



# High Throughput Sequencing Facility



02/12/2022

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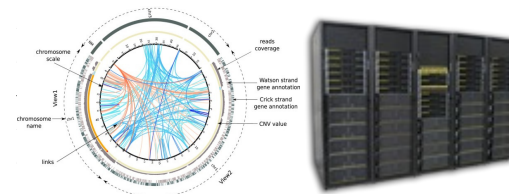
Molecular  
biology



Sequencing



Bioinformatics



- Created at the CGM in February 2010 (IMAGIF)
- I2BC since 2015
- Open to both academic and industrial users, whether in France or abroad

**IBiSA** • Infrastructures  
en Biologie  
Santé et  
Agronomie

 **FRANCE  
GENOMIQUE**



# Services in molecular biology

**Molecular  
biology**

Sequencing

Bioinformatics

## Illumina library preparation

- DNA-seq, ChIP-seq
- RNA-seq
- Small RNA-seq, miRNA-seq
- Single cell sequencing (10X genomics)



## ONT library preparation

- DNA-seq
- RNA-seq (direct, cDNA)
- tRNA sequencing
- Cas9 enrichment
- ULR



**Custom-made libraries if needed**

# Sequencing platforms

Molecular  
biology

**Sequencing**

Bioinformatics

Versatile set of equipment:



Illumina NS 550



Illumina NS 2000



ONT GridION



ONT PromethION P2 solo  
(spring 2023)

# Bioinformatics services

Molecular  
biology

Sequencing

**Bioinformatics**

- Differential gene expression
- Genome assembly
- Genome annotation
- SNPs, indels, rearrangement detection
- DNA/RNA modifications (ONT)
- TE detection (ONT)
- **On-demand analyses**

