

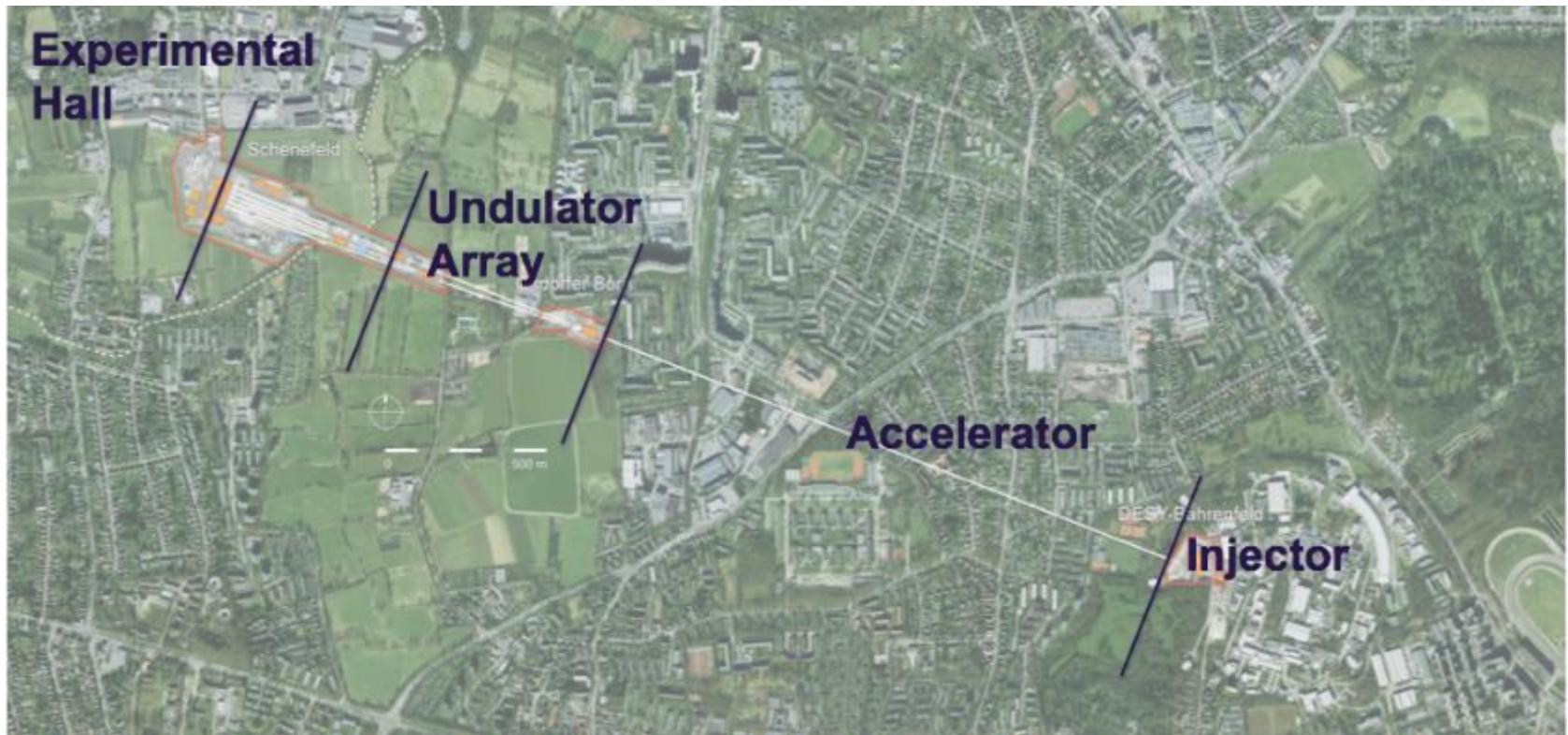
Seeing elusive protein structures at high resolution: Instrumentation for XFEL crystallography and imaging

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SPB/SFX
Postdoctoral Scientist

Hamburg DESY, 09 June 2017



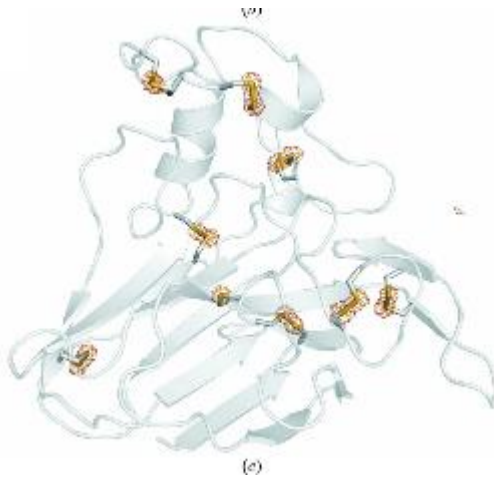
The European XFEL



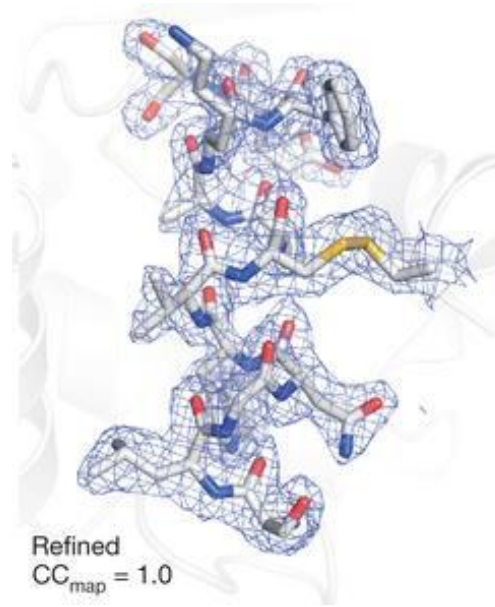
Traditional X-ray sources



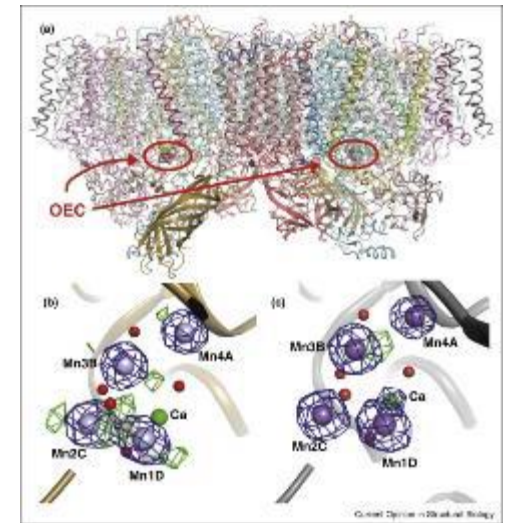
Protein structure



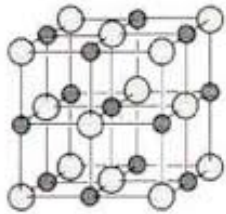
Nass et al. (2016)



TRM Barends et al. (2014)



R Neutze (2015)



Scattered x-rays is proportional to N^2
(~ 100 x 100 x 100 elements)

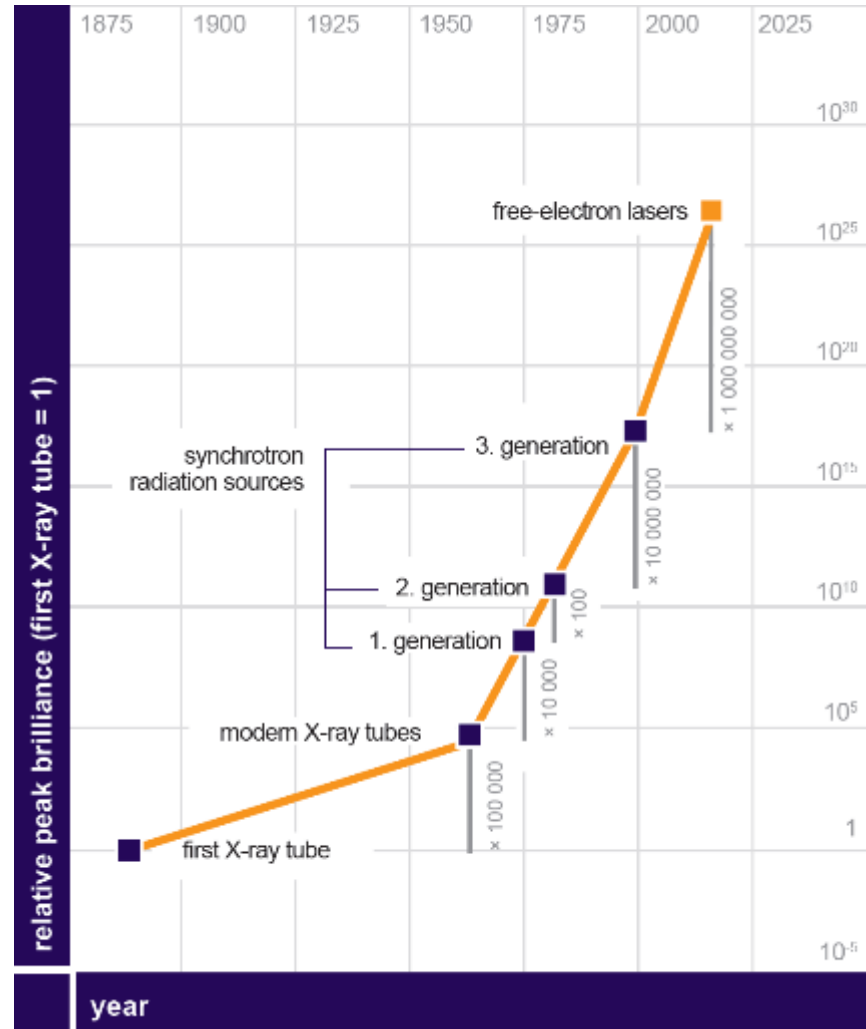
- If $N=1$, then scattering is proportional to 1
(~ a million times less than above)

