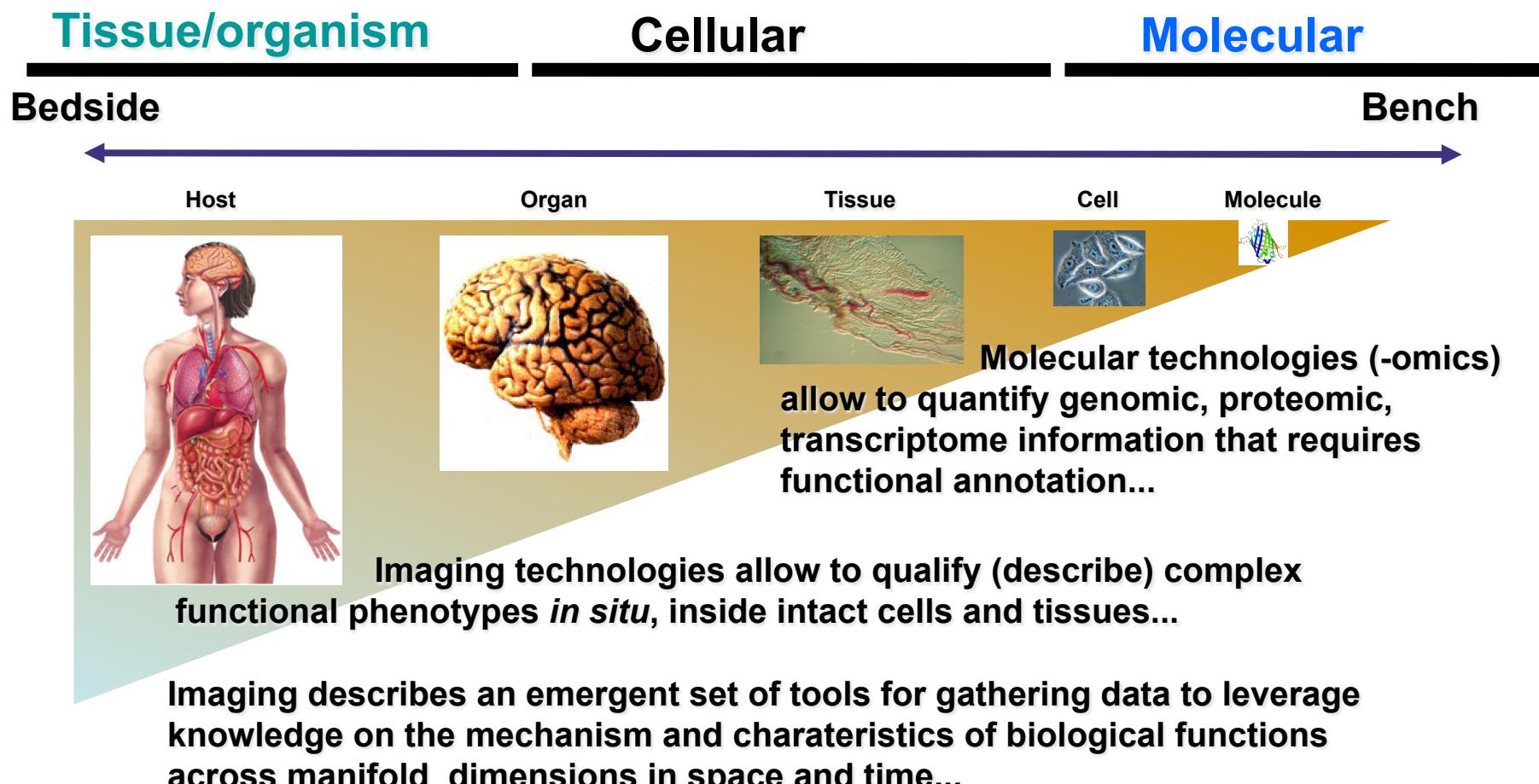


Imagopole

EuCAI founding meeting

Trinity College, Dublin, Feb. 18th 2013
Organized by Peter Hovarth & Anthony Davies





Biology is a scale problem...

Tissue/organism

Cellular

Molecular

Intravital imaging-2Pi/SHG

Multi-D fluorescence

Cryo-Electron-Microscopy

In situ cell imaging/tracking

Bioluminescence

CLEM- Correlatoire EM

Ex-vivo tissue imaging

Imaging Flow Cytometry

FRET/FLIM/FCS/FRAP/FLIP

Histopathology/QWBA/MALDI

Tomo-Electron microscopy

STORM/PALM/STED/+???

In Vivo BL/FL/FMT/Spectral imaging

Hyper-Spectral

AFM/FL

Relative Sizes and Detection Devices

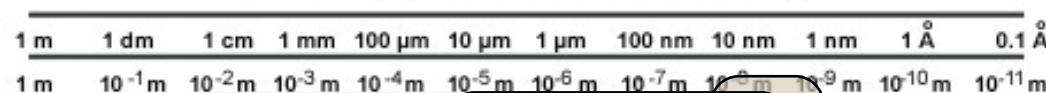
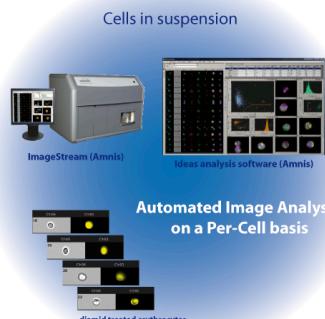


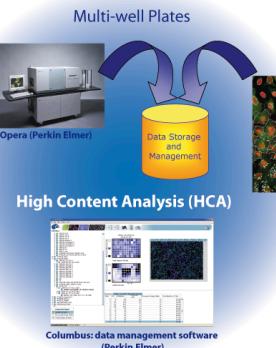
Figure adapted from
Florida State University
"Expressions" web-site

Integrative Experimental Biology

Imaging Cytometry



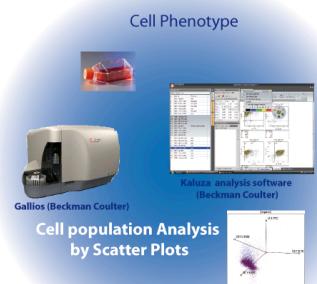
High Content Screening (HCS)



In Vivo Optical Analysis



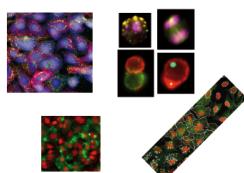
Cytometry



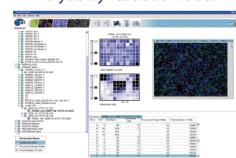
Electronic microscopy



Quantification



Analysis by readout model

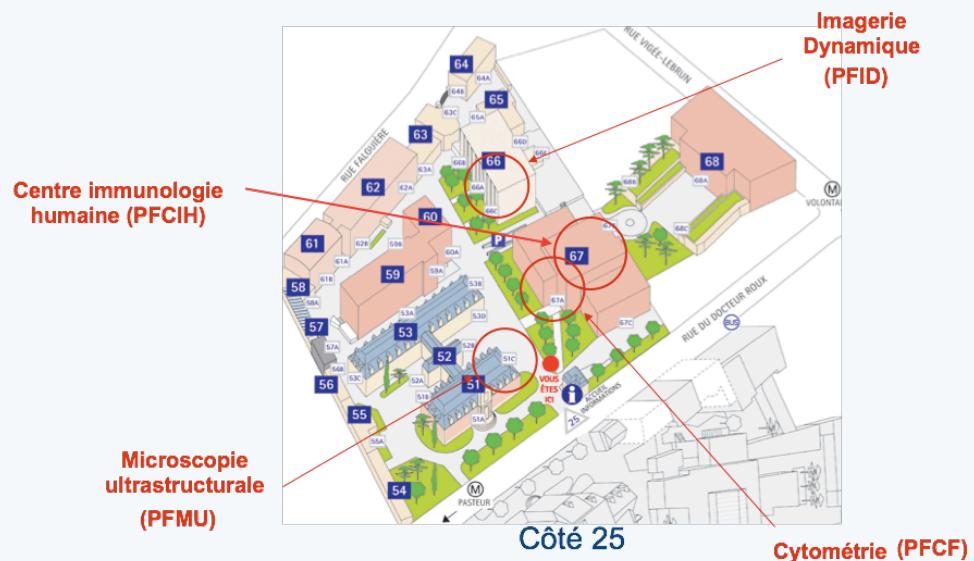


Validation & Statistics

- b Correlation analysis (Relationship between HCA readouts)
 - b Normalisation
 - b Multivariate Analysis
 - b Cluster analysis of readouts (Classification)
 - b Hit/Event selection Procedure
-

