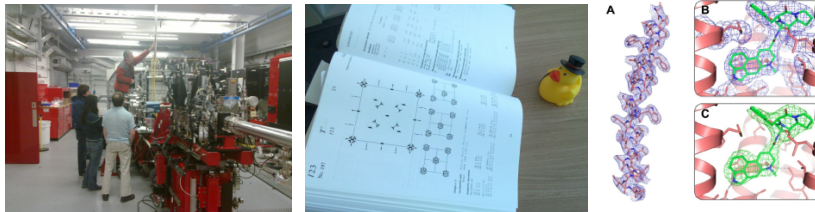


Crystallography and molecular imaging using X-ray lasers.

Fundamentals and applications

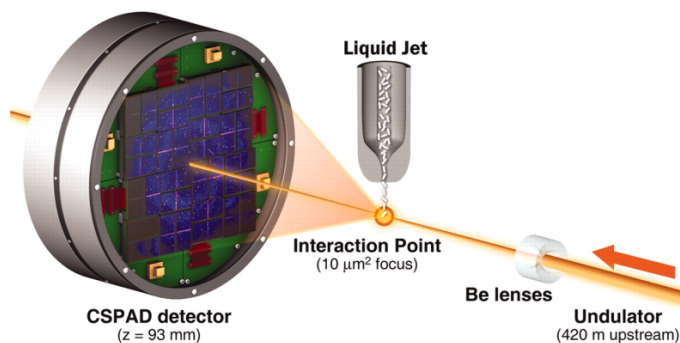


Thomas A. White
CERN Accelerator School
Hotel Scandic Emporio, Hamburg
9th June 2016



What is this all about?

- > Study the **structure and function of proteins** ...
- > ... using a **hard X-ray free-electron laser** ...
- > ... by using **many small crystals** of the protein.



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What is a protein?

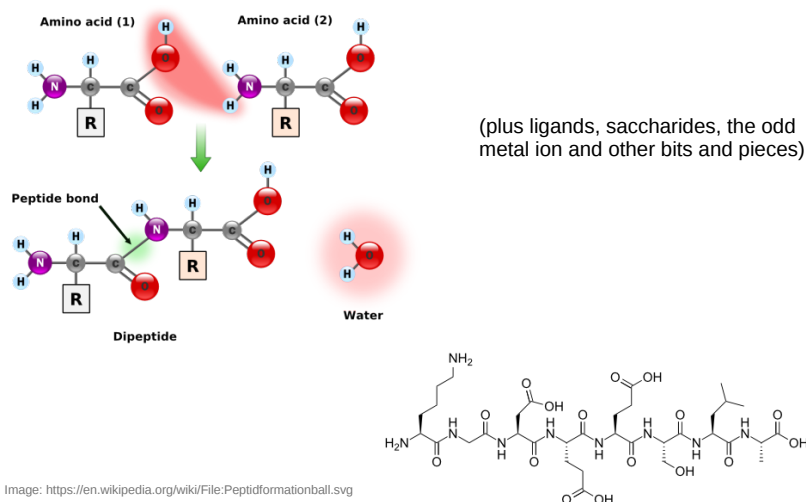


Image: <https://en.wikipedia.org/wiki/File:Peptidformationball.svg>

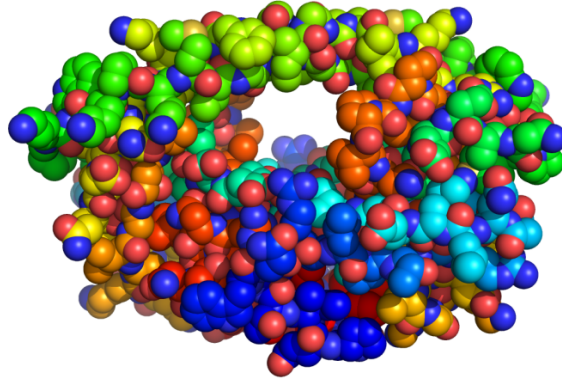


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Why study the structure of proteins?

- > The function of proteins is determined by their structure, not elemental composition.
- > All proteins have the same empirical formula (ratio of elements).



