

Ne = 30000

Density of segregating sites

Nucleotide diversity (π)

Number of mutations

Number of trees

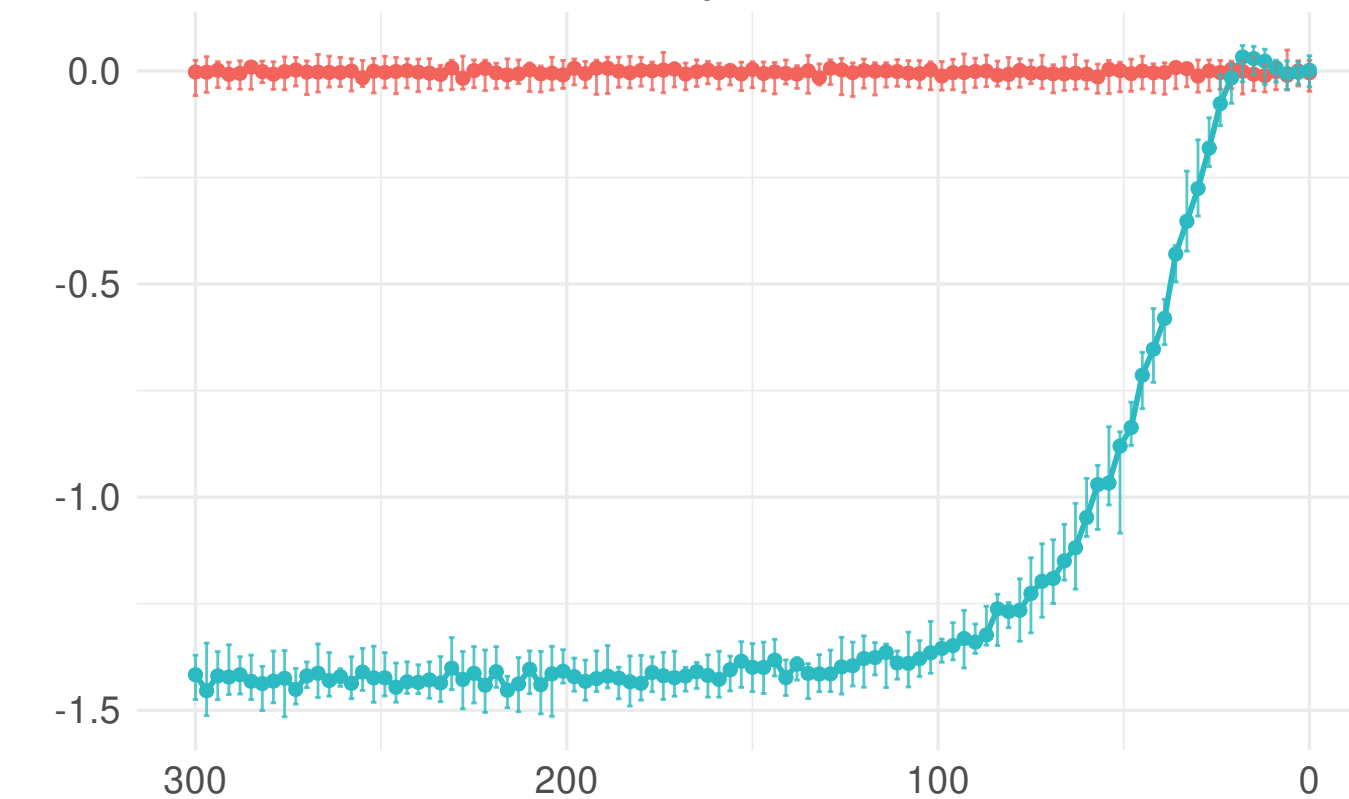
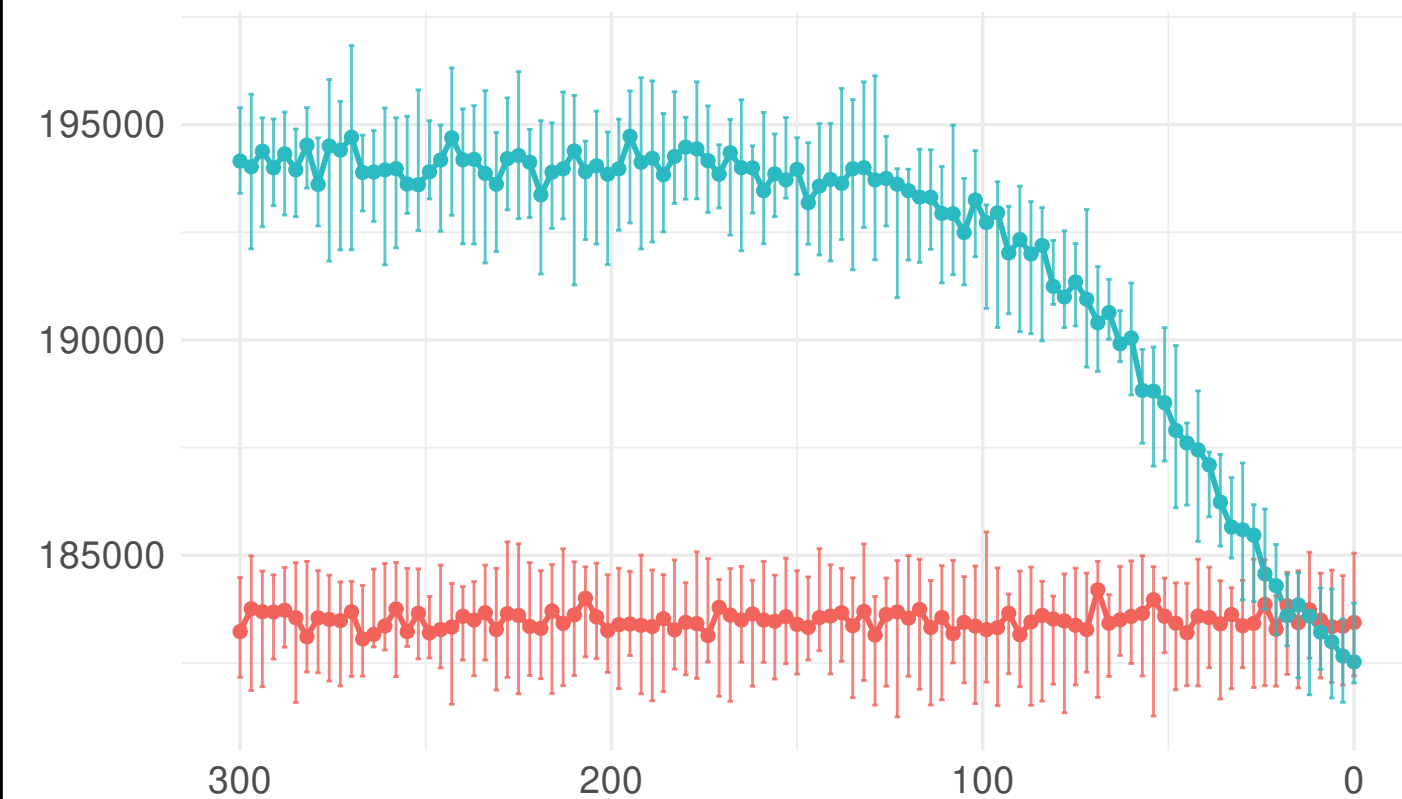
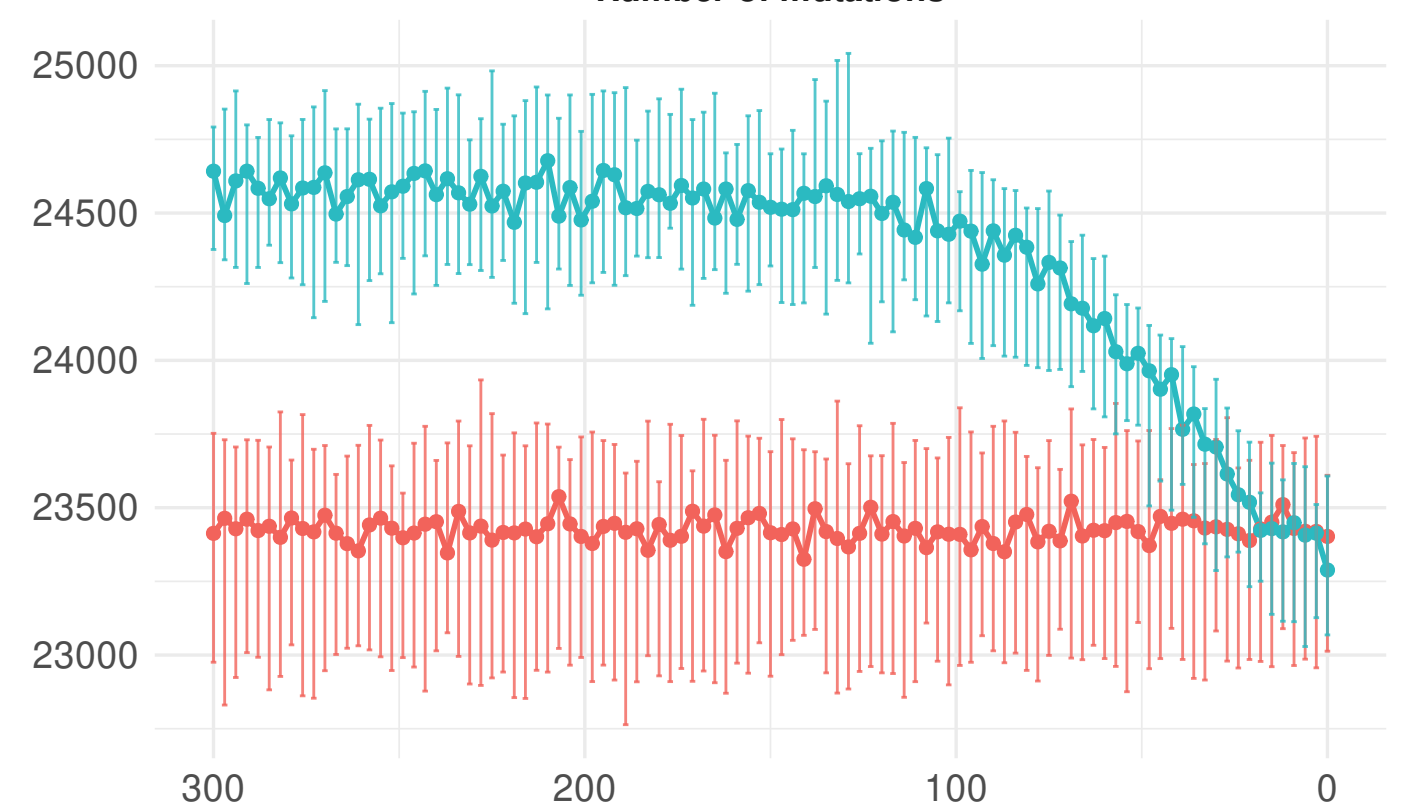
