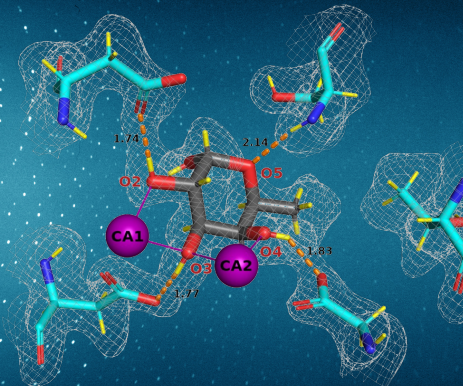
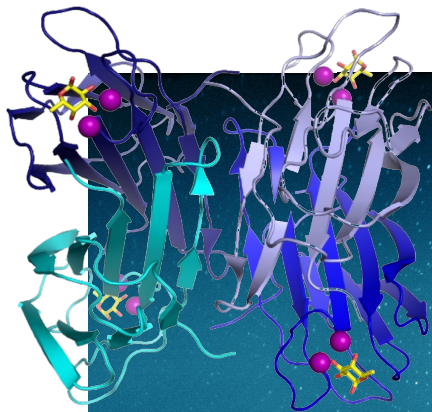


Seeing the chemistry in biology using neutron crystallography



Matthew Blakeley

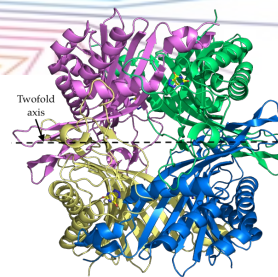
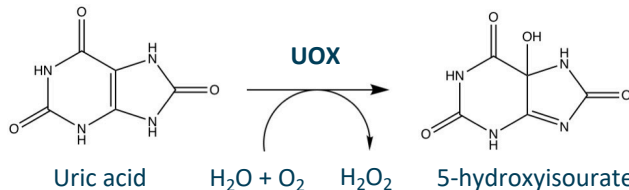
LADI-DALI instrument scientist

Large-Scale Structures Group, Institut Laue-Langevin

Urate oxidase enzyme mechanism studies

PhD project of Lindsay McGregor, ILL/King's College London (R. Steiner)

- **Urate oxidase (UOX)** is the archetypal co-factor free enzyme, catalyzing the reaction of uric acid to 5-hydroxyisourate.
- Neutron diffraction studies of UOX complexes can help determine the **exact catalytic pathway**.

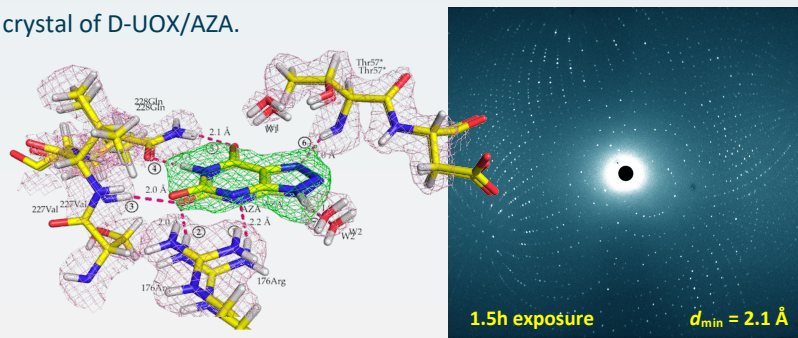
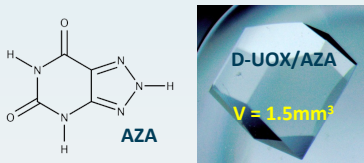


UOX tetramer (81, 96, 105 Å)

Perdeuterated UOX in complex with 8-azaxanthine (AZA).

McGregor *et al.*, (2021) *IUCr* **8**, 46-59.

LADI: Neutron data to 2.1 Å (>80%) at RT in 16.5h using a 1.5mm³ crystal of D-UOX/AZA.

