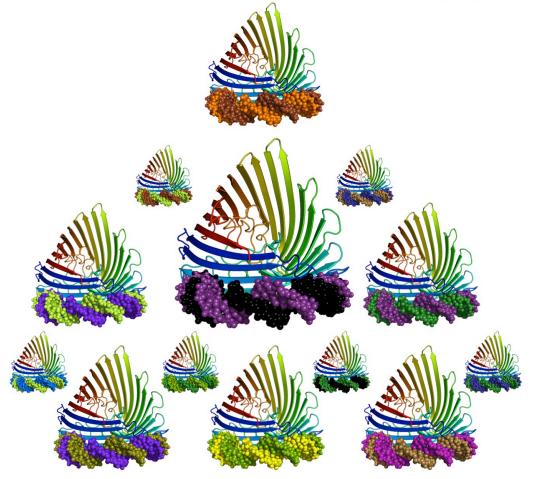
\triangle Gibbs₃₇



Prime Time Thermo

Organizers - Tonya Zeczycki & Krishna Mallela 37th Annual Gibbs Conference on Biothermodynamics Touch of Nature Outdoor Education Center • Carbondale, IL, USA October 14-17, 2023

Sponsors - Applied Photophysics, Biophysical Society, Daedalus Innovations, Genentech, Horiba Scientific, ISS, JASCO, LUMICKS, Malvern Panalytical, New England Biolabs, Nicoya, National Science Foundation, OLIS, Protein Society, Refeyn, Texas State University, University of Iowa

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Image Credits Front cover - Gibbs Delta Campfire logo & Ring of Fires by Madeline Shea (porin (3jty) fire and nucleosome DNA (6c0w) logs). The book includes images by Greg DeKoster (Gibbs campfires with the "37 cluster" of stars), Dan Herschlag (Keynote Lecture), Karen Fleming (Gary K. Ackers Lecture in Biothermodynamics) & Touch of Nature SIUC.

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Touch of Nature (ToN) Outdoor Education Center Southern Illinois University – Carbondale



Image courtesy of Touch of Nature

Driving Directions https://ton.siu.edu/about-us/

GPS Address: The easiest way to get to Touch of Nature is to use this address in a GPS navigation: **1206 Touch of Nature Road, Makanda, IL 62958**

From the north: Take Interstate 57 South to the Route 13 exit (Carbondale - Exit #54B), turn west (right) onto Route 13 toward Carbondale, go approximately 15 miles. Look for Wal-Mart on your right once you enter Carbondale. Turn left (south) at this light, Giant City Road. Continue 8 miles south on Giant City Road. Touch of Nature Road will be on your left (east). Turn left and continue straight on Touch of Nature Road. Administration building is the first left

From the south: Follow I-24 to I-57 North, or stay on I-57, to the Route 148 exit (Exit #45), head northwest on 148 for approximately 2 miles. Turn left on Grassy Road. There is a convenience store on this corner. Follow Grassy Road for about 7 miles. Stay left when you come to a Y. Continue southwest on Grassy Road. Grassy Road will come to a T at Giant City Road. Turn south (left) on Giant City Road. Continue on Giant City Road approximately 1 mile. Touch of Nature Road will be on your left. Turn left and continue straight on Touch of Nature Road. Administration building is the first left.

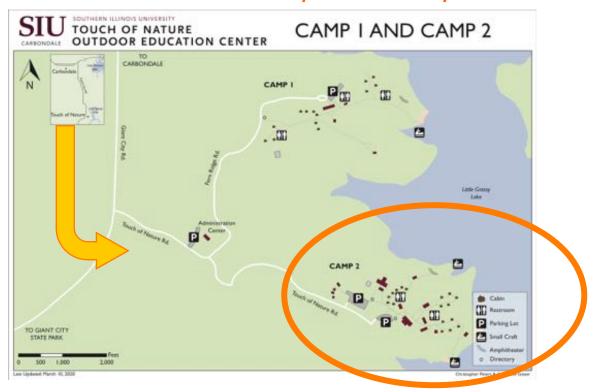
From the east: Head west on I-70 or, if you are further south, take I-64 to I-57 South. Continue approximately 50 miles south from I-64 to the Route 13 exit (Marion - Exit #54B), head west on Route 13 toward Carbondale, go approximately 15 miles. Look for Wal-Mart on your right once you enter Carbondale. Turn left (south) at this light, Giant City Road. Continue 8 miles south on Giant City Road. Touch of Nature Road will be on your left (east). Turn left and continue straight on Touch of Nature Road. Administration building is the first left.

