

WIR SCHAFFEN WISSEN – HEUTE FÜR MORGEN

Oliver Bunk :: Laboratory for Macromolecules and Bioimaging (LSB) :: Paul Scherrer Institute

PX I, II, III, S-/I-TOMCAT, cSAXS @ SLS 2.0

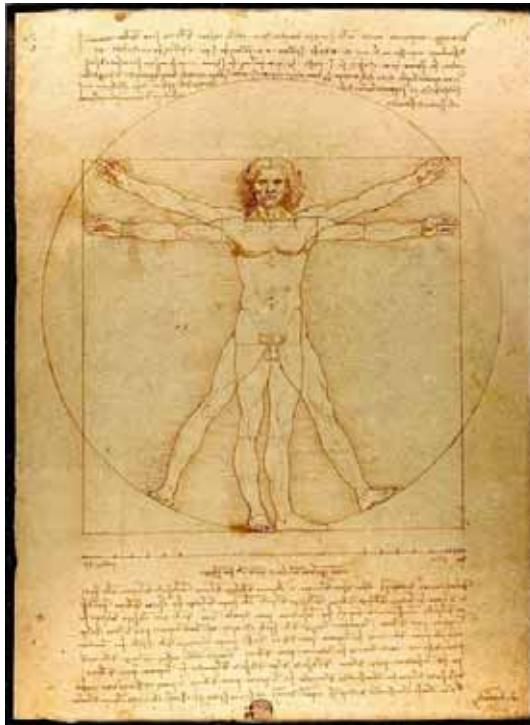
Seeing is believing.

Crystallography and imaging in life and materials science at
synchrotron radiation facilities

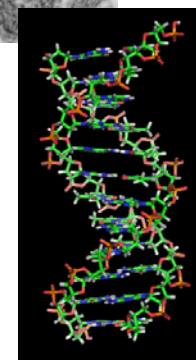
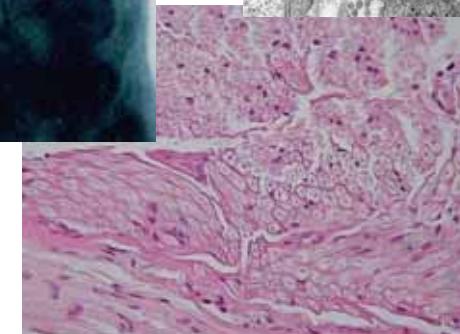
SLS 2.0 upgrade workshop, Feb. 28th 2022



Hierarchically structured materials



1 m



1 Å

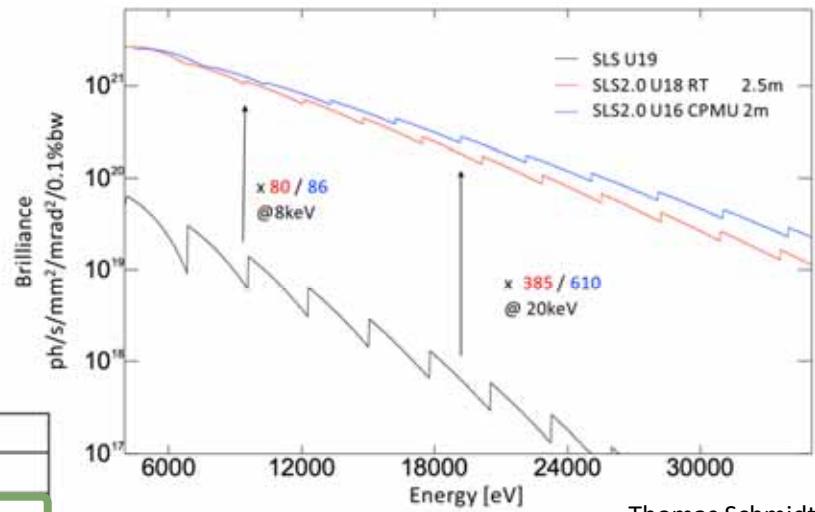
(from Wikipedia)

