

Uncertainty in NMR based 3D protein models

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Keywords

Pdb, Crystal, Pdb, Cif

Abstract

While I was working on implementing proper author-given chain IDs in PDB structures for Jmol's mmCIF reader today, I thought it was interesting to mention the recent article Traditional Biomolecular Structure Determination by NMR Spectroscopy Allows for Major Errors by Nabuurs (DOI:10.1371/journal.pcbi.0020009, open access), working at the CMBI, two floors away from my former working location at the Radboud University Nijmegen.

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