### PAUL SCHERRER INSTITUT

### Life Science Research at Paul Scherrer Institute



### ETH

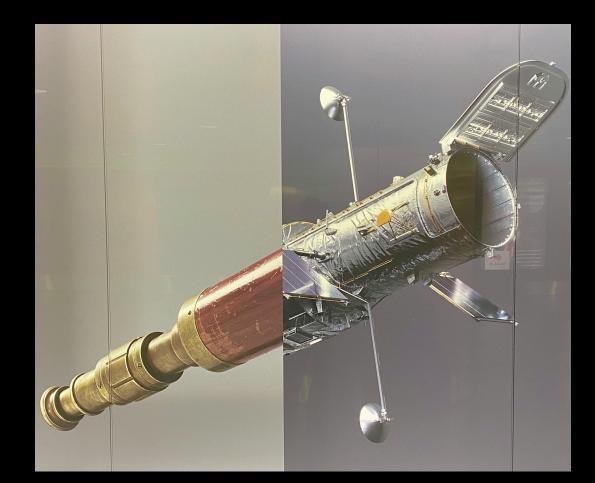
Eidgenössische Technische Hochschule Zürich Swiss Federal Institute of Technology Zurich

#### Prof. Gebhard F. X. Schertler

Head, Division of Biology and Chemistry, PSI Professor for Structural Biology ETH Zürich Laboratory of Biomolecular Research Switzerland

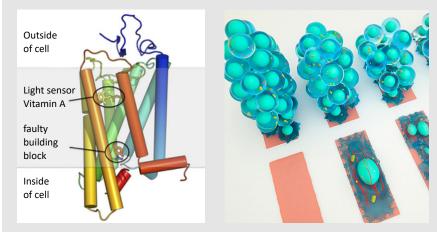


#### **Technology Drives Science and Science Drives Technology**





### Biology and Health at PSI



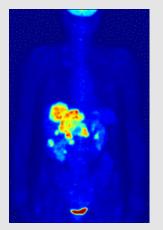
#### **Structure of Proteins**

for the targeted development of new drugs Prof. G. Schertler Prof. M. Steinmetz

 $\Rightarrow$  Accelerators: SwissFEL, SLS

Nanoscale biology of molecular structure and dynamics in the living cell Prof. G. Shivashankar, Prof. J.P. Abrahams

 $\Rightarrow$  Detectors: EM



#### **Radio Pharmaceuticals**

for the diagnosis and therapy of tumours Prof. R. Schibli, Dr. R. Eichler

**Proton Therapy** for destruction of tumours and protection of healthy tissue Prof. D. Weber

 $\Rightarrow$  Accelerators: HIPA, SINQ  $\Rightarrow$  Isotopes, Radiochemistry, CRS



 $\Rightarrow$  Accelerators: COMET

### The Complexity of Biology

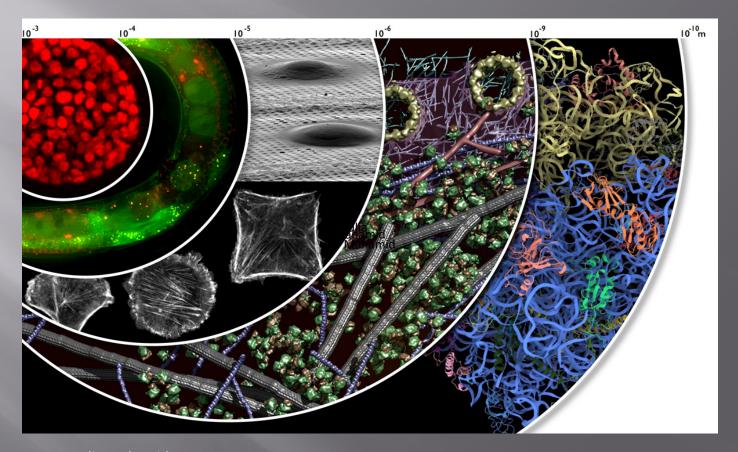
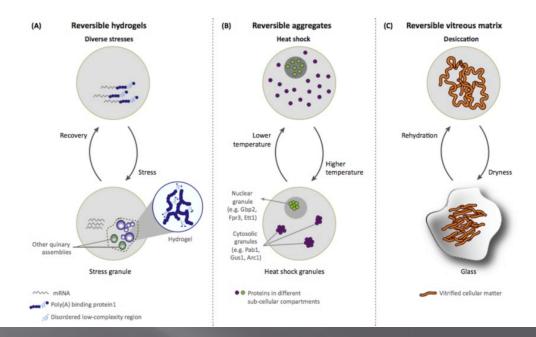


