

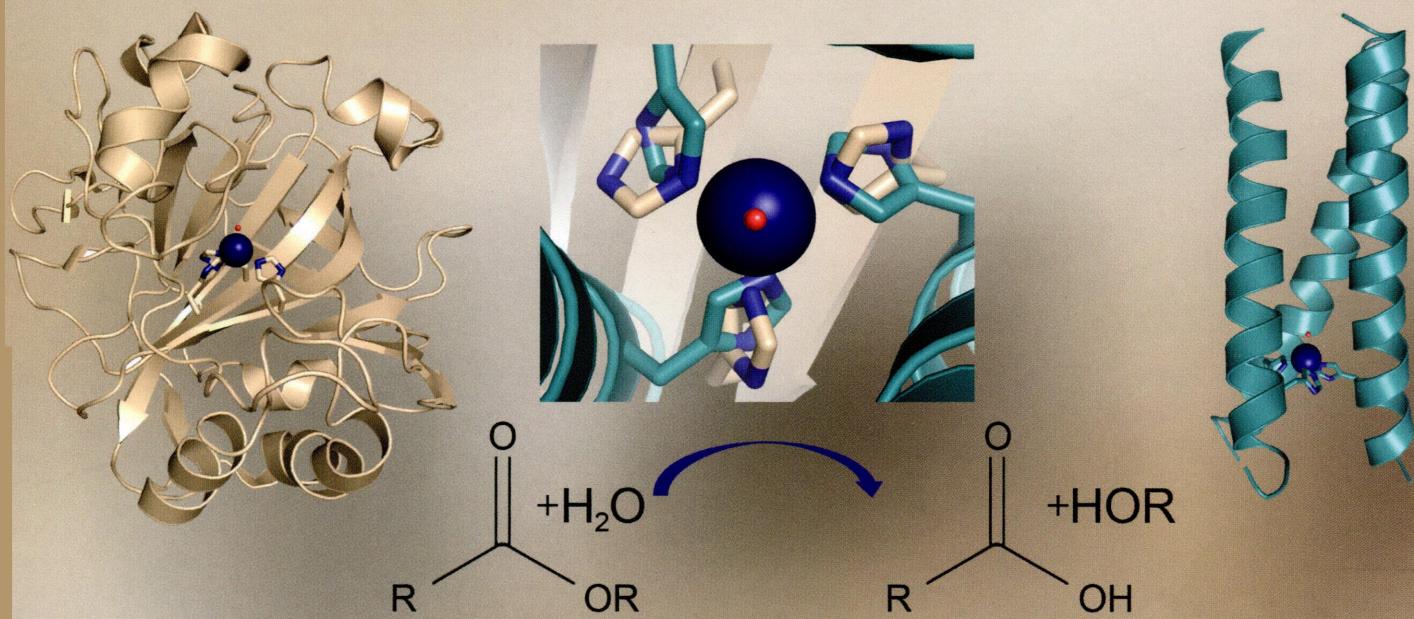
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ON THE COVER: Using de novo and redesign approaches toward the preparation of structural and functional models of hydrolytic zinc metalloenzymes. [Zastrow, M. L., and Pecoraro, V. L. (2014) *Biochemistry* 53, 957–978]

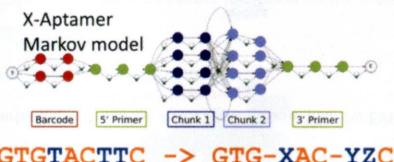
Rapid Reports

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[dx.doi.org/10.1021/bi500443e](https://doi.org/10.1021/bi500443e)

Aptaligner: Automated Software for Aligning Pseudorandom DNA X-Aptamers from Next-Generation Sequencing Data
Emily Lu, Miguel-Angel Elizondo-Riojas, Jeffrey T. Chang,* and David E. Volk*

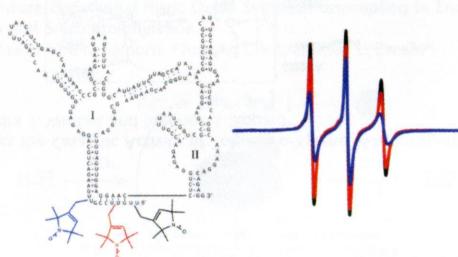


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[dx.doi.org/10.1021/bi500404b](https://doi.org/10.1021/bi500404b)

