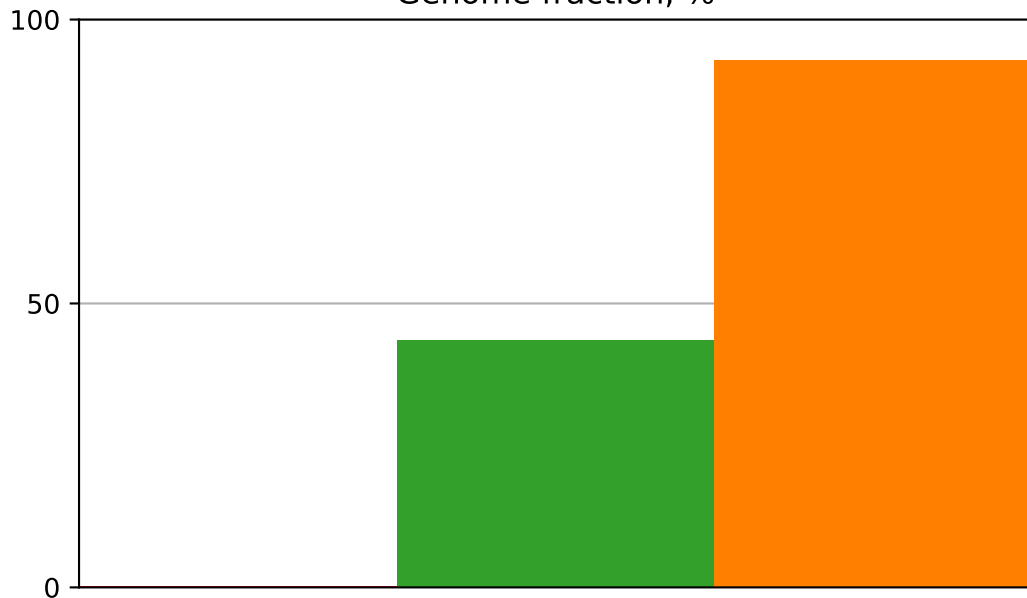


Genome fraction, %



sortie_reference1



sortie_sequencage1



sortie_sequencageserreur