More? <http://pdb.rcsb.org/motm/6>

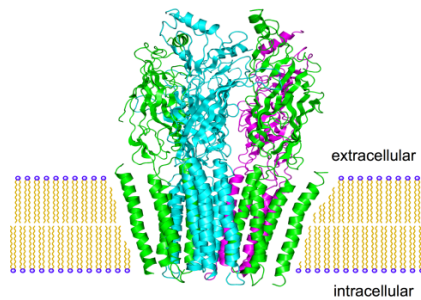
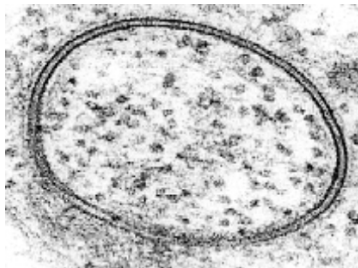


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Why study the structure of proteins?

- > Membrane proteins are particularly difficult to study.



Nicotinic acetylcholine receptor (nAChR)

Images: https://en.wikipedia.org/wiki/File:Annular_Gap_Junction_Vesicle.jpg
https://en.wikipedia.org/wiki/File:NACHR_2BG9.png

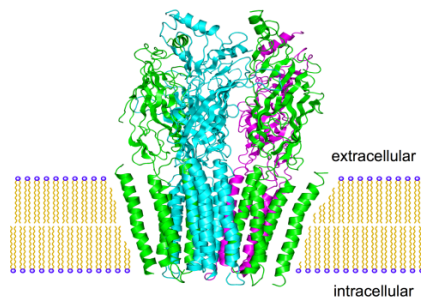


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Why study the structure of proteins?

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Nicotinic acetylcholine receptor (nAChR)

Image: https://en.wikipedia.org/wiki/File:NACHR_2BG9.png

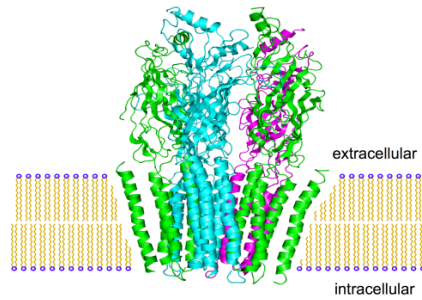
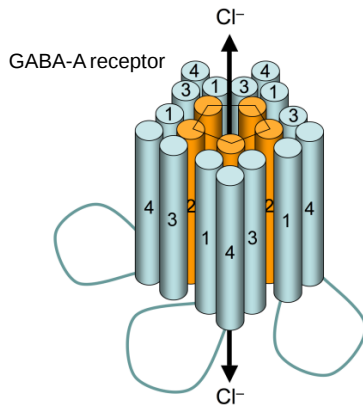


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Why study the structure of proteins?

- > Membrane proteins are particularly difficult to study.



Nicotinic acetylcholine receptor (nAChR)

Images: https://en.wikipedia.org/wiki/File:NachR_2BG9.png
https://en.wikipedia.org/wiki/File:NachR_2BG9.png



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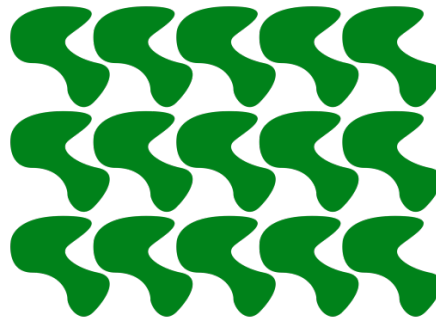


Studying proteins using X-rays

- > What is a crystal?



A molecule of "blob"



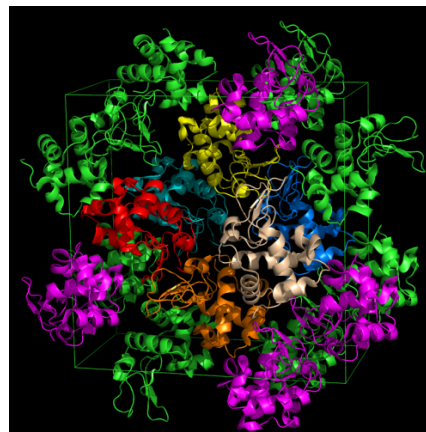
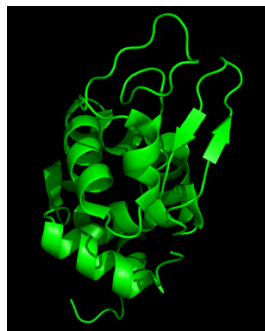
A crystal of "blob"



