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The 27th Gibbs Conference on Biothermodynamics

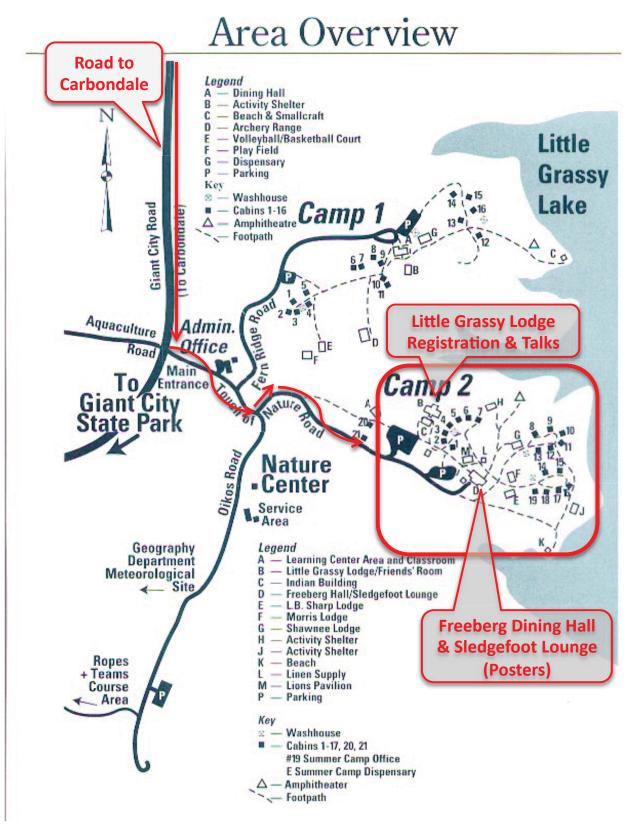


2013 Gibbs Conference	on
Biothermodynamics	

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Map of Touch of Nature Environmental Center

Most of the Gibbs Conference activities will be held in "Camp 2" as shown in the map below. Cell phone reception is extremely limited; parking lots are popular places for making calls.



The Gibbs Conference on Biothermodynamics

History

Fall, 1986

Discussion of the discipline: Thermodynamics in Biological Systems At the Gill residence in Vail, Colorado Gary Ackers, Wayne Bolen, Ernesto Freire, Stan Gill, Jim Lee

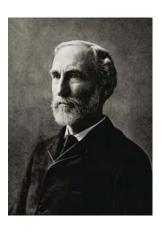
February, 1987

Discussion of the discipline: Thermodynamics in Biological Systems
The Gumbo Shop, New Orleans, LA during the 31st annual Biophysical Society
Meeting - Gary Ackers, Norma Allewell, Wayne Bolen, Ken Breslauer, Ken Dill,
Ernesto Freire, Stan Gill, Jim Lee

A history of the first ten years of the meeting was provided by Ackers GK and Bolen DW, The Gibbs Conference on Biothermodynamics: Origins and Evolution. *Biophysical Chemistry* **64** (1997) 3-5 (doi:10.1016/S0301-4622(96)02246-6)

An update is provided by Shea, MA, Correia, JJ, and Brenowitz, MD Introduction: Twenty five years of the Gibbs Conference on Biothermodynamics *Biophysical Chemistry* **159** (2011) 1-5 (doi:10.1016/j.bpc.2011.07.002)





Meetings

All meetings have been held at the Touch of Nature Environmental Center associated with Southern Illinois University – Carbondale. From 1987 through 1993, all of the speakers in the scientific sessions were students or postdoctoral fellows.