Data Storage & Management





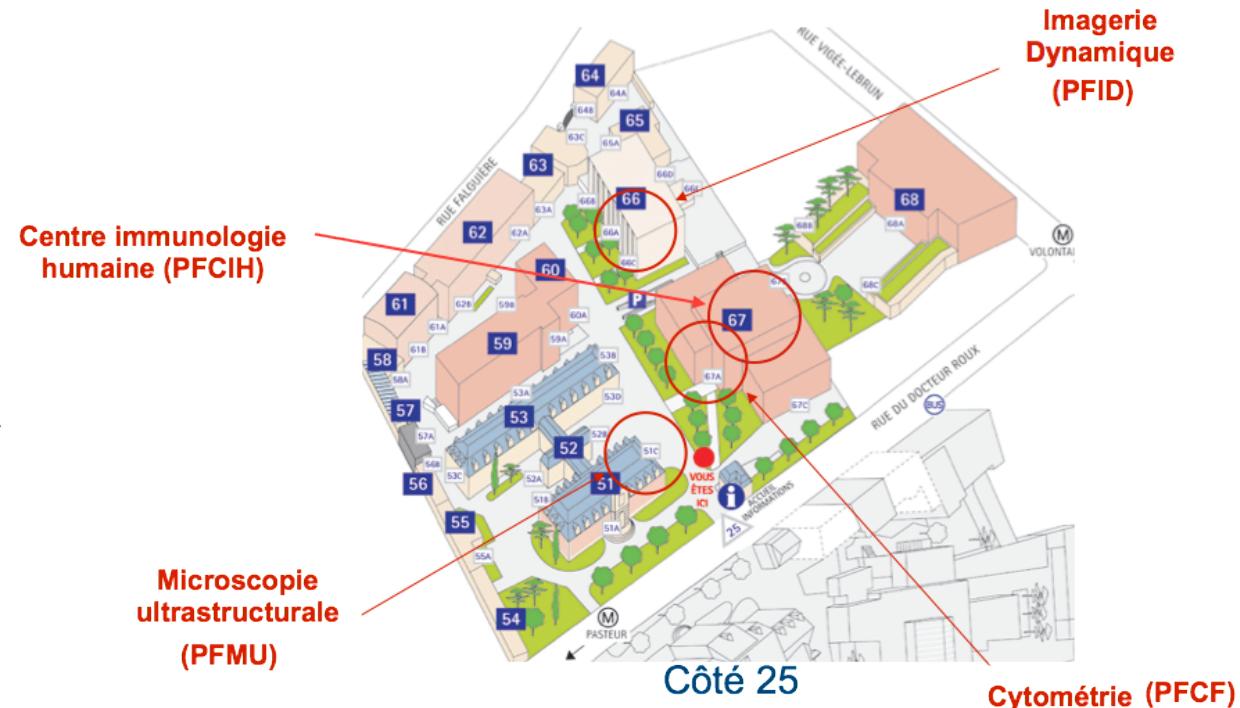
Imagopole mission: *Develop and apply scientific imaging technologies, in the context of experimental frameworks aimed at understanding of biological processes, and their usurpation by infectious disease*

www.imagopole.org

Technology Platform philosophy: platforms have a different focus to research units



- Ultrastructural Microscopies
- Imaging & Flow Cytometry
- Dynamic optical imaging
- Intravital & *in vivo* imaging
- Image processing & analysis
- Bio-informatics, & statistics
- Molecular imaging
- Translational research



• Reagents, Contrast Agents

35 Permanent scientific, engineering, technical staff including 7 temporary staff (post-docs, students etc); 40 major equipment installations (imaging microscopes, scanning & flow cytometry, cryoEM etc); Informatics comprising 100 active clients, numerous file-, web-, and calculation servers, 10TB local RAID storage; 120TB FAS 6070 NetApp SAN storage a variety of “imaging” softwares: Huygens-SVI, Metamorph, Imaris, Osirix, imageJ, Definiens, 3i etc



Imagopole

Key facts & figures



- ISO 9001 Certification for quality in service standards (Dec 2007)
- IBiSA national scientific platform label (Nov 2009)
- >700 registered users, >45,000 hours/year “burn-time”
- 10% External users
- 4 Patents, 1 Software copyright (1 spin-out Stratocore.com)
- >25 peer reviewed high-impact scientific articles a year
- Global annual budget millions euros (salaries, consumables, equipment)

IMAGOPOLE
Pôle de Dynamique Moléculaire et Fonctionnelle

*Plate-forme de Cytométrie en Flux
Plate-forme d'Imagerie Dynamique
Plate-forme de Microscopie Ultrastructurale
Centre d'Immunologie Humaine*

Upon request...



