

Serial crystallography – recent results, challenges and prospects

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MPI Medical Research Heidelberg



Serial crystallography

Serial collection of partial datasets of different crystal(position)s

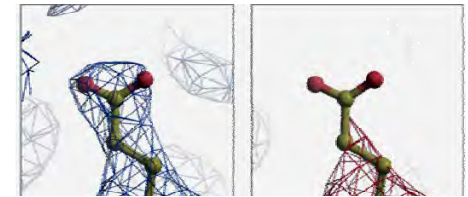
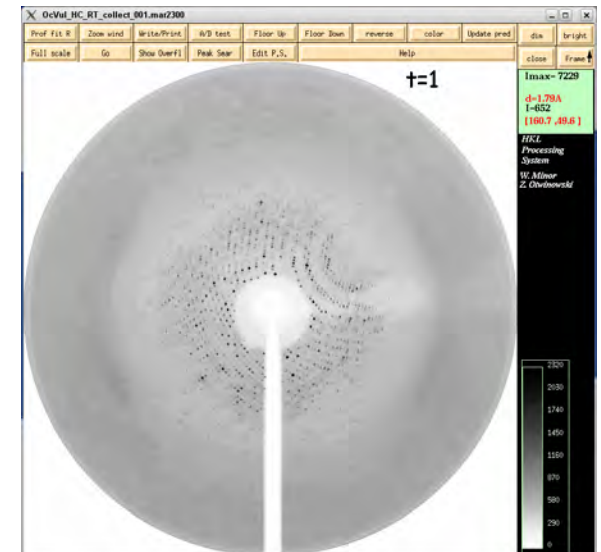
- mitigate the effects of **radiation damage**

Global damage

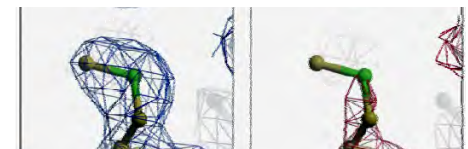
- Loss of resolution
- Increase of mosaicity
- Change in unit cell constants

Local damage

- Decarboxylations, S-S bond breakage
- Photo-reduction of redox systems, e.g. metal centers, flavins,...



Local damage to structure



Serial crystallography

Serial collection of partial datasets of different crystal(position)s

- mitigate the effects of **radiation damage**
- **time-resolved** pump probe measurements

has been performed for a very long time ...

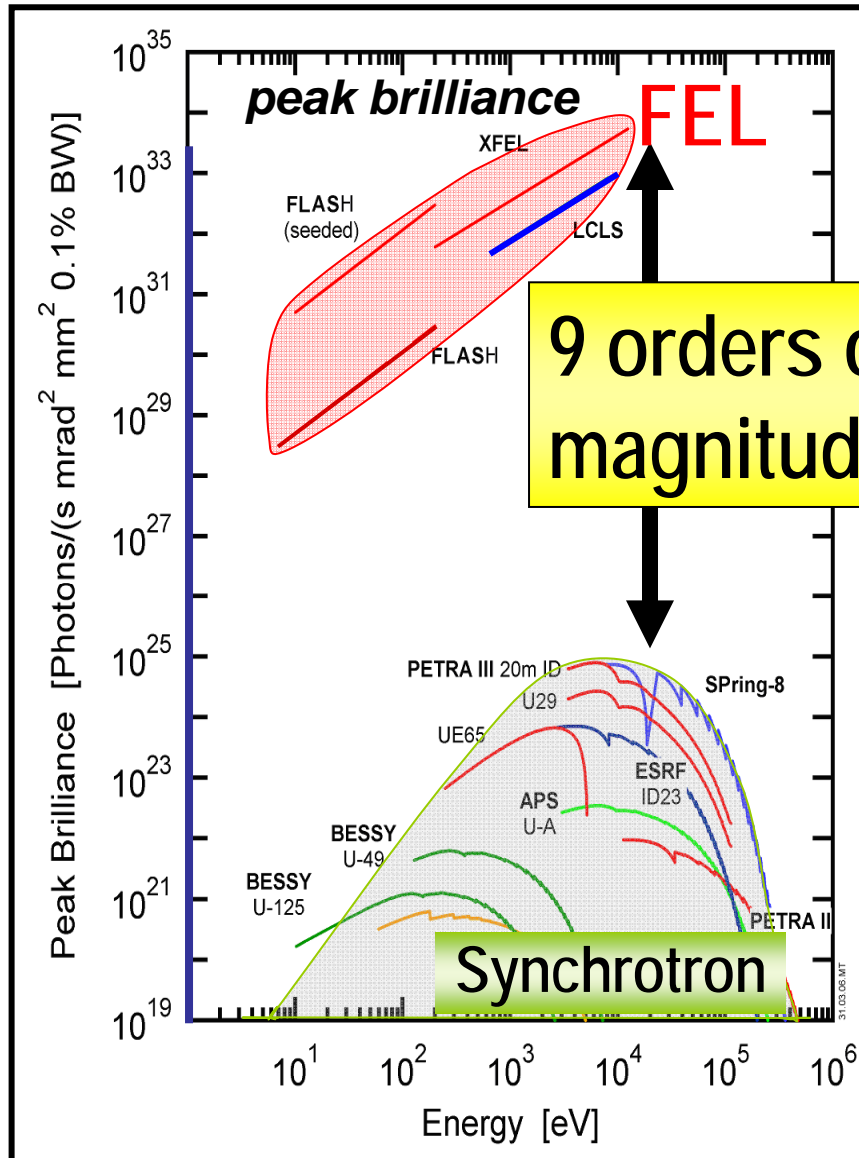
- New light sources require much faster sample change due to short lifetime of sample

XFELs

10^{12} photons/pulse, few fs exposures

- Time-resolved measurements on irreversible reactions

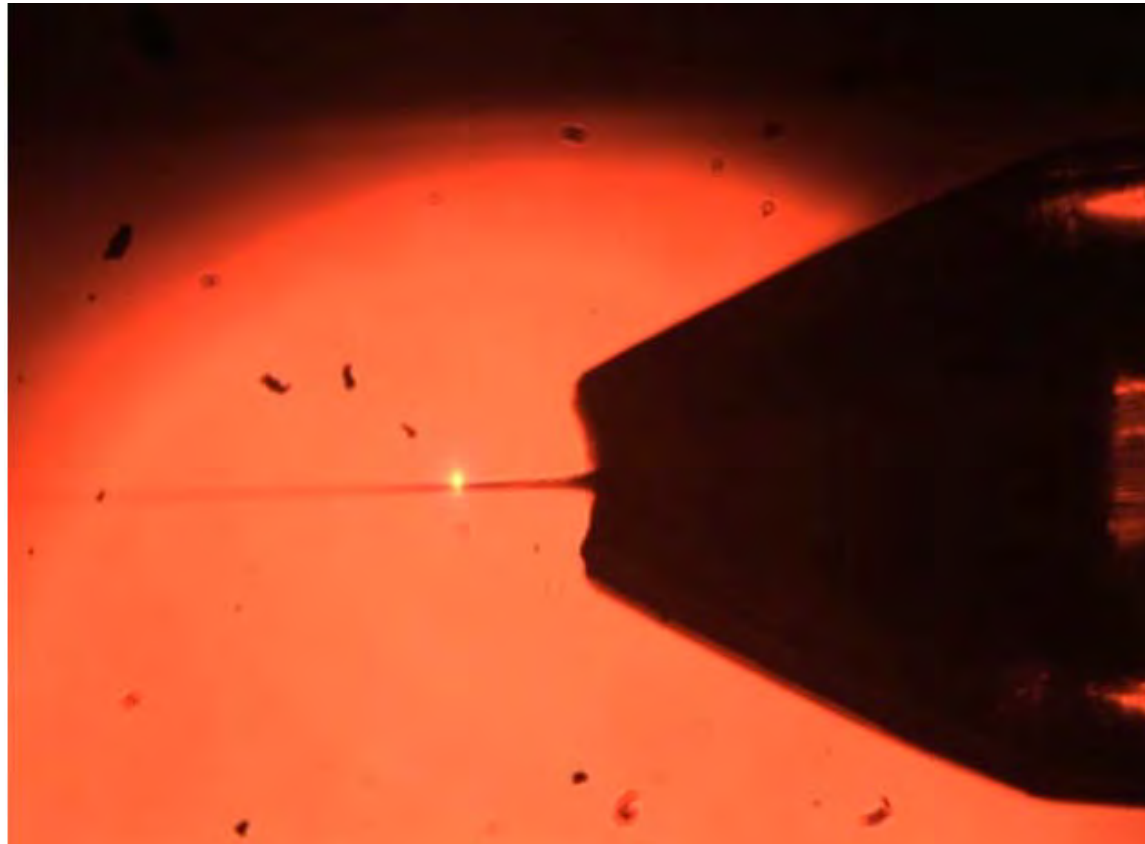
X-ray free-electron lasers



- FLASH: 2005
- Fermi: 2009
- LCLS: 2009
- SACLA: 2011
- Fermi 2011
- XFEL: 2017
- PSI, LCLSII, KVI, Shanghai,...

