# European Synchrotron Radiation Facility

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## **BAG Beam time Progress Report**

This represents a summary of the BAG progress in the reporting period, and is **in addition to** the standard ESRF report sheet for each project which will be used for the Review of the BAG.

#### **BAG** Title

#### **Allocation Period**

### List of publications resulting from ESRF beam time

- \*. M.J.Mate, M.Zamocky, P.M.Alzari, L.M.Nykyri, C.Betzel, F.Koller, **I.Fita** "Refined crystal structure of catalase-A from *Saccharomyces cerevisiae*". J.Mol.Biol. **286**, (1999), 135-149.
- \*. M.J.Maté, M.Serdal-Sevinc, B.Hu, J.Bujons, J.Bravo, J.Switala, W.Ens, P.C.Loewen I.Fita
  "Mutants that alter the covalent structure of Catalase-Hydroperoxidase II from Escherichia coli"
  J.Biol.Chem. 274, (1999), 27717-27725.
- \*. P.Nicholls, **I.Fita** and P.C.Loewen
  "Enzymology and Structure of Catalases"
  Advances in Inorganic Chemistry. Ed.G.Mauk. (In press).
- \*. M.J.Maté, G.Murshudov, J.Bravo, W.Melik-Adamyan, P.C.Loewen and I.Fita "Heme-Catalases" Handbook of Metalloproteins, John Wiley& Sons, Ltd. (In press).

### Global Summary

Catalases are heme-containing tetramers, with molecular weights ranging from about 200 till 350 kDa, that degrade hydrogen peroxide. Despite the large number of studies done on catalases belonging to the Clade I and II groups virtually no structural information exists from a most distinctive group of catalases (Clade III) that includes all plants and some bacterial enzymes. To overcome this situation data was collected at the ESRF from catalase crystals of *Pseudomonas*, of *Listeria* (space group I222 (or I212121) and unit cell parameters a=75 Å, b=122 Å and c=370 Å). Crystals are also available from the hemeless plant catalase from pea (space group P4<sub>1</sub>2<sub>1</sub>2, or its enantiomorph P4<sub>3</sub>2<sub>1</sub>2, with unit cell parameters a=111 Å, c=204 Å). The three catalases analysed belong to the Clade III group. Structure determination is now in progress including a search for heavy atom derivatives for the apparently very different structure adopted by the hemeless protein.

## Visits made to the ESRF

Date(s) of visits	Beamline	No. of Shifts	Short Summary of each Visit
1. 20-Nov-99/22-Nov-99	ID14-2	·	Data col.lection of native crystals of Listeria catalase at 2.5 A resolution
13-Feb-00/15-Feb-00	ID14-2		Data collection of native crystals of Pseudomonascatalase at 2.2 A resolution
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