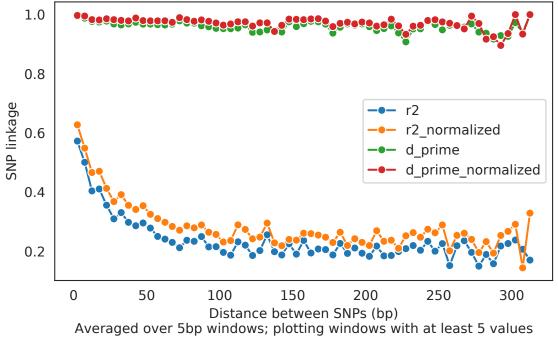
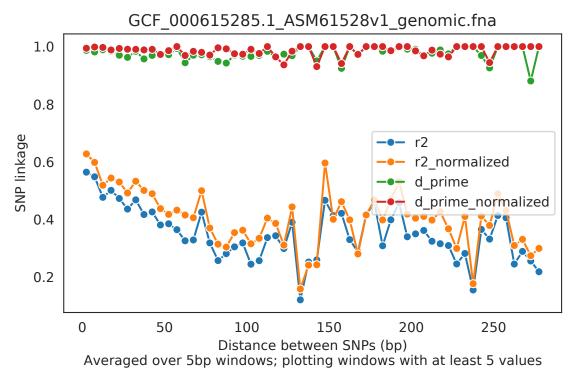
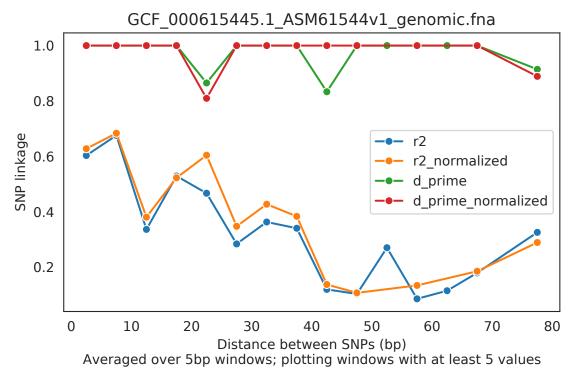
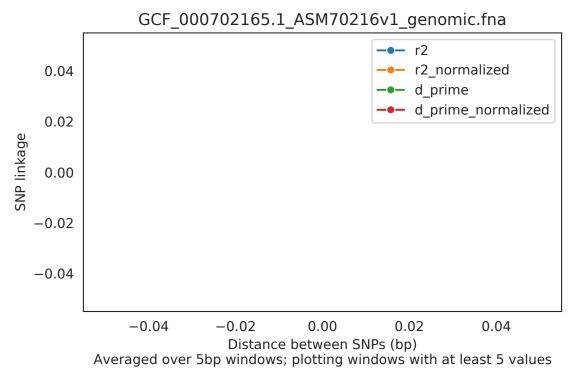


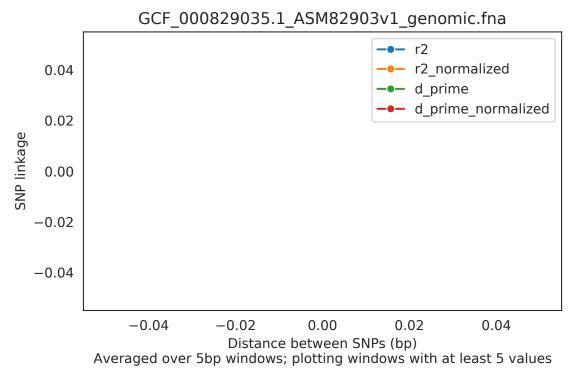
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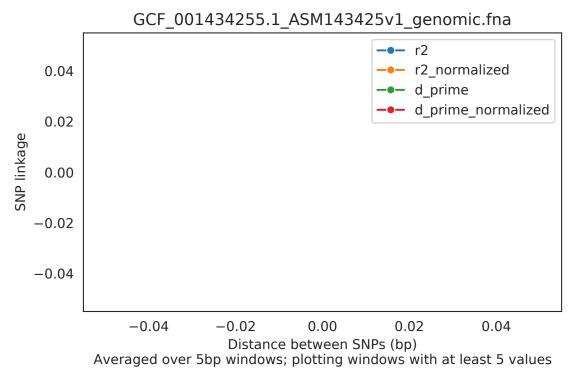