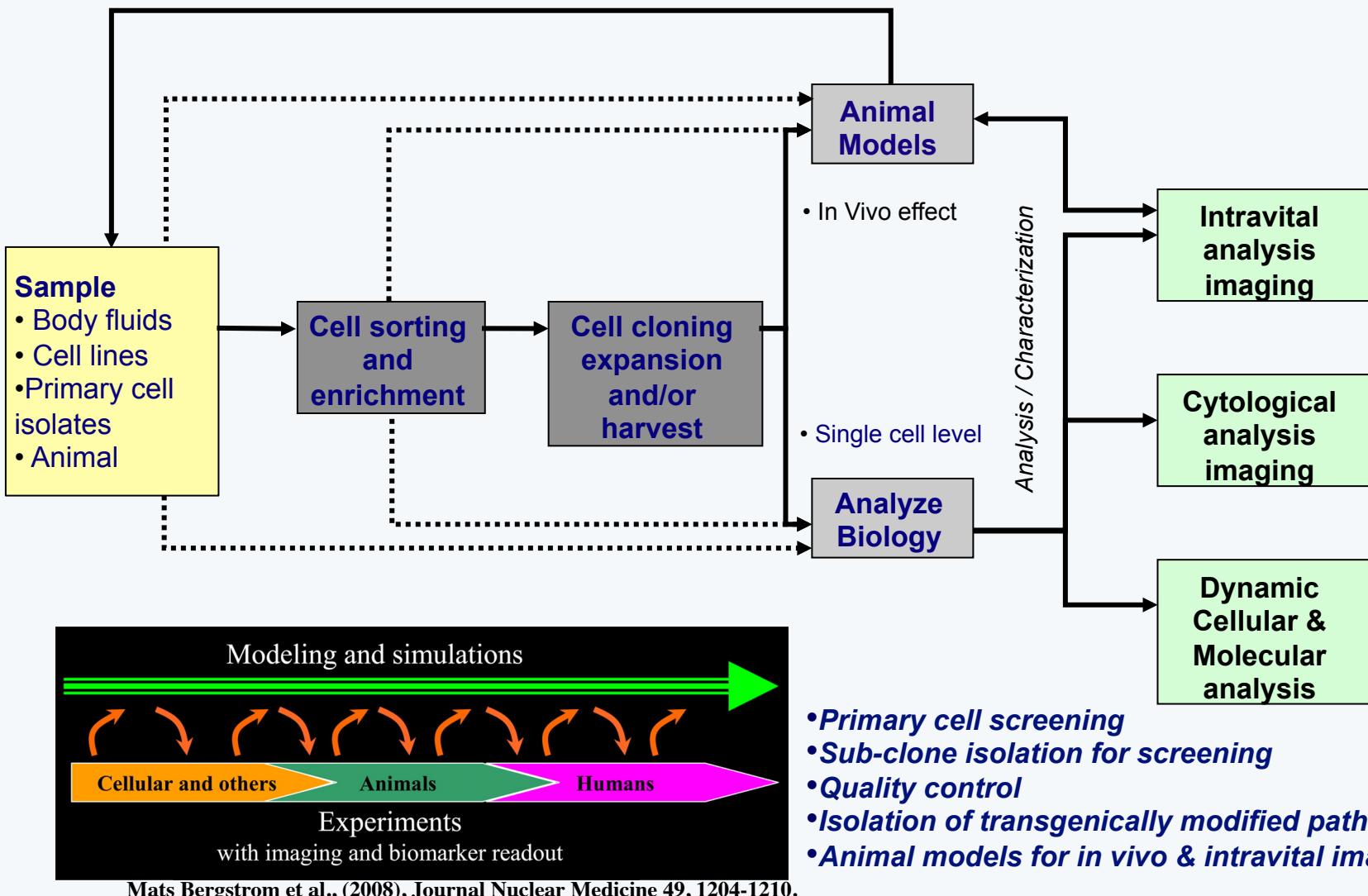
Stratocore

Core facility resource management
<http://stratocore.com>

Integrative Experimental Biology

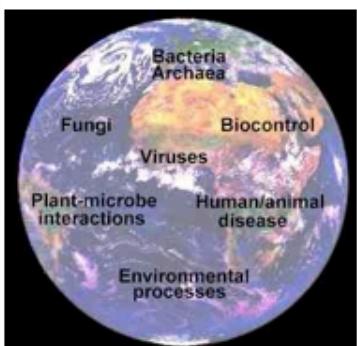
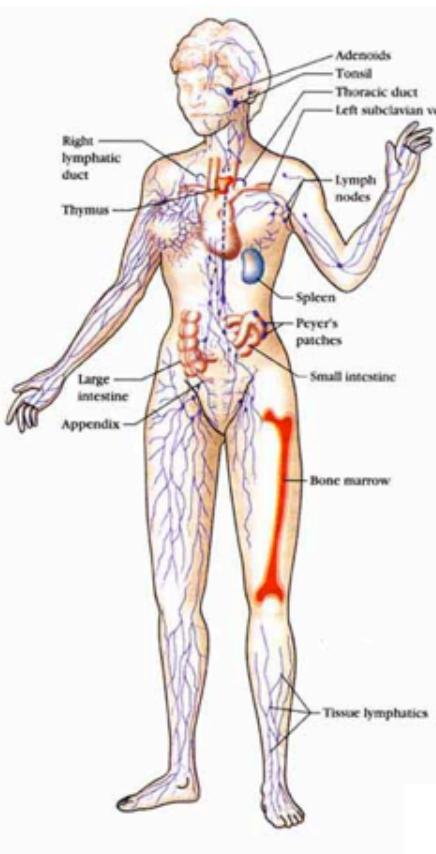


Institut Pasteur



A systems approach to understanding the Immune System

Genetic & Environmental Determinants of Immune Phenotype Variance

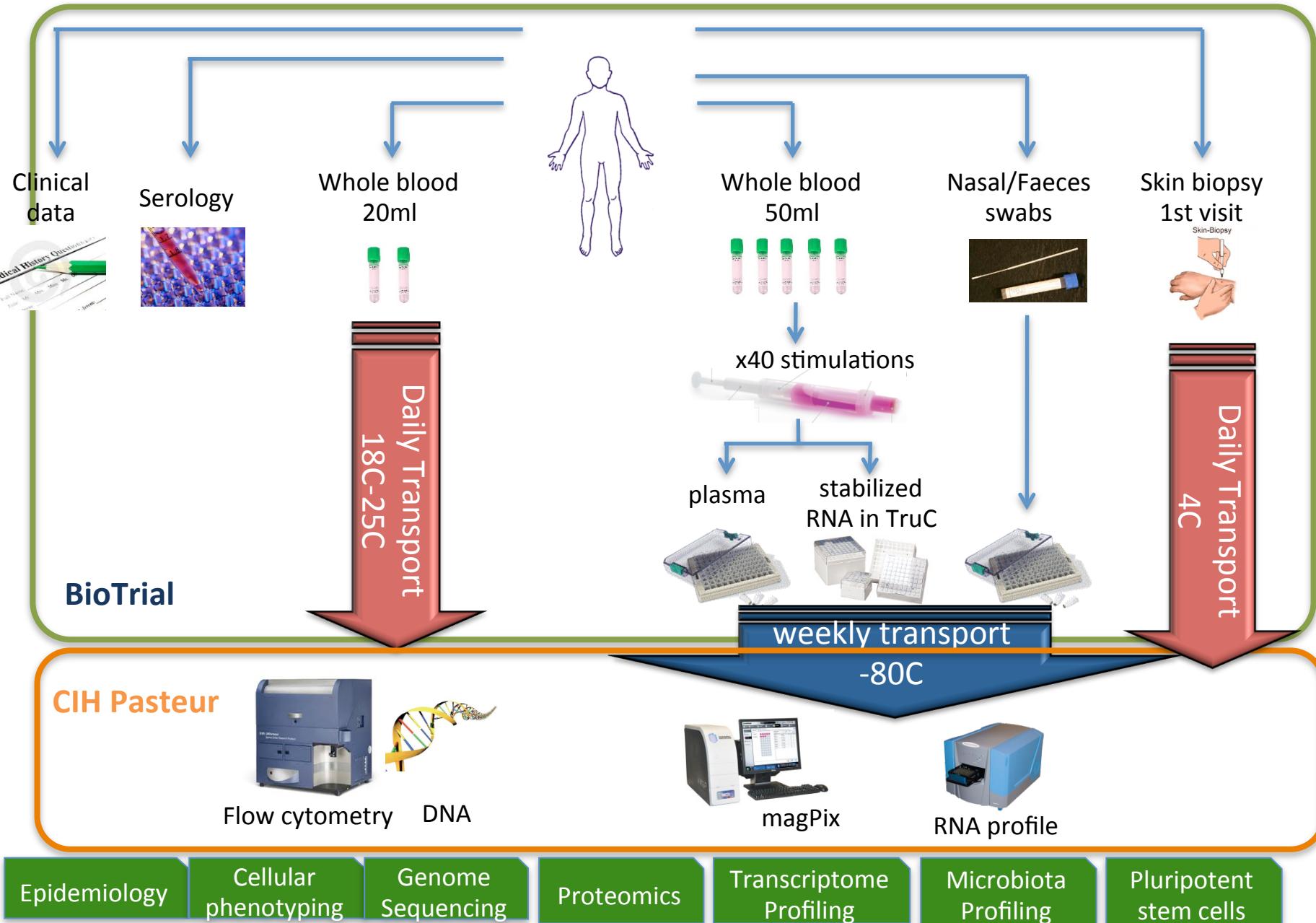


- How variable is the « immune response » across a population of healthy individuals?
- How can we account for this variation – genetic, epigenetic *versus* environmental control?
- How have genetically-controlled immune mechanisms contributed to our survival?
- How do perturbation of these mechanisms contribute TODAY to immune-related pathologic conditions?

