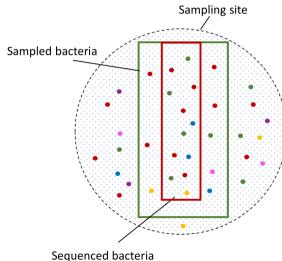
b)
Sampling site
Number of si



Number of sequences (Sequencing depth)

