

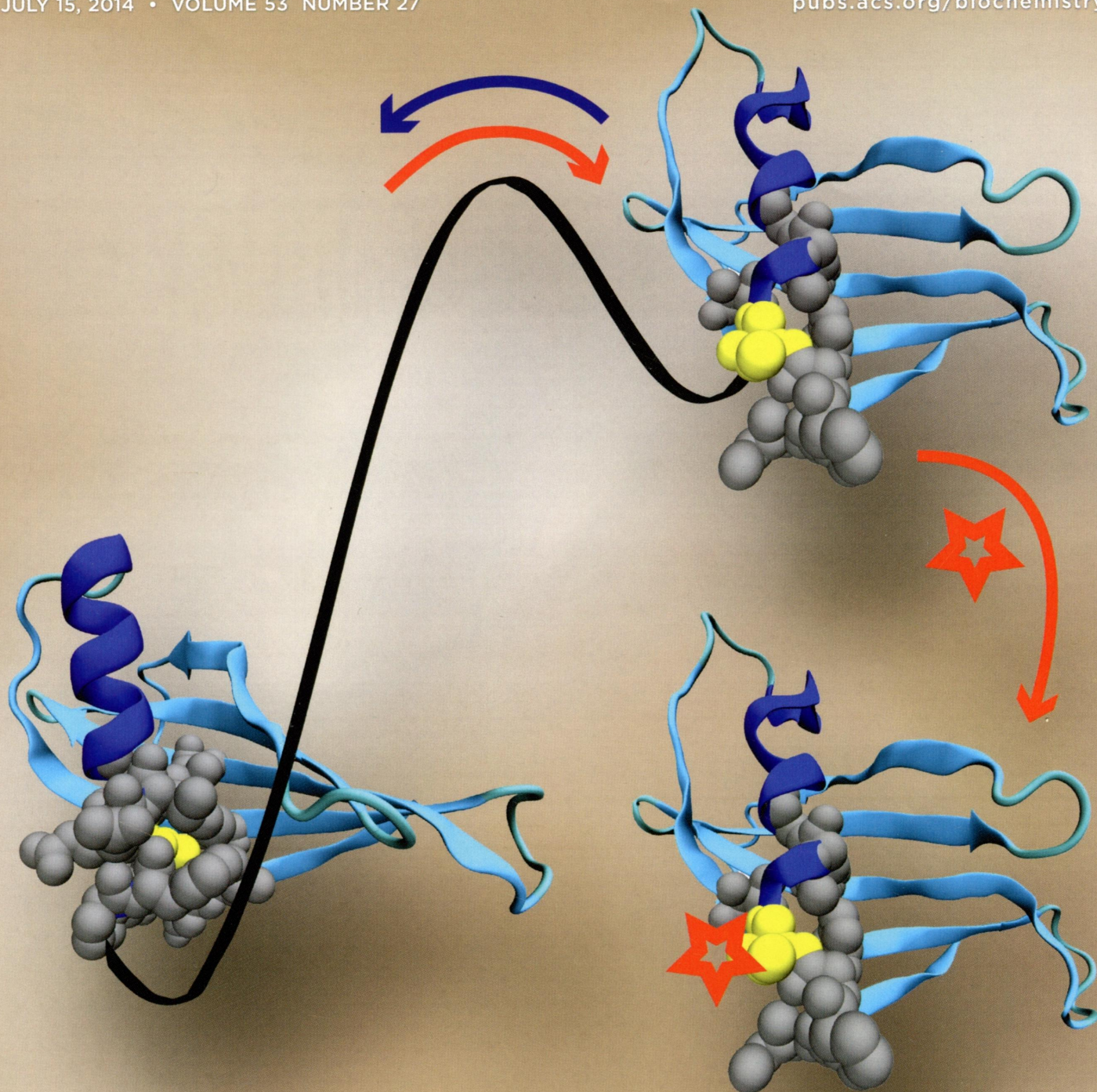
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JULY 15, 2014 • VOLUME 53 NUMBER 27

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ON THE COVER: Thiol labeling of a single cysteine residue (yellow spheres) has been used to monitor rare unfolding events in a protein under natively like conditions. The residues (gray spheres) surrounding the buried cysteine move apart, resulting in solvent exposure and hence labeling of the side chain thiol. This deprotection of the side chain is associated with an energy barrier between the native state and a partially unfolded, labeling-competent intermediate. Such intermediates have been mapped onto the unfolding energy landscape of the protein monellin using the kinetic and thermodynamic information obtained from thiol labeling. [Malhotra, P., and Udgaonkar, J. B. (2014) *Biochemistry* 53, 3608–3620]

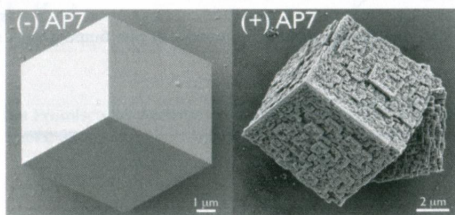
Rapid Reports

4317 **S**

[dx.doi.org/10.1021/bi500664w](https://doi.org/10.1021/bi500664w)