A path towards personalized medicine

- Host response to infectious agents
- Response to adjuvants and vaccines
- Relationship between commensals and immunity

1000 healthy donors included: 3 visits (V0, V1, V2)

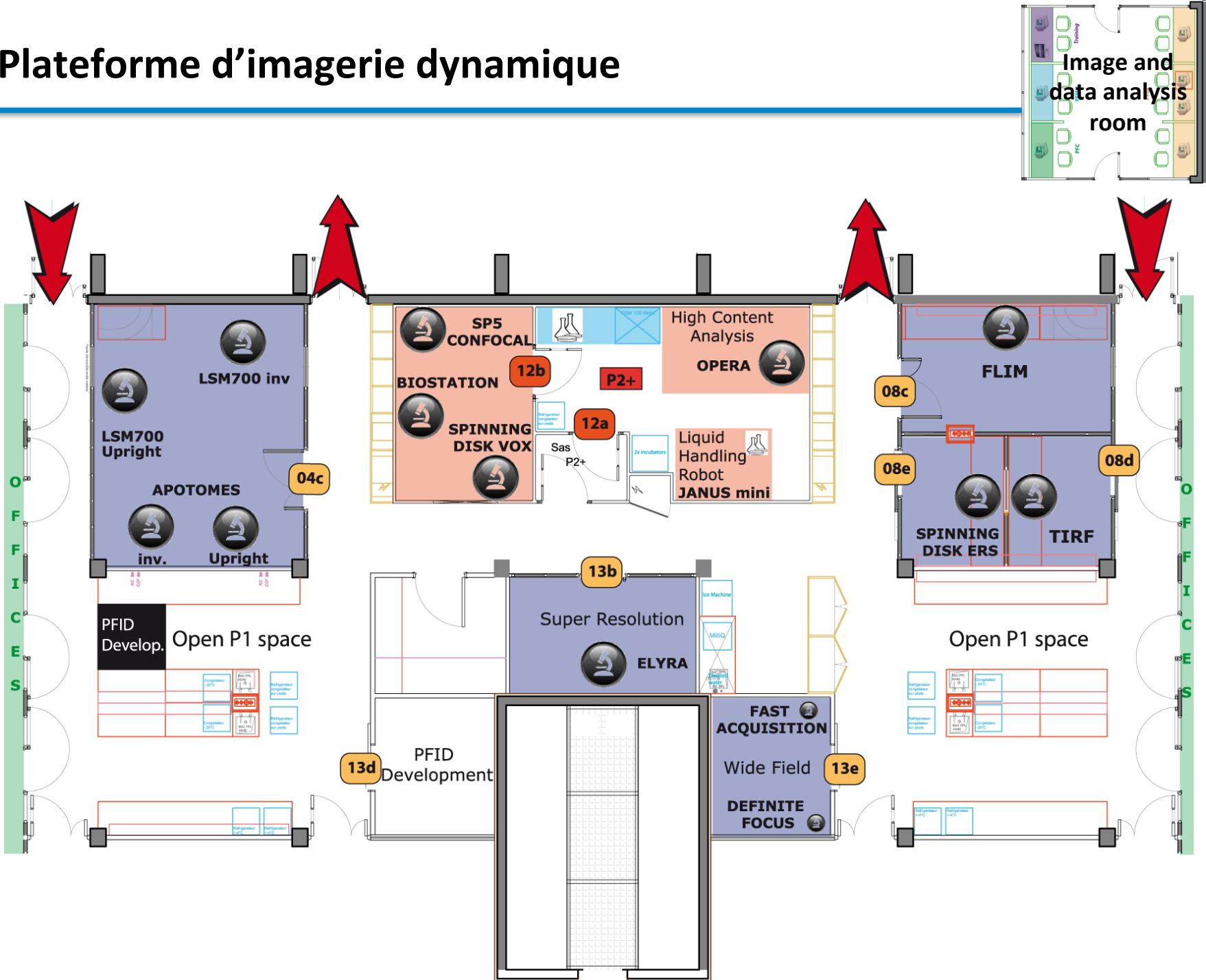


Imagopole mission:

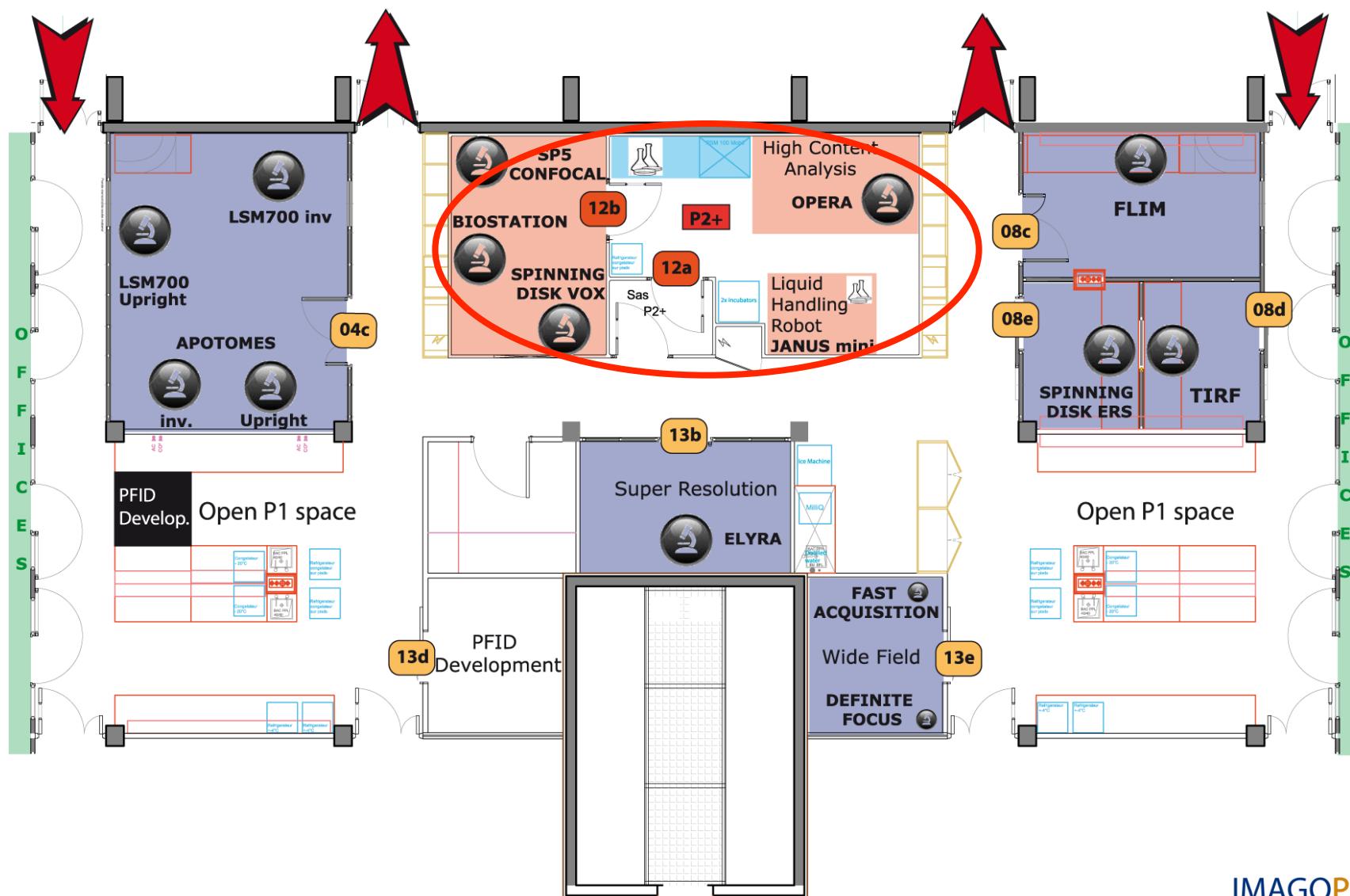
Apply cutting edge “imaging” science & technologies to studies on the dynamics of biological processes and their usurpation by infectious disease

“Imaging” describes an emergent set of experimental tools...