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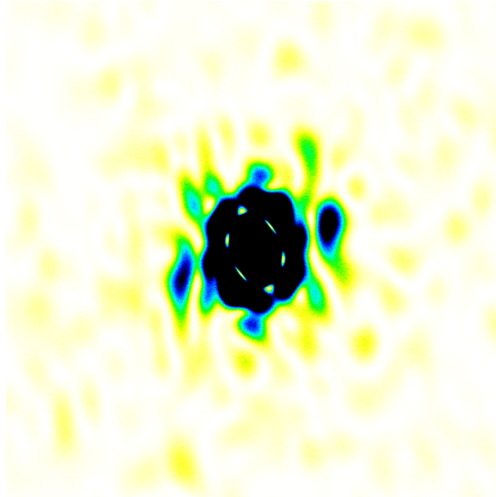
Studying proteins using X-rays



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Studying proteins using X-rays



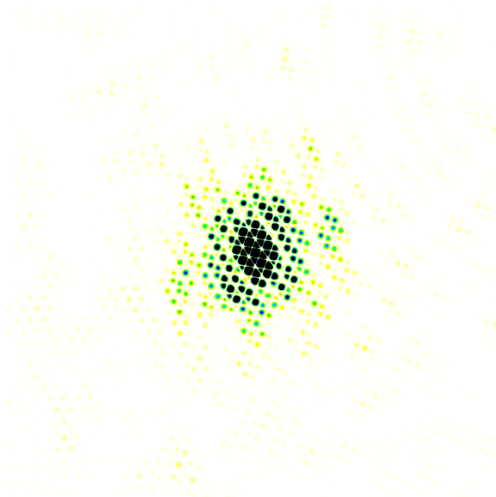
1 molecule



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Studying proteins using X-rays



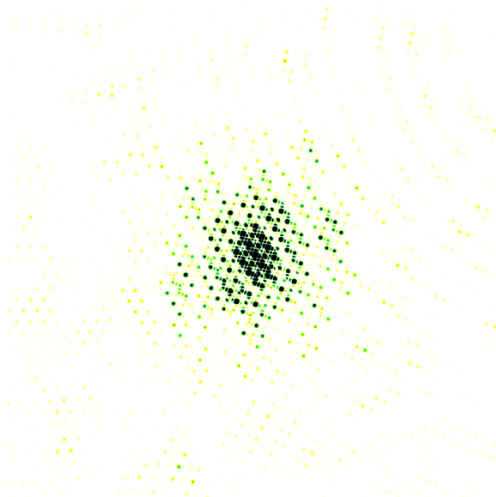
2x2x2 lattice



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Studying proteins using X-rays



3x3x3 lattice



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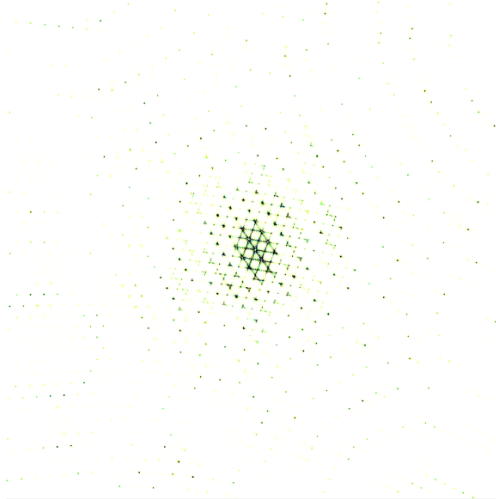


Studying proteins using X-rays

10x10x10
lattice



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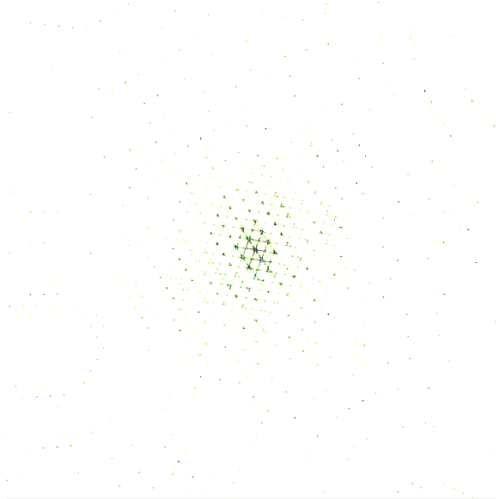


Studying proteins using X-rays

20x20x20
lattice



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Studying proteins using X-rays

1 molecule



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Studying proteins using X-rays

