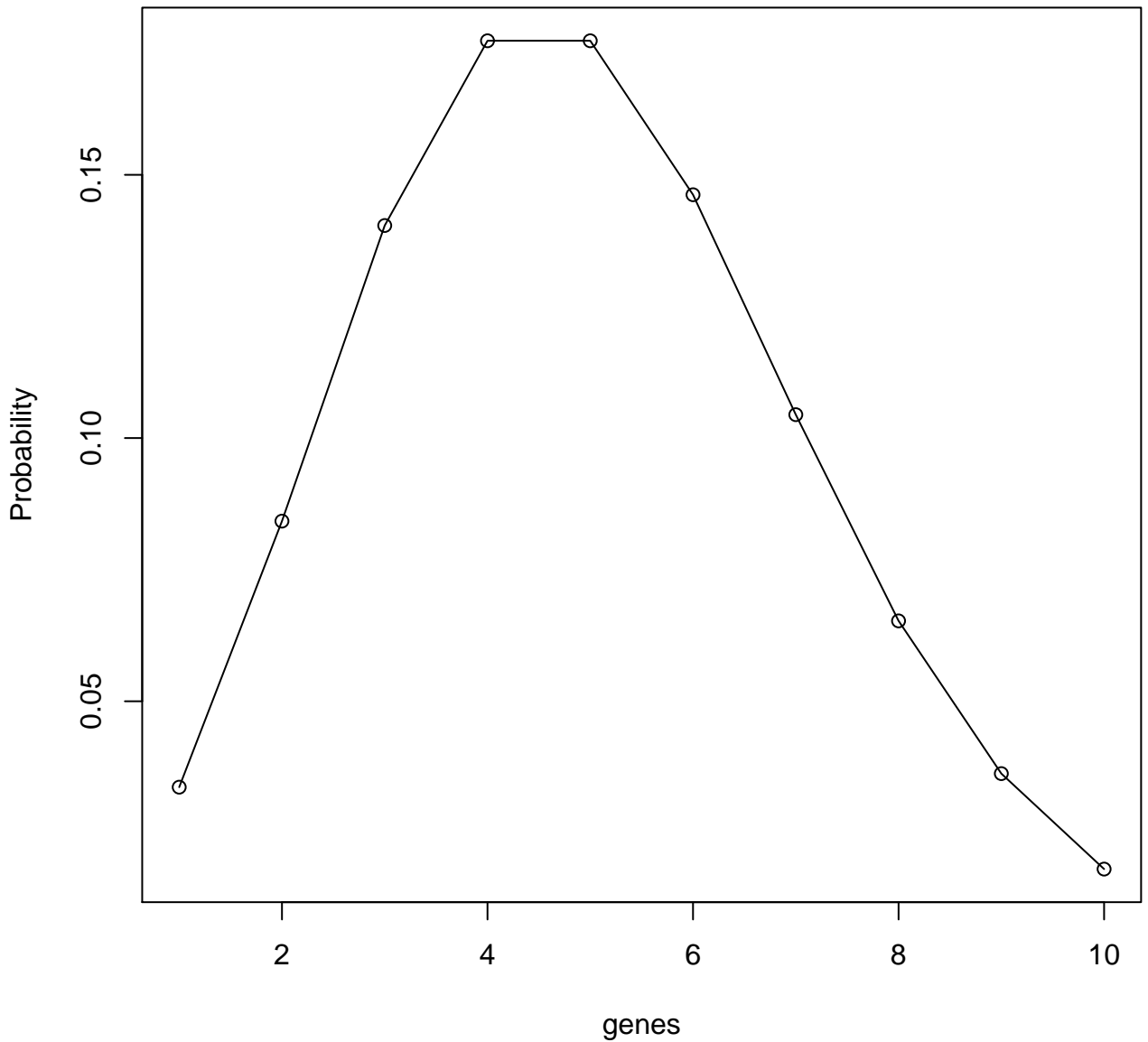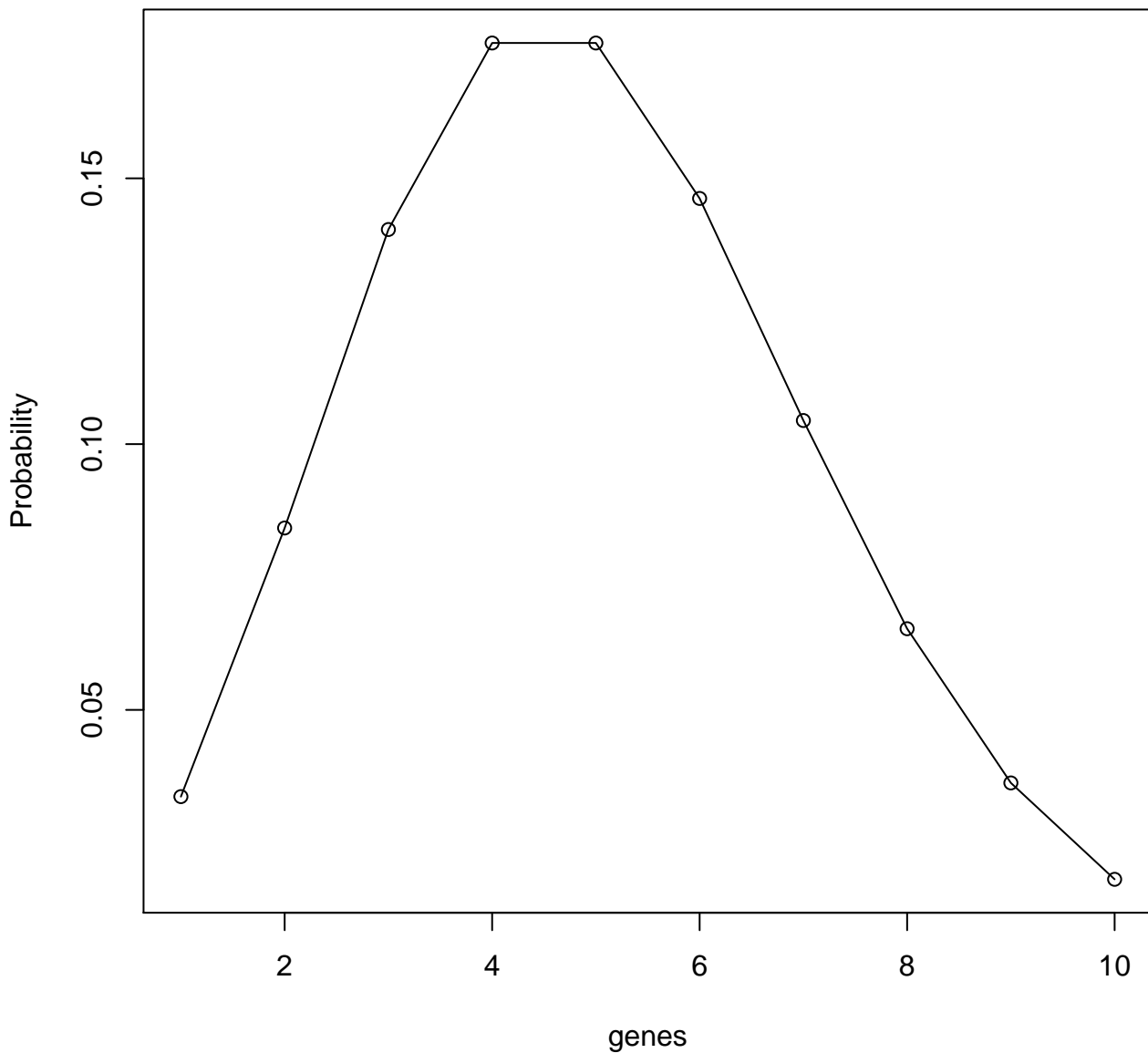# Parameters

genes
chr
nSpecies
averageEvents
minimalEventsOnBranch
averageInversionLength
averageDuplicationLength
averageDeletionLength
probInversion
probTranslocation
probDuplication
probDeletion
probFusion
probFission
probWGD
averageDeletionRateWGD
outDirectory
phyChroFormat
phyChroInstallationDirectory
save
nSimulations
seed

Inversion length theoretical distribution

Duplication length theoretical distribution

Deletion length theoretical distribution

Probability

genes