SLS 2.0 is highly benefiting all hard X-ray techniques

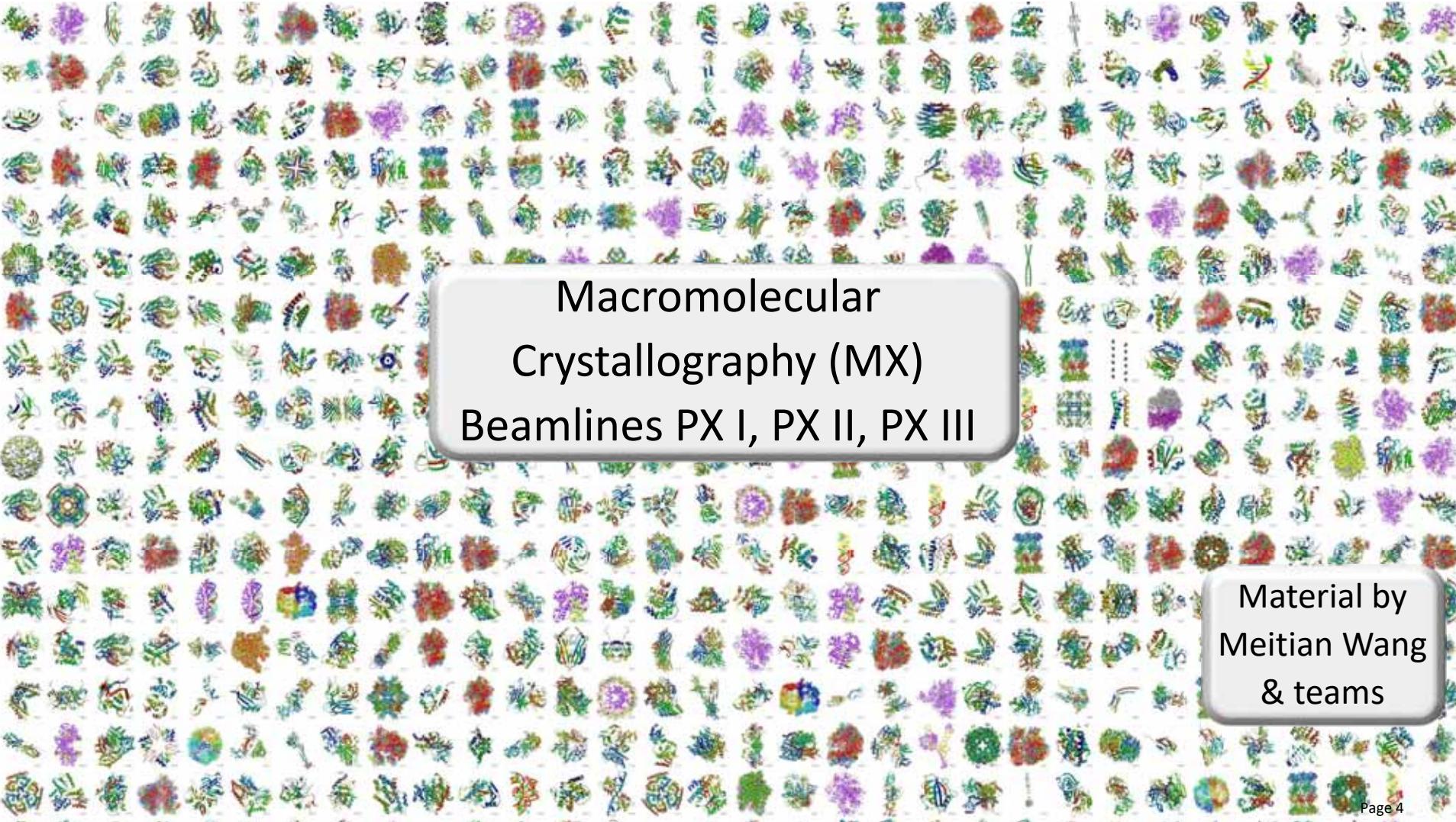


	SLS today	SLS 2.0
Circumference [m]	288.007'289	288.000'000 → 288.000'222
Energy [GeV]	2.411	2.700
Working point Q_x / Q_y	20.43 / 8.74	39.37 / 15.22
Natural chromaticity ξ_x / ξ_y	-67.3 / -21.0	-99.0 / -33.4
Emittance [pm.rad]	5630	149 → 157 → 136
Energy spread [10^{-3}]	0.878	1.103 → 1.147 → 1.038
Radiation loss per turn [keV/turn]	549	666 → 689 → 892
Momentum compaction factor [10^{-4}]	+6.04	+1.06 → +1.04 → +1.04
Horizontal damping partition J_x	1.00	1.85 → 1.82 → 1.64
Damping times [ms]		4.2/7.8/6.8 → 4.1/7.5/6.4 → 3.6/5.8/4.3

Andreas Streun, SLS 2.0 Baseline Lattice, SLS2-SA81-004-14 (1.9.2020)



Thomas Schmidt,
SLS 2.0 beamline
CDR advisory
committee

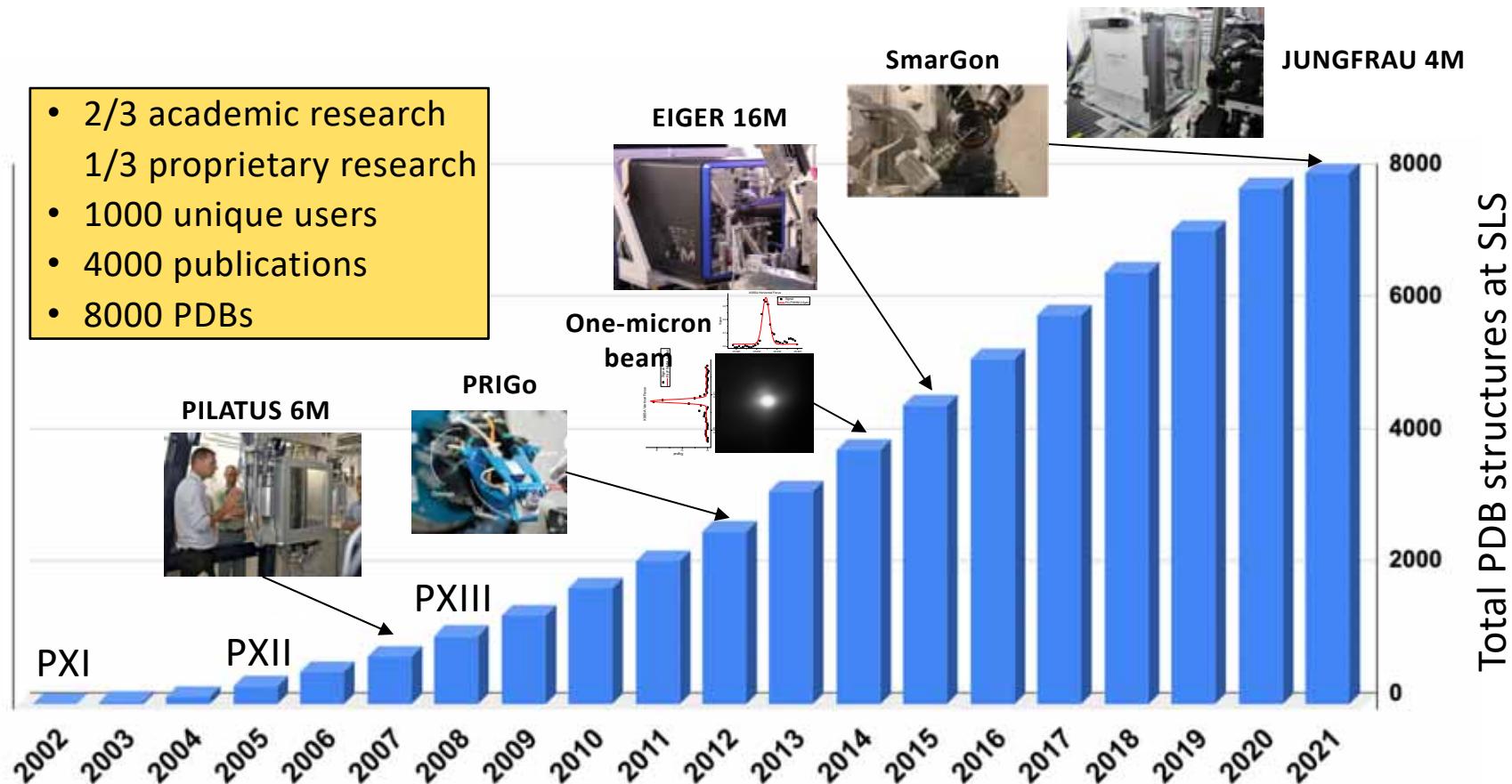


Macromolecular
Crystallography (MX)
Beamlines PX I, PX II, PX III

Material by
Meitian Wang
& teams

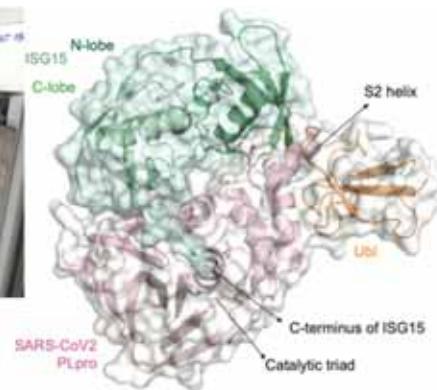
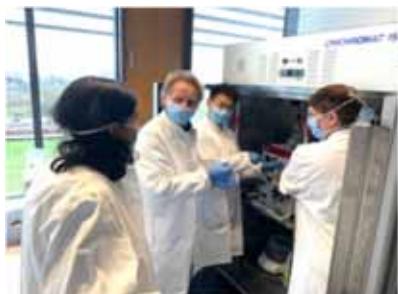
SLS MX Beamlines: 20 years

- 2/3 academic research
1/3 proprietary research
- 1000 unique users
- 4000 publications
- 8000 PDBs

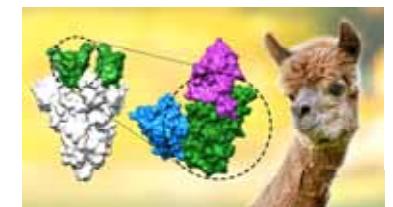


COVID-19, and COVID-19 related research

- Continuous user operation through the pandemic
- 100% remote mode since 2020
- > 50'000 sample mounts in 2021
- SLS Special call for COVID-19 research

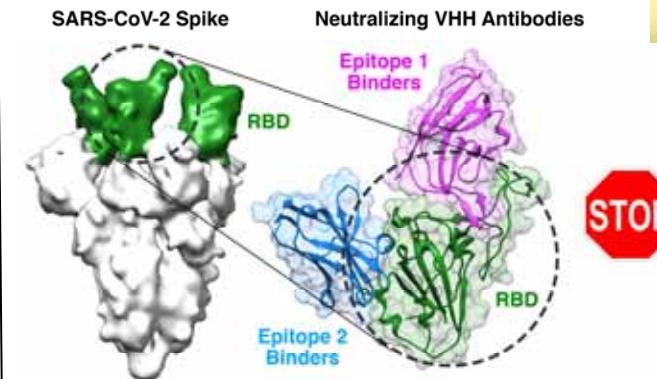


Papain-like protease regulates SARS-CoV-2 viral spread and innate immunity.
Ivan Dikic / University Frankfurt, *Nature* (2020)
Four months from X-ray data collection to publication



