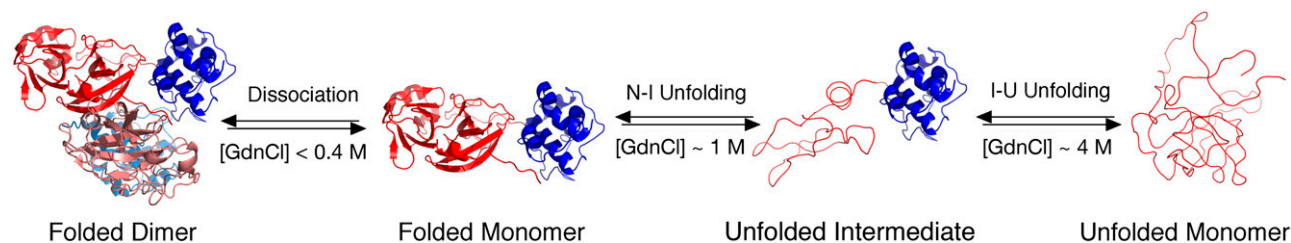


Corrections

Hui-Ping Chang, Chi-Yuan Chou, and Gu-Gang Chang. 2007. Reversible unfolding of the severe acute respiratory syndrome coronavirus main protease in guanidinium chloride. *Biophys. J.* 92:1374–1383

Fig. 8 was printed in black and white. It should have been in color. Here is the correct image:



doi: 10.1529/biophysj.107.0900157

Marcin Imielinski, Calin Belta, Harvey Rubin, and Adam Halász. 2006. Systematic analysis of conservation relations in *Escherichia coli* genome-scale metabolic network reveals novel growth media. *Biophys. J.* 90:2659–2672.

Part of the text incorrectly cited references instead of equations. The corrected text reads:

Employing Lemma 2, we observe that the nonemptiness of the set in Eq. 7 with $T = \emptyset$,

$$\left\{ u \in \mathbb{R}^{|U|}, v \in \mathbb{R}^n \mid \hat{S} \begin{bmatrix} u \\ v \end{bmatrix} \geq 0, \left(\hat{S} \begin{bmatrix} u \\ v \end{bmatrix} \right)_i > 0 \right\}, \quad (10)$$

is equivalent to the emptiness of the set

$$\{g \in \mathbb{R}^m \mid g^T \hat{S} = 0, g \geq 0, g_i > 0\}, \quad (11)$$

which, using Eqs. 8 and 9, can be written as

$$\{g \in G \mid g_i > 0, g_U = 0\}. \quad (12)$$

doi: 10.1529/biophysj.107.0900158

Jeroen van Gestel and Simon W. de Leeuw. 2007. The formation of fibrils by intertwining of filaments: Model and application to amyloid A β protein. *Biophys. J.* 92:1157–1163.

In Figs. 1, 2, 3, and 7, on the vertical axis of each graph, the length scale should read micrometer, rather than millimeter. Similarly, on the horizontal axis of the graph depicted in Fig. 7, the pitch is given in micrometers.

doi: 10.1529/biophysj.107.0900159

Benjamin R. G. Johnson, Richard J. Bushby, John Colyer, and Stephen D. Evans. 2006. Self-assembly of actin scaffolds at ponticulin containing supported phospholipid bilayers. *Biophys. J.* 90:L21–L23.

An error in the axis units has been noticed on Fig. 3, which has affected the published value for the dissociation constant for the binding of F-actin filaments to a ponticulin containing tethered bilayer. The corrected dissociation constant value should read $11 \pm 5 \mu\text{M}$. The corrected value suggests similar affinity binding as for the actin binding protein profilin (13).

doi: 10.1529/biophysj.107.0900160