- > We would prefer to use single molecules rather than crystals, but the signal level is very low:

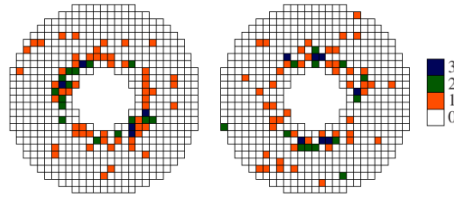


FIG. 1. (Color online) The same or different? Two simulated measurements (noisy diffraction patterns) in a single-particle imaging experiment, where color (white, orange, green, and blue) represents recorded photon counts (0, 1, 2, and 3). Are the differences in the measurements purely statistical, or do they reflect a different view (orientation) of the particle?

→ Loh and Elser, Phys. Rev. E 80 (2009) 026705

Studying proteins using X-rays

1 molecule

Studying proteins using X-rays

2x2x2 lattice

Studying proteins using X-rays

3x3x3 lattice



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Studying proteins using X-rays

10x10x10
lattice



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Studying proteins using X-rays

20x20x20
lattice

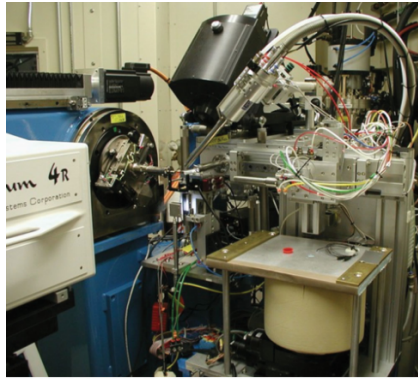


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Crystallographic data collection

- > Goniometer, one crystal (or a few crystals) for entire dataset



© Garland Science 2010



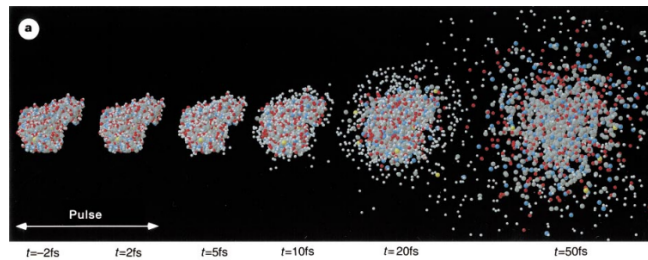
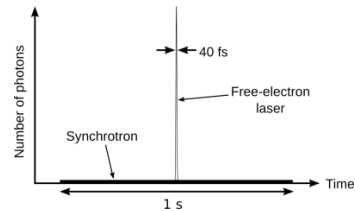
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Why use a free-electron laser?

Very short, very intense pulses of X-rays mean we can "side-step" radiation damage.

R. Neutze et al., "Potential for biomolecular imaging with femtosecond X-ray pulses", Nature 406 (2000) p752.

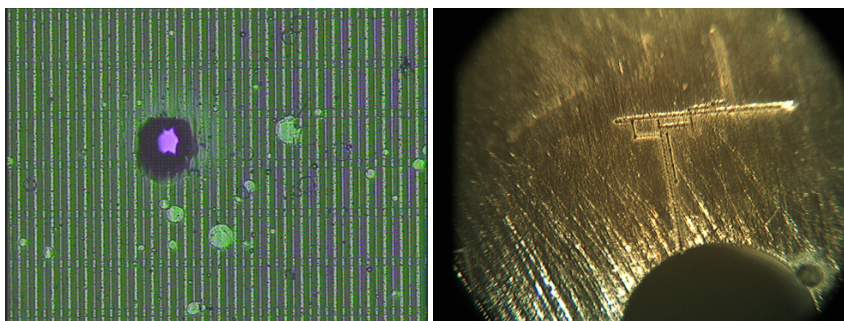


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Radiation damage

- > X-rays damage crystals
(... and the detector, and the shroud, and ...)