"wild-type" SARS-CoV-2

 Alpha (N501Y)
 Beta (K417N, E484K, N501Y)
 Gamma (K417T, E484K, N501Y)
 Delta (L452R, T478K)
 Epsilon (L452R)
 Zeta (E484K)
 Eta (E484K)
 Theta (E484K, N501Y)
 Iota (E484K)
 Kappa (L452R, E484Q)



Nanobodies (blue and magenta) bind to the receptor-binding domain (green) of the coronavirus spike protein (grey), thus preventing infection with SARS-CoV-2 and its variants.
Thomas Gütter / MPI for Biophysical Chemistry, *EMBO J* (2021)



Positioning MX on the international landscape



PX I beamline at the SLS (PSI).

Rules of thumb:

- Ribosome Nobel Prize would be based on cryo EM data nowadays.
- The phase problem is solved.
- **MX excels at high spatial and time resolution in combination with high throughput.**

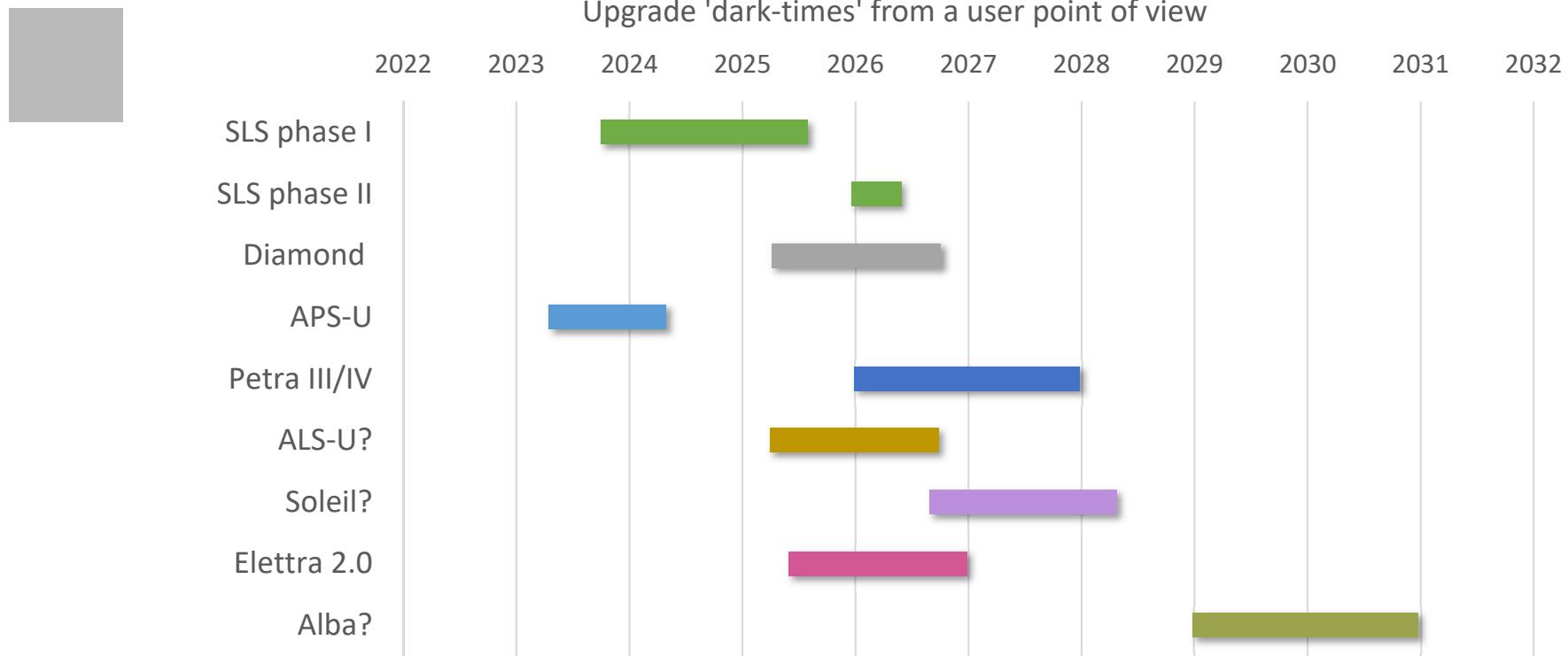
AlphaFold
Protein Structure Database

Developed by DeepMind and EMBL-EBI

<https://www.epfl.ch/labs/lbem/instruments/>

<https://alphafold.ebi.ac.uk/>

Dark-times from a user point of view



MX Team @ SLS



Meitian
Wang

MX application group

<u>Katherine McAuley</u>	Takashi Tomizaki	
Florian Dworkowski	Anuschka Pauluhn	
Sylvain Aumonier	Vincent Olieric	John Beale

MX instrumentation group

<u>Wayne Glettig</u>	Dominik Buntschu
Nathalie Meier	Roman Schneider
Sonia Reber	Tomislav Marijolovic

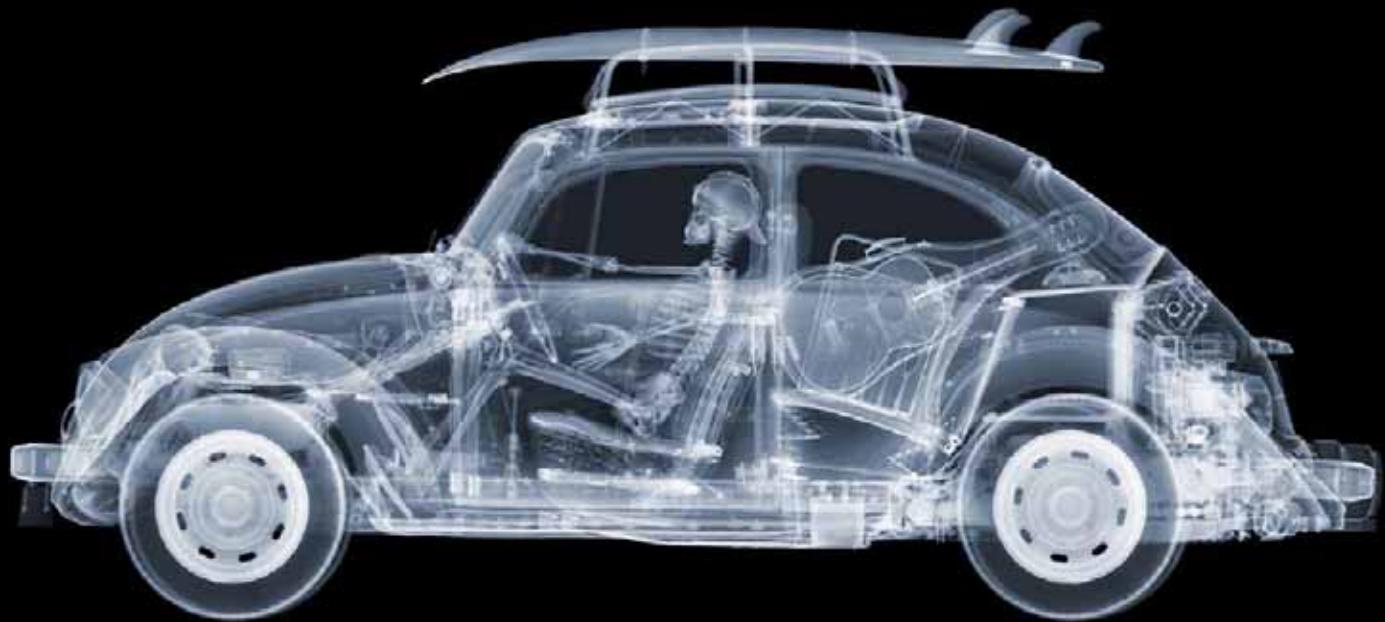
MX data group

<u>Justyna Wojdyla</u>	Kate Smith
Ezequiel Panepucci	Filip Leonarski
Greta Assmann	