Plateforme d'imagerie dynamique



BSL2+ Equipment for live cell imaging in relevant models



New animal facility: BIME level -2 (BSL3)

2x IVIS Spectrum

- Filters 20nm band pass from 400nm to 800 nm for fluorescence and bioluminescence
- 5 mice at a time
- second system equipped with a CT scanner



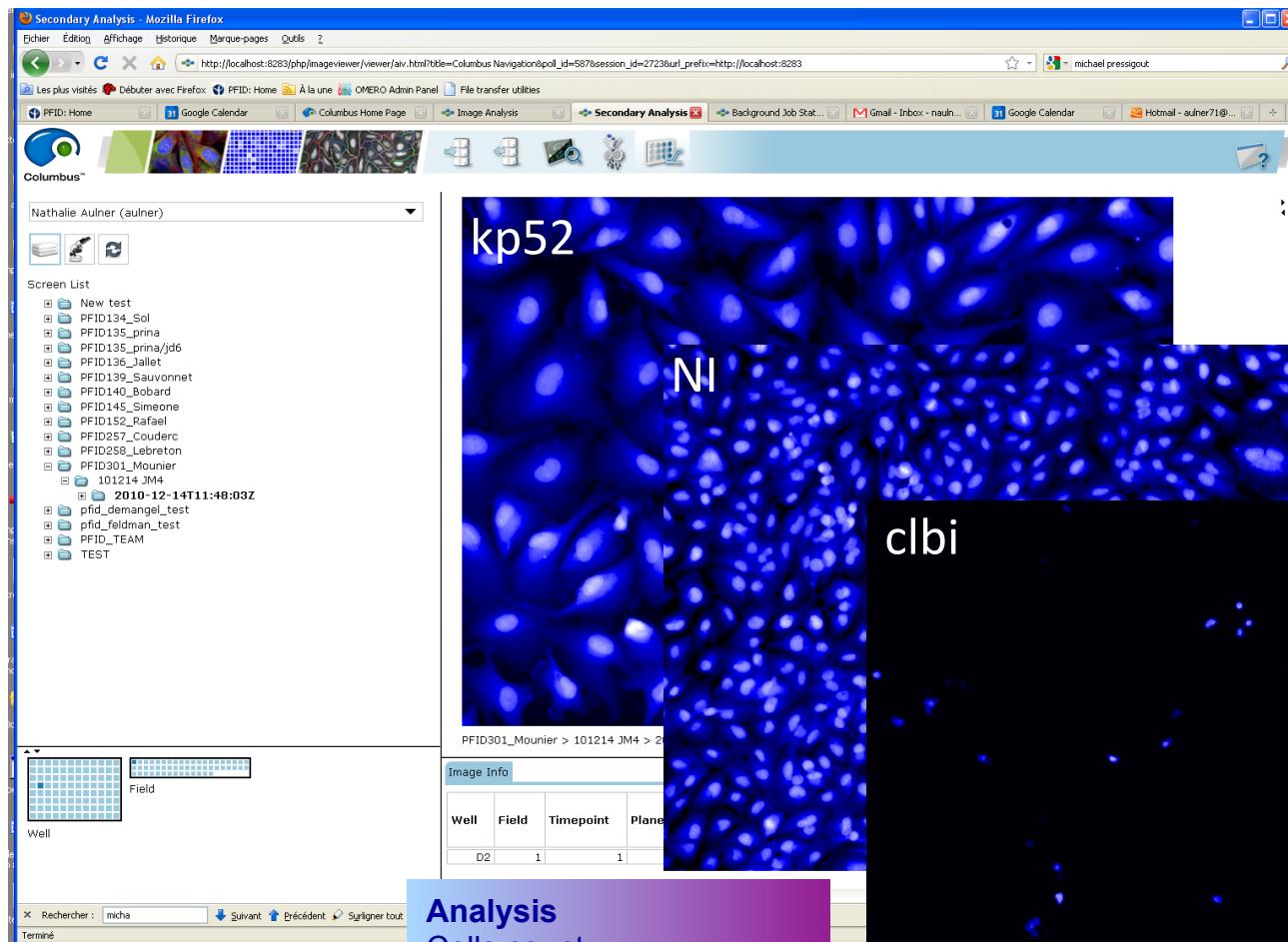
1 FMT (PerkinElmer)

- 3D NIR fluorescence
- 1 animal at a time



Possible under development system (longer term)

- Combined Fluorescence and bioluminescence



Secondary Analysis - Mozilla Firefox

Bilier Édition Affichage Historique Marque-pages Quits

http://localhost:8283/php/imageviewer/viewer/aiw.htm?tblle=Columbus Navigation&spoll_id=587&session_id=2723&url_prefix=http://localhost:8283

michael pressigout

PFID: Home Google Calendar Columbus Home Page Image Analysis Secondary Analysis Background Job Stat... Gmail - Inbox - nauln... Google Calendar Hotmail - aulner71@...

Columbus™ Nathalie Aulner (aulner)

Screen List

- New test
- PFID134_Sol
- PFID135_prina/d6
- PFID136_Jallet
- PFID139_Sauvionnet
- PFID140_Bobard
- PFID145_Simeone
- PFID152_Rafael
- PFID257_Couderc
- PFID258_Lebreton
- PFID301_Mounier
- 101214_JM4
- 2010-12-14T11:48:03Z
- pfid_demangel_test
- pfid_feldman_test
- PFID_TEAM
- TEST

Well

Rechercher : micha

Terminé

kp52

NI

clbi

PFID301_Mounier > 101214 JM4 > 2010-12-14T11:48:03Z

Well	Field	Timepoint	Plane
D2	1	1	

Analysis

Cells count

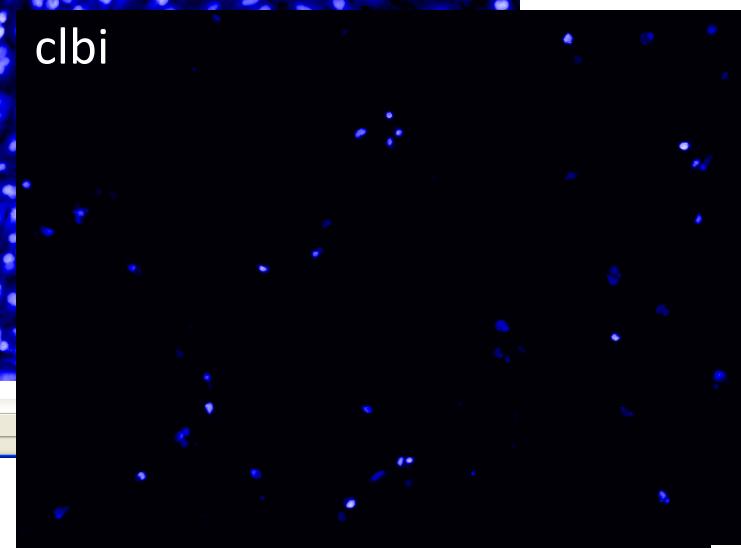
Fitness of cell population

Cells Size

Subpopulation analysis

...