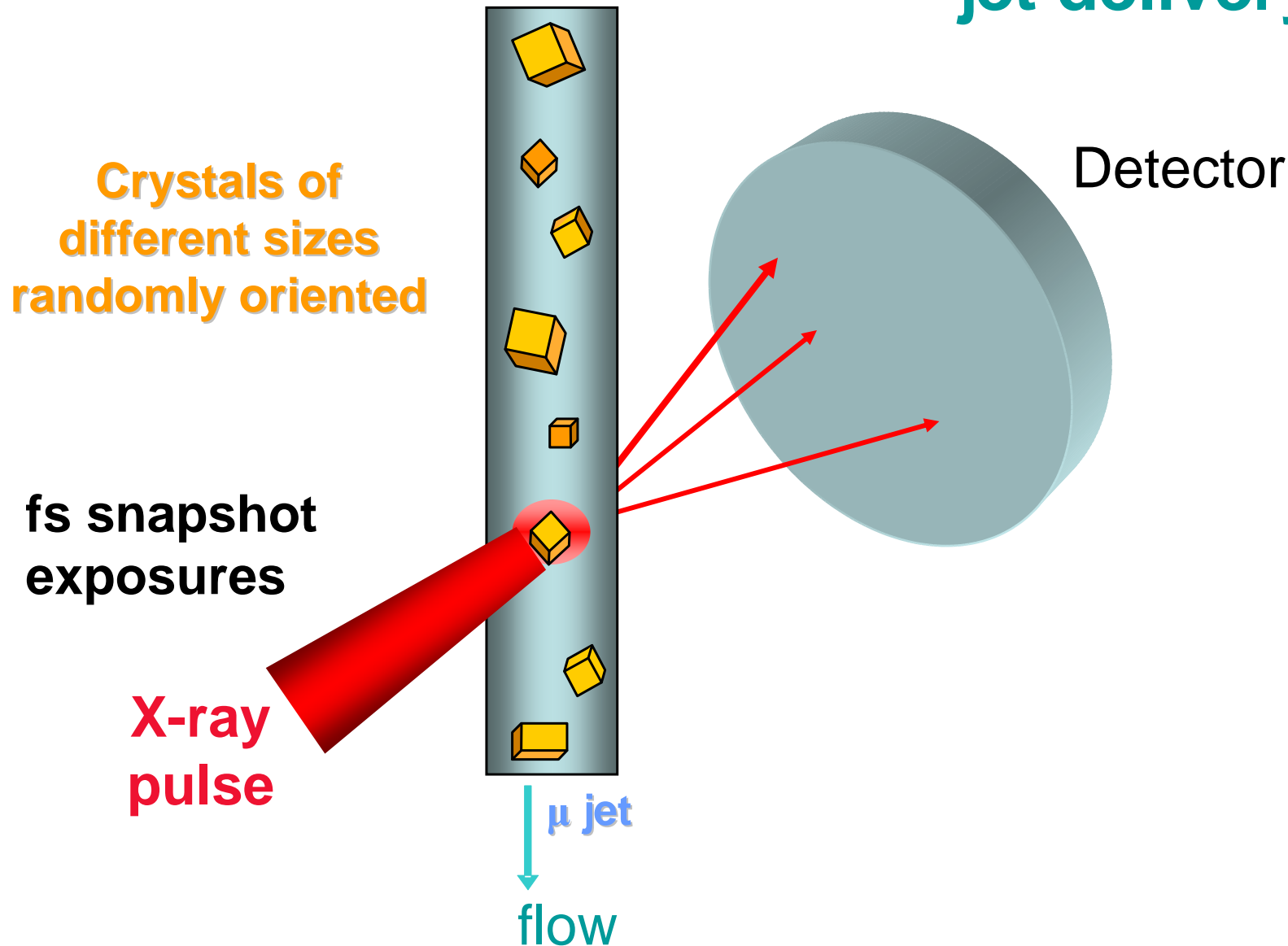
- 10^{12-13} photons \sim 3-500 fs pulses
- repetition rate: 120 Hz
- photon energy: 300 eV-10 keV
- transversally fully coherent

Samples are destroyed upon interaction with the full FEL beam

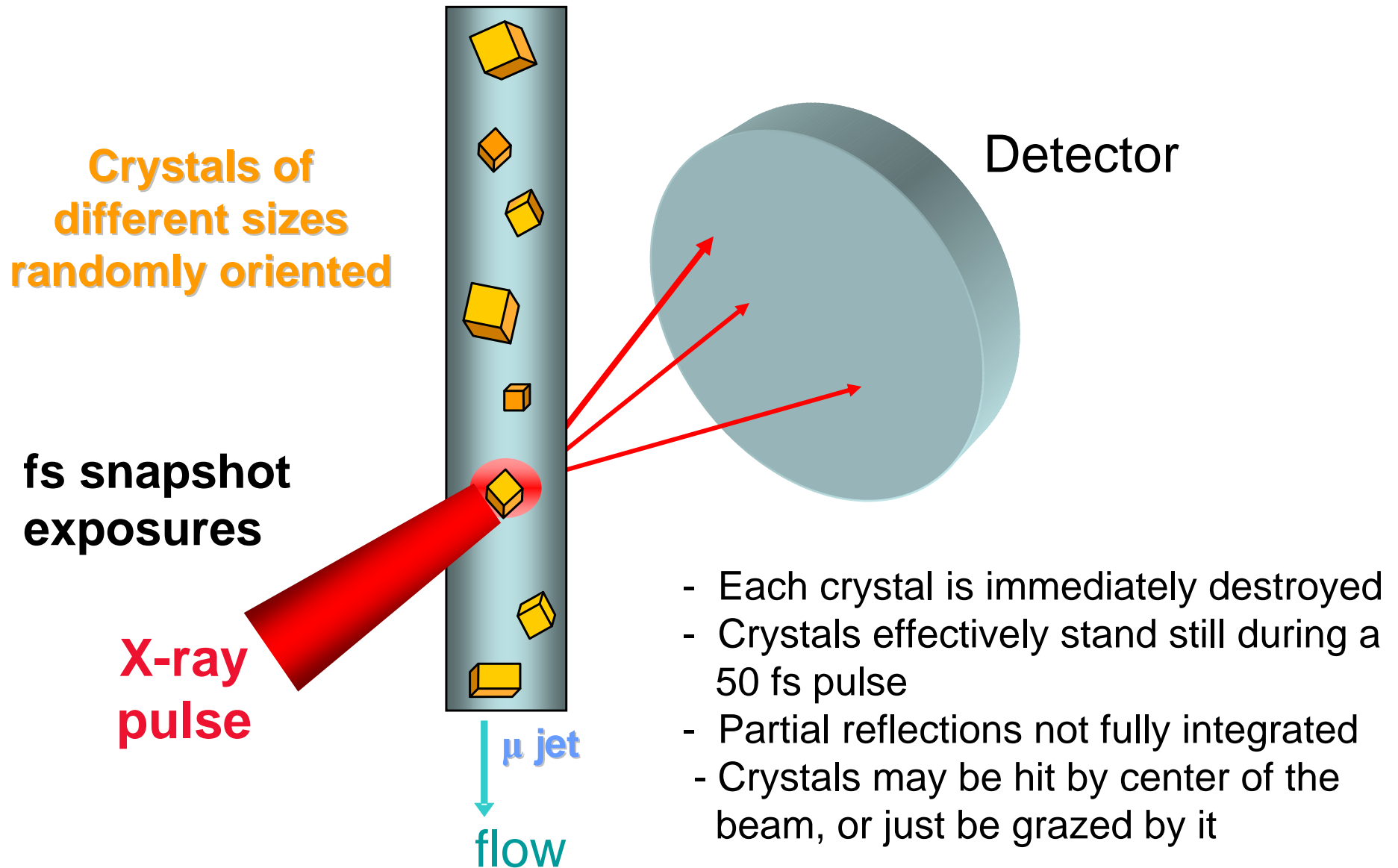


Plasma glow at FEL liquid jet interaction zone

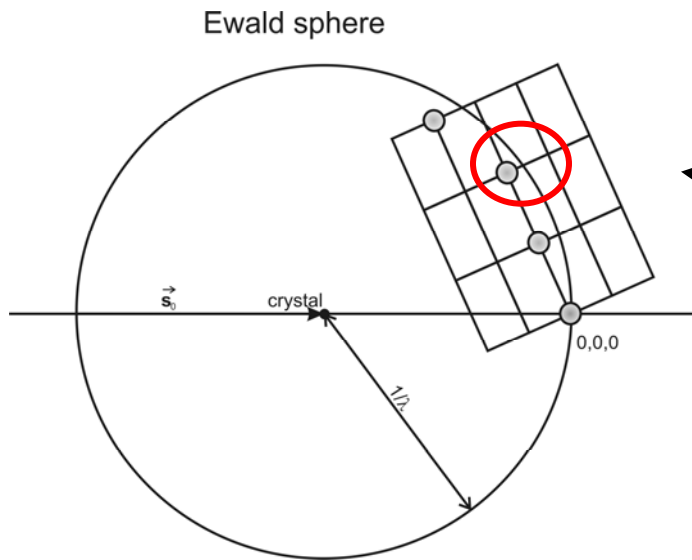
Serial femtosecond crystallography jet delivery



