Corrections

Solution NMR and Computer Simulation Studies of Active Site Loop Motion in Triosephosphate Isomerase, by Francesca Massi, Chunyu Wang, and Arthur G. Palmer, III*, Volume 45, Number 36, September 12, 2006, pages 10787−10794.

Page 10789. The equation in the legend of Figure 3 should be $((10\Delta\delta_H)^2 + \Delta\delta_N^2)^{1/2}$.

Page 10792. In the caption for Figure 8, the a and b panels are inverted. The first sentence should be: Distance between the C_α atoms of the residues of loop 6 and Y208 is represented as a function of time for trajectories starting from the open (pdb 1YPI) (a) and closed (pdb 7TIM) (b) states of TIM.

Page 10794. The following reference describing the computer program used for simulations should have been included: Brooks, B. R., Bruccoleri, R. E., Olafson, B. D., States, D. J., Swaminathan, S., and Karplus, M. (1983) CHARMM: A program for macromolecular energy, minimization, and dynamics calculations, *J. Comput. Chem. 4*, 187−217.

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