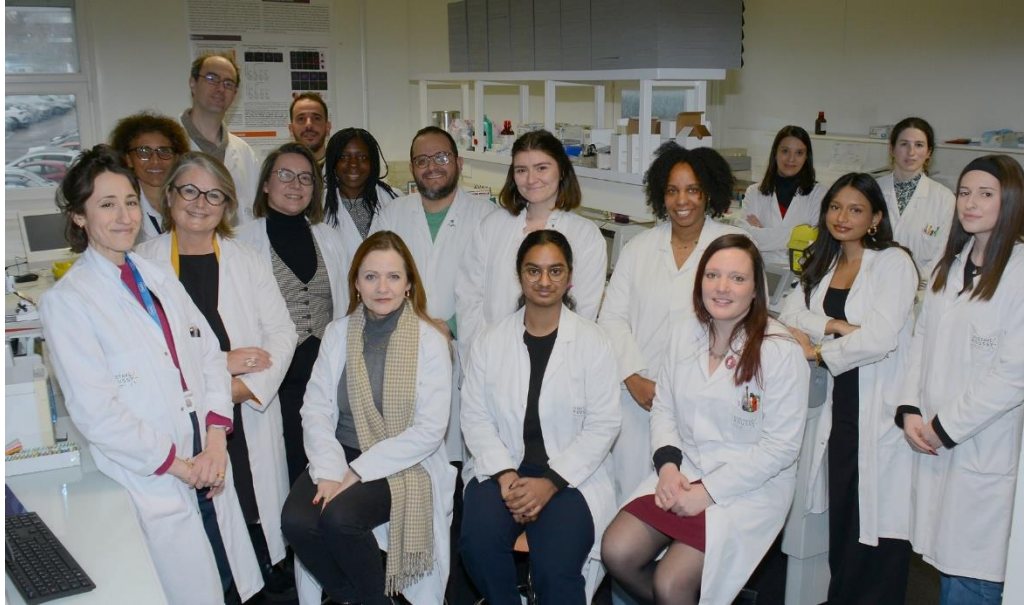


PETRA Core Facility

Experimental and Translational Pathology



*PETRA Open Day
Tuesday, 16 December 2025*

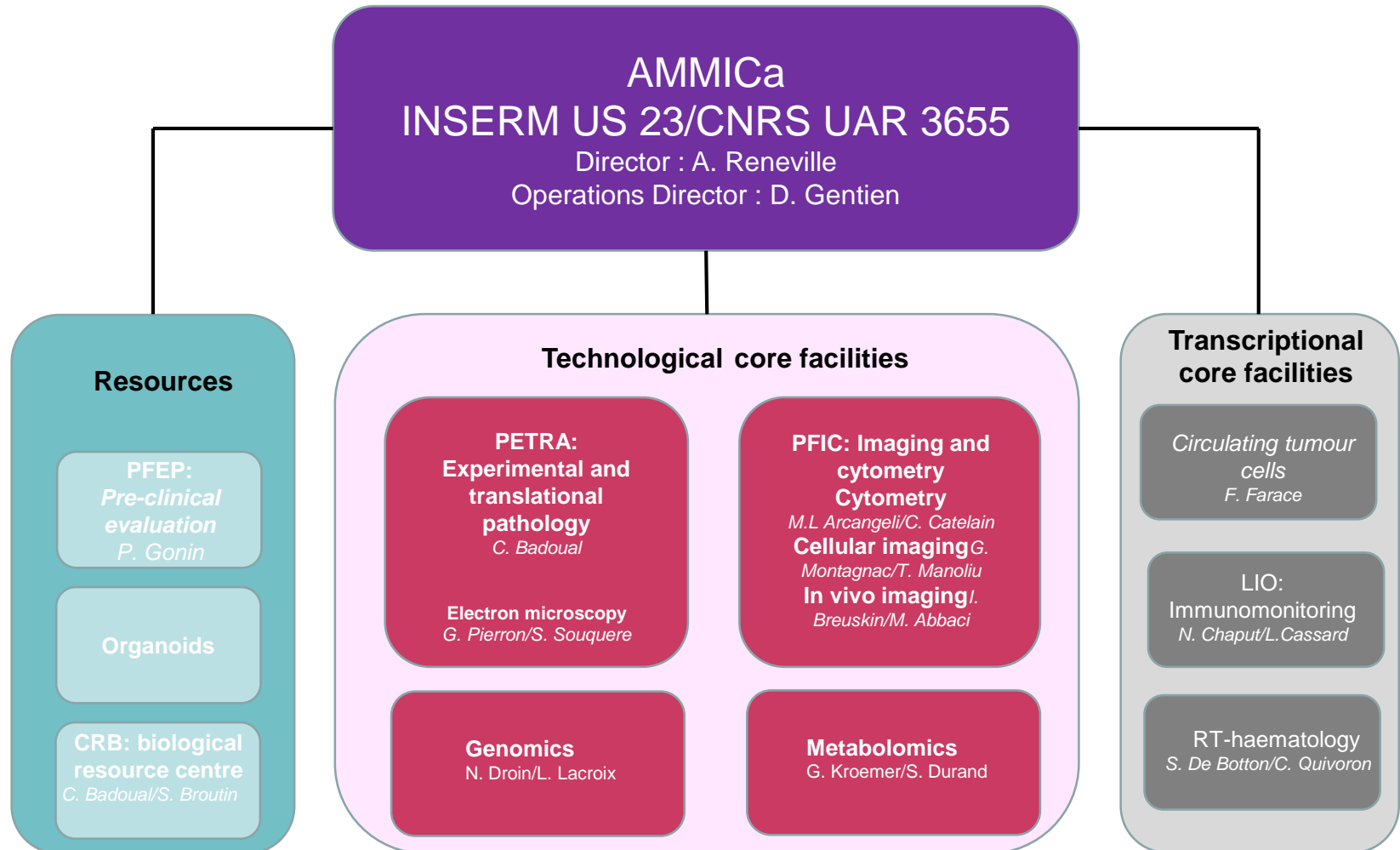
*Prof. Cécile Badoual-Virginie Marty-Olivia Bawa
Plateforme Pathologie Expérimentale et TRAnslationnelle (PETRA)
Gustave Roussy*

OUTLINE

- Who we are about us?
 - PETRA
 - Team organization
 - Module I : Research program
 - Module II : Experimental pathology
 - Module III : Clinical research
 - Module IV : Digitization, image analysis
- What do we offer?
 - Spatial Omics
 - Proteomics
 - Transcriptomics
 - MultiOmics (integrated)
 - How to contact us?

Who we are about us?