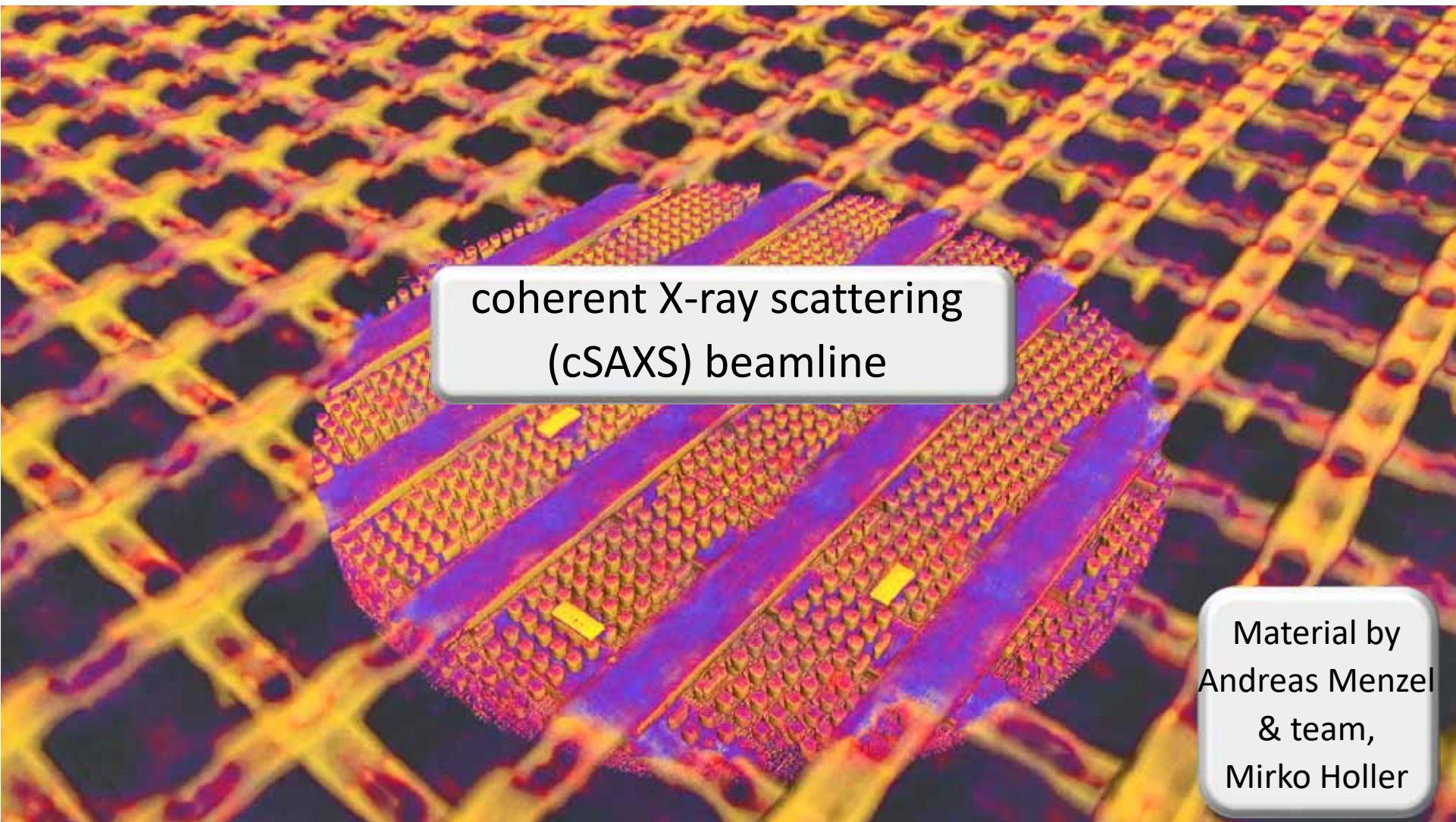
MX sample group

<u>May Sharpe</u>	Eric Plichta
Chia-Ying Huang	Deniz Eris

X-ray imaging
Beamlines cSAXS,
S-TOMCAT, I-TOMCAT



Nick Veasey, 1971 California Bug (May 2015)



coherent X-ray scattering
(cSAXS) beamline

Material by
Andreas Menzel
& team,
Mirko Holler

Main Activities at cSAXS

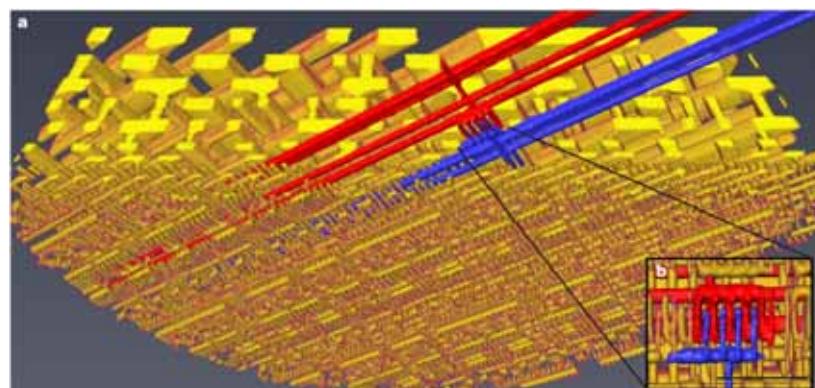
High-Resolution Imaging Ptychography

LETTER

doi:10.1038/nature21409

High-resolution non-destructive three-dimensional imaging of integrated circuits

Mirko Holler¹, Manuel Guizar-Sicairos², Esther H. R. Tsai¹, Roberto Dinapoli¹, Elisabeth Müller¹, Oliver Bunk¹, Jörg Raabe² & Gabriel Aepli^{1,2,3}



Holler et al., Nature 543, 402 (2017)

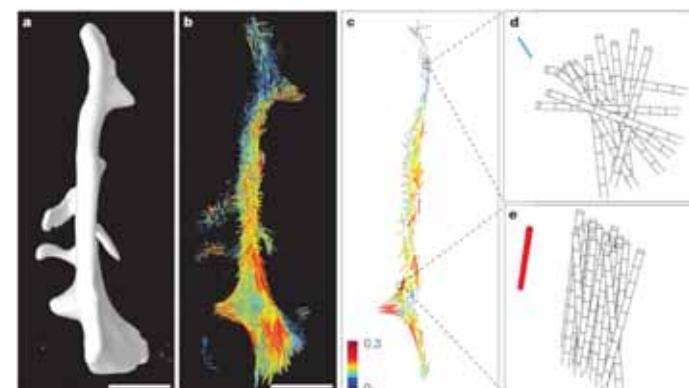
SAXS Imaging, Tomography, and other High-Brilliance Applications