From the west: Head east on I-64 approximately 50 miles from St. Louis. Take the Route 127 (Nashville) exit south. Continue south for approximately 50 miles to Murphysboro. At Murphysboro, go left (east) for approximately 10 miles. You will pass through the town of Carbondale. Watch for the mall on your right. When you see Wal-Mart on your left, this is your light. Turn right (south) at this light, Giant City Road. Continue 8 miles south on Giant City Road. Touch of Nature Road will be on your left (east). Turn left and continue straight on Touch of Nature Road. Administration building is the first left.

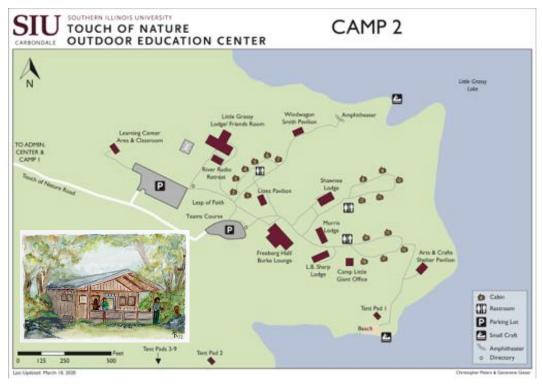
History https://ton.siu.edu/about-us/history.php https://news.siu.edu/2018/03/032718-special-olympics-history.php https://hee-journal.uni-koeln.de/sites/hee-journal/user_upload/Ritzel_2018.pdf

Map of TON Camps 1 & 2 with Carbondale Inset

Going south on Giant City Road, turn left onto Touch of Nature Rd. and proceed to Camp 2.



Camp 2 – Most activities will take place here.



Selected Touch of Nature Facilities

River Radio Retreat Building

Session Breaks and Business Meeting will be held here.



Image courtesy of Touch of Nature

Friends Room in Little Grassy Lodge

Saturday Night Thermo talks/panel and Platform Sessions will be held here.



Image courtesy of Touch of Nature

Lodging ranges from "dormitory-like" to rustic cabins



Image courtesy of Touch of Nature



Introduction History & Governance

History of the Gibbs Conference on Biothermodynamics

Fall, 1986

Discussion of the discipline:

Thermodynamics in Biological Systems.

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.

1997

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



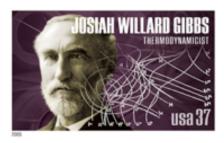


J. Willard Gibbs

2005

On May 4, 2005, the United States Postal Service issued a stamp in honor of J. Willard Gibbs (1839 – 1903) (see https://news.yale.edu/2005/04/27/u-s-postage-stamp-series-unveiling-yale-celebrate-exceptional-scientists).

More of his history may be found in "The greatest mind in American history" (https://yalealumnimagazine.org/articles/4496-josiah-willard-gibbs) and NAS online biographies (https://www.nasonline.org/publications/biographical-memoirs/memoir-pdfs/gibbs-josiah.pdf).



2011

An update on the history of the Gibbs conference was provided by Madeline A. Shea, John J. Correia and Michael D. Brenowitz in "Introduction: Twenty five years of the Gibbs Conference on Biothermodynamics" available in *Biophysical Chemistry* 159 (2011) 1-5 (doi:10.1016/j.bpc.2011.07.002).

Conference Organizers & Keynote Speakers

All in-person meetings have been held at the Touch of Nature Outdoor Education Center (prior to 2022, it was called the Touch of Nature Environmental Center) associated with Southern Illinois University–Carbondale.

From 1987 through 1993, all speakers in scientific sessions were students or postdoctoral fellows. PI speakers were introduced at the 8th conference held in 1994. In 2020 and 2021, the conference was held virtually (*via* Zoom) because of the COVID-19 pandemic. A list of Conference Organizers and Keynote Speakers follows.

