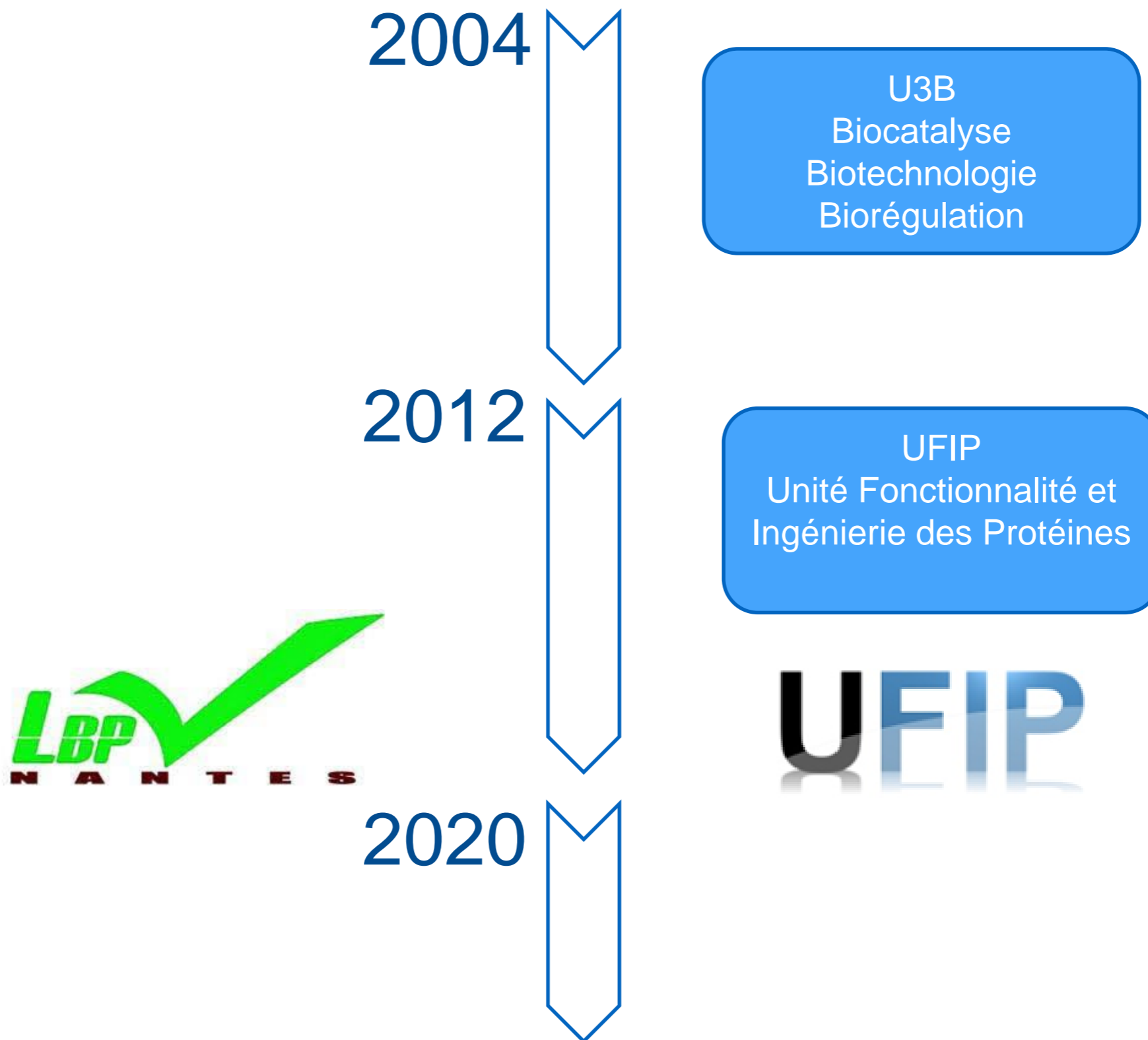
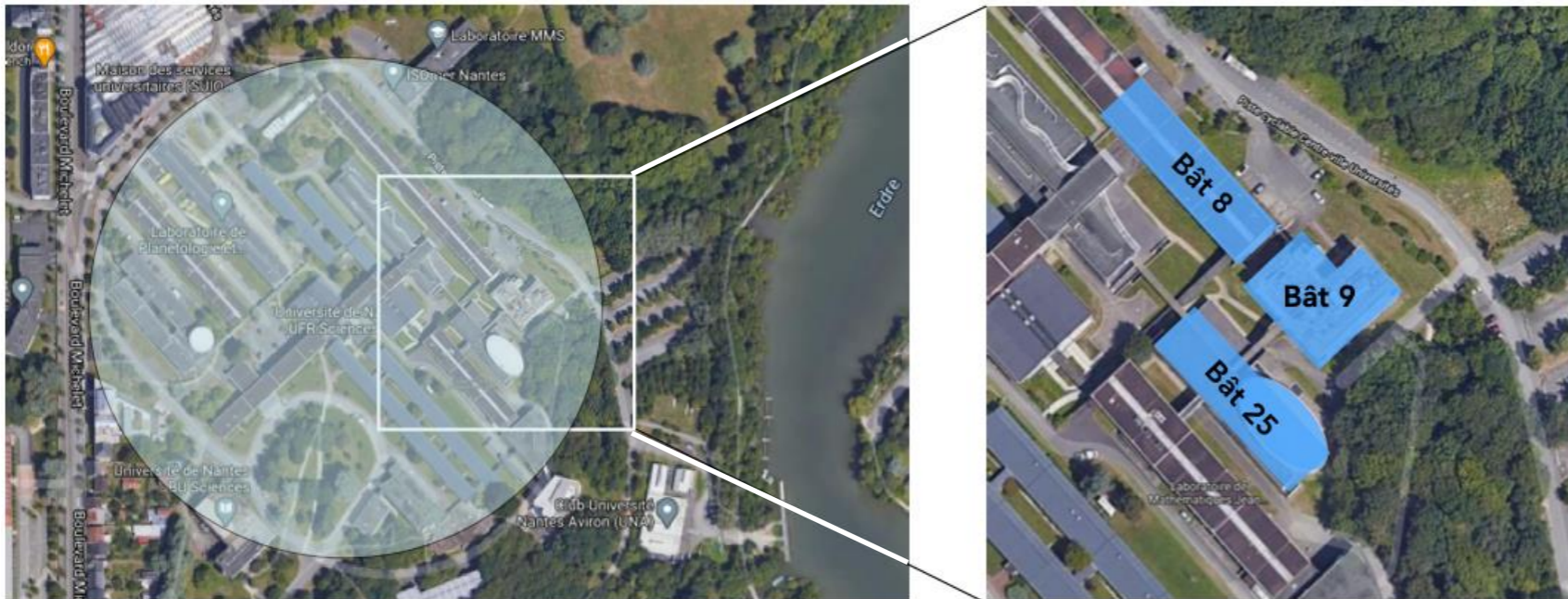