G. Weidenspointner et al., Proc. SPIE (2011) 80780U

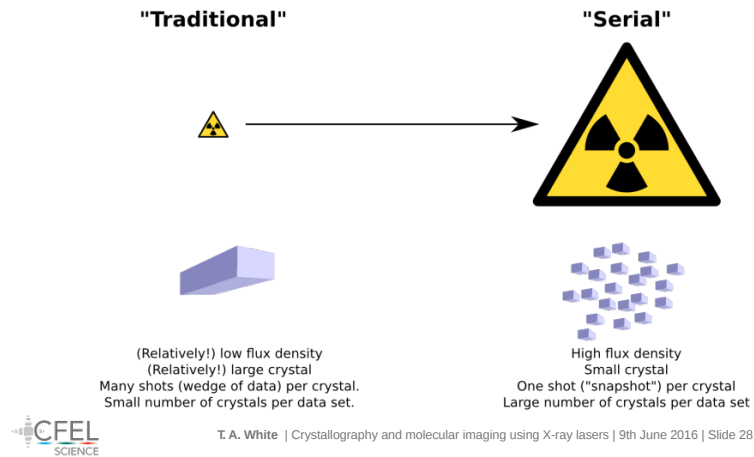


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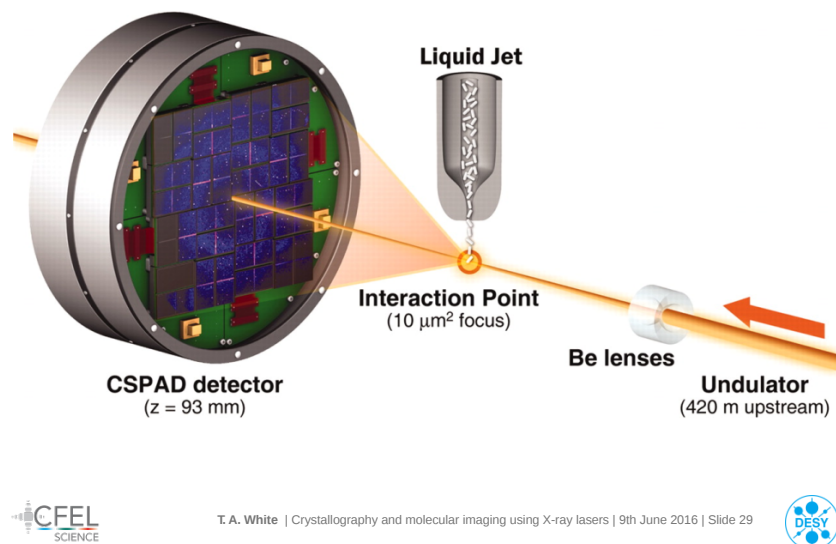


Multi-crystal data collection

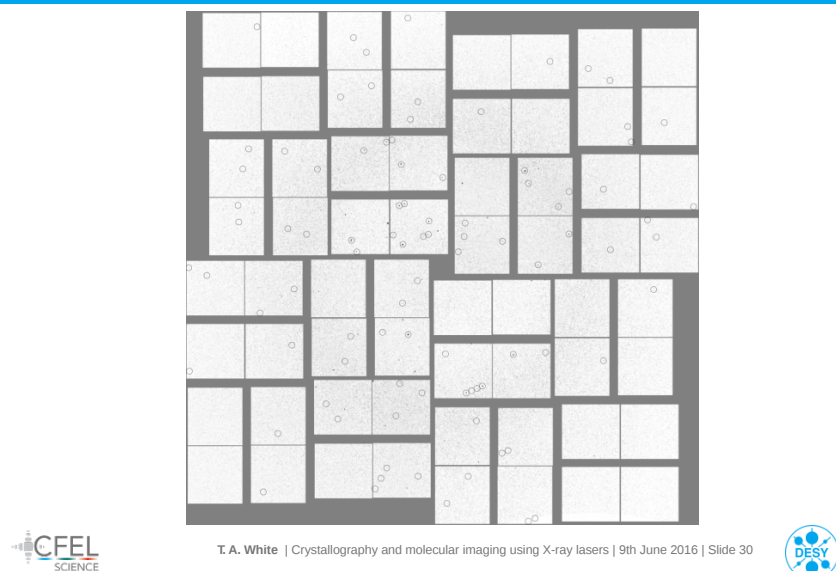
- > We want to study more and more difficult proteins.
- > Size of crystals down, intensity up, damage up, number of crystals up.
- > Limit: one diffraction pattern per crystal: "serial crystallography".



