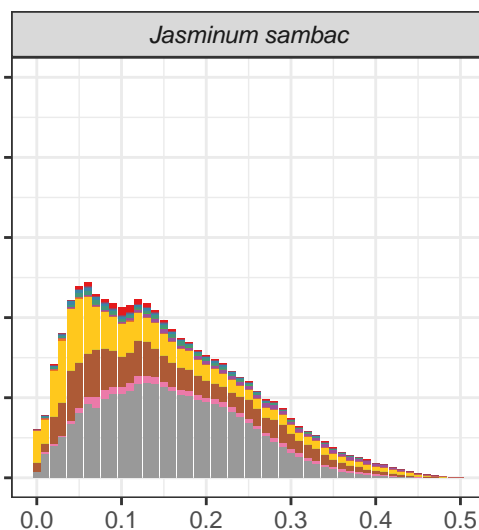
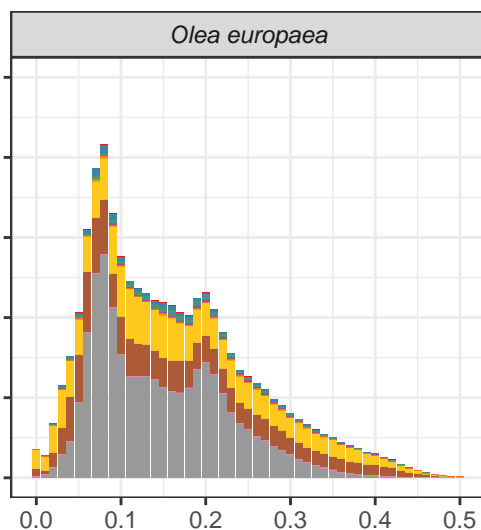
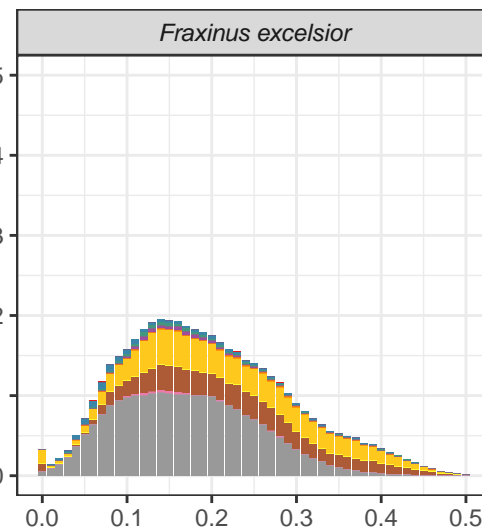
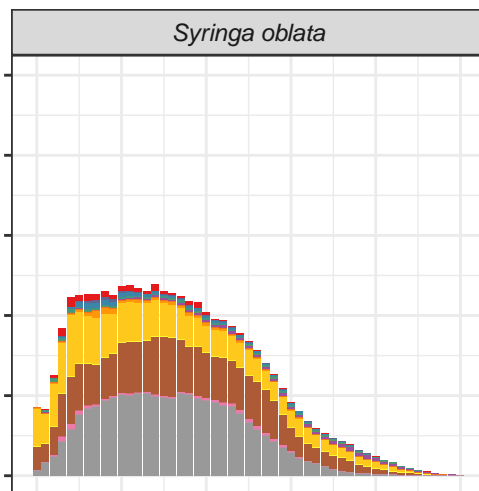
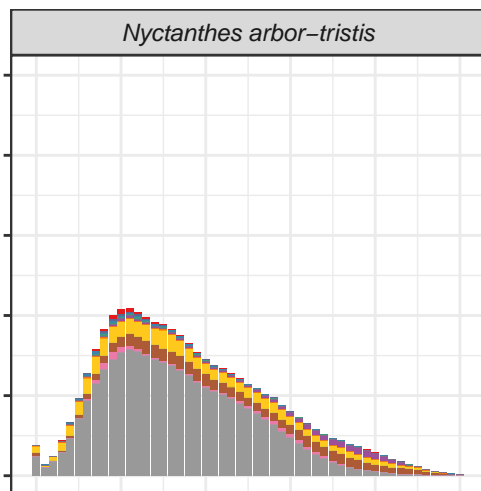


Genome coverage [%]



Kimura distance from TE family consensus sequence

Repeat Type