Conclusion: Need a lot more x-rays

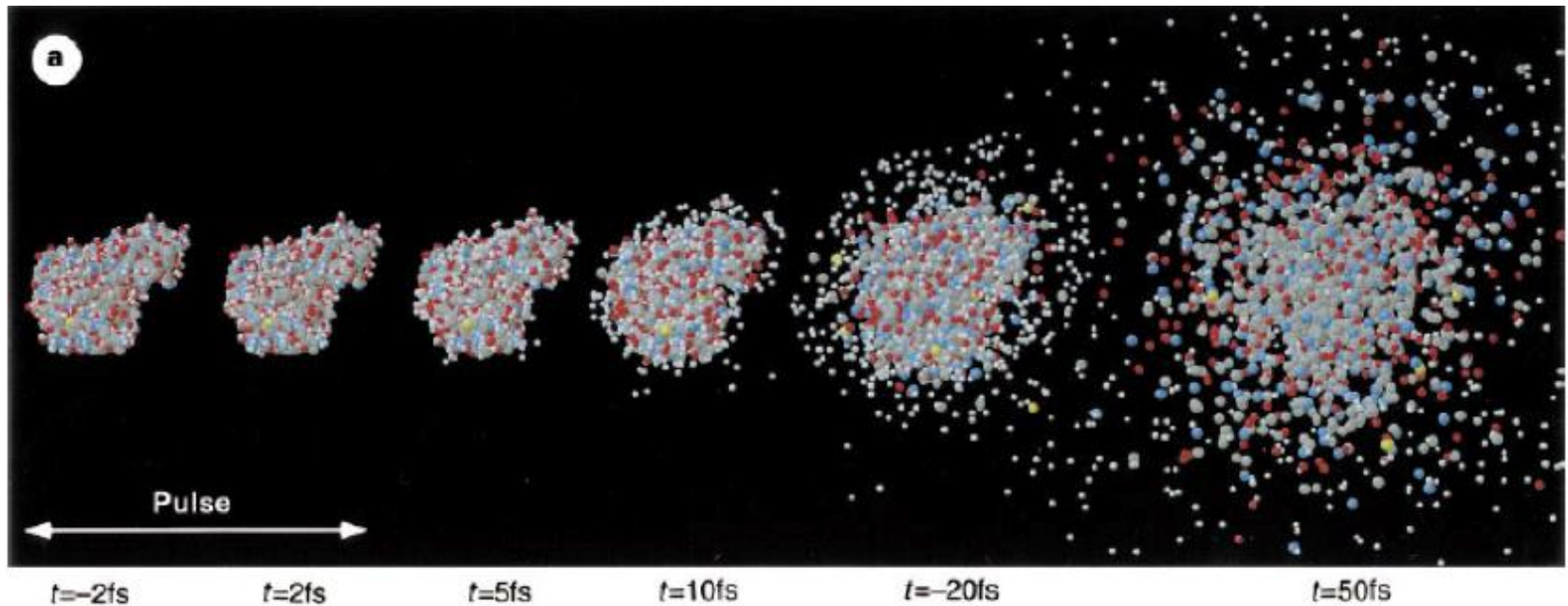
SASE radiation

FELs offer improvements over 3rd generation synchrotrons

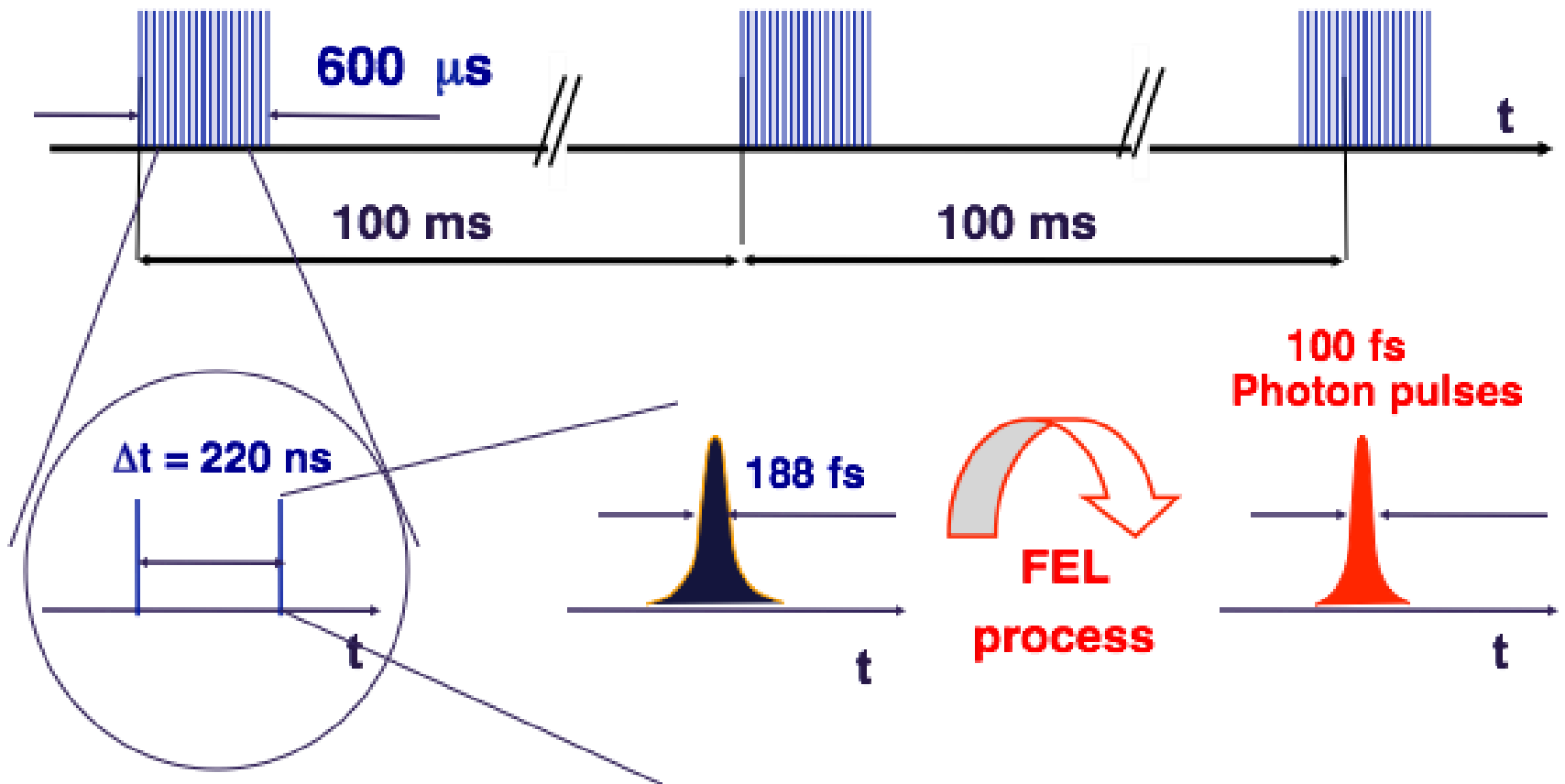
- Orders of magnitude brighter
- Extremely short pulse duration
- Both μm and nm focus