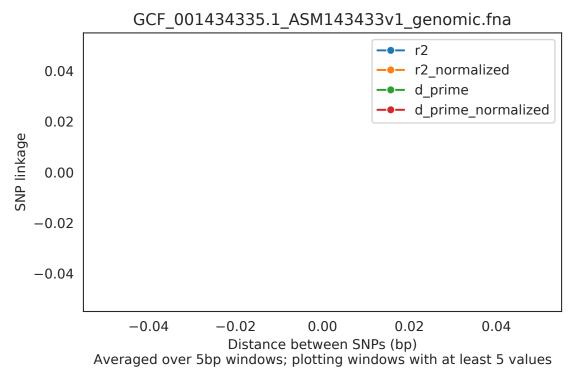


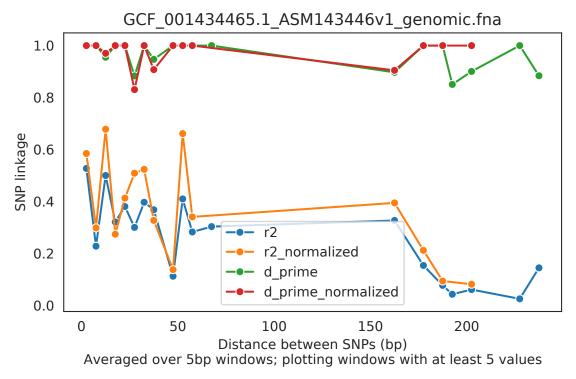


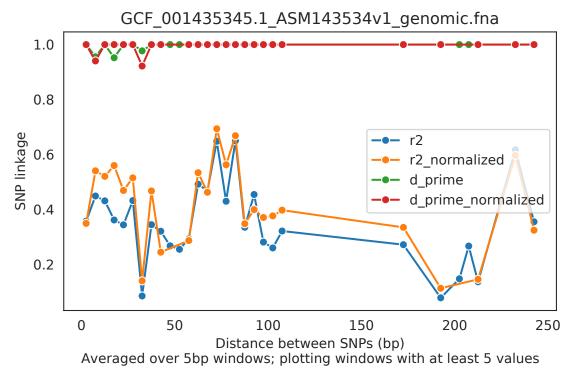


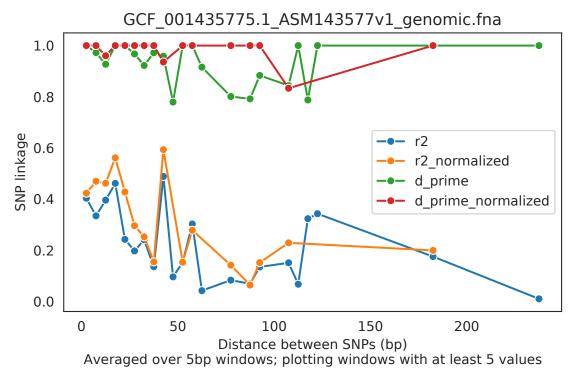
GCF 001433875.1 ASM143387v1 genomic.fna 1.0 0.8 **SNP** linkage 0.6 0.4 r2\_normalized d prime 0.2 d prime normalized 50 100 150 200 250 300 Distance between SNPs (bp) Averaged over 5bp windows; plotting windows with at least 5 values

