

Cloud Computing  
NextFlow  
Data reproducibility



EUROPEAN  
GENOME-PHENOME  
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EMBL-EBI



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## The Cloud Computing Pilot

In recent years, the technological advances in genomics have resulted in the massive generation of data and the development of various tools for their analysis. Nowadays, a limited number of computational tools represent the core of data analysis pipelines, but their continuous updating poses new challenges that the scientific community needs to face.

### Challenges

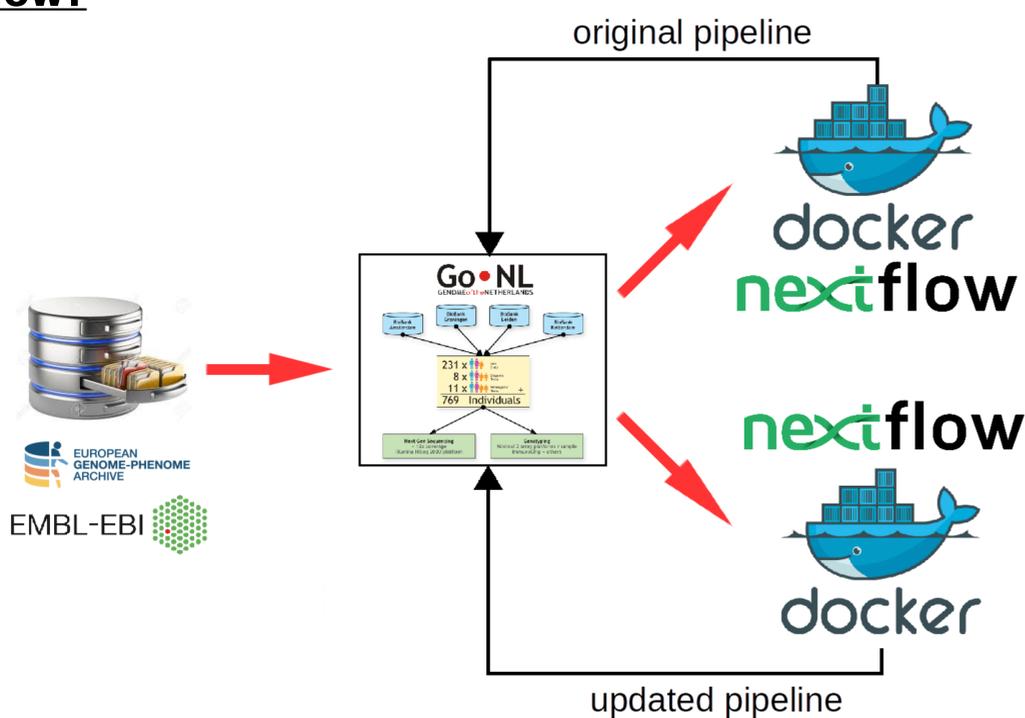
**Resource obsolescence:** Data are processed with reference genomes and analysis pipelines that were current at the time, but become obsolete quite fast

**Resource portability:** Software and tools require specific environment and dependencies that limit pipeline portability and maintaining them represents a time consuming task

### Goals

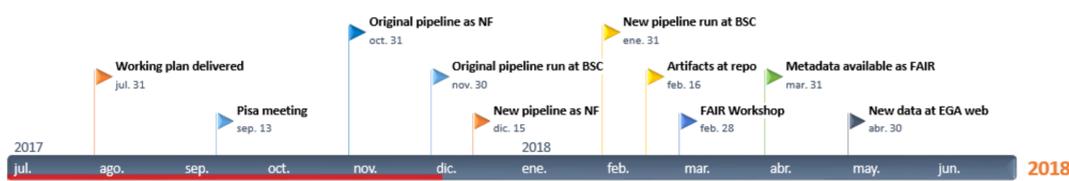
- Enable reproducibility → through the reconstruction of original analysis pipelines and the use of software containers which guarantee invariant executions and consistent results over time
- Allow portability → by packaging the original pipeline by using Nextflow workflow framework which enables the deployment across different computational environments in a portable manner
- Data re-analysis → by reproducing original pipelines and the use of updated resources

### How?



- ◆ A third part dataset (GoNL project) as use case
- ◆ Reproduction of the original pipeline
- ◆ Production of an updated pipeline
- ◆ Containerized versions of both pipelines
- ◆ Test both pipelines on the use case dataset

### State of the art



- ✓ Retrieve and install softwares and resources used in the original pipeline
- ✓ Generate a pipeline using the same commands and options used in the original pipeline
- ✓ Run the pipeline on a testing GoNL dataset
- ✓ Convert the pipeline as NextFlow pipeline

### Technical challenges

- ✗ Availability of original softwares and resources (GATK 1.0, ancillary files from 1000Genomes Project)
- ✗ Custom files and resources used in the original pipeline
- ✗ Openness to share sensitive data within the scientific community and the associated security from resource providers