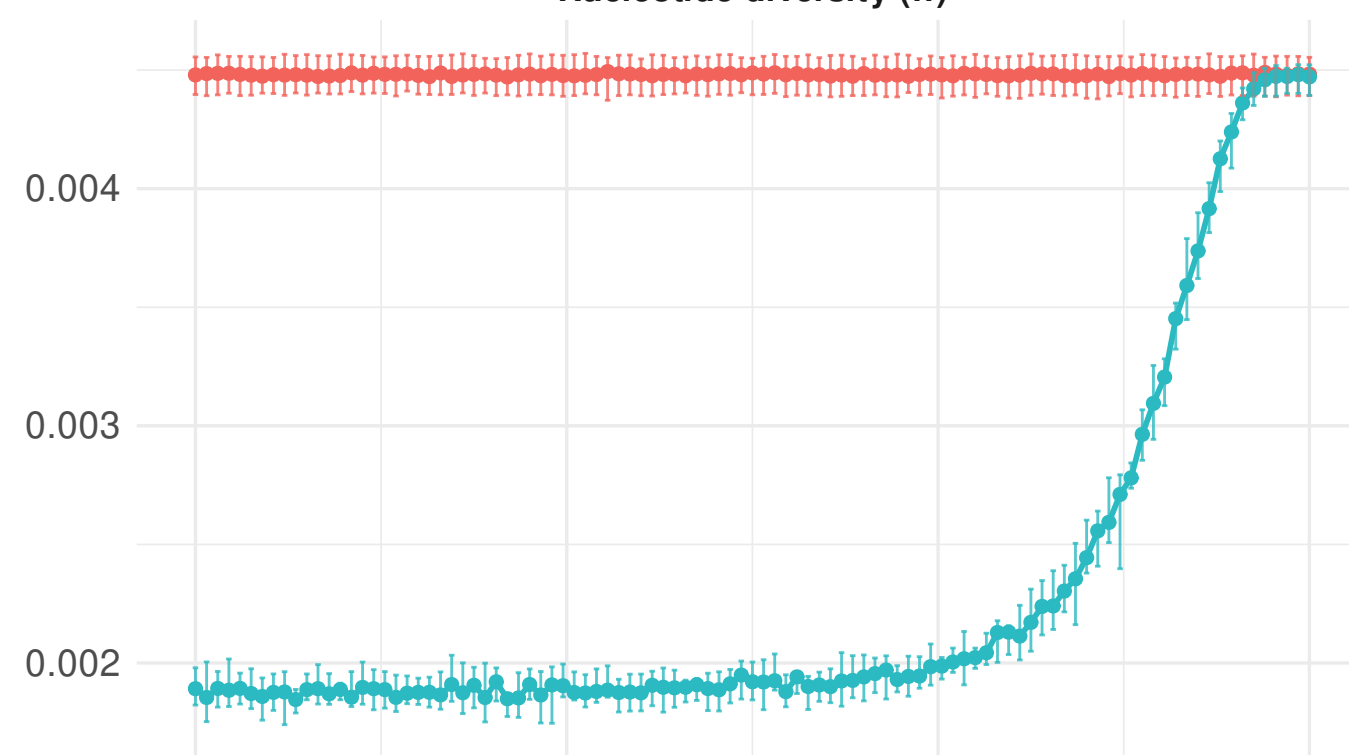
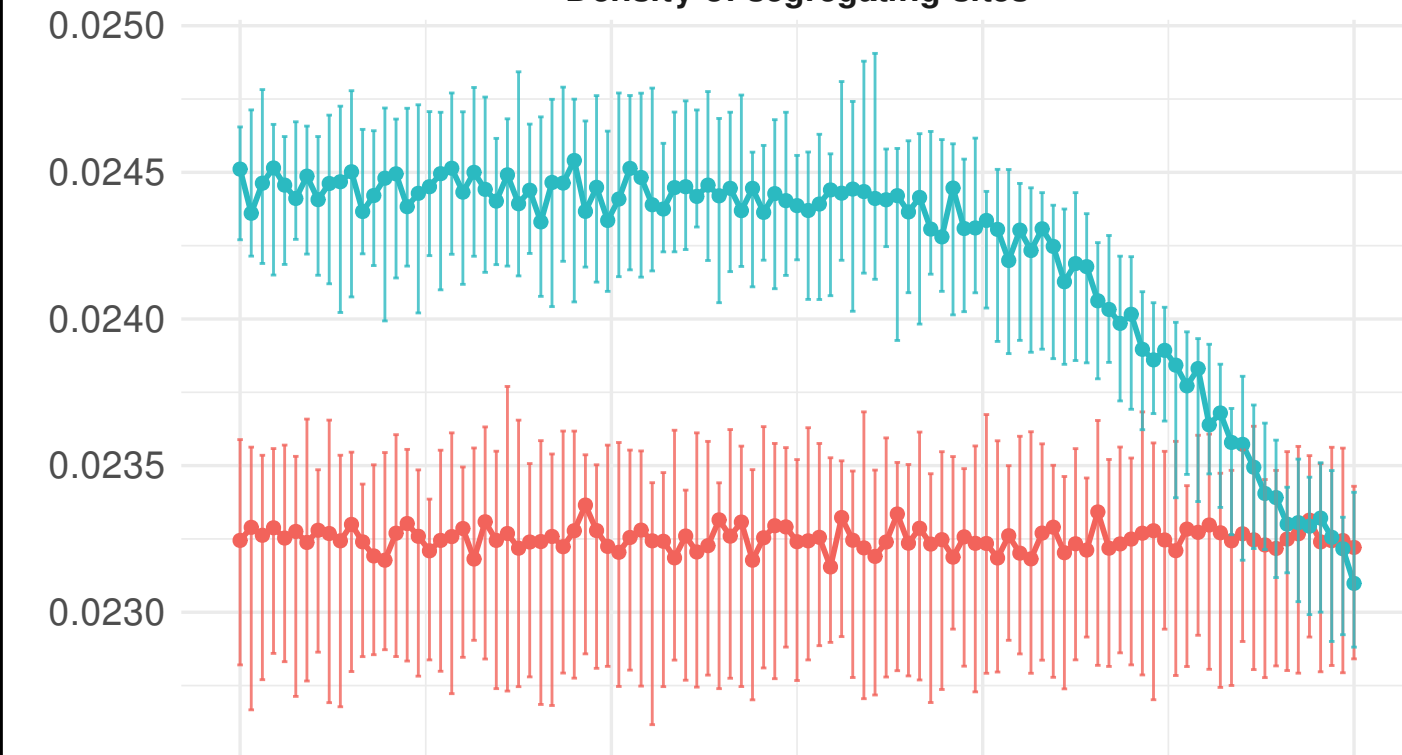
Tajima's D