Unité en Sciences Biologiques et Biotechnologies

UMR 6286
CNRS / Nantes University

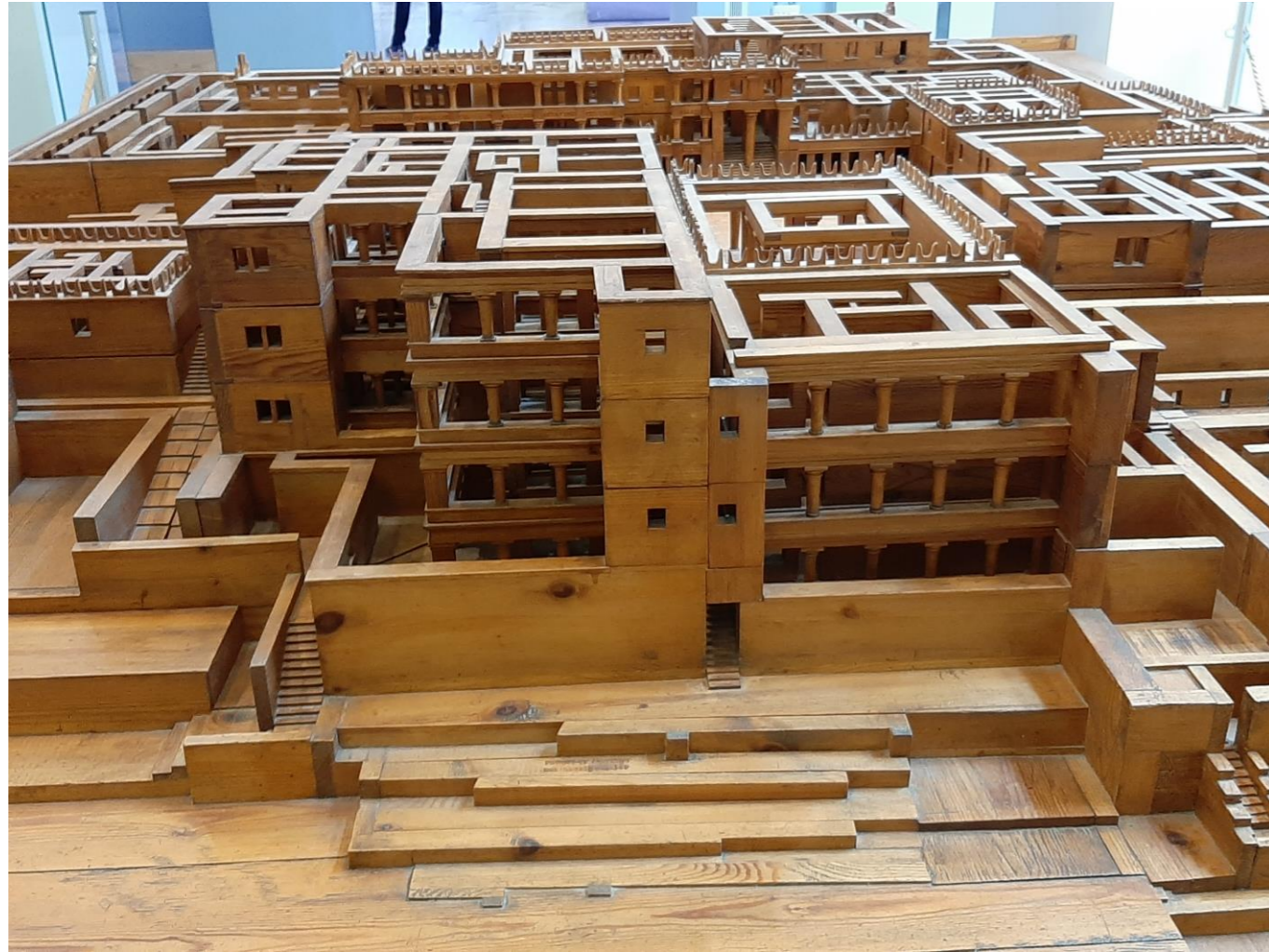


- Nantes University, Faculty of Sciences & Techniques, on the borders of Erdre river
- Institut de Cancérologie de l'Ouest (ICO)





Implantations

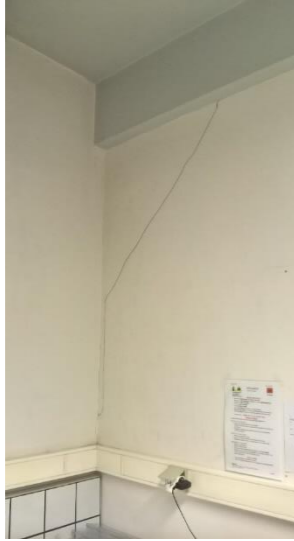
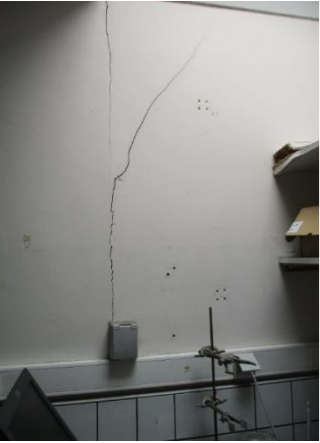






Implantations

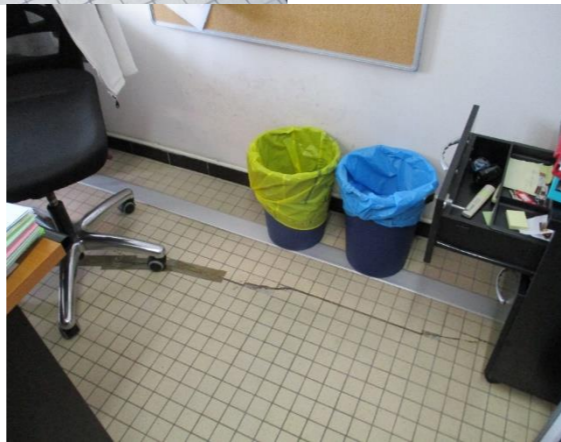
Laboratoires



Secrétariat

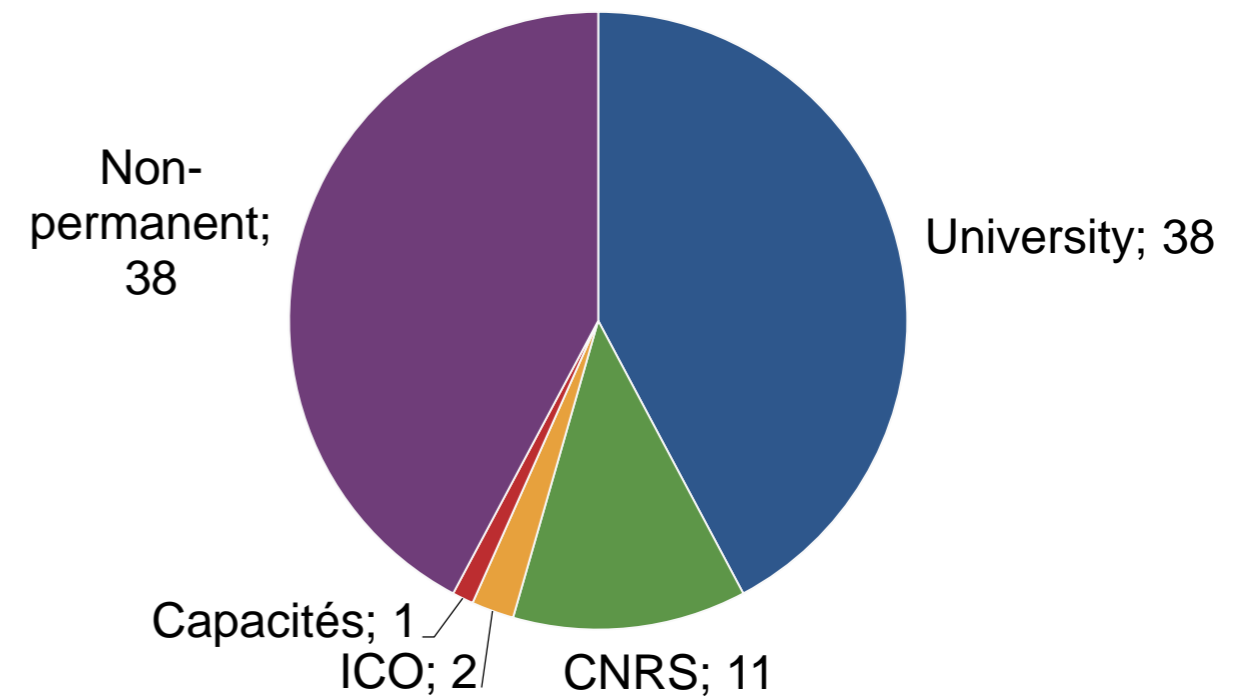


Bureau





- **Staff: 92 members**
(as from Sept 25, 2023)
 - **54 permanent members**
 - 25 university faculties
 - 4 CNRS researchers
 - 2 ICO researchers
 - 13 ITRF (1 admin)
 - 7 ITA CNRS
 - 1 Capacités
 - 2 COB
 - **38 non-permanent members**
 - 22 PhD students (13 foreigners)
 - 4 Post-Docs (2 foreigners)
 - 12 ITRF or ITA (2 admin)



Highly involved in training

Unité en Sciences Biologiques et Biotechnologies (US2B)

Directeur : Bernard Offmann (PU Univ) - Directeurs adjoints : Cyrille Grandjean (DR CNRS) et Philippe Simier (PU Univ)

Équipe de
Bioinformatique
Structurale
(YH Sanejouand, DR
CNRS)

Ingénierie moléculaire et
glycobiologie
(C. Grandjean, DR CNRS)

Mécanismes et
régulation de la
réparation de l'ADN
(F. Fleury, PU Univ)