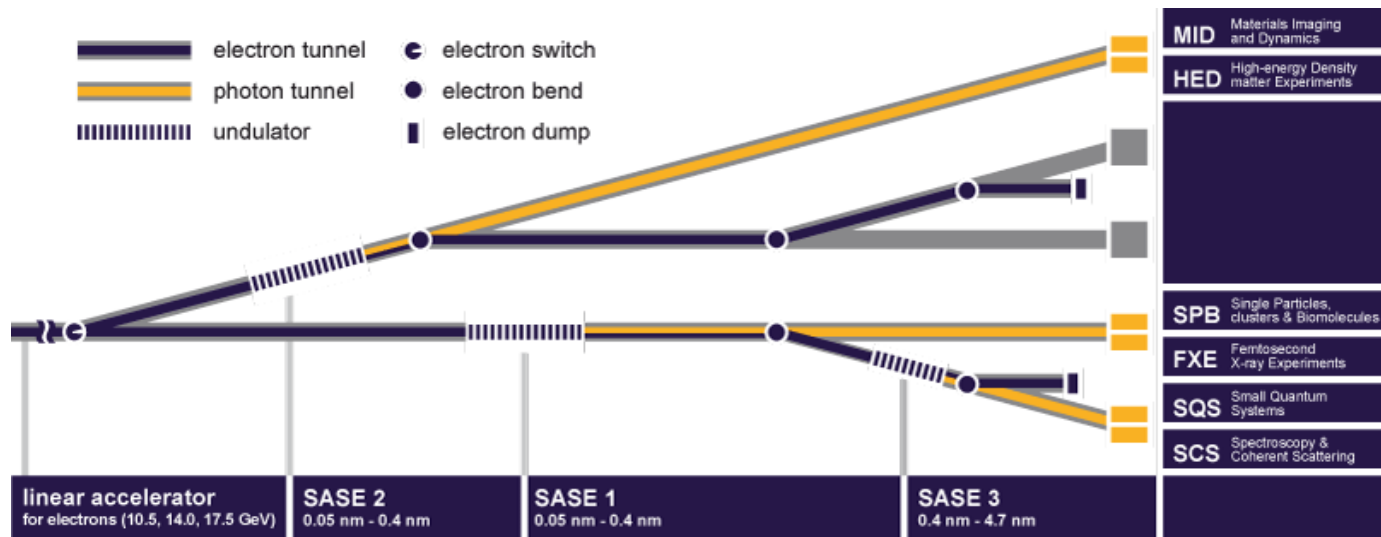
Diffraction before destruction



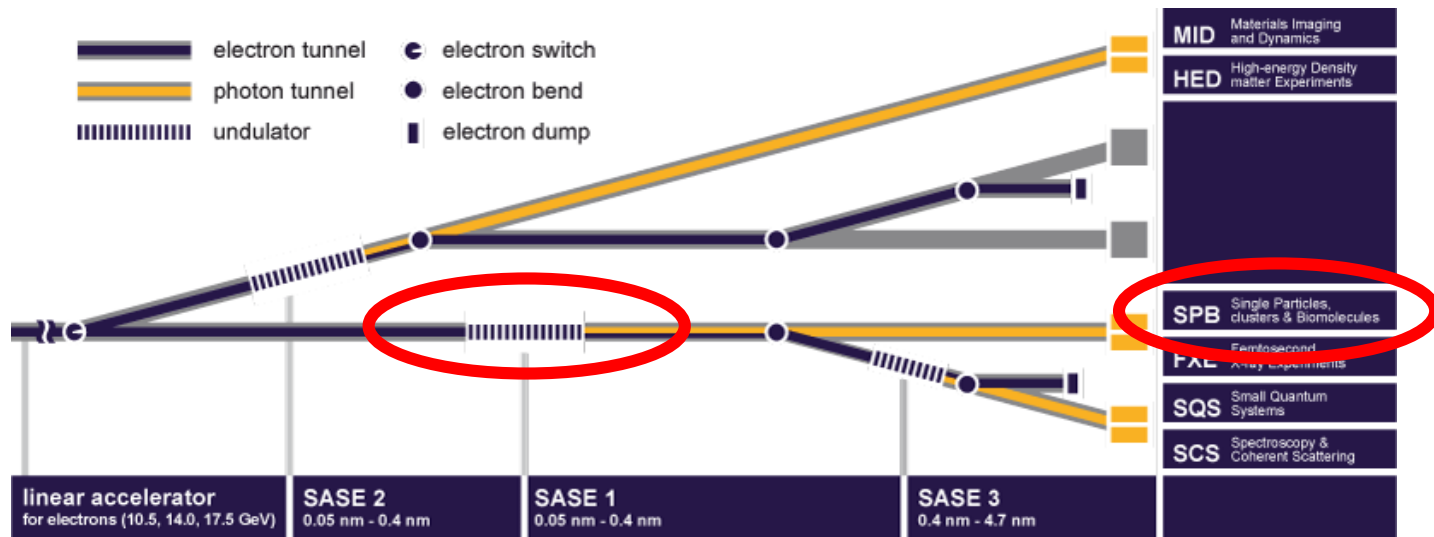
SASE pulse train



Undulator



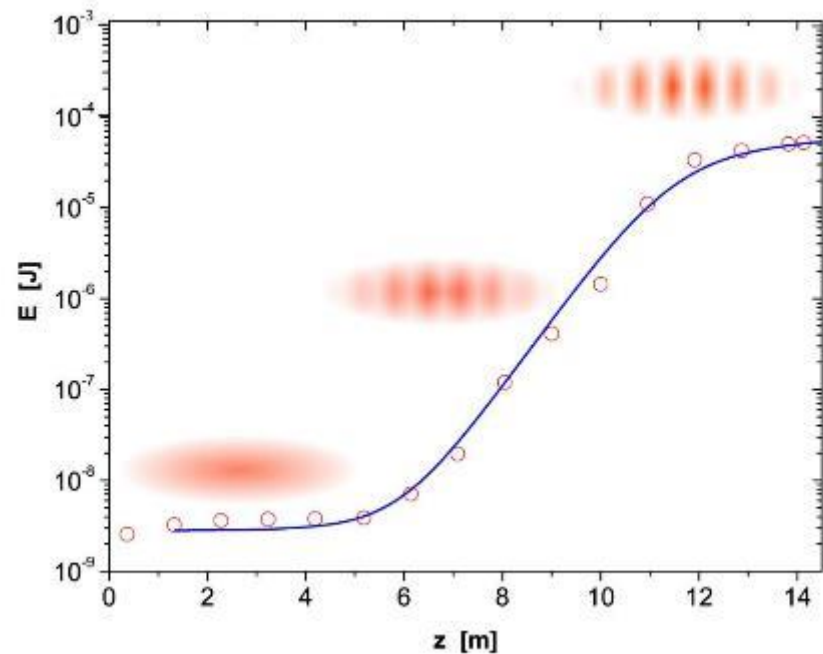
Undulator



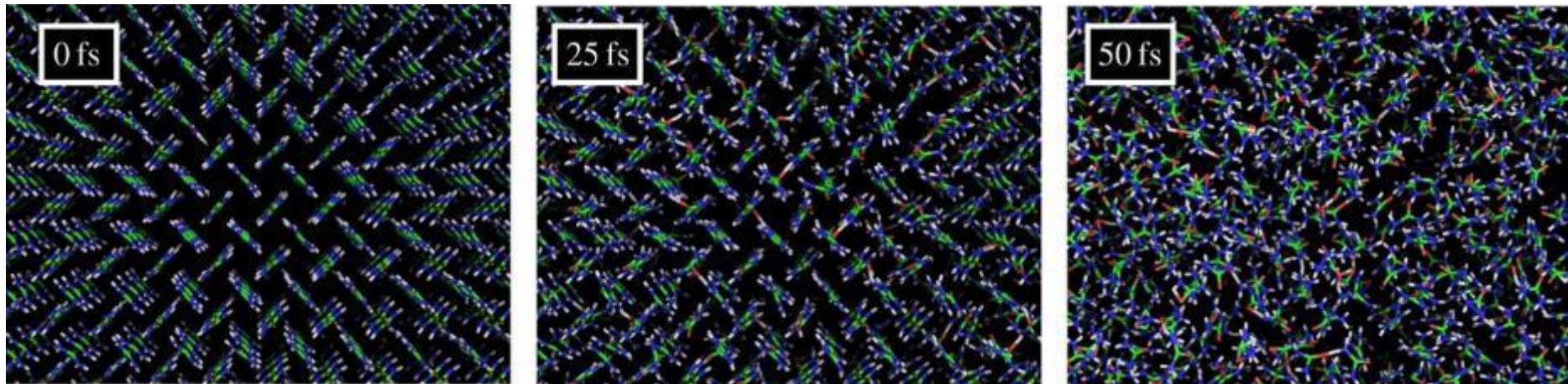
Undulator

Coherent radiation is emitted from 175 m of SASE undulator

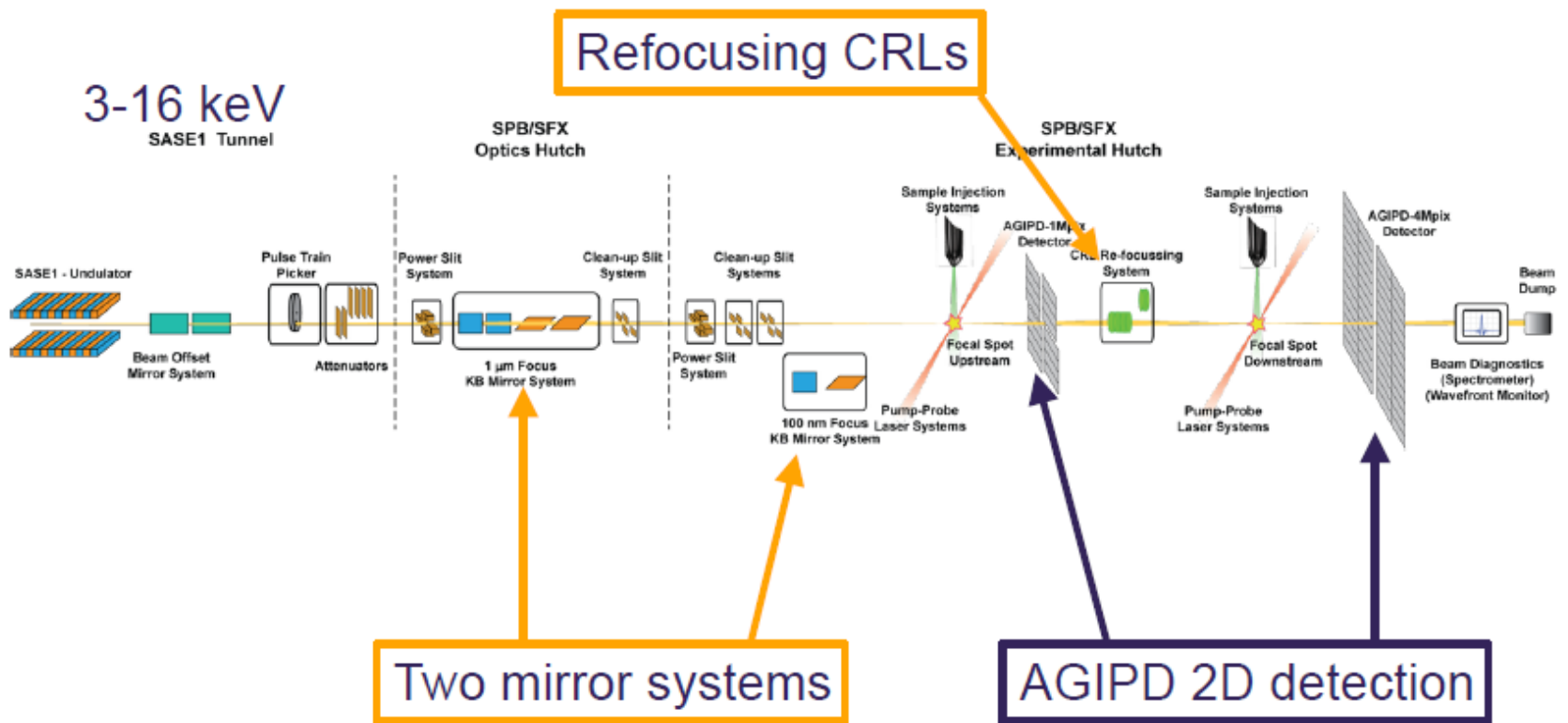
- Initial section of undulator produces radiation
- The radiation couples with the electron's to bunch the electrons
- Saturation occurs when the electrons are completely microbunched
- This process greatly amplifies the radiation emitted