LETTER

doi:10.1038/nature18056

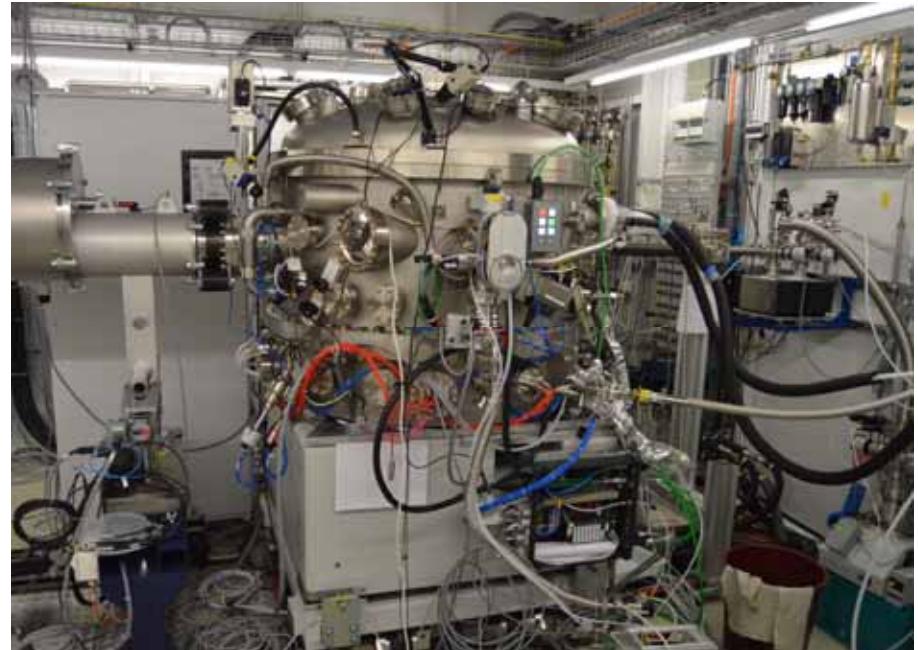
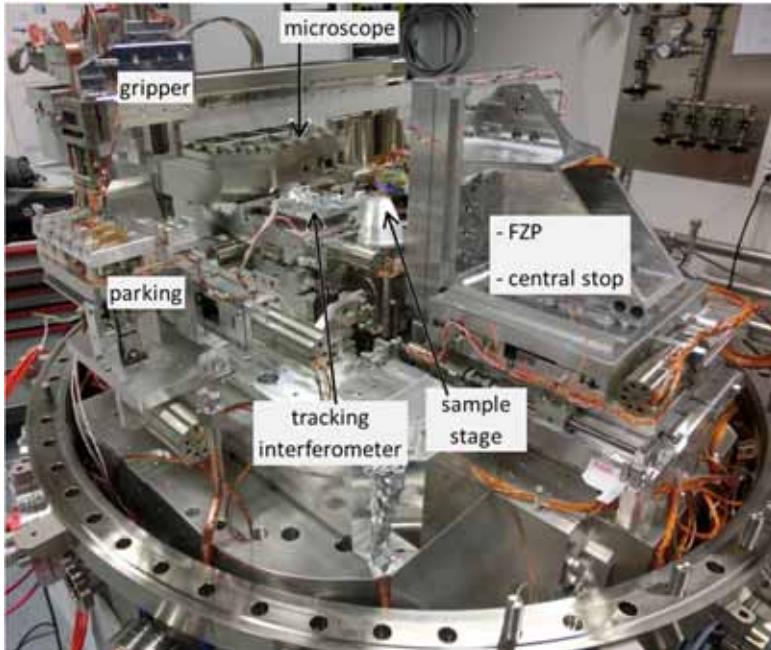
Nanostructure surveys of macroscopic specimens by small-angle scattering tensor tomography

Marianne Liebi¹, Marius Georgiadis², Andreas Menzel², Philipp Schneider³, Joachim Kohlbecker², Oliver Bunk¹ & Manuel Guizar-Sicairos²



Liebi et al., Nature 527, 349 (2015)

Unique instrumentation for ptychographic tomography under cryo conditions



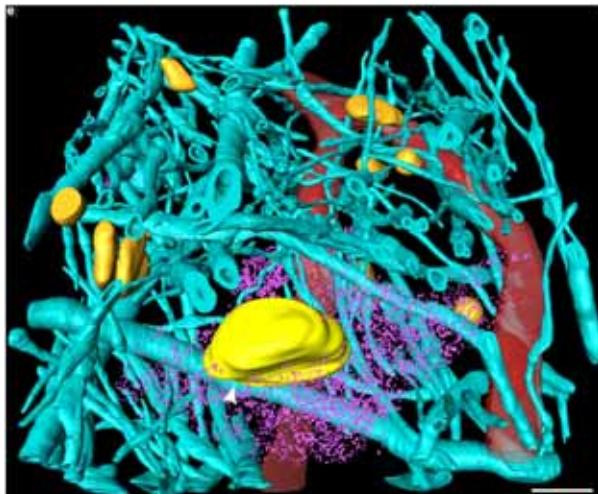
OMNY: Ptychographic nano-tomography for samples under cryo conditions.

Mirko Holler et al., Rev. Sci. Instrum. 83, 073703 (2012). DOI: 10.1063/1.4737624

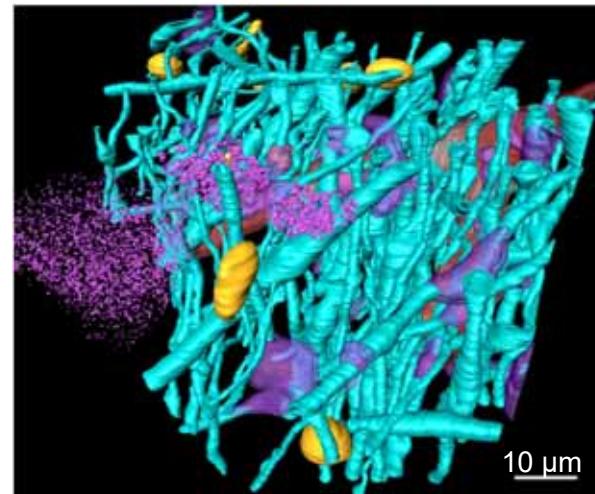
Mirko Holler and Jörg Raabe, Opt. Eng. 54, 054101 (2015). DOI: 10.1117/1.OE.54.5.054101

Mirko Holler et al., Rev. Sci. Instrum. 89, 043706 (2018). DOI: 10.1063/1.5020247