1987	Organizers: Jim Lee and Wayne Bolen Philosophical Talks: Gary K. Ackers and Ken Dill
1988	Organizers: Gary Ackers and Michael Johnson
1989	Organizers: Susan G. Frasier and Michael Johnson
1990	Organizers: Michael Johnson and Marty Straume
1991	Organizers: Gary Ackers and Tim Lohman Keynote Speaker: Ernesto Freire
1992	Organizers: Jim Lee and Tomasz Heyduk Keynote Speakers: Serge Timasheff and John Schellman
1993	Organizers: Maurice Eftink and Glen Ramsay Keynote Speakers: Peter von Hippel and Julian Sturtevant
1994	Organizers: Enrico Di Cera and Madeline Shea Keynote Speakers: Gary Ackers and Kathleen S. Matthews
1995	Organizers: Kenneth P. Murphy and Michael D. Brenowitz Keynote Speakers: Victor Bloomfield and Mario Amzel

1996	Organizers: Jonathan B. Chaires and Michael L. Doyle Keynote Speakers: J. Michael Schurr and Allen Minton
1997	Organizers: Dorothy Beckett and Jack Correia Keynote Speaker: Adrian Parsegian
1998	Organizer: Andy Robertson Keynote Speaker: David Draper
1999	Organizers: Bertrand Garcia-Moreno E. and John Shriver Keynote Speakers: Wayne Bolen and Gary Ackers
2000	Organizers: George Turner and Kim Sharp Keynote Speaker: Steve White
2001	Organizers: Margaret A. Daugherty and Luis A. Marky Keynote Speaker: George Rose
2002	Organizers: Michael Mossing and George Makhatadze Keynote Speaker: Rodney Biltonen
2003	Organizers: Vince Hilser and Dick Sheardy Keynote Speaker: Jim Lee
2004	Organizers: Doug Barrick and Kathleen Hall Keynote Speaker: Nacho Tinoco
2005	Organizers: Trevor Creamer and Clay Clark Keynote Speaker: Carl Frieden
2006	Organizers: Karen Fleming and Rohit Pappu Keynote Speakers: Madeline A. Shea and Timothy Lohman
2007	Organizers: Brian M. Baker and Michael T. Henzl Keynote Speaker: Jamie Williamson
2008	Organizers: Jannette Carey and David Bain Keynote Speakers: Dorothy Beckett and Ken Dill
2009	Organizers: Nathan Baker and Liskin Swint-Kruse Keynote Speaker: Linda Jen-Jacobson The Gary K. Ackers Lecture in Biothermodynamics: Michael Brenowitz
2010	Organizers: Elisar Barbar and Vince LiCata Keynote Speaker: C. Nick Pace The Gary K. Ackers Lecture in Biothermodynamics: Timothy Lohman
2011	Organizers: Gibbs Society of Board of Directors Keynote Speaker: Bertrand Garcia-Moreno E. The Gary K. Ackers Lecture in Biothermodynamics: Madeline Shea Editors of Special Issue of <i>Biophysical Chemistry</i> – Enrico Di Cera, Tim Lohman, Jack Correia
2012	Organizers: Aaron L. Lucius and Patricia L. Clark Keynote Speaker: Terry G. Oas The Gary K. Ackers Lecture in Biothermodynamics: Enrico Di Cera
2013	Organizers: James L. Cole and Aron W. Fenton Keynote Speaker: Doug Barrick The Gary K. Ackers Lecture in Biothermodynamics: Bertrand Garcia-Moreno E.

Gibbs Society Governance

Incorporation

In 2002, the *Gibbs Society of Biological Thermodynamics* incorporated in the Commonwealth of Virginia, under the guidance of Michael L. Johnson, then Treasurer of the Society and originator of the Society website. Articles of Incorporation and By Laws are available here: http://www.jhu.edu/~gibbs.

Current Officers

- President: David Bain, 2012-2013
- ❖ Vice President: Michael L. Johnson, 2011 2013
- Secretary: Margaret A. Daugherty, 2004 2013
- ❖ Treasurer: John J. Correia, March 2011 October, 2016

Board of Directors, listed alphabetically

- David Bain, President
- Douglas Barrick, Past President
- John J. Correia, Treasurer
- Margaret Daugherty, Secretary
- Michael L. Johnson, Vice President
- George Makhatadze, President Elect
- Madeline Shea