Radiation damage

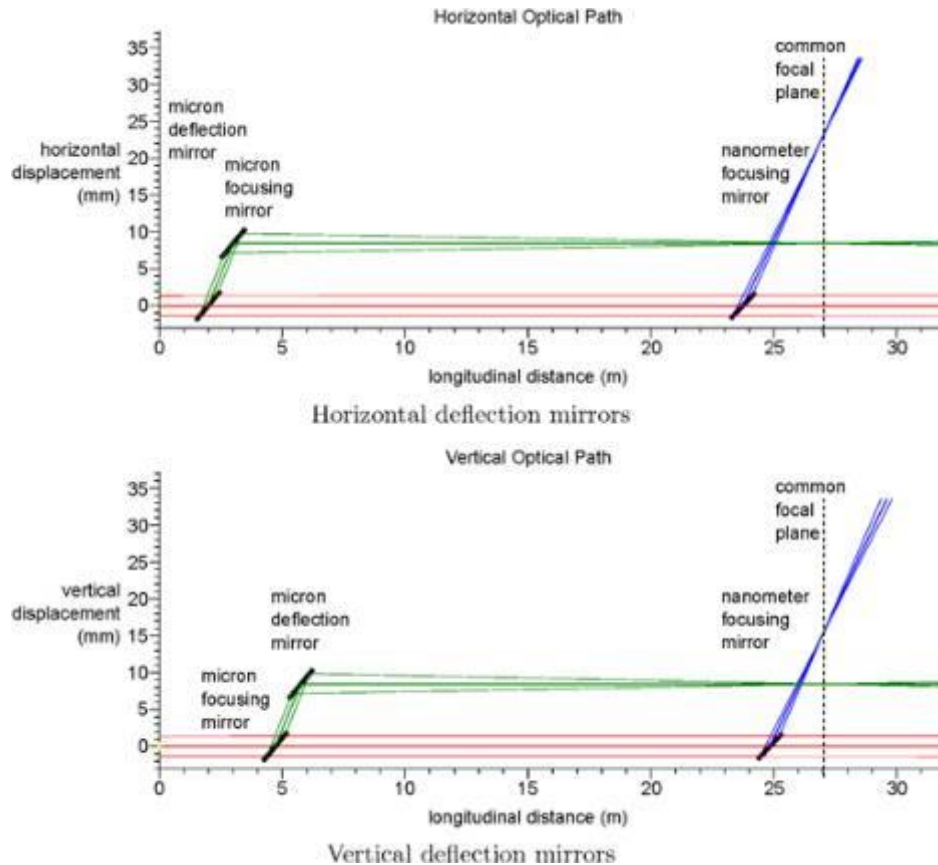


Imaging biomolecules and X-ray radiation



Technical Design Report: Scientific Instrument SPB, (2013)
A. P. Mancuso, et al.

KB mirror systems



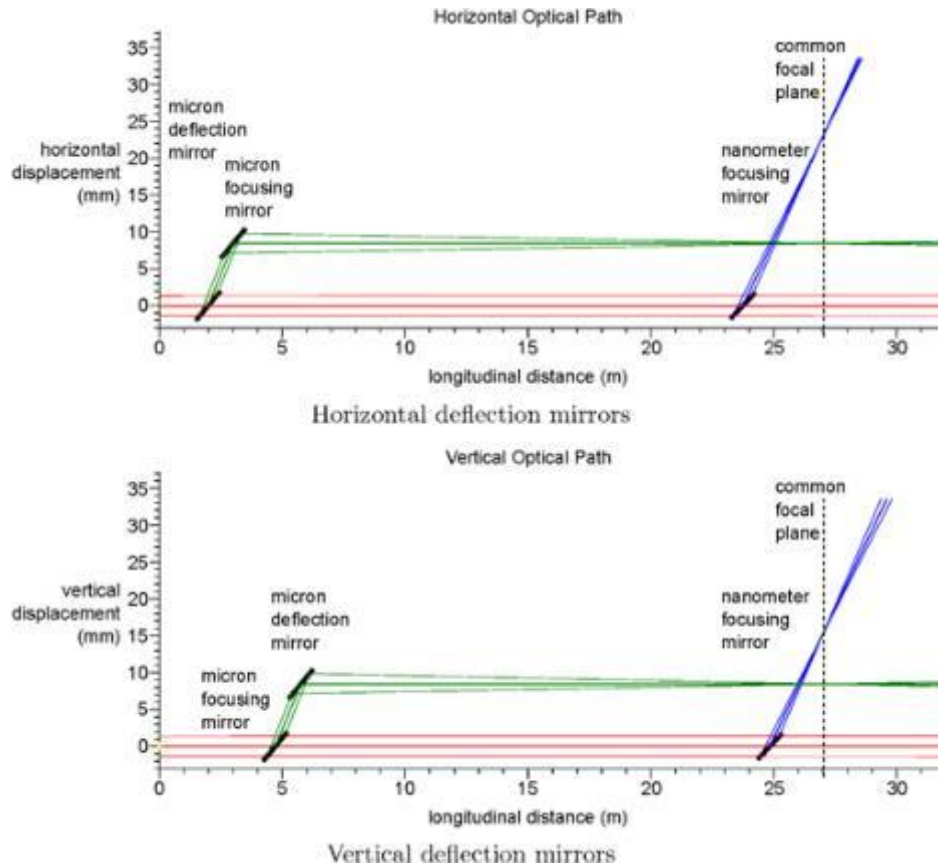
Kirkpatrick-Baez mirrors will produce a 1 micron and 100 nanometer focused beam

■ 4 bounce system (1 μm) in green

■ 2 bounce system (100 nm) in blue

■ B₄C and Ru coatings

KB mirror systems



Kirkpatrick-Baez mirrors will produce a
1 micron and 100 nanometer focused
beam

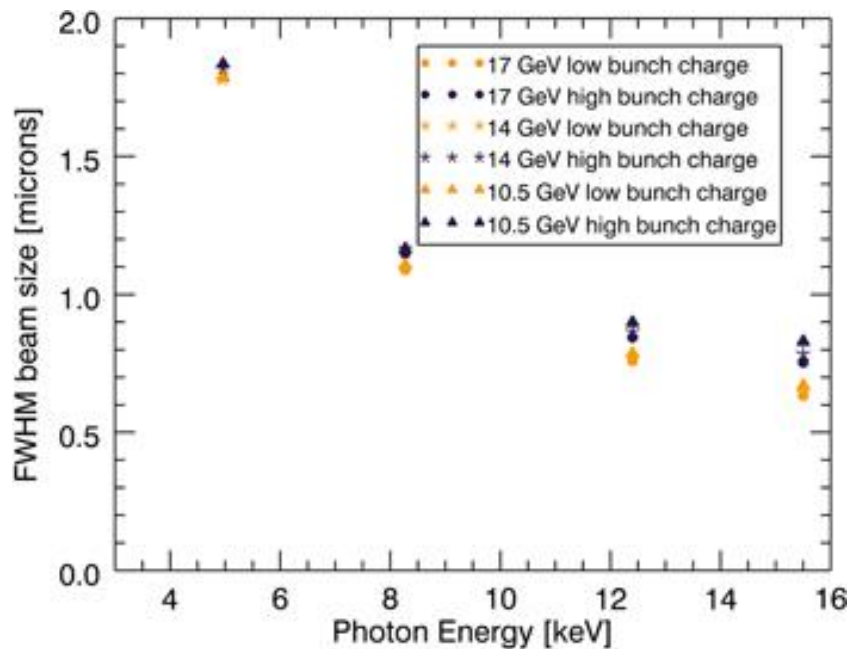
■ 4 bounce system (1 μm) in green

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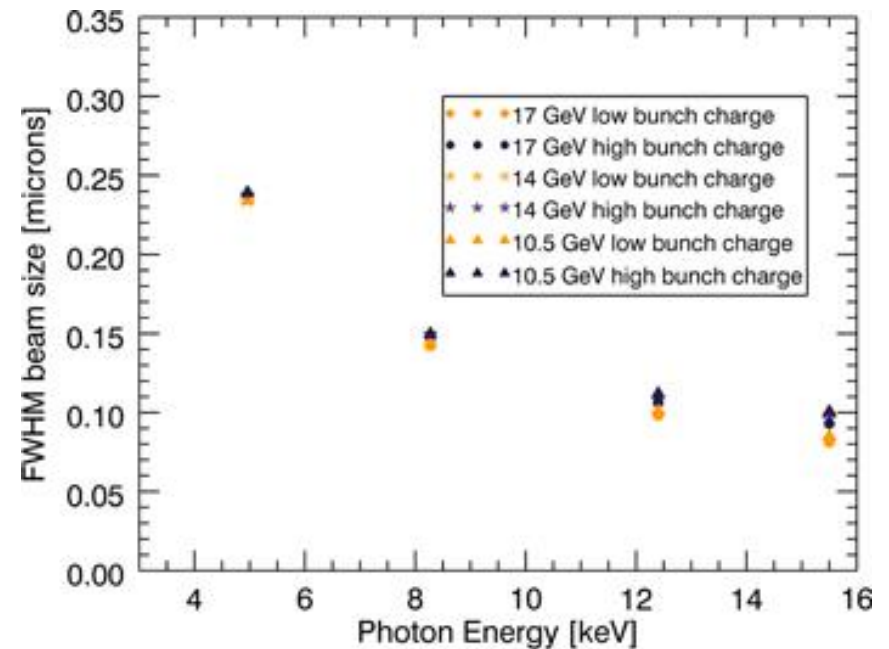
■ B₄C and Ru coatings