AMMICA : PETRA

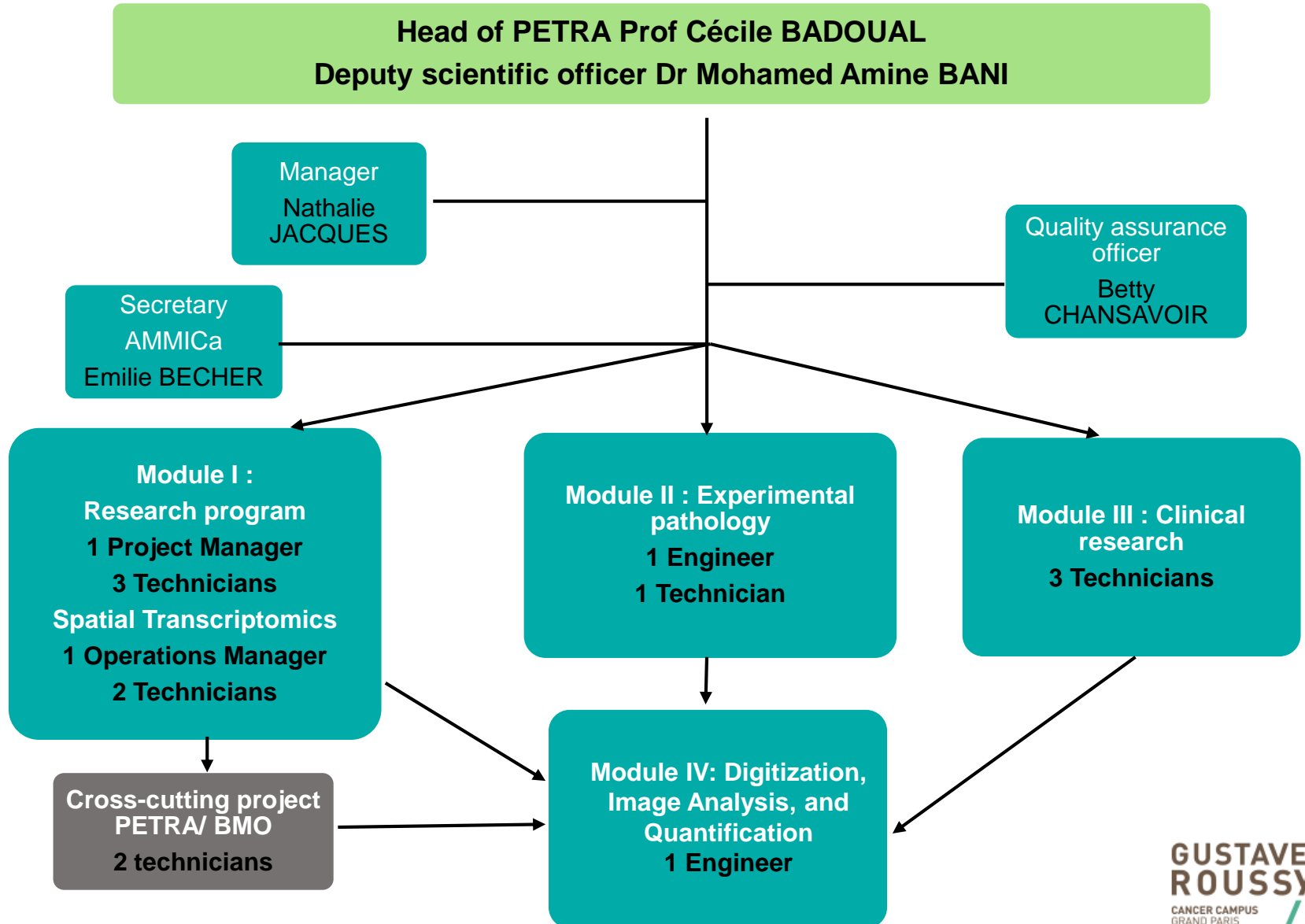


AMMICA : Molecular analysis, modeling, and imaging of cancer

Who we are about us?

- ISO9001 certified since 2015
- Member of the Rés'Hist network since 2023
- Our missions: To perform conventional histology, multiplex immunohistochemistry, spatial transcriptomics and image analysis techniques
- Our objectives: Consulting, monitoring and innovation in morphological and molecular in situ analysis techniques for research

Team organization



Module I: Research program 1/2

- Field of activity: Research projects and projects ancillary to therapeutic trials (82 projects in 2024)
- Team:
 - Project manager Virginie Marty: provides assistance in setting up research projects, cross-disciplinary expertise in transcriptomics and spatial proteomics, implementation of new technologies, training of interns
 - Technicians
 - ✓ Laëtitia Bordelet : conventional histology and multiplex IHC, PLA specialist
 - ✓ Elodie Edmond : conventional histology and multiplex IHC, FISH specialist
 - ✓ Karine Godefroy : conventional histology, expert in pre-analytical spatial transcriptomics techniques