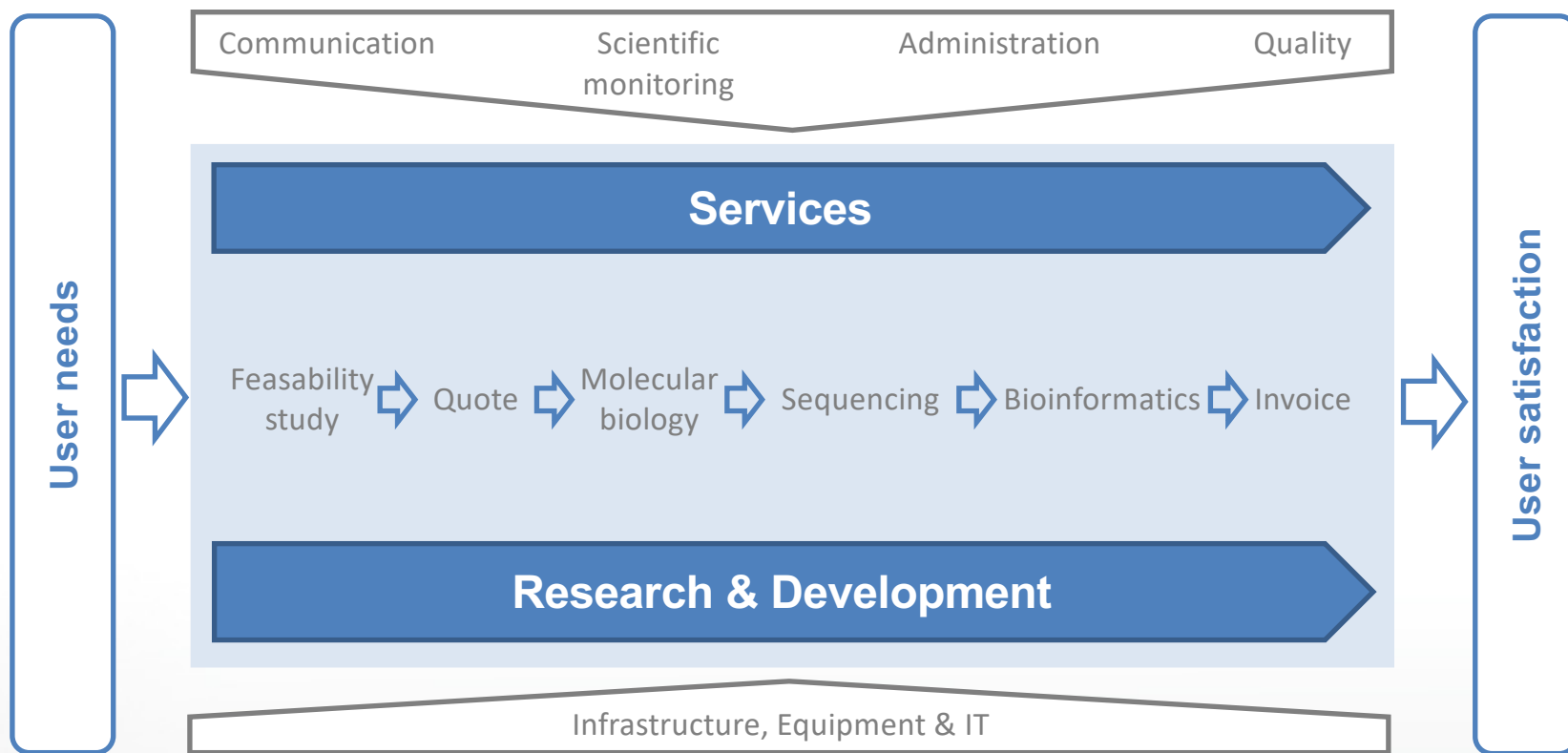


I2BC cluster

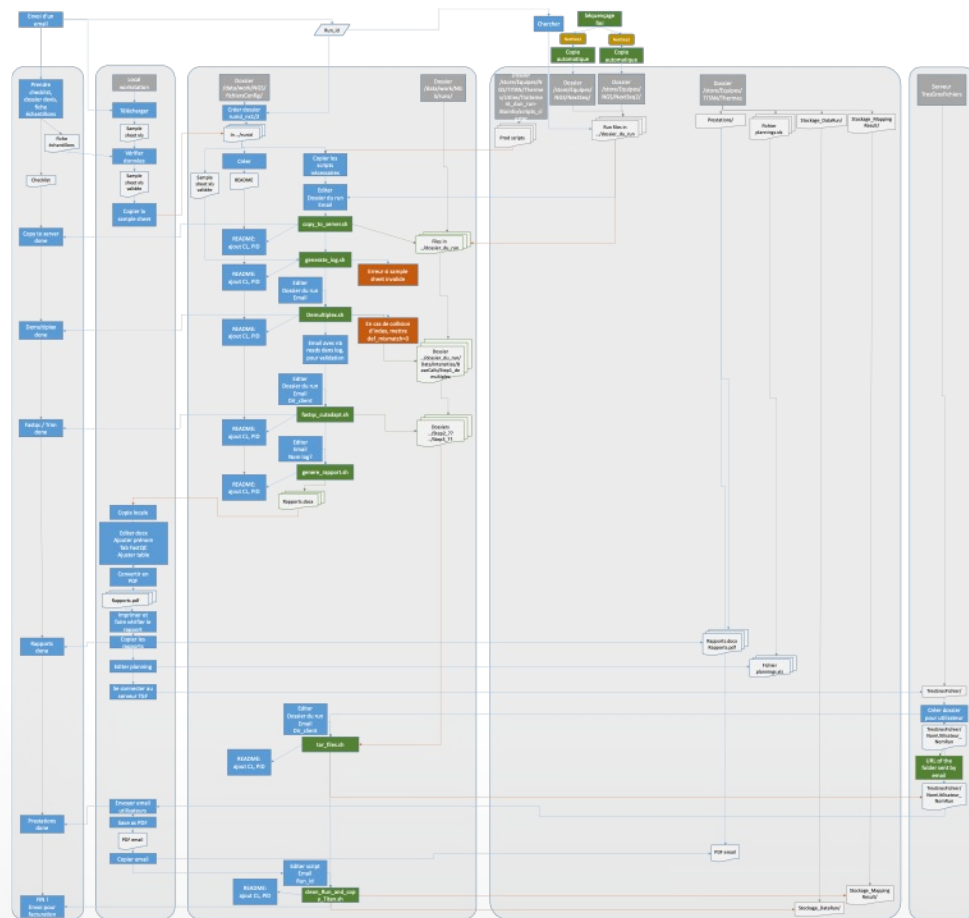


GPU

# « Theoretical » data flow

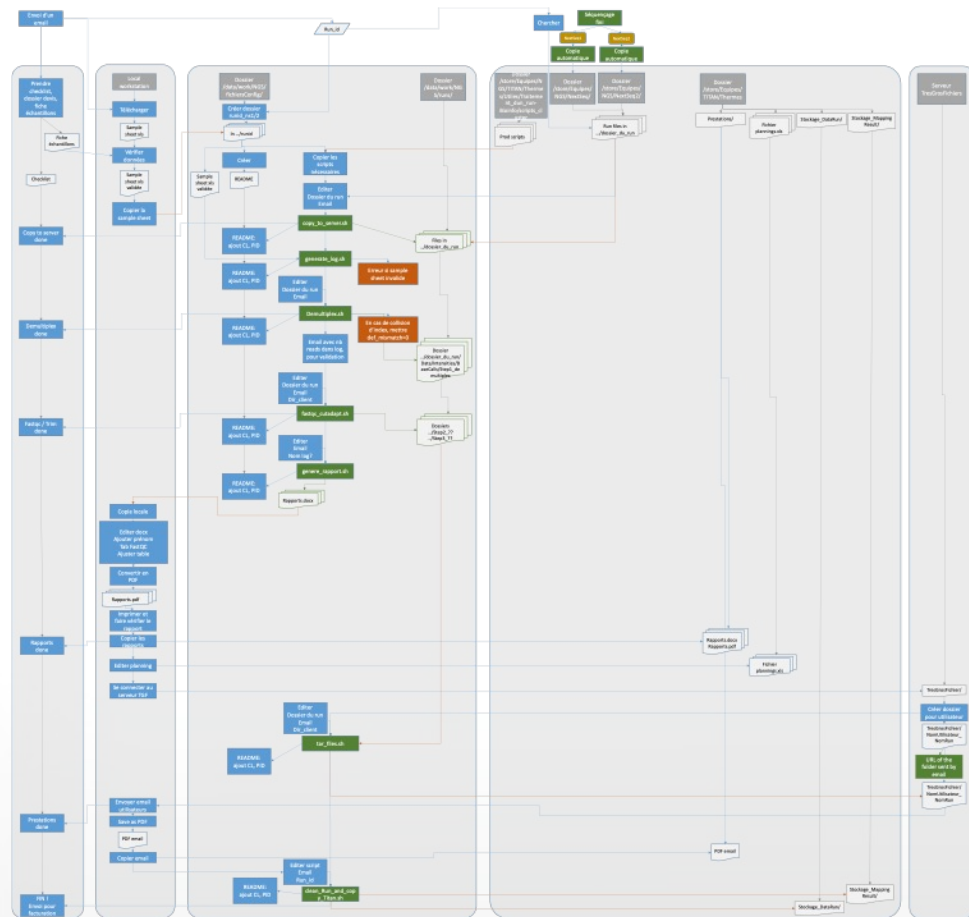


# Real data flow





# Real data flow



- (Who?)
- What?
- Where?
- When?
- How?







Who?

- Who is in charge of the current step?

What?

- What is being processed? What is the outcome?

Where?

- Where is the data, before/after the step?

When?

- When is that step performed?

How?

- How is the data processed?



## User needs

“Produce new data!”



Feasibility  
study

## User satisfaction



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## Metadata !

User metadata

- Request form
- Number of samples
- Species, ...



Feasibility  
study



## User satisfaction



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PF's metadata

- Date of arrival
- Person receiving the samples...