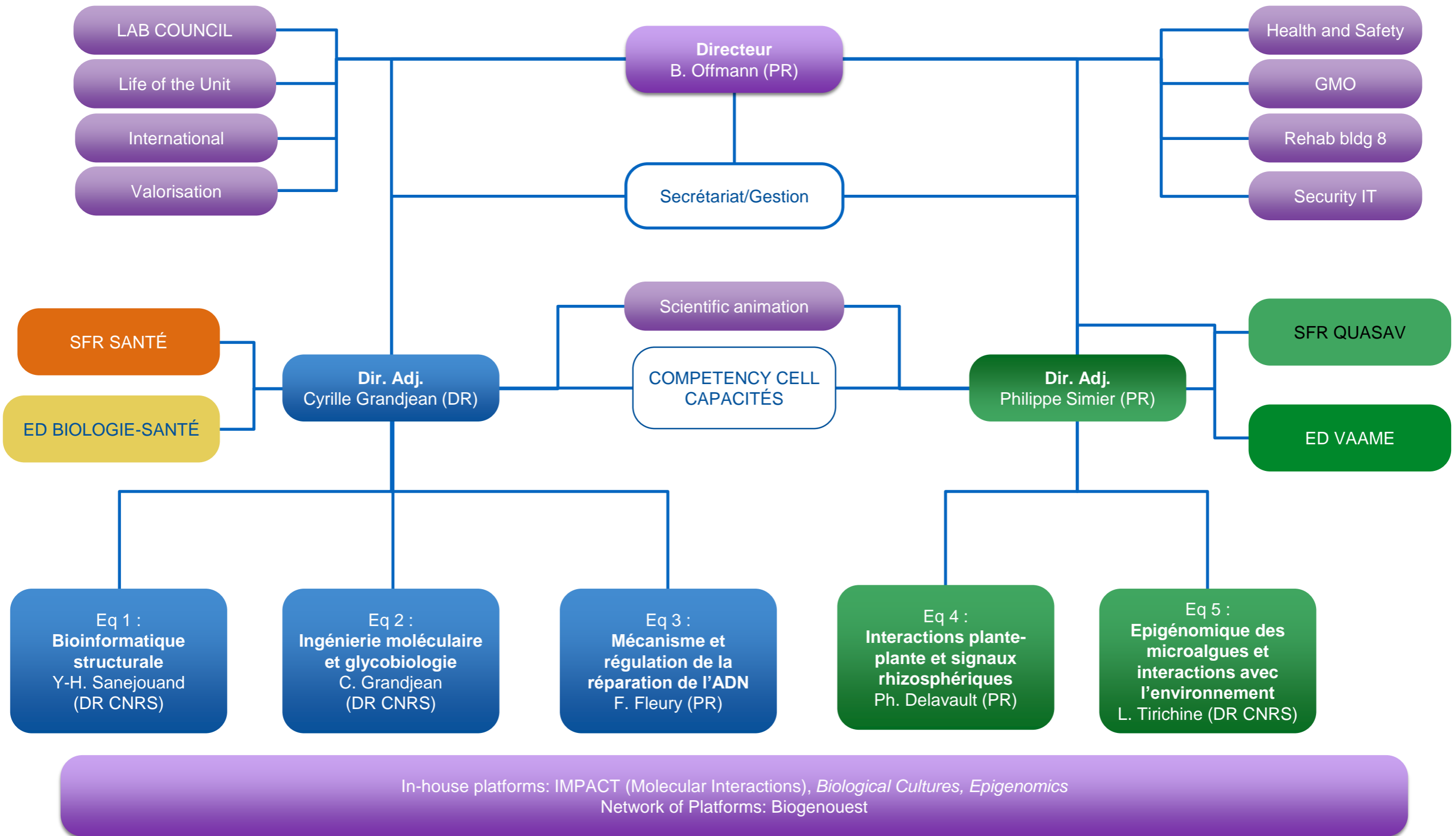
Épigénomiques des
microalgues et
interactions avec
l'environnement
(L. Tirichine, DR CNRS)

Interactions plante-plante
et signaux
rhizosphériques
(Ph. Delavault, PU Univ)

Plateforme IMPACT
(C. Charlier)

Cellule de compétences DZYME
(C. Grandjean)

Functional organisation

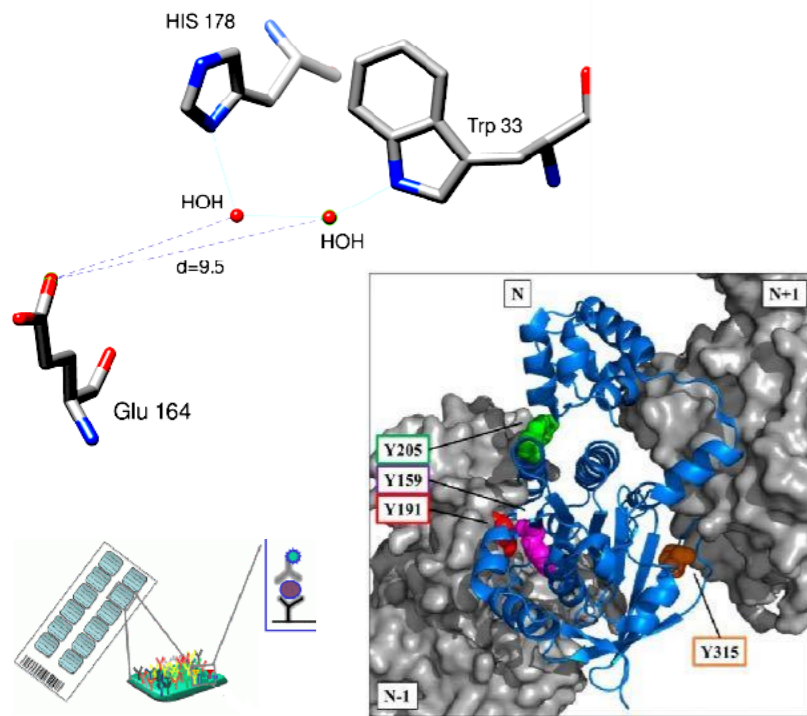


- **Our vision:** develop a high level **basic research** in **biochemistry** and **biology** (*in silico, in vitro, in cellulo & in vivo*) and keep a strong multidisciplinary expertise.
- **Our aim within next 5 yrs:** develop a high level **integrative biochemistry** and **biology** research for
 - deciphering how **protein function** and **biological activity** are regulated
 - developing **innovative biotechnologies**

Main topic Functions of proteins and bio-regulation

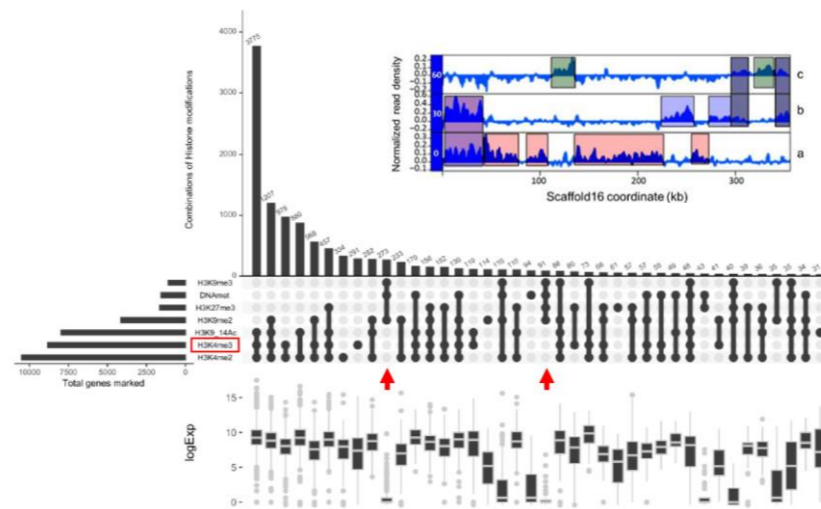
Protein biochemistry

regulation of protein function, structure prediction, molecular interactions, protein design