Module I: Research program 2/2

- Field of activity: Spatial transcriptomics (17 projects since the first machines were installed) in association with molecular biology and bioinformatics core facilities

« Possibility of mapping gene expression and improving understanding of tissue biology. »

- Team:

- Operations Manager: Valérie Camara-Clayette: management of transcriptomics projects implementation

- Technicians

- ✓ AbdelHafidh Belakermi : expert in GeoMx and Visium technologies

- ✓ Adsaya Rathakrishnan : Merscope technology expert

Module II: Experimental pathology

- Field of activity: Pre-clinical studies (animal models, PDX, organoids and spheroids) (81 projects in 2024)
- Team :
 - Engineer Olivia Bawa: provides support in setting up experimental pathology projects with expertise in conventional histology and multiplex IHC (murine samples and organoids), development of multi-omics techniques (RNA + proteins), training of interns
 - Technicienne Lily Camus : specialist in conventional histology and development of new markers

Module III: Research clinical

- Field of activity: Phase I, II and III therapeutic trials (600 projects in 2024)
- Their missions :
 - Respond to institutional requests from Gustave Roussy's "Guichet unique"
 - Validate feasibility reports
 - Register samples belonging to a trial
 - Bio-banking
 - Perform conventional histology techniques and immunohistochemistry requests
 - Provide tutoring
- Team :
 - Technicians:
 - ✓ Betty Chansavoit : Quality Assurance Manager
 - ✓ Laura Cheron
 - ✓ Mailys Rolland

Module IV: Digitization, image analysis

- Field of activity: all PETRA modules (100 projects in 2024)
- Engineer Nicolas Signolle: expert in digitisation, simple image analysis, multiplex, hiplex, LS multiomics, algorithm creation
- Training courses offered: Olympus slide scanner, QuPath



HISTOLOGY & SPATIAL BIOLOGY

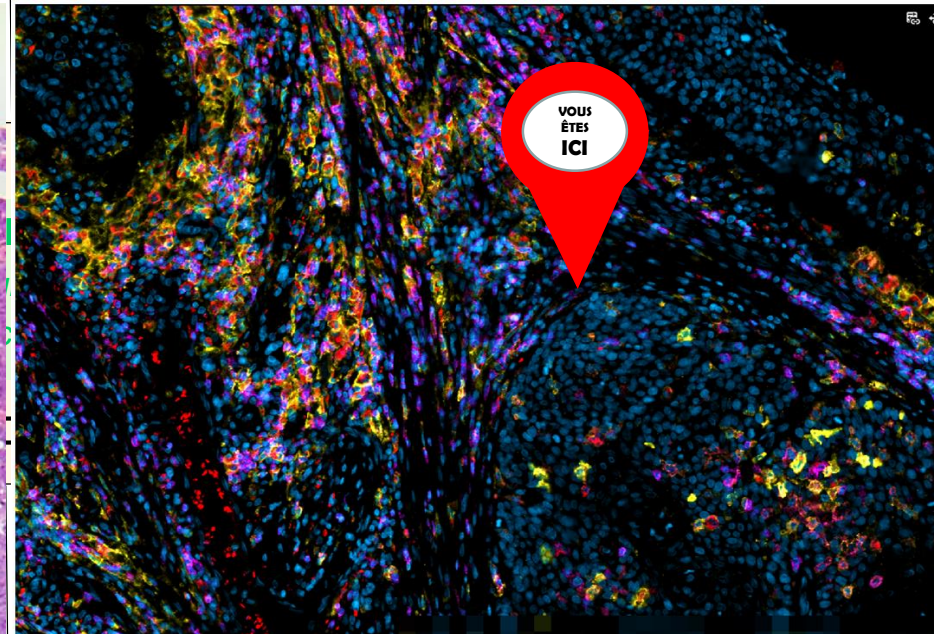
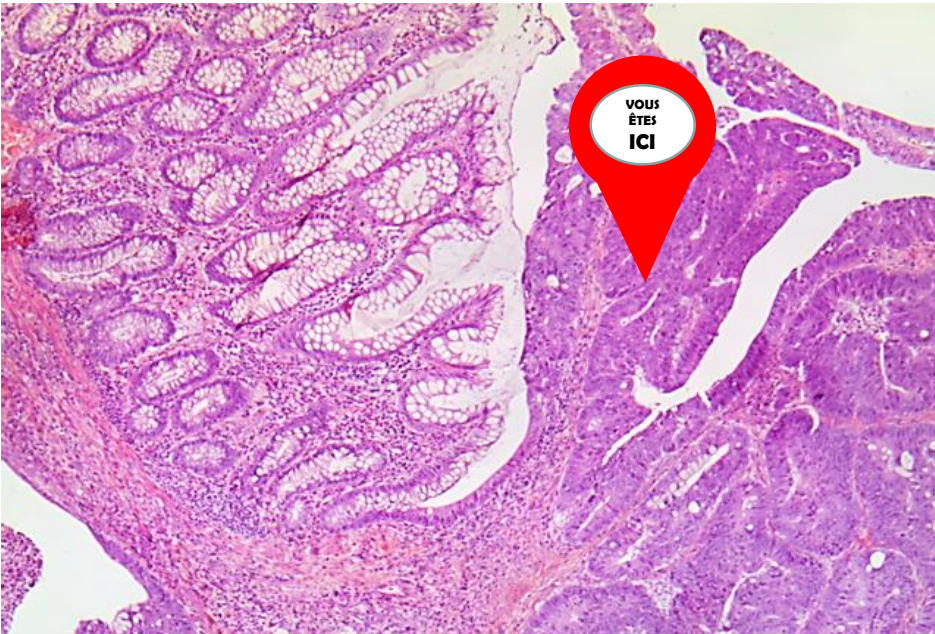
PRE-ANALYTICAL

