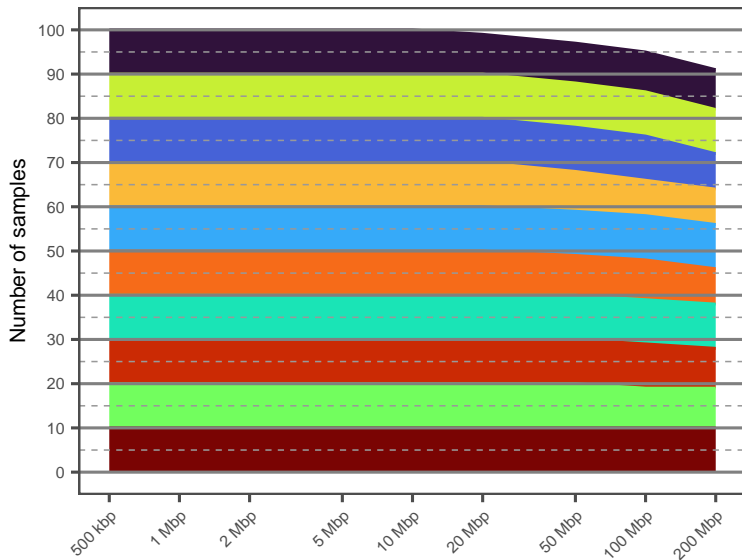
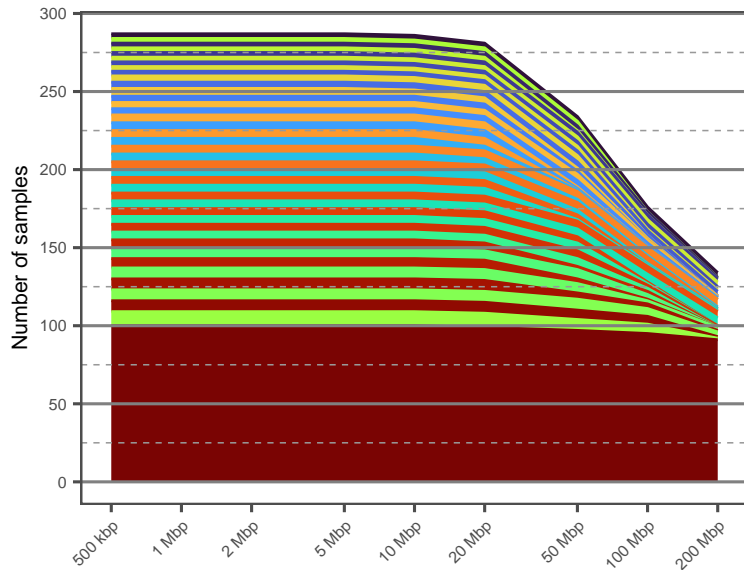


# Number of samples available for different data amounts

*Stigmaphyllon* species



Malpighiales genera



Post-cleaning base pairs available