—————ERRATA—————

A Protein Folding Degree Measure and Its Dependence on Crystal Packing, Protein Size, Secondary Structure, and Domain Structural Class [*J. Chem. Inf. Comput. Sci. 44,* 1238−1250 (2004)] By E. Estrada. Molecular Informatics, X-ray Unit, RIAIDT, Edificio CACTUS, University of Santiago de Compostela, 15782 Santiago de Compostela, Spain
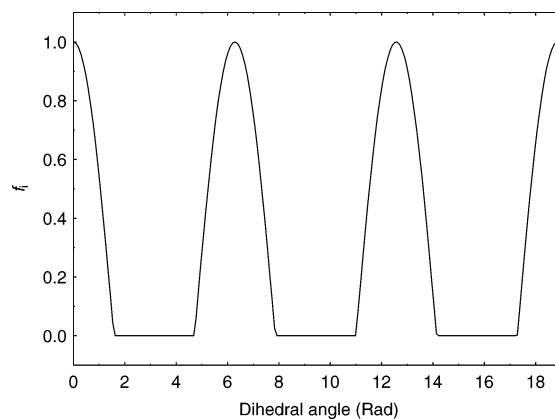
Pages 1242−1248. The function f_i used in the main diagonal of the matrices for the calculation of the I_3 index is not $f_i = \cos\theta_i$ as might be inferred from the text below eq 1 on page 1241. Instead we have used in all calculations the function

$$f_i = \frac{1}{2}[1 + \text{sgn}(\cos\theta_i)]\cos\theta_i \qquad (1)$$

where $\text{sgn}(x)$ is the sign function defined as

$$\text{sgn}(x) = \begin{cases} -1, x < 0 \\ 0, x = 0 \\ 1, x > 0 \end{cases} \qquad (2)$$

The plot of this function is given below for illustration.

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