Serial crystallography in practice

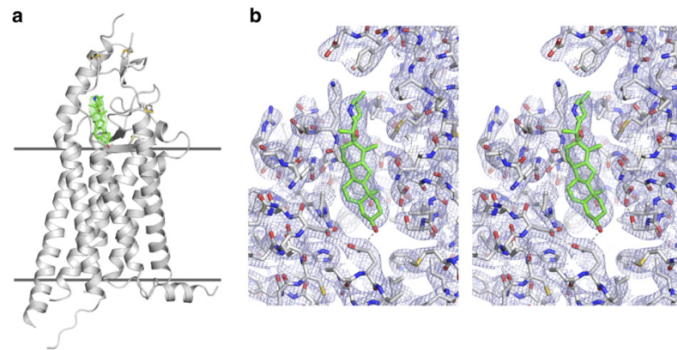


Serial crystallography in practice



Example: Human receptors

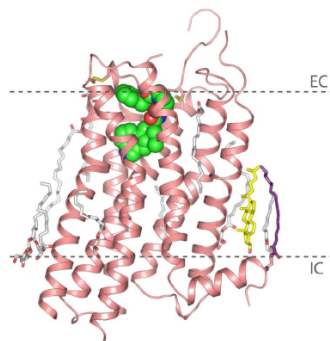
> Weierstall et al., Nature Communications 5:3309 (2014)



"Smoothened" receptor

Example: Human receptors

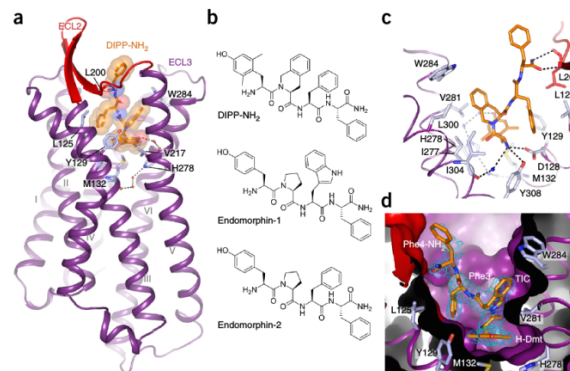
> Liu et al., Science 342 (2013) p1521



5HT2B receptor

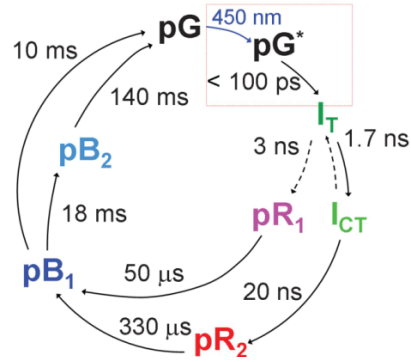
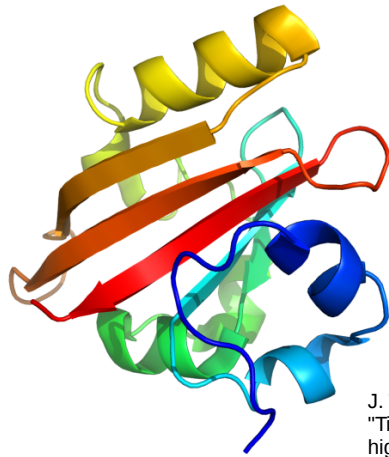
Example: Human receptors