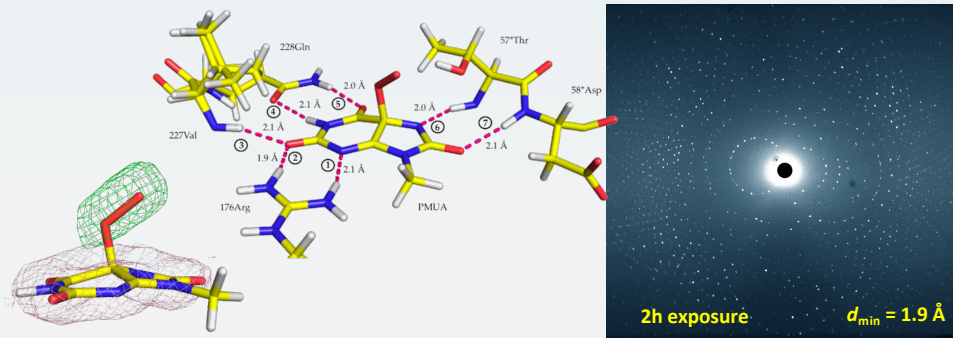
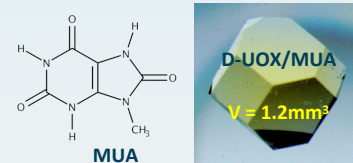


Identified the protonation states of active-site residues, and the W1 site as a neutral D₂O water molecule.

Perdeuterated UOX in complex with 9-methyl uric acid (MUA).

Peroxide intermediate formed in aerobic conditions.

LADI: Neutron data to 1.9 Å (>92%) at RT in 3d using a 1.2mm³ crystal of D-UOX/MUA.



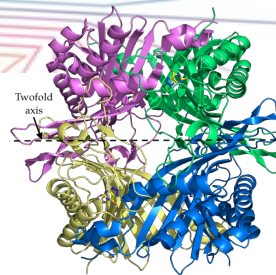
Identified the peroxide as deprotonated hence negatively charged. Thr57* is ideally positioned to protonate the peroxide Op2 atom.

Urate oxidase enzyme mechanism studies

PhD project of Lindsay McGregor, ILL/King's College London (R. Steiner)

Neutron data from the UOX complexes was used in the development of new tools for neutron crystallographic refinement using REFMAC5, CCP4.

Catapano *et al.*, (2023) *Acta Cryst.*, D79, in press.

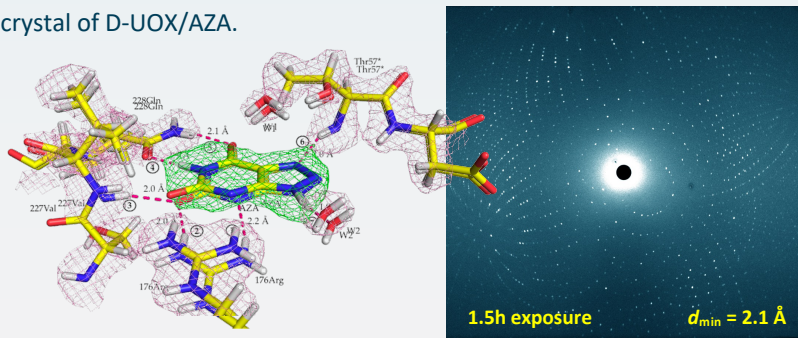
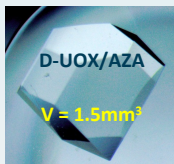
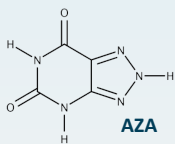