Human brain tissue



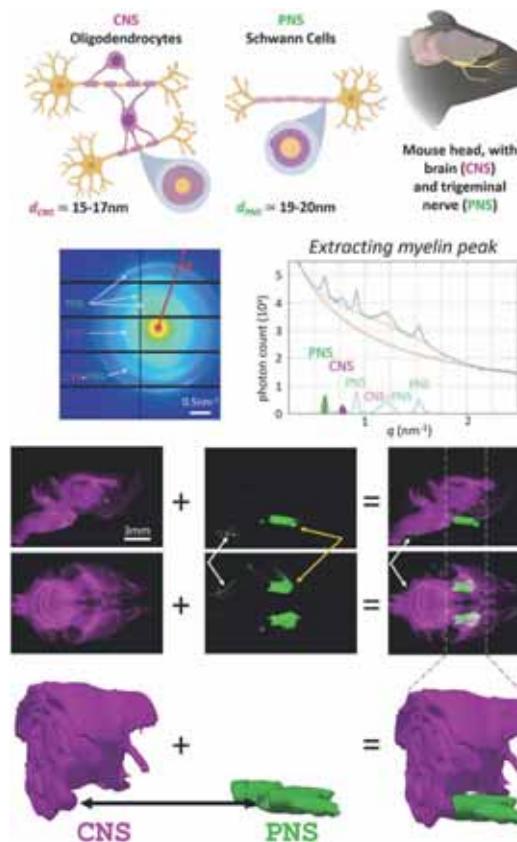
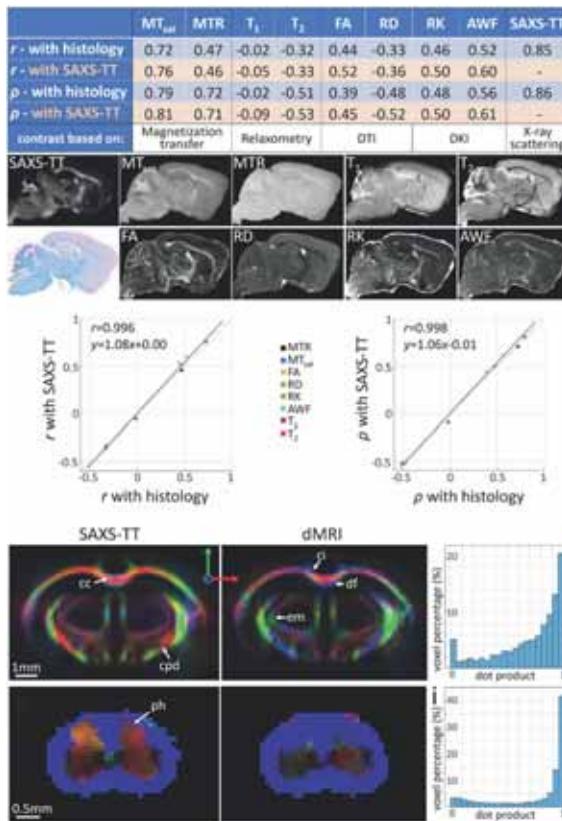
5 specimens imaged from a healthy individual as control



4 specimens imaged from a Parkinson-affected individual

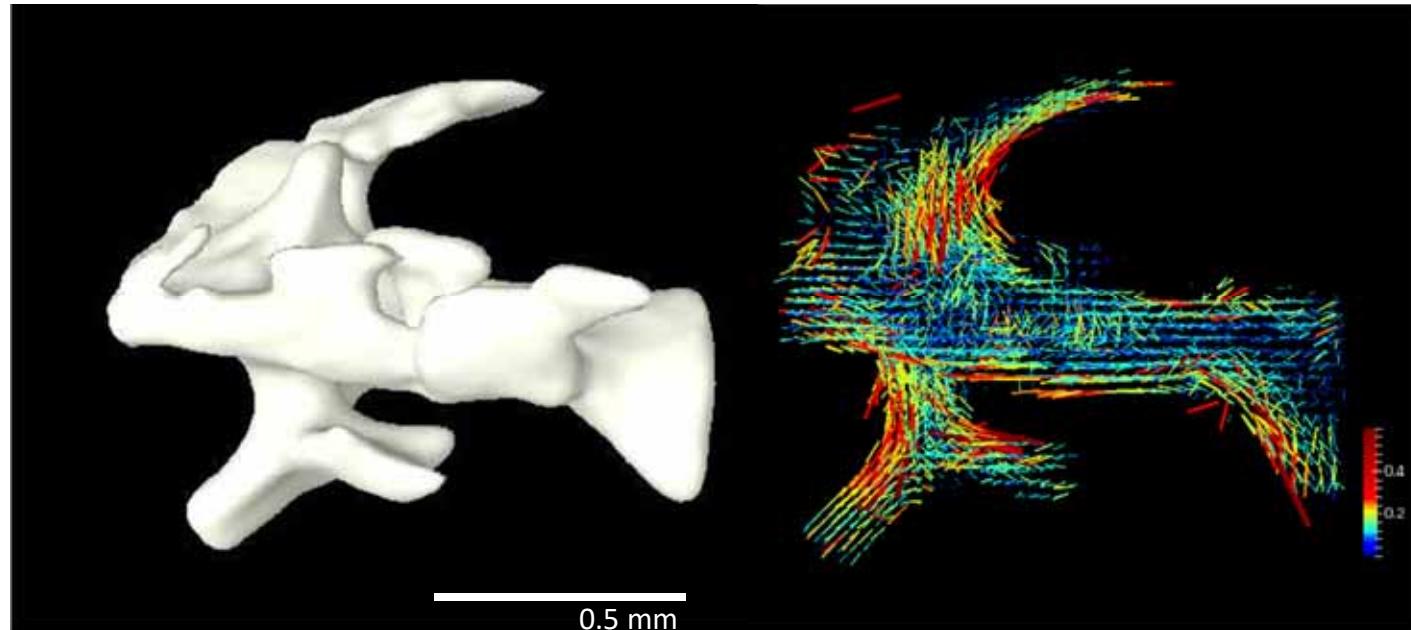
Myelinated axons
Swellings along axons
Cell nuclei
Neuromelanin-containing organelles
Blood vessels
Blood cells

SAS TT complementing established techniques for instance, in neuroscience



Marios Georgiadis *et al.*, Nature Comm. **12**, 2941 (2021)

Scanning SAXS in 3D (SAS TT): Trabecular bone from human vertebra



Marianne Liebi *et al.*, *Nature* **527**, 349 (2015).

Marianne Liebi *et al.*, *Acta Cryst. A* **74**, 12 (2018).