Model

Hudson

WF

Generations ago (past -> present)



Ne = 3000

Density of segregating sites

Nucleotide diversity (π)

Number of mutations

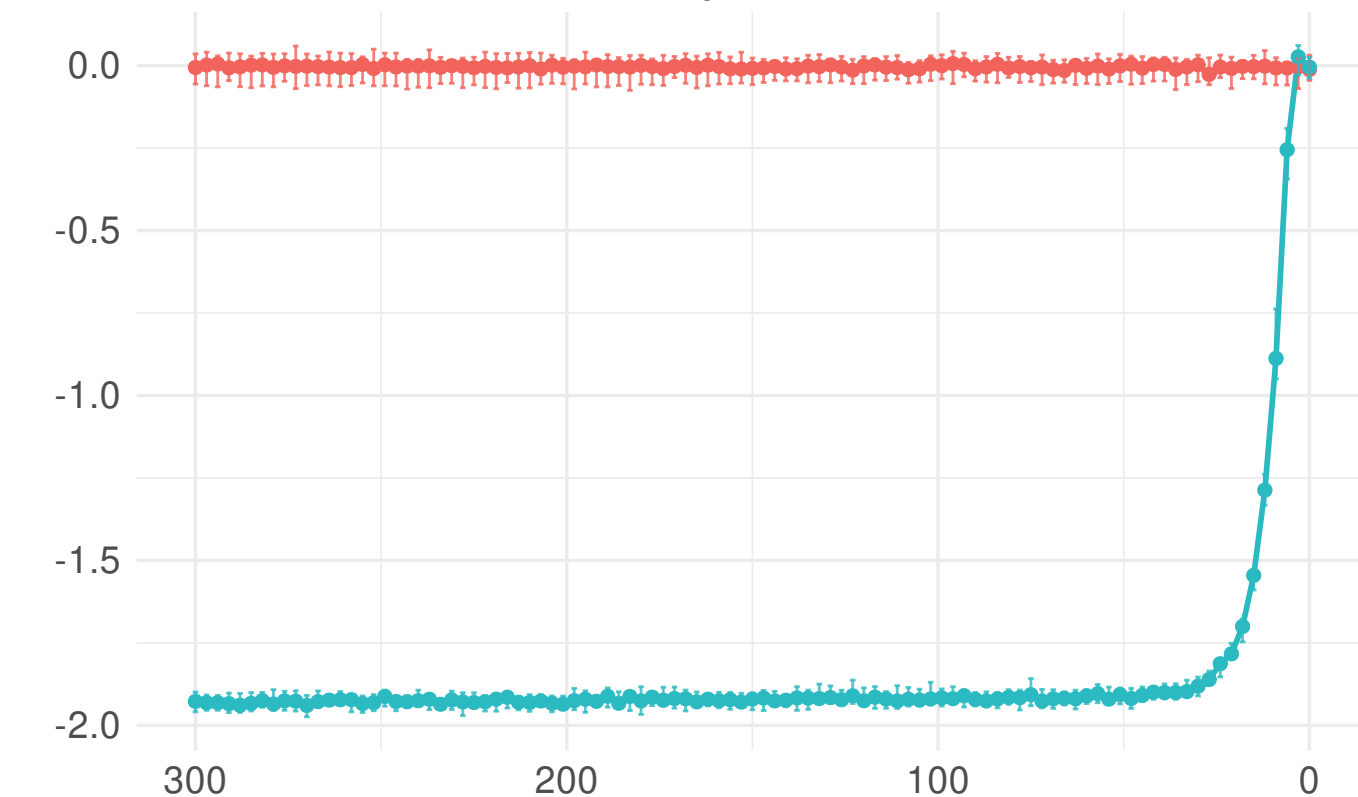
