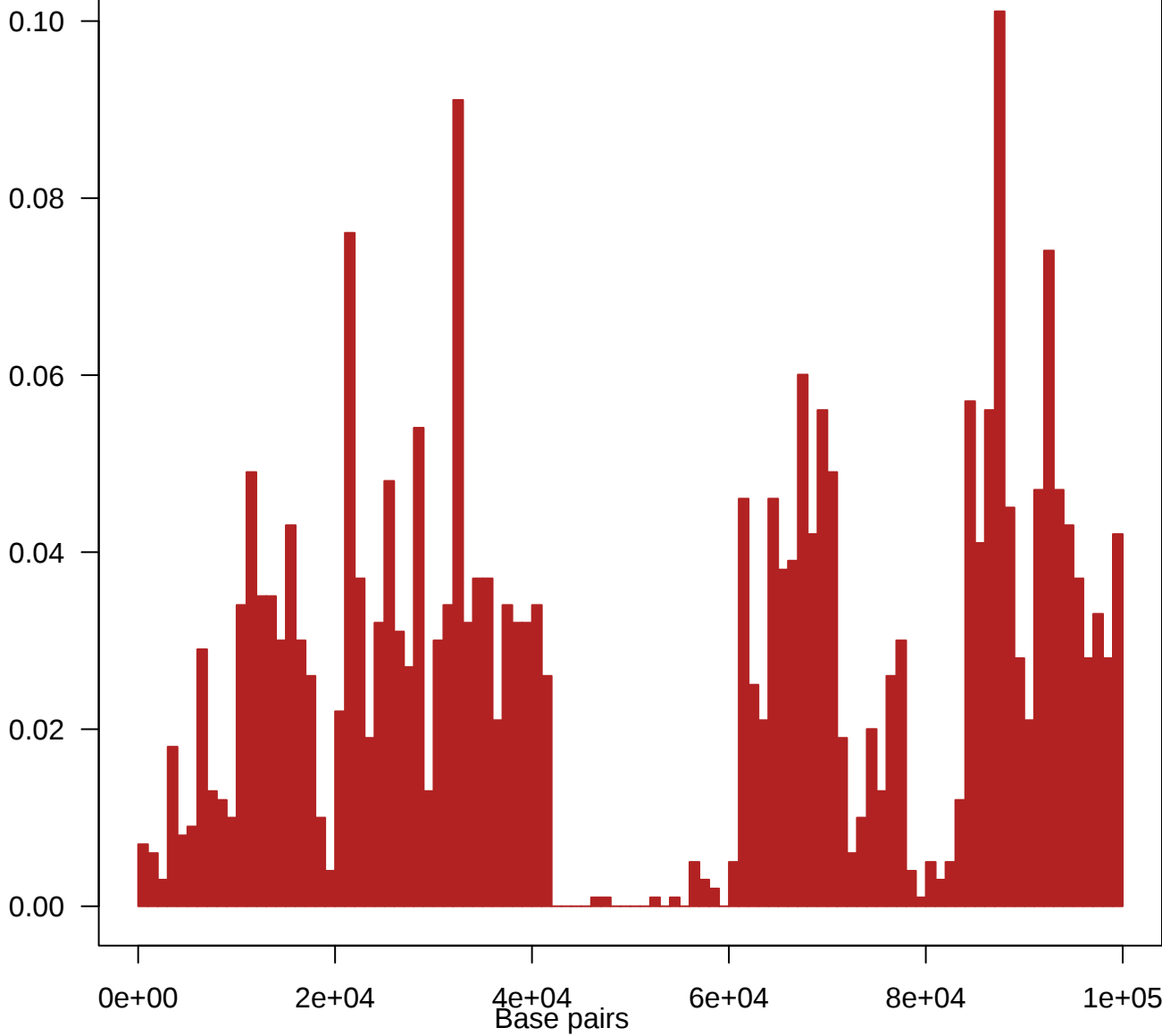


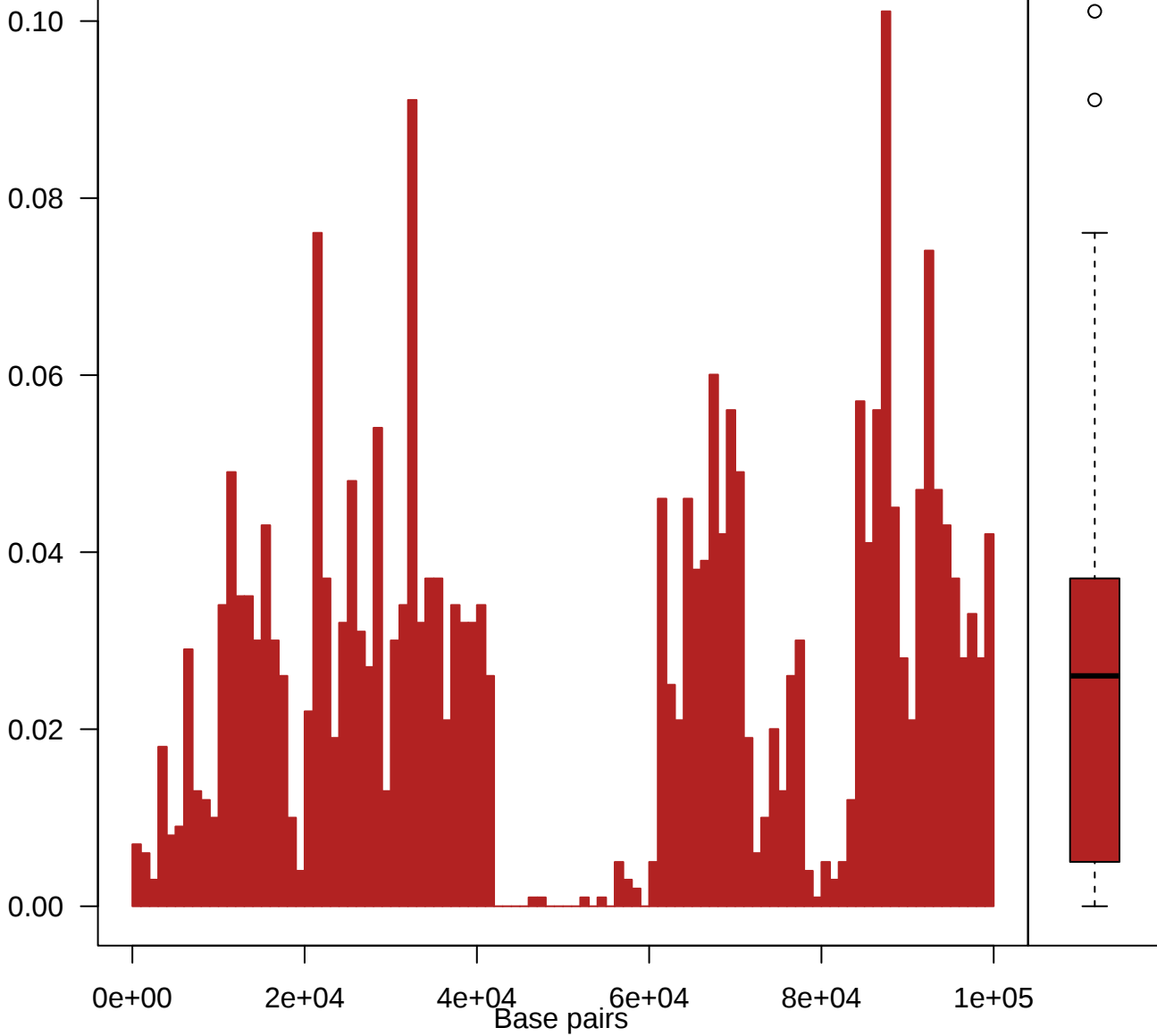
Supercontig

Variants per Site

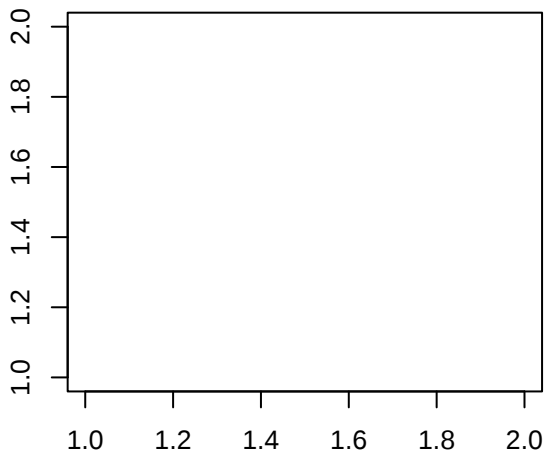


Supercontig

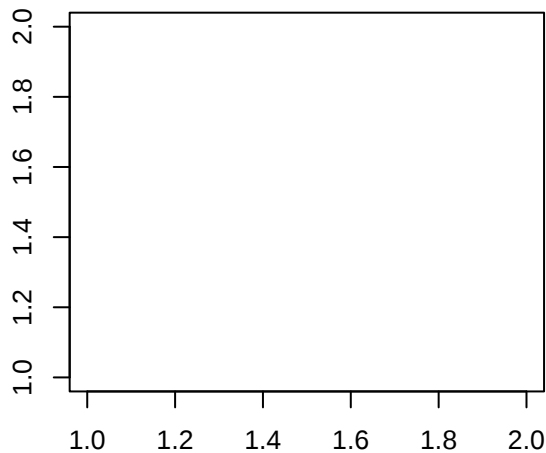
Variants per Site



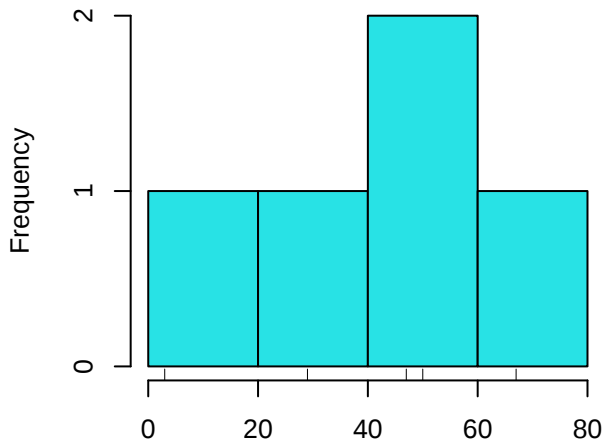
No depths found



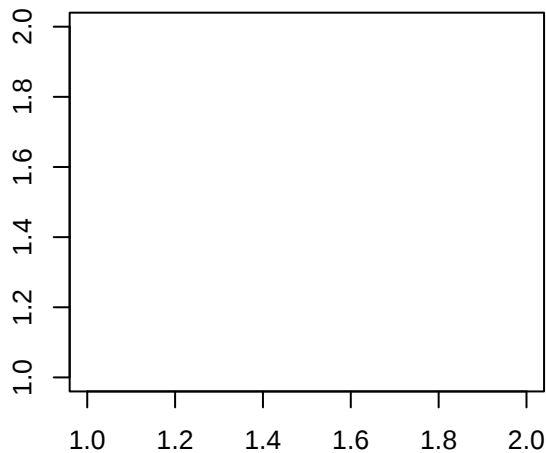
No mapping qualities found