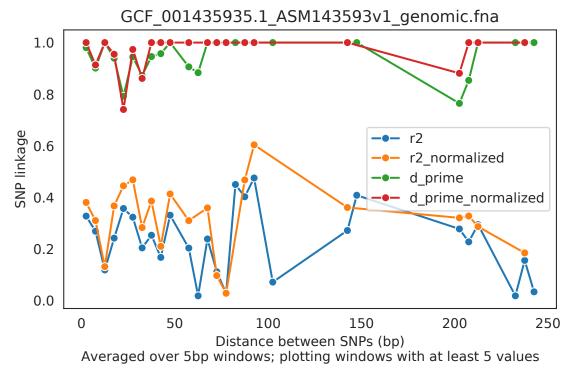


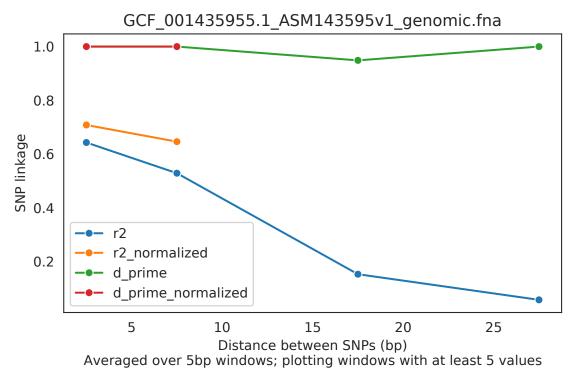


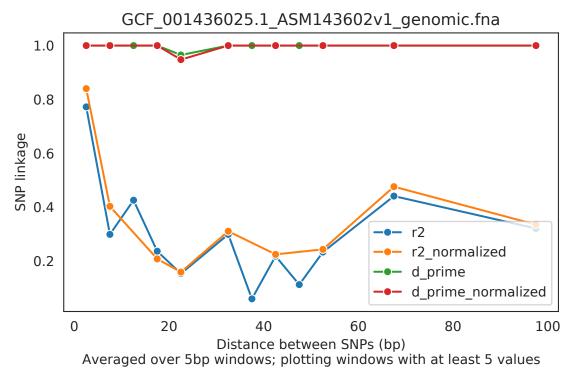


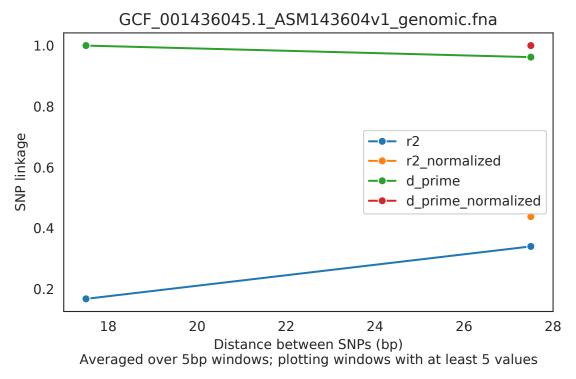


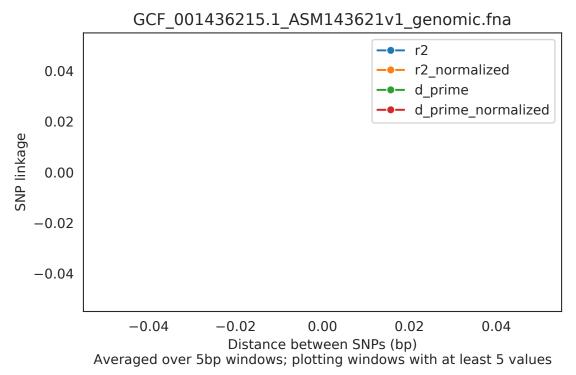


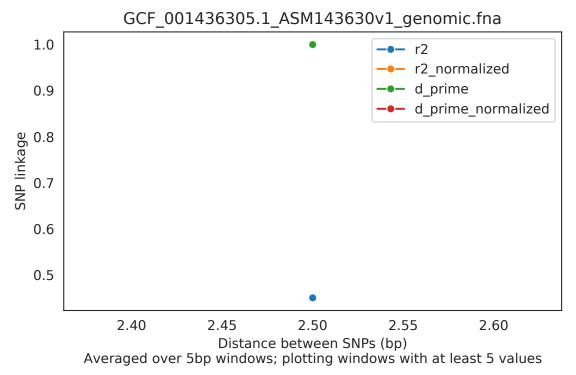












GCF 001436315.1 ASM143631v1 genomic.fna r2\_normalized d\_prime\_normalized 50 100 150 200 250 300 350

1.0

0.9

0.8

0.6

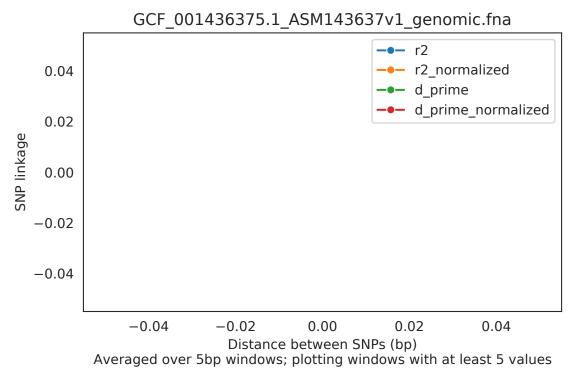
0.5

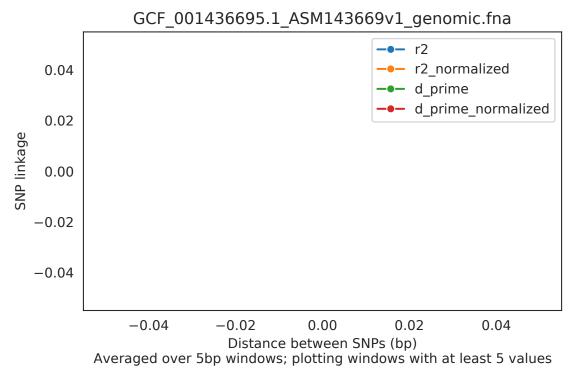
0.4

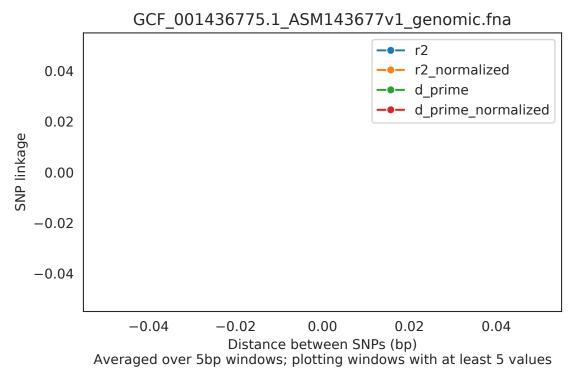
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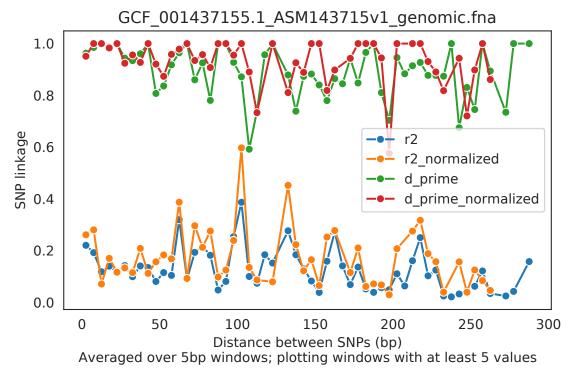
SNP linkage 0.7

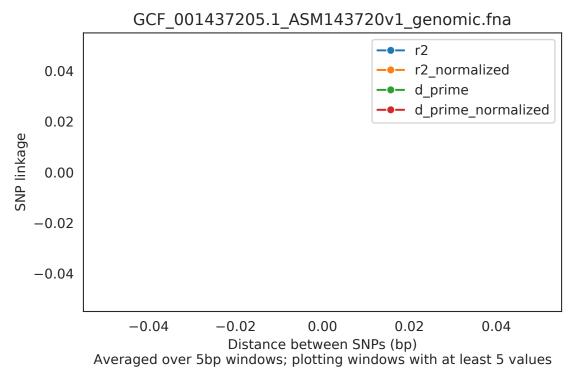
> Distance between SNPs (bp) Averaged over 5bp windows; plotting windows with at least 5 values