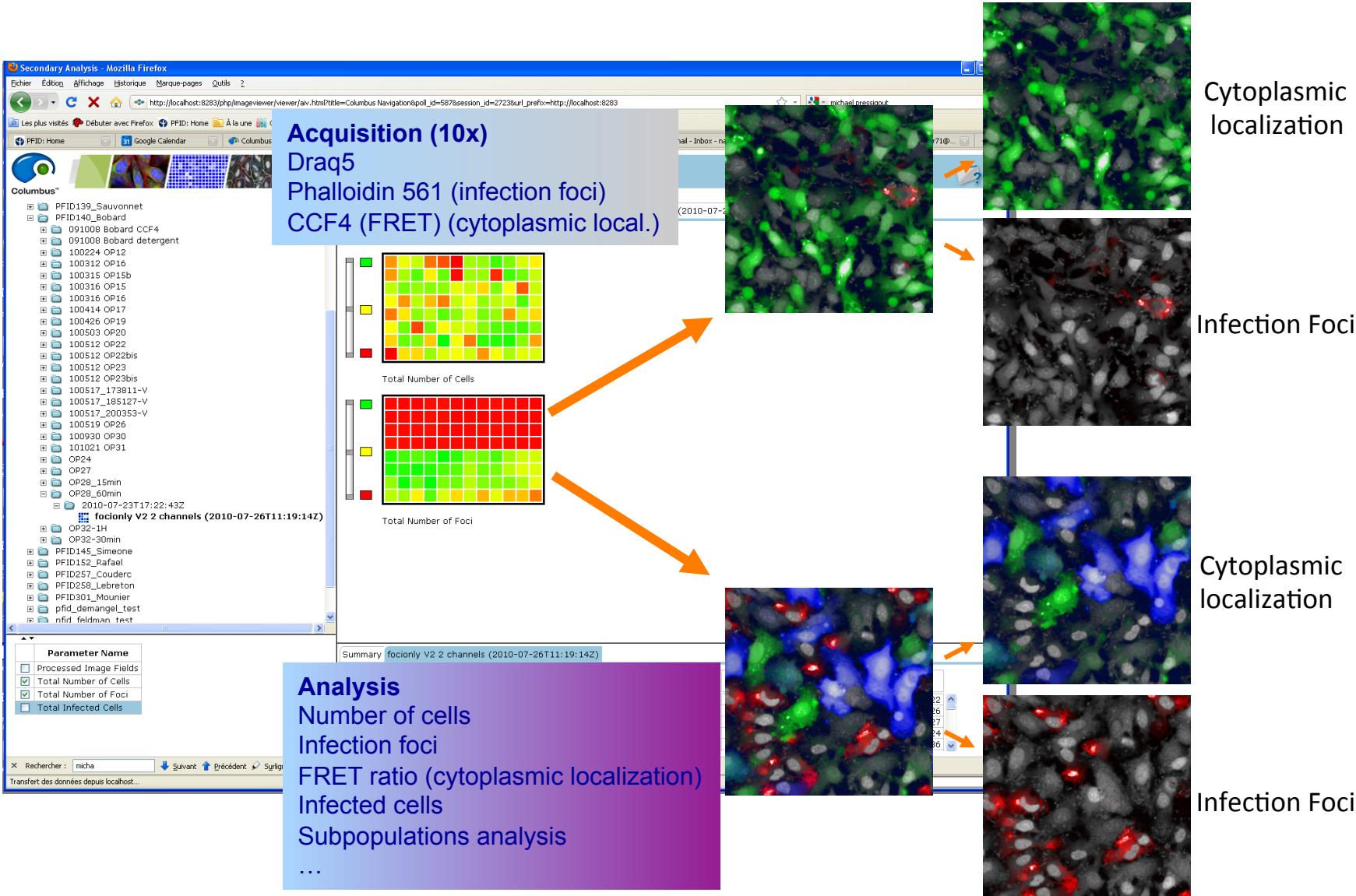
Acquisition 10x bin2
Hoechst
HC Cell Mask Blue



IMAGOPOLE
Pôle de Dynamique Moléculaire et Fonctionnelle

Plate-forme de Cytométrie en Flux
Plate-forme d'Imagerie Dynamique
Plate-forme de Microscopie Ultrastructurale
Plate-forme Centre d'Immunologie Humaine

Development of a high throughput siRNA screening for deciphering shigella's infection behaviour



Research & Development

Assembly and signalling mechanism of IL-7 receptor signalosome regulating CD4+ T-cell homeostasis and its alteration in HIV-infected patients

Blanche Tamari, Florence Gajdaj, Nadège Lefebvre, Céline Cullinan, Pauline Jarry, Marine Thiebaud, Thierry Rose
Unité d'Immunologie Cellulaire, Biopuces, Institut Curie, Paris, France

Audrey Sales, Isabelle Dragavon, Jean-Yves Timmer, Spencer Shorte, FIDM-C, Gérald Monneret, David DiGregorio, Imagine de l'Institut du Radium, Paris, France

1- Biological project

a) Direct electron microscopy (DPIE) of T cell

b) Immunogold labelling

c) Model and molecular pathway

d) Schematic diagram of the IL-7 receptor signaling pathway showing the assembly of the signalosome.

e) Topographic representation of the architecture of signaling molecules.

f) Schematic diagram of the IL-7 receptor signaling pathway showing the assembly of the signalosome.

Legend: Membrane, cytoskeleton, actin, or core of signaling molecules; Topographic architecture of signaling molecule; Systemic & functional.

2- Super-resolution microscopy

a) Fluorescence microscopy image of a T cell nucleus.

b) Schematic of super-resolution microscopy.

c) Schematic of super-resolution microscopy.

3- SIM Structured illumination Microscopy

a) Schematic of structured illumination microscopy.

b) Fluorescence microscopy images of T cells stained for Vimentin, Colocalization, and Structured illumination.

4- PALM PhotoActivation Localization Microscopy

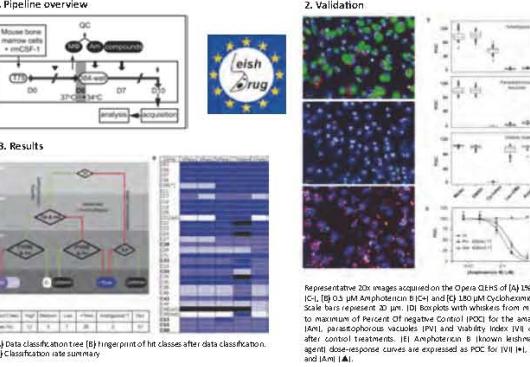
a) Fluorescence microscopy images of T cells stained for TH11 and PALM-TH11.

b) Schematic of photoactivation localization microscopy.

c) Fluorescence microscopy images of T cells stained for TH11 and PALM-TH11.