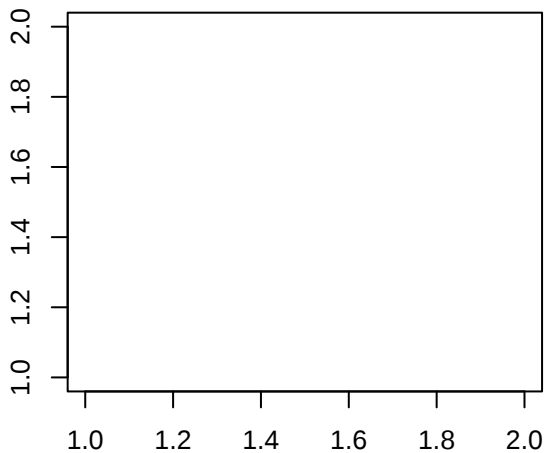
Quality (QUAL)



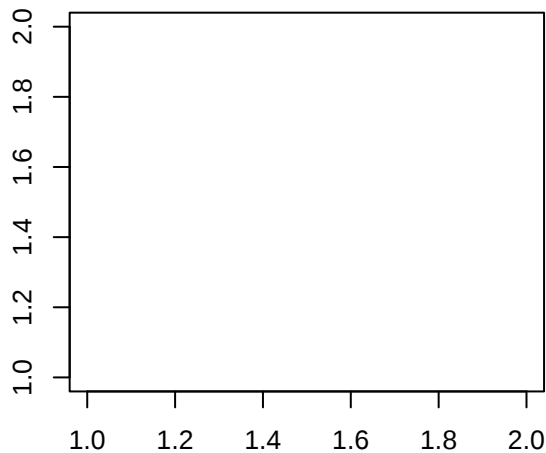
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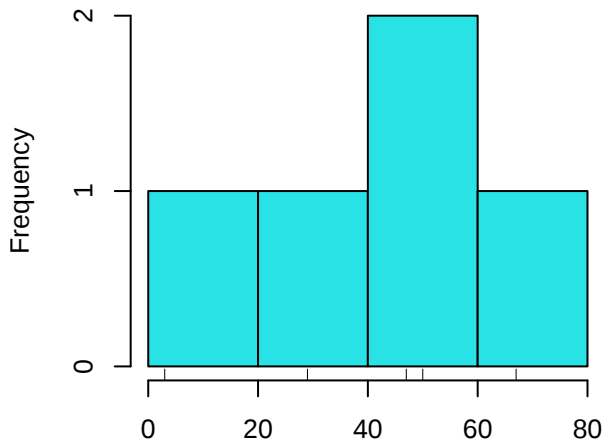
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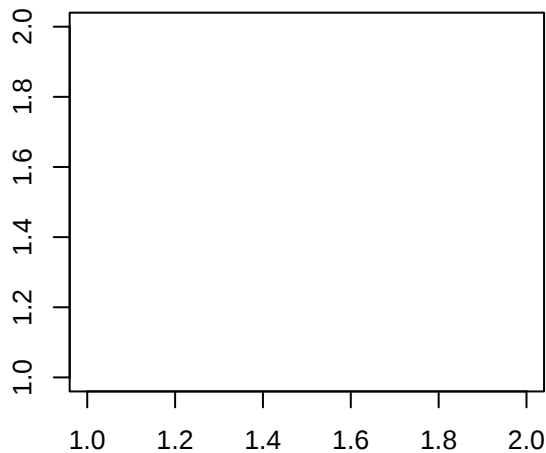
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Quality (QUAL)