> Fenalti et al., Nature Structural and Molecular Biology 22 (2015) p265



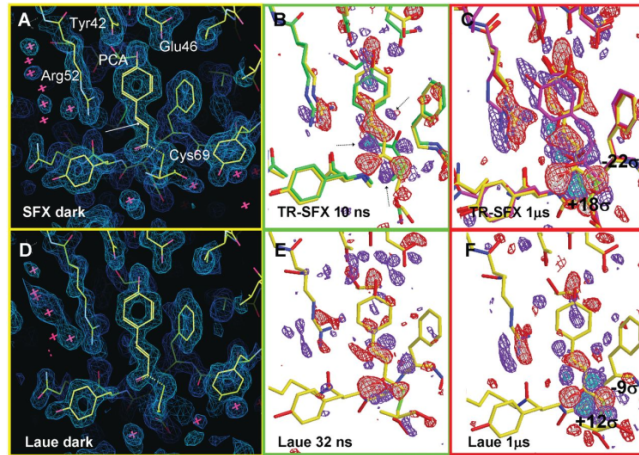
Delta-opioid receptor

Photoactive Yellow Protein

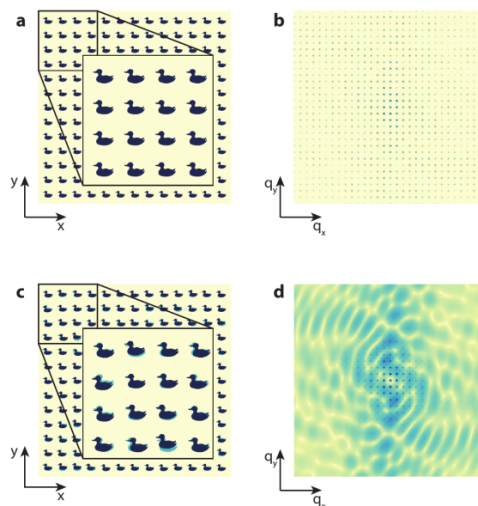


J. Tenboer, S. Basu, N. Zatsepin, K. Pande et al. "Time-resolved serial crystallography captures high-resolution intermediates of photoactive yellow protein". Science 346 (2014) p1242

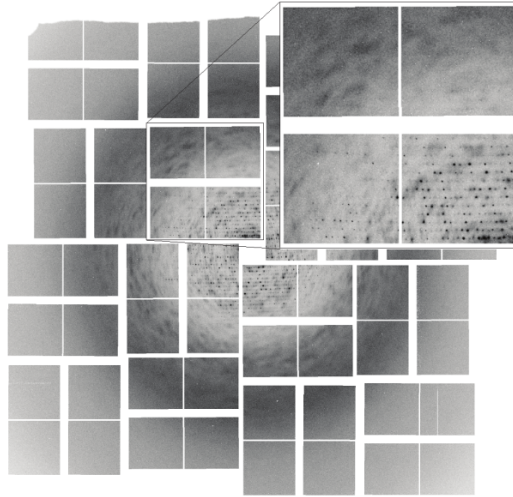
Photoactive Yellow Protein



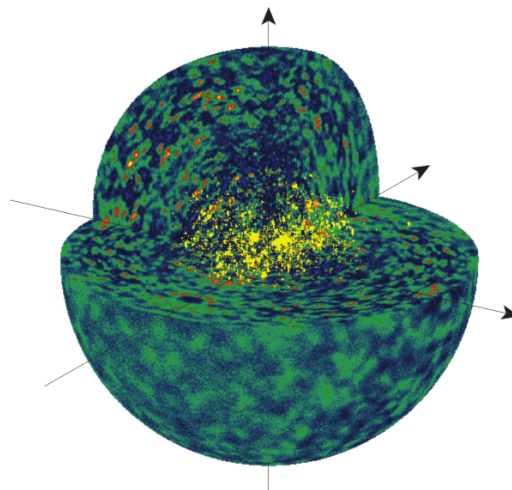
Molecular Imaging



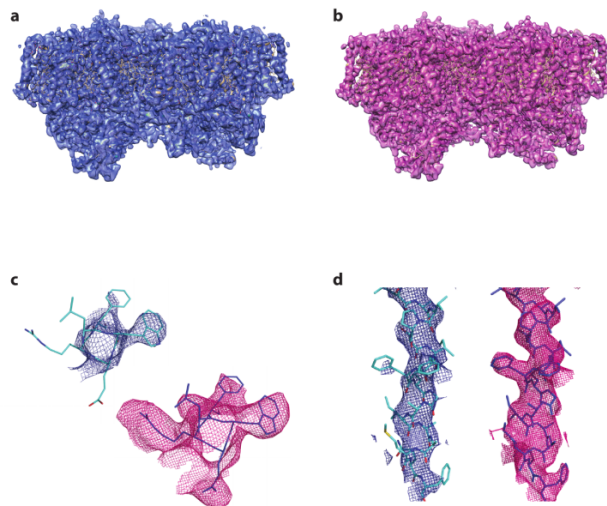
Molecular Imaging



Molecular Imaging



Molecular Imaging



Useful references

First FEL crystallography experiment:

H. N. Chapman, P. Fromme et al., Nature 470 (2011) p73

Liquid jet sample injector ("Gas dynamic virtual nozzle"):

D. P. DePonte, U. Weierstall et al., J. Phys. D: Appl. Phys. 41 (2008) 195505

Lipidic cubic phase extrusion injector:

U. Weierstall, D. James, C. Wang et al., Nature Communications 5:3309 (2014)

Molecular imaging using imperfect crystals:

K. Ayyer, O. M. Yefanov, D. Oberthür et al., Nature 550 (2016) p202

Loads more references on CrystFEL website: <https://www.desy.de/~twhite/crystfel>

Email me: taw@physics.org