Image: Julia Mahamid EMBL https://www.embl.org/groups/mahamid/wp-content/uploads/2021/05/mahamid-group-fig1.jpg

### Intrinsically Disordered Proteins Adaptively Reorganize Cellular Matter During Stress

Intrinsically disordered proteins (IDPs) can protect cells from diverse stresses by forming higher order assemblies such as reversible aggregates or granules. Recently, Boothby et al. show that IDPs protect tardigrades against desiccation by forming a glass-like amorphous matrix, highlighting that material properties of disordered proteins can confer adaptation during stress.

The paper by Chavali et al can be found here.







### Large-volume, high-resolution tomography

**22-nm & 100-nm Intel processor** Holler *et al*.

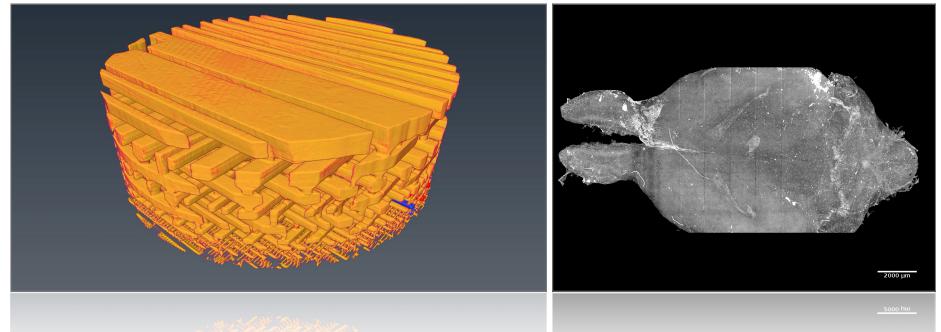


**3D mouse brain** Miettinen *et al*.





Human Brain Project





### Correlative multi-scale analysis in biology and medicine

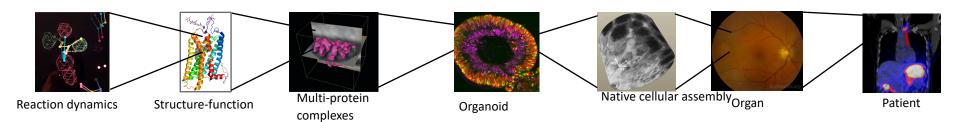
Integrating atomic scale and low-resolution structural and dynamic information in physiological context at cellular and multicellular levels to understand living systems

- Expertise in photon- and electron-based imaging methods across multiple length scales, as well as *in silico* modelling of biological and biochemical processes, is highly relevant for basic and applied biomedical research.
- These approaches help the development of non-invasive diagnostics and therapies for cancer, neurodegenerative states and infectious diseases.
   Collaboration is fostered with Swiss hospitals, biotech, big pharma and with institutes involved in biomedical research.

D-ITET; D-HEST D-BIOL; D-BSSE



Materials Science and Technology Empa Platform for Image Analysis; Biomedical Imaging Technologies

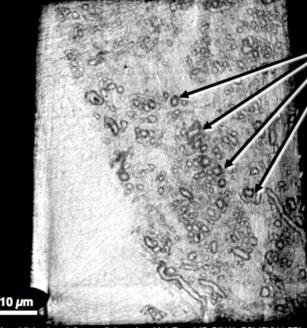




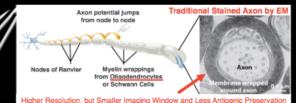
### OMNY ptychography of brain tissue

OMNY (tOMography Nano crYo) Technology Developed at PSI Delivers Unmet Need for Efficient Nanoscale Imaging of Large Windows of Unstained Cryo Tissues

Unstained cryo-preserved mouse brain Reconstructed OMNY Tomogram – Movie



Example of Identifiable Structures: "Highways of Communication" Myelinated Axons in Brain



#### "Non-Destructive Approach"

No cutting, no milling; unique to others. Imaging does not destroy specimen.

#### "Precise Navigation System"

Localise elusive cellular hallmarks for correlative imaging to reveal higher resolution **ultrastructural details** and **molecular identities** (as for disease).