Characterizing the Dynamics of the Leader–Linker Interaction in the Glycine Riboswitch with Site-Directed Spin Labeling
Jackie M. Esquiaqui, Eileen M. Sherman, Sandra A. Ionescu, Jing-Dong Ye,* and Gail E. Fanucci*



Articles

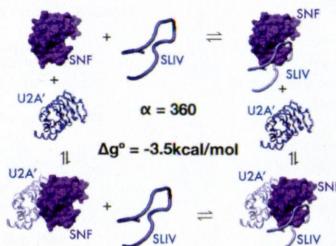
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Linkage and Allostery in snRNP Protein/RNA Complexes

Sandra G. Williams and Kathleen B. Hall*

dx.doi.org/10.1021/bi500192a



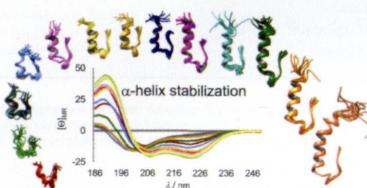
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Rational Design of α -Helix-Stabilized Exendin-4 Analogs

Petra Rovó, Viktor Farkas, Pál Stráner, Mária Szabó, Ágnes Jermendy, Orsolya Hegyi, Gábor K. Tóth, and András Perczel*

dx.doi.org/10.1021/bi500033c

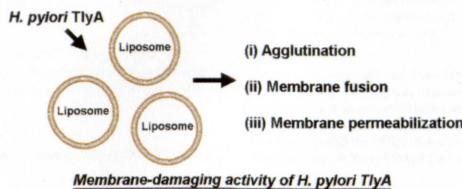


3553

dx.doi.org/10.1021/bi500152n

Helicobacter pylori TlyA Agglutinates Liposomes and Induces Fusion and Permeabilization of the Liposome Membranes

Kusum Lata and Kausik Chattopadhyay*

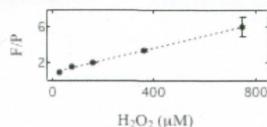


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dx.doi.org/10.1021/bi401078b

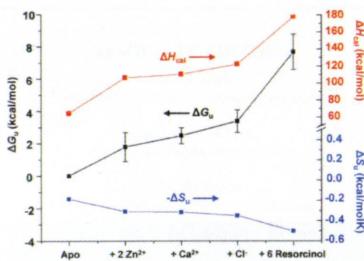
How Hydrogen Peroxide Is Metabolized by Oxidized Cytochrome c Oxidase

Daniel Jancura,* Jana Stanicova, Graham Palmer, and Marian Fabian*



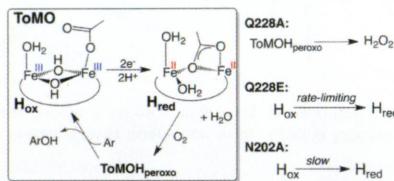
Thermodynamic Contributions to the Stability of the Insulin Hexamer

George P. Lisi, Chien Yi M. Png, and Dean E. Wilcox*



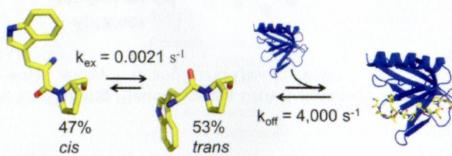
A Flexible Glutamine Regulates the Catalytic Activity of Toluene o-Xylene Monooxygenase

Alexandria Deliz Liang, Alexandra T. Wrobel, and Stephen J. Lippard*



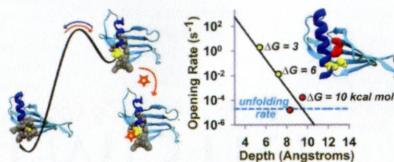
Isomerase-Catalyzed Binding of Interleukin-1 Receptor-Associated Kinase 1 to the EVH1 Domain of Vasodilator-Stimulated Phosphoprotein

Alexander I. Greenwood, Jeahoo Kwon, and Linda K. Nicholson*



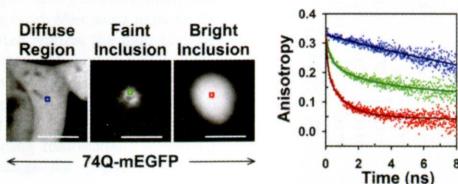
High-Energy Intermediates in Protein Unfolding Characterized by Thiol Labeling under Nativelike Conditions