Sample inclusions

Media for analysis: Slides / Shavings

BMO Core facility

*Bulk genomic & transcriptomic
analytical techniques*



POST-ANALYTICAL

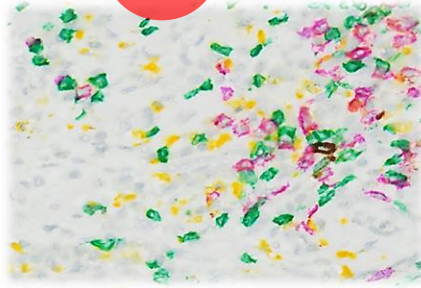
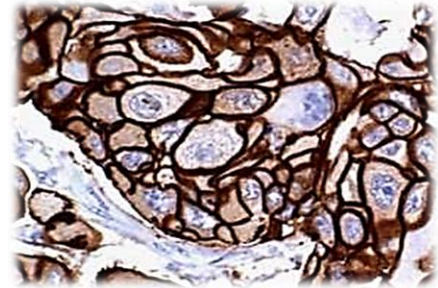
Scanning, image analysis: quantification...

Spatial Proteomic Landscape

Plexing : from 2 to 1 200 proteins

Chromogenic

Single cell

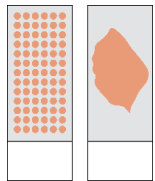


1 protein

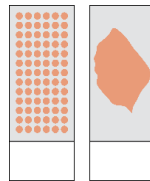
Benchmark UltraPlus
& *Bond-RX*

Up to 5 proteins - Full custom

Discovery Ultra



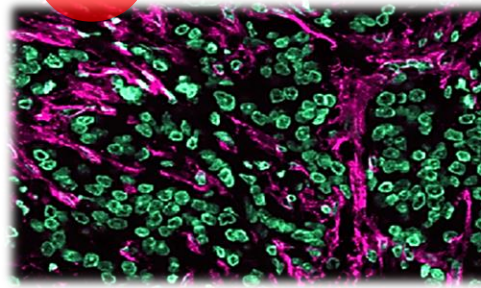
whole slide



whole slide

Fluorescence

Single cell



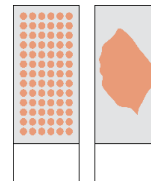
Up to 6 proteins

Bond-RX

Full custom

Up to 40 proteins

COMET



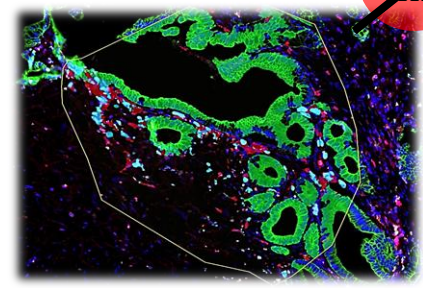
whole slide



12.5 x 12.5 mm

Proteomic Profile per ROI

Single cell



Up to 1 200 proteins - Fixed panel

GeoMx



35 x 14 mm

THROUGHPUT

30 slides = 4h

30 slides = 17h

30 slides = 17h

4 slides = +/-48h

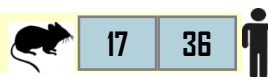
4 slides = 2j

(without sequencing)

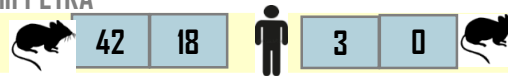
AVAILABLE in PETRA