KB mirror systems

micron focus



nano focus



Manufacturing achievements

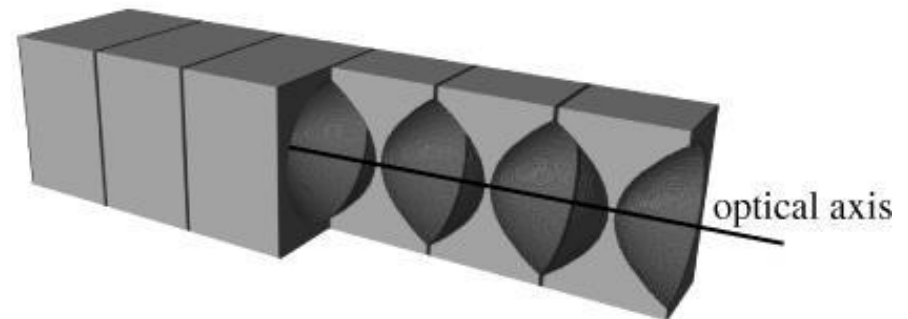
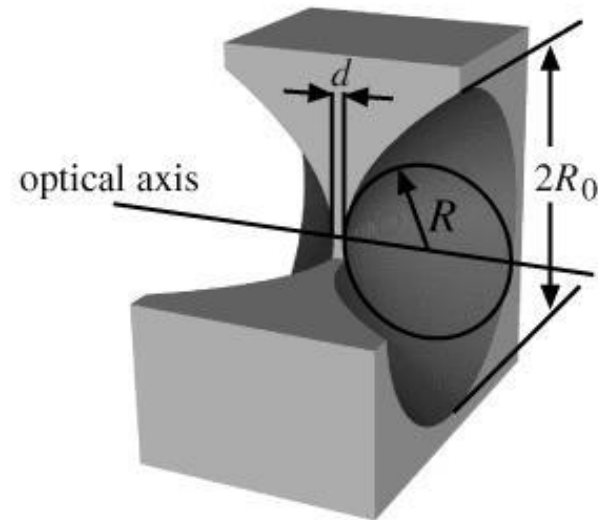


- 970 x 25 mm clear aperture
- 1.8 nm peak to valley variation
- 2x 50 nm thick polished coatings
 - B₄C and Ru
 - 950 x 10 mm

Compound refractive lenses

Initial operation will use the CRL in the tunnel

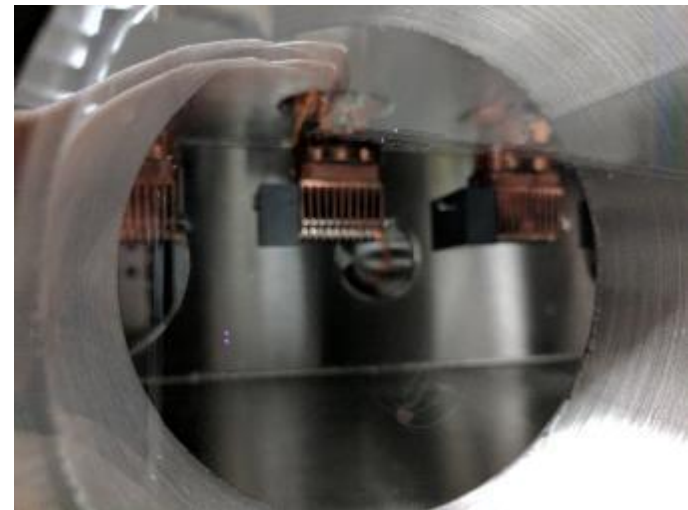
- CRL is a repeating structure of parabolic surfaces
- 12 mm diameter
- 2 mm thickness
- Focus to 2.5 μm diameter



Cassette ordering

Initial operation will use the CRL in the tunnel

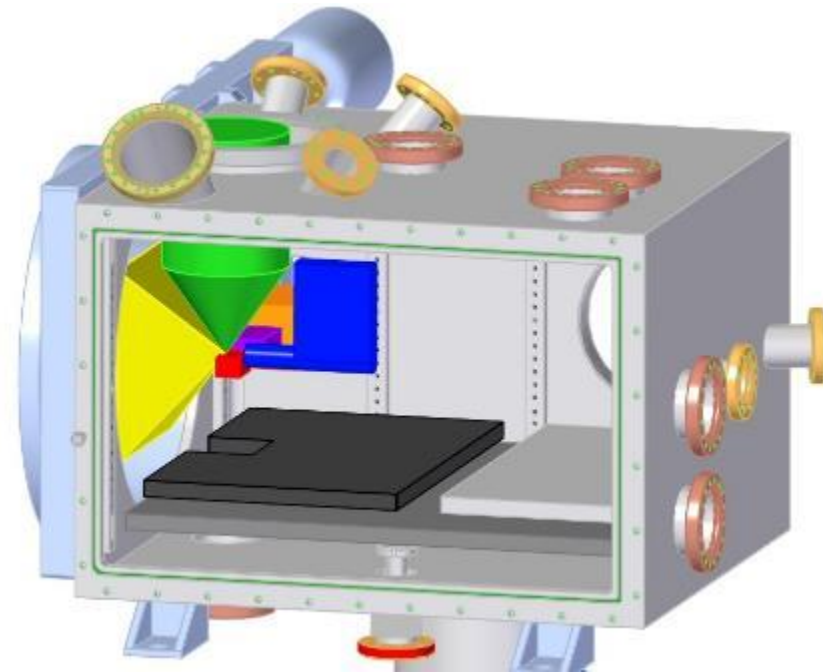
- 1, 2, 4, 8, 10, 10
- Allows the change of X-ray energy while still maintaining focal location



Interaction region

There are two inline interaction regions, the 1st upstream region is available day 1

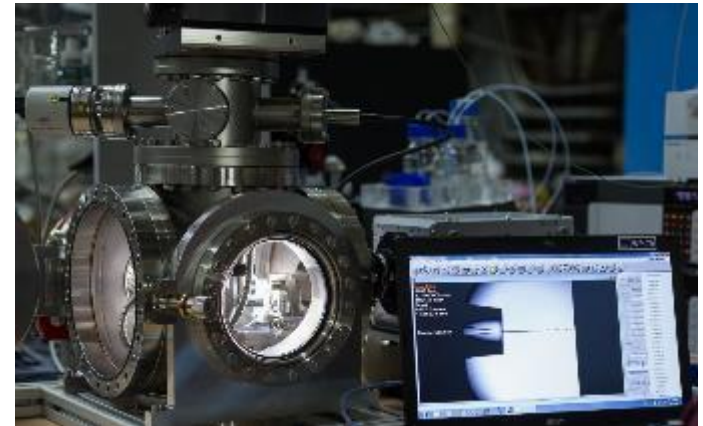
- Day 1 will provide 60 pulses per train
10 Hz
- Photons at 8.4 keV
- Pulse energy at 2 mJ
- Pulse duration 43 fs
- 120° open scattering cone



Sample injection

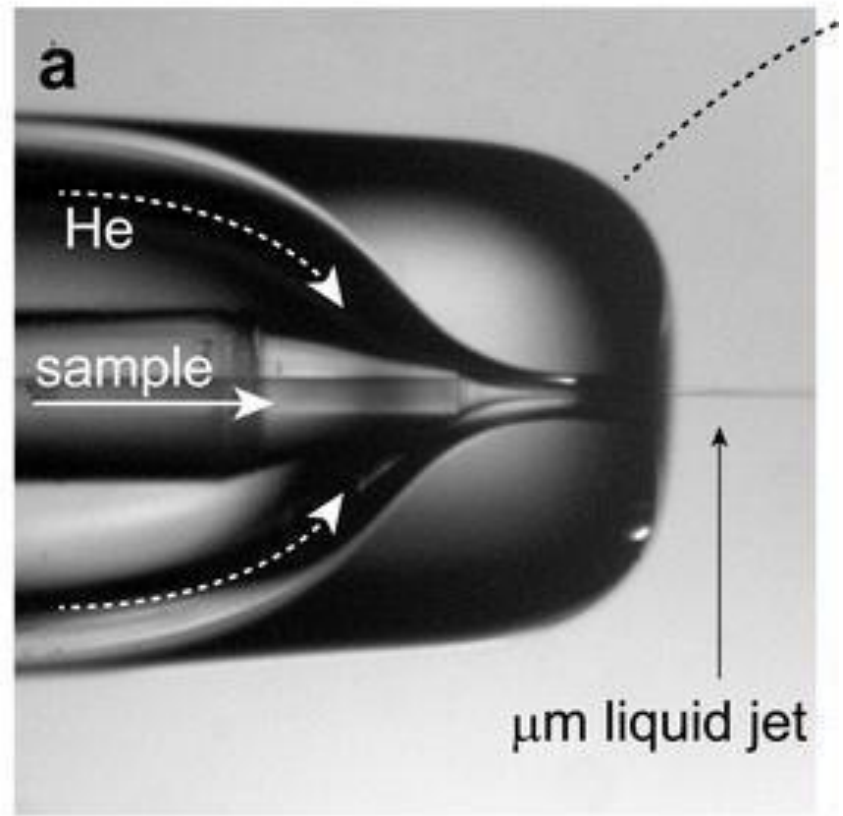
Many methods have been adopted to deliver sample into the interaction region