Serial femtosecond crystallography



Indexing and integrating reflections: rotation/oscillation method

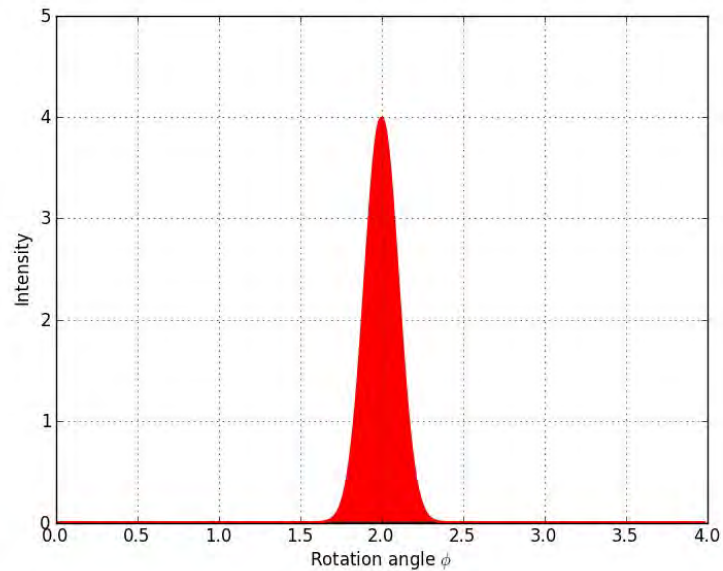


Accurate determination of profile \rightarrow accurate I and $\sigma(I)$

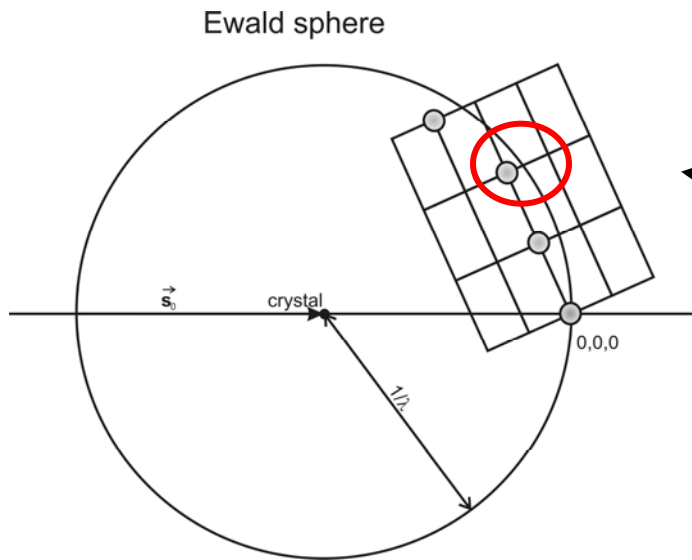
Rotation method

- rotate xtal over finite range
- calculate orientation matrix from observed spot positions

Can fully integrate whole reflections!



Indexing and integrating reflections: rotation/oscillation method



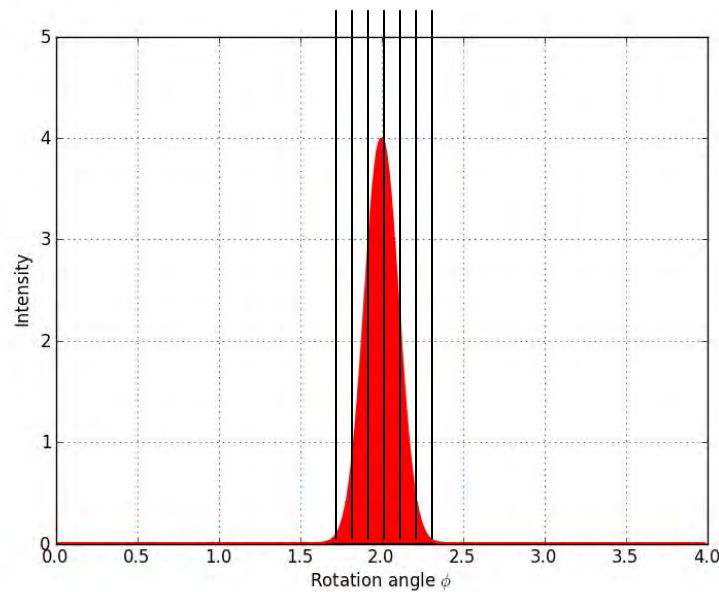
Accurate determination of profile \rightarrow accurate I and $\sigma(I)$

(fine) slicing

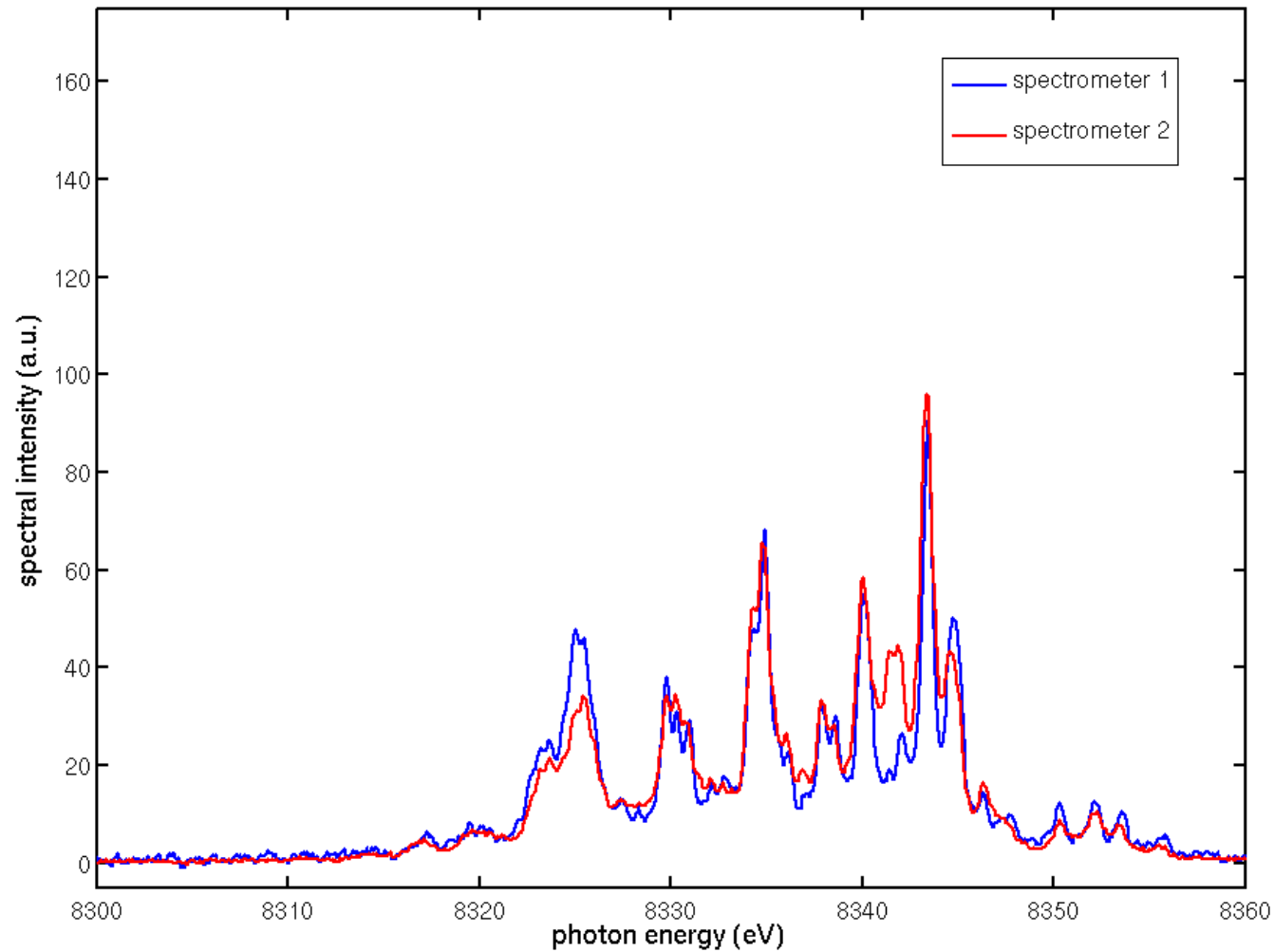
Rotation method

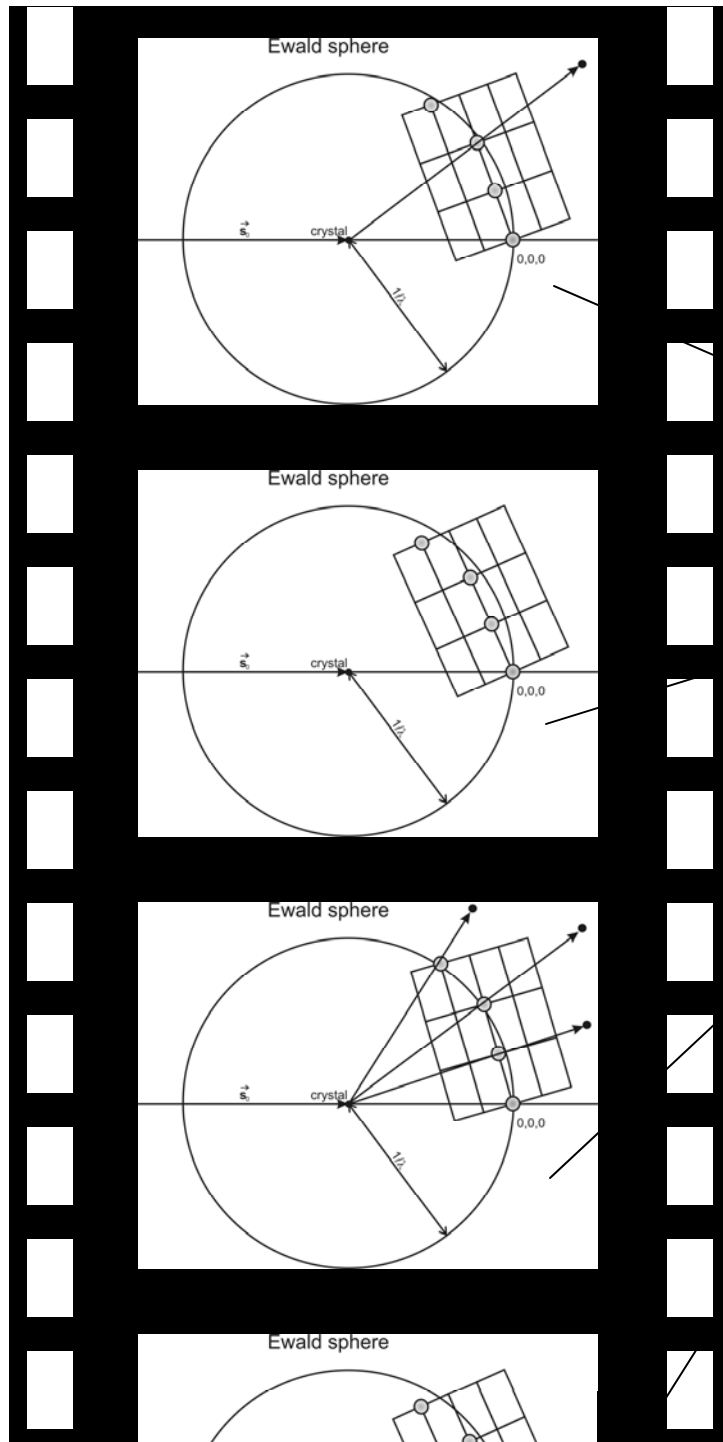
- rotate xtal over finite range
- calculate orientation matrix from observed spot positions

Can fully integrate whole reflections!