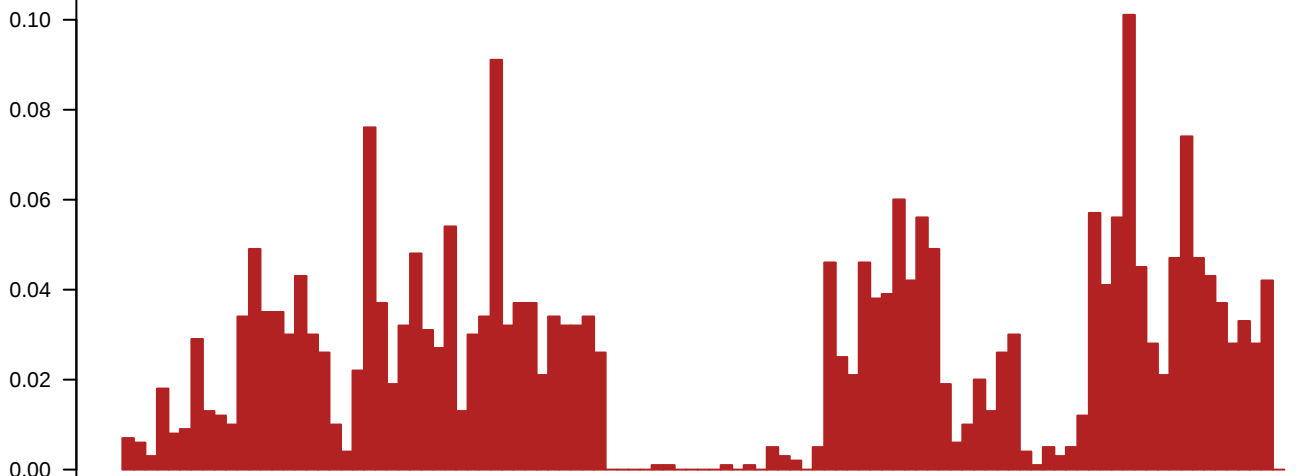


No SNP densities found



Supercontig

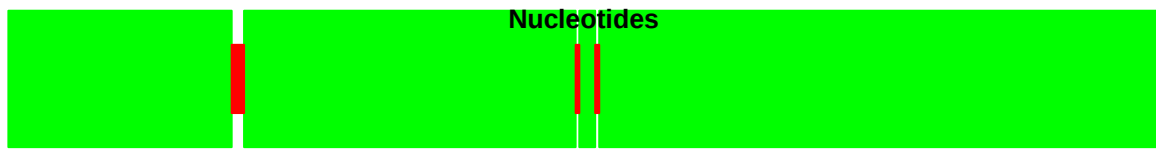
Variants per Site



Nucleotide Content

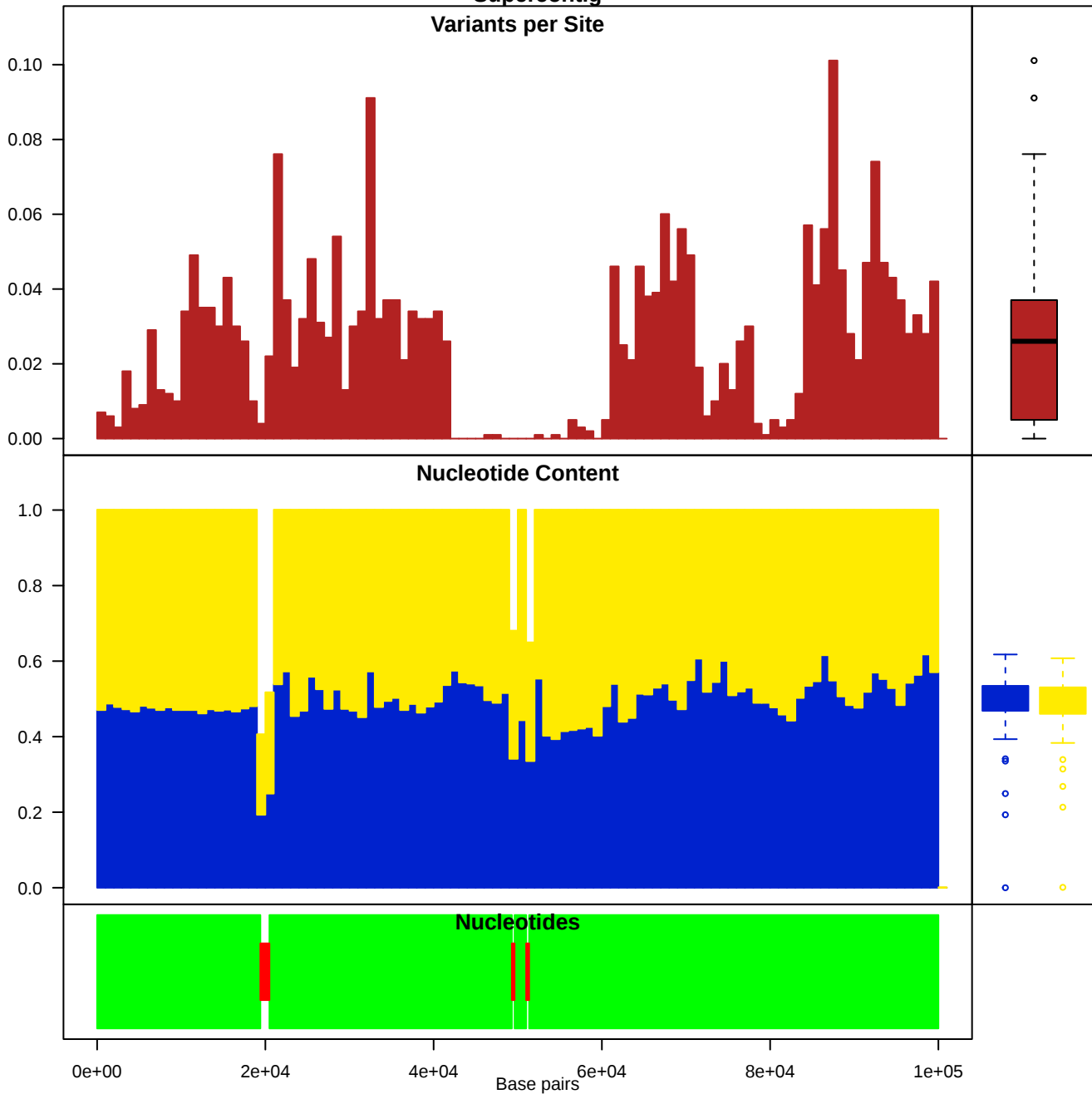


