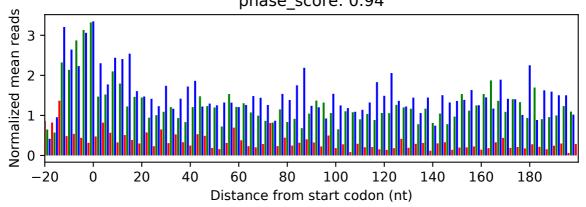


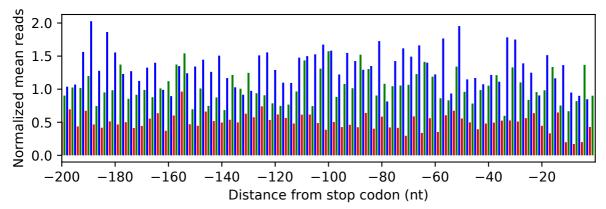
32 nt reads, proportion: 10.20% phase score: 0.97



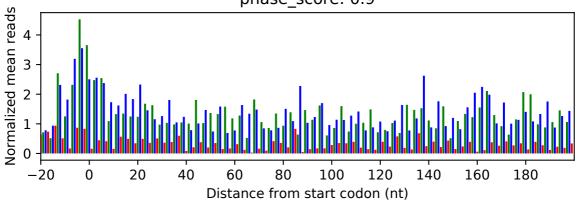


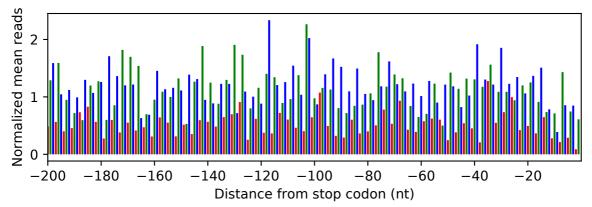
33 nt reads, proportion: 7.82% phase score: 0.94



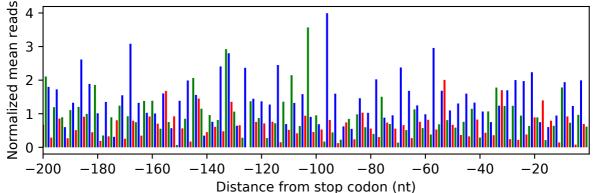


34 nt reads, proportion: 3.31% phase score: 0.9





35 nt reads, proportion: 2.67% phase score: 0.76 Distance from start codon (nt)



Normalized mean reads

-20