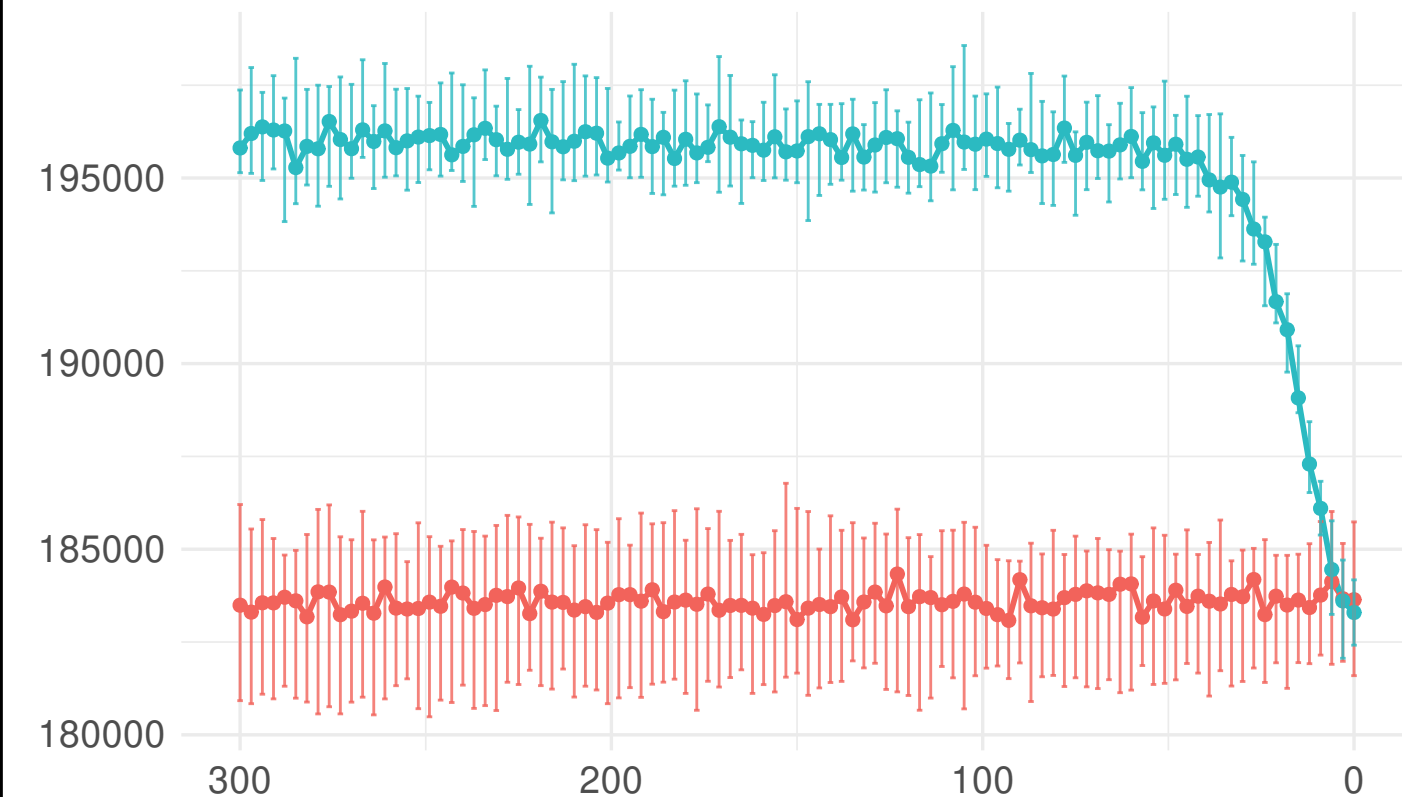
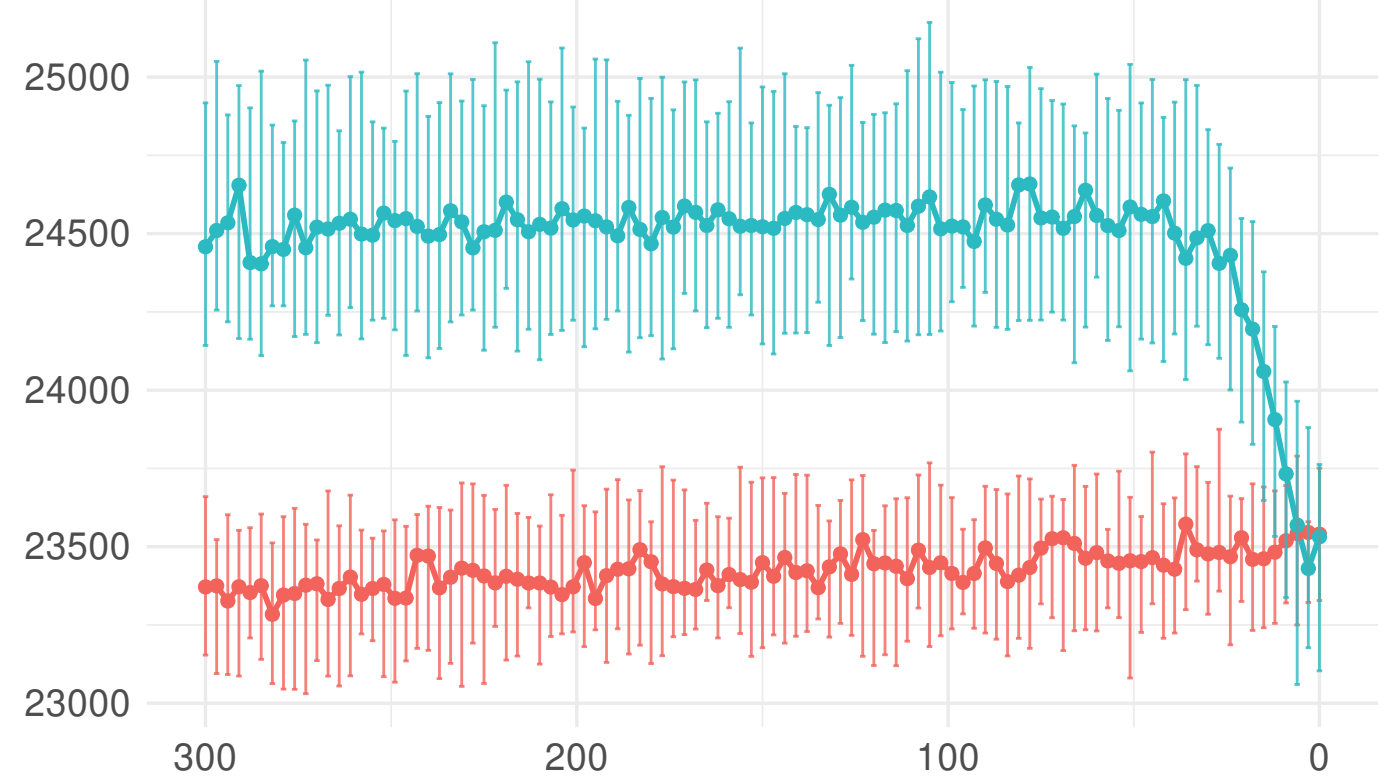
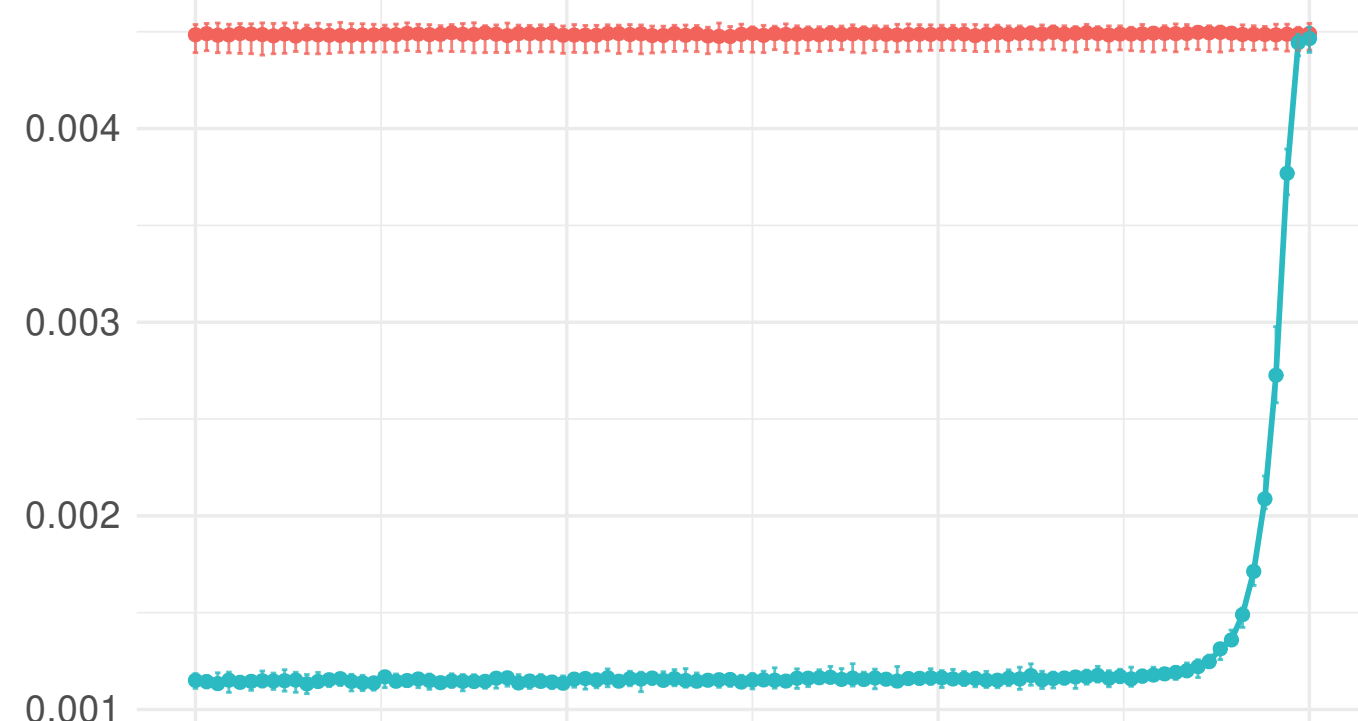
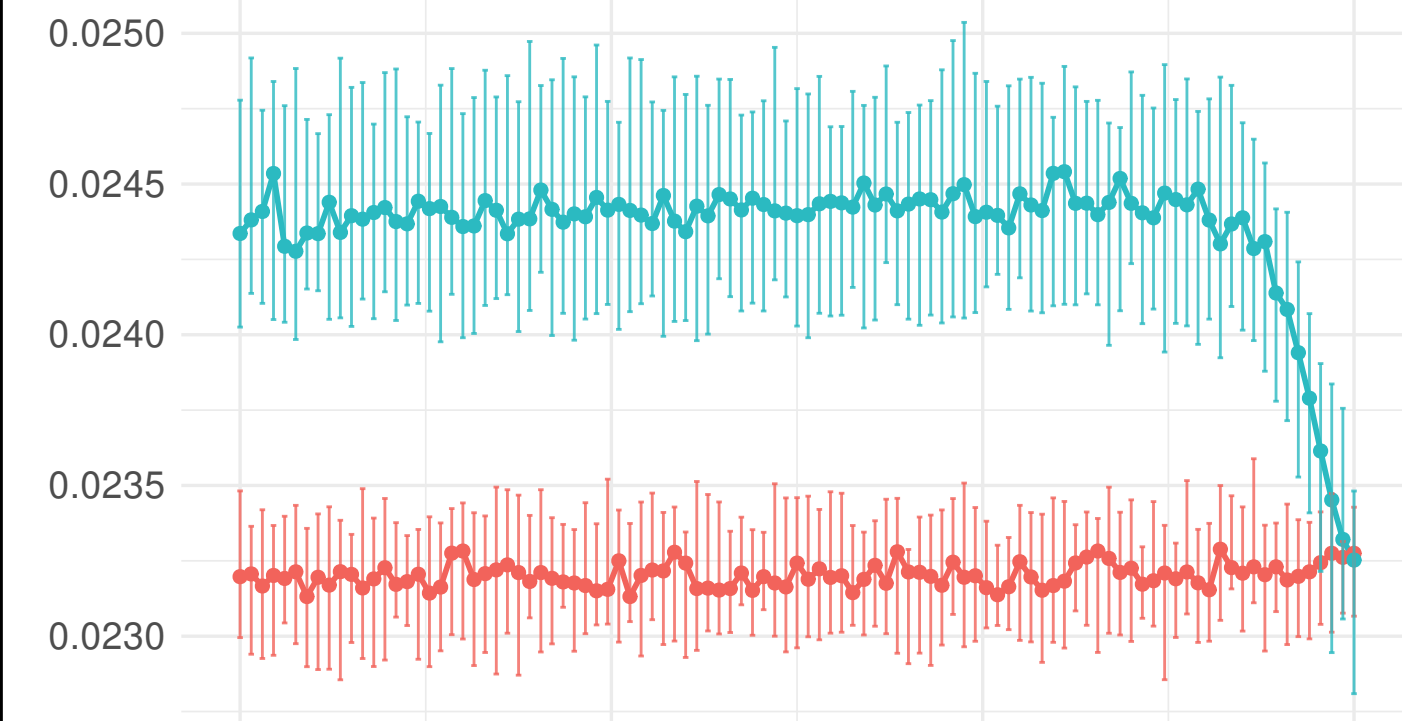
Number of trees

