































36 nt reads, proportion: 32.88% phase score: 0.097 Normalized mean reads 15 10 5 20 -2040 60 80 100 120 160 180 0 140 Distance from start codon (nt) Normalized mean reads 15 10 5 0 -200-180-160-120-80-60 -20-140-100-40Distance from stop codon (nt)