The Coherent X-Ray Scattering Group



Ana
Diaz



Andreas
Menzel



Christian
Appel



Irene
Rodriguez-Fernandez



Johannes
Ihli



Manuel
Guizar-Sicairos



Nicholas Williams
Phillips



Xavier
Donath



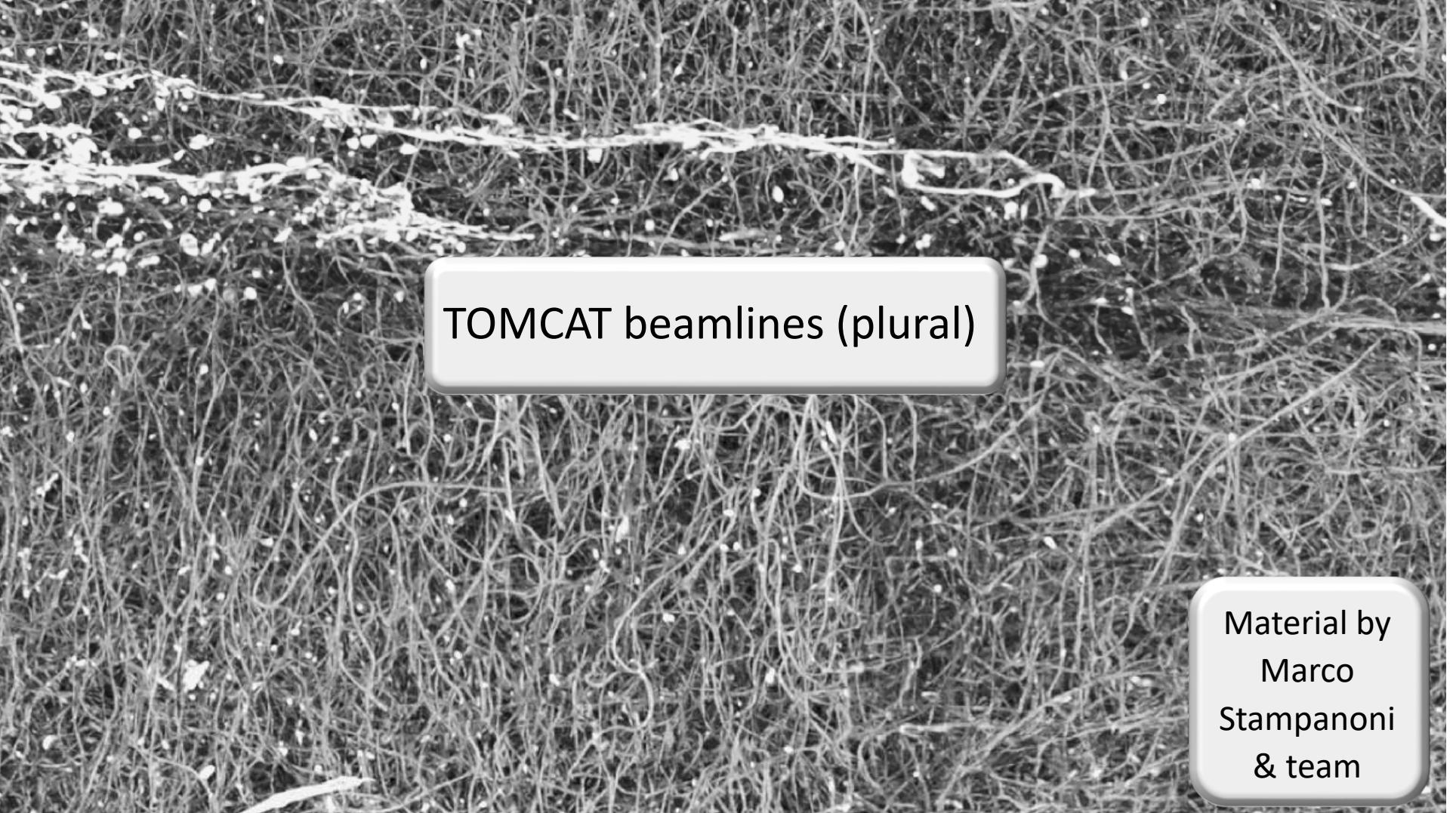
Zirui
Gao

Mirko Holler
Tomas Aidukas

Benedikt Rösner

Bernd Schmitt
Lars Erik Fröjd
...

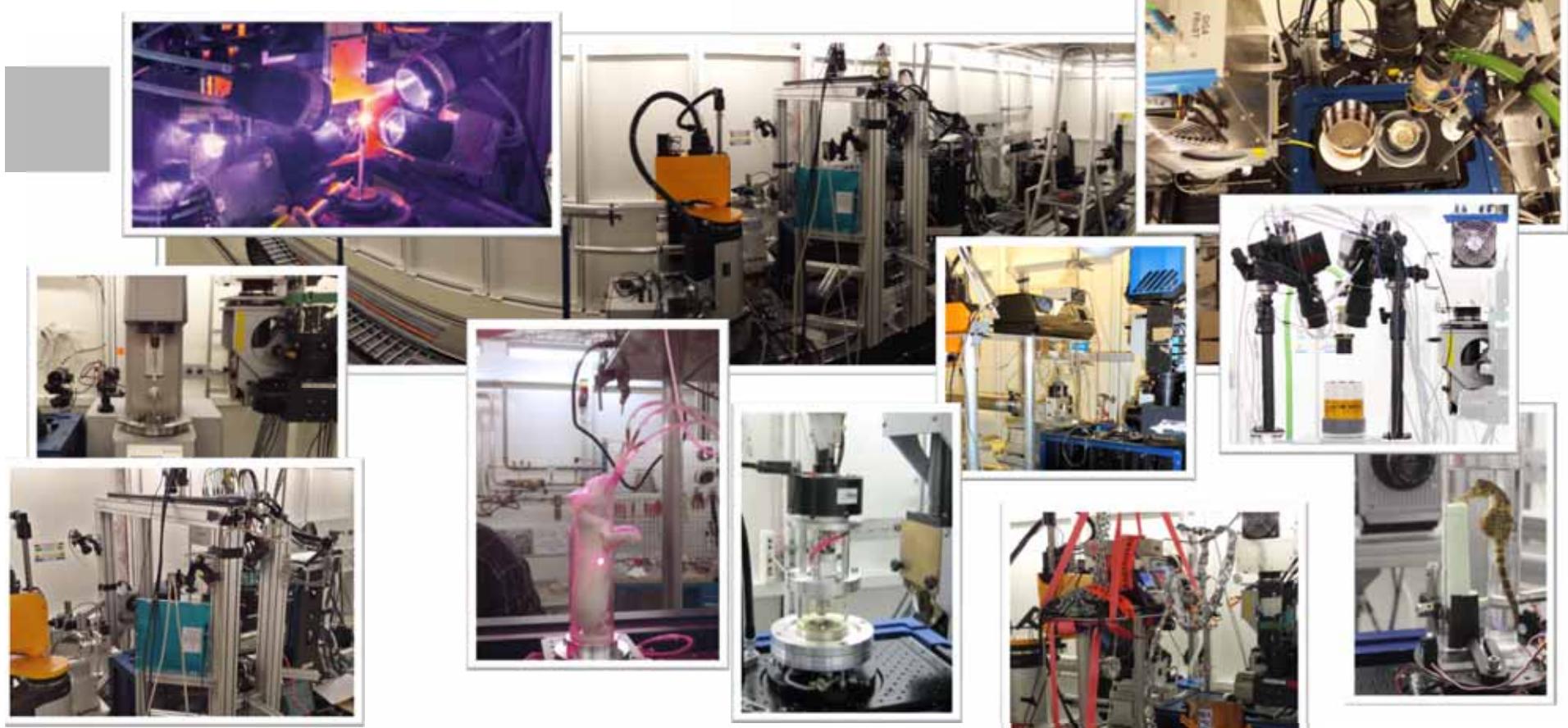
Alun Ashton
Markus Janousch
Klaus Wakonig
...



TOMCAT beamlines (plural)