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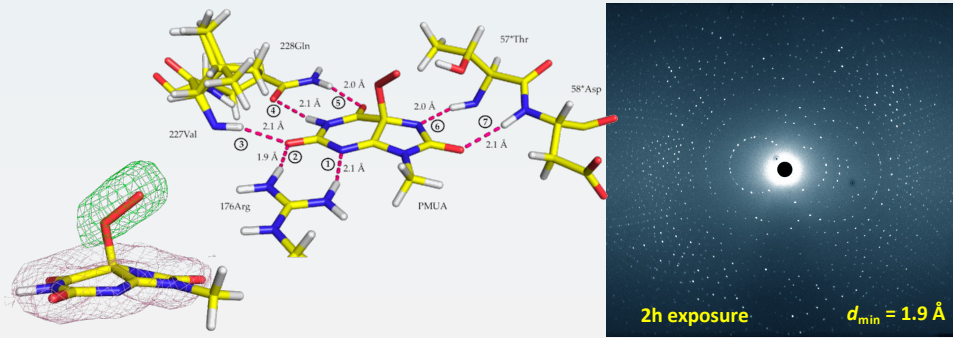
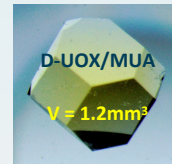
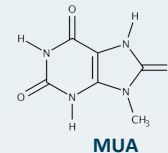


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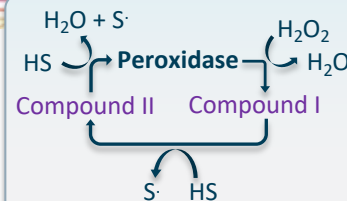


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Heme peroxidases cryo-crystallography studies

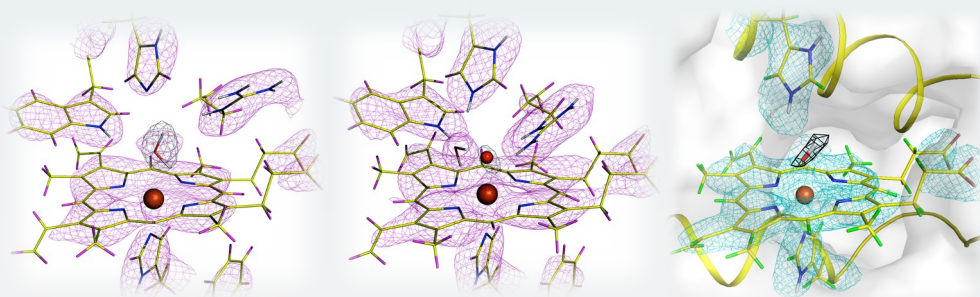
PhD project of Cecilia Casadei, ILL/University of Leicester (H. Kwon, P. Moody, E. Raven)

- **Heme peroxidases**, such as **cytochrome c peroxidase (CcP)** and **ascorbate peroxidase (APX)** carry out a wide range of oxidations using highly reactive ferryl intermediates. Reaction proceeds via **two transient intermediate states Compound I (CI) and Compound II (CII)**. Longstanding debate regarding the actual chemical species of **CI** and **CII**.



Cytochrome c peroxidase CI and ascorbate peroxidase CII at 100K

Cryo-trapped (at 100K) the transient reaction intermediates **CI in CcP** (by reaction with H_2O_2) and **CII in APX** (by reaction of *m*-CPBA), and collected neutron diffraction data at 100K. As a control, RT neutron data collected for **ferric CcP** (ground state).



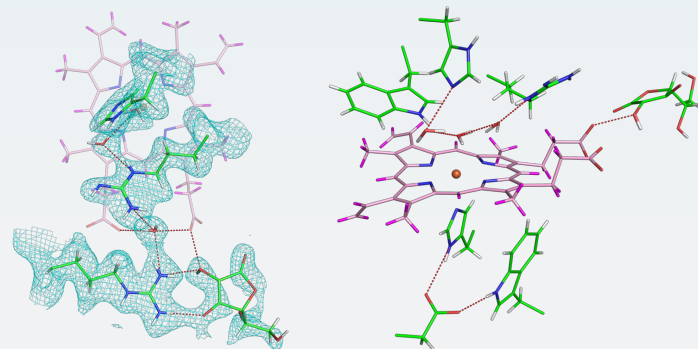
Ferric CcP at RT (LADI, 2.4 Å) CcP CI at 100K (BioDiff, 2.5 Å) APX CII at 100K (LADI, 2.2 Å)

Revealed the chemical species of CcP CI as Fe(IV)=O and APX CII as Fe(IV)-OH .