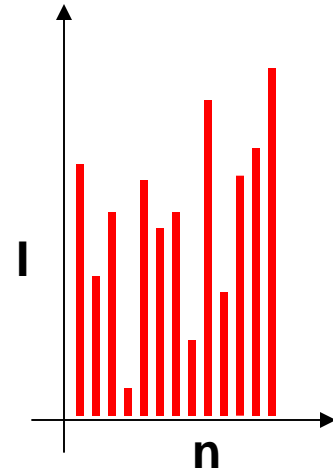
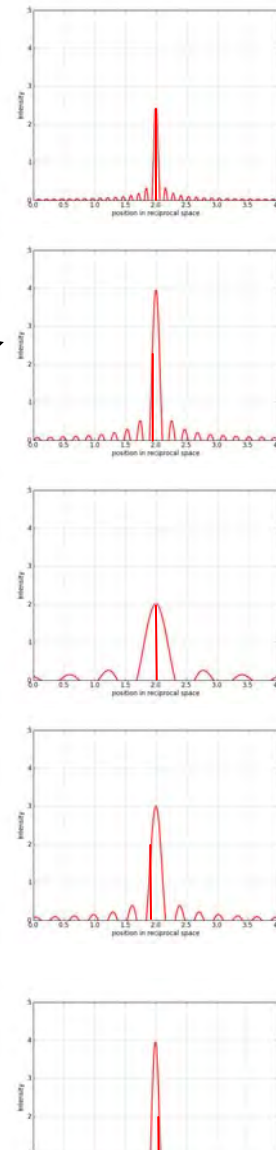
More fluctuations: SASE FEL spectrum varies from shot to shot, too





It is possible to do a *Monte Carlo* integration over multiple *indexed* femtosecond images and obtain a dataset of fully integrated reflections

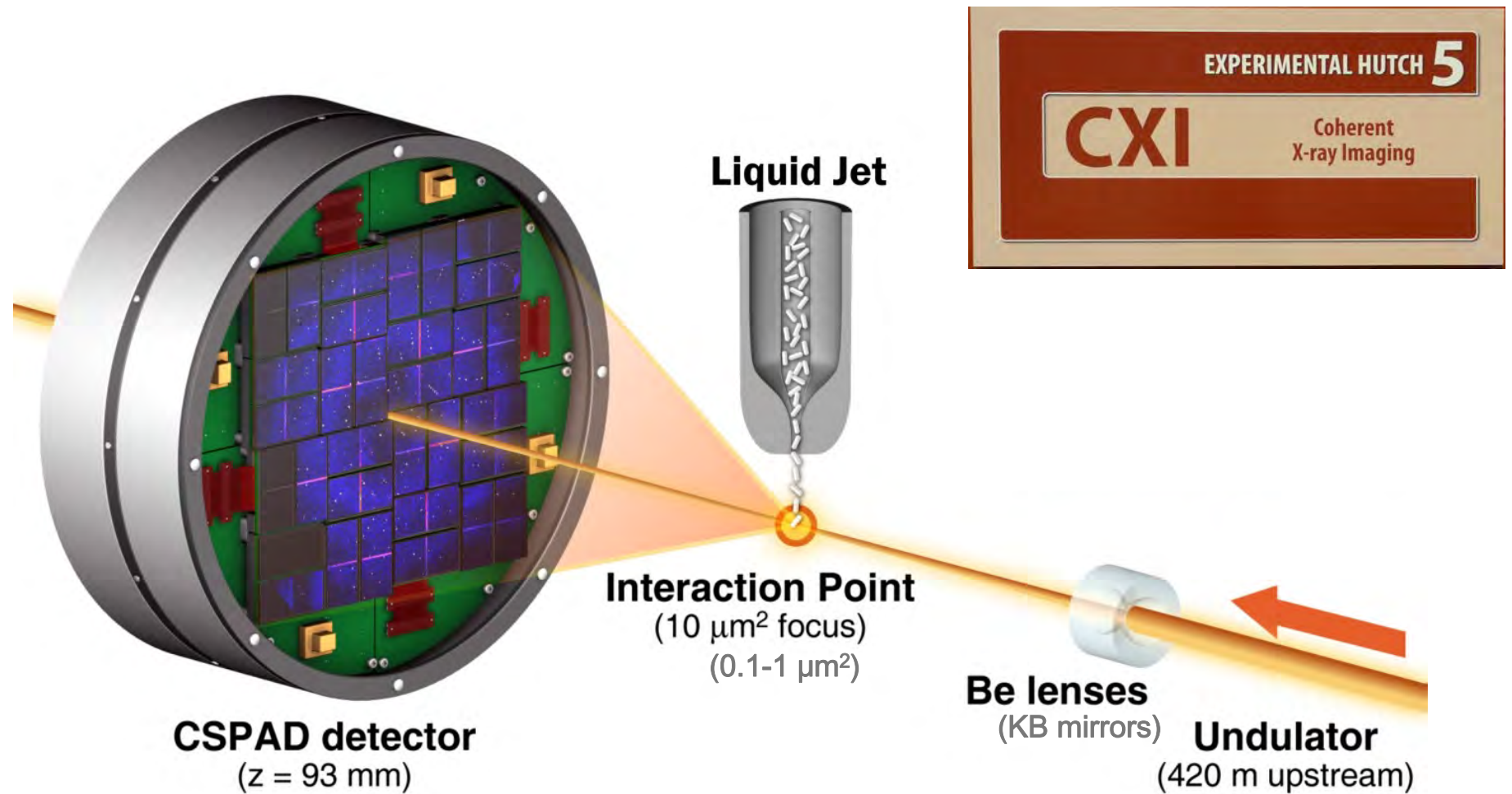
Kirian *et al* (2010), *Optics Express*, **18**, 5713-5723:



$$\langle I[hkl] \rangle = \sum I_n / n$$

...how good are these I? What are their sigma's?

High resolution serial femtosecond crystallography



Cornell-SLAC Pixel Array Detector

Boutet et al Science **337**:362 (2012)

