| ESRF | **Experiment title:**<br>**Structural and functional studies of: a) Arrestin, b) β-arrestin...** | **Experiment number**:<br>LS-1816 BAG |
|---|---|---|
| **Beamline**:<br>ID14-3 | **Date of experiment**:<br>from: 28.10. to: 30.10.2000 | **Date of report**:<br>16.02.2001 |
| **Shifts:**<br>6 | **Local contact(s)**: Steffi Arzt | *Received at ESRF:* |

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**Report:**

Overall we collected 13 datasets at the beamline ID14-3, especially bacteriorhodopsin (bR) datasets. The quality of the bR datasets depend on a low twinning ratio (usable only if the twinning factor is <15% ) and the ratio of the intermediate to ground state (ratio intermediate/ground must be >60% in the crystal).

The dataset of arrestin soaked with a phosphorylated peptide has a usable resolution of 3.3 Å. The reduced diffraction quality is induced by the strong reaction of the peptide with the protein.

For β-arrestin co-crystallized with inositolhexaphosphate ($IP_6$) we recorded a datasets of 4.0 Å resolution. The $IP_6$ molecule is clear visible in the electron density.

A publication about the structure of arrestin complexed with inositolhexaphosphate is submitted to the Journal of Molecular Biology.