19th Annual Gibbs Conference on Biothermodynamics

Saturday, October 15 - Tuesday, October 18, 2005

Touch of Nature Environmental Center Southern Illinois University Carbondale, Illinois

Organizers

Clay Clark Trevor Creamer

Sponsors

Abbott Laboratories

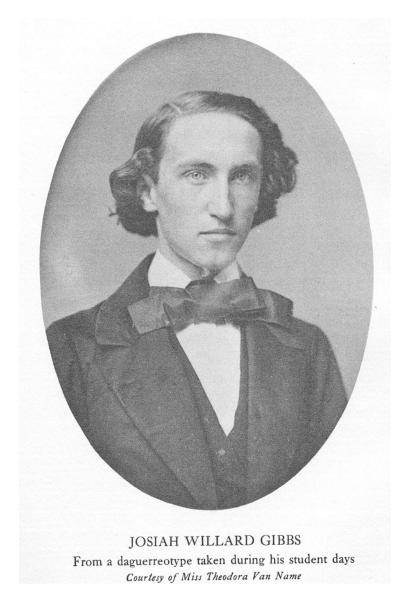
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<u>Cover figure:</u> Structural alignment of complexes of Calmodulin (CaM) with Trifluoperazine (TFP) in 1:1, 1:2, and 1:4 CaM:TFP ratios, RMSD = 0.59Å. Alignment and graphics created with Pymol ™. Kindly provided by **Michael Feldkamp** (U. lowa).

1. The Gibbs Conference on Biothermodynamics is only possible because of:



Let's face it, without Mr. Gibbs we'd all be doing something else, like genetics, or (gasp!) kinetics...

History of the Gibbs Conference on Biothermodynamics

Touch of Nature Environmental Center, Carbondale, Illinois

Fall, 1986	1 st discussion of the discipline of Thermodynamics in Biological Systems. held in Vail, Colorado: Gary Ackers, Wayne Bolen, Ernesto Freire, Stan Gill, Jim Lee
Feb, 1987	2 nd discussion of the discipline of Thermodynamics in Biological Systems. held in New Orleans: Gary Ackers, Norma Allewell, Wayne Bolen, Ken Breslauer, Ken Dill, Ernesto Freire, Stan Gill, Jim Lee
	Annual Gibbs Conference on Biothermodynamics
1 st – 1987	Organizers: Jim Lee & Wayne Bolen
2 nd - 1988	Organizers: Gary Ackers & Michael Johnson
3 rd - 1989	Organizers: Susan G, Frasier & Michael Johnson
4 th - 1990	Organizers: Michael Johnson & Marty Straume
5 th - 1991	Organizers: Gary Ackers & Tim Lohman Keynote: Ernesto Freire
6 th - 1992	Organizers: Jim Lee & Tomasz Heyduk Keynotes: Serge Timasheff & John Schellman
7 th - 1993:	Organizers: Maurice Eftink & Glen Ramsay Keynotes: Peter von Hippel & Julian Sturtevant
8 th - 1994:	Organizers: Enrico Di Cera & Madeline Shea Keynotes: Gary Ackers & Kathleen S. Matthews
9 th - 1995:	Organizers: Kenneth P. Murphy & Michael D. Brenowitz Keynotes: Victor Bloomfield & Mario Amzel
10 th - 1996:	Organizers: Jonathan B. Chaires & Michael L. Doyle Keynotes: J. Michael Schurr & Allen Minton
11 th - 1997:	Organizers: Dorothy Beckett & Jack Correia Keynote: Adrian Parsegian

12th - 1998: Organizer: Andy Robertson

Keynote: David Draper

13th - 1999: Organizers: Bertrand Garcia-Moreno & John Shriver

Keynotes: Wayne Bolen & Gary Ackers

14th – 2000: Organizers: George Turner & Kim Sharp

Keynotes: Steve White (replaced Rodney Biltonen)