- GDVN
- Aerosol
- Fixed targets

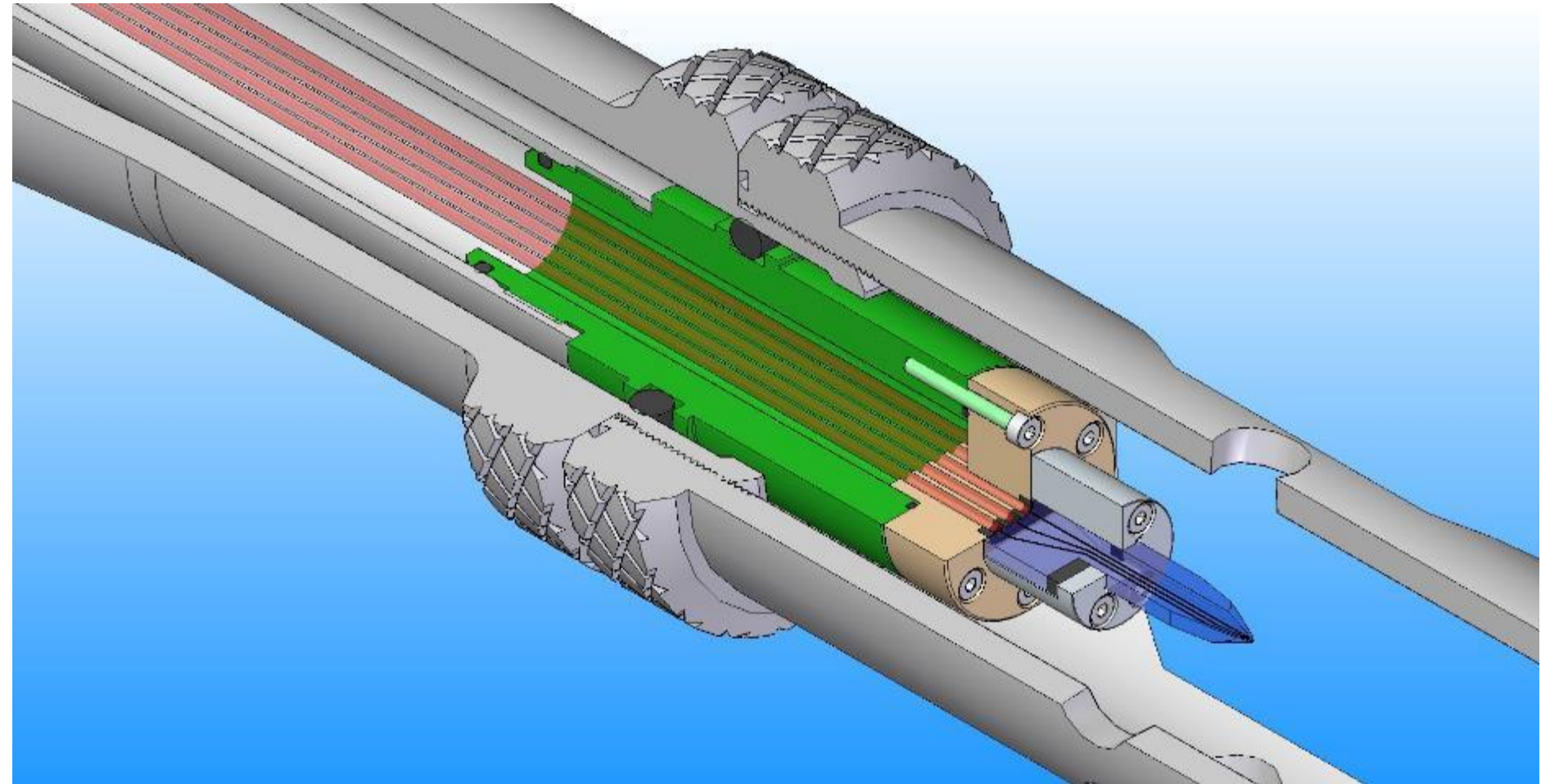


Gas Dynamic Virtual Nozzle

- Capillary guides the sample into the interaction region
- Inert gas focuses the jet into a thin stream
- A μm thin jet can be maintained over more than 100 μm distance
- A modified system can also handle viscous solutions



Design study of a microfluidic mixing nozzle in 25 mm nozzle rod



Aerosol injectors

- Injector creates particle beam
- Pressures decrease through the injector nozzle
- Rapid motion of sample

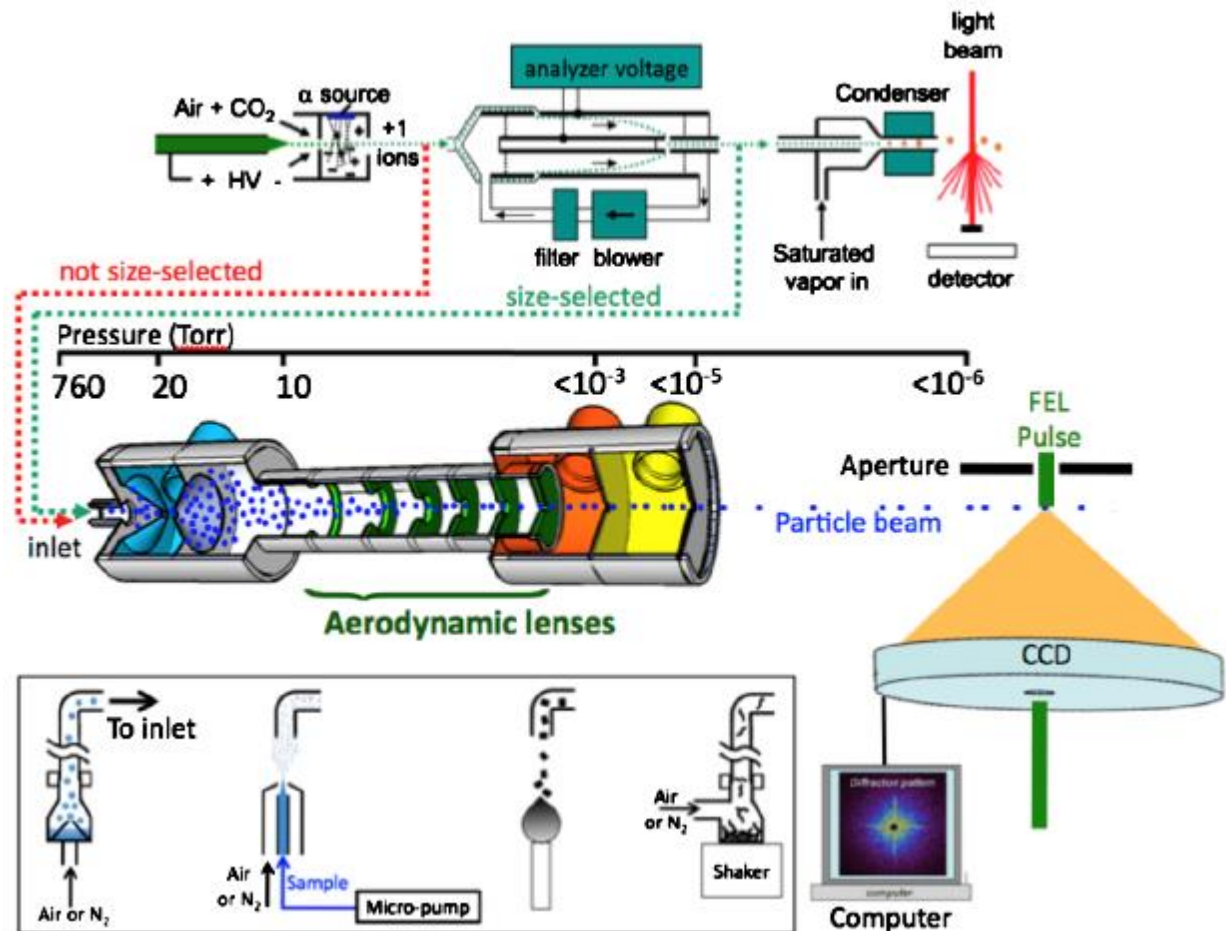
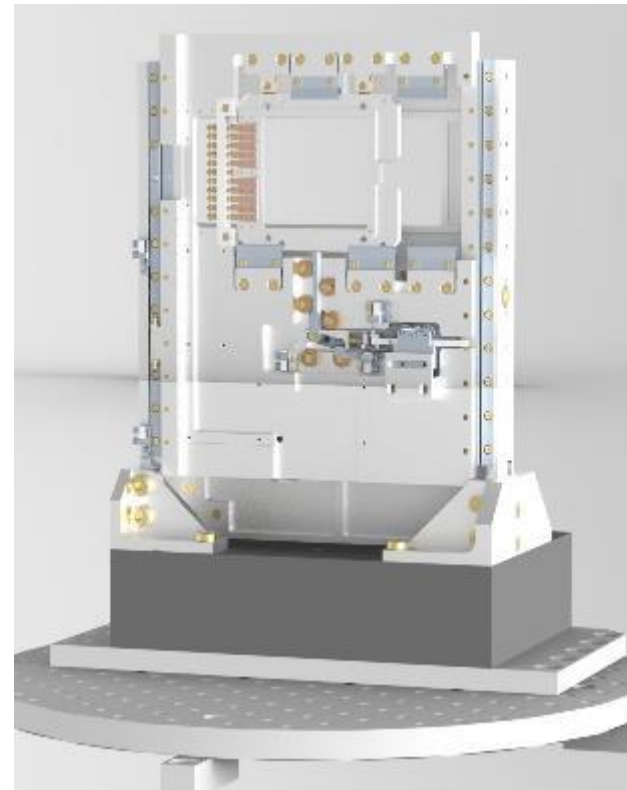
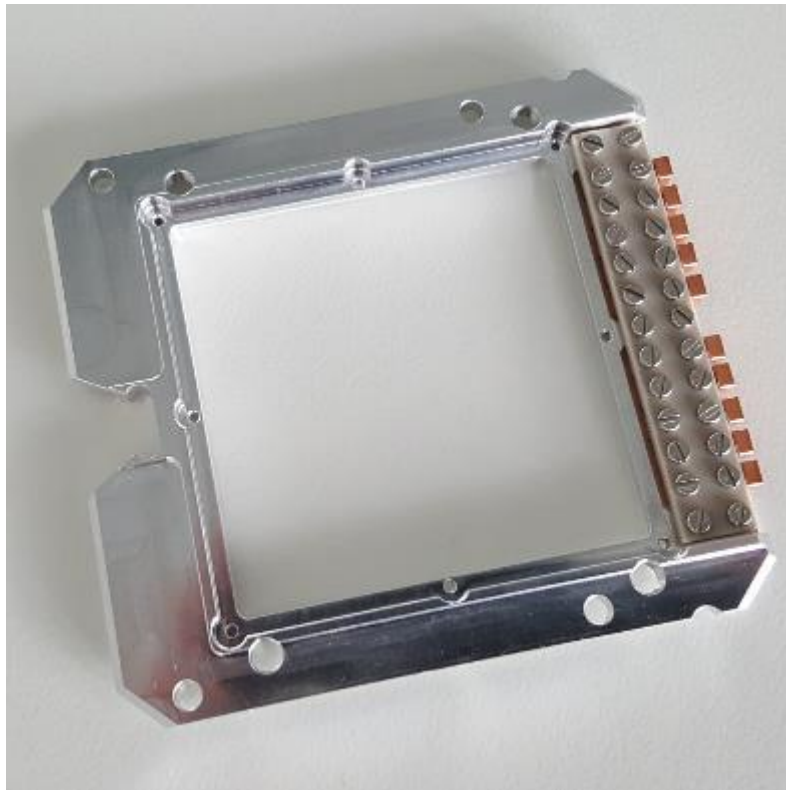


