

Genomic & Transcriptomic applications for microorganisms studies

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GeT (Genome & Transcriptome) is a cutting-edge multi-site core facility in Toulouse (France), providing the academic and private scientific communities with technologies and expertise in the area of genomics and transcriptomics.

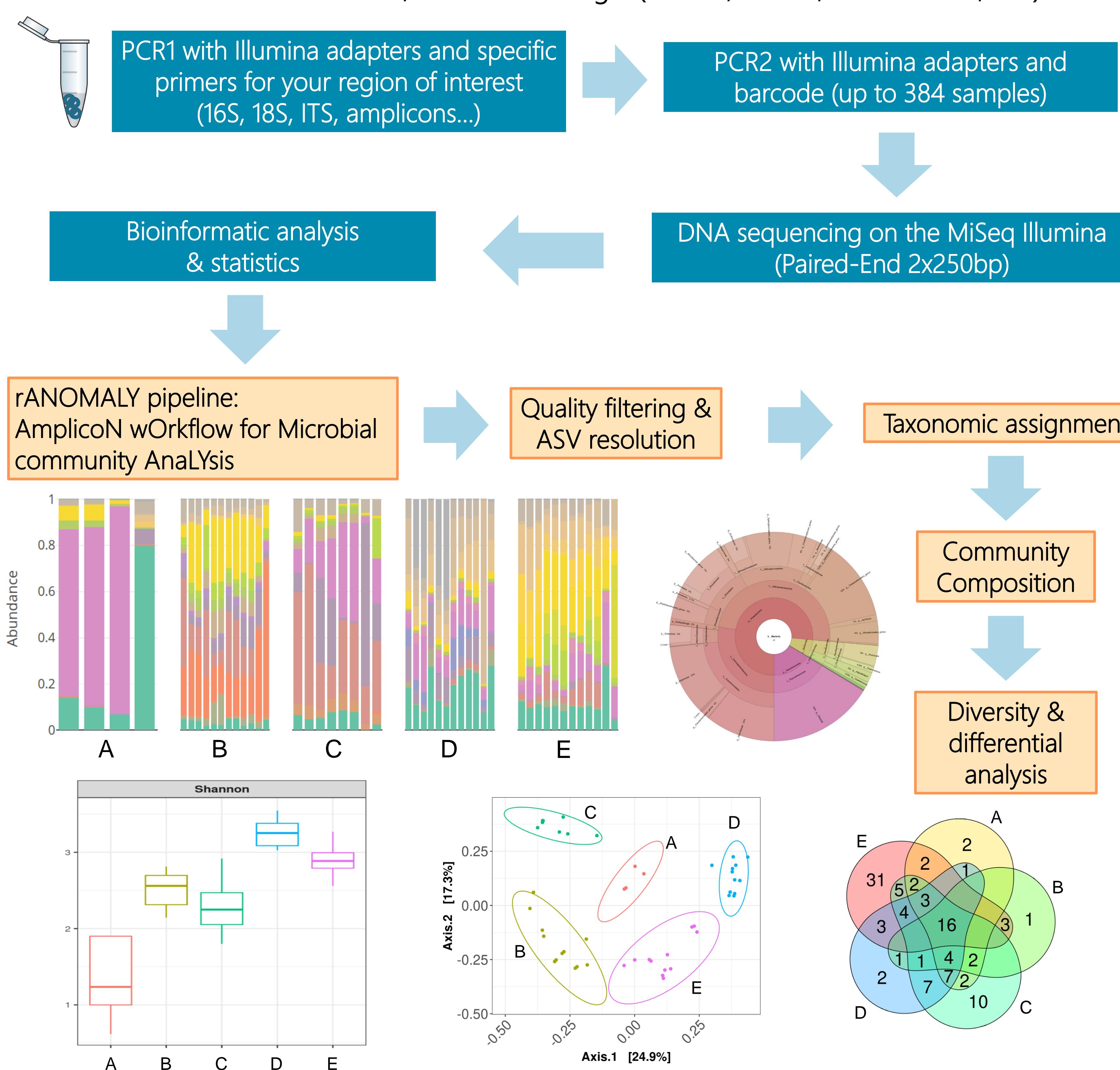
GeT-Biopuces is one of the four GeT sites. Located at INSA Toulouse, within the Toulouse Biotechnology Institute (TBI), the platform is ISO9001 & NFX50-900 certified and specialized in biotechnology and in the study of micro-organisms. Its missions are:



Applications	Technologies	Input	Bioinfo Biostats analysis
Amplicons, 16S, 18S, ITS Small genomes (bacteria and yeast) RNAseq with ribodepletion	Illumina MiSeq ThermoFisher GeneStudio S5	10 ng DNA	✓
		10 ng DNA	✓
		500 ng total RNA	✓
DNAseq Plasmids/Fosmids	Oxford Nanopore MinION	1 µg DNA	✓
		100 ng DNA	✓
		1 ng DNA	✓
Copy Number Variation Microbe detection Gene expression	Qiagen QiaQuity One	< 1 ng DNA	✓
		1 pg total RNA	✓
		1 pg total RNA	✓

Metagenomic Sequencing

By using our comprehensive metagenomic sequencing and analysis pipeline, start analyzing DNA extracted from complex samples of interest without the need to cultivate micro-organisms. This approach provides you with a global view of the diversity and functions of microbial communities present in a given environment. Explore the composition, abundance and interactions in a microbiota, whatever its origin (human, animal, environment, etc.).



Digital PCR

This flexible, solid partitioning-based new technology allows you to split your sample into several thousands elementary PCR reactions (8,500 or 26,000 partitions), to search for rare events, work on difficult samples rich in inhibitors, or precisely determine differences as low as 10% between samples. No need for standards, quantification is absolute and up to five different targets can be multiplexed.