Material by
Marco
Stampanoni
& team

Needs for In-situ :: In-vivo :: Operando

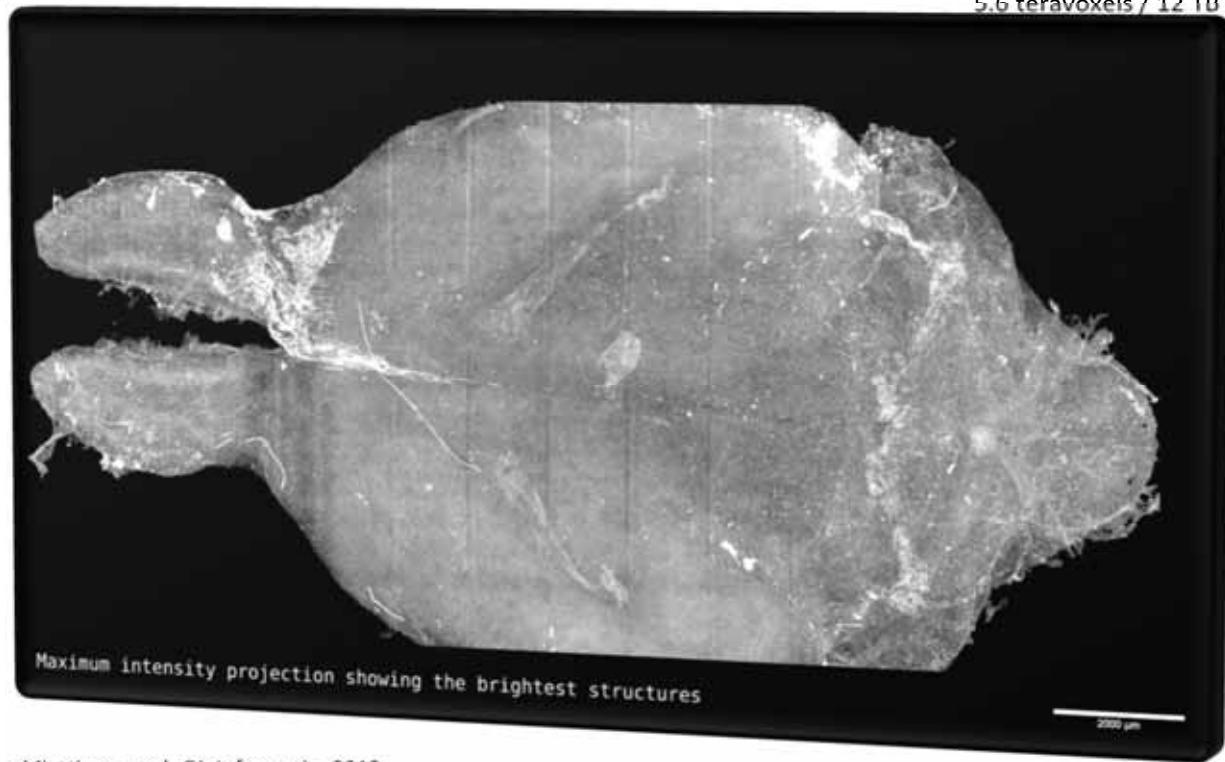




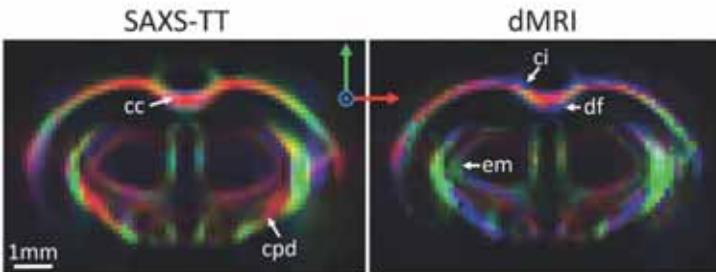
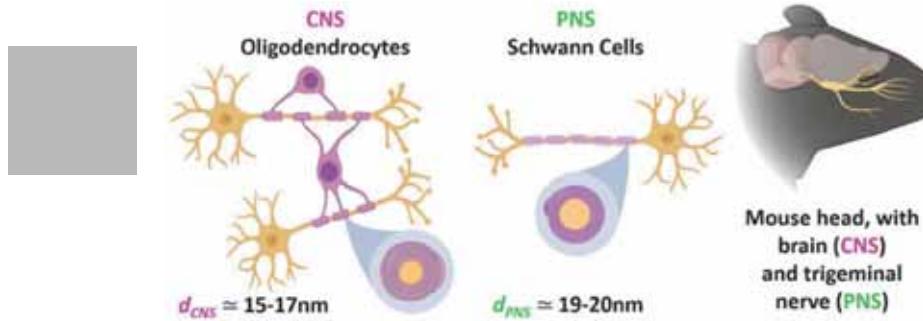
Hunting the complete mouse brain microvascular architecture

This is a stitched tomogram of more than 1000 single volumes, assembled non-rigidly.

5.6 teravoxels / 12 TB



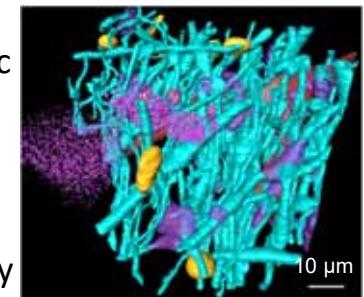
X-ray neuroimaging: 3+ complementary methods



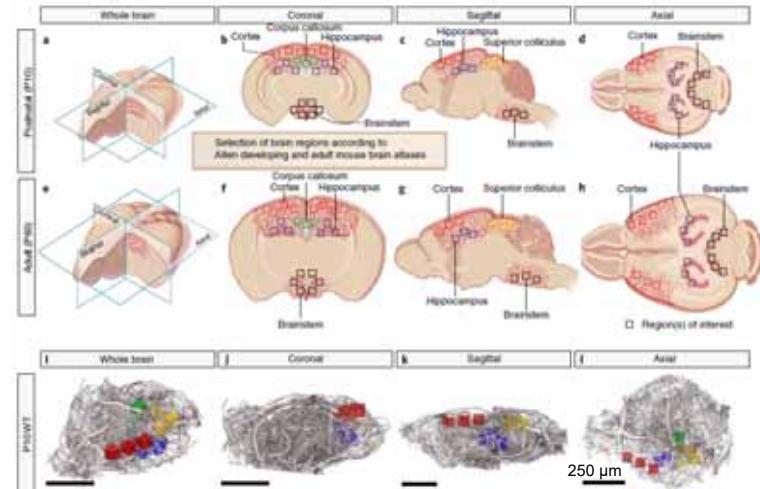
Scanning SAXS tomography at cSAXS or PX I

- M. Georgiadis *et al.*, Nature Comm. **12**, 2941 (2021)
 H.T. Tran *et al.*, Front. Neurosci. **14**, 570019 (2020)
 T. Wälchli *et al.*, Nature Protocols **16**, 4564 (2021).

Cryo ptychographic tomography using OMNY at cSAXS



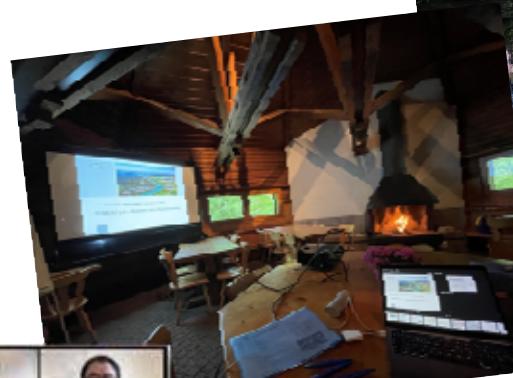
Full-field tomography (with 'stitching') at S- and I-TOMCAT



The TOMCAT crew is working on it!

2021

2019...



2020-2021...



TOMCAT2.0 - Brainstorming

Summary

The future for your science at SLS hard X-ray crystallography and imaging beamlines is bright.

We are looking forward to collaborate with you on your challenging and fascinating scientific projects.



[https://www.luzern.com/en/highlights/listicals/
top-five-sunrise-locations-in-central-switzerland/](https://www.luzern.com/en/highlights/listicals/top-five-sunrise-locations-in-central-switzerland/)

LSB Mission

Excellence from source to detector / excellence in life and materials science

We are at the forefront of method and instrumentation developments to render otherwise inaccessible information accessible via X-ray based techniques. In collaborations with Swiss and international scientists we employ these strengths for state-of-the-art research with focus on life and materials science. We are competent partners for academic and industrial users.