Image: Joachim Schulz

Fixed targets

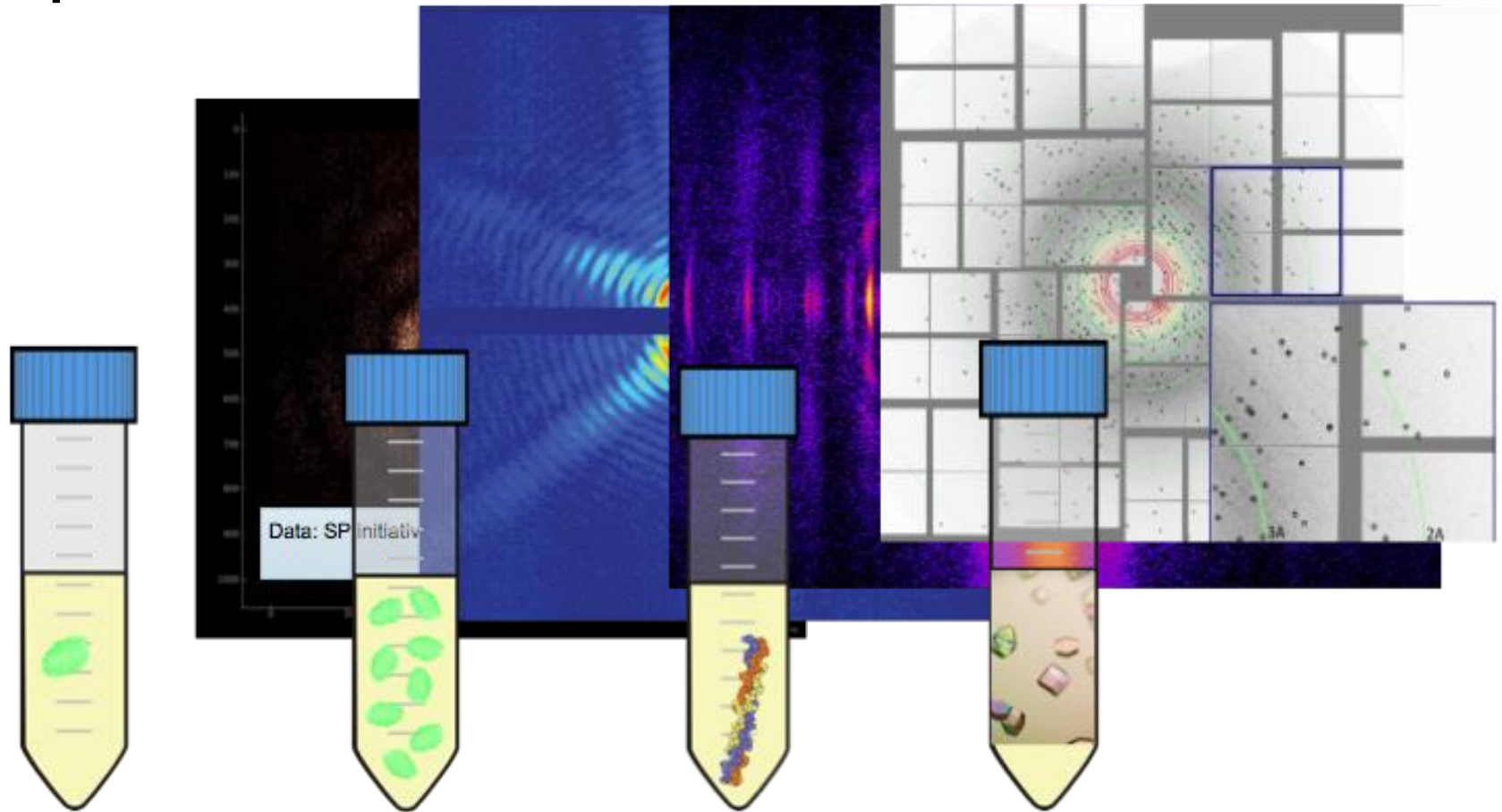


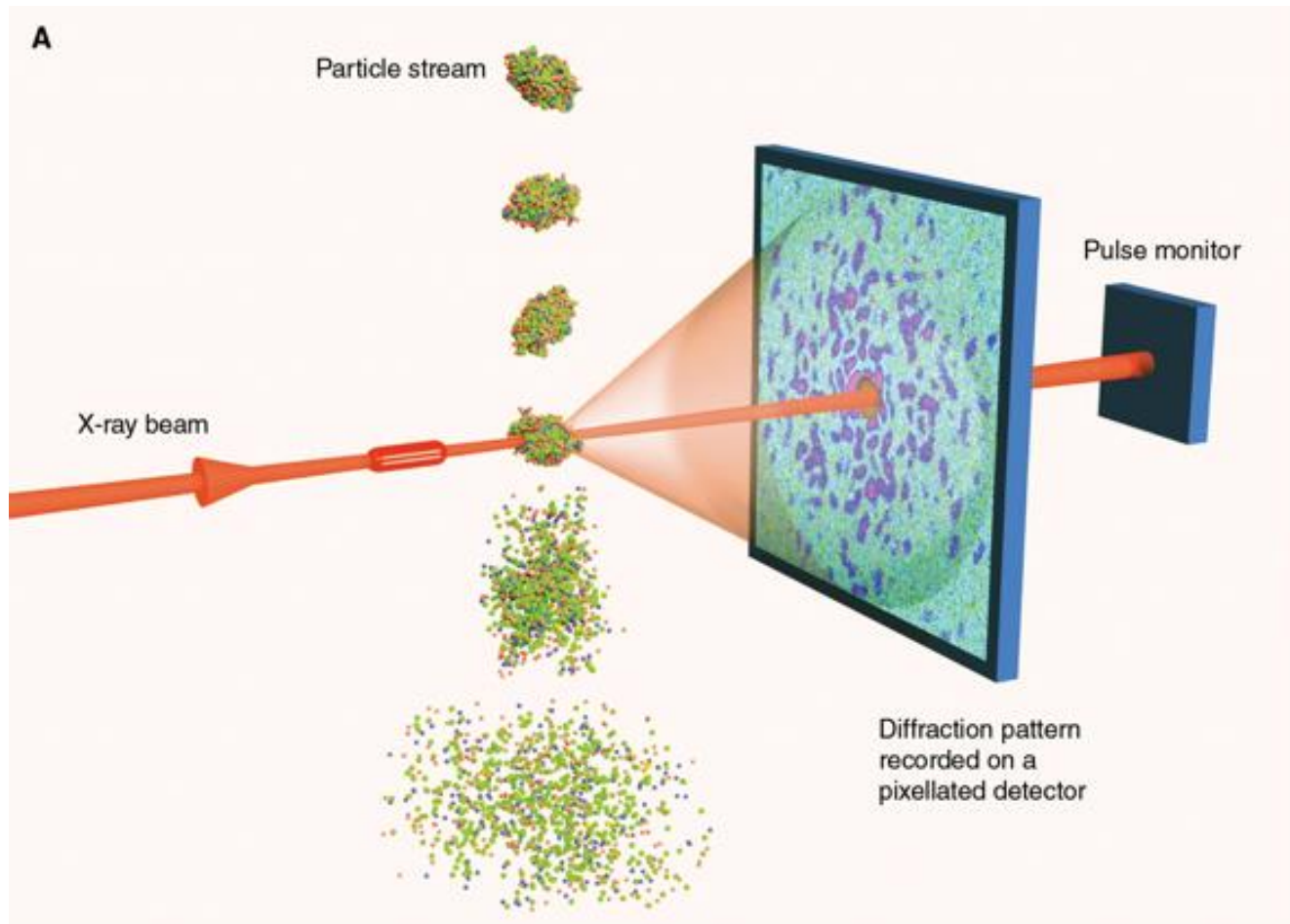
AGIPD Detectors



- 6 – 13 keV
- 1×10^4 at 12 keV dynamic range
- 5σ at 12 keV sensitivity
- Number of storage cells 352
- Can store images at 4.5 MHz