Nucleotides



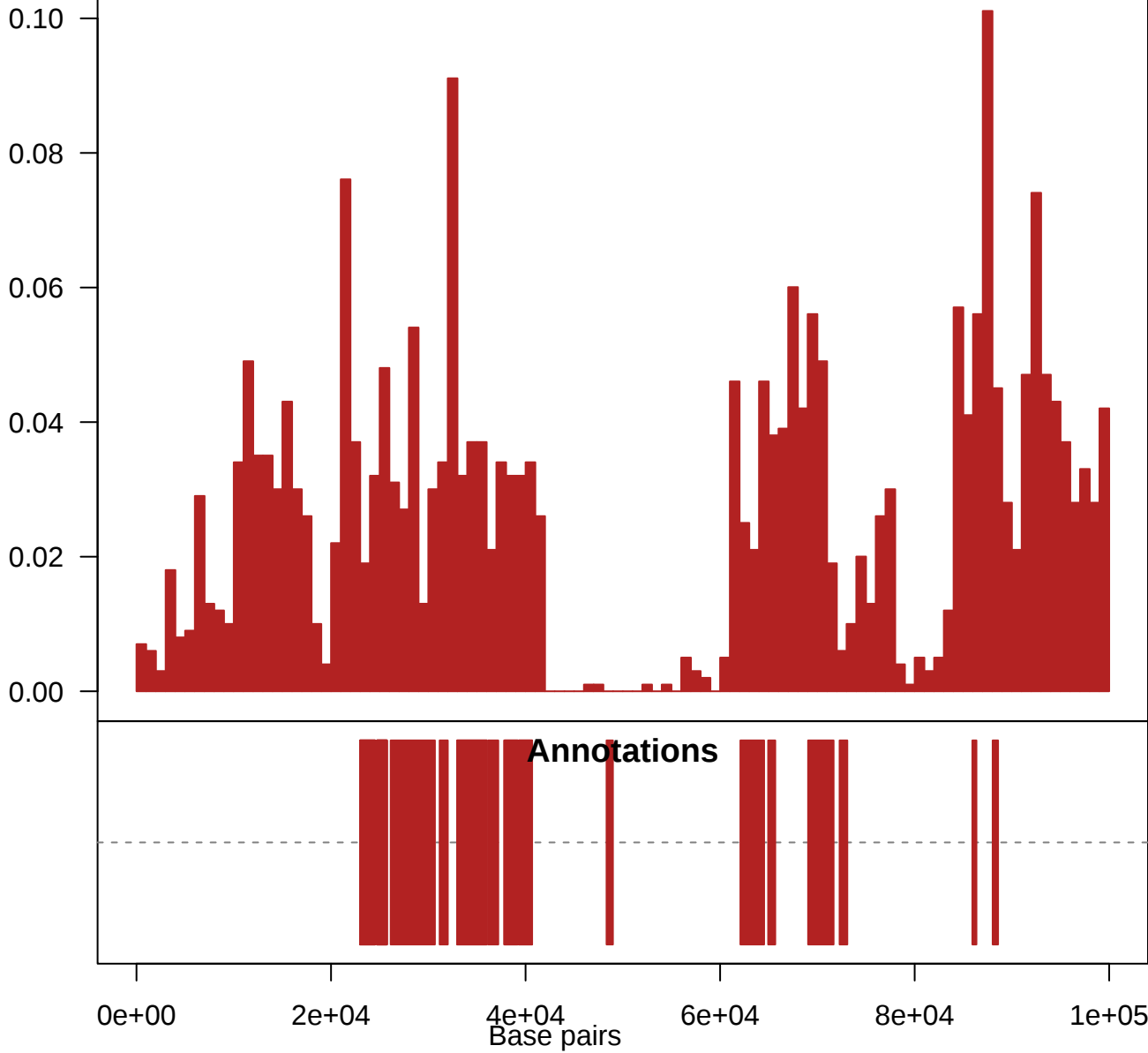
Base pairs

Supercontig



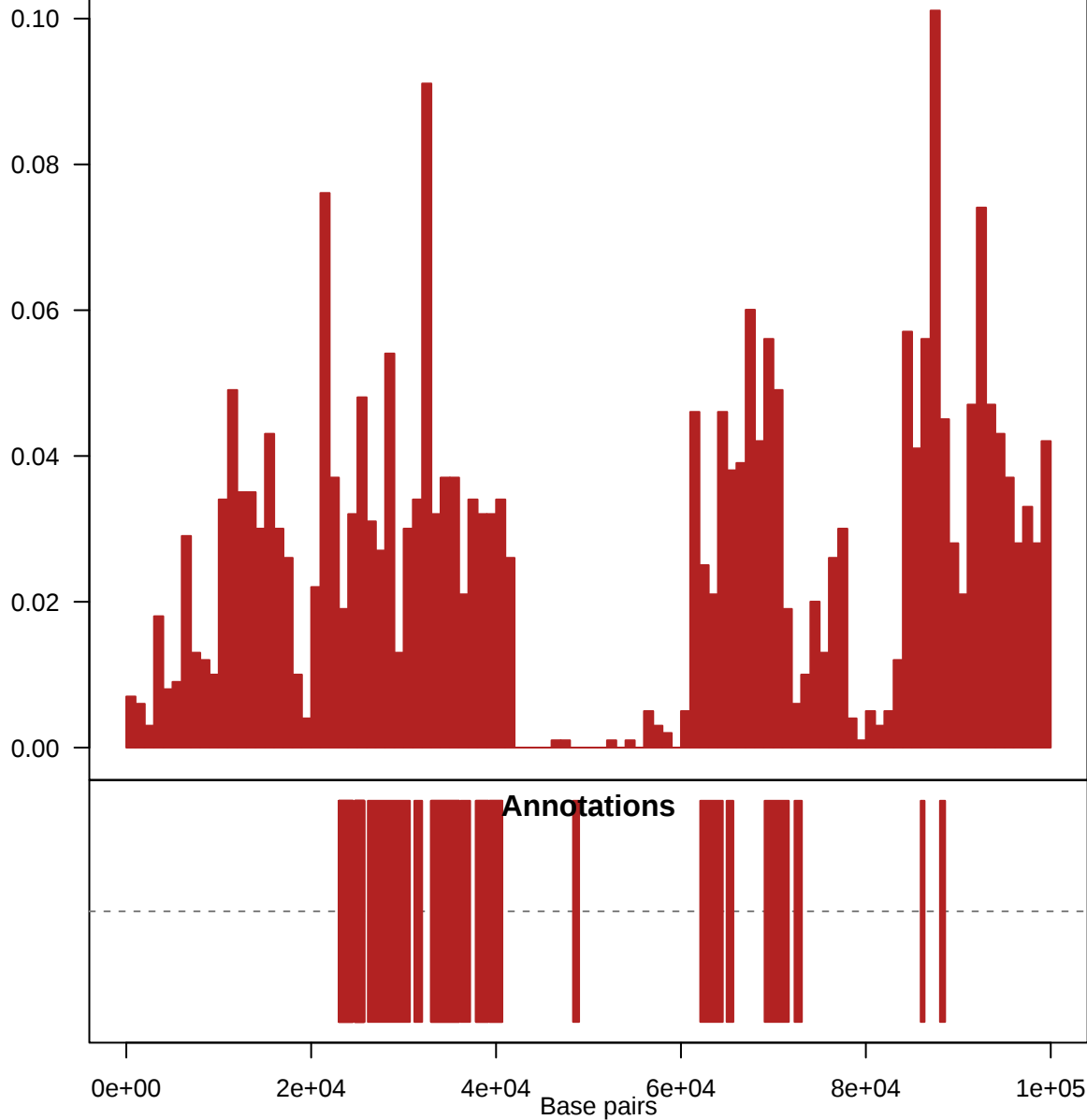
Supercontig

Variants per Site

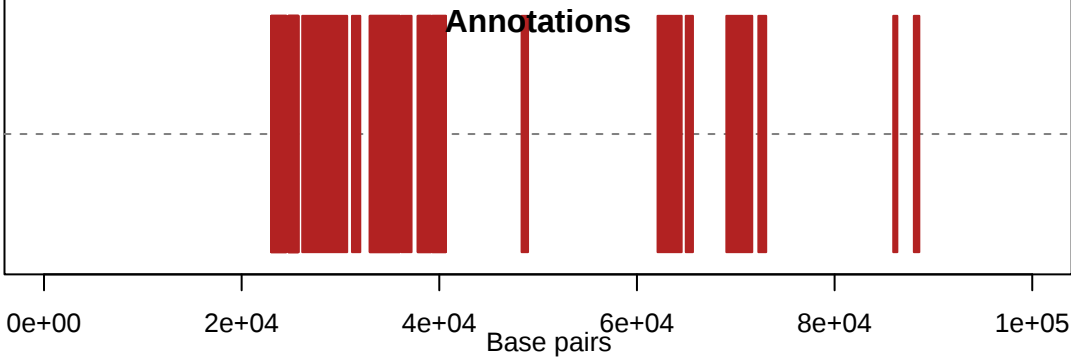