Gas-focused liquid microjets



Daniel DePonte, Uwe Weierstall, John Spence, Bruce Doak



S. Botha, R. Shoeman, R.B. Doak Heidelberg

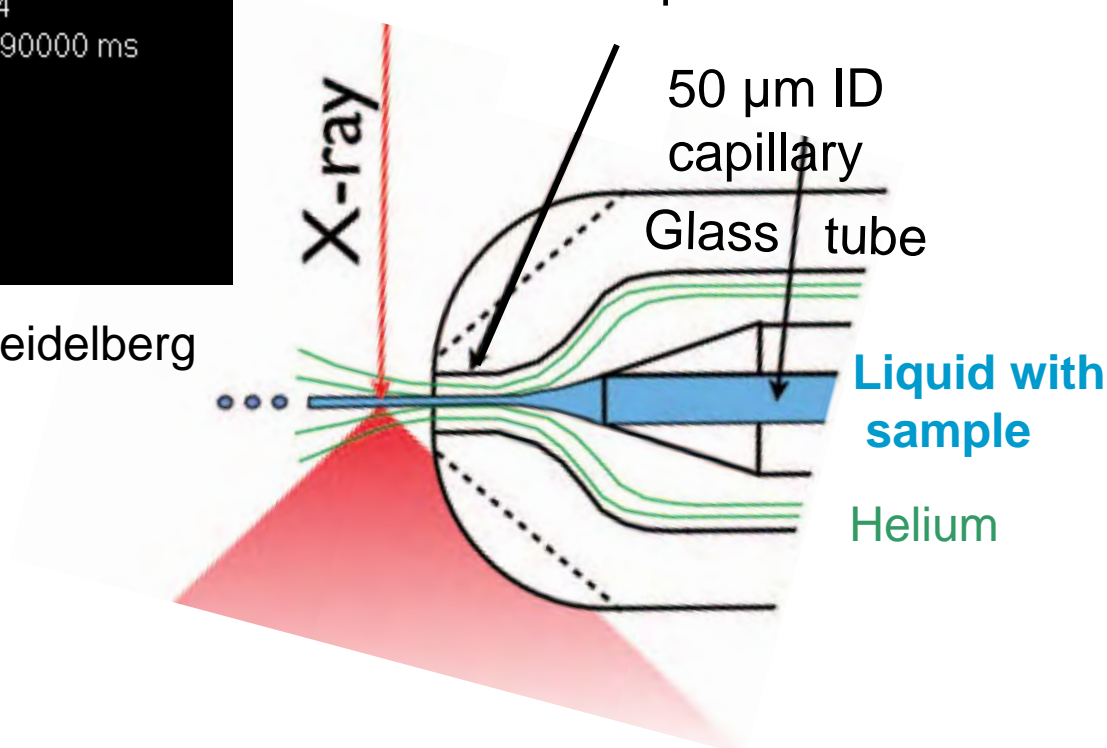
Q=30 μ L/min

P_{Helium}=30 psi

Frame Rate= 300 000 fps

Play Back= 30 fps

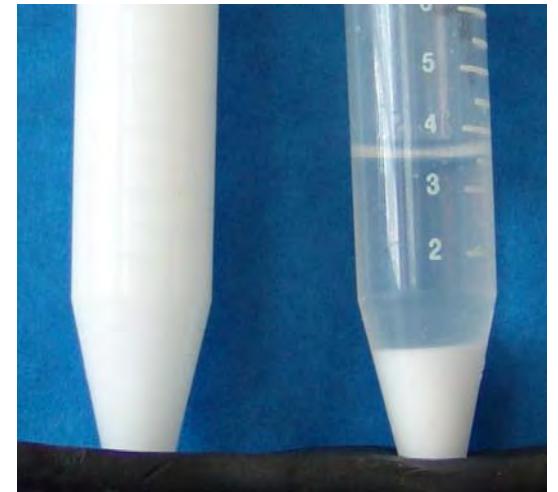
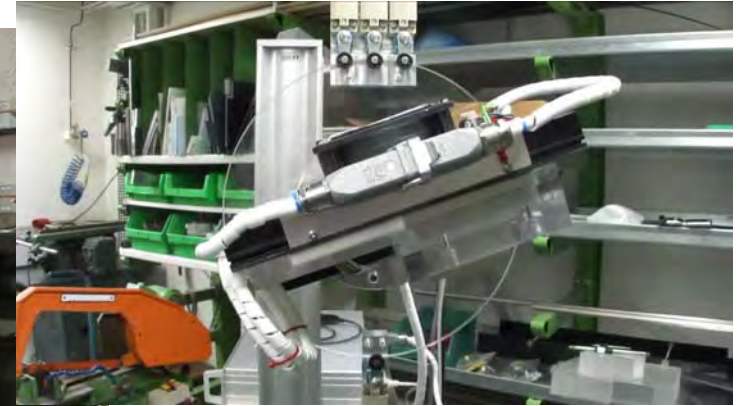
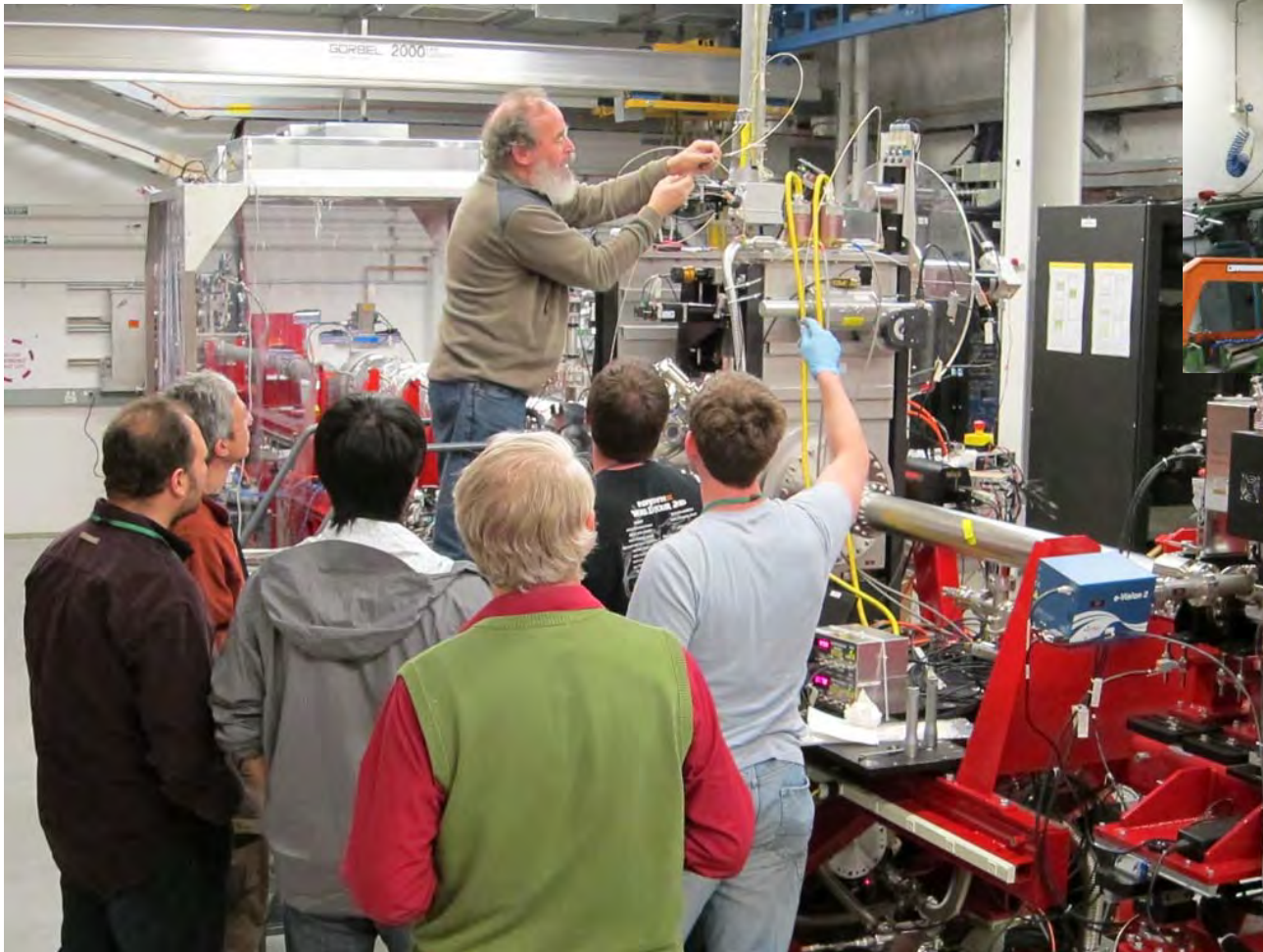
Convergent 50-100 μ m diameter aperture



Gas-dynamic virtual nozzle
Sample at room temperature,
in solution

Weierstall, Spence, Doak, Rev. Sci. Inst **83**, 035108 (2012)

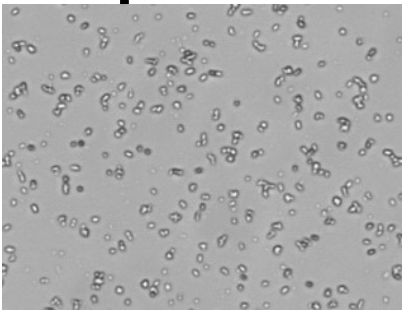
Samples are injected vertically, crystal settling prevented by rotating temperature-controlled syringe pump



Lomb *et al.* J. Appl. Cryst. **45**, 674 (2012).

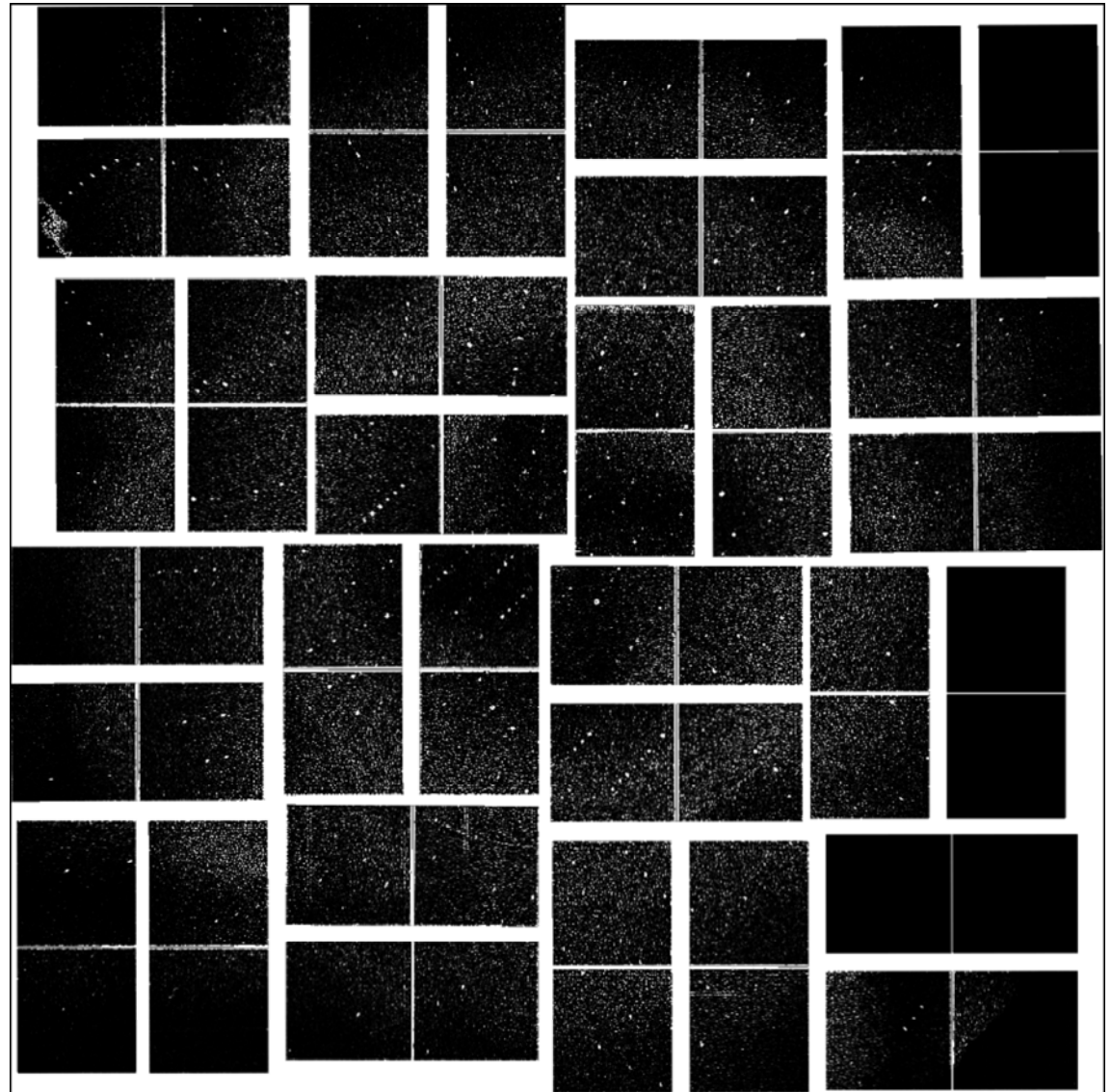
High resolution femtosecond diffraction of micron-sized lysozyme crystals