Tajima's D

Model

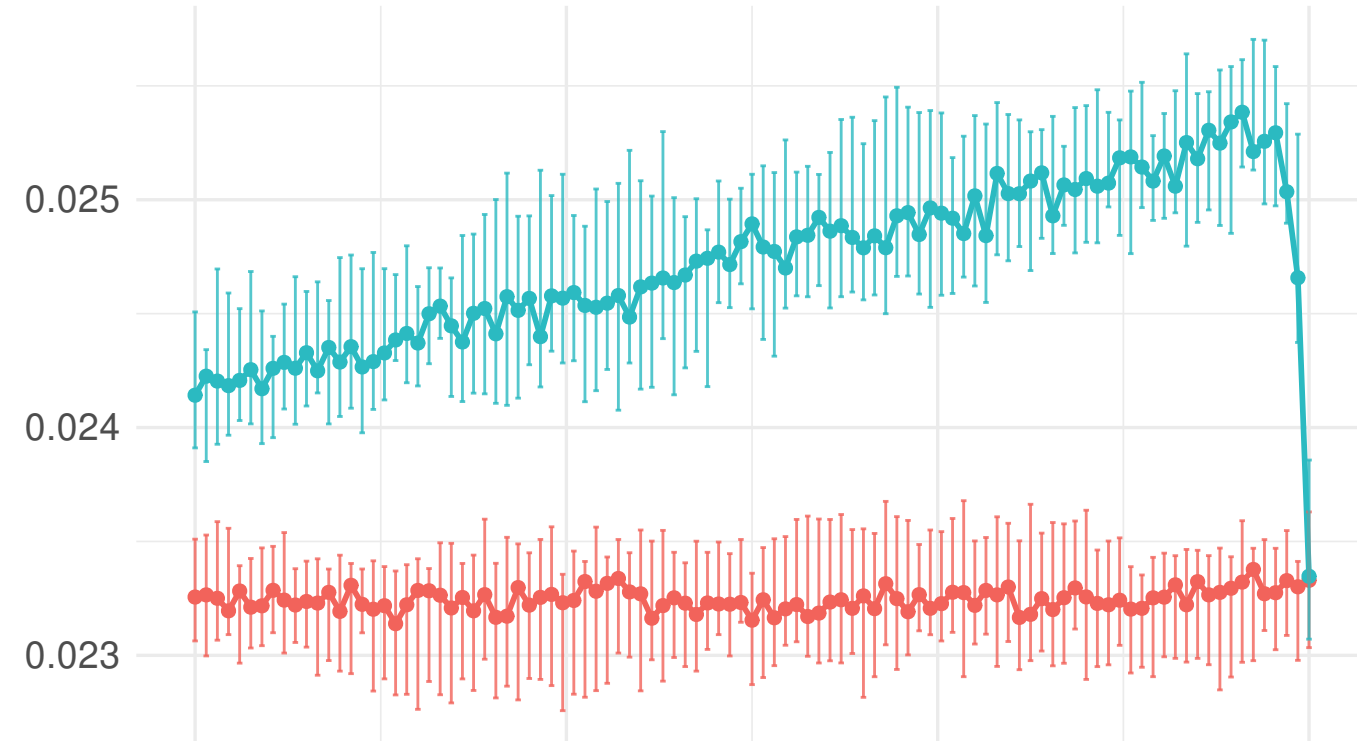
Hudson
WF

Generations ago (past -> present)