Pooja Malhotra and Jayant B. Udgaonkar*



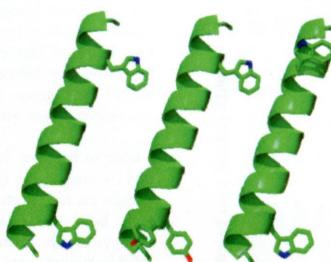
Fluorescence Anisotropy Uncovers Changes in Protein Packing with Inclusion Growth in a Cellular Model of Polyglutamine Aggregation

Vishal Bhardwaj, Mitradas M. Panicker, and Jayant B. Udgaonkar*



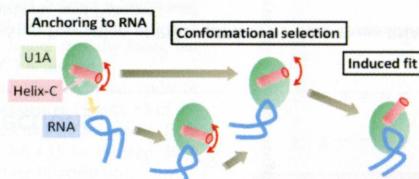
Comparisons of Interfacial Phe, Tyr, and Trp Residues as Determinants of Orientation and Dynamics for GWALP Transmembrane Peptides

Kelsey A. Sparks, Nicholas J. Gleason, Renetra Gist, Rebekah Langston, Denise V. Greathouse, and Roger E. Koeppe II*



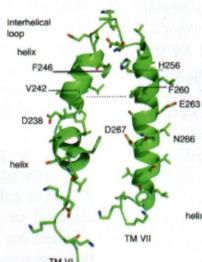
Combined Mechanism of Conformational Selection and Induced Fit in U1A–RNA Molecular Recognition

Ikuo Kurisaki, Masayoshi Takayanagi, and Masataka Nagaoka*



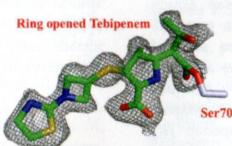
Structural and Functional Analysis of the Transmembrane Segment Pair VI and VII of the NHE1 Isoform of the Na^+/H^+ Exchanger

Claudia Alves, Brian L. Lee, Brian D. Sykes, and Larry Fliegel*



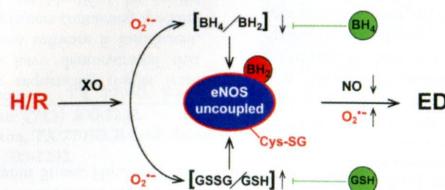
Tebipenem, a New Carbapenem Antibiotic, Is a Slow Substrate That Inhibits the β -Lactamase from *Mycobacterium tuberculosis*

Saugata Hazra, Hua Xu, and John S. Blanchard*



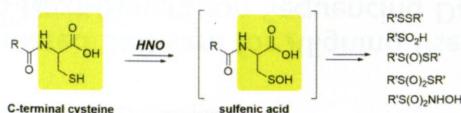
Hypoxia and Reoxygenation Induce Endothelial Nitric Oxide Synthase Uncoupling in Endothelial Cells through Tetrahydrobiopterin Depletion and S-Glutathionylation

Francesco De Pascali, Craig Hemann, Kindra Samons, Chun-An Chen, and Jay L. Zweier*



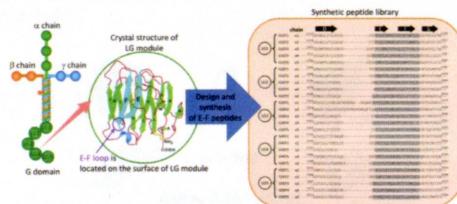
Reactivity of C-Terminal Cysteines with HNO

Gizem Keceli and John P. Toscano*



Biological Activities of the Homologous Loop Regions in the Laminin α Chain LG Modules

Fumihiko Katagiri, Toshihiro Hara, Yuji Yamada, Shunsuke Urushibata, Kentaro Hozumi, Yamato Kikkawa, and Motoyoshi Nomizu*

**Additions and Corrections****Addition to Entropy-Driven One-Step Formation of Phi29 pRNA 3WJ from Three RNA Fragments**

Daniel W. Binzel, Emil F. Khisamutdinov, and Peixuan Guo*

The authors would like to add the following information to the manuscript:

In the original manuscript, we described the formation of the Phi29 pRNA 3WJ from three RNA fragments (Binzel et al., 2015). We have now determined the sequence of the third fragment, which is 20 nucleotides long and contains a hairpin loop. This addition increases the total length of the pRNA 3WJ by 20 nucleotides. The addition of the third fragment does not change the overall structure of the pRNA 3WJ, which remains a single hairpin loop. The addition of the third fragment also does not change the overall entropy of the system, as the entropy of the system is determined by the entropy of the individual fragments. The addition of the third fragment does not change the overall entropy of the system, as the entropy of the system is determined by the entropy of the individual fragments.



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