High content analysis of primary macrophages hosting *Leishmania* amastigotes: application to leishmanicidal agents identification

Nathalie Aulier, Anne Dancrae, Eline Rouault-Hardouin, Julie Desnivs, Olivier Helyck, Philippe Henri-Commeire, Hélène Munier-Lehmann, Gerald F. Spahl, Spencer L. Shantz, Olivier Milon et Eric Prina
Unité Pasteur-23-28 du Docteur Roux - 75013 Paris, Unité Institut Pasteur, (Inrapo)glo - Institut Pasteur, Laboratoire d'Immunobiologie et Pathologie, Département Pasteur-23 et Mycologie, Institut Pasteur, Unité Chimie et Bactéries, Département de Biologie Structurale et Chimie, CNRS UMR 3252, Institut Pasteur, Unité Pathologie moléculaire et Signalisation, Département de Biologie Structurale et Mycologie, CNRS URA 2581, Present address: Génopole SaS, Parc Brochet, 102 Avenue Gaston Rocard 92130 Nanterre, France



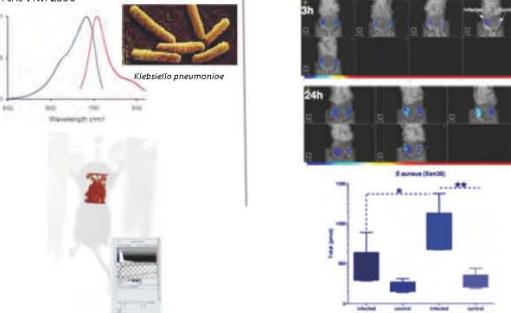
4. Conclusion
The present study, we developed a scalable throughput high content approach to select chemicals acting exclusively on L. monocytogenes amastigotes which are the causative agent of Leishmaniasis. The approach uses a library of 1000 compounds and a high content image analysis pipeline to identify active compounds without the need for Bead-Blanching which highly purifies populations of amastigotes, compound and fluorophore reporters are sequentially added to adherent amastigotes without any washing or addition of fresh medium. Applying these procedures allows reproducibly benefiting from high number of amastigotes having a unique lifetime, in absence of any extracellular amastigote. We also demonstrated that the presence/absence of PV was a valid indicator for anti-leishmania activity by correlating the PV and amastigote numbers per well in Amphotericin B dose-response experiments.
This approach can be used for drug screens over assays that have been described recently in the literature by allowing real-time monitoring and kinetic studies on living cells. A kinase inhibitor identification study has been recently launched under the auspices of the Leishmania project sponsored by the EU framework program.

FUEL takes advantage of an epifluorescent interaction between a living luminescent light source (i.e., bioluminescent bacteria) and a fluorophore (Figure 1) as a means to significantly increase the production of red photons (Figure 2). When applied to *in vivo* conditions, an overall increase in both the total and red photons is observed (Figure 3), increasing the efficiency and the contrast of bioluminescence imaging.

Fluorescence Tomography Imaging of bacterial infection

Abdessalem Rekiki¹, Samantha Blazquez¹, Marie-Anne Nicla¹, Joe Dragavon¹, Spencer L. Shorte¹, Philippe Sansonet², Régis Tournebise^{2,3}
(¹) Imappage, PFIID, Institut Pasteur, Paris
(²) Unité de Pathogénie Microbienne Moléculaire, Institut Pasteur, Paris
(³) Unité INSERM U788, Institut Pasteur, Paris

- | Infection : <i>Klebsiella pneumoniae</i> | Subcutaneous infection : <i>Staphylococcus aureus</i> |
|--|---|
| PLA.4 is a near-infrared fluorescent protein | Bald C female mice (7-8 weeks) |
| PLA.4 is expressed by <i>K. pneumoniae</i> under the control of a Lac promoter | Injected iIM with 1×10^7 CFU/mouse |
| Sub C mice were infected via the intranasal route with IFP-KP | 3 and 24h post infection, mice were injected with 5 nmoles of Bacterinsense |
| Image are imaged 1 and 2 days post-infection on the FMT2500 | 1h after agent injection, mice were imaged on the FMT2500 |
| | BLT mice were imaged at 0, 3 and 24h PI |



Fluorescence Tomography Imaging quantitatively permits 3D imaging of bacteria infection *in vivo*

Research & Development

Pilus structure in *Streptococcus agalactiae* : Immuno-labeling in SEM and TEM

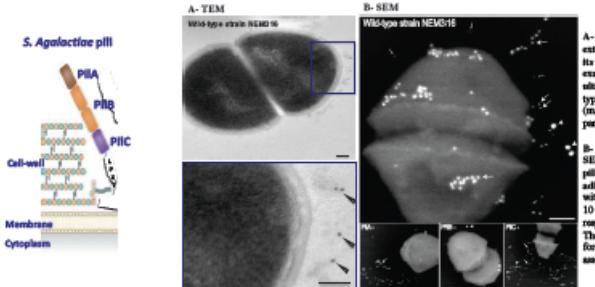
A. Mellot, Y Koste-Omairghz, E Meunier, G Daniellou, E Cézard, P Tissé-Castet, S Daniellou*

¹Institut Pasteur, Imaginepole, Département de Cell Biology and Infection, Paris, France

²Institut Pasteur, Unité de Recherche Birkhoff des Bactéries pathogènes et piliers, Département of Microbiology, Paris, France

³Institut National de la Santé et de la Recherche Médicale (INSERM) U570, Université Paris, Faculté de Médecine René Descartes, UMR5370, Paris, France

Adherence to host epithelial cells is the first critical step of the infectious process. Most bacterial pathogens have long filamentous structures known as pili which are often involved in the initial adhesion of bacteria to host tissues. The pilus of *S. agalactiae* is composed of three structural subunit proteins : Gba1478 (pIIA), Gba1077 (pIIB) the major component and Gba1474 (pIIC) analysed by immuno-electron-microscopy (IEM) using antibody conjugated with gold particles. We analysed a capsular type III polysaccharide (mAb 89) by TEM and we examined the pilus arrangement by SEM.



By using IEM we showed that the pilus is essential for optimal display of the pilus associated adhesin and overcomes the masking effect of the capsule. This method is a process currently used to localize proteins of interest and combine with the high resolution of SEM or TEM permitted to associate an immunolocalisation with a surface morphology or ultrastructural details.

*Dual role for pili in adhesion to epithelial cells and biofilm formation in streptococci epidermidis. Mellot A, Koste-Omairghz Y, Mellot E, Daniellou G, Cézard E, Tissé-Castet S, Daniellou S. PLoS Pathog. 2009 May 8;

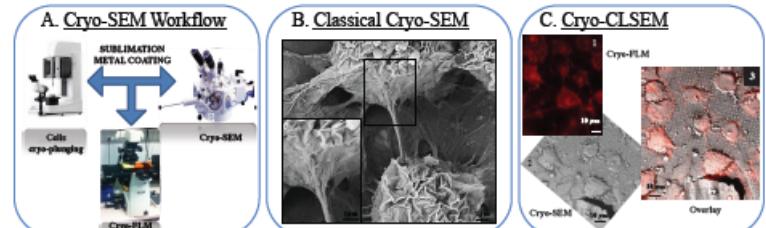
Cryo-CLSEM: Cryo-correlative Light & Scanning Electron Microscopy: Application to Tunneling Nanotubes

A. Mellot, S. Guératgeau*, P. Bourne*, C. Schmitt*, K. Ossouef*, C. Zanchi*, A. Sartori*

¹Institut Pasteur, Imaginepole, Department Cell Biology and Infection, Paris, France

²Institut Pasteur, Unité de recherche Toxicité métabolique et Pathogénie, Paris, France

Recently Tunneling Nanotubes (TNT) have emerged as a new way of inter-cellular communication for the transfer of various proteins, organelles & pathogens between cells. TNTs are transient, thin, very fragile membrane structures, detached from the substrate, that do not easily survive conventional electron microscopy preparation methods involving dehydration and resin embedding. Thus we used cryo-preparation and imaging methods to preserve their structural integrity and to observe them close to their native state. Specifically in this part of the project we use a Cryo-CLSEM approach, that combines fluorescent and surface ultrastructural information in cryo-conditions, to characterise the surface morphology of TNTs and their connection to the cell bodies.



Cryo-SEM workflow: osteohemangioma renal (CAD) cells were grown on EM grids and then cryo-fixed in liquid ethane. For cryo-SEM observations, the grid was submitted and coated with Au/Pd inside the cryo transfer module installed on a JEOL 6700F and then imaged at -150°C. For Cryo-CLSEM, cells were first imaged using a cryo-kidney which permit to image cells with fluorescence in cryo conditions.

