

Erratum to: Solution NMR structures reveal unique homodimer formation by a winged helix-turn-helix motif and provide first structures for protein domain family PF10771

Alexander Eletsky · Donald Petrey · Qiangfeng Cliff Zhang · Hsiau-Wei Lee · Thomas B. Acton · Rong Xiao · John K. Everett · James H. Prestegard · Barry Honig · Gaetano T. Montelione · Thomas Szyperski

Published online: 6 March 2012
© Springer Science+Business Media B.V. 2012

Erratum to: J Struct Funct Genomics
DOI 10.1007/s10969-011-9121-3

In addition to the financial support mentioned in the original publication, this work was also supported by the National Institutes of Health, grant number: GM030158-30 (to BH).

The online version of the original article can be found under doi:[10.1007/s10969-011-9121-3](https://doi.org/10.1007/s10969-011-9121-3).

A. Eletsky · T. Szyperski (✉)
Department of Chemistry, The State University of New York at Buffalo, and Northeast Structural Genomics Consortium,
Buffalo, NY 14260, USA
e-mail: szypersk@buffalo.edu

D. Petrey · Q. C. Zhang · B. Honig
Department of Biochemistry and Molecular Biophysics, Howard Hughes Medical Institute, Center for Computational Biology and Bioinformatics, Columbia University, New York, NY 10032, USA

H.-W. Lee · J. H. Prestegard
Complex Carbohydrate Research Center, University of Georgia, and Northeast Structural Genomics Consortium, Athens, GA 30602, USA

T. B. Acton · R. Xiao · J. K. Everett · G. T. Montelione
Department of Molecular Biology and Biochemistry, Center of Advanced Biotechnology and Medicine, Rutgers, The State University of New Jersey, Piscataway, NJ 08854, USA

T. B. Acton · R. Xiao · J. K. Everett · G. T. Montelione
Department of Biochemistry, Robert Wood Johnson Medical School, UMDNJ, and Northeast Structural Genomics Consortium, Piscataway, NJ 08854, USA