15th - 2001: Organizers: Margaret A. Daugherty & Luis A. Marky

Keynote: George Rose

16th - 2002: Organizers: Michael Mossing & George Makhatadze

Keynote: Rodney Biltonin

17th - 2003: Organizers: Vince Hilser & Dick Sheardy

Keynote: Jim Lee

18th – 2004: Organizers: Doug Barrick & Kathleen Hall

Keynote: Ignacio Tinoco

19th – 2005: Organizers: Clay Clark & Trevor Creamer

Keynote: Carl Frieden

In 2002 The Gibbs Conference on Biothermodynamics incorporated as a mechanism of preserving the philosophy and spirit of the meeting. For a published 10 year history see Ackers and Bolen, "The Gibbs Conference on Biothermodynamics: Origins and evolution." Biophys. Chem. **64**, p 3-5, 1997.

President: Madeline Shea, Oct, 2004 – Oct, 2005.

President Elect: Dorothy Beckett.

Secretary: Margaret A. Daugherty, Oct, 2002 – Oct, 2008.

Treasurer: Michael Johnson, Oct, 2002 – Oct, 2007.

Past Presidents:

Gary Ackers, Oct, 2001 – Oct, 2002. Jack Correia, Oct, 2002 – Oct, 2003. Wayne Bolen, Oct, 2003 – Oct, 2004.

2. Meeting Schedule

Saturday, October 15

4:00-7:00 Check-in

7:00-10:00 Reception in Indian Room

Sunday, October 16

7:30 - 8:30 Breakfast in Freeberg Hall

Session I: Nucleic Acids

	Moderator: Melanie Bozza (Seton Hall U., R. Sheardy lab)
9:00 9:10	Welcome (Clay Clark, Trevor Creamer and Madeline Shea) Brad Chaires (U. Louisville) Uncovering the energetic basis of G-quadruplex stability.
9:50	Mikhail Karymov (U. Nebraska, Y. Lyubchenko lab) Single molecule analysis of Holliday junction dynamics and branch migration.
10:20	Break
10:45	Matt Freyer (Northen Arizona U., E. Lewis lab) Thermodynamic studies of the binding of netropsin and DAPI to several A_2T_2 containing DNA hairpins.
11:15	Kathleen Hall (Washington U.) Thermodynamics of the conformational change of a Mg ²⁺ - sensing Riboswitch.
12:00	Lunch in Freeberg Hall

Session II: Stability & Evolution

	Moderator: Brian Cannon (Johns Hopkins U., B. García-Moreno lab)
2:30	Matt Cordes (U. Arizona) Mutationally induced changes in protein structure during evolution.
3:10	Chris Olsen (U. Nebraska, L. Marky lab) Unfolding of G-quadruplexes: Thermodynamic contributions for the stacking of two G-quartets with $T\rightarrow U$ substitutions.
3:40	Break
4:00	Mirco Junker (U. Notre Dame, P. Clark lab) The pertactin β -helix folding mechanism suggests common themes for the secretion and folding of autotransporter proteins.
4:30	Sarah Woodson (Johns Hopkins U.) How RNA gets in shape.
5:15	Keynote Address – Carl Frieden (Washington U.) <i>Protein folding: Are we there yet?</i> Introduction by George Rose
6:30	Dinner in Freeberg Hall
8:00	Posters & Beer/Wine I in Sledgefoot Hall

Monday, October 17

7:00 - 8:30 Breakfast in Freeberg Hall

Session III: Denatured States

	Moderator: Veronique Chellgren (U. Kentucky, T. Creamer lab)
9:00	Dan Raleigh (Stony Brook U.) The role of the unfolded state in protein folding and stability.
9:40	Nick Fitzkee (Johns Hopkins U., G. Rose lab) How sterics and solvation reduce the size of protein conformational space.
10:10	Break
10:40	Hoang Tran (Washington U., R. Pappu lab) Towards an accurate model for protein denatured states.
11:10	Neville Kallenbach (New York U.) Structure in peptide models of unfolded proteins.
12:00	Lunch in Freeberg Hall
12:00	Gibbs business meeting: All previous organizers please attend.