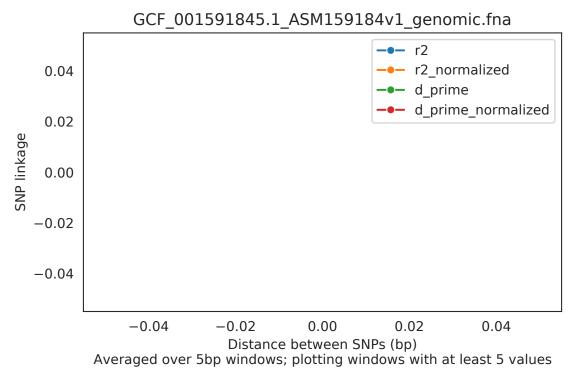




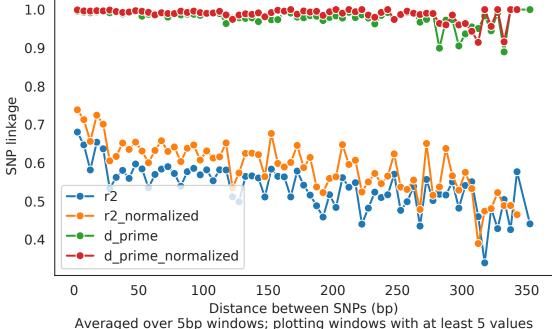


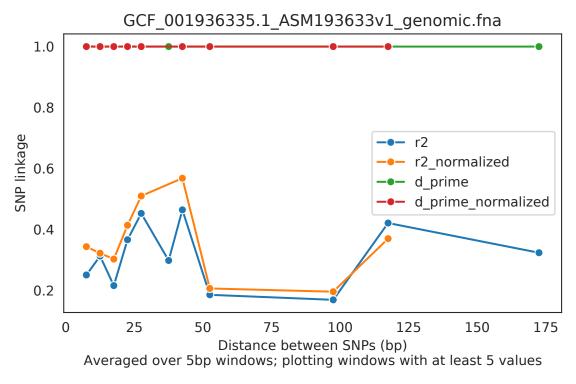


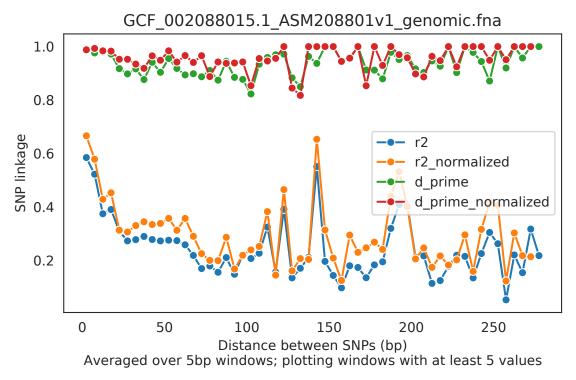


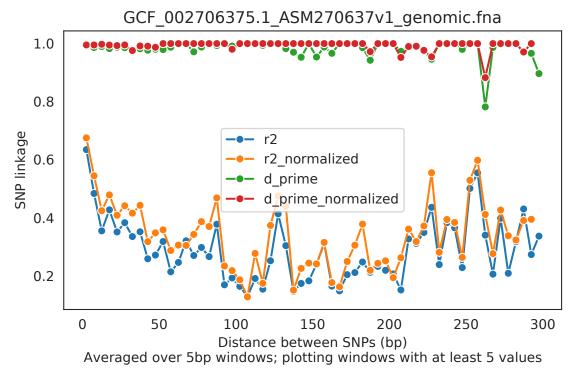


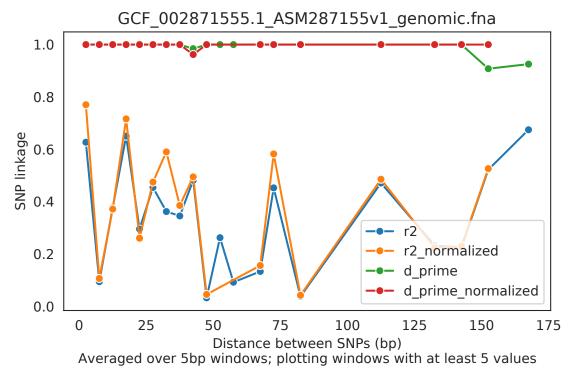
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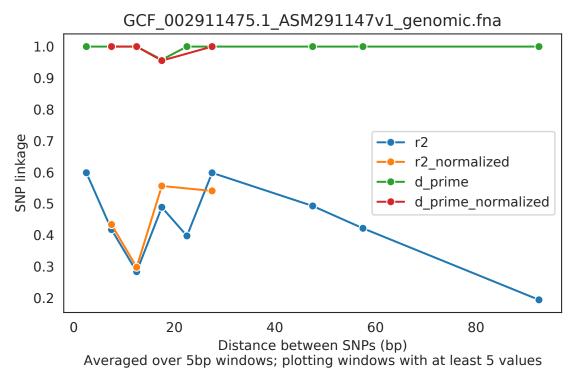