Lysozyme crystals
1-2 μm \emptyset



40 fs pulse*, 3 mJ/pulse
10 μm^2 focus
Transmission 15%
0.6 mJ/sample
33 MGy/pulse
9.4 keV , $\lambda = 1.32 \text{ \AA}$
Resolution 1.9 \AA

*electron bunch length



Comparison of FEL and synchrotron data

	40 fs	5 fs	Synchrot.
Dose / crystal	33 MGy	3 MGy	0.02 MGy
Dose rate [Gy / s]	8.3×10^{20}	5.8×10^{20}	9.6×10^2
Number of DP	$\sim 1.5 \times 10^6$	$\sim 2 \times 10^6$	100
Hits	66442	40115	100
Indexed DP	12247	10575	100
B-factor [\AA^2]	28.3	28.5	19.4
R/R _{free} [%]	19.2 / 22.09	18.5 / 22.7	16.8 / 20.0

Resolution limit: 1.9 \AA

R-factor vs resolution

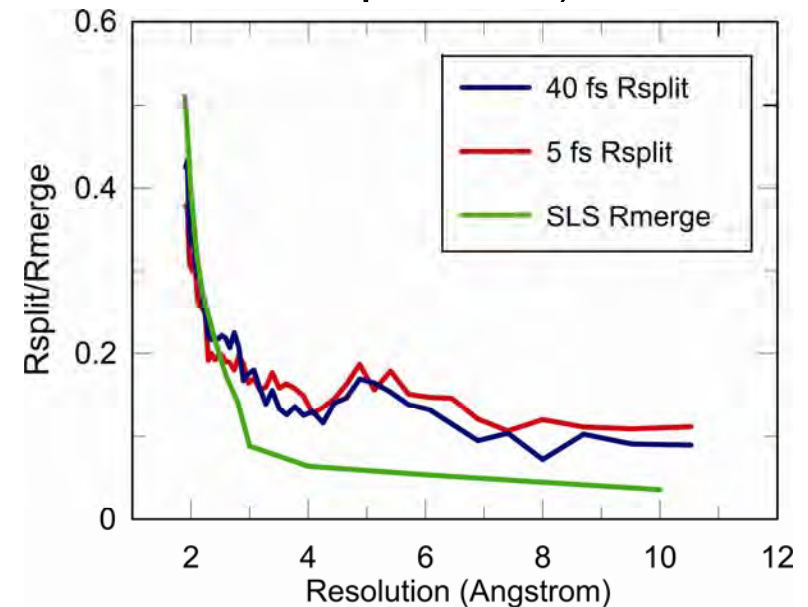
40 fs LCLS data

(1 μm lysozyme crystals)

and

SLS synchrotron data

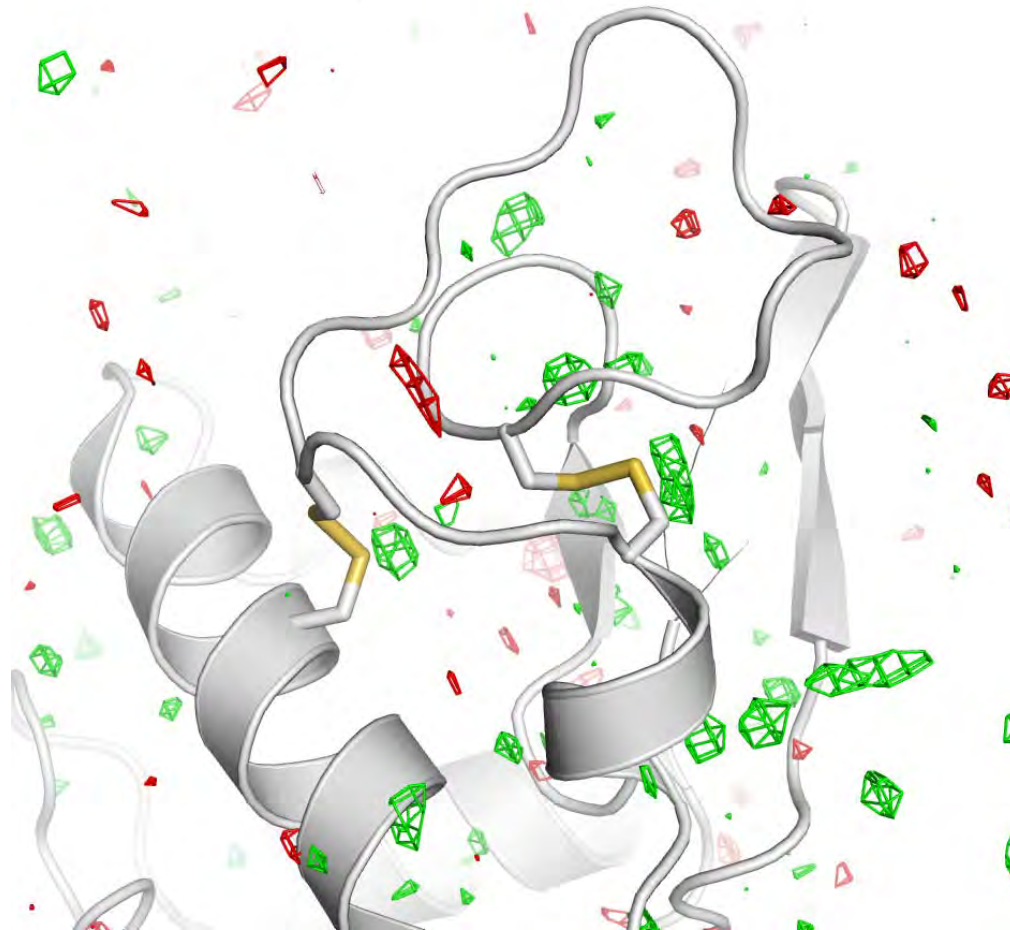
(200 μm lysozyme crystal, room temperature)



Boutet et al Science **337**:362 (2012)

Serial femtosecond crystallography yields undamaged high resolution structures

No difference density Fobs (synchrotron (SLS) – Fobs (LCLS))

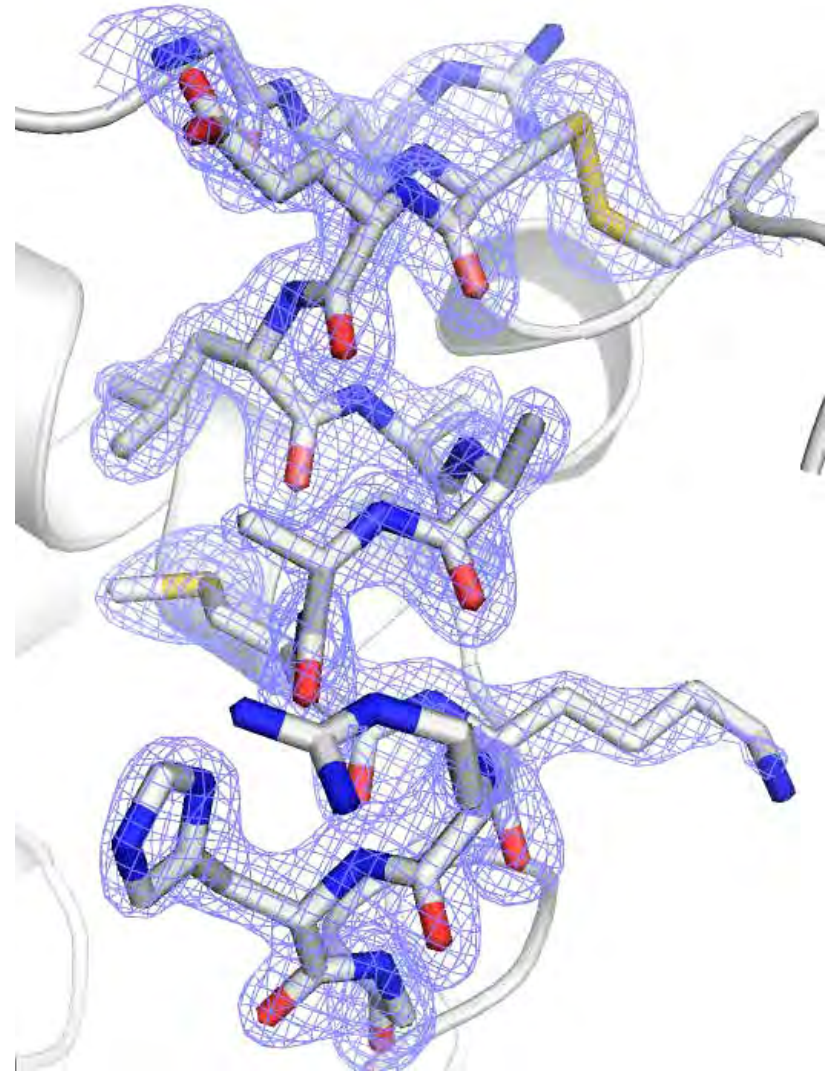


Boutet et al Science **337**:362 (2012)