The Intrinsically Disordered C-RING Biominerization Protein, AP7, Creates Protein Phases That Introduce Nanopatterning and Nanoporosities into Mineral Crystals

Eric P. Chang, Jennie A. Russ, Andreas Verch, Roland Kröger, Lara A. Estroff, and John Spencer Evans*

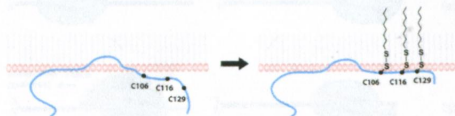


4320 **S**

[dx.doi.org/10.1021/bi5005238](https://doi.org/10.1021/bi5005238)

Modest Effects of Lipid Modifications on the Structure of Caveolin-3

Ji-Hun Kim, Dungeng Peng, Jonathan P. Schleich, Arina Hadziselimovic, and Charles R. Sanders*

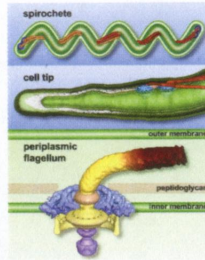


4323

[dx.doi.org/10.1021/bi500059y](https://doi.org/10.1021/bi500059y)

Molecular Architecture of the Bacterial Flagellar Motor in Cells

Xiaowei Zhao, Steven J. Norris, and Jun Liu*



Articles

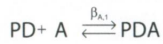
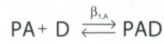
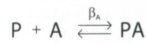
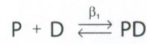
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[dx.doi.org/10.1021/bi500224t](https://doi.org/10.1021/bi500224t)

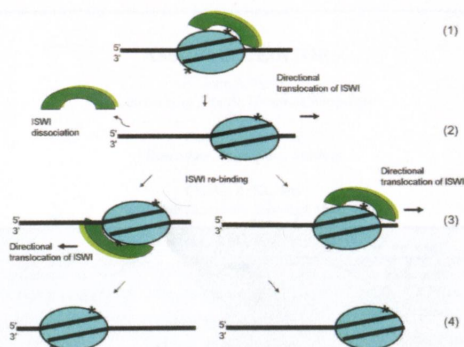
Quantitative Determination of Binding of ISWI to Nucleosomes and DNA Shows Allosteric Regulation of DNA Binding by Nucleotides

Gada Al-Ani, Koan Briggs, Shuja Shafi Malik, Michael Conner, Yoshiaki Azuma, and Christopher J. Fischer*



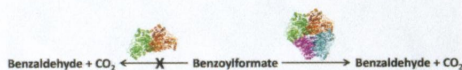
ISWI Remodels Nucleosomes through a Random Walk

Gada Al-Ani, Shuja Shafi Malik, Allen Eastlund, Koan Briggs, and Christopher J. Fischer*

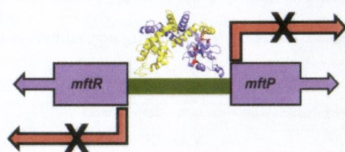


Perturbation of the Monomer–Monomer Interfaces of the Benzoylformate Decarboxylase Tetramer

Forest H. Andrews, Megan P. Rogers, Lake N. Paul, and Michael J. McLeish*

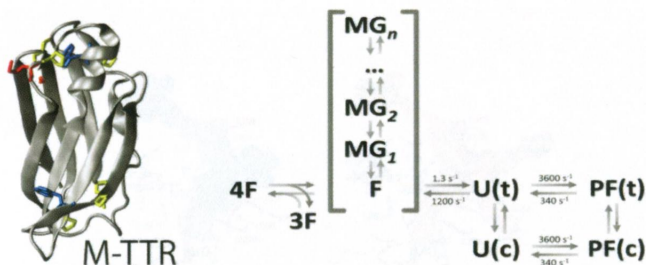
Ligand-Binding Pocket Bridges DNA-Binding and Dimerization Domains of the Urate-Responsive MarR Homologue MftR from *Burkholderia thailandensis*

Ashish Gupta and Anne Grove*



A Complex Equilibrium among Partially Unfolded Conformations in Monomeric Transthyretin

Simona Conti, Xinyi Li, Stefano Gianni, Seyyed Abolghasem Ghadami, Joel Buxbaum, Cristina Cecchi, Fabrizio Chiti, and Francesco Bemporad*

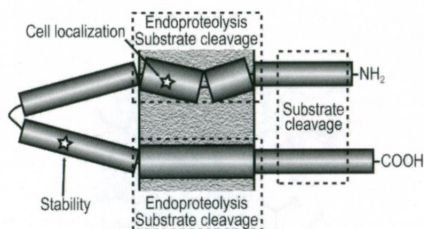


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dx.doi.org/10.1021/bi500489j