Supercontig

Variants per Site

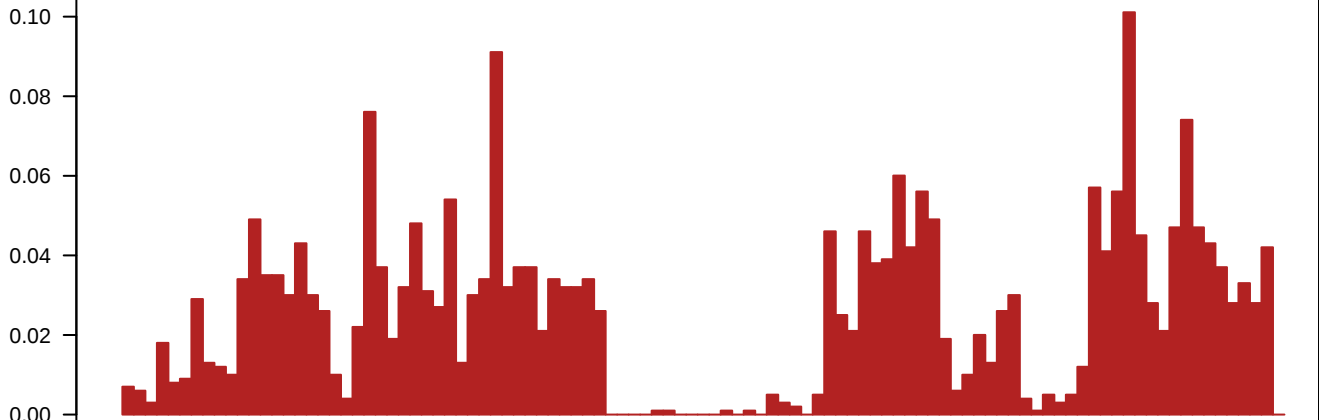


Annotations



Supercontig

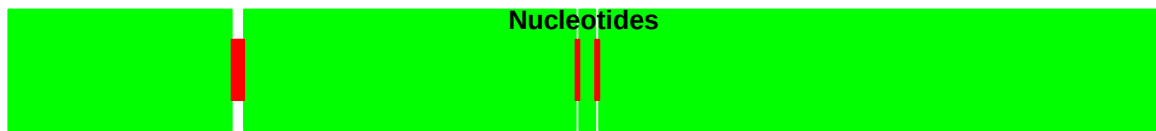
Variants per Site



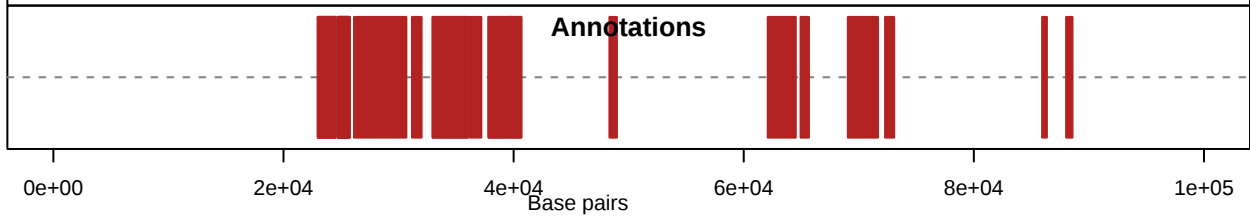