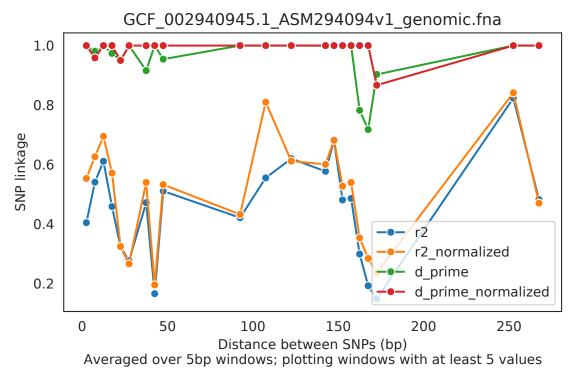


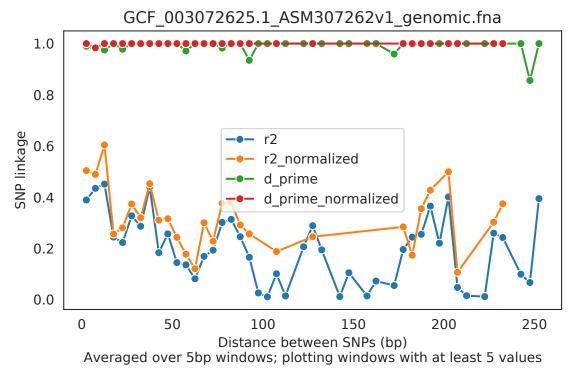


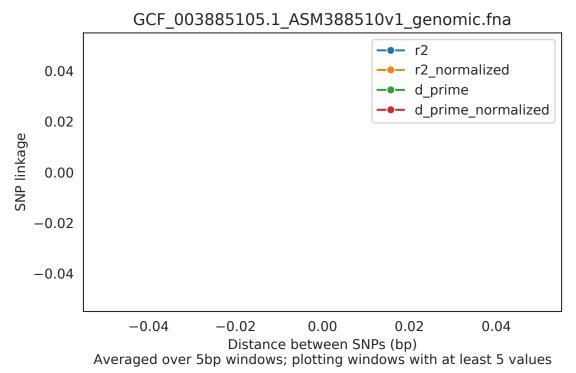


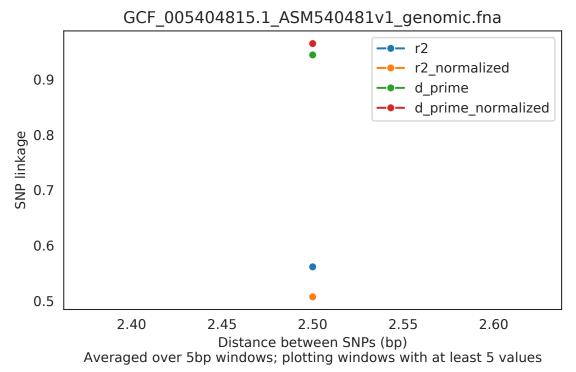


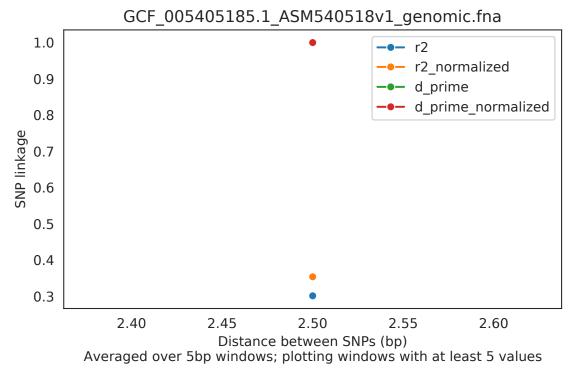


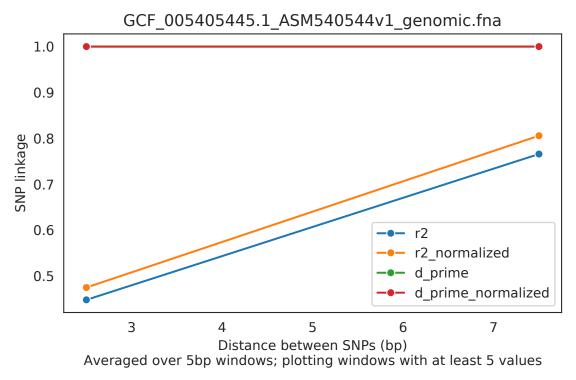


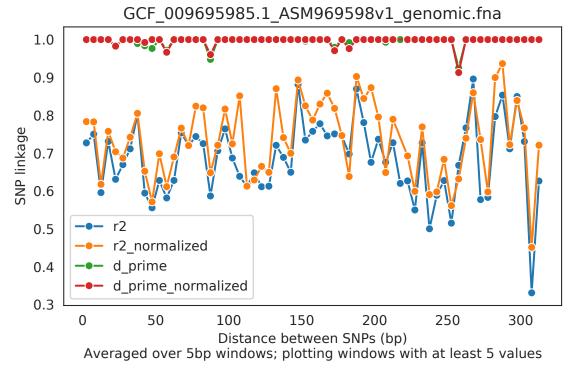


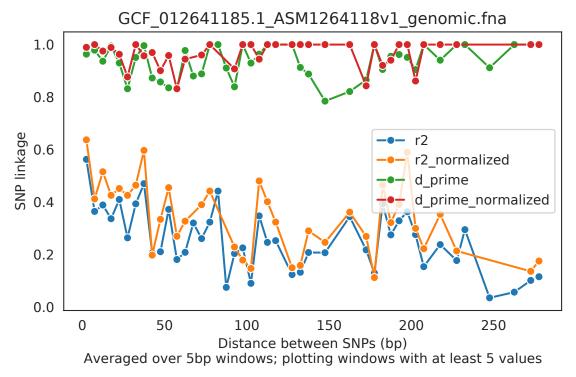


