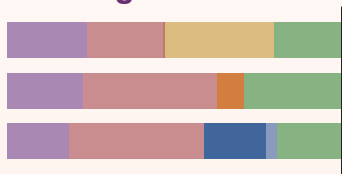
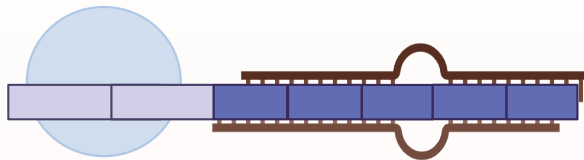


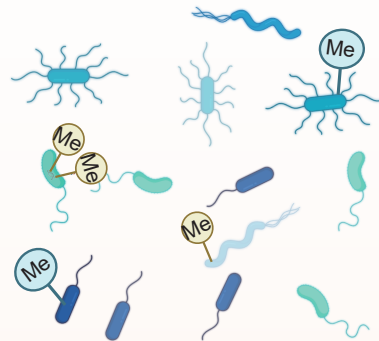
Assembly circled genome



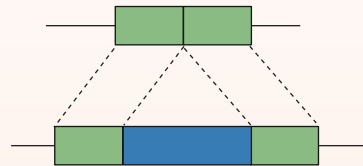
Rapid identification of community structure



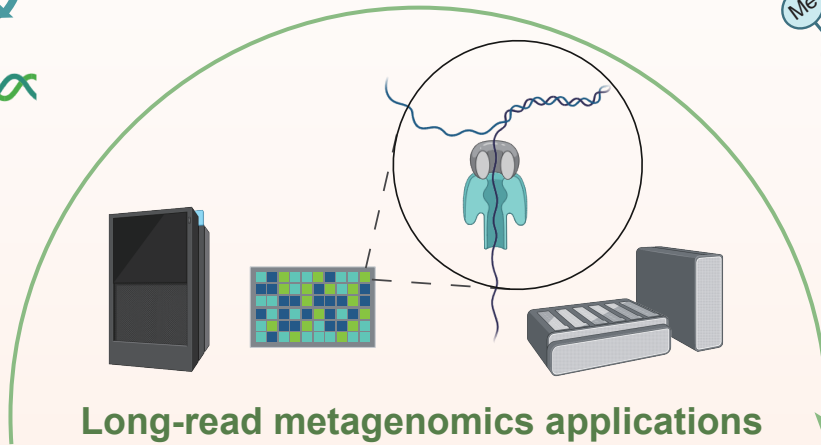
Predict gene and elements



Epigenetic and association



Microbial variation and population heredity



Long-read metagenomics applications