Nucleotide Content



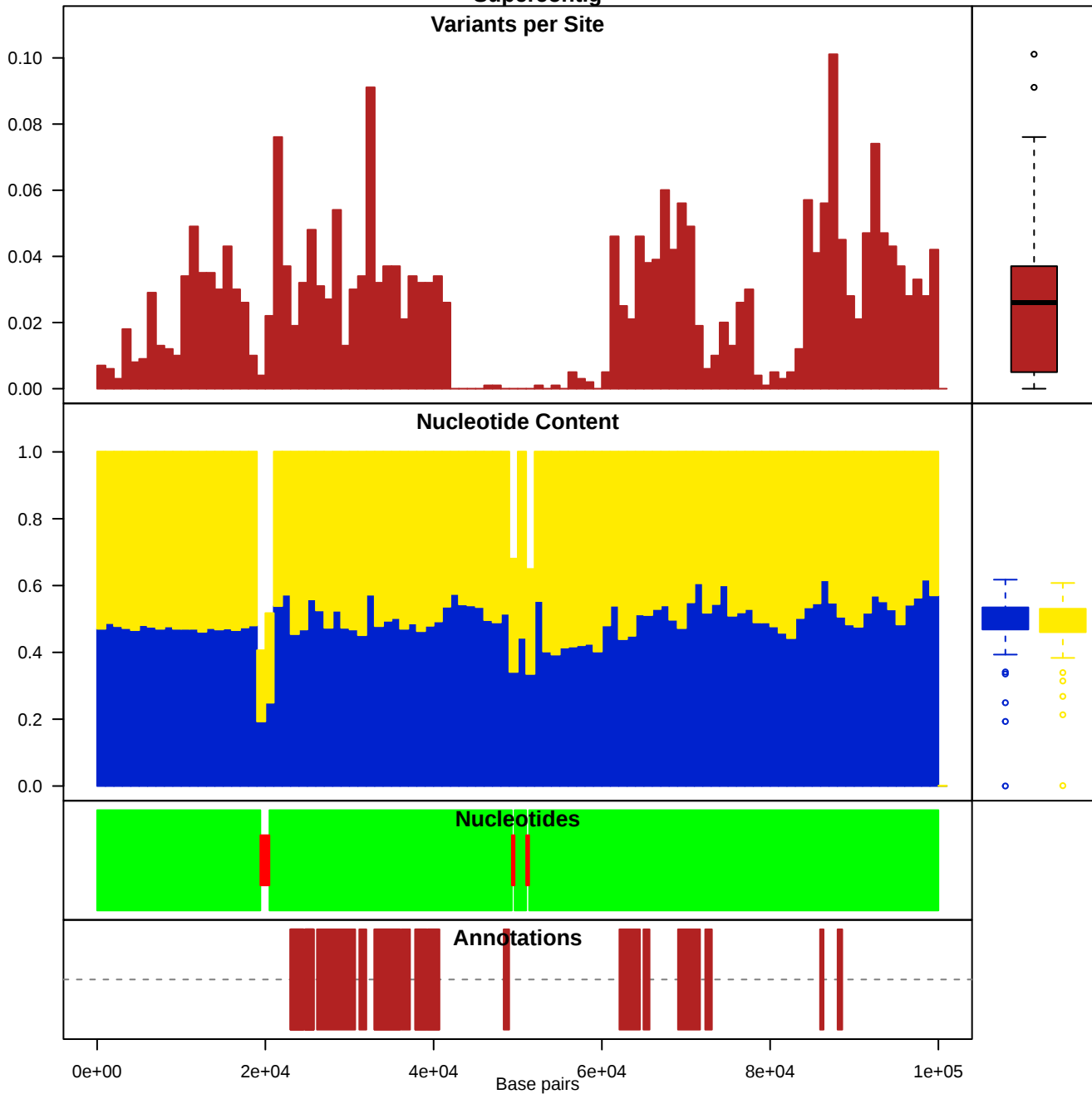
Nucleotides



Annotations

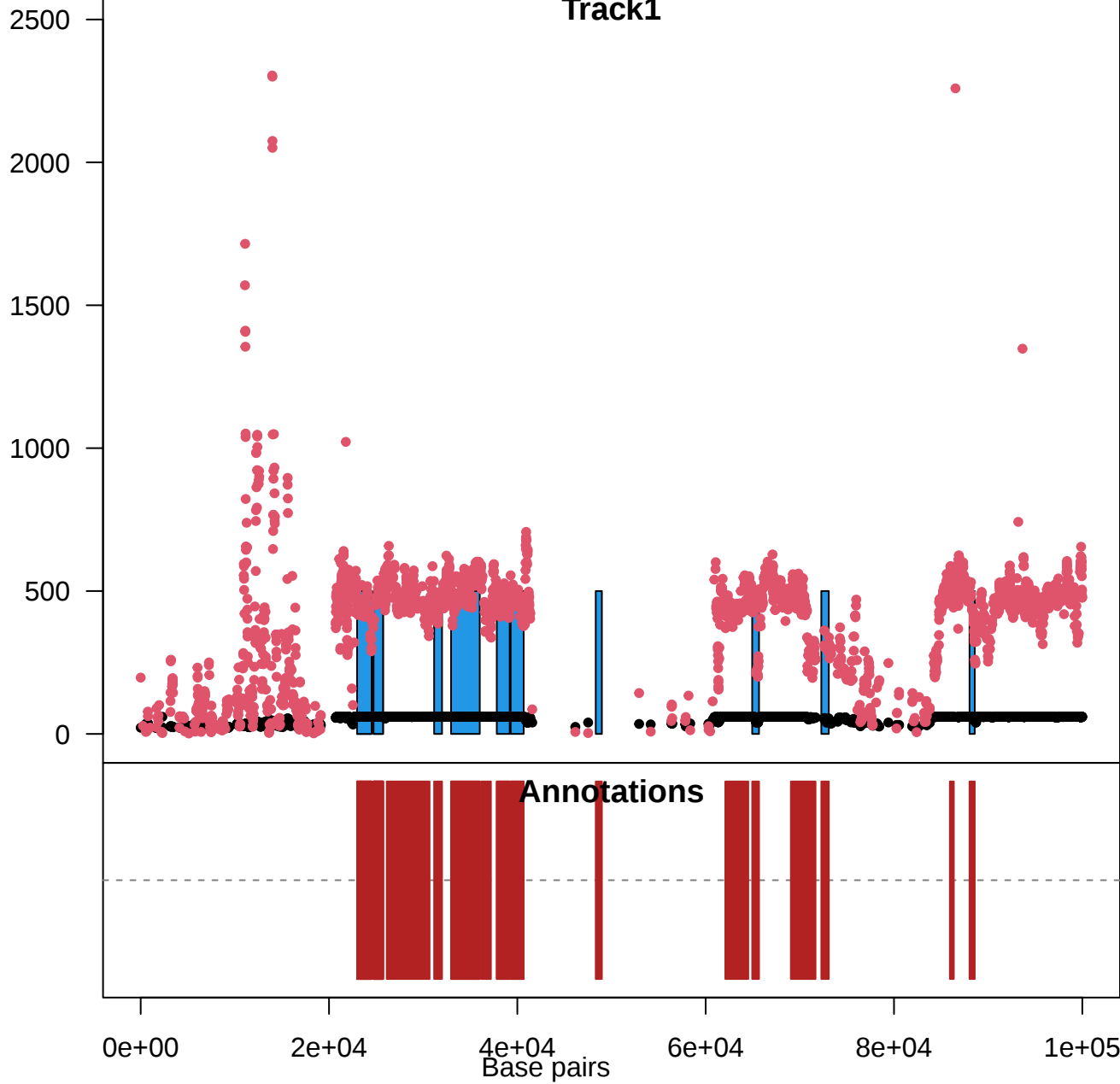


Supercontig



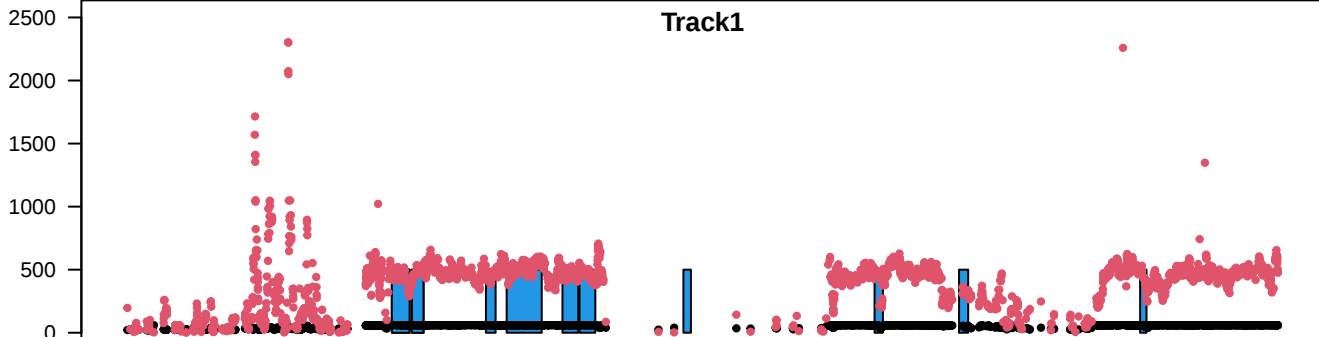
Supercontig