Year	Conference Organizers	Keynote Speaker(s)
1987	Jim Lee and Wayne Bolen Philosophica	I Talks by Gary K. Ackers and Ken Dill
1988	Gary Ackers and Michael Johnson	•
1989	Susan G. Frasier and Michael Johnson	
1990	Michael Johnson and Marty Straume	
1991	Gary Ackers and Tim Lohman	Ernesto Freire
1992	Jim Lee and Tomasz Heyduk	Serge Timasheff and John Schellman
1993	Maurice Eftink and Glen Ramsay	Peter von Hippel and Julian Sturtevant
1994	Enrico Di Cera and Madeline Shea Gar	y K. Ackers and Kathleen S. Matthews
1995	Kenneth P. Murphy and Michael D. Brenow	ritz Victor Bloomfield and Mario Amzel
1996	Jonathan B. Chaires and Michael L. Doyle	J. Michael Schurr and Allen Minton
1997	Dorothy Beckett and Jack Correia	Adrian Parsegian
1998	Andy Robertson	David Draper
1999	Bertrand García-Moreno E. and John Shrive	
2000	George Turner and Kim Sharp	Steve White
2001	Margaret A. Daugherty and Luis A. Marky	George Rose
2002	Michael Mossing and George Makhatadze	
2003	Vince Hilser and Dick Sheardy	
2004	Doug Barrick and Kathleen Hall	
2005	Trevor Creamer and Clay Clark	
2006	Karen Fleming and Rohit V. PappuN	
2007	Brian M. Baker and Michael T. Henzl	
2008	Jannette Carey and David Bain	-
2009	Nathan Baker and Liskin Swint-Kruse	
2010	Elisar Barbar and Vince J. LiCata	
2011	Gibbs Society Board of Directors	
	Aaron L. Lucius and Patricia L. Clark	
2013	James L. Cole and Aron W. Fenton	3
2014	Andrew B. Herr and Steven T. Whitten	3
2015	Ernesto J. Fuentes and James R. Horn	
2016	Sarah Bondos and Nick Fitzkee	
2017	Scott Showalter and Ana Maria Soto	•
2018 2019	Chiwook Park and David Draper	
2019	Matthew Auton and Carlos Castañeda Roberto Galletto and Karen A. Lewis	-
2020	Gibbs Society Board of Directors	
2021	Aaron L. Lucius and James R. Horn	
2022	Tonya Zeczycki and Krishna Mallela	
2023	Tonya Z60Zyoni and Misilia Malicia	Daniei rieischlag

The Annual Gary K. Ackers Lecture in Biothermodynamics

Since 2009, the Gibbs Society Board of Directors sponsors an annual lecture honoring the scientific contributions of founding organizer Gary K. Ackers (1939-2011) to the field of biothermodynamics.

Lecture	Year	Ackers Lecturer
1	2009	Michael Brenowitz
2	2010	Timothy Lohman
3	2011	Madeline Shea
4	2012	Enrico Di Cera
5	2013	Bertrand García-Moreno E.
6	2014	David E. Draper
7	2015	Walter S. Englander
8	2016	Ken Dill
9	2017	Dorothy Beckett
10	2018	James C. Lee
11	2019	Jack Correia
12	2020	Douglas Barrick
13	2021	David Bain
14	2022	Tobin Sosnick
15	2023	Karen Fleming

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. Current bylaws are available online (https://www.gibbssociety.org/) and in this document. Articles of Incorporation may be requested from the Treasurer.

Current Officers and Terms

President	James R. Horn, 2022 – 2023
President-Elect	Sarah E. Bondos, 2022 – 2023
Past-President	Aaron L. Lucius, 2022 – 2023
Vice President	Aron Fenton, 2022 –
Secretary	Andrew Herr, 2019 – 2025
Treasurer	Karen Lewis, 2022 – 2027

Current Board Members-at-Large

Vince J. LiCata (2017) current term expires 2027 Madeline A. Shea (2006) current term expires 2024

2022 – 23 Board of Directors (listed alphabetically by last name)

Sarah E. Bondos, Andrew Herr, James R. Horn, Karen Lewis, Vince J. LiCata, Aaron L. Lucius, Madeline A. Shea

Former Presidents

2001 – 2002 2002 – 2003 2003 – 2004	Jack Correia
2004 – 2005 2005 – 2006	Madeline A. Shea
2006 – 2007 2007 – 2008	J. Brad Chaires
2008 – 2009	
2010 – 2011 2011 – 2012	Karen G. Fleming
2012 – 2013 2013 – 2014	David L. Bain
2014 – 2015 2015 – 2016	Patricia Clark
2016 – 2017 2017 – 2018	James Cole
2018 – 2019 2019 – 2020	Brian Baker
2020 – 2021 2021 – 2022	

Former Treasurers

2001 – 2011 Michael L. Johnson 2011 – 2017 Jack Correia 2017 – 2022 Aron Fenton

Former Secretaries

2004 – 2013 Margaret A. Daugherty 2013 – 2019 Liskin Swint-Kruse

Former Vice President

2010 - 2022 Michael L. Johnson

Former Board Members-at-Large

2003 - 2017 Jack Correia

Committees and Other Contributors to Conference Organization

Ackers Lecturer Selection Committee

Vince J. LiCata (Chair) 2021 – and current members of the Gibbs Society Board of Directors