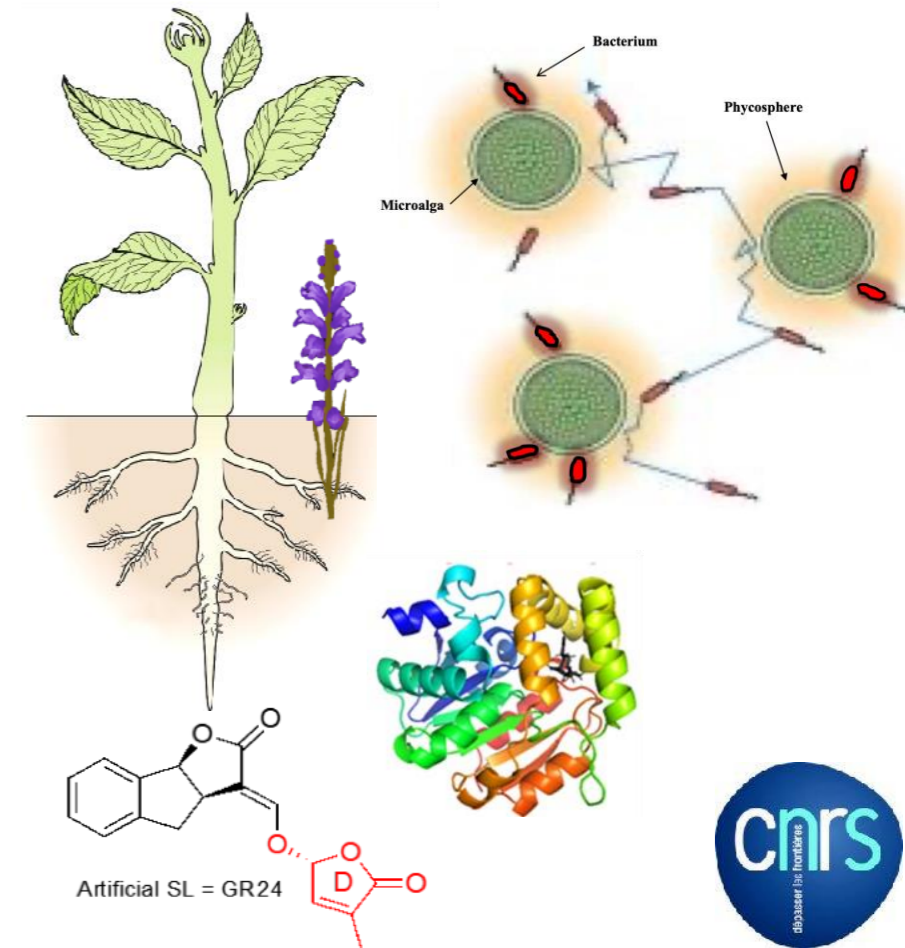


Epigenetic regulation

Epigenomics



Molecular basis of biological interactions



#4: Plant-Plant interactions and rhizospheric signals

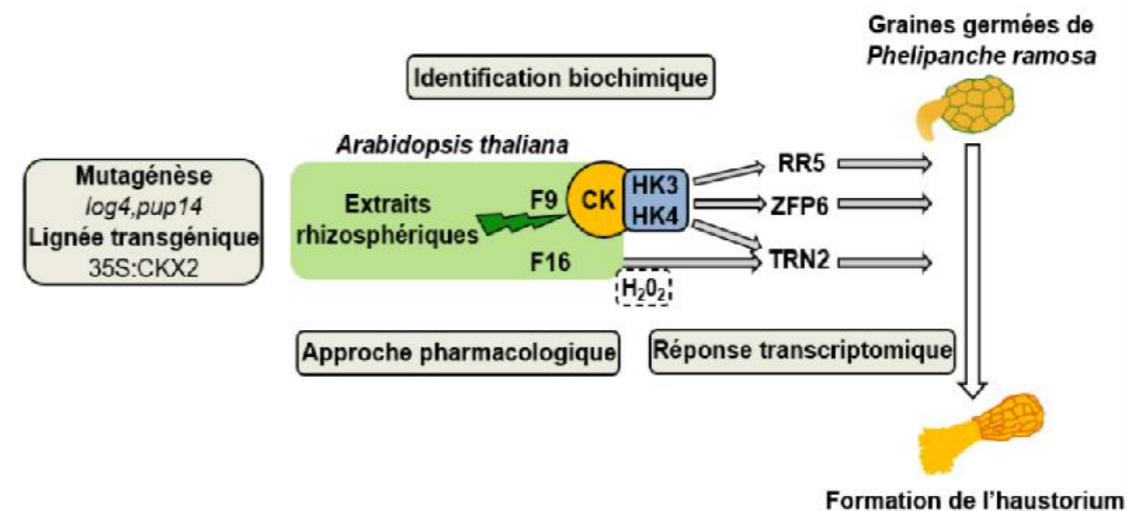
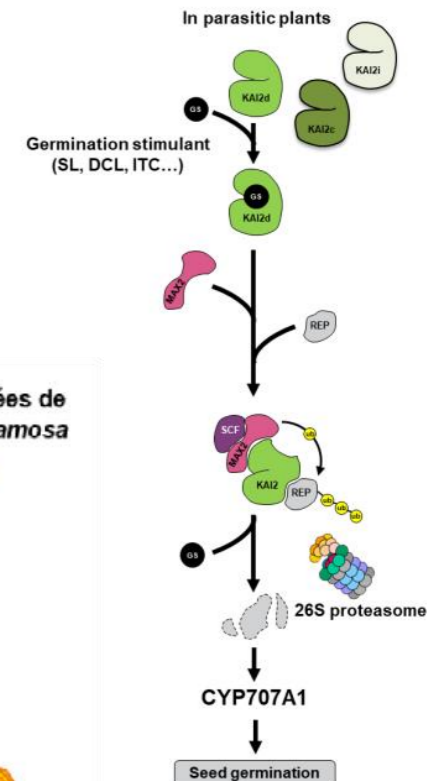
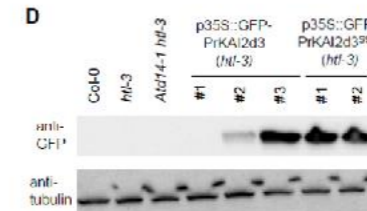
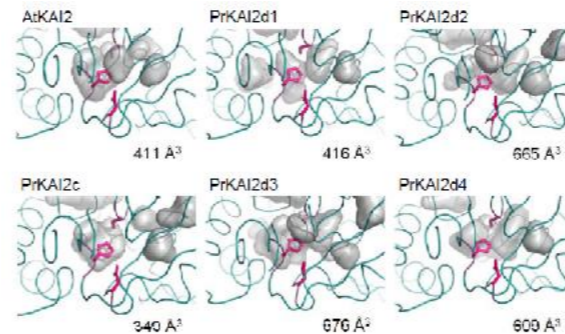
Philippe Delavault (11 permanent + 5 PhD members)

Philippe Delavault (PR)
 Hocine Benharrat (MCU)
 Grégory Montiel (MCU)
 Lucie Poulin (MCU)
 Jean-Bernard Pouvreau (MCU)
 Philippe Simier (PR)
 Séverine Thoiron (MCU)
 Christophe Véronesi (MCU)
 Sabine Delgrange (AJT)
 Adeline Dintheer (TCH CDD)
 Johannes Schmidt (TCH)

OBJECTIVES

- **Molecular determinisms of plant - parasitic plant interactions:** chemical factors and receptors involved
- **Involvement of rhizospheric microbiota:** reconstitution of tripartite model interactions
- **Genetic resources and applications**

- **Main expertises:**
 - Protein biochemistry, metabolomics, transcriptomics, plant biology and physiology, molecular biology, genetic engineering



#5: Epigenomics of micro algae and interactions with the environment

Leila Tirichine (4 permanent + 11 non-permanent members)

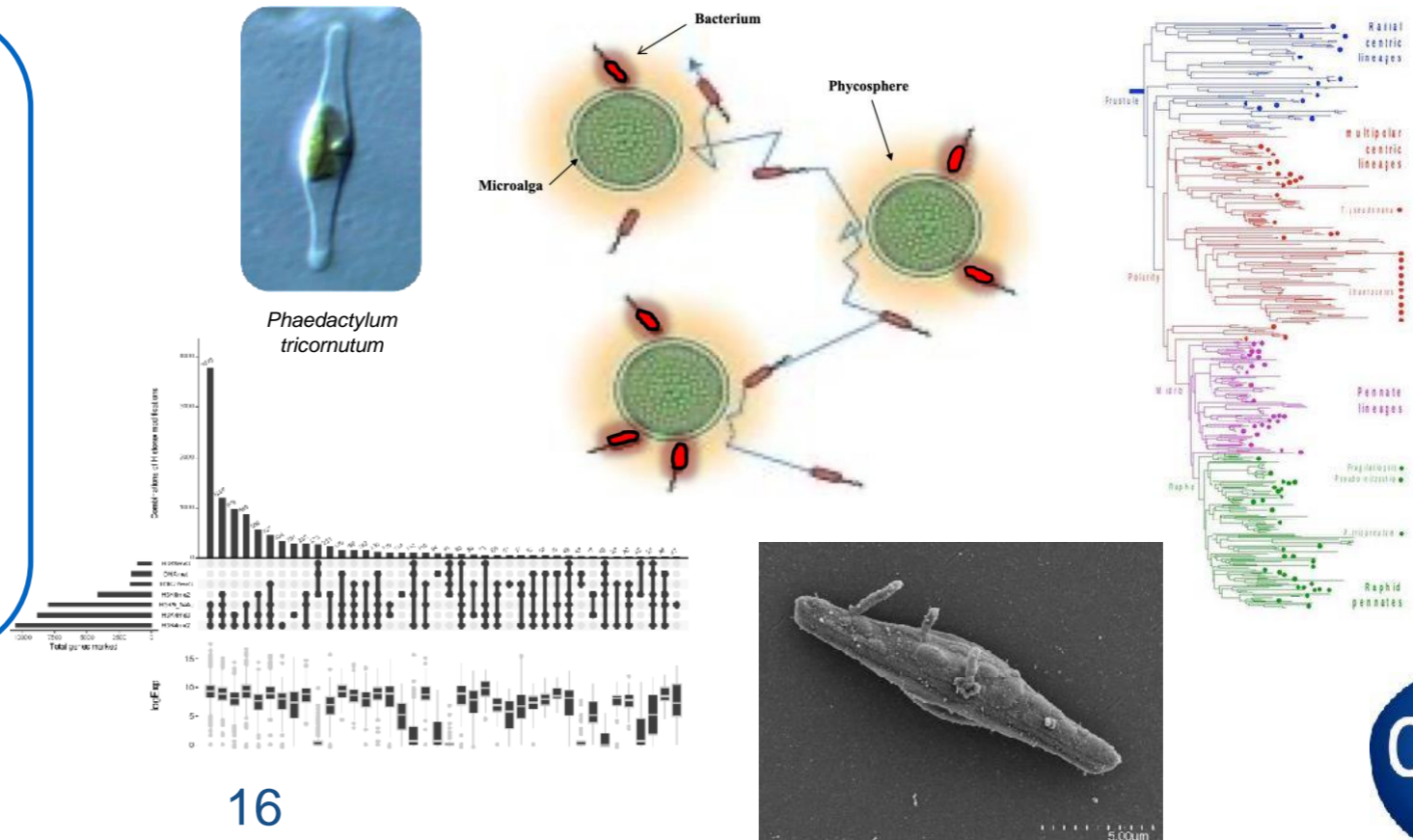
Leila Tirichine (DR CNRS)
Céline Duc (MCU)
Agnès Sebart-Groisillier (IR CNRS)
Carine Pruvost (AJT) 20%

