

Effects of the Arctic (E²²→G) Mutation on Amyloid β -Protein Folding: Discrete Molecular Dynamics Study [*Journal of the American Chemical Society* **2008**, *130*, 17413–17422. DOI: 10.1021/ja804984h]. A. R. Lam,* D. B. Teplow, H. E. Stanley, and B. Urbanc

Page 17416. In the first paragraph of section 3.1, we reported that the strength of the effective hydrophobic interactions, E_{HP} , is by definition equal to the absolute value of the potential energy between two residues at a distance of $<7.5 \text{ \AA}$. The correct value for this distance used in the simulations was $<5 \text{ \AA}$.

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