FEL derived intensities provide high resolution structures

**Molecular replacement-
phased density, 1.9 Å
resolution**

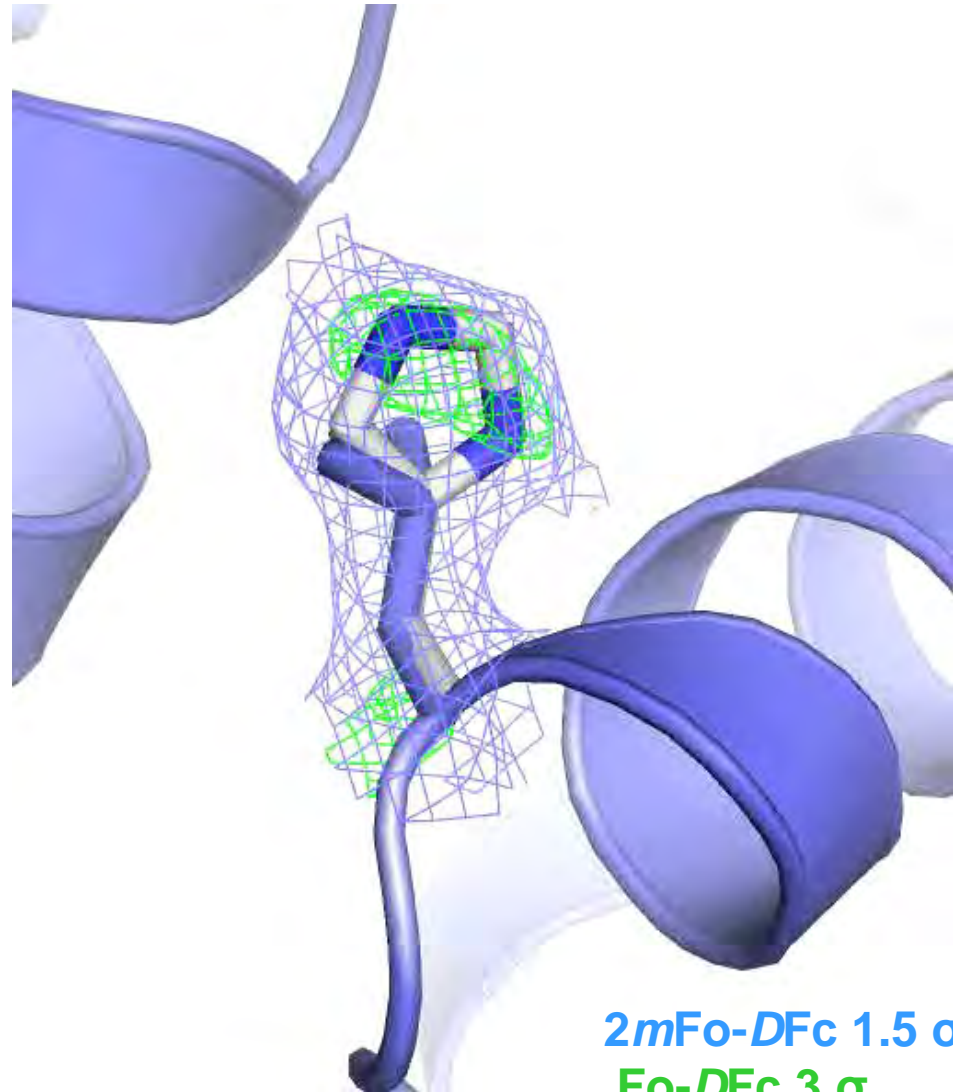
- Resolution better than 2 Å because S-atoms in disulfides can be resolved separately, S-S distance is 2 Å
- Good definition of side chains



1.5 σ 2mFo-DFc

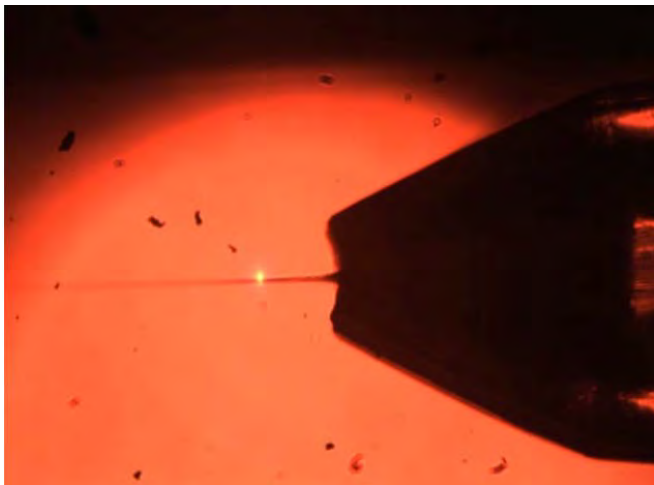
FEL derived intensities are good enough to see small differences

Molecular replacement
with turkey lysozyme
(Valine where there should
be histidine)



Applications of serial femtosecond crystallography

- Analysis of (sub)micron crystals, including membrane proteins in sponge (Nature Meth. **9**: 263 (2012)) or lipidic cubic phase (Science **342**: 1521 (2013))
- SAXS and WAXS measurements
- Time-resolved pump-probe studies on light-sensitive systems



Placement of pump laser beam determines time delay

FEL beam hitting LCP-jet at 1 Hz

PIs: Raymond Stevens, Vadim Cherezov *The Scripps Research Institute*

LCP Jet: Uwe Weierstall, Bruce Doak, John Spence *Arizona State University*

Serial ~~femtosecond~~ crystallography at synchrotrons

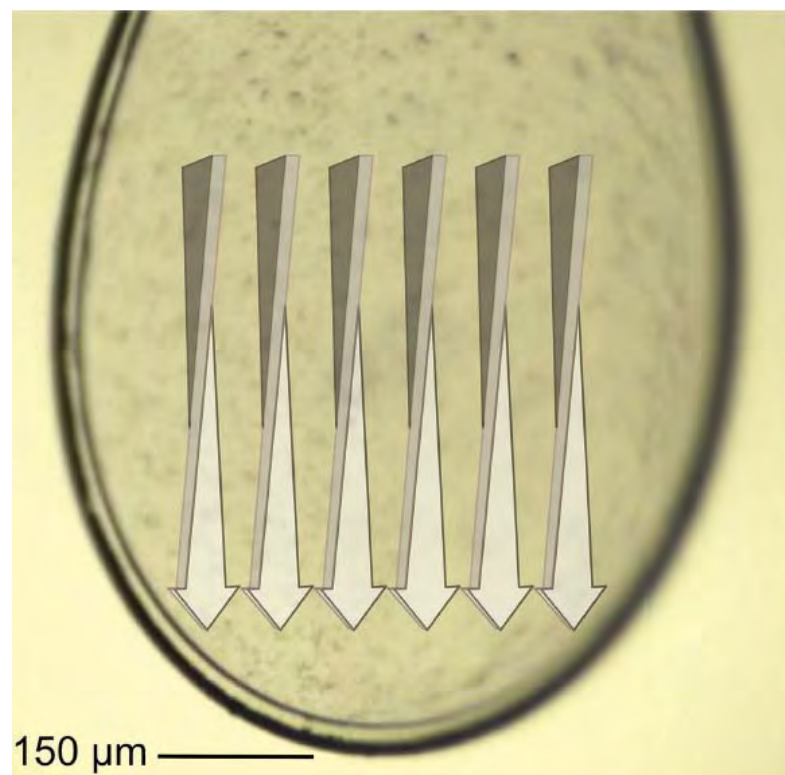
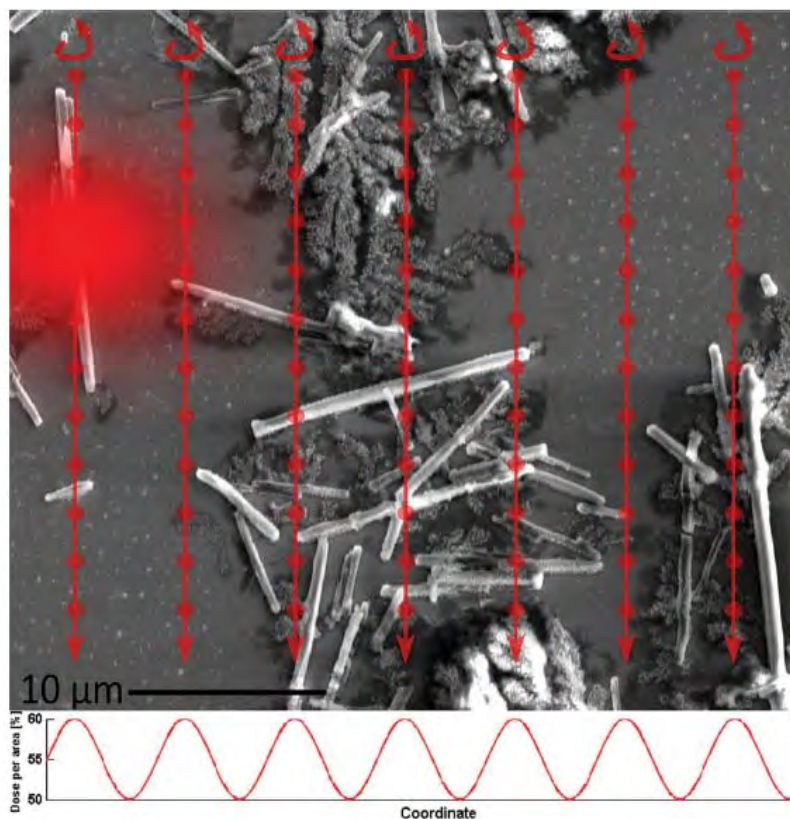
Longer exposure times