Experiments



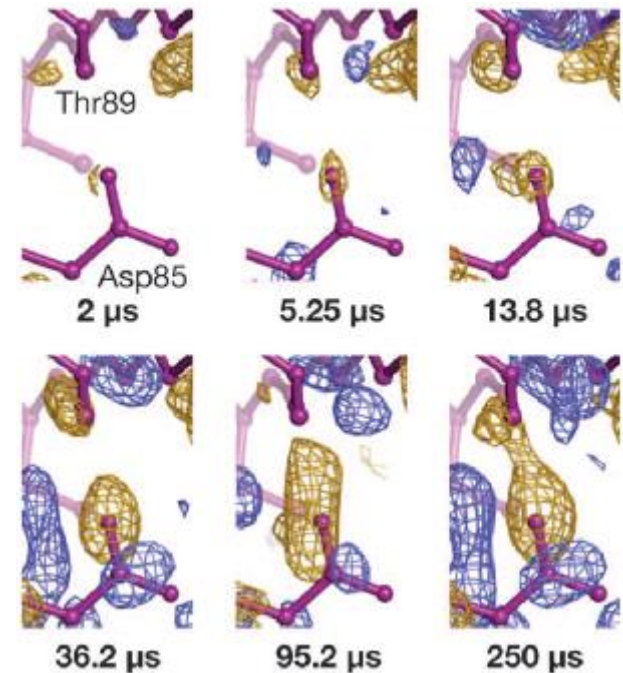


Experiments

STRUCTURAL BIOLOGY

A three-dimensional movie of structural changes in bacteriorhodopsin

- Time-resolved SFX
- Bacteriorhodopsin conformational changes
- 10 fs long pulses at SACLA

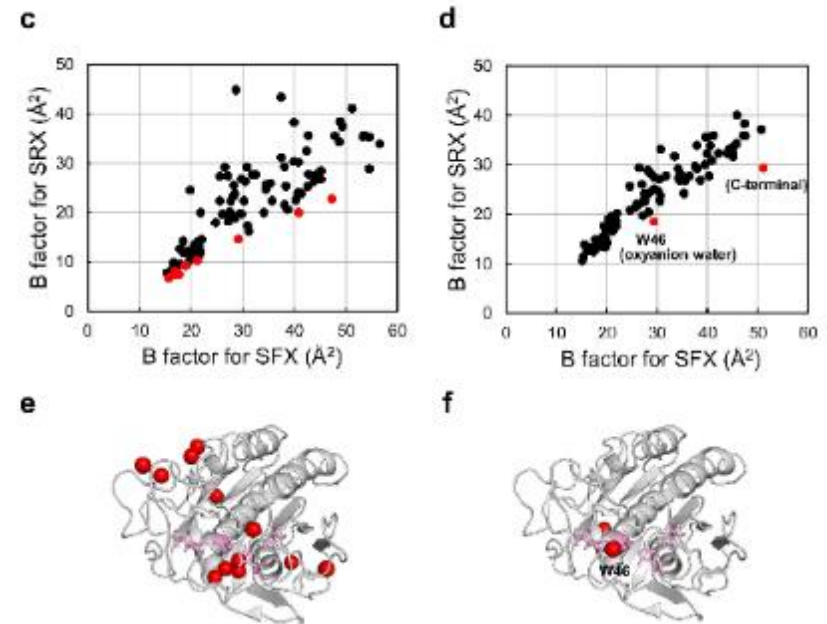


Eriko Nango, et al 2017, Science

Experiments

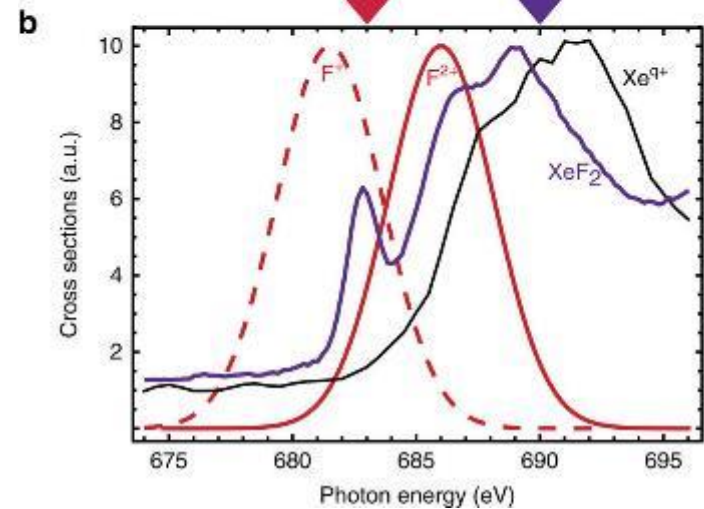
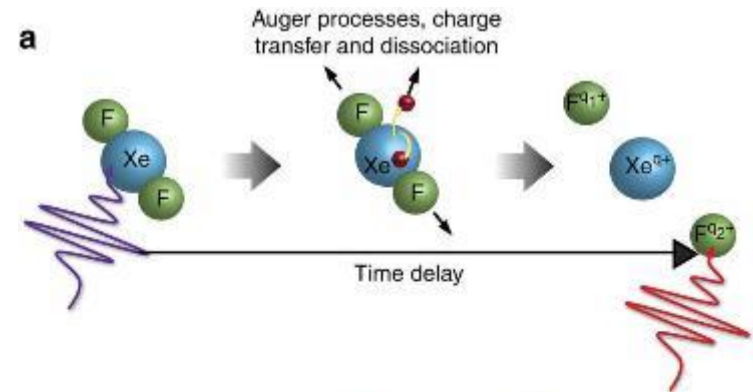
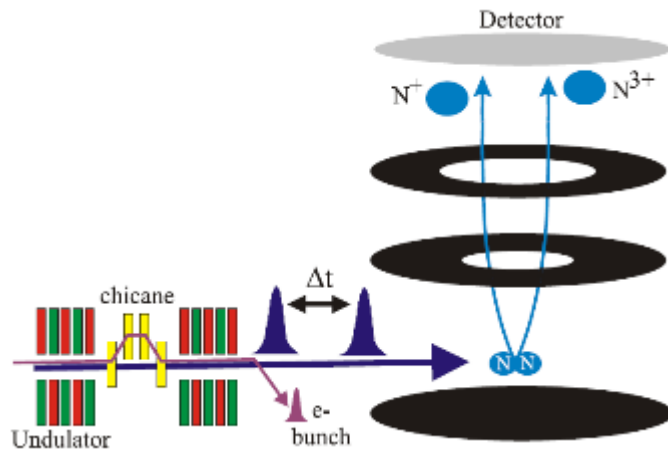
Atomic resolution structure of serine protease proteinase K at ambient temperature

- High resolution SFX
- 1.2 Å res proteinase K
- Comparison of SFX and SRX
B-factors



Experiments

- Two-colour X-ray-pump/X-ray-probe scheme used for studying the molecular fragmentation of XeF_2
- These reactions occur on the fs timescale, perfect for the European XFEL



Experiments



[Home](#) [Mission](#) [CXI File Format](#) [Browse Data](#) [Resources](#) [Sponsors](#) [Contact Us](#)

Browse Data

- [ID 1](#) - Single mimivirus particles intercepted and imaged with an X-ray laser
- [ID 2](#) - Single mimivirus particles intercepted and imaged with an X-ray laser
- [ID 3](#) - Femtosecond diffractive imaging with a soft-X-ray free-electron laser
- [ID 4](#) - High-resolution x-ray diffraction microscopy of specifically labeled yeast cells
- [ID 5](#) - High-resolution x-ray diffraction microscopy of specifically labeled yeast cells
- [ID 6](#) - High-resolution x-ray diffraction microscopy of specifically labeled yeast cells
- [ID 7](#) - High-resolution x-ray diffraction microscopy of specifically labeled yeast cells
- [ID 8](#) - High-resolution x-ray diffraction microscopy of specifically labeled yeast cells
- [ID 9](#) - Cryptotomography: reconstructing 3D Fourier intensities from randomly oriented single-shot diffraction patterns
- [ID 10](#) - Femtosecond free-electron laser x-ray diffraction datasets for algorithm development-

Deposit Data

If you are interested in depositing data please [contact us](#).

In the future we hope to have a more automated way to deposit data.

Acknowledgments: People—the most important components

Sample Environment

AGIPD consortium
Adrian Mancuso
Zunaira Ansari
Richard Bean
Johan Bielecki
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