Past Presidents

2001-2002	Gary Ackers
2002-2003	Jack Correia
2003-2004	D. Wayne Bolen
2004-2005	Madeline Shea
2005-2006	Dorothy Beckett
2006-2007	Jonathan (Brad) Chaires
2007-2008	Tim Lohman
2008-2009	Luis Marky
2009-2010	Bertrand Garcia-Moreno E.
2010-2011	Karen Fleming
2011-2012	Doug Barrick

Past Treasurer

2001-2011 Michael L. Johnson

Committees & Other Contributions

Ackers Lecturer Selection Committee – Micheal Brenowitz, Chair Gibbs Society Website Hosting – Karen Fleming (2010 -) GoogleDocs Application/Registration & PayPal – Nathan Baker and Jack Correia Mailing List – Margaret Daugherty Saturday Night Thermo Organizers – Liskin Swint-Kruse and Vincent J. LiCata

With thanks to Alan Teska at the Touch of Nature Conference Center

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5th Annual Gary K. Ackers Lecture

2013 Lecturer – Bertrand Garcia-Moreno E., Department of Biophysics, Johns Hopkins University

This lecture honors the scientific contributions of Gary K. Ackers (1939-2011) to the field of Biological Thermodynamics. He served on the faculty at the University of Virginia, Johns Hopkins University and the Washington University School of Medicine. He was a Fellow of the Biophysical Society and was one of the founding organizers of the Gibbs Conference.

Gary demonstrated a lifelong commitment to the growth and development of an intellectual community of scholars devoted to furthering the field of biothermodynamics. Gary was an active member of the Biophysical Society throughout his career and served as President of the Society, as well as Organizer of the annual meeting. While on the faculty of the University of Virginia, he was a leader in the graduate biophysics training program. When on the faculty in the Department of Biology at the Johns Hopkins University, he conceived and organized the *Institute for Biophysical Studies of Macromolecular Assemblies*, a university-wide training program in molecular biophysics that has continued for decades. While at Johns Hopkins, he also played a leading role in the establishment of the Gibbs Conference on Biothermodynamics, an annual meeting organized to promote innovative development of biophysical principles applied to current problems in biology and to train the next generation of molecular biophysicists to tackle hard problems rigorously. After moving to St. Louis to chair the Department of Biochemistry and Molecular Biophysics at Washington University, he spearheaded a new graduate program in biophysics and hired many faculty who have joined the community of regular contributors to the Gibbs Conference.

Gary was a pioneer in the development of methods and application of principles of equilibrium thermodynamics to the study of linkage in complex macromolecular assemblies. Studies from his laboratory on the energetics of self-association and ligand binding in human hemoglobin proved unequivocally that the classic and elegant MWC model of intersubunit allostery was insufficient to explain cooperative oxygen binding: the position, as well as the number, of ligands matters. His contributions in this area greatly enhanced our understanding of the relationship between structure, energy and function in hemoglobin, and in multimeric allosteric systems in general. By probing ever more deeply into the molecular mechanism of cooperativity, he demonstrated a beautiful, useful, and general strategy for dissecting functional energetics in macromolecular assemblies.

His quantitative study of the interactions between proteins and nucleic acids in the bacteriophage lambda system included the development of quantitative DNase footprinting methods for measuring free energies of repressor-operator interactions. The footprinting assay remains an effective tool for measuring the extremely tight binding constants that are often encountered in site-specific interactions between proteins and nucleic acids. Those studies paved the way for similar methods to study protein-nucleic acid interactions in more complex systems, including time-resolved studies of the kinetics of RNA folding. Based on his experimental studies of phage lambda, his group developed statistical thermodynamic models to simulate the lysogenic-to-lytic growth switch: the series of macromolecular events that determine the fate of bacteriophage lambda during infection of *E. coli*. This work demonstrated how a complex biological function could be predicted quantitatively, strictly from the kinetics of transcription and translation, and the Gibbs free energy of interactions between the key macromolecular components in the genetic switch.