Session IV: Macromolecular Interactions

	Moderator: Daniel Deredge (LSU, V. LiCata lab)
2:30	Elizabeth Komives (U. California, San Diego) Biophysics of the NF- κ B/I κ B α interaction.
3:10	Olga Lubman (Johns Hopkins U., D. Barrick lab) In vitro dissection of Notch/RBPjk interaction: Insights into the mechanism of Notch mediated transcriptional switch.
3:40	Rhonda Newman (U. Iowa, M. Shea lab) Interactions of calmodulin with regulatory regions of the ryanodine receptor type 1: Distinct roles of domainsin protein allostery.
4:10	Break
4:40	Aaron Lucius (UTMB Galveston, W. Bujalowski lab) Allosteric interactions between the nucleotide-binding sites and the ssDNA-binding site in the PriA helicase-ssDNA complex.
5:10	Jack Correia (U. Mississippi) The use of multi-wavelength and fluorescence AUC: Case studies of tubulin regulatory factors and antimitotic drugs.
6:30	Dinner in Freeberg Hall – Buffalo Tro
8:00	Posters & Beer/Wine II in Sledgefoot Hall

Tuesday, October 18

7:00 - 8:30 Breakfast in Freeberg Hall

Session V: Membranes

	Moderator: Horia Petrache (NIH, A. Parsegian lab)
8:30	Announcements
8:40	Lukas Tamm (U. Virginia) Structural dynamics and thermodynamics of the gating of the OmpA ion channel.
9:20	Yuhua Song (Washington U., N. Baker lab) Molecular dynamics simulations of salicylate effects on the micro- and mesoscopic properties of a dipalmitoylphosphatidylcholine bilayer.
9:50	Break
10:10	Damien Thevenin (U. Delaware, C. Robinson lab) Folding and assembly of adenosine receptors.
10:40	AnnMarie Stanley (Johns Hopkins U., K. Fleming lab) Thermodynamics of dimerization of the outer membrane phospholipase A transmembrane β -barrel.
11:10	Stephen White (U. California, Irvine) Translocons, membranes, and the folding of membrane proteins.
11:50	Wrap Up & Lunch

3. Posters

Poster Information:

Posters will be presented in one of two evening sessions in Sledgefoot Hall (next to Freeberg Dining Hall). Session I will be held Sunday evening and Session II on Monday evening. Both sessions will start at 8:00pm.

Session I: First authors* whose last names begin with A through I. Posters may be mounted starting on Sunday morning. Please take your poster down by the break on Monday morning.

Session II: First authors* whose last names begin with J through Z. Posters may be mounted starting on Monday morning. Please take your poster down before the talks start on Tuesday morning.



"Look! The bar is open!"

"Yesss!"

* Note that we are using the last name of the first author listed on the poster. If you are presenting a poster but are not first author, please present in the session appropriate for the last name of the first author.

Poster Session I: List of Presentations (A - I)

Oxygenation of normal human hemoglobin contains both cooperative and noncooperative binding steps. Why?

Gary K. Ackers, Alexandra Klinger, Connie Yarian, Varsha Keelara and Jo M. Holt

Domain interactions of yeast TBP.

Claire A. Adams and Michael G. Fried

Structure and mechanism of calmodulin binding to a cell surface targeting region of the N-methyl-D-aspartate receptor.

Zeynep Akyol, Johannes W. Hell and Madeline A. Shea

A method for jointly fitting fluorescence and absorbance AUC data.

P. Holland Alday, Eric George, David Brown, Tom Laue, Walter Stafford, Peter Sherwood and John J. Correia

5' coding region cis-acting determinants of bop-gene expression: secondary structure. Michael Arent and George J. Turner

Calorimetric characterization of a TCR-peptide/MHC interaction. Kathryn M. Armstrong and Brian M. Baker

UV resonance Raman studies of polyproline II structure in isotope edited alanine peptides.

Wendy Barber-Armstrong, Sean Decatur and Ishita Mukerji

Increased immunogenicity of an anchor-modified tumor-associated antigen is due to the enhanced stability of the peptide/MHC complex: implications for vaccine design.