OBJECTIVES

- ***P. tricornutum* epigenomics:** reference epigenome, natural accessions diversity
- **Molecular mechanisms of microalgae-bacteria cross talk:** for nitrogen fixation
- **Socio-microbiology:** phenotypic heterogeneity of nitrogen fixing bacteria in the phycosphere
- **100 diatoms genomes**
- **μAlgae factories**

• Main expertises:

- environmental epigenetics & epigenomics, bioinformatics, microbiology, cell biology, molecular biology, genetic engineering



Equipe 3

Mécanisme et régulation de la réparation de l'ADN

Fabrice Fleury (PR)

Houda Benhelli-Mokrani (MCU)
Christine Bobin-Dubigeon (MCU-PH)
Alain Defontaine (COB)
Yvonnick Chéraud (MCU)
Antoine Chalopin (PH ICO)
Dominique Heymann (PU-PH)
Marie-Christine Heymann (PU-PH ICO)
Dimitri Levitsky (COB)
Javier Munoz-Garcia (Ch ICO)
Vincent Potiron (Ch ICO)
Axelle Renodon-Cornière (CR CNRS)
Stéphane Supiot (PU-PH)
Pierre Weigel (MCU)

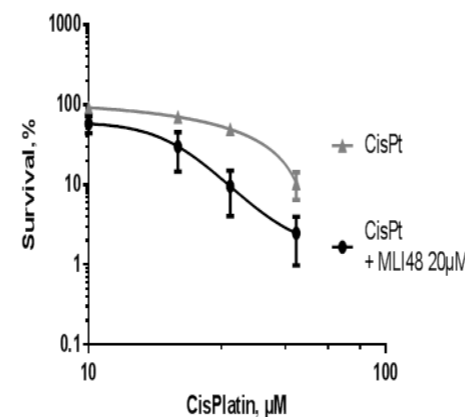
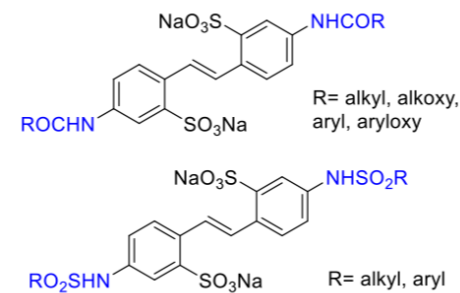
Franck Bertorelle (IR CNRS) 50%
Damien Marquis (TCH)
Vanessa Ménil (TCH CNRS)
Carine Pruvost (AJT) 40%
Céline Robiou du Pont (AJT)

OBJECTIVES

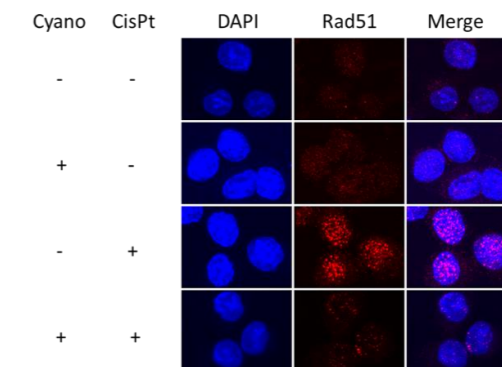
- **Modulation of Rad51 activity:** chemical inhibitors, (synthetic and natural molecules), *in vivo* screening
- **Role of a new phosphorylation of Rad51:** *in cellulo* studies, new cellular models (CRISPR/Cas9), appl. to TNBC screening
- **Biotechnological development:** carbon dots, biosensors, xenograft

Expertises

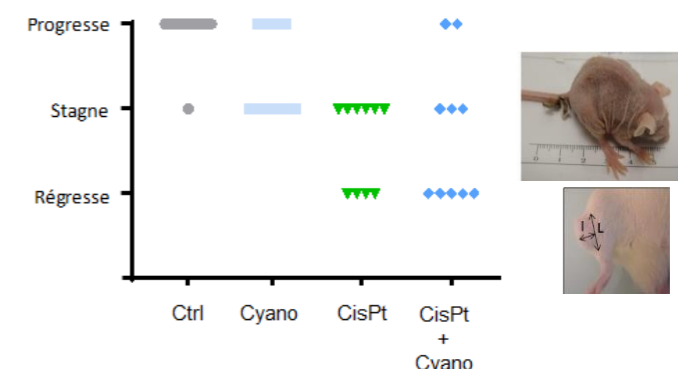
- protein biochemistry,
- cell biology, protein arrays,
- drug screening, biophysics
- nanotechnology



Clonogenic assay (CisPt treatment +/- MLI48 at 20µM)



RAD51 foci formation (CisPt treatment)



Tumoral progression (CisPt treatment +/- MLI48)

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Mécanisme et régulation de la réparation de l'ADN

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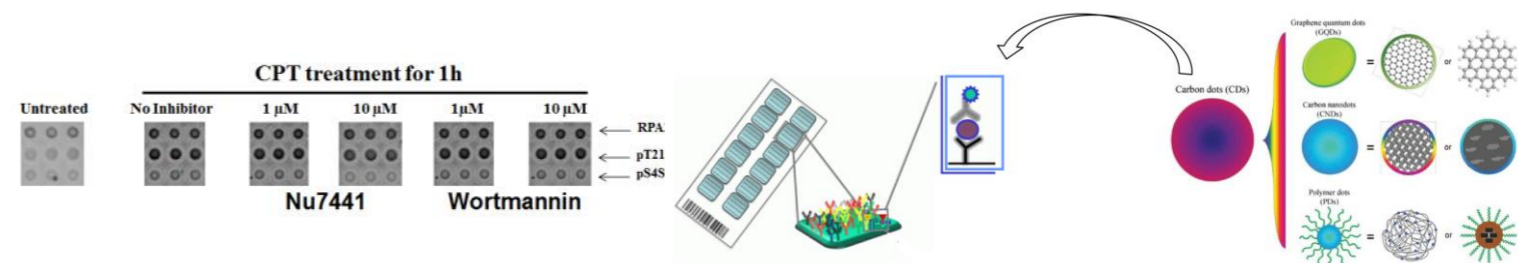
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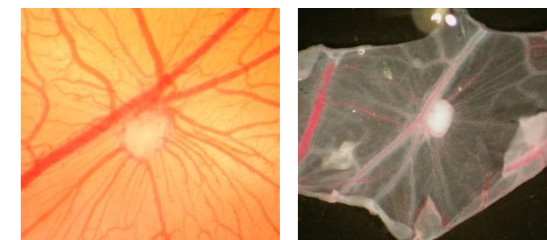
Expertises

- protein biochemistry,
- cell biology, protein arrays,
- drug screening, biophysics
- nanotechnology



Day 0 Day 8 Day 10 Day 17

Incubate eggs Graft DRG Graft cancer cells Harvest CAM tissue



EXPERTISE : PROTÉOMIQUE FONCTIONNELLE

PROFILING

Profils d'abondance protéiques

Profils des modification de protéines cibles (MPT)

CRIBLAGE MOLECULAIRE

Identification de modulateurs d'interaction : protéines, peptides, petites molécules

CARACTERISATION/VALIDATION DES INTERACTIONS

Affinité, K_{on} , K_{off} , stœchiométrie, contributions enthalpique et entropique

ANALYSE ET INTEGRATION DE DONNEES

Développement de nouveaux outils ✧Galaxy

LOCALISATION

2 sites

Unité de Biochimie et Ingénierie des Protéines
UFIP - UMR 6286 CNRS

Centre de Recherche en Cancérologie et Immunologie Nantes-Angers
CRCINA
INSERM UMR1232 CNRS
ERI.6001

17/03/2020



IMPACT
Interactions Moléculaires Pucés ACTIVités

Faculté des Sciences
Campus Lombarderie 2, rue de la Houssinière
44 000 Nantes



SFR Santé F. Bonamy
FFD 4203/TMS Inserm 016/CNRS 35565
IRS 2 - 22 Boulevard Benoni Goullin
44200 Nantes



Spotter seFLEXARRAYER S3
(Scienton)



Scanner Odyssey CLX
(LICOR)



SPR - Biacore 3000
Biacore 1200 (GE)



Technologies Alpha et Label Free
EnSpire (Perkin Elmer)

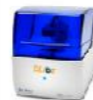
Protein arrays

SPRiplex II (Horiba)



- Profiling
- Criblage
- Caractérisation
- Contrôle qualité

Technologie BLI BLITZ System (FortéBIO - Full Life Sciences)



Fluorescence/anisotropie fluorescence
Spectrofluorimètre FP-6500
Photomètre de polarisation de fluorescence FP-715 (Jasco)



Dichroïsme Circulaire et linéaire
Spectropolarimètre J-810 (Jasco)



Microcalorimètre Auto ITC200 (GE)



HIRF / BRET / FRET / AF
Mithras LB940

INNOVATION

BIOPUCES ANALYTIQUES / DIAGNOSTIC

Diagnostic & recherche pharmaceutique

Nouveaux systèmes d'analyse

BIOPUCES FONCTIONNELLES

Interactions protéine-protéine, protéine-ligands

Etudes des MPT (Phosphorylation, glycosylation ..)

