ERRATA

Ternary complex structure of human HGPRTase, PRPP, Mg²⁺, and the inhibitor HPP reveals the involvement of the flexible loop in substrate binding

GANESARATNAM K. BALENDIRAN,¹ JOSÉ A. MOLINA,^{2,6} YIMING XU,^{3,7} JAN TORRES-MARTINEZ,⁴ ROBERT STEVENS,⁵ PAMELA J. FOCIA,⁴ ANN E. EAKIN,⁴ JAMES C. SACCHETTINI,¹ AND SYDNEY P. CRAIG III⁴

¹Department of Biochemistry and Biophysics, Texas A&M University, College Station, Texas 77843-2128

²Department of Biochemistry, University of Puerto Rico-Medical Sciences Campus, San Juan, Puerto Rico 00936

³Department of Biochemistry, Temple University School of Medicine, Philadelphia, Pennsylvania 19140

⁴Laboratory of Molecular Parasitology and Drug Design, School of Pharmacy, CB#7360,

Research Triangle Park, North Carolina 27709

The PDB accession number for the structure published in the above referenced article (Volume 8, Number 5, 1999, pp. 1023–1031) is 1D6N. We regret the omission in the original article.

Probing the conformation of a human apolipoprotein C-1 by amino acid substitutions and trimethylamine-N-oxide

OLGA GURSKY

Department of Biophysics, Boston University School of Medicine, 715 Albany Street, Boston, Massachusetts 02118-2526

An NIH Grant number was inadvertently omitted from the acknowledgments printed on p. 2063 of the above referenced article (Volume 8, Number 10, 1999, pp. 2055–2064). The complete text of the Acknowledgments is reprinted below (the online edition of the article has been amended to correct for the omissions).

Acknowledgments

The author is indebted to Dr. David Atkinson for his support and help at all stages of this work and for a thorough reading of the manuscript; to Dr. Mary T. Walsh for her encouragement, help with the CD maintenance, and helpful comments on the manuscript; to Dr. G. Graham Shipley for critical reading on the manuscript; and to Michael Gigliotti for technical support. This work was supported by Grant-in-Aid 13–512–956 from the American Heart Association (Massachusetts Affiliate) and NIH Grant HL61429 to O.G., and by NIH Grant HL26335 (D.M. Smith, P.I.).

University of North Carolina at Chapel Hill, Chapel Hill, North Carolina 27599

⁵Mass Spectrometry Facility, Section of Biochemical Genetics, Duke University Medical Center,