Tiffany K. Baxter, Oleg Y. Borbulevych, Jeffrey M. Carney, Paul Helquist, Nicholas P. Restifo and Brian M. Baker

Thermodynamics and kinetics of the NH- κ B/I κ B α and NF- κ B/DNA interactions. Simon Bergqvist, Carries Hughes, Thomas Huxford, Gourisankar Ghosh and Elizabeth Komives

Effect of surfactant properties on the stability of the solubilized outer membrane protein ompX.

C. L. Bianco, E. W. Kaler, A. M. Lenhoff and C. R. Robinson

Transition state thermodynamics of the low- to physiological-temperature nondenaturational conformational change of bovine adenosine deaminase by slow scan rate differential scanning calorimetry.

Melissa A. Bodnar and B. Mark Britt

Unfolding dynamics of lima bean trypsin inhibitor by differential scanning calorimetry. *Melissa A. Bodnar and B. Mark Britt*

Synthetic porphyrazines: a new class of DNA binding molecules.

Melanie Bozza, Carlos Ramirez and Richard D. Sheardy

Determining the mechanism of allosteric regulation of NikR binding to DNA activated by Ni²⁺ binding.

Michael Bradley, Peter Chivers and Nathan Baker

Outer membrane protein F demonstrates strong self-association.

Nancy K. Burgess and Karen Fleming

Nonlinear regression fittiing algorithm for multiple independent binding equilibria, Monte Carlo error analysis, and Saroff distributions.

R. Buscaglia, M. W. Freyer and E. A. Lewis

pKa values of internal ioniazable residues in ribonuclease H.

Brian Cannon and Bertrand García-Moreno

Oligoproline effects on the structural properties of polyglutamine.

Veronique Chellgren, Brian Chellgren, Ronald Wetzel and Trevor Creamer

SNF: a bifunctional RNA binding protein.

Alan Chen and Kathleen Hall

Molecular determinants of pKa values: role of backbone flexibility and of short-range interactions in networks of polar and ionizable residues.

Michael S. Chimenti, Kelli Baran, Carolyn Fitch, Jamie Schlessman, Katie Herbst and Bertrand García-Moreno

Testing functional selection in TCR recognition of ligand.

John R. Clemens, Rebecca L. Davis-Harrison, Susan J. Gagnon, William E. Biddison and Brian M. Baker

A computational analysis of progesterone receptor-DNA binding isotherms: insight into transcriptional regulation.

K. D. Connaghan-Jones, A. F. Heneghan, M. T. Miura and D. L. Bain

Biophysical investigation of the TCR-peptide/MHC interaction: a comparative study of two receptors that bind a common ligand.

Rebecca L. Davis-Harrison and Brian M. Baker

Structural, thermodynamic, and kinetic comparisons of the binding of full-length yTBP and C-terminal core TBP with DNA.

Roberto Delgadillo, JoDell Whittington, Michael Brenowitz and Larry Parkhurst

Dissimilar "glutamate effects" on DNA binding by the type I DNA polymerases from *E. coli* and *T. aquaticus*.

Daniel J. Deredge, Gregory S. Thompson, Carmen R. Ruiz, Ke Jiang, Clayton F. Runfalo and Vince J. LiCata

DNA binding properties and activation of the ATPase activity of adenoassociated virus Rep40 helicase.

John David Dignam, Susan S. Dignam, Roy F. Collaco, Patrick Needham and James P. Trempe

Lanthanide spectroscopic studies of Mg(II)-dependent *Pvu*II restriction endonuclease. *Cynthia M. Dupureur, Lori M. Bowen, Gilles Muller and James P. Riehl*

Transmembrane domain interactions contribute to Erythropoietin Receptor activation. Alex Ebie and Karen G. Fleming

Structure and stability of Sso10b2, a hyperthermophile RNA and DNA binding protein. Stephen P. Edmondson, Andrew T. Clark, Tracy L. Armstrong and John W. Shriver

Chain length dependence of *de novo* tailspike folding. *Michael S. Evans, Mary C. Finn and Patricia L. Clark*

Molecular basis of CaMKII regulation by calmodulin: calcium-dependence of kinase binding.

