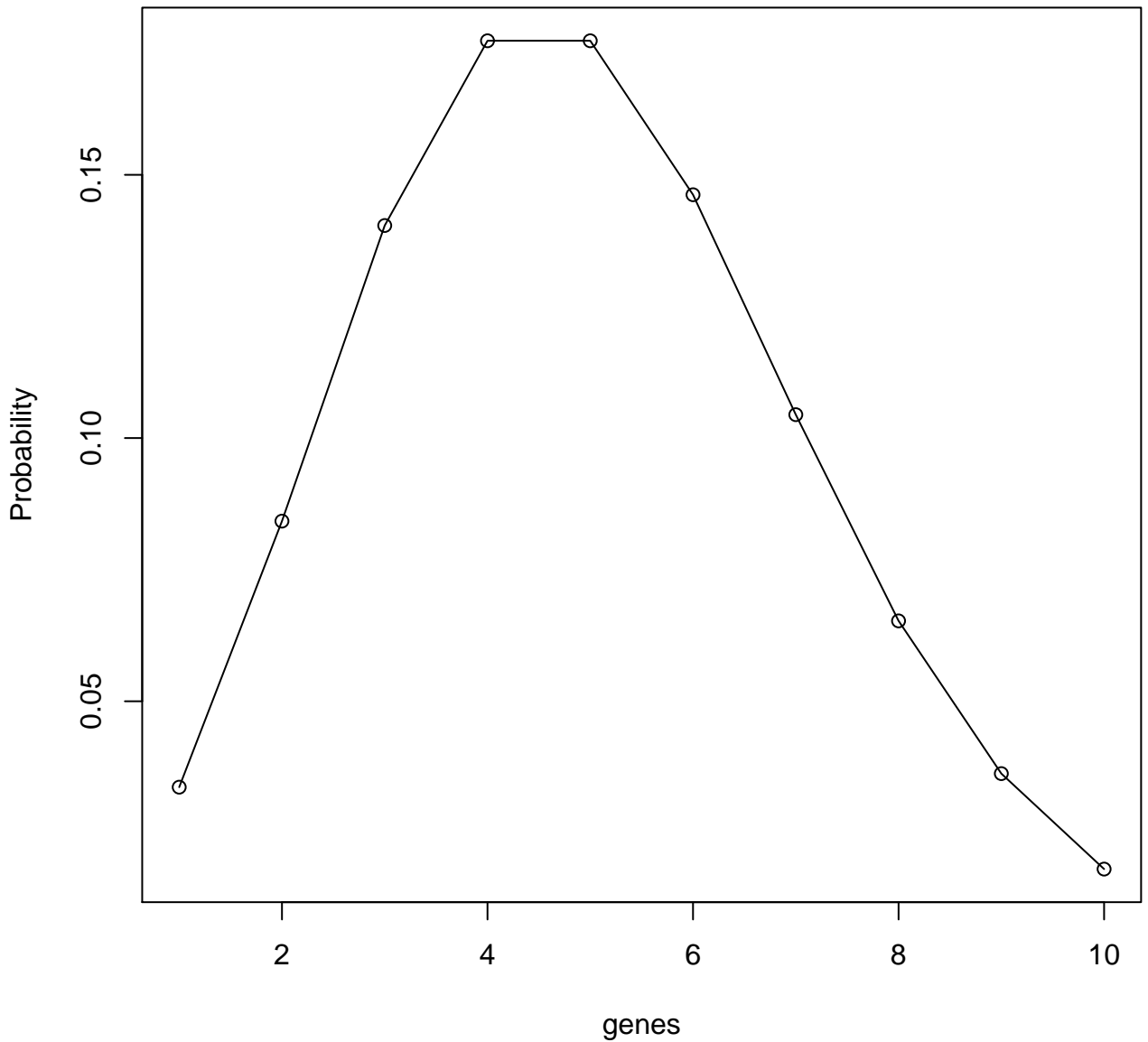


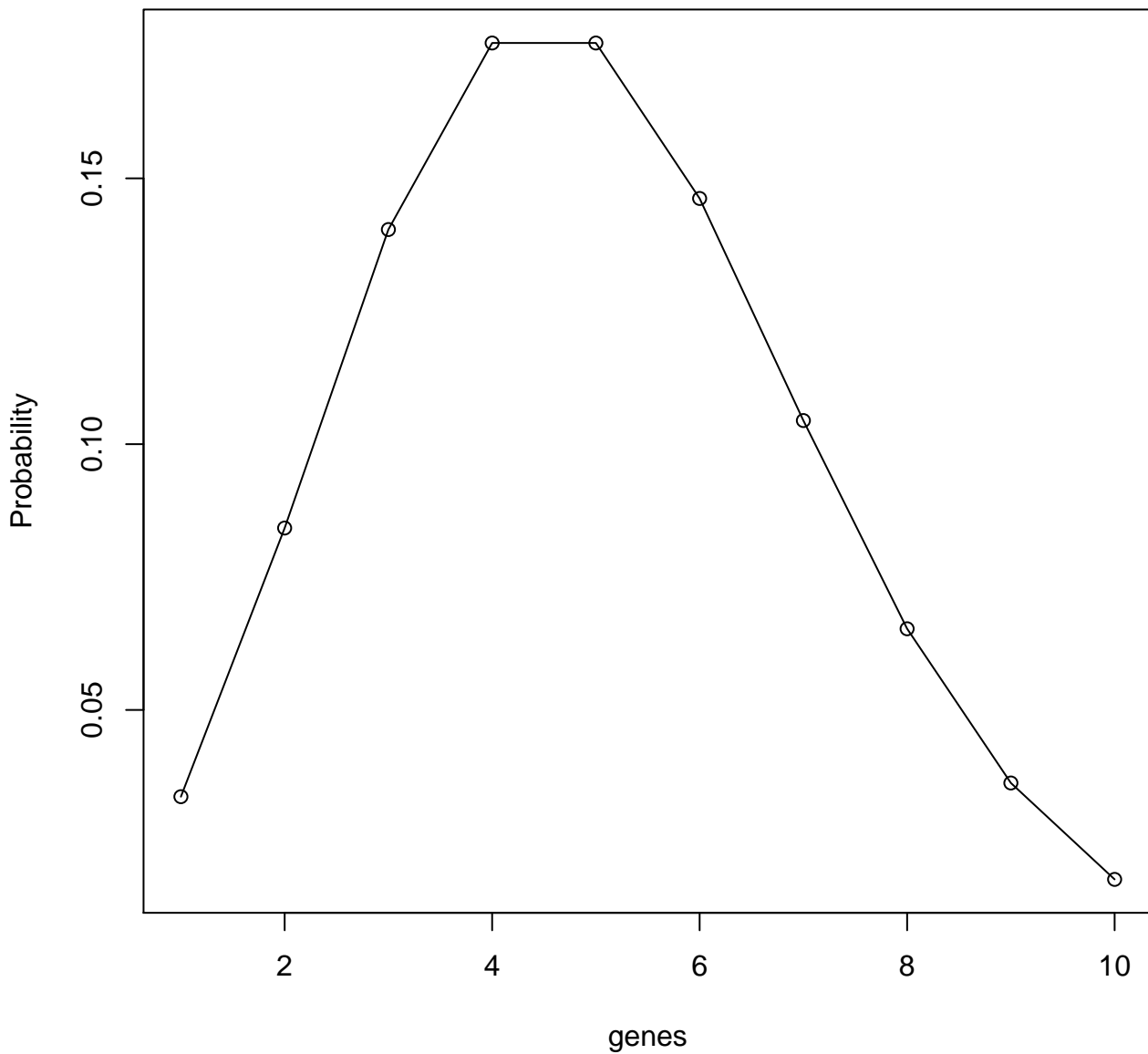
Parameters

genes	18000
chr	23
nSpecies	13
averageEvents	1000
minimalEventsOnBranch	1
averageInversionLength	5
averageDuplicationLength	5
averageDeletionLength	1
probInversion	0.6
probTranslocation	0.2979
probDuplication	0.05
probDeletion	0.05
probFusion	0.001
probFission	0.001
probWGD	1e−04
averageDeletionRateWGD	0.8
outDirectory	results_vert1
phyChroFormat	TRUE
save	TRUE
nSimulations	10
seed	0

Inversion length theoretical distribution



Duplication length theoretical distribution



Deletion length theoretical distribution

