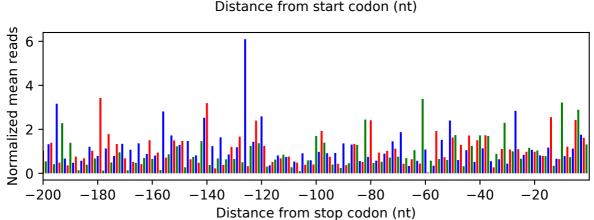


29 nt reads, proportion: 20.39% phase score: 0.13 Distance from start codon (nt)



Normalized mean reads

-20