During Gary's early career, he developed methods to measure association constants in self-associating systems based on analytical gel permeation chromatography. Those methods have since become standard tools in the field. His group was also responsible for modifications of the cryo-gel electrophoresis methods, moving from applying them to hemoglobin to protein-DNA interactions. These contributions focused on developing the capacity to quantify intermediate states that are only transiently populated during the course of a biochemical process. His more than 200 articles and chapters changed our view of the molecular mechanisms that govern complex biochemical reactions.

△Gibbs₂₇ • Saturday Evening • October 5, 2013

4:00 – 10:00 pm Check-in at Little Grassy Lodge

7:30 – 10:00 pm Open Reception in Indian Lodge – Light refreshments, beer, wine and soft drinks

Participants are expected to make dinner arrangements independently

Gibbs Mugs Bazaar – pick up mugs purchased at registration

Saturday night thermo – Event for trainees only

Faculty Organizers

Vince LiCata, Louisiana State University & Liskin Swint-Kruse, University of Kansas Medical Center

Trainee Moderators: Jaycob Warfel, LiCata Lab, Louisiana State University

Kate Henderson, Emerson Lab, Mississippi State University

5:30 pm Freeberg Hall – Dinner for trainees who registered in advance

6:00 – 7:00 pm Flash Talks (Poster Introductions) – Session open to all trainees

 The annexin-lipid party line: Communication between protein and membrane

Anika Rannikko, Hinderliter Lab, University of Minnesota - Duluth

- Structural and dynamic features underlie the switch of ligand binding specificity in a Tiam1 PDZ domain mutant Xu Liu, Fuentes Lab, University of Iowa
- 3. Cyclipostin derivatives as inhibitors of rat hormone sensitive lipase Elena Vasilieva, Dupureur Lab, University of Missouri St. Louis
- 4. **Domain interactions in VA I RNA mediate high-affinity PKR binding** Katherine Launer-Felty, Cole Lab, University of Connecticut
- 5. **Distinctive solvation patterns make renal osmolytes diverse** Prem Sinha, Rösgen Lab, Penn State College of Medicine
- 6. Are proton binding reactions in proteins coupled to local conformational fluctuations?

Daniel Richman, Garcia-Moreno Lab, Johns Hopkins University

- Calmodulin recognition of anti-parallel nested binding sites in the neuronal voltage-dependent sodium channel Na_V1.2 Mark Miller. Shea Lab. University of lowa
- 8. Contributions of alanine residues to the hydrodynamic radius of p53(1-93) Romel Perez, Whitten Lab, Texas State University

7:00 – 7:15 pm Refreshment break

7:15 – 8:15 pm Career panel – Session open to all trainees

Andrew Robertson, National Psoriasis Foundation

Ana-Maria Soto, Towson University

Catherine Royer, Rensselaer Polytechnic Institute

8:15 pm Adjourn to reception in Indian Lodge

Δ Gibbs₂₇ • Sunday Morning • October 6, 2013

Breakfast served in Freeberg Hall

7:00 - 8:15 am

Magramalagular Folding and Stability

Macromolecular Folding and Stability

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8:30 - 8:35 am	Welcome by David Bain, Gibbs Society President
Moderator:	Giselle Jacobson, Clark Lab, University of Notre Dame
8:35 - 8:50 am	Introduction to the 27 th Annual Gibbs Conference Keynote Speaker George Makhatadze, Rensselaer Polytechnic Institute
8:50 – 9:30 am	Keynote Lecture Measuring coupling energies in simple proteins and extrapolating to complicated proteins Doug Barrick, Johns Hopkins University
9:30 – 9:45 am	Capped G-quadruplexes: A model for a looped out motif flanked by duplex DNA Vu Le, Lewis Lab, Mississippi State University
9:45 - 10:00 am	Probing N-H N hydrogen bond "strain" in a conserved N-capping motif Matthew Preimesberger, Lecomte Lab, Johns Hopkins University
10:00 - 10:30 am	Break – Refreshments in Indian Lodge
10:30 – 11:00 am	Studying the folding of individual residues in a small protein Nicholas Fitzkee, Mississippi State University
11:00 – 11:15 am	Exchange of putrescine ²⁺ and Mg ²⁺ in the ion atmosphere of RNA: Contributions to folding and gene regulation Robert Trachman, Draper Lab, Johns Hopkins University
11:15 – 11:45 am	Tug-of-war between thermodynamic stability and actin-binding function of tandem calponin-homology domains Krishna Mallela, University of Colorado Anschutz Medical Campus
11:45 am	Conference photo near Freeberg Hall
11:50 am	Lunch in Freeberg Hall