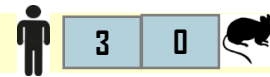
190 492



17 36



42 18



3 0



5 morpho Abs

DEVELOPPEMENT TIME

15 days

2 mois

5 mois

3 mois

1 mois

Spatial Transcriptomic Landscape

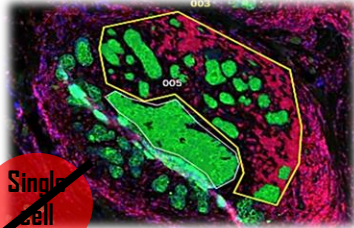
Panels : from 140 to 18 000 genes

Up to 12 RNA

Discovery tools

Hypothesis-driven tools

NGS-based Spatial Transcriptomics

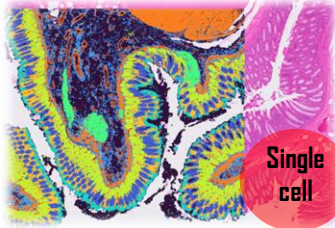


Up to 18 000 genes
Fixed panels

GeoMx



35 x 14 mm



WT - 18 000 genes

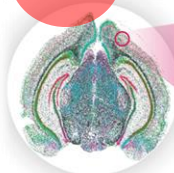
Visium-HD



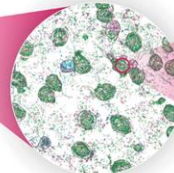
6,5 x 6,5 mm

Imaging-based Spatial Transcriptomics

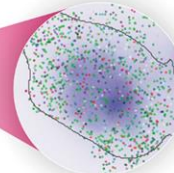
Single cell



WHOLE SECTION



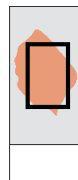
WIDE FIELD OF VIEW



SUB-CELLULAR

Up to 480 genes
100% custom
& mutations targeting

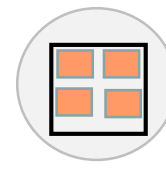
Xenium



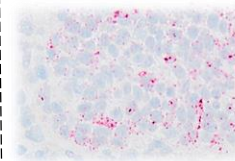
10 x 22 mm

Up to 960 genes
Full custom

Merscope



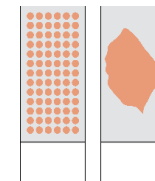
10 x 10 mm



Single cell

From 1 to 5 RNA
chromo / fluo

Bond Rx



whole slide

Up to 12 RNA

fluo

COMET



12.5 x 12.5 mm

4 slides = 2 days

(without sequencing)