Unpublished work from collaboration: Uni. Basel (C-CINA), PSI (SYN-LSB), Roche Innovation Center Basel (MDCB, NORD DTA)

#### Human brain tissue

#### healthy

Parkinson-diseased

Myelinated axons Swellings along axons Cell nuclei Neuromelanincontaining organelles Blood vessels Blood cells

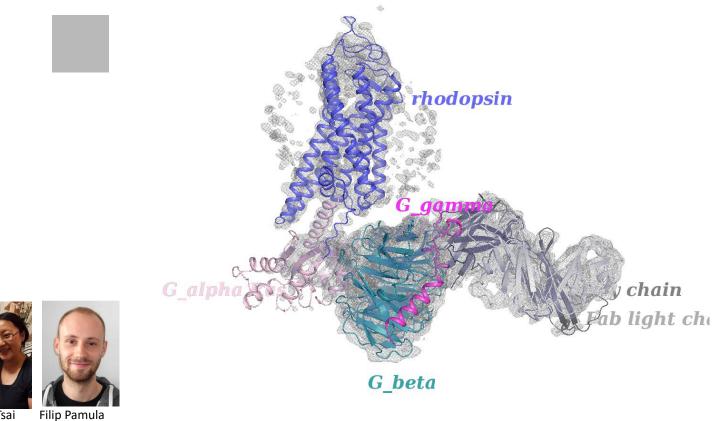
> H.Tri Tran et al., Front. in Neuroscience

5 specimens imaged from a healthy individual as control

4 specimens imaged from a Parkinson-affected individual



Cry0-EM Rhodopsin Gi Protein Complex isolated from a living tissue



ChingJu Tsai Filip Pa



rhodopsin

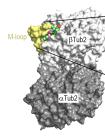
mini-Go

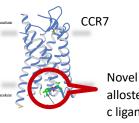
#### Integrative Structural Biology with pharmacological relevance

*Exploring membrane proteins and the microtubule* system using X-rays and cryo electron microscopy

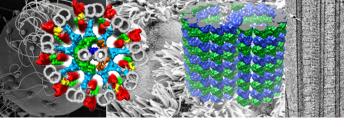
- → All methods in structural biology are critical to remain highly competitive.
  → Both research fields are highly relevant for drug discovery enabling:
  - Collaborations with pharma, biotech and spin-offs
  - In-house structure-based drug discovery projects
  - Generation of *spin-off* companies

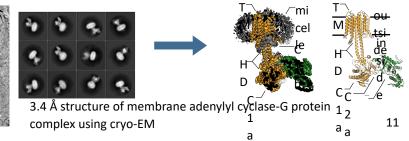
Tubulin









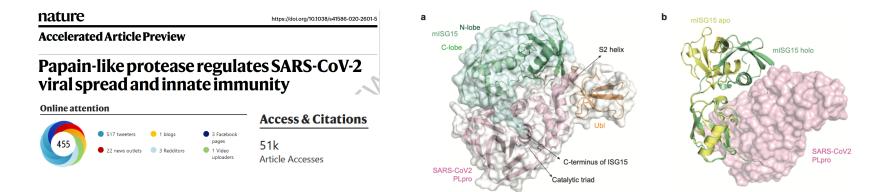




## Protein structures of COVID-19 virus from SLS

The structural biology work was performed at the macromolecular crystallography beamline **X06SA-PXI at the SLS** during the "**PRIORITY COVID-19 Call**". The crystallographic **data were collected on 9 April 2020**.

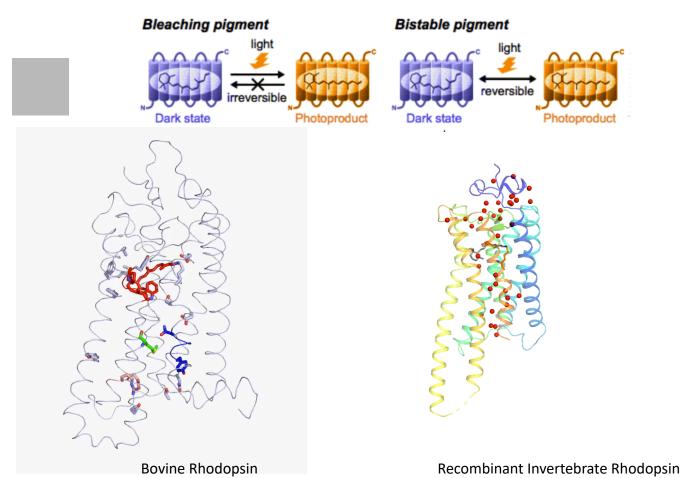
The planned **Easter shutdown of the SLS was cancelled for this specific experiment**. The paper was submitted within one month after answering the proposal call and **published in Nature on 29 July 2020**.



