Genomes/transcriptomes added= 28 replicates= 2
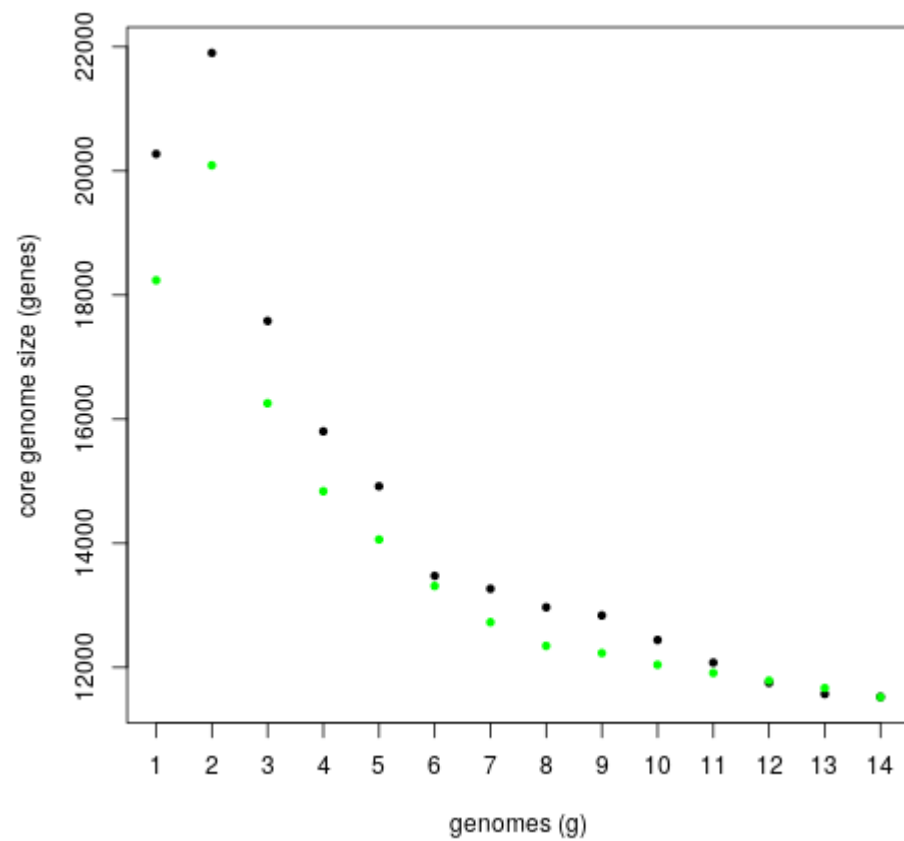
$coregenes(g) = 11175 + 13301 \, exp\left(\frac{-g}{3.66}\right)$

residual standard error = 848.08