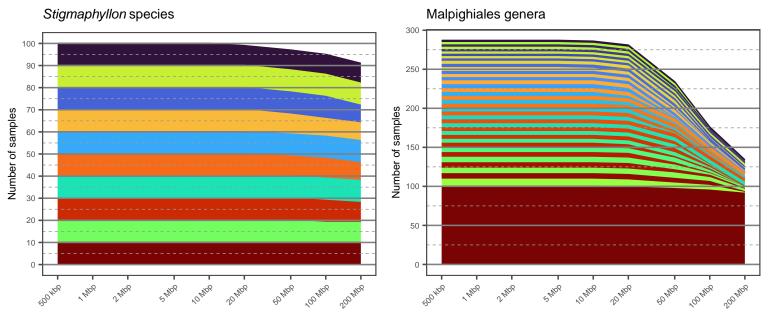
Number of samples available for different data amounts



Post-cleaning base pairs available