Track1

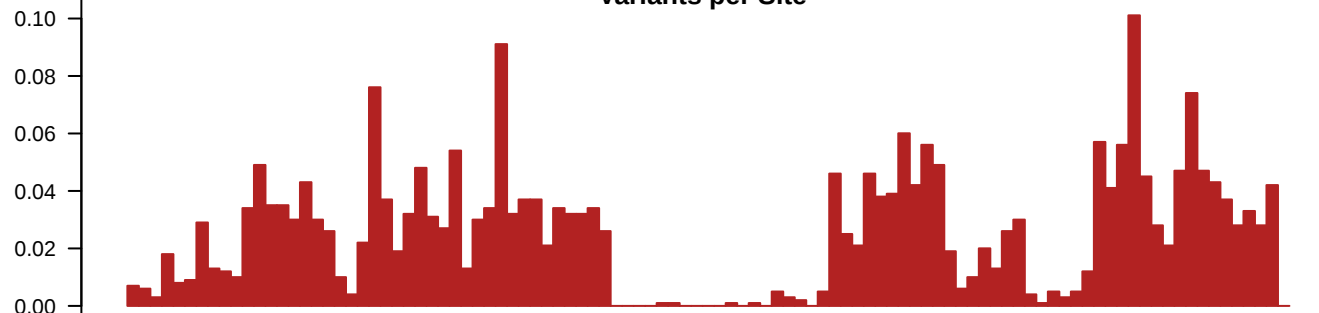


Supercontig

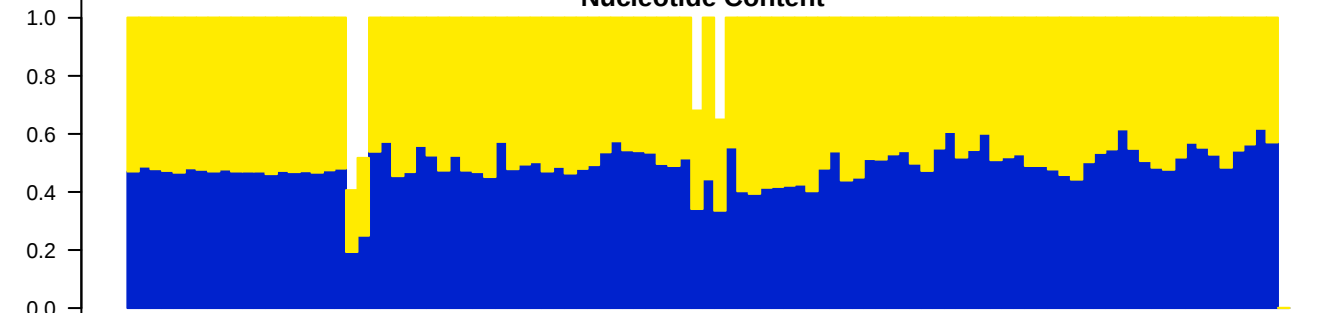
Track1



Variants per Site



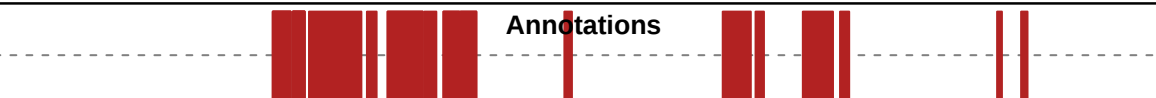
Nucleotide Content