Shin, D., Mukherjee, R., Grewe, D. *et al.* Papain-like protease regulates SARS-CoV-2 viral spread and innate immunity. Nature (2020). <u>https://doi.org/10.1038/s41586-020-2601-5</u> Structural analysis of SARS-CoV-2 PLpro in complex with full length ISG15



### **Visual Pigments**





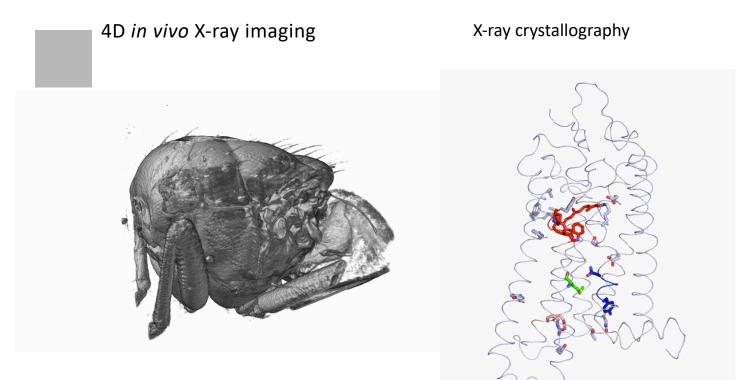
Eshita Mutt Elena Leska Niranjan Varma Imaging methods are dominating all of life science.
 Atomic structure drives mechanistic understanding and drug discovery.

- Large multicomponent molecular «machines» drive many cellular processes
- Intrinsically disordered protein domains are at the centre of nano and cellular organisation Optical readouts have been developed for signaling dynamics
- Computational methods for networks describe non linear responses in regulated processes
- Life sciences are inherently interdisciplinary and depend on novel method development



Dynamics in Molecular Biology

Visualizing the motion of an object helps to understand its function



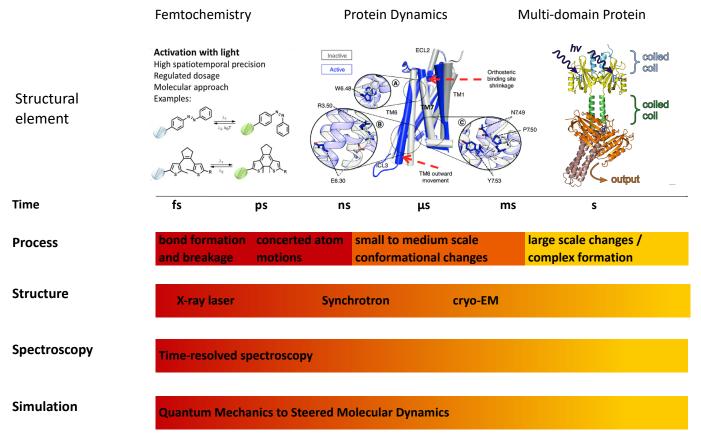
Mokso et al., Sci. Rep., 2015

Standfuss et al., Nature, 2011



**Time-resolved Structural Biology** 

Driving Structural biology from molecular snapshots towards molecular movies





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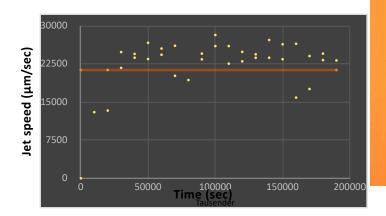
### Jetting tests with high speed camera