T. Idil Apak Evans and Madeline A. Shea

Interactions of the anti-psychotic drug trifluoperazine with calmodulin and its effects on Ca²⁺ binding affinity.

Michael D. Feldkamp, Susan E. O'Donnell and Madeline A. Shea

Differentiating the ligand's chemical requirements for allosteric interactions from those for protein binding; phenylalanine inhibitionof pyruvate kinase.

Aron W. Fenton and Gissel McDonald

Thermodynamics of paperclip DNA triplexes.

Daniel P. Flaherty and Luis A. Marky

Investigating the heat capacity effect associated with RNA processes. Leslie Frank, Brian Doctrow and David Draper Unfolding thermodynamics of a DNA decamer duplex containing dG→deaza-dG substitutions.

Manjori Ganguly, Luis A. Marky and Barry I. Gold

Predicting interactions between cytoskeletal proteins. Shatadal Ghosh and David Sept

Increasing protein stability through rational redesign of surface charge-charge interactions.

Alexey V. Gribenko, Samantha S. Strickler, Alexander V. Gribenko, Timothy R. Keiffer and George I Makhatadze

Evidence for an evolutionary conserved DNA-recognition code at work in the Cro protein family of transcription factors.

Branwen M. Hall, Kelly R. LeFevre and Matthew H. J. Cordes

Osmotic pressure probes interactions in lipid-DNA complexes.

Daniel Harries, Horia I. Petrache, V. Adrian Parsegian, Dganit Danino and Ellina Kesselman

Quantitative analysis of progesterone receptor function: role of salt and salt-type in regulating self-association.

Aaron F. Heneghan, Keith Connaghan-Jones, Michael T. Miura and David L. Bain

Site-specific mutagenesis of the rat β -parvalbumin CD site.

Michael T. Henzl and Kelly Ndubuka

Protein stabilization and the effect of mixed osmolyte solutions: an analysis of the intrinsic stabilizing ability of osmolytes.

Luis Marcelo F. Holthauzen and D. W. Bolen

Electrostatic side-chain couplings in the gating of the OmpA ion channel suggest a mechanism for pore opening.

Heedeok Hong, Gabor Szabo and Lukas K. Tamm

A general and efficient way of expressing peptide for NMR studies of class I MHC/peptide complexes.

Francis K. Insaidoo and Brian M. Baker

Poster Session II: List of Presentations (J - Z)

Dynamics of *E. coli* RecA filament formation on ssDNA tail of a ssDNA-dsDNA junction. *C. Joo, M. Nakamura, S. A. McKinney, I. Rasnik and T. Ha*

Effects of DNA structure DNA binding of O^6 -alkylguanine-DNA alkyltransferase (AGT). Sambit R. Kat, Anthony E. Pegg and Michael G. Fried

Molecular dynamics simulations of mixed lipid bilayers. Stephen P. Kelty, Roger McMullen and George Turner

Unfolding of intramolecular triplexes with exclusive TAT base-triplets. *Irine Khutsishvili, Ronald Shikiya and Luis A. Marky*

Electrostatic effects in repeat proteins: salt dependence of unfolding of YopM, a leucine-rich-repeat protein.

Ellen Kloss and Doug Barrick

Spectroscopic and molecular dynamics evidence for a sequential mechanism for the DNA A→B transition.

Kelly M. Knee, Colin E. Aitken, Sergei Y. Ponomarev, Surjit Dixit, David L. Beveridge and Ishita Mukerji

Effects of monovalent salt on the observed enthalpy and heat capacity change for *E. coli* SSB binding to ssDNA.

Alexander G. Kozlov and Timothy M. Lohman

Study of DNA-Sfi I complex stability using AFM force spectroscopy.

Alexey V. Krasnoslobodtsev, Luda S. Shlyakhtenko, Egor V. Ukraintsev and Yuri L. Lyubchenko

Thermodynamics of yeast (*Saccharomyces cerevisiae*) replication protein A binding to single-stranded DNA.