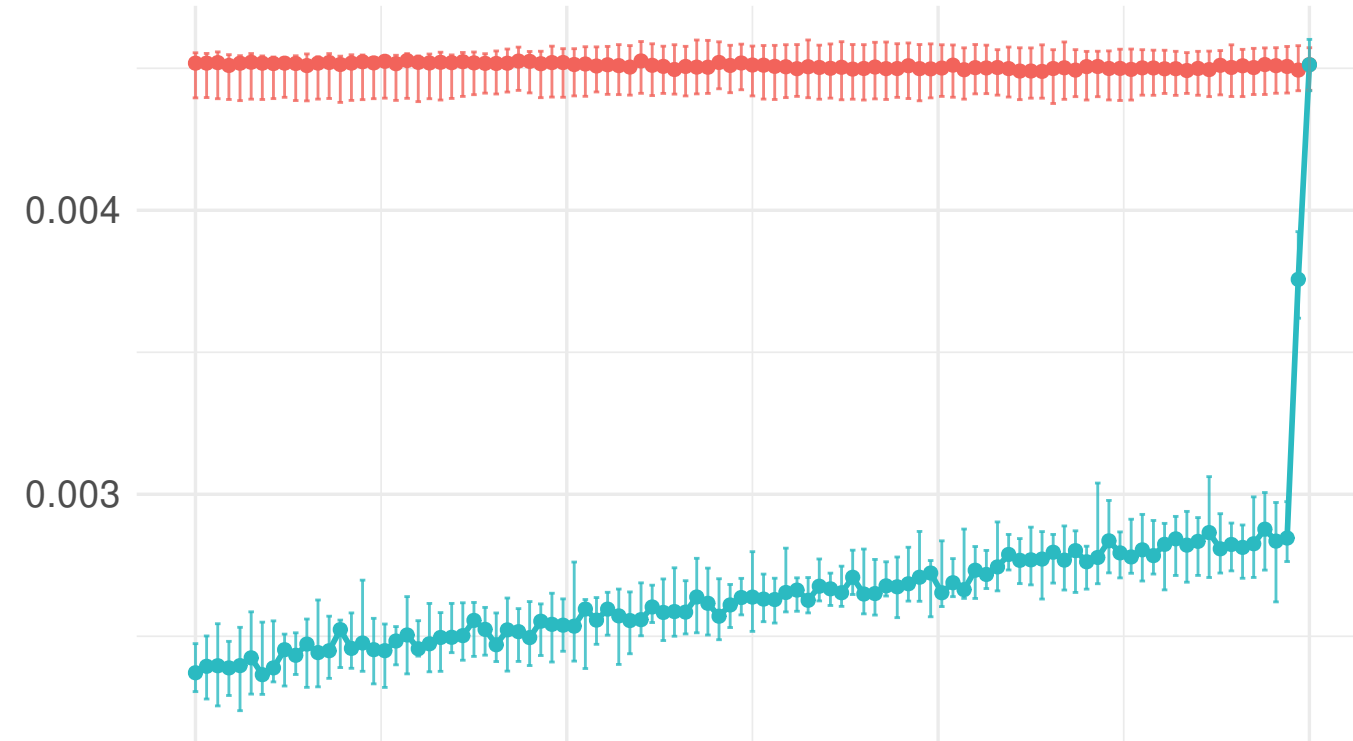


Ne = 300

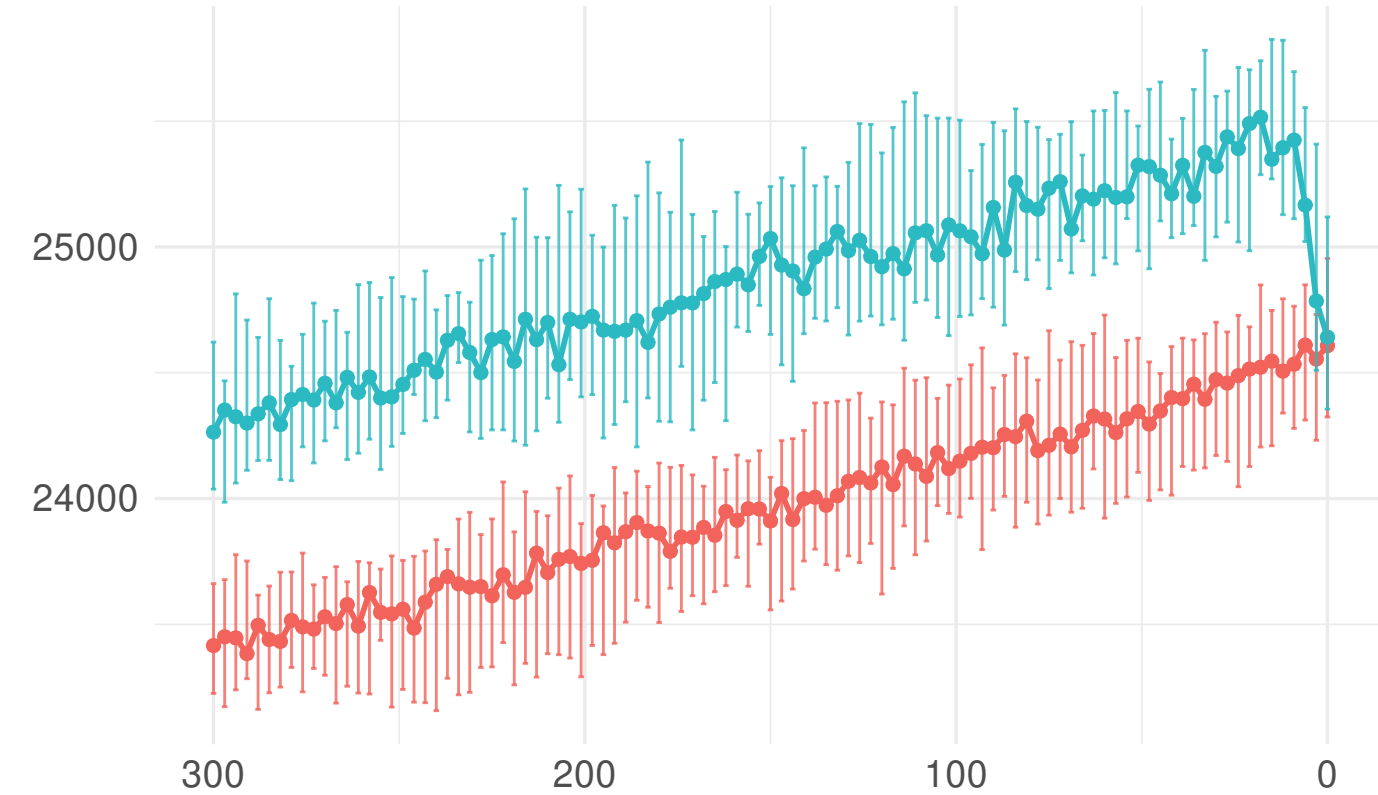
Density of segregating sites



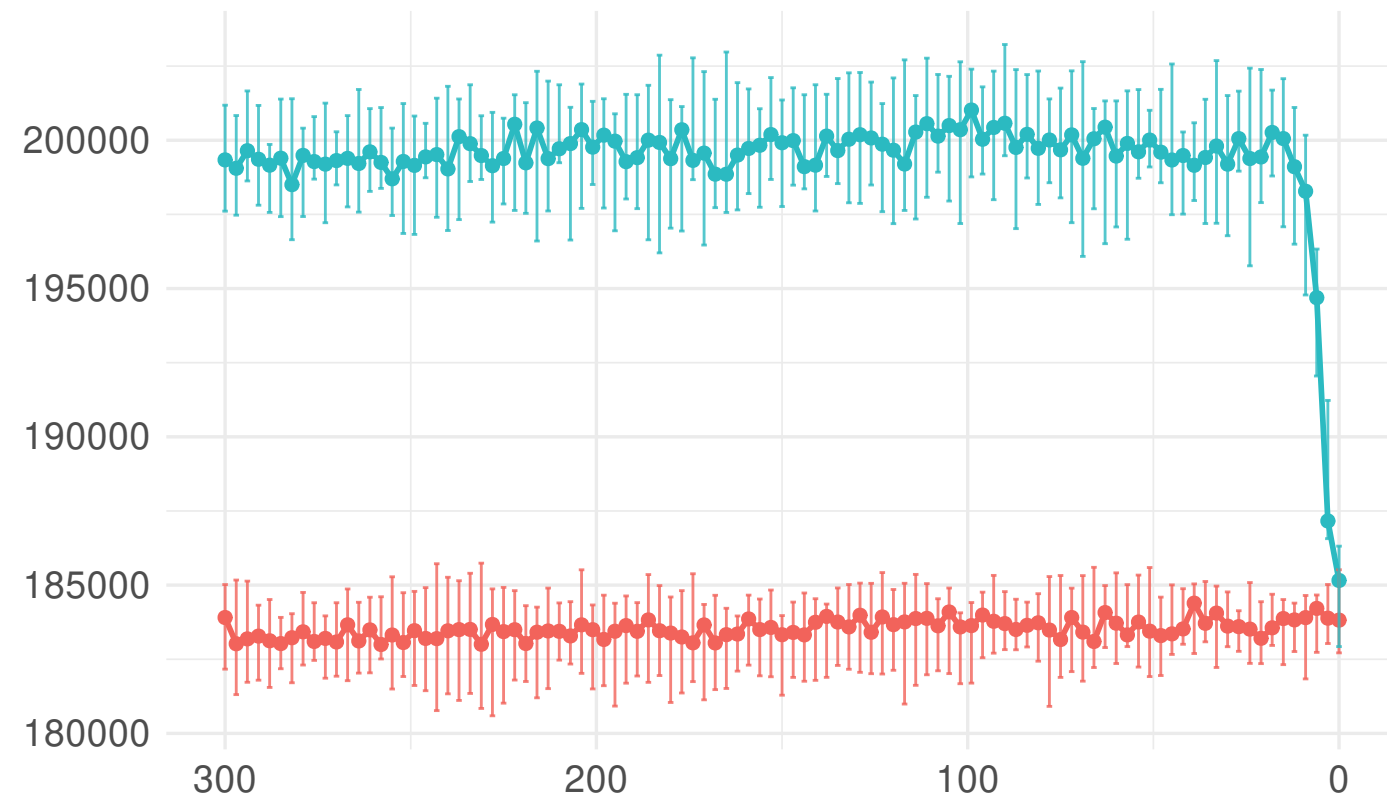