A smooth injection improves data quantity and quality



Dan James

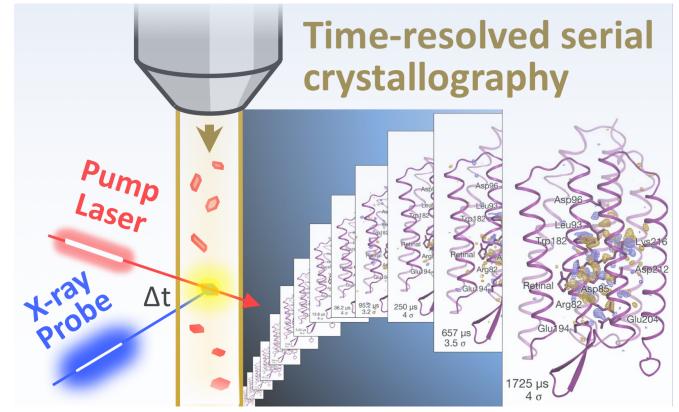
- Video of the jet is taken at 1000-2000 fps
- The video is analysed by measuring the distance a crystal (or other feature) travels over a given number of frames







Bringing time-resolved measurements to the molecular scale : Swiss FEL



Standfuss, Curr. Opin. Struc. Biol., 2019

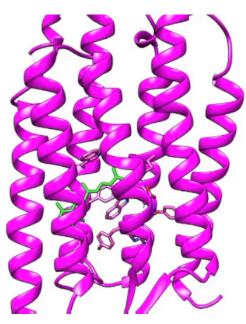


#### Time resolved structural analysis of bacteriorhodopsin: Structural basis of photocatalysis

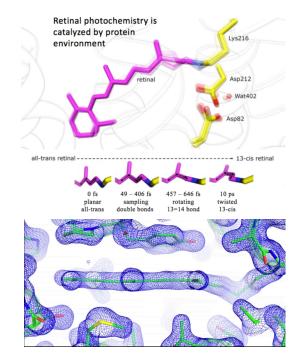


Joerg Standfuss





The chemical basis of optogenetics

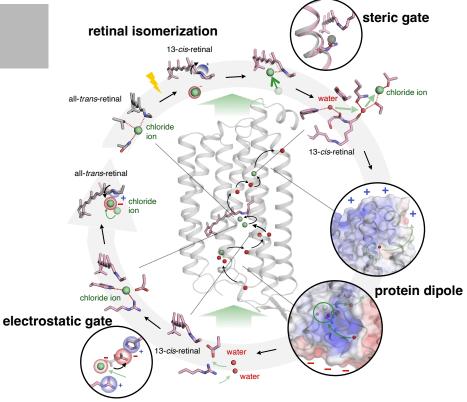


Nogly et al. Science 2018; DOI: 10.1126/science.aat0094



## Movie of Chloride Pump through Cell Membrane

#### **ETH** zürich



S. Mous et al., Science 10.1126/science.abj6663 (2022).

Molecular analysis of **full activation cycle of lightdriven chloride pump mechanism** using time resolved, serial X-ray crystallography and multiscale simulation.

- **Time-resolved** detection of **transient anion binding** across the cavity (time range: 10 ps to 55 ms).
- Light-induced isomerization of retinal provides energy for chloride transport.
- Driving force created by protein dipole moment.
- Unidirectional transport ensured by steric and electrostatic molecular gates.





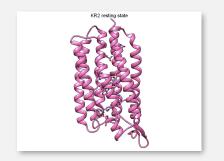


Sandra Mous, ETHZ

G. Gotthard, PSI P Nogly, ETHZ

**Examples of biological experiments at SwissFEL** 

#### Rhodopsins pave the way into a dynamic future for structural biology



Skopintsev et al., 2020, Nature

#### Sodium pumping rhodopsin

- > Ten molecular snapshots of sodium transport out of the cell
- > Next-generation **optogenetic** tool

#### Chloride pumping rhodopsin

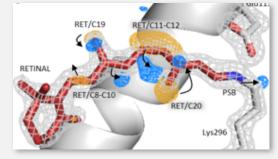
Mous et al., 2022 Science

dark state

- > SwissFEL and SLS resolves **chloride** > Molecular snapshots of **early** transport into the cell
- > Electrostatic gates **explain** transport

#### **Visual GPCR rhodopsin**

- events in vision
- > **GPCR** activation



Gruhl et al., 2022, Nature, accepted



### Potential applications in pharmacology

Dynamic atomic information is necessary for understanding pharmacology

This includes fundamental processes like induced-fit, conformational selection or lock-and-key mechanisms of ligand binding

Understanding catalytic effect of proteins requires measuring the changes to the energetic landscapes introduced by the bound substrate or ligand

Quantification of the dynamic mixture of different conformations of Receptors, Channels and Transporters in solution can explain drug action