Casadei *et al.*, (2014) *Science*, **345**, 193; Kwon *et al.*, (2016) *Nat. Commun.*, **7**, 13445.

Perdeuterated APX (Ferric and ascorbate complex)

Neutron data at 100K using **D-APX** for (i) **ferric APX** (LADI, 2.2 Å, $V = 0.1\text{mm}^3$, 7.5d) and (ii) the **APX/ascorbate complex** (BioDiff, 2.1 Å). The ascorbate complex is stable and non-catalytic in absence of peroxide.



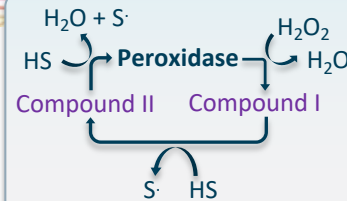
Revealed the protons in a PCET pathway, including an unexpected neutral Arg species in the APX/ascorbate complex.

Kwon *et al.*, (2020) *Proc. Natl. Acad. Sci., U.S.A.* **117**, 6484.

Heme peroxidases cryo-crystallography studies

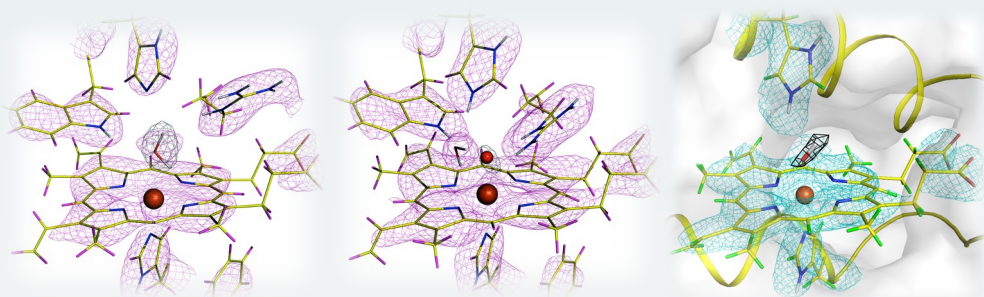
PhD project of Cecilia Casadei, ILL/University of Leicester (H. Kwon, P. Moody, E. Raven)

This work led to the development of improved low-temperature sample environment capabilities for LADI and the results led to revision of the textbook enzyme mechanism. (Moody & Raven, (2018) *Acc. Chem. Res.*, **51**, 427.)



Cytochrome c peroxidase CI and ascorbate peroxidase CII at 100K

Cryo-trapped (at 100K) the transient reaction intermediates **CI** in **CcP** (by reaction with H_2O_2) and **CII** in **APX** (by reaction of *m*-CPBA), and collected neutron diffraction data at 100K. As a control, RT neutron data collected for **ferric CcP** (ground state).



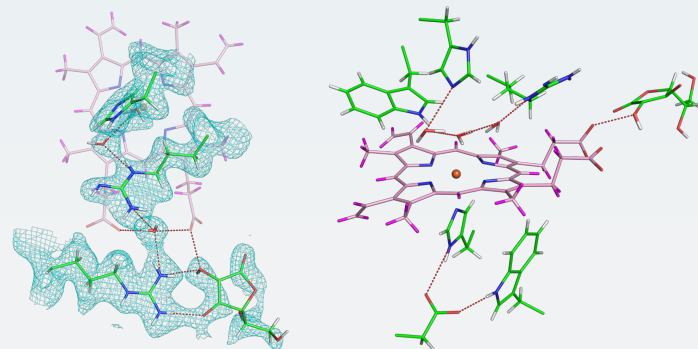
Ferric CcP at RT (LADI, 2.4 Å) CcP CI at 100K (BioDiff, 2.5 Å) APX CII at 100K (LADI, 2.2 Å)

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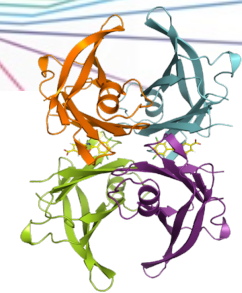


Revealed the protons in a PCET pathway, including an unexpected neutral Arg species in the APX/ascorbate complex.