Puces à cellules/interactions aux interfaces

Etudes de complexes multimoléculaires

ANALYSE ET INTÉGRATION DE DONNÉES

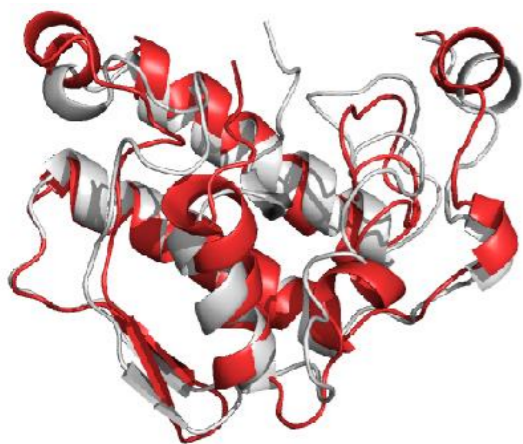
Données MTS et HTS issues des profils d'expression ou du criblage

Données publiques (ou analyses de la plateforme) et données virtuelles

Equipe 1

Bio-informatique structurale

Yves-Henri Sanejouand (DR CNRS)
 Bernard Offmann (PR)
 Stéphane Téletchéa (MCU)
 Johann Hendrickx (IE CNRS)
 Lionel Hoffmann (IE CNRS)



OBJECTIVES

- **Development of methods** to understand
 - the **sequence-structure relationship** (fold recognition, protein design)
 - **structure-flexibility-function relationship** (sampling methods, role of the zero-point energy)
- **Applications of standard methods:** molecular recognition, drug design, protein engineering

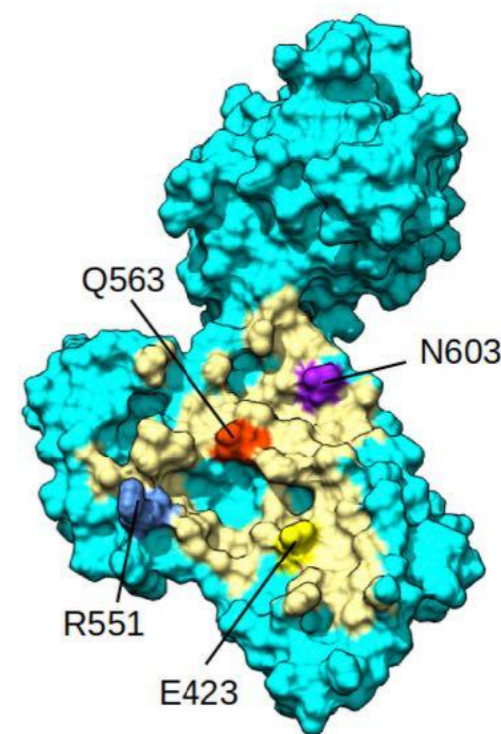
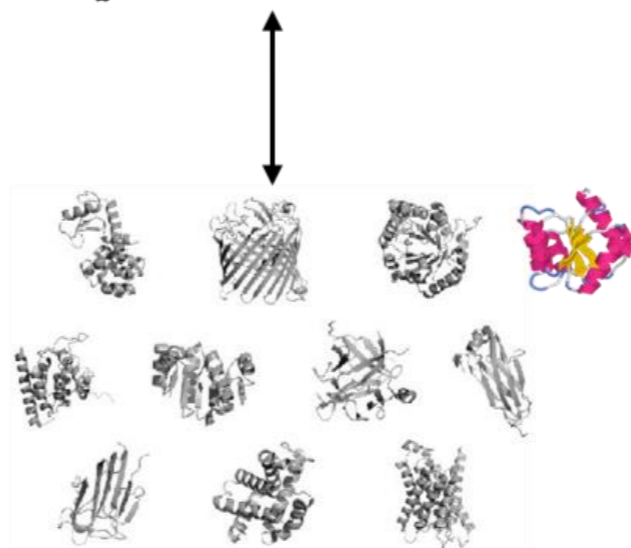
• Main expertises

- structural bioinformatics
- molecular modelling
- machine learning
- software development

$$A_{\text{vib}} = \sum \frac{1}{2} h\nu_i + \sum kT \ln \left(1 - e^{-\frac{h\nu_i}{kT}} \right)$$

Fold recognition

SNISRQAYADMFGPTVGDVKVRLADTELWIEVED
 DLTTYGEEVKFGGGKVIIRDGMGQGQMLAADCVD
 LVLTNALIVDHWGIVKADIGVKDGRIFAIGKAG
 NPDIDIQPNVTIPIGAATEVIAAEGKIVTAGAS



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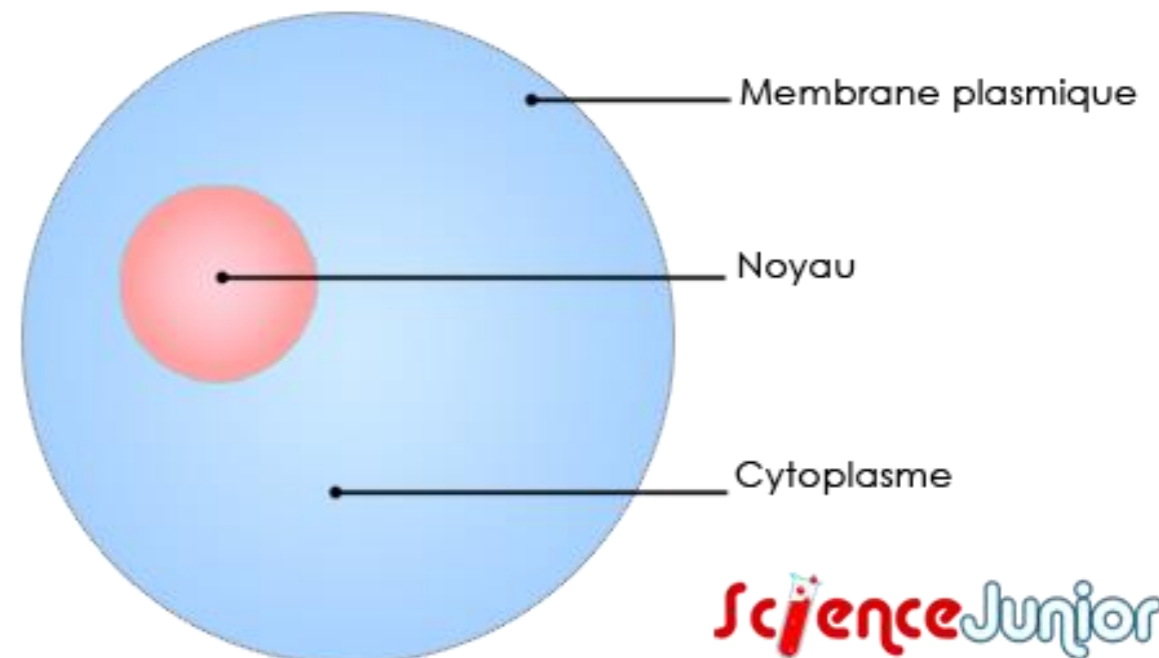
Claude Solleux (TCH)

OBJECTIVES

- **Glyco-enzymology:** monitoring and controlling enzyme activity
- **Glyco-recognition engineering:** inhibitors for galectins, lectin engineering
- **Anti-infectious glyco-technology:** glycoconjugate vaccines, bacterial sialidase inhibitors

• Main expertises:

glyco-enzymology, glyco-chemistry, protein biochemistry, molecular & metabolic engineering, nanotechnology