=> Traceability

User satisfaction

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**Secured storage** => Risk management  
(automatic, local, short/medium term)

Feasibility  
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Molecular  
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**Data production**  
(ultra raw data)



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- Software, version, CL => Reproducibility

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Primary  
treatment

**Raw data**



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Feasibility study

Molecular biology

**Data production**  
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Primary treatment

**Raw data**

## User satisfaction

**Shared storage**  
(automatic, very short term)

**Metadata**  
(reports)

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- Software, version, CL => Reproducibility

**Secured storage**  
(automatic, long term)

**Remote copy**  
(automatic, short term)

- Software, version, CL

**Secured storage**  
(automatic, medium term)

Feasibility study

Molecular biology

**Data production**  
(ultra raw data)

Primary treatment

**Raw data**

Bioinformatic analysis

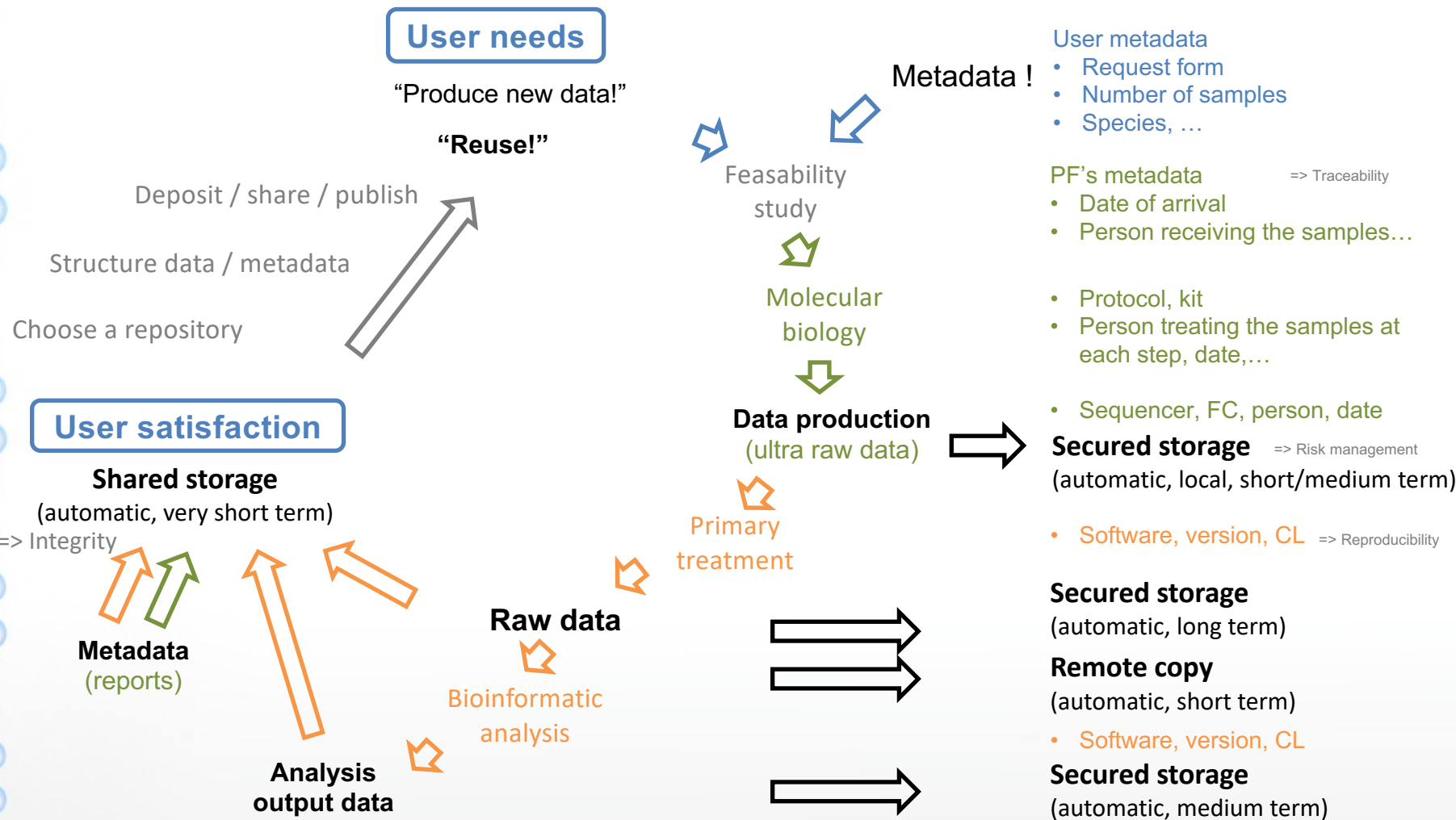