Nucleotides



Annotations



0e+00

2e+04

4e+04

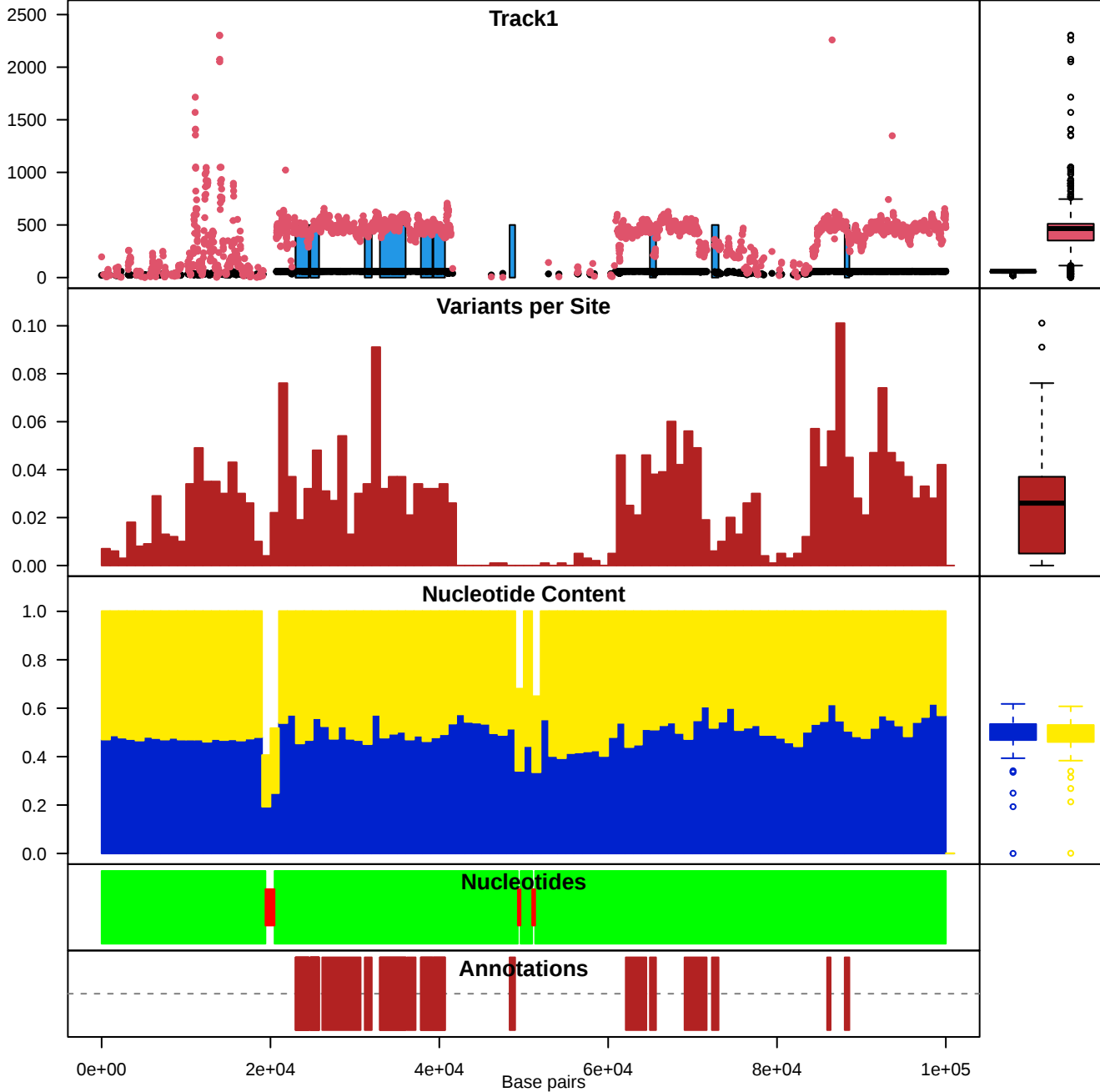
6e+04

8e+04

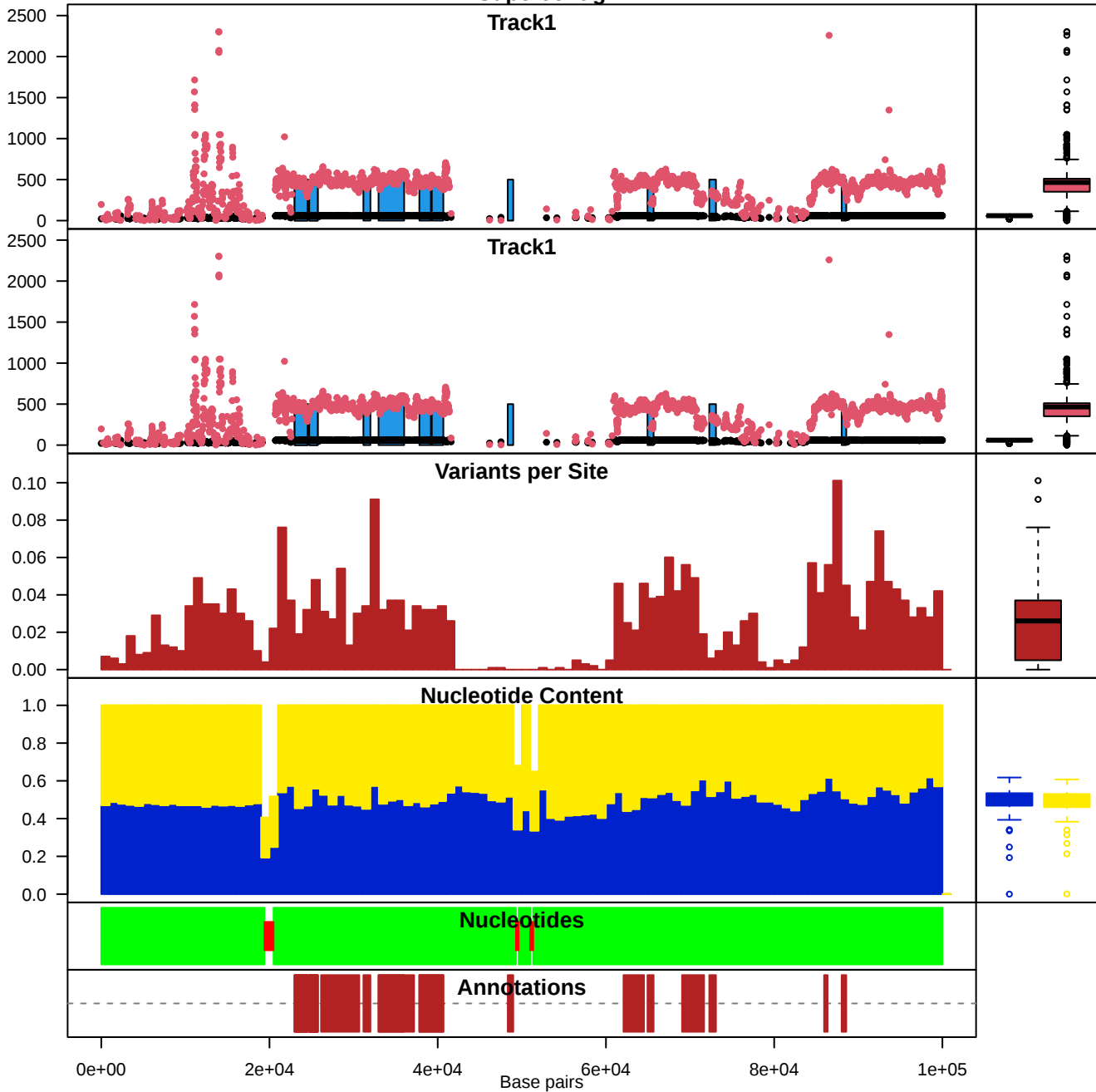
1e+05

Base pairs

Supercontig



Supercontig



Supercontig