Tokuyasu method applied to label an autophagy receptor and Chikungunya virus

N Cayet¹, D Judith^{2,3}, T Coudert^{2,3}, M Lescuit^{2,3} and MC Prévost⁴

¹Institut Pasteur, Imaginepole, Plate-Forme de Microscopie Ultrastructurale, Paris,

²Institut Pasteur, Microscopie et cellules souches Group, Paris,

³Inserm,夙寧 U624, Paris,

⁴Université Paris Descartes, Hôpital Necker-Enfants malades, Centre d'Infectiologie Necker-Pasteur, Service des Maladies Infectieuses et Tropicales, Paris-F

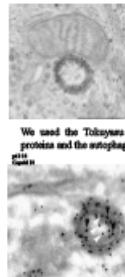
The Tokuyasu method uses a mild aldehyde fixation, which retains both morphology and antigenicity. The use of sucrose prevents ice crystal formation during the subsequent freezing procedure which hardens the sample and it can be cut into thin sections. The protein labelling at room temperature with antibodies allows the study of protein distribution at high resolution.

Essential steps of the Tokuyasu procedure for sectioning and immunolabelling.

- 1 Tissue processing
Aldehyde fixation followed by sucrose infiltration and freezing
- 2 Cryosectioning
Frozen sections are cut with dry knife (gliss or diamond)
- 3 Section retrieval
Sections are picked up, thawed on a droplet surface, and deposited onto grids
- 4 Immunolabelling
Grids, section side down, are passed over successive drops of solution to perform immunolabelling
- 5 Fixative contrasting, embedding and drying
Grids are contrasted with uranyl acetate and lead citrate, dried in a vacuum oven, and the sections become embedded in the thin film after drying *in situ*

(Liozu et al., 1996)

Morphology of the intracellular Chikungunya replication sites



Chikungunya virus (CHIKV) is a recently re-emerged arbovirus responsible in 2005 for a massive outbreak in India and islands of the Indian Ocean.

HeLa cells were infected with CHIKV for 20h and prepared for classical TEM. Small vesicles containing and surrounded by macroscopic

virions were detected. Scale bar = 100nm.

We used the Tokuyasu technique to study the spatial relationship of viral proteins and the autophagy marker p62.
p62
p62
HeLa cells were infected with CHIKV for 20h and fixed for immuno-EM. Small vesicles containing and surrounded by macroscopic were detected. p62 (black arrowheads) and capsid (white arrowheads) were labelled with appropriate primary antibody followed by protein-A-gold of different sizes respectively.
Scale bar = 100nm.

By immuno-EM we showed co-localisation of the autophagy receptor p62 and the capsid of Chikungunya virus.

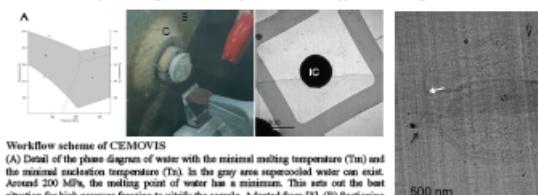
Virus Associated Pyramids in *Sulfolobus islandicus*: A TEM study for their formation

M Sadiou¹, T Quat², S Lucas², P Portere², D Prenguelli²

¹Institut Pasteur, Imaginepole, Department Cell Biology and Infection, Paris, France

²Institut Pasteur, Biologie Moléculaire du Génome des Extrémophiles, Department of Microbiology, Paris

The Archeal virus *Sulfolobus islandicus* rod shaped virus 2 (SIRV2) induces lysis of the host cell to release their virions. In the case of SIRV2 the virions assemble in the cytoplasm, where the major capsid protein forms the outer structure of the linear virus with the double stranded viral DNA in the interior. Parallel to virion assembly the formation of Virus Associated Pyramids (VAPs) takes place at the membrane of the infected cells [1]. We used Cryo-Electron Microscopy Of Vitreous Sections (CEMOVIS) to visualise the morphological changes during this infection process. CEMOVIS consists of cryo-immobilisation, cryo-sectioning, and low dose cryo-electron microscopy and offers a higher resolution than chemical prepared samples.



SIRV2 clusters in the cytoplasm
S. islandicus 10 p.i. with SIRV2. Before lysis of the cell and release of the viral particles (VP), the VP's are densely packed in the cytoplasm. Concentrally (white arrow) VAPs (black arrow) are induced. The VAPs protrude through the S-layer (arrowhead) of the cell. The inset shows a higher magnification of a VAP.

Workflow scheme of CEMOVIS
(A) Detail of the phase diagram of water with the minimal melting temperature (T_m) and the minimal nucleation temperature (T_n). In the gray area supercooled water can exist. Around 200 MPa, the melting point of water has a minimum. This sets out the best situation for high pressure freezing to vitrify the sample. Adapted from [3]. (B) Sectioning of the frozen sample. With an eyelash the sections are pulled over the grid. (C) The sections are observed in the cryo-microscope close to -180°C under low dose conditions. The picture shows an overview of the boundary between two sections. The different grey level between the two sections reflects the thickness of them. IC = ice crystal.

References:

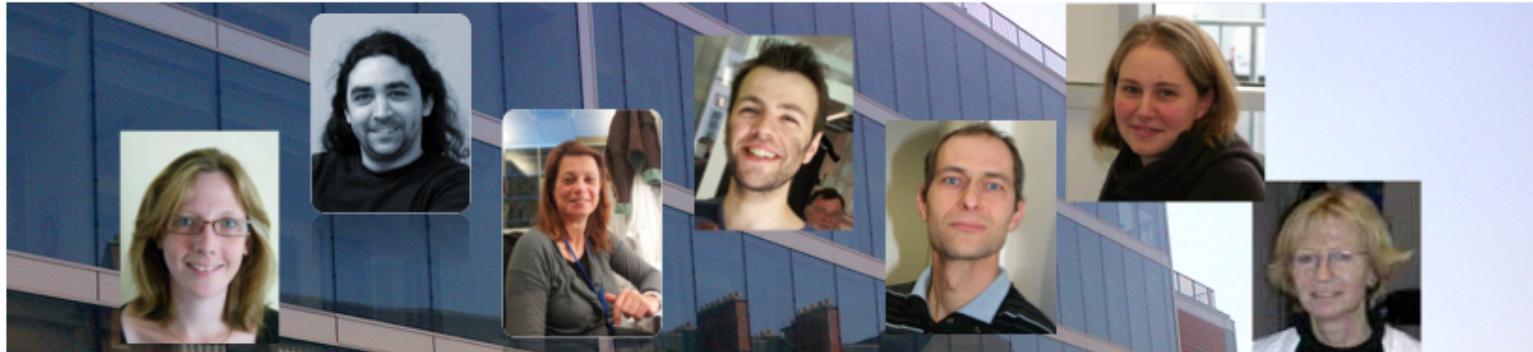
[1] Sims A, Kelman EA, Sheldrick K, Orme TE, Pines M, Prevost MC, Fontaine P, Tardieu G, Bernander R, Prenguelli D. Proc. Natl. Acad. Sci. (2009) 106:11396-11.

[2] This work was supported by the Programme Blanche of Agence Nationale de la Recherche (Grant ANR-09-BLAN-0288-01). T.Q. is supported by a fellowship from the Ministère de l'Enseignement Supérieur et de la Recherche of France.

[3] Verdicchio D, Gruber W, Stader D. Methods Cell Biol. (2008) p. 151-164.



Institut Pasteur



Federal
Ministry
of Education
and Research



FRANCE-BIOIMAGING



IMAGOPOLE
Pôle de Dynamique Moléculaire et Fonctionnelle

Plate-forme de Cytométrie en Flux
Plate-forme d'Imagerie Dynamique
Plate-forme de Microscope Ultrastructural
Centre d'Immunologie Humaine