

Data phasing using the Molecular Replacement method or the Single/Multiple Anomalous Diffraction (SAD or MAD) method, including EXAFS

Friday, 20 March 2009 11:30 (1:00)

Content

Summary

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Session Classification : Data phasing using the Molecular Replacement method or the Single/Multiple Anomalous Diffraction (SAD or MAD) method, including EXAFS