Kwon *et al.*, (2020) *Proc. Natl. Acad. Sci., U.S.A.* **117**, 6484.

Transthyretin Amyloidosis

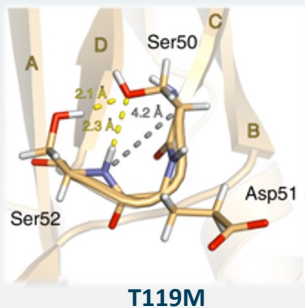
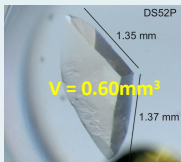
PhD project of Alycia Yee, ILL/University of Keele (T. Forsyth/E. Mitchell)



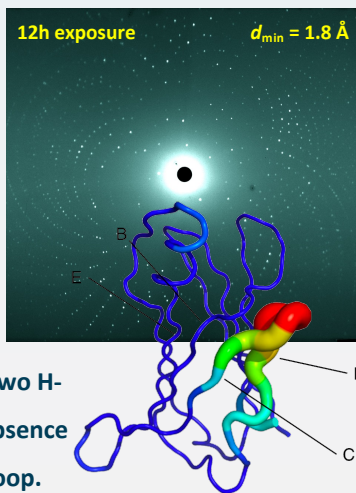
- **Transthyretin (TTR)** is a transport protein that carries thyroxine and retinol-binding protein bound to retinol.
- Wild-type TTR and certain mutants (e.g. **TTR_S52P**) are unstable, while other mutants (e.g. **TTR_T119M**) are stable. Unstable forms have a propensity to dissociate to form **amyloid fibrils** leading to a fatal hereditary disease, familial amyloid polyneuropathy (FAP).

Perdeuterated WT TTR, TTR_S52P & TTR_T119M

Analysed the H-bond networks in wild-type and mutant forms of TTR. **LADI**: Neutron data to 1.8 \AA at RT using a 0.6 mm^3 crystal of D-TTR_S52P.

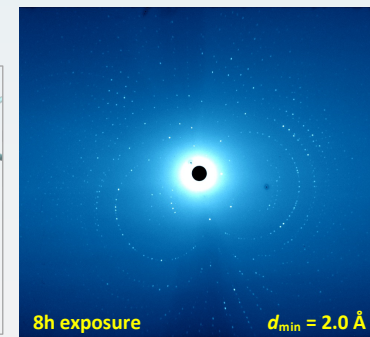
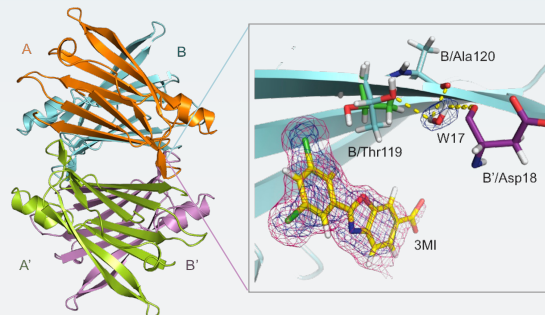
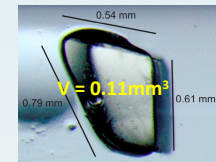


In T119M (and WT TTR), Ser52 forms two H-bonds with Ser50, while in S52P the absence of these H-bonds creates a looser CD loop.



Perdeuterated TTR_S52P/tafamidis complex

One strategy to inhibit TTR dissociation is to stabilise the tetramer by small molecule drug binding to the thyroxine (T4) binding sites. **LADI**: Neutron data to 2.0 \AA at RT in **4.5d** using a 0.11 mm^3 crystal of D-TTR_S52P/tafamidis complex.



Binding of tafamidis results in a change of orientation of Thr119, which creates additional water-mediated H-bonds between chains B and B'.

Yee *et al.*, (2019) *Nat. Commun.*, **10**, 925; Yee *et al.*, (2017) *J. Appl. Cryst.*, **50**, 660