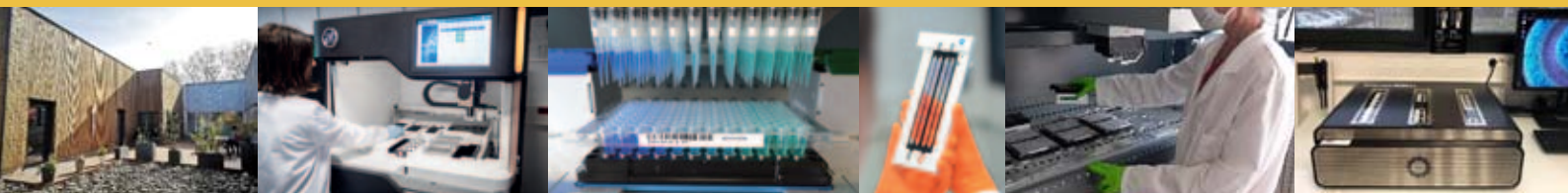
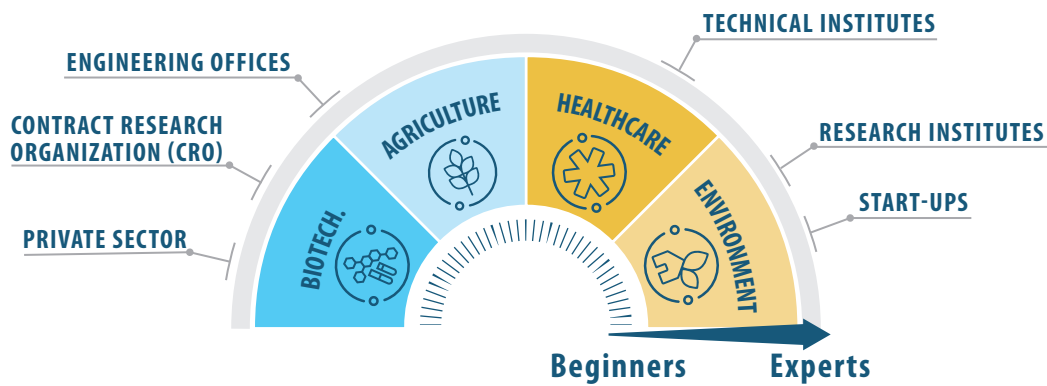


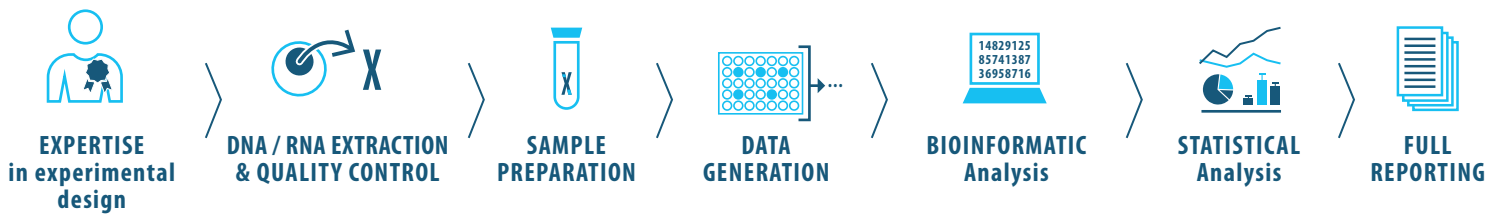
GeT-IT Services : Expertise for your needs in genomic analysis



CUSTOMERS AND MARKETS



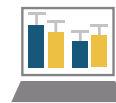
COMPLETE WORKFLOW from samples to data analysis



SOLUTIONS we deliver



WHOLE GENOME SEQUENCING
For *de novo* or re-sequencing applications



BIOINFORMATIC & BIostatistic SOLUTIONS
Delivering comprehensive and high quality data analyses



TRANSCRIPTOMICS
Differential expression or annotation



METAGENOMIC SOLUTIONS
Analyzing diversity and functions of microbial communities



GENOTYPING BY SEQUENCING
Allegro® Targeted genotyping



TRAINING COURSES
Targeted interventions and supports



RESEARCH AND DEVELOPMENT
Support in setting up collaborative project



EXPERTISE AND ADVICE

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Expertise for your needs in genomic analysis

**AGILITY & CLOSE
CUSTOMER RELATIONSHIP**

SHOTGUN METAGENOMICS



MICROBIAL COMMUNITY STRUCTURE

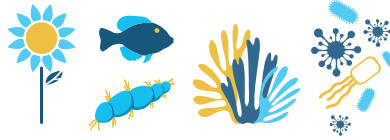
Diversity and abundance

FUNCTIONAL ANALYSIS

Gene prediction
and metabolic pathways

DIVERSITY OF ANTIBIOTIC RESISTANCE GENES

LONG READ SEQUENCING



COMPLETE AND ACCURATE VIEWS OF ALL TYPES OF GENOMIC VARIATION

HIGH-QUALITY REFERENCE GENOMES

IDENTIFICATION OF GENES
associated with toxicity, virulence,
and antimicrobial resistance

GENOTYPING BY SEQUENCING

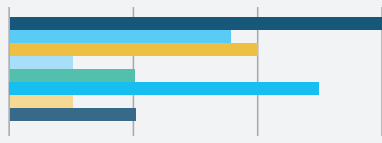


DISCOVERY OF GENOTYPE-PHENOTYPE ASSOCIATIONS

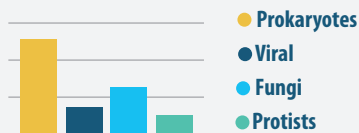
**ASSESSMENT OF
POPULATION STRUCTURE
& PATTERNS OF ADAPTATION**

CHOOSE YOUR LEVEL OF ECOSYSTEM CHARACTERIZATION

Species level



Domain

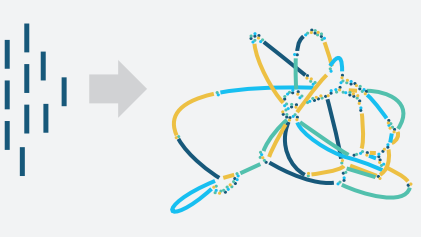


Functional profiles

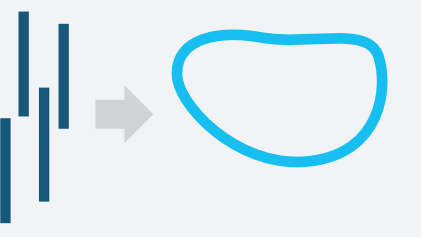


CREATE EXCEPTIONNAL DE NOVO ASSEMBLY

Short read assembly



Long read assembly



INTERROGATE MASSIVE TARGETED SNP USING NGS

Genetic relationship
among individuals
based on SNP datasets