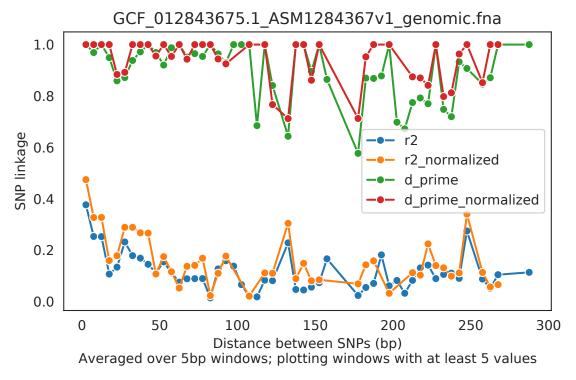


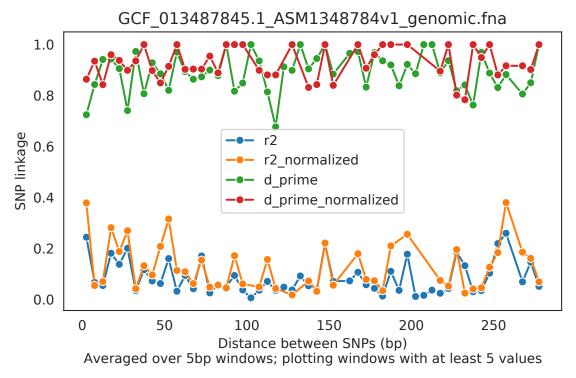


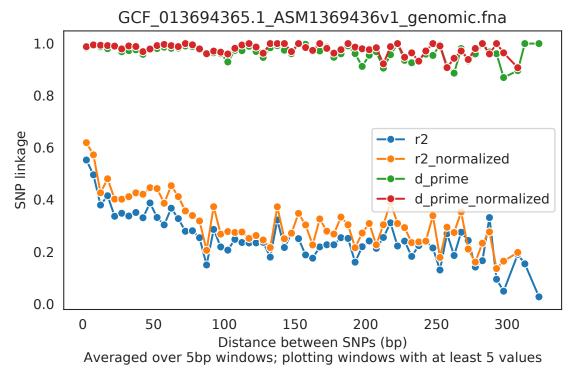


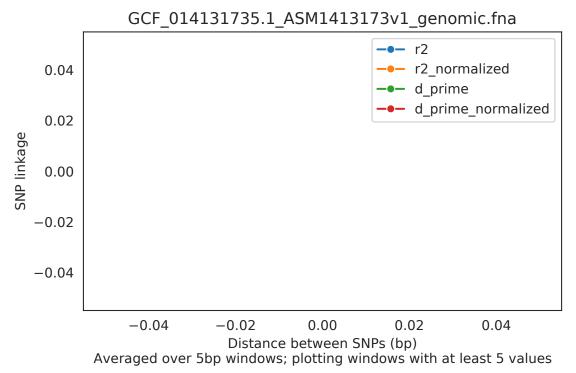


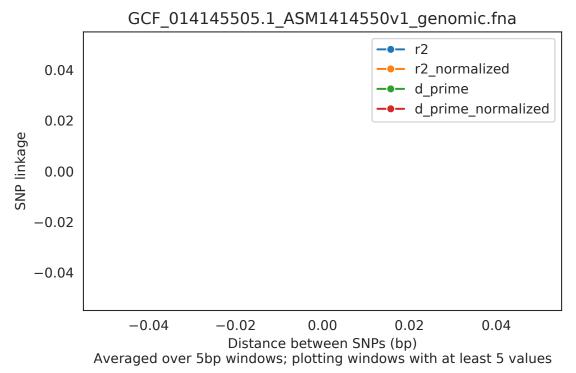


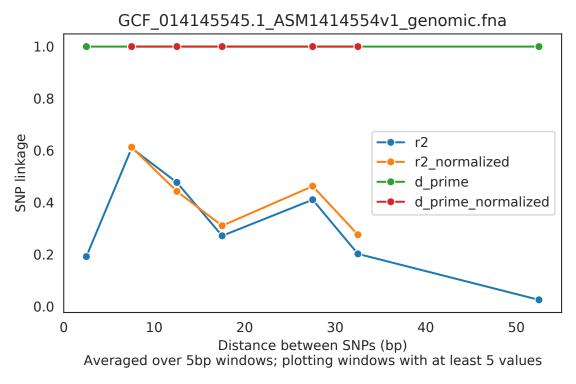


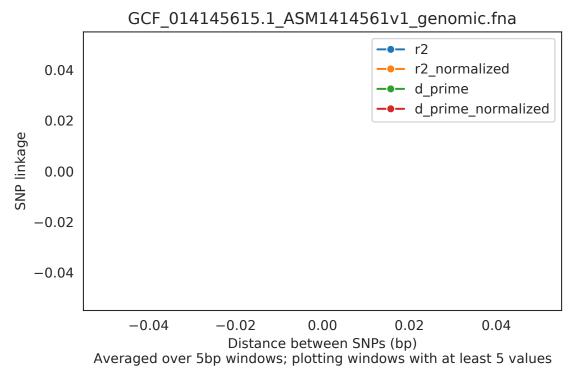


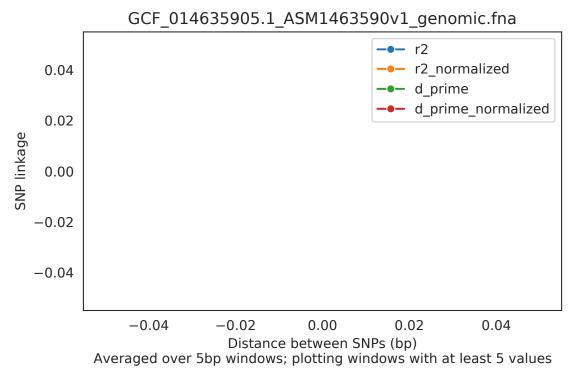