Pen-2 Is Essential for γ -Secretase Complex Stability and Trafficking but Partially Dispensable for Endoproteolysis

Oliver Holmes, Swetha Paturi, Dennis J. Selkoe, and Michael S. Wolfe*

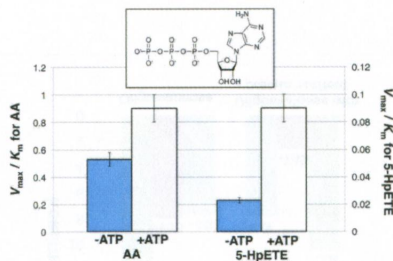


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dx.doi.org/10.1021/bi401621d

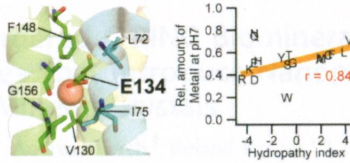
ATP Allosterically Activates the Human 5-Lipoxygenase Molecular Mechanism of Arachidonic Acid and 5(S)-Hydroperoxy-6(E),8(Z),11(Z),14(Z)-eicosatetraenoic Acid

Christopher J. Smyrniotis, Shannon R. Barbour, Zexin Xia, Mark S. Hixon, and Theodore R. Holman*



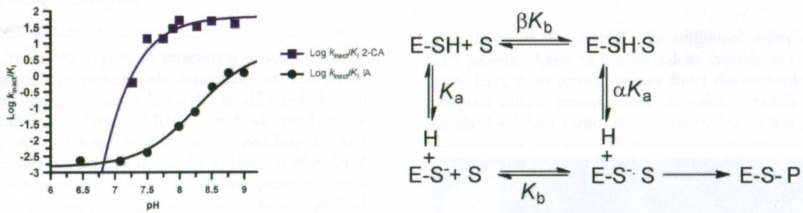
Contribution of Glutamic Acid in the Conserved E/DRY Triad to the Functional Properties of Rhodopsin

Keita Sato, Takahiro Yamashita, and Yoshinori Shichida*



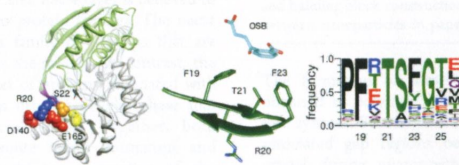
Mechanistic Studies of Protein Arginine Deiminase 2: Evidence for a Substrate-Assisted Mechanism

Christina J. Dreyton, Bryan Knuckley, Justin E. Jones, Daniel M. Lewallen, and Paul R. Thompson*



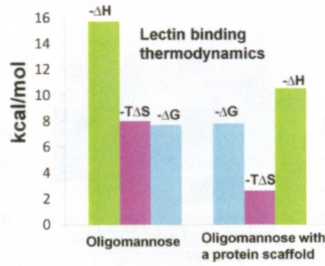
Role of an Active Site Loop in the Promiscuous Activities of *Amycolatopsis* sp. T-1-60 NSAR/OSBS

Andrew W. McMillan, Mariana S. Lopez, Mingzhao Zhu, Benjamin C. Morse, In-Cheol Yeo, Jaleesia Amos, Ken Hull, Daniel Romo, and Margaret E. Glasner*



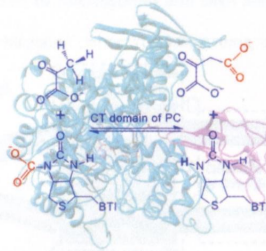
Significant Other Half of a Glycoconjugate: Contributions of Scaffolds to Lectin–Glycoconjugate Interactions

Melanie L. Talaga, Ni Fan, Ashli L. Fueri, Robert K. Brown, Yoann M. Chabre, Purnima Bandyopadhyay, René Roy, and Tarun K. Dam*



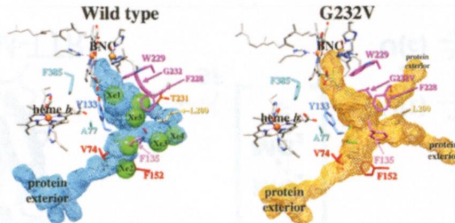
QM/MM Study of the Reaction Mechanism of the Carboxyl Transferase Domain of Pyruvate Carboxylase from *Staphylococcus aureus*

Xiang Sheng and Yongjun Liu*



Conserved Glycine 232 in the Ligand Channel of b_3 Cytochrome Oxidase from *Thermus thermophilus*

William McDonald, Chie Funatogawa, Yang Li, Ying Chen, Istvan Szundi, James A. Fee, C. David Stout, and Ólóf Einarsdóttir*



Novel Human Butyrylcholinesterase Variants: Toward Organophosphonate Detoxication

Mary Dwyer, Sacha Javor, Daniel A. Ryan, Emily M. Smith, Beilin Wang, Jun Zhang, and John R. Cashman*