**Analysis output data**

## User satisfaction

**Shared storage**  
(automatic, very short term)

**Metadata**  
(reports)







# Difficulties

## Support

- Paper vs numerical (meta)data

## Storage

- Financial, ecological aspects
- Avoid redundancy (discussions with our institute)
- For how long ? (storage vs archive)
- Users still contact us for old data

## Bioinformatic analysis

- Reports, integrity verification
- FAIR for bioinformatics: versioning system, Docker images...
- How to facilitate deposition on repositories?



# Conclusions

(Meta)data producers



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- starts before anything is done
- is (part of) our Quality system



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A data management plan is (part of) our Quality system

- Traceability, risk management, reproducibility...





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Far from perfect, but improving

- FAIR at all levels (inside and outside the facility)
- Improve output reports and data structure



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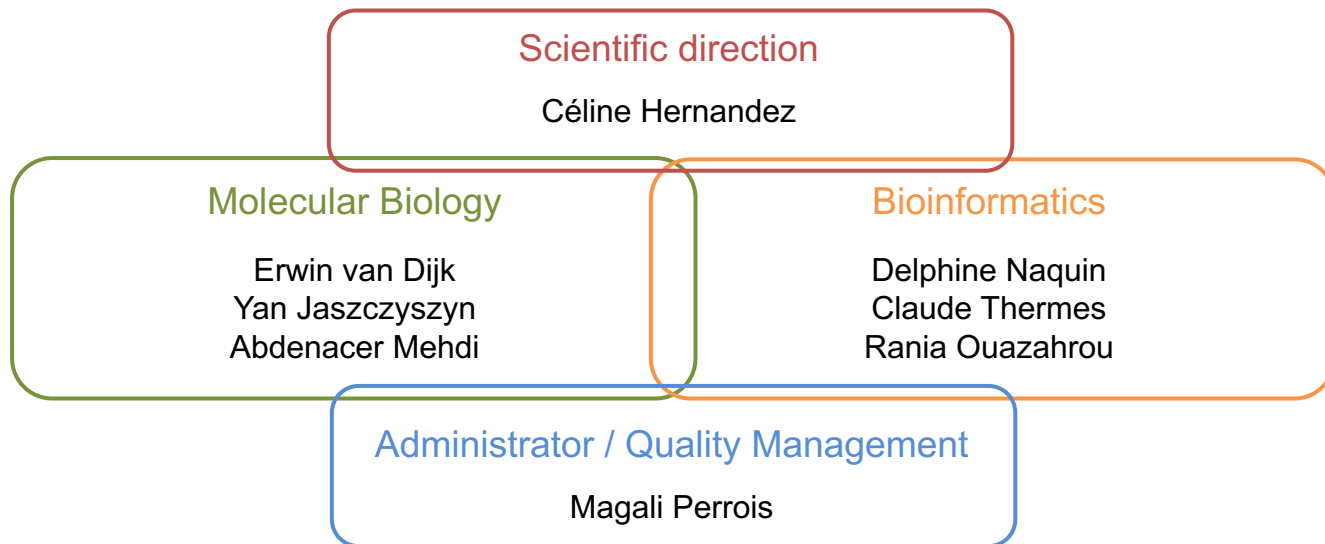
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“Data producer” or “Research output transformer”?

# High Throughput Sequencing Facility





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