1 slide = 2 areas = 3 days

2 slides = 1 week

1 slide = 1 week

30 slides = 17h

4 slides = 50h

THROUGHPUT

AVAILABLE in PETRA



5 morpho Abs



9

0



DEVELOPPEMENT TIME

1 mois

NO

NO

1 mois / new tissu type

15 jours

1 mois

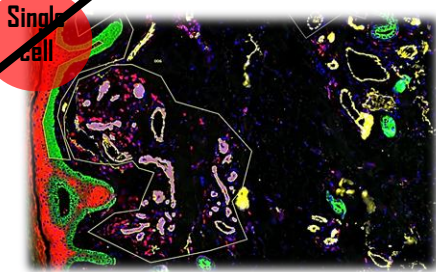
Spatial MultiOmics Landscape



Spatial MultiOmics Landscape

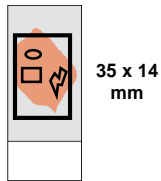
Discovery tools

NGS-based Spatial Transcriptomics & Proteomics



Up to 18 000 genes
+ 1 200 proteins
Fixed panels

GeoMx



Imaging-based Spatial Transcriptomics & Proteomics

Single cell

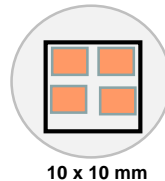
Up to 480 genes
+ 27 proteins
Fixed & Custom panels

Xenium



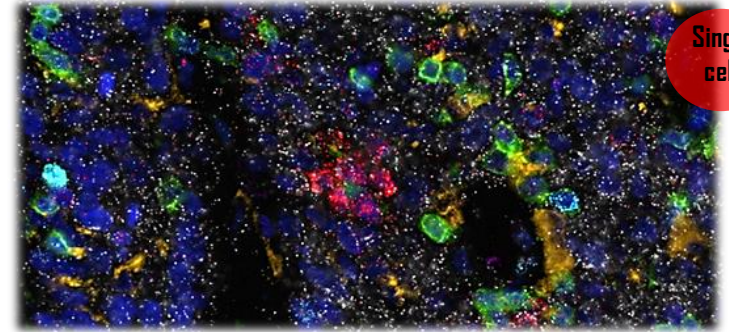
Up to 960 genes
+ 5 proteins
Full custom

Merscope



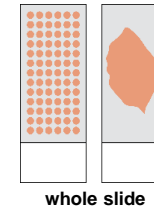
Hypothesis-driven tools

Imaging-based RNAscope and multiplex IF



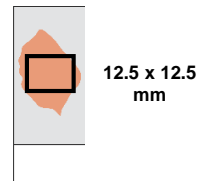
Up to 4 RNA
+ 2 proteins

Bond-RX



Up to 12 RNA
+ 24 proteins

COMET



THROUGHPUT

4 slides = 2j

(without sequencing)

2 slides

1 slide = 1 week

30 slides = 20h

4 slides = 66h

AVAILABLE in PETRA



5 morpho Abs



0 0



0 1



1 0



DEVELOPPEMENT TIME

1 mois

NO

NA

3 mois

3 mois

How to contact us? 1/2



Student,
Researcher,
Oncologist,
Pathologist

Fill in the contact form

Send to the following address :
contact-petra@gustaveroussy.fr

Research Project Module:
complete a project form

Experimental Pathology
Module: complete a
project form

Image digitization and
analysis module:
complete a project form

Signing the quotation in
accordance with the charter

How to contact us? 2/2



Preparation of the quotation

Presentation of the project if necessary at the PETRA meeting: Wednesday in the Crystal Room at B2M, attended by Professor Cécile Badoual and Dr Mohamed Bani.

Technical deadlines, choice of technologies, advice from our expert pathologists

Implementation of techniques, reporting of results, invoicing