- Crystals do not necessarily stand still during exposure
 - Tumbling potential problem
 - Rotation for data collection possible
- Radiation damage

Slurry of cryocooled microcrystals on a loop

Serial crystallography on *in vivo* grown microcrystals using synchrotron radiation

Cornelius Gati,^{a,‡} Gleb Bourenkov,^{b,‡} Marco Klinge,^c Dirk Rehders,^c Francesco Stellato,^a Dominik Oberthür,^{a,d} Oleksandr Yefanov,^a Benjamin P. Sommer,^{d,e} Stefan Mogk,^e Michael Duszynski,^e Christian Betzel,^d Thomas R. Schneider,^{b,*} Henry N. Chapman^{a,f,*} and Lars Redecke^{c,*}



IUCRJ, 2014

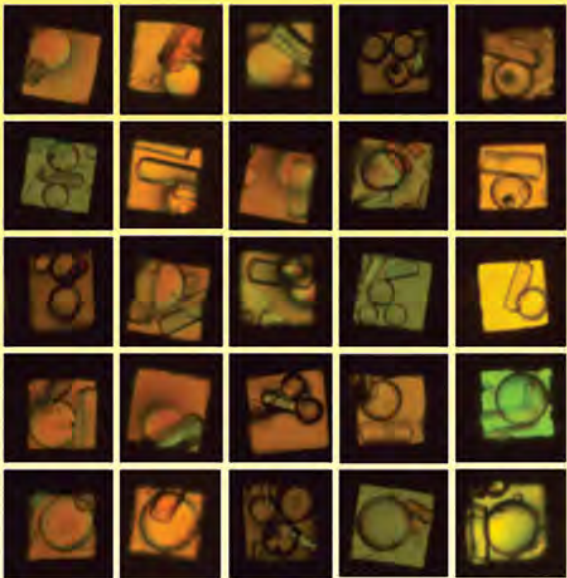
High throughput crystallography at room temperature

ISSN 0907-4449

Volume 68
Part 3
March 2012

Cytoplasmic domain of *Yersinia pestis* YscD
Epithelial adhesin 1 from human pathogen *C. glabrata*
Plant S-adenosyl-L-homocysteine hydrolase complexes
DNA (6-4) photoproduct dTT(6-4)TT in complex with Fab fragment
B. cereus adenosine phosphorylase
S. maltophilia 5'-deoxy-5'-methylthioadenosine phosphorylase II
Catechol-O-methyltransferase inhibitor complex
Grid-enabled web service for low-resolution refinement
Effects of cryoprotectants on human blood group A and B glycosyl transferases
CYP108D1 from *N. aromaticivorans* DSM12444
Stability of *Clostridium thermocellum* enzymes
3-Isopropylmalate dehydrogenase under high pressure

Acta Crystallographica Section D
**Biological
Crystallography**
Editors: E. N. Baker and Z. Dauter



Crystallography on a chip, p. 321

journals.iucr.org
International Union of Crystallography
Wiley-Blackwell

Chip mounts

- control over solution, all crystals can be handled (after appropriate surface chemistry is established)
- with or without bottoms
- diffraction images +/- laser for time resolved studies
- 100% hit rates, chip can be pre-scanned for occupied positions

Practical aspects of the chip

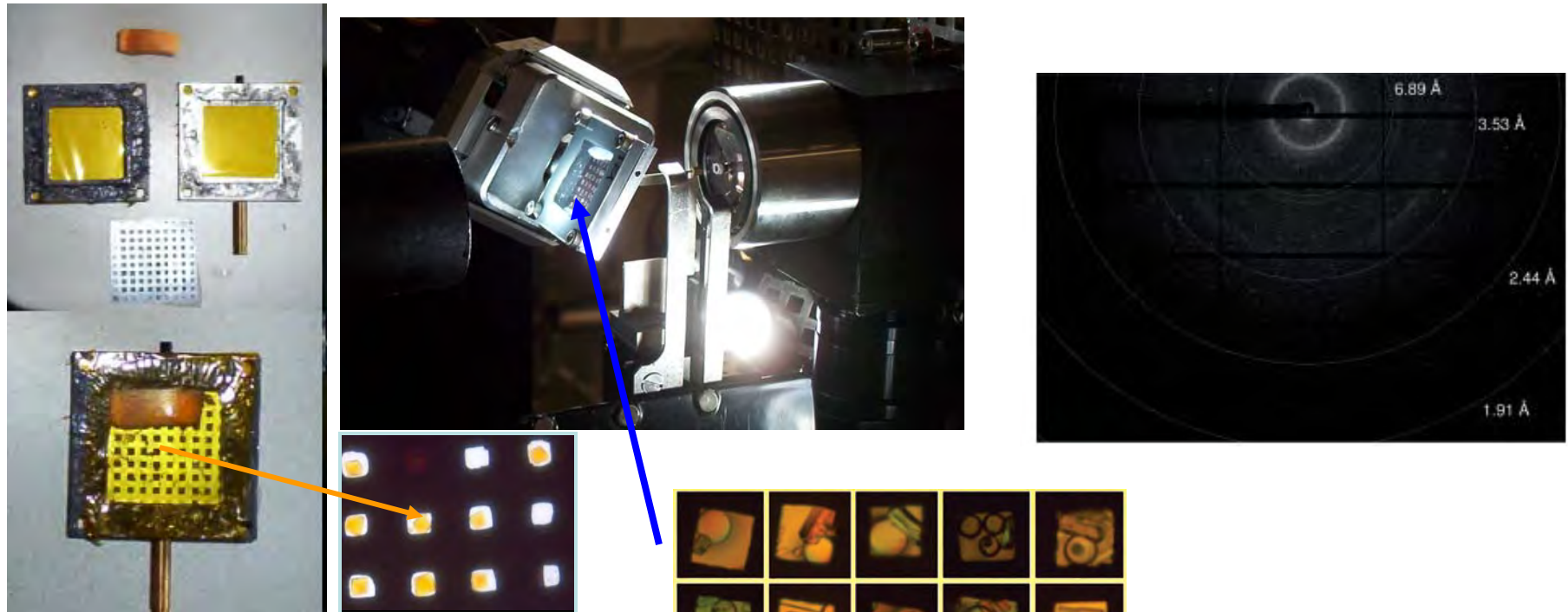
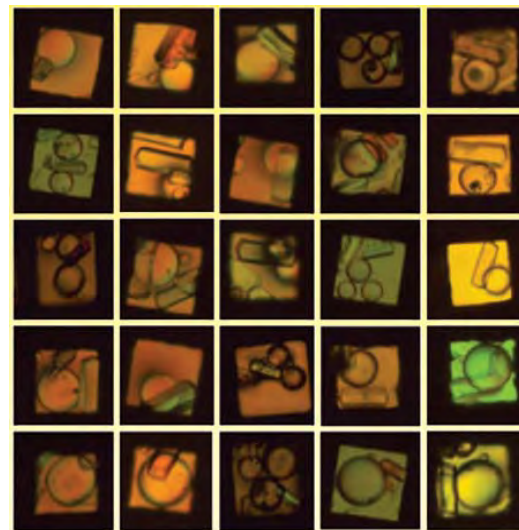
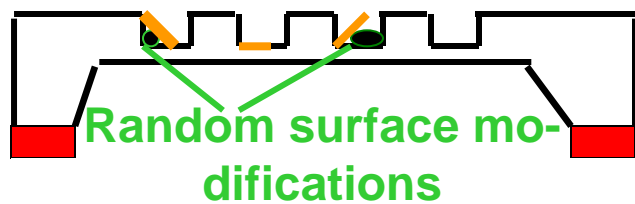


Plate like crystals with random orientations



Surface roughing of well-bottoms can induce random xtal orientations

Zarrine-Afsar et al Acta Cryst. D 68: 321-3 (2012)

Room temperature serial crystallography at the SLS using a high viscosity extrusion injector at atmospheric pressure

Data removed

Collection of still and rotation data possible

Different high viscosity media possible

accounting for required flow rate, crystal preference

**Room temperature serial crystallography at the Swiss Light Source
using a high viscosity extrusion injector at atmospheric pressure**

Data removed

Serial crystallography

Serial collection of partial datasets of **many many different crystals**

- mitigate the effects of **radiation damage**
- **time-resolved** measurements

in high throughput fashion

to provide fast sample change to account for short lifetime of sample

ID29 (Phase II) 10^{14} photons/s/ μm^2 Henderson limit 200 μs
XFELs 10^{12} photons/pulse, few fs exposures

Background

Radiation damage (mechanisms)

Jet@SLS, LCLS

Sabine Botha,
Thomas Barends
Karol Nass
Robert Shoeman
R. Bruce Doak
Lutz Foucar
Wolfgang Kabsch



Chip@SLS

Asrash Zarrine-Afsar
Thomas Barends
Christina Mueller
Lukas Lomb
RJ Dwayne Miller (MPI SD)

Florian Dworkowski
Meitian Wang



Martin Fuchs

LCLS

Henry Chapman and group, in particular Thomas White, DESY
John Spence, Uwe Weierstall and group, ASU
Petra Fromme and group, ASU

Sebastien Boutet, Marc Messerschmidt, Garth Williams

