



The PSB in 4 dates

- November 2002: The PSB was established by a Memorandum of Understanding by the EMBL, the ESRF, the ILL and the IBS
- January 2006: Inauguration of the joint laboratory building (Carl-Ivar Brändén Building (CIBB)) for the PSB and Institute for Structural Virology, University Joseph Fourier (IVMS).



- January 2007: Merge of the EMBL and the CNRS/University virology unit into UVHCI (Unit of Virus Host Cell Interactions).
- October 2013: The IBS moved onto the EPN campus





The Partnership for Structural Biology - Grenoble 350 scientists within walking distances

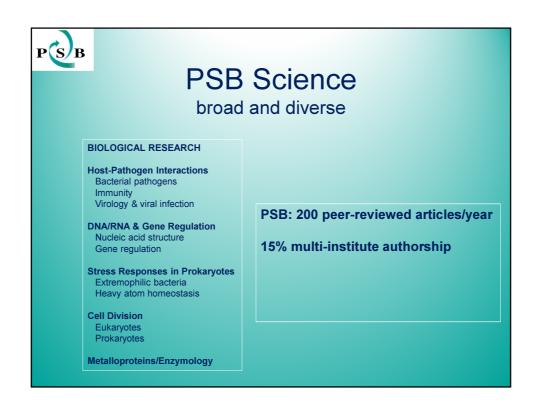




PSB Training activities

- Crystallography tutorials and SANS/SAXS courses for PSB Students
- EMBO Courses co-organised by the partners
- HERCULES
- Erasmus Mondus Program
- Master in Structural Biology (Univ J. Fourier, Grenoble)







PSB Science

broad and diverse

BIOLOGICAL RESEARCH

Host-Pathogen Interactions

Bacterial pathogens Immunity

Virology & viral infection

DNA/RNA & Gene Regulation

Nucleic acid structure Gene regulation

Stress Responses in Prokaryotes

Extremophilic bacteria
Heavy atom homeostasis

Cell Division

Eukaryotes Prokaryotes

Metalloproteins/Enzymology

TECHNOLOGY DEVELOPMENT

Methodologies for Structural Biology

Protein Expression Crystallisation Functional Studies Structural Methods

Instrumentation

Synchrotron Neutron scattering



PSB: a unique palette of **23** technological platforms for integrated structural biology studies

Protein Expression

Cell Free ESPRIT

Eukaryotic Expression Facility

Deuteration Lab Isotopic Labeling Robiomol

Sample Characterization

Surface Plasmon Resonance

Analytical Ultra Centrifugation Biophysics Cell imaging Mass Spectrometry Membrane Protein Purification Platform NMR Quality Control Protein Sequencing

High Resolution Studies

Cryobench FIP Beamline (BM30)

High Field Nuclear Magnetic Resonance HT Crystallisation

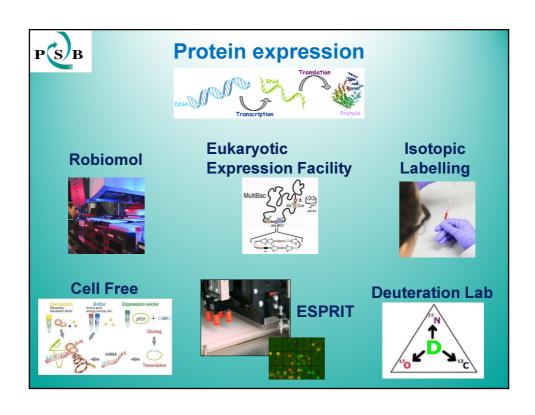
HT Membrane Protein Crystallisation

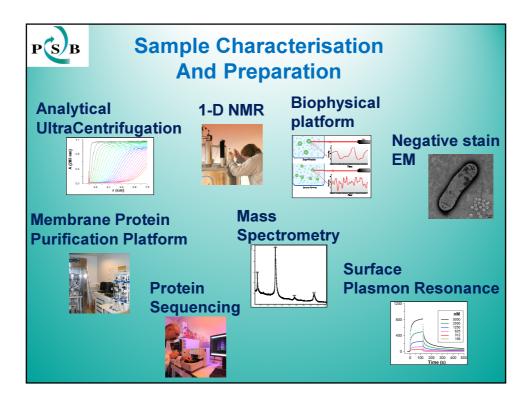
Neutron Diffraction

Structural Biology Beamlines

Supramolecular Structures

Electron microscopy SANS/ SAXS







Crystallisation

High Throughput Crystallisation (HTX)



- 200 Regular Users (Over 681 registered scientists)
- 1900 samples processed/year (~1000000 crystallisation experiments)
- Granted access to 110 Scientists in 15 countries (many of which are users of ESRF or ILL)



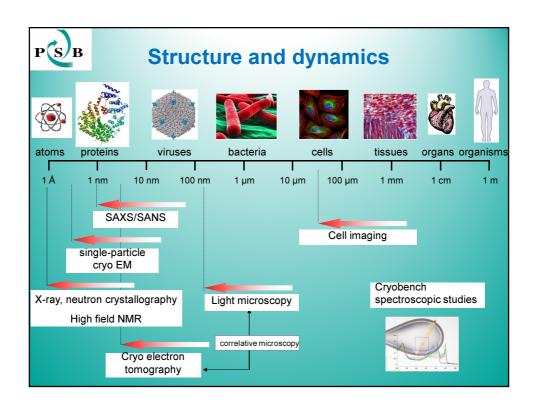
Crystallisation

High Throughput Crystallisation (HTX)



High Throughput Membrane Protein Crystallisation Platform (HTMPC)









High Field NMR Facility



- 6 NMR Spectrometers from 600 to 950 MHz
- State of the art detection probes
- ¹H, ²H, ¹³C, ¹⁵N, ¹⁹F & ³¹P NMR
- Solution & Solid State NMR applications

P(S)B

Neutron diffraction beamlines

LADI-III: quasi-Laue neutron diffractometer



- Data collection at RT or cryogenic temperatures using Cobra cryostream:
- using Cobra cryostream; e.g. cryo-trapping studies of enzymatic reaction intermediates
- Optimized for high- to medium-resolution (1.5 2.5A) studies of large (50 120A on edge) unit-cell systems using perdeuterated crystals (0.05mm3 0.5mm3)

D19: thermal neutron monochromatic diffractometer



- Monochromatic data collection (λmono 0.8 – 2.4Å, e.g. λ = 1.46Å)
- Large (120° x 30°) PSD 'banana' detector
- Optimized for high-resolution (better than 1A) studies of smaller (30 – 70A on edge) unit-cell systems using crystals > 1mm3



How to access PSB platforms?

Peer Review

next deadlines ESRF: 1st March 2015 ILL: 17th February 2015

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- Transnational access (INSTRUCT, BioStruct-X, BIO-NMR,...)
- Collaboration
- Proprietary/paid-for access mode



The Future

• EPN Structural Biology Access Hub web page



The Future

- EPN Structural Biology Access Hub web page
- Integrated access between PSB platforms



The Future

- EPN Structural Biology Access Hub web page
- Integrated access between PSB platforms
 - SAXS/SANS
 - HTX/MX beamlines