Free Time until Late Afternoon Session

Information about local parks & attractions is available near the entrance to Little Grassy Lodge.

∆Gibbs₂₇ • Sunday Afternoon • October 6, 2013

Nucleic Acids and Their Binding Partners

Moderator:	Irine Khutsishvili, Marky Lab, University of Nebraska Medical Center
3:00 - 3:30 pm	Structured mRNA induces the ribosome into a hyper-rotated state Peter Cornish, University of Missouri
3:30 – 3:45 pm	How do the different DNA-binding modes of <i>E. coli</i> SSB affect function? Vincent Waldman, Lohman Lab, Washington University
3:45 – 4:15 pm	Looking at multicomponent systems with small angle x-ray scattering: lons around DNA; DNA around proteins Lois Pollack, Cornell University
4:15 - 4:35 pm	Break – Refreshments in Indian Lodge
4:35 – 4:50 pm	Binding and salt linkages of HIV-1 reverse transcriptase with different DNA and RNA substrates reflect polymerization versus RNase H compatible modes of interaction Hiromi Brown, LiCata Lab, Louisiana State University
4:50 – 5:20 pm	DNA binding behavior of sequence specific polyamides Cynthia Dupureur, University of Missouri - St Louis
5:20 – 5:35 pm	Yeast Rap1 interacts with telomeric sequences with multiple binding modes Erik Feldmann, Galletto Lab, Washington University
5:35 - 6:05 pm	Substrate recognition by human O6-alkylguanine DNA alkyltransferase Michael Fried, University of Kentucky
6:05 - 6:15 pm	General Discussion
6:15 pm	Dinner in Freeberg Hall

∆Gibbs₂₇ • Sunday Evening • October 6, 2013

8:00 - 10:00 pm Poster Session I in Sledgefoot (lower level) & Freeberg (upper level)

Presenters with last name A to L

Please remove posters before midnight to make room for Monday presenters.

Sponsors displays in Freeberg (upper level) – near beer, wine and soft drinks

∆Gibbs₂₇ • Monday Morning • October 7, 2013

Posters to be presented on Monday night may be mounted as soon as space is available on Sunday night.

Airport Ride Board will be available in Little Grassy Lodge, near check-in window

7:00 - 8:15 am **Breakfast served in Freeberg Hall**

Structure / Thermodynamic Correlations

8:30 - 8:35 am	Announcements by organizers
Moderator:	Keith Connaghan, Bain Lab, University of Colorado Anschutz Medical Campus
8:35 – 8:50 am	Introduction to the Gary K. Ackers Lecture in Biothermodynamics David Bain, University of Colorado Anschutz Medical Campus
8:50 - 9:30 am	5th Annual Gary K. Ackers Lecture pH-Dependent processes in proteins Bertrand Garcia-Moreno E., Johns Hopkins University
9:30 - 9:45 am	Impact of pH on the structure and function of neural cadherin Matt Dukes, Pedigo Lab, University of Mississippi
9:45 - 10:00 am	Revisiting thermodynamic cooperativity: Studying TRBP interactions with double stranded RNA Roderico Acevedo, Showalter Lab, Pennsylvania State University
10:00 - 10:20 am	Break – Refreshments in Indian Lodge
10:20 - 10:50 am	Molecular switches in signaling and disease Linda Nicholson, Cornell University
	Molecular switches in signaling and disease
10:50 – 11:05 am	Molecular switches in signaling and disease Linda Nicholson, Cornell University Thermodynamics of cationic and anionic surfactant interaction
10:50 – 11:05 am	Molecular switches in signaling and disease Linda Nicholson, Cornell University Thermodynamics of cationic and anionic surfactant interaction Vytautas Petrauskas, Matulis Lab, Vilnius University Structure, dynamics and inhibition of Tiam1 PDZ domain complexes
10:50 - 11:05 am 11:05 - 11:35 am	Molecular switches in signaling and disease Linda Nicholson, Cornell University Thermodynamics of cationic and anionic surfactant interaction Vytautas Petrauskas, Matulis Lab, Vilnius University Structure, dynamics and inhibition of Tiam1 PDZ domain complexes Ernesto Fuentes, University of Iowa