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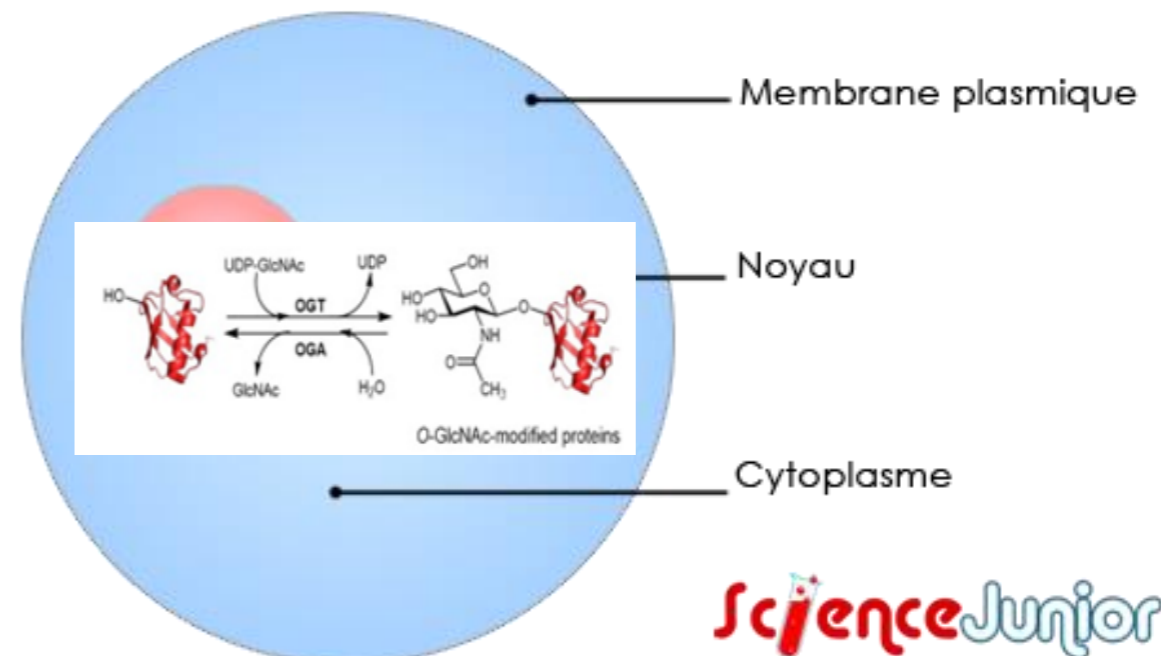
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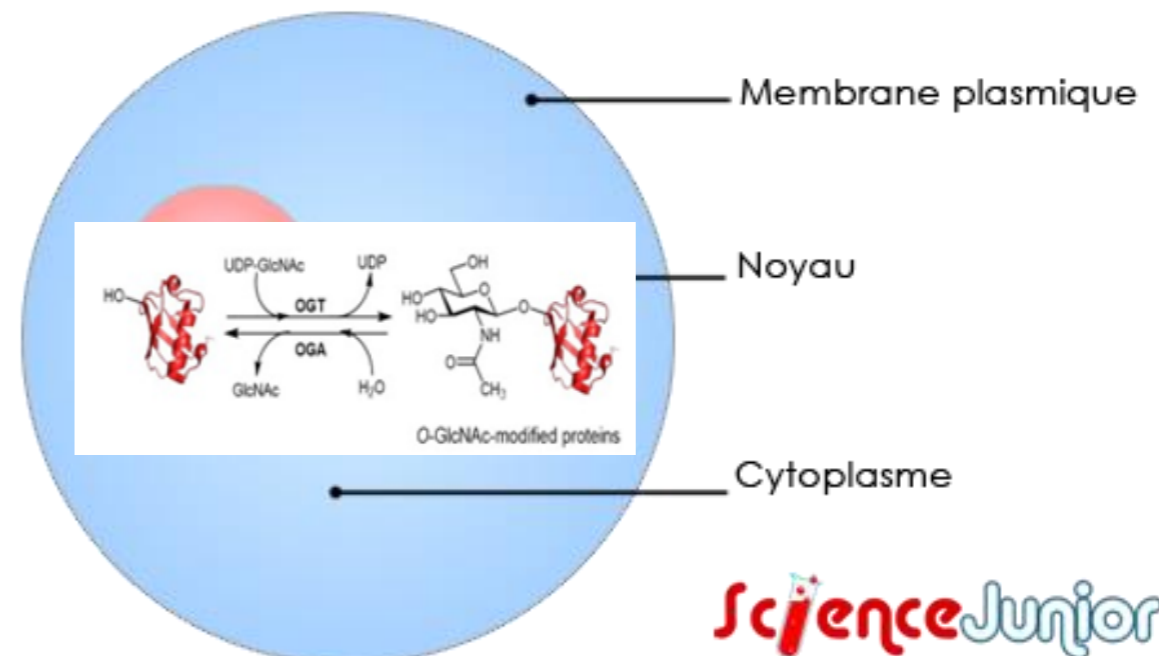
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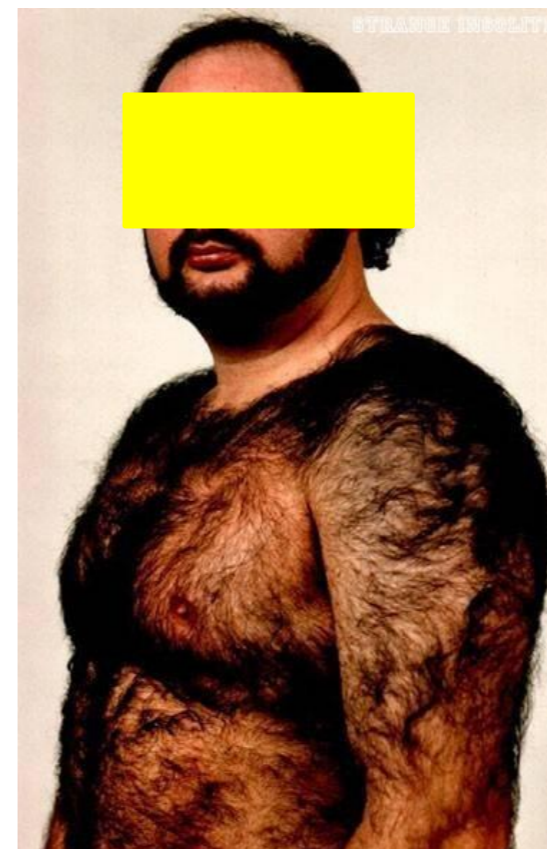
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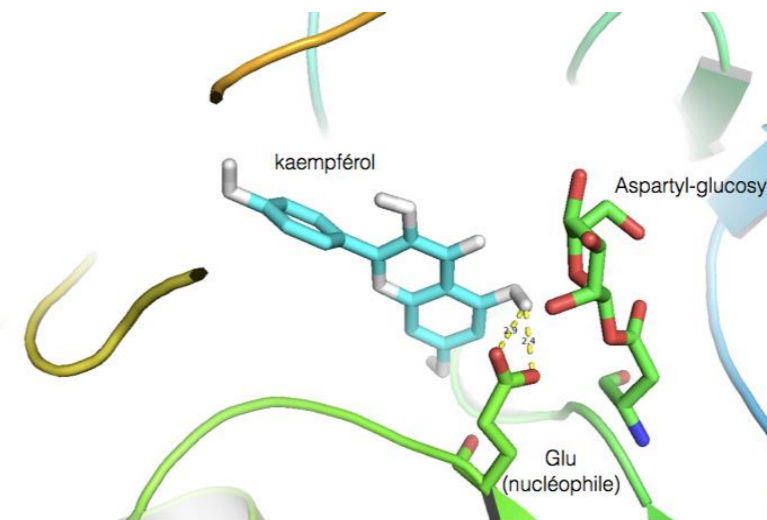
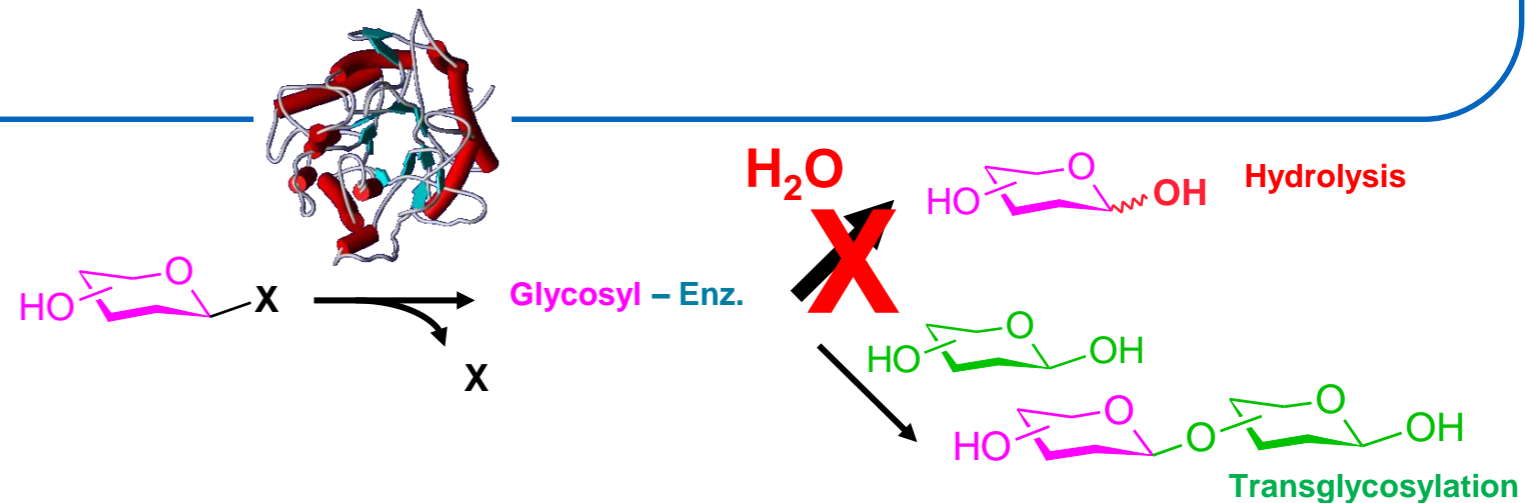
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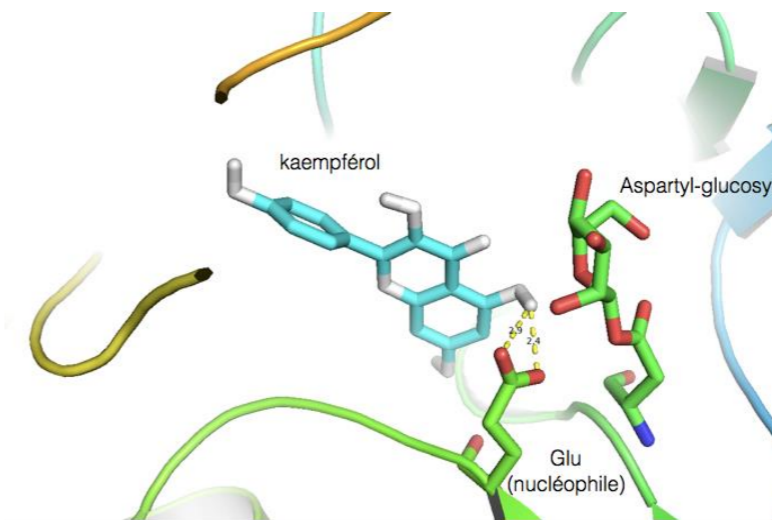
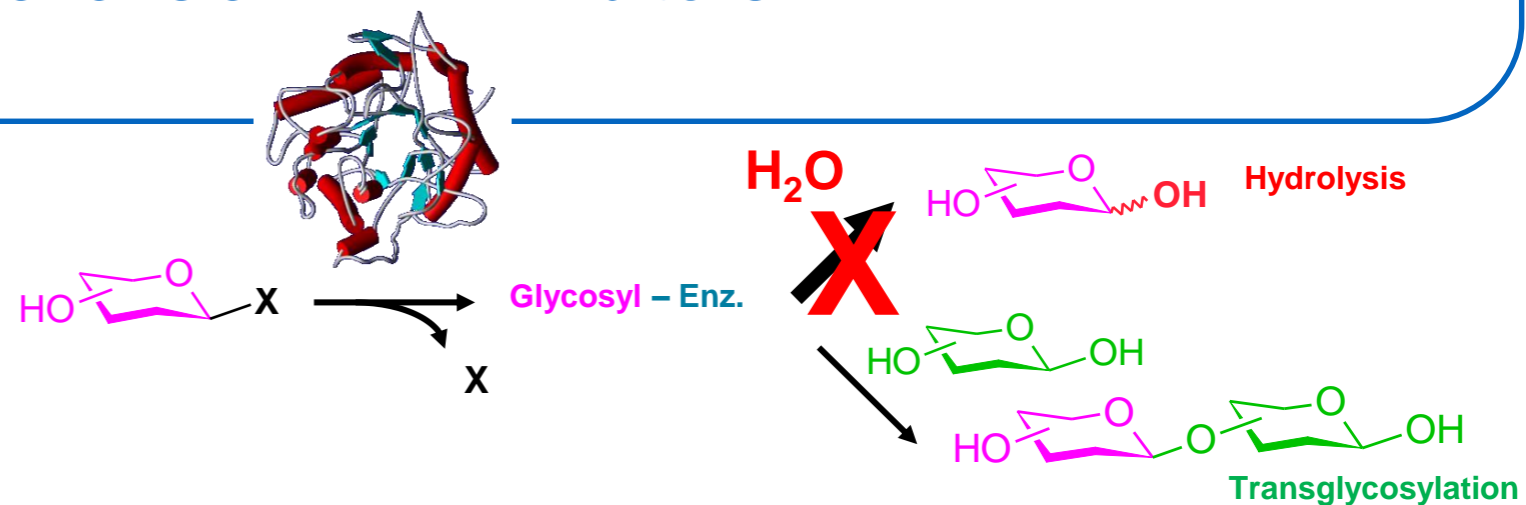
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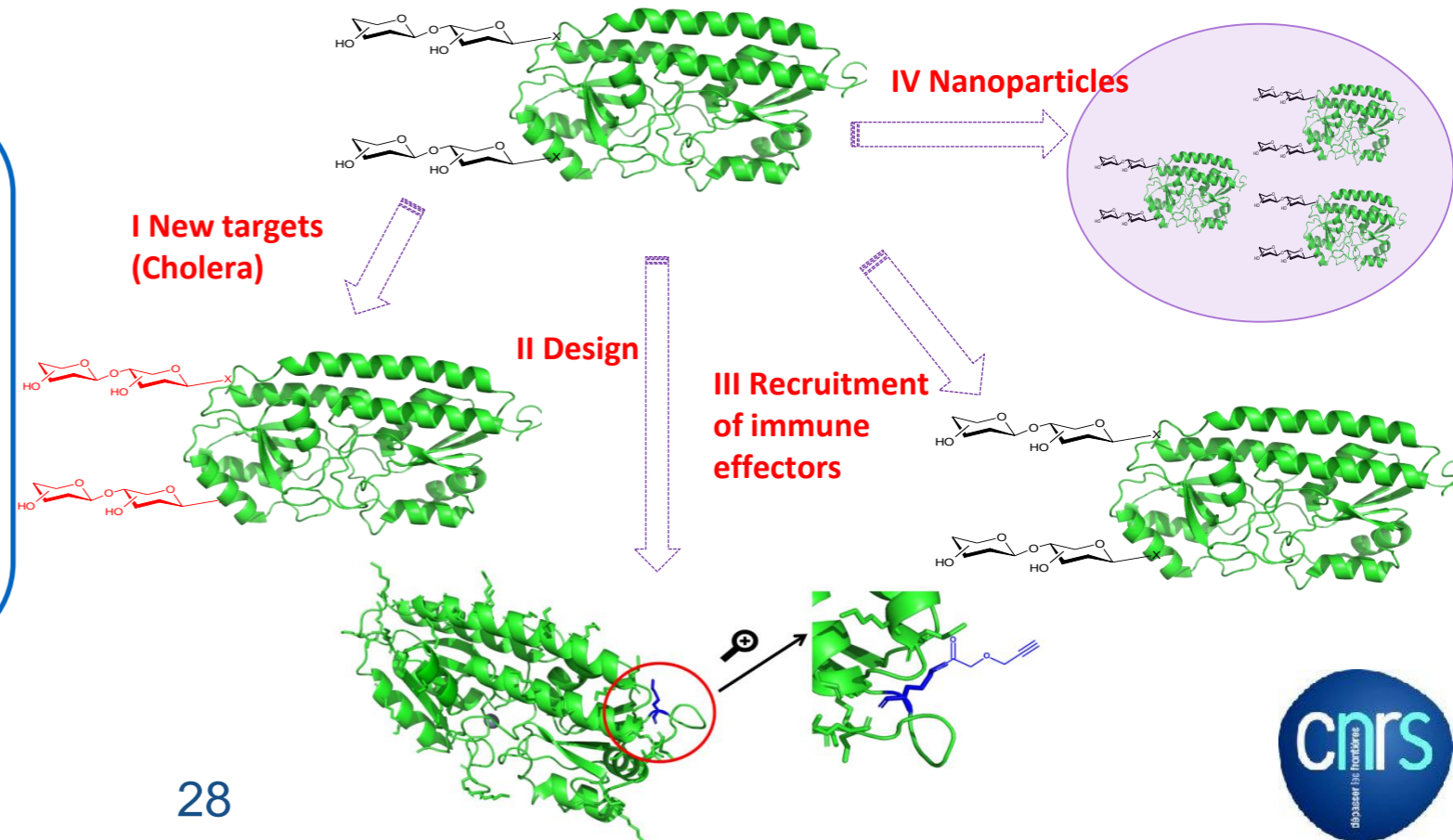
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DZYME (valorisation of our expertise)

Snapshot:

Enzymology & glycochemistry

Our services are based on the association of our expertise in enzymology and glycosciences

Supporting labs

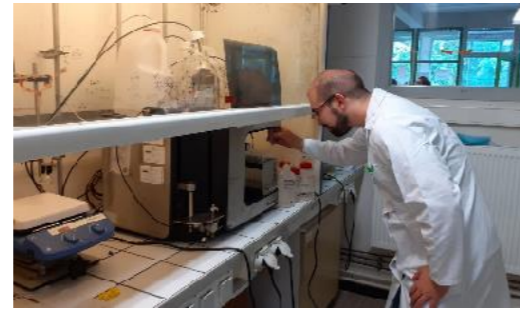


Enzymology



Ingénierie enzymatique | Évolution dirigée | Mutagenèse aléatoire | Optimisation de procédés | Synthèse enzymatique

Glycochemistry



Synthèse de glycosides | Elaboration de glycoconjugués | Dosage de glycomolécules

DZYME's clients:

Glycoconjugation



Modélisation d'interactions | Production de lectines/glycosidases | Mesures d'affinité et d'activité

Applied glycotecnologies



Cosmétique | Santé | Nutraceutique | Feed & food



**Thank you for your
attention**

