O.D.I.N.

Omics Data Integration Network

Kévin LEBRIGAND, PhD Institut de Pharmacologie Moléculaire et Cellulaire Université Côte d'Azur, CNRS, France.

- lebrigand@ipmc.cnrs.fr
- @kevinlebrigand
- @ @kevinlebrigand@genomic.social

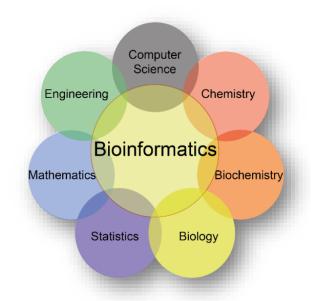




- Inter-UCA institutes of biology project (IRCAN Jipunca (Minstitut de Biologie Volrose (Archallorech
- ☐ Inter-disciplinary multi-omics: genomics, proteomics, metabolomics, cytometry, imaging
- □ Document and structure expertise's and resources dedicated to omics data analysis
- □ ODIN will be a multi-omics research project data manager (FAIR)
- □ ODIN will store and share reference data analysis guidelines
- □ ODIN will allow easy-to-start data analysis through 4D-SMICS infrastructure
- Optimize bioinformatics for UCA research teams scientific production

Bioinformatics

Discipline at the crossroads of biology, computer science and new technologies

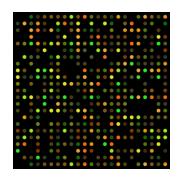


Characterized by the application of mathematical, statistical, computational methods to the analysis of biological, clinical, biochemical and biophysical data.

- development and implement tools to **store** and **manage** information manage data flows and infrastructure, database storage, web services implementation
- analysis and interpretation of data to identify relevant information analysis workflows setup, scripting analysis, figures production for publications, PI support
- development of new algorithms and statistical tools (computational biology) complex programing language, new signal mining, academics research in mathematics (AI)

Example of 20 years of transcriptomics

Driven by microfluidics technological developments

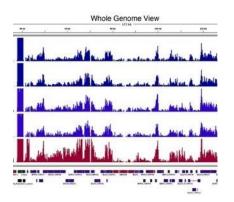


Early 2000's: DNA microarray

- Large-scale transcriptome
- Oligonucleotide probe tilling
- Fluorochrome signal analysis
- Bulk resolution



Cost : 4k€ 20 samples 25k genes **0,5M matrix**

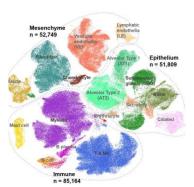


Late 2000's: RNA sequencing

- Whole transcriptome
- Next Generation Sequencing
- Full-transcript coverage
- Bulk resolution



Cost: 4k€ 20 samples 50k genes 1M matrix

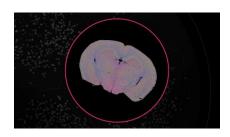


Mid 2010's: Single-cell

- Whole transcriptome
- Microfluidics + NGS
- 3p-end gene signal (UMI)
- Sensitivity (6%)
- Single-cell / state resolution



Cost : 4k€ 5k cells 50k genes 250M matrix



2020's: Spatial

- 500-1000 gene targets
- Imaging analysis
- Multiplexing FiSH (single molecule)
- Sensitivity (30-80%)
- Sub-cellular resolution



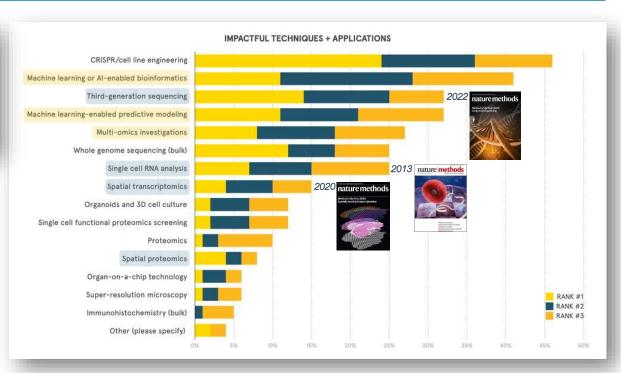
Cost: 4k€ 250k cells 1k genes 250M matrix

+ Spatial dimension

Academics labs scientific priorities

Requires expertise in bioinformatics and computational biology





Are UCA core facilities and research teams ready to process and analyze all those data?

4D-Omics (Equipex+, PIA3)

Instrument numérique pour la biologie quantitative multi-échelle en région Sud



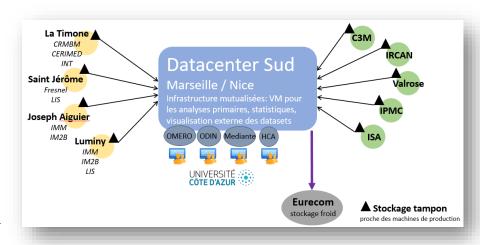




Structurer la composante biologique de DATASUD

Université Côte d'Azur, Université d'Aix Marseille

- ☐ Coordinateur: Pascal Barbry
- Projet: oct.2021 jun.2029
- Budget: 6.595.999 €
- √ Convaincre les laboratoires de biologie des avantages d'une migration vers des ressources numériques distribuées:
- ☐ Économie de moyens, optimisation des moyens en personnel
- ☐ Recentrage des laboratoires sur l'exploitation de la donnée
- ✓ Illustration de la démarche sur quelques programmes phares:
- ☐ <u>Human Cell Atlas:</u> un consortium international fortement structuré, basé sur la distribution rapide et large de données potentiellement sensibles
- □ *Omero*: un système d'information pour les données d'imagerie
- ☐ <u>Mediante:</u> un système d'information pour les données de génomique
- □ <u>ODIN</u>: centralisation des scripts et objets d'analyses des données de biologie
- ✓ Entrepôts de données:
- ☐ Assurer la pérennité des données du site (5 Pb bandes Eurecom, may 23)
- ☐ Travail sur les données de santé (CHU)