Nucleotide diversity (π)



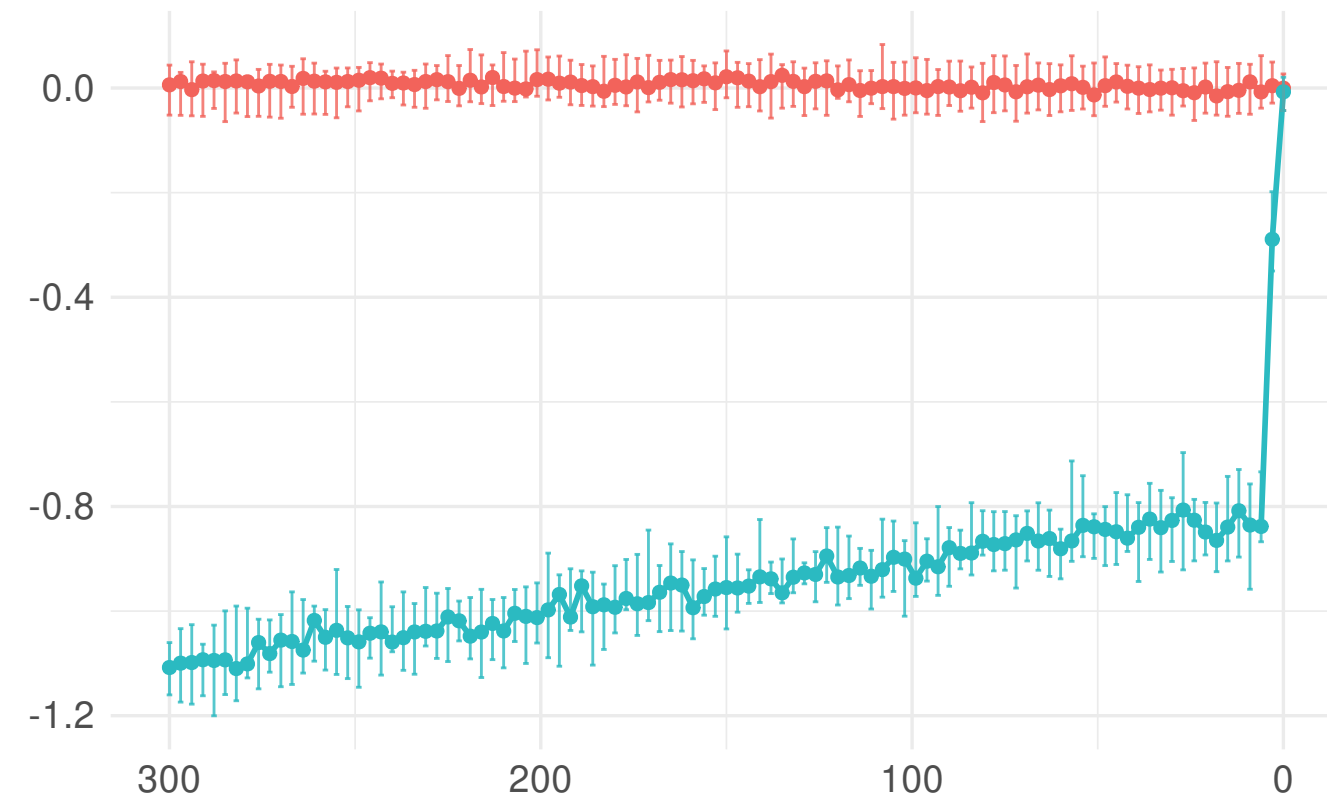
Number of mutations



Number of trees



Tajima's D



Generations ago (past -> present)

Model

- Hudson
- WF