

# 転写制御タンパク質構造解析研究チーム

## Transcription Regulation Protein Crystallography Team

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2004年度、我々は引き続き *Thermus thermophilus* HBS および *Pyrococcus horikoshii* OT3 由来タンパク質の構造研究を行い、さらにこれらの機能メカニズムを解明するため、リガンドおよび補酵素との複合構造の解析も行った。現在、我々は転写因子および転写後調節に関与するタンパク質に注目している。(12月1日より研究チーム名を、構造解析第3研究チームから転写制御タンパク質構造解析研究チームに変更)

During the 2004–2005 fiscal year our team continued structural studies of proteins from the *Thermus thermophilus* HB8 and *Pyrococcus horikoshii* OT3. In order to understand the functional mechanisms of these proteins, the structures of ligand and/or cofactor bound forms were also determined. Now we are particularly interested in proteins involved in transcription and post transcriptional regulation of gene expression.

### Research Subjects

1. Current state and future plans for high-throughput protein crystallography

### Staff

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### 誌上発表 Publications

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