Number of samples available for different data amounts
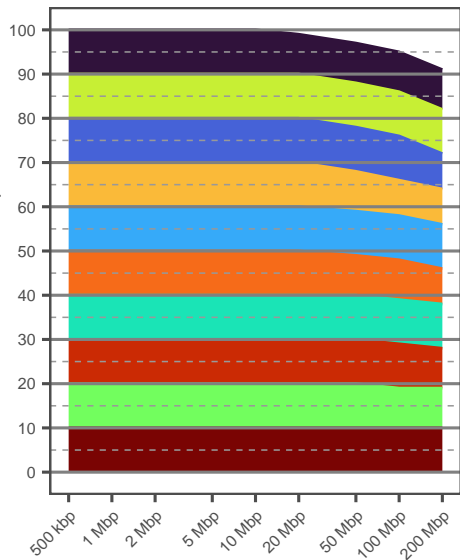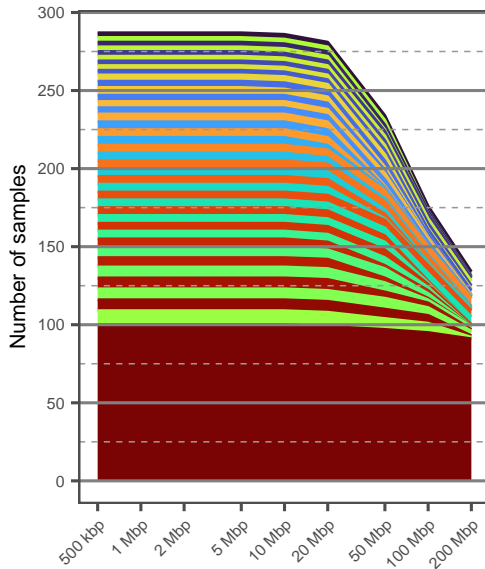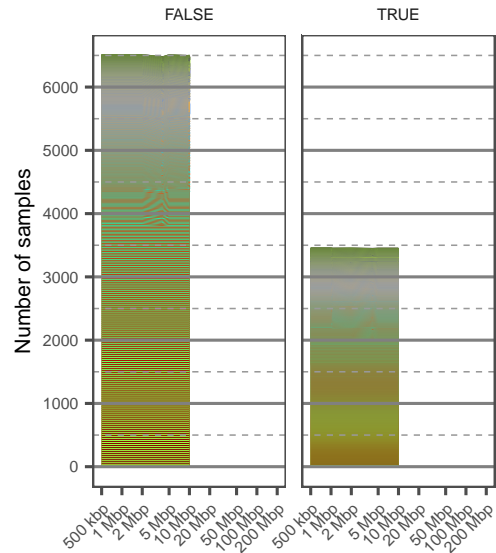
Stigmaphyllon species

Malpighiales genera

Eukaryotic families

FALSE

TRUE

Number of samples

Post−cleaning base pairs available