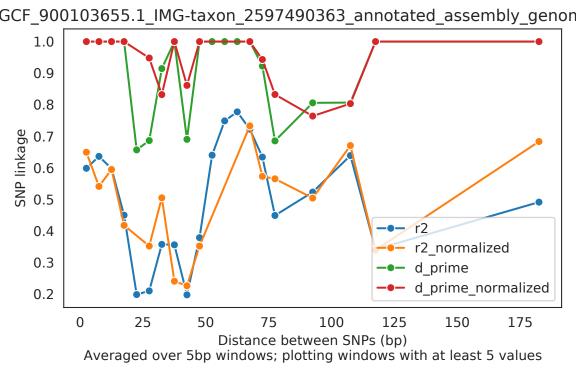




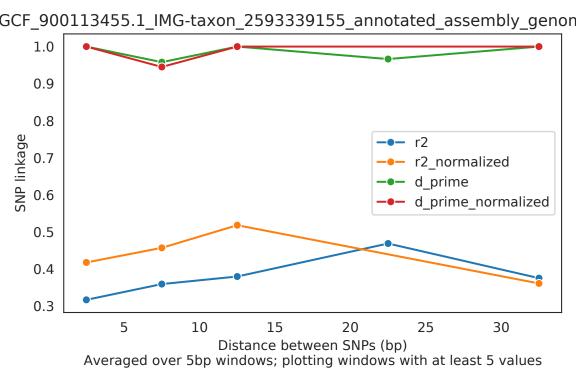


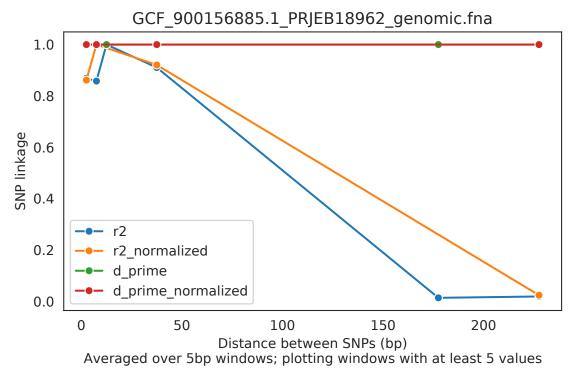


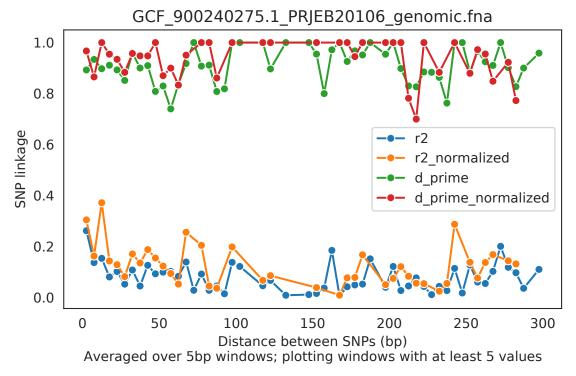


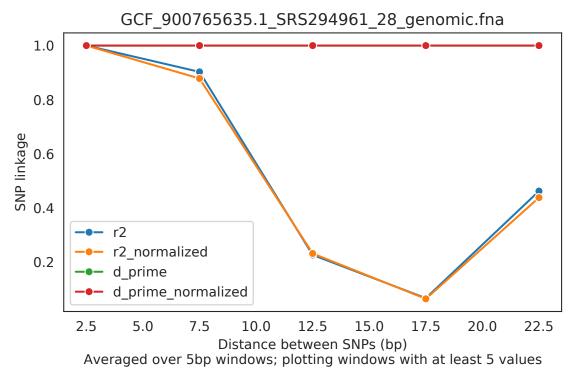


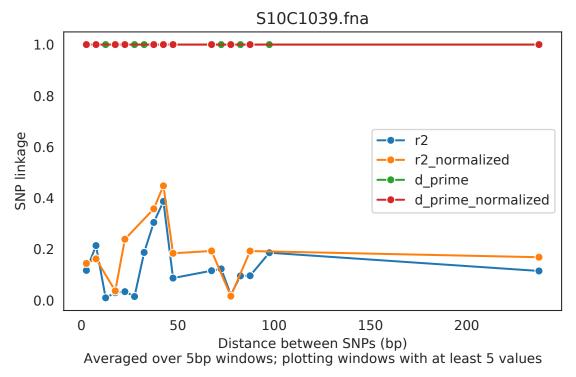
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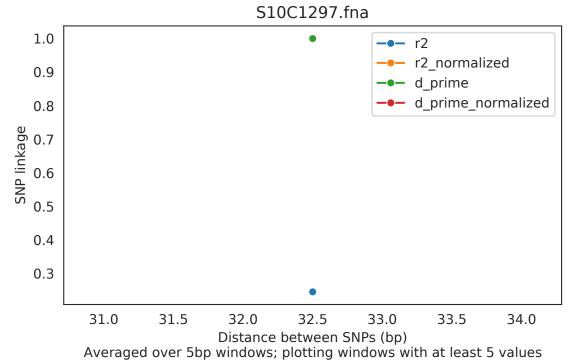


