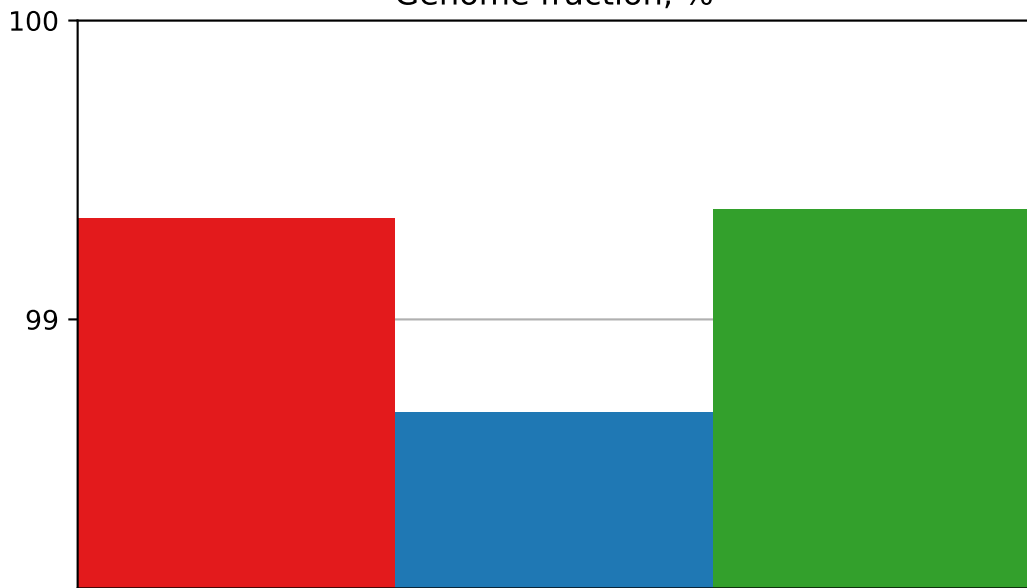


Genome fraction, %



SRR33190466_closedgap



SRR33190466.consensus



ref_SRR33190466