Data analysis sustainability and expertise sharing between UCA research teams





Step-by-step



1. Draw the UCA omics resources map (production systems and expertises)

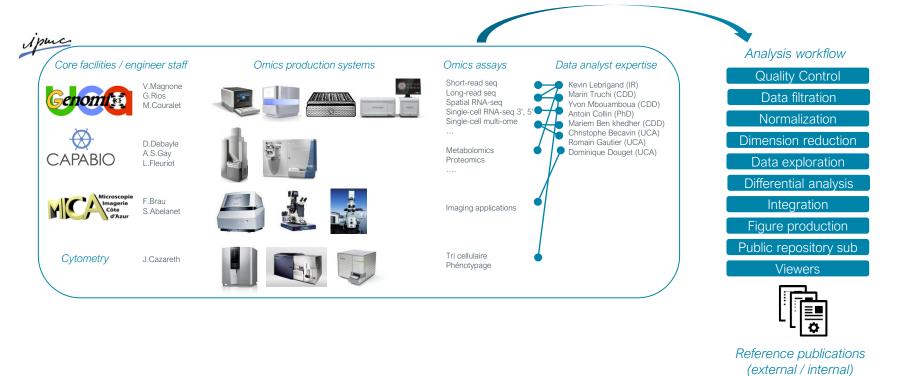


Step-by-step



Guidelines

2. Document internal or external reference analysis guidelines



Step-by-step



3. Implement a multi-omics research project FAIR data manager system

ODIN web portal



- Java J2EE development
- Hosted by Tomcat app server
- PostgreSQL database

Storing / Managing

- User/team-oriented research project manager (FAIR)
- Reference guidelines for "omics" data analysis

Step-by-step



4. Start bioinformatics analysis session based on new data and shared guidelines

ODIN web portal





- Java J2EE development
- Hosted by Tomcat app server
- PostgreSQL database

Storing / Managing

- User/team-oriented research project manager (FAIR)
- Reference guidelines for "omics" data analysis

Bioinformatics analysis













- RStudio server
- Jupyter Notebook server

Technical solutions

- Local export archives (data + analysis scripts)
- Container: Kubernetes, Docker, Singularity
- Virtual machines





UCA Research Engineer - IDEX academy 4 (September 2023)

Acknowledgments

Omics Data Integration Network





Olivier Croce Gaël Cristofori Boris Meyer Claude Philippe Erwan Simon



Arnaud Hubstenberger Thierry Lepage Antoine Fortuné Luc Martin

Kevin Lebrigand
Marin Truchi



Etienne Danchin Corinne Rancurel Martine Da-Rocha Arthur Péré



Michele Trabucchi Maria Stathopoulou

IPMC bioinformatics platform

Structure and organize bioinformatics resources and developments @ IPMC level



Committed to support IPMC research teams and facilities

- ☐ IR recruitment (CDD, 1 year, march 2023)
- ☐ Support experimental design setup, funding requests and paper methods writing
- ☐ Constant technological watch in the priority technological areas defined by the institute
- Support bioinformatics @ IPMC (internal seminars)
- ☐ French Institute of Bioinformatics (IFB) subscription (visibility, ease for future recruitment)
- ☐ Dedicated space to group bioinformaticians recruited by teams and facilities
 - Group work emulation
 - Define standardized workflows for IPMC equipment in Genomics, Proteomics, Cytometry and Imaging
 - Mentoring, best practices and guidelines sharing
 - Expertise / methodological developments shared between teams
 - Provide an optimal working environment, IT infrastructure (4D-\$\infty\text{MICS}\)

UCA omics ecosystem

Cartography of UCA resources and expertise in omics data production and analysis



Cartographie des ressources et expertises omiques des instituts de biologie UCA

En relation avec les différents partenaires du réseau, réaliser une cartographie des ressources des instituts partenaires UCA en lien avec la production et l'analyse des données de biologie quantitatives: bioinformaticiens, expertises, systèmes de production des plateaux techniques, infrastructure informatique, équipes de recherches potentielles partenaires.



Cytometry J.Cazareth

Omics production systems













Omics assays

Short-read seg Long-read seg Spatial RNA-seq Single-cell RNA-seg 3' Single-cell multi-ome

Metabolomics Proteomics

Imaging applications

Kevin Lebrigand (IR) Marin Truchi (CDD) (von Mbouamboua (CDD) Antoin Collin (PhD) Mariem Ben khedher (CDD) Christophe Becavin (UCA) Romain Gautier (UCA) Dominique Douget (UCA)

Data analyst expertise

Informatics infrastructure

Caire (serveur web) Intel Xeon - 8 cores 48 Gb RAM 29 To storage

Joyo (calcul cluster)

Intel Xeon - 8 cores 48 Go de RAM, 15 + 22 To storage SGE x12 nodes (20 cores, 64 Gb RAM)

Bego (stats server)

2 x Intel Xeon Gold 6248 - 40 cores GPU A100 Ampere 2 To RAM 37 To storage

Réalisation du même travail dans les autres instituts de biologie UCA