Trainee funding panel

2:00 - 3:00 pm Discussion of trainee fellowship applications - open to all attendees

Trevor Creamer, University of Kentucky

Sarah Bondos, Texas A&M Health Science Center

Dorothy Beckett, University of Maryland

∆Gibbs₂₇ • Monday Afternoon • October 7, 2013

Macromolecular Interactions

Moderator:	Clarissa Weaver, Lucius Lab, University of Alabama at Birmingham
3:15 – 3:45 pm	Transition metal homeostasis: Allosteric switching by metal sensor proteins David Giedroc, Indiana University
3:45 – 4:00 pm	Heterotropic linkage between protein-protein and protein RNA interactions determines protein partitioning in the snRNPs Sandra Williams, Hall Lab, Washington University
4:00 – 4:30 pm	Engineering linked-equilibria into protein interfaces to allow regulation of protein interactions James Horn, Northern Illinois University
4:30 - 4:45 pm	Break – Refreshments in Indian Lodge
4:45 – 5:15 pm	Biophysical studies on the A1 and A3 domain homologues of von Willebrand factor Matthew Auton, Mayo Clinic
5:15 – 5:45 pm	Structure and function of a novel phosphopeptide binding domain from the transcriptional corepressor MINT
	Rhett Kovall, University of Cincinnati College of Medicine
5:45 - 6:00 pm	

∆Gibbs₂₇ • Monday Night • October 7, 2013

8:00 - 10:00 pm Poster Session II in Sledgefoot (lower level) & Freeberg (upper level)

Presenters with last name M to Z

Sponsors displays in Freeberg (upper level) – near beer, wine and soft drinks

\triangle Gibbs₂₇ • Tuesday Morning • October 8, 2013

Check-out: Please leave your room keys at the counter in the lobby of Little Grassy Lodge. Airport Ride Board will be available in Little Grassy Lodge, near the check-in window

7:30 – 8:30 am **Breakfast in Freeberg Hall**

Modeling and Computation

8:40 - 8:45 am	Closing announcements by organizers
Moderator:	Catie Shelton, Herr Lab, University of Cincinnati College of Medicine
8:45 - 9:15 am	Computation and calorimetry: What can we learn from each other? Michael Gilson, University of California-San Diego
9:15 - 9:30 am	The tertiary architecture of LacI/GalR proteins accommodates multiple networks of functionally important positions Daniel Parente, Swint-Kruse Lab, University of Kansas Medical Center
9:30 - 10:00 am	Temperature effects on the hydrodynamic radius of intrinsically disordered proteins Steven Whitten, Texas State University
10:00 – 10:20 am	Break – Refreshments in Indian Lodge
10:20 - 10:40 am	Rational design of two proteins with 80% sequence identity but different folds using sequence transplantation Lauren Porter, Potomac Affinity Proteins
10:40 - 10:55 am	Deletion of the connecting linker between kringle-1 and kringle-2 in prothrombin stabilizes the zymogen in a collapsed conformation Nicola Pozzi, Di Cera Lab, Saint Louis University
10:55 – 11:25 am	Molecular modeling of the ligand binding free energy and conformational rearrangements in AMPA type glutamate receptors Maria Kurnikova, Carnegie Mellon University
11:25 am	Box lunch available in Freeberg Hall
Check-out	Please leave your keys at the counter in Little Grassy Lodge