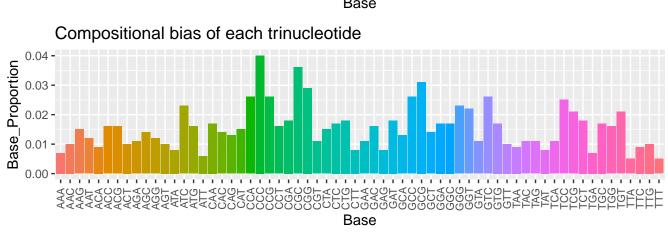
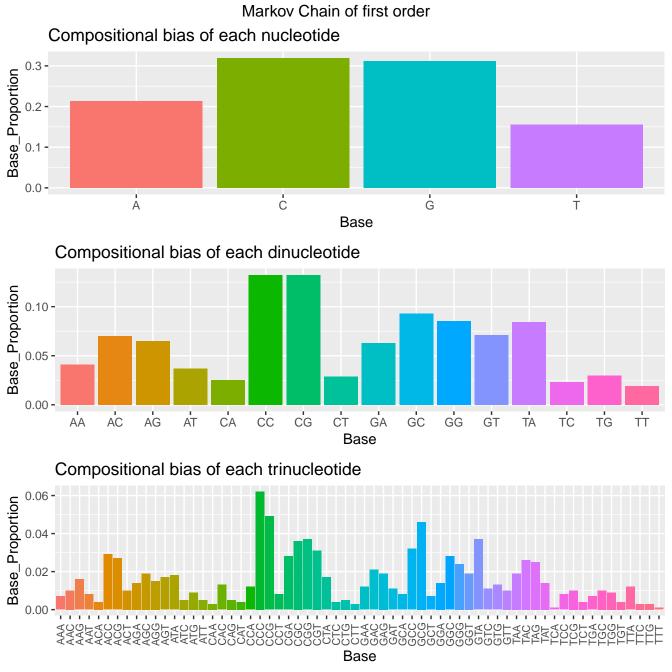
Multinomial Model of DNA Evoloution Compositional bias of each nucleotide 0.3 -Base\_Proportion 0.2 -0.0 -Ċ Ġ ÷ Α Base Compositional bias of each dinucleotide Base\_Proportion 0.09 -0.06 -0.03 -0.00 -AC AG ΑT СA СC СĠ СТ ĞA ĠС ĠĠ ďТ ΤΆ TC ΤĠ Τ̈́Τ ΑA Base



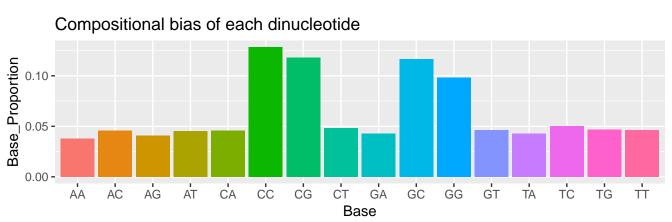


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