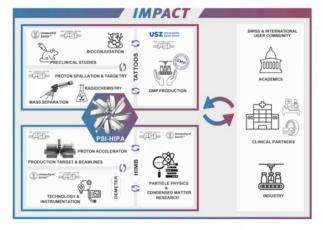
Using the right kind of experiments all these questions can be addressed



Das Ziel von IMPACT (engl.: Isotope and Muon Production using Advanced Cyclotron and Target technologies) SBFI-Infrastrukturantrages (Periode 2025-2028) ist der Aufbau von zwei neuen Target-Stationen sowie Strahllinien an der Protonenbeschleunigeranlage HIPA (Eng.: High Intensity Proton Accelerator) des Paul Scherrer Instituts. Dies wird zu einer Erhöhung der Myonen-Intensität um einen Faktor 100 führen (HIMB) und neue Radionuklide für die fortgeschrittene Krebsbehandlung verfügbar machen (TATTOOS).





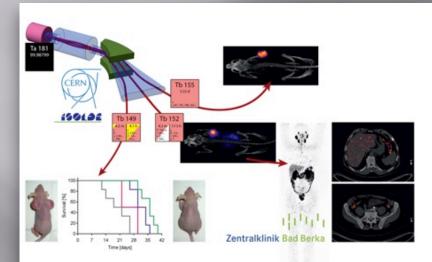


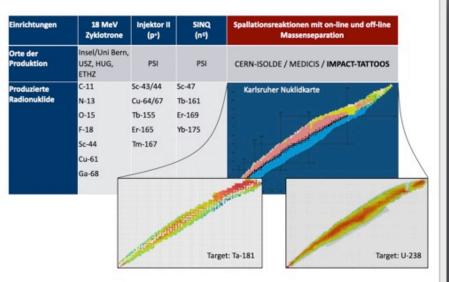
IMPACT@HIPA Die geplante Erweiterung der bestehenden Protonenbeschleunigerinfrastruktur zu Gunsten von HIMB (in grün; engl.: High-Intensity Muon Beams) und TATOOS (in blau; engl.: Targeted Alpha Tumor Therapy and Other Oncological Solutions). Strategie IMPACT Erklärtes Ziel ist eine substanzielle Erweiterung der bestehenden Protonenbeschleunigerinfrastruktur am PSI, die es erlaubt eine weltweit führende Anlage weiter auszubauen. Dies eröffnet eine Vielzahl einzigartiger Möglichkeiten für die weitere Erhöhung der Attraktivität des Forschungsstandortes Schweiz.



#### IMPACT: TATTOOS







Die "Terbium-Schwestern" Die Terbiumisotope Tb-149 (Therapie), Tb-152 (Diagnose) und Tb-155 (Diagnose) wurden in Zusammenarbeit mit ISOLDE (CERN) am PSI auf deren radiopharmazeutische Anwendbarkeit erforscht. Diese Vorstudien waren äusserst vielversprechend und sind einer der entscheidenden Treiber hinter TATTOOS. Produktion von Radionukliden für die Radiopharmazie Verschiedene Produktionsmöglichkeiten von Radionukliden für die Radiopharmazie (Bildgebung und Therapie) in der Schweiz. Mit TATTOOS wird die Anzahl verfügbarer Radionuklide (inkl. therapeutischen Dosen) schlagartig um ein Vielfaches vergrössert.



Park Innovaare – Ready end of 2023 (innovation platform for startups and industry)

