# Galaxy for long-read ONT data analysis and public education

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### Nanopore data analysis

#### Advantages

Realtime sequencing Simplified and less ambiguous genome assembly Capability to span repetitive genomic regions Identification of large structural variation

#### **Trends**

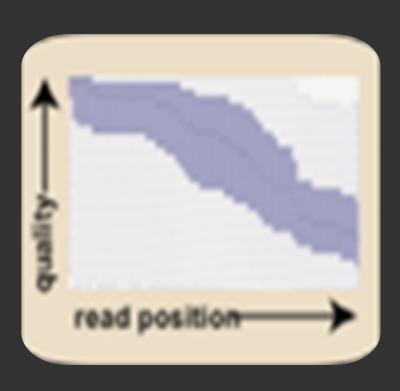
More accessible and easier/simpler: higher throughputs Rapid and constant tool development and upgrade of algorithms and software

#### Challenges

Bioinformatics data analysis

### NanoGalaxy: A Galaxy toolkit, workflows & training





Polishing, QC & preprocessing Porechop Filtlong Nanopolish Poretools



**Genome Assembly** Minimap2 Miniasm Racon Flye Unicycler



Wtdbg2 Canu Mapping Minimap2

GraphMap **Visualisation** Nanoplot

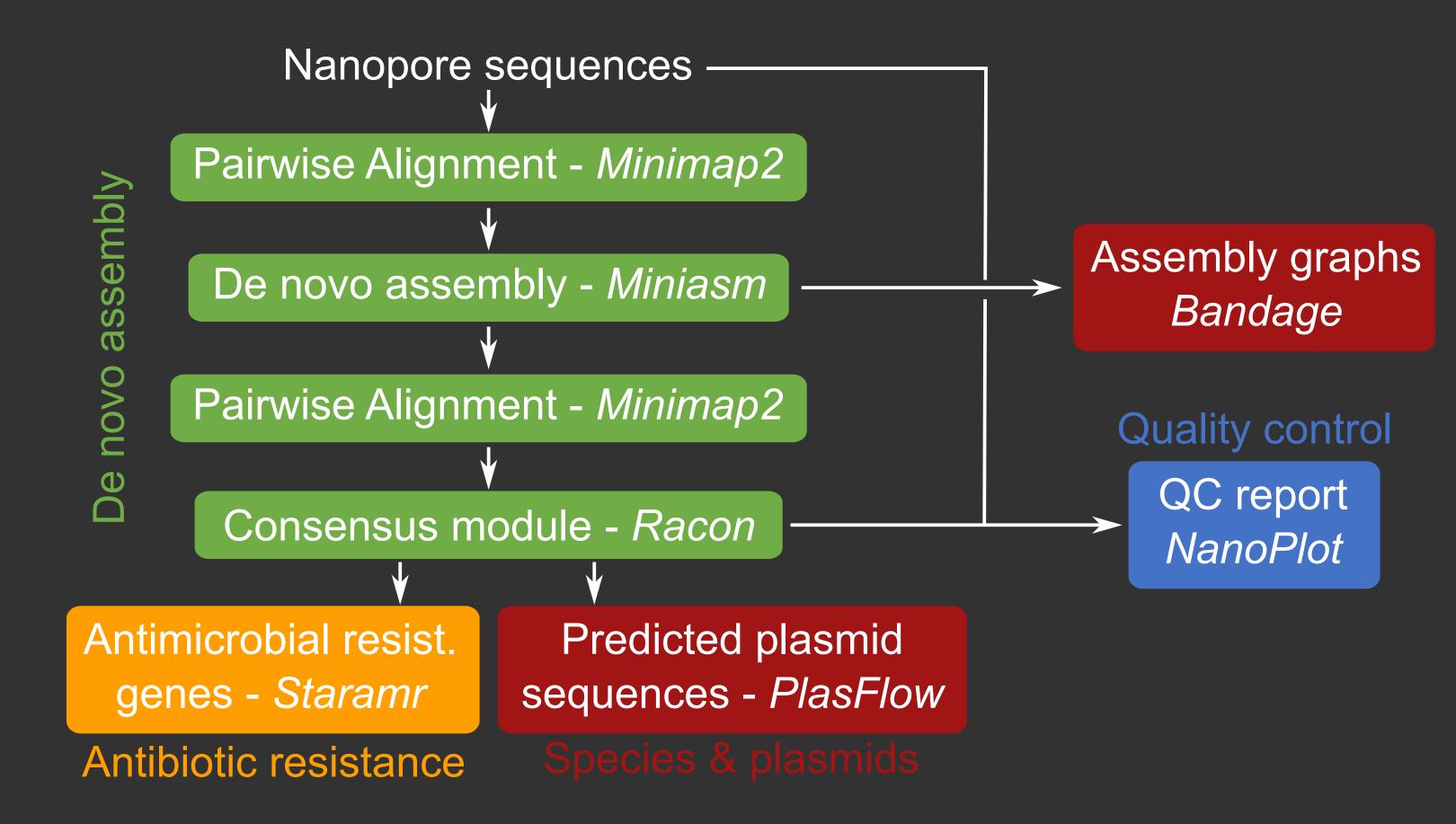
Bandage



**Taxonomy & metagenomics PlasFlow** Staramr Kraken2

### Worflows

Genome assembly using Flye for highly repetitive genomes Genome assembly using Unicycler (Illumina and ONT) for K. pneumoniae Antibiotic resistance detection



### Training material

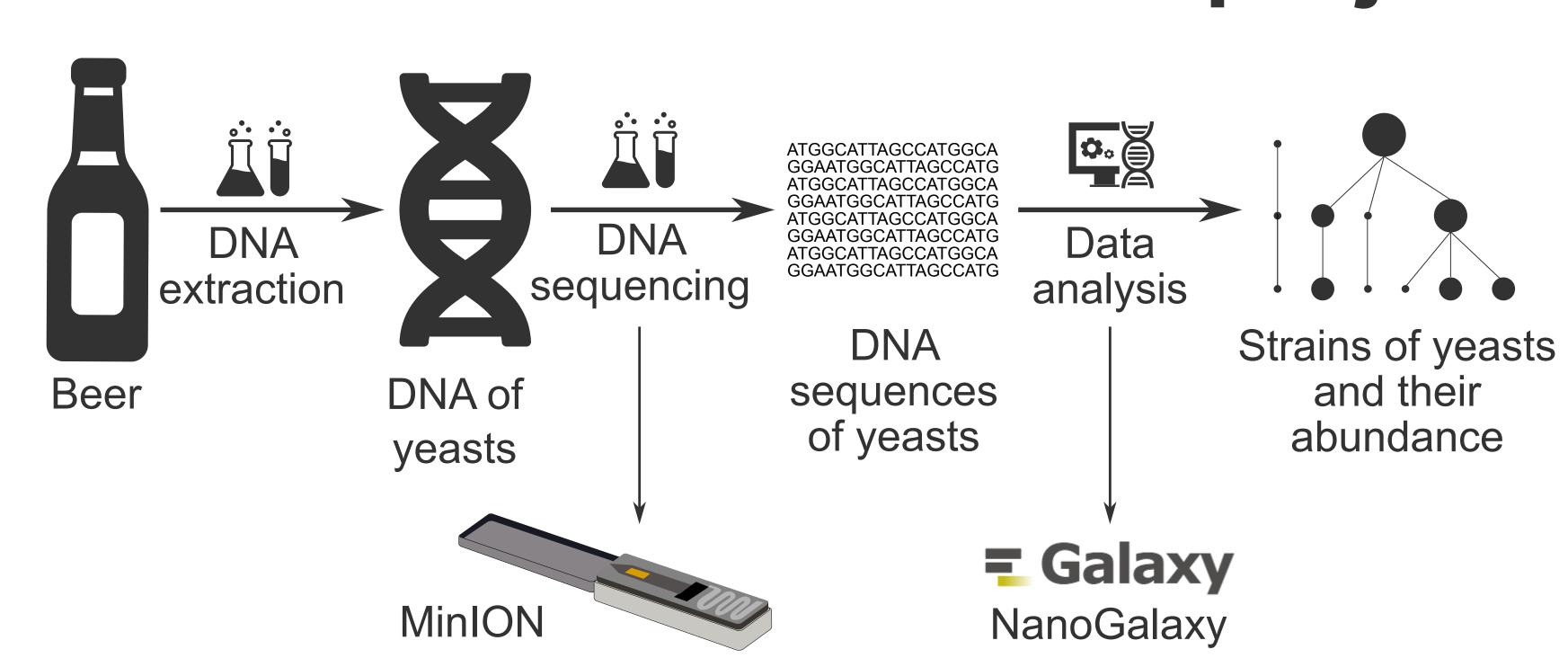
Assembly of plasmids and determination of antibiotic resistance following the previous workflow

Available on training.galaxyproject.org

### Application to public education: the BeerDEcoded project

Workshops where pupils, students and citizens extract, sequence and analyze the "DNA of beer" to bring them in contact with molecular biology, data-analysis, and open science.





## nanopore.usegalaxy.eu

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