S. Kumaran, Alexander G. Kozlov and Timothy M. Lohman

In vitro targeting of DNA triplexes with oligonucleotides. Hui-Ting Lee, Ronald Shikiya and Luis A. Marky

Structure of the Sfil-DNA complexes analyzed by atomic force microscopy. Alexander Y. Lushnikov, Vladimir N. Potaman, Elena A. Oussatcheva, Richard R. Sinden and Yuri L. Lyubchenko Communications between distant sites in envelope protein domain III of neutralization-resistant mutants of West Nile virus.

Rodrigo Maillard, Matthew Jordan and J. Ching Lee

Assembly of λ terminase protomers into a viral DNA processing and packaging machine.

N. Karl Maluf, H. Gaussier, M. Feiss and C. E. Catalano

Predicting the binding location of phospholipids on capping protein. *Michelle McCully and David Sept*

The effect of cations on the structure and function of NikR proteins. Alyssa A. Meheen, Erin L. Benanti and Peter T. Chivers

Allosterism in the binding of AGT to short single-stranded DNAs. *Manana Melikishvili and Michael G. Fried*

The effect of changing the detergent concentration on the free energy of transmembrane helix-helix association: Wyman linkage or dilution? *Naveen Michaud-Agrawal, Thomas B. Woolf and Karen G. Fleming*

The effect of dinitroanilines on parasitic protozoan tubulin dynamics. Arpita Mitra and David Sept

The biotin repressor: complexibility in the function of a flexible loop in an allosteric response.

Saranga Naganathan and Dorothy Beckett

Pulse proteolysis: a simple method for quantitative determination of protein stability and ligand binding.

Chiwook Park and Susan Margusee

Measured depletion of ions at the biomembrane interface: implicationson intermembrane interactions.

Horia I. Petrache, Daniel Harries and V. Adrian Parsegian

Unfolding thermodynamics of DNA duplex-hairpin motifs.

Karim P. Ramos, Ronald Shikiya and Luis A. Marky

Thermal stability landscape for Klenow DNA polymerase as a function of pH and salt concentration.

Allison J. Richard, Alexandra L. Klinger, Matthew J. Todd and Vince J. LiCata

Pressure-modulation temperature-scanning calorimetry. *Jörg Rösgen, Kristian Boehm and Hans-Jürgen Hinz*

Osmolyte dependence of biochemical reactions quantified by the phase diagram method.

Jörg Rösgen, Allan C. Ferreon, Josephine Ferreon and David Wayne Bolen

Allosteric mechanism of the EPD mutation in BK_{Ca} channel gating. *Akansha Saxena and David Sept*

A multiscale model for actin filaments. David Sept and Aravind Rammohan

Thermal unfolding of the β -hairpin peptide trpzip-4. Werner W. Streicher and George I. Makhatadze

Determination of the ATP coupling stoichiometry during *E. coli* UvrD monomer ssDNA translocation.

Eric J. Tomko, Christopher J. Fischer and Timothy Lohman

RNA-binding domain of human U1A: expression, purification and characterization of the wild type and engineered variant.

Jessica M. Tomlinson, Samantha S. Strickler and George I. Makhatadze

5' coding region cis-acting determinants of *bop* gene expression: primary structure. *George J. Turner and Michael Arent*

Evaluating the conformational flexibility of stalled nascent chains by fluorescence anisotropy.

Krastyu G. Ugrinov, David A. Johnson and Patricia L. Clark

Conformation-dependent interprotein interactions studied by AFM force spectroscopy. *Egor V. Ukraintsev, Tatiana O. Zaikova, John F. W. Keana and Yuri L. Lyubchenko*

Characterization of HU-cruciform DNA interaction. *Iulia Vitoc and Ishita Mukerji.*

The importance of solvent detail in modeling biomolecular solvation. Jason A. Wagoner and Nathan A. Baker

What role does the denatured sub-ensemble play in determining the native fold of a protein?

Suwei Wang, Steven T. Whitten, Tong Liu and Vincent J. Hilser