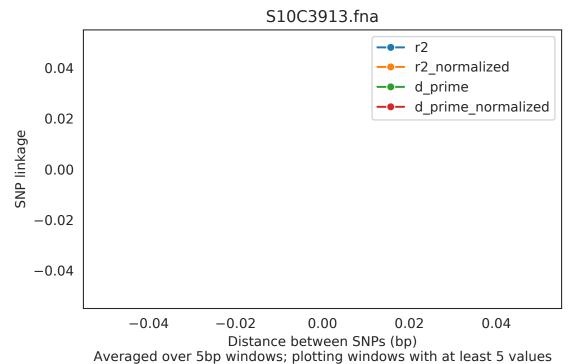




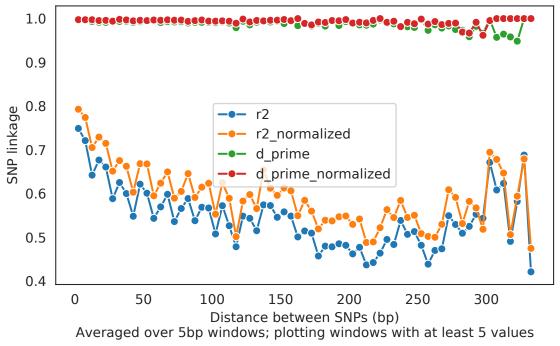


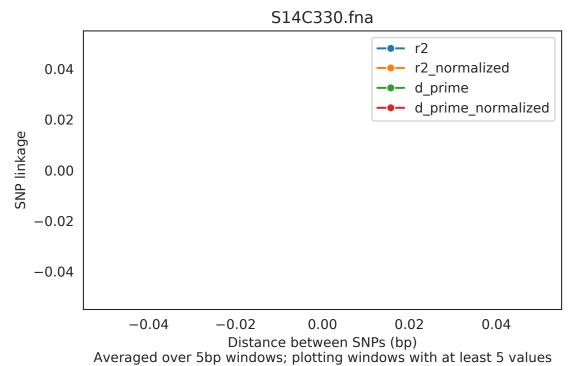


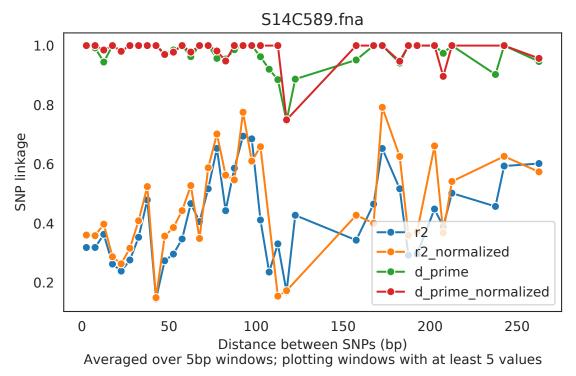


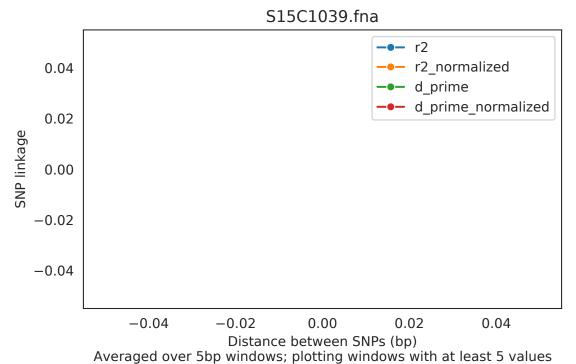


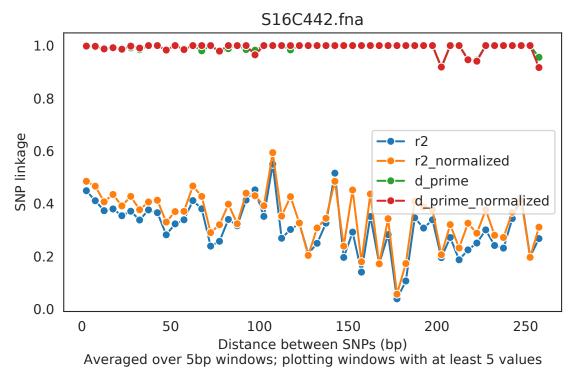
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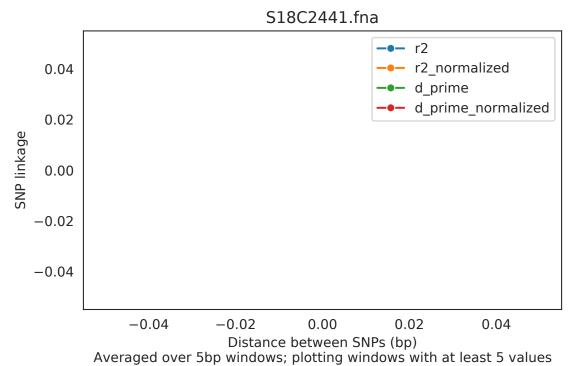




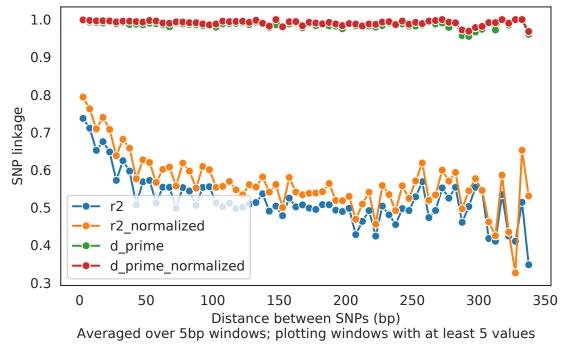








S20C3157.fna



S24C362.fna