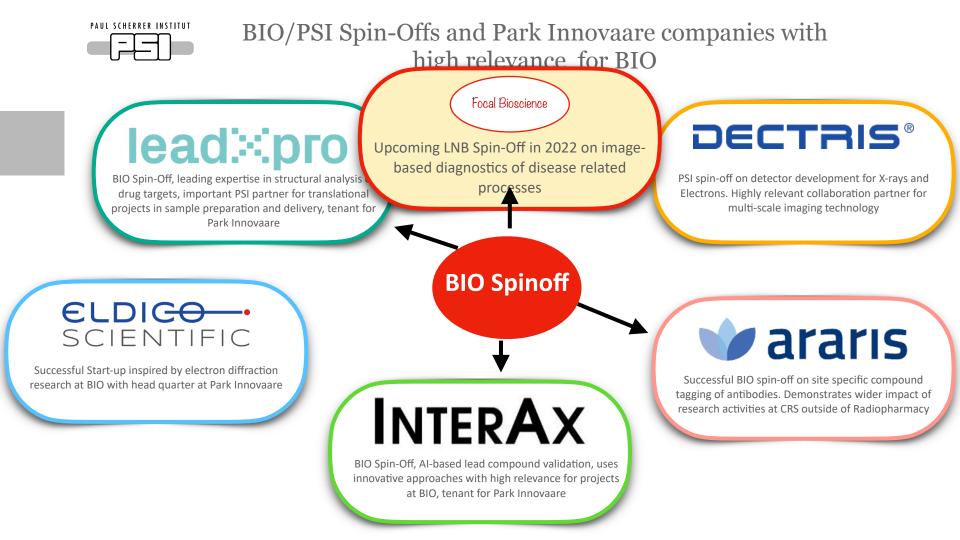




Debiopharm

*`eaTecr* 

- large-scale research facilities (SLS, SwissFEL, SINQ, SμS)
- precision workshop
- vibration-free laboratories
- clean room for nanofabrication



### Life sciences are

- > Changing rapidly
- Demand a very different user environment than large scale physics
- Are inherently interdisciplinary and driven by method development
- Need large facilities to catalyse these developments better

**Biology & Life Sciences are major drivers for large high** energy facilities in Europe

# PAUL SCHERRER INSTITUT

PSI offers a world-wide unique combination of large-scale facilities: X-ray sources SLS and SwissFEL, neutron source SINQ and muon source SµS and technology platforms for a broad range of research and development in science and medicine, education and collaboration with industry. Infrastructure for science and innovation.

Own science and technology program and combination with world-leading schools EPFL and ETH and three other Swiss national labs expands **know-how** and **talent pool**. **Park Innovaare** of Switzerland Innovation is a new innovation park connected directly to the institute, setting the stage for much larger **economic impact**.



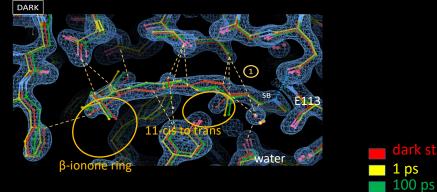
### Wir schaffen Wissen – heute für morgen

Thank you for your attention

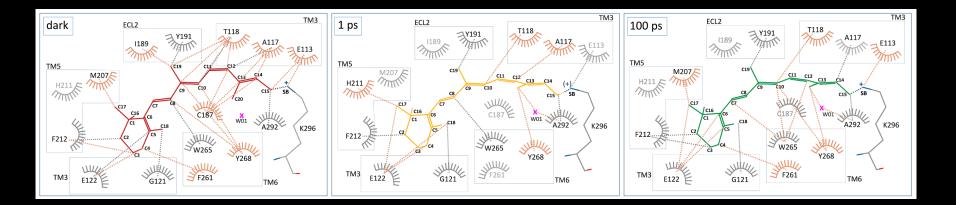


#### Ultrafast changes in the rhodopsin binding pocket.

weakening of retinal interactions in the ligand binding pocket :

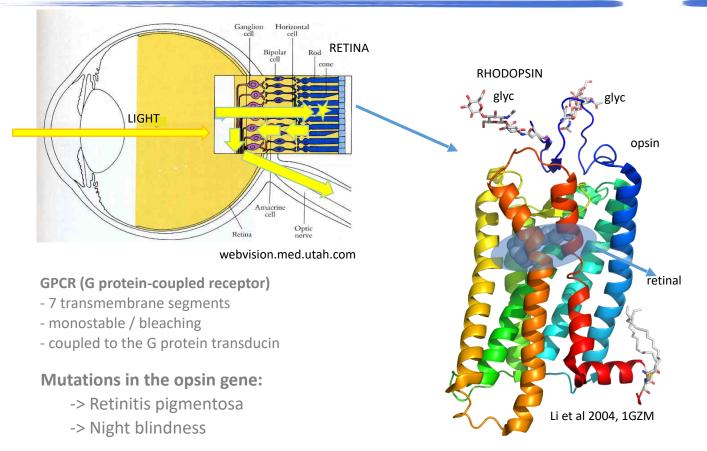


Environment distances < 3.7 Å





### Mammalian rhodopsin and the first step of vision





#### PICOSECOND DYNAMICS OF THE FIRST STEP OF VISION : Rhodopsin TR-SFX



