

## Corrections

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Solution Structure and Dynamics of the Human–*Escherichia coli* Thioredoxin Chimera: Insights into Thermodynamic Stability, by Bindi Dangi, Anatoliy V. Dobrodumov, John M. Louis, and Angela M. Gronenborn\*, Volume 41, Number 30, July 30, 2002, pages 9376–9388.

Page 9376. The coordinates for the structure have been submitted to the Protein Data Bank (entry 1M7T).

BI0251687

10.1021/bi0251687

Published on Web 09/10/2002

Monounsaturated PE Does Not Phase-Separate from the Lipid Raft Molecules Sphingomyelin and Cholesterol: Role for Polyunsaturation? by Saame Raza Shaikh, Michael R. Brzustowicz, Noah Gustafson, William Stillwell, and Stephen R. Wassall\*, Volume 41, Number 34, August 27, 2002, pages 10593–10602.

Page 10598. The values for  $M_1$ ,  $\bar{S}_{CD}$ ,  $\langle L \rangle$ , and area quoted in Table 3 for POPE- $d_{31}$ , POPE- $d_{31}$ /SM (1:1 mol), and POPE- $d_{31}$ /SM/CHOL (1:1:1 mol) were calculated from  $^2\text{H}$  NMR spectra that were not baseline corrected after Fourier transformation. The conclusions remain the same. The corrected values appear below.

	$M_1$ ( $\times 10^4 \text{ s}^{-1}$ )	$\bar{S}_{CD}$	$\langle L \rangle$ ( $\text{\AA}$ )	area ( $\text{\AA}^2$ )
POPE- $d_{31}$	6.00	0.197	13.3	68.0
POPE- $d_{31}$ /SM (1:1 mol)	5.42	0.167	12.7	71.3
POPE- $d_{31}$ /SM/CHOL (1:1:1 mol)	8.12	0.266	14.6	62.0

BI0251721

10.1021/bi025172i

Published on Web 09/14/2002