Fundraising Committee:

Faculty - Ernesto Fuentes (Chair), Lisa Warner and Allyn Schoeffler

Local Arrangements Committee

Faculty - Carlos Castañeda (Chair), Emma Morrison, Aron Fenton Trainees – Whitney Bond, Jasmine Cubuk, Taylor Devlin, Jerry Dinan, Emery Usher

Poster Committee

Faculty - Steven Whitten (Chair) and Sean Fanning

Presentation Logistics Committee

Faculty - Kurt Piepenbrink (Chair), Sukesh Bhaumik, Elizabeth Duran, Stephen Fried Trainees - Pritam Chakraborty, Alexandra Lucas, Xinzhe Ren, Yingzi Xia, Yafan Yu

Program Book Committee

Madeline Shea (Chair), David Bain, Gregory DeKoster

Registration – JotForm and PayPal

Nicholas C. Fitzkee and James R. Horn

Saturday Night Thermo Co-Organizers

Faculty: Vince J. LiCata and Ana Maria Soto

Trainees: Govinda Hancock, Kacey Mersch, Kristen Young

SharePoint Guru

James R. Horn

Speaker Nominating Committee

Organizers Tonya Zeczycki and Krishna Mallela, Scott Showalter, Clarissa Durie

Webmasters for Gibbs Society - https://www.gibbssociety.org

Sarah E. Bondos and Chiwook Park (2020 – present)

Gibbs Mission and Policies

Mission Statement (approved January 20, 2023)

The Gibbs Society of Biological Thermodynamics is committed to:

- advancing the development and application of thermodynamic principles to understand biological systems;
- fostering the professional growth of early-career trainees;
- promoting an equitable, accessible, and inclusive biothermodynamics community;
 and
- hosting an annual conference that includes significant opportunities for trainee oral presentations.

During registration online, all participants have accepted the following policies of the Gibbs Conference on Biothermodynamics.

Confidentiality Statement

Please remember that the content of all presentations (both talks and posters) for this meeting are confidential material and may contain unpublished results. Abstracts will not be posted online.

Please ask permission from the authors before taking photos or screenshots of posters or poster material.

Please do not record or take screenshots of the talks, posters, or any presented material unless a speaker has given you express permission.

Code of Conduct

The Gibbs Society of Biological Thermodynamics is committed to providing a safe and productive environment that fosters open dialogue and the exchange of scientific ideas, promotes equal opportunities and treatment for all participants, and is free of harassment and discrimination.

Harassment includes speech or behavior that is not welcome or is personally offensive, whether it is based on ethnicity, gender, religion, age, body size, disability, veteran status, marital status, sexual orientation, gender identity, or any other reason. It includes stalking, unnecessary touching, and unwelcome attention.

Behavior that is acceptable to one person may not be acceptable to another, so use discretion to be sure that respect is communicated. Harassment intended in a joking manner still constitutes unacceptable behavior. Anyone experiencing conduct that violates this Code should report this conduct to any member of the Gibbs Society Board.

Bylaws to Articles of Incorporation for Gibbs Society of Biological Thermodynamics

January 20, 2023

Mission Statement (edited January 20, 2023)

The Gibbs Society of Biological Thermodynamics is committed to:

- advancing the development and application of thermodynamic principles to understand biological systems;
- fostering the professional growth of early-career trainees;
- promoting an equitable, accessible, and inclusive biothermodynamics community; and
- hosting an annual conference that includes significant opportunities for trainee oral presentations.

Bylaw 1 (established December 11, 2020; edited January 20, 2023):

Henceforth, the Board of Directors shall consist of seven (7) rotating members, comprising (1) the Past-President, (2) the current President, (3) the President-Elect, (4) Secretary (3-year renewable term), (5) Treasurer (5-year renewable term), and (6-7) two Members-at-Large (6-year renewable terms each, staggered 3 years apart). All board members will be listed publicly on the Gibbs Society of Biological Thermodynamics website.

The positions of advisory Vice President, Assistant Treasurer, and Webmaster will be appointed by the Board of Directors and publicly posted on the website. Vacancies and renewals on the Board of Directors will be filled by a majority vote of the Board of Directors. As needed, other positions may be created and appointed by the Board of Directors and publicly posted on the website. All members of the Board of Directors must have served as an Organizer of the Gibbs Conference on Biothermodynamics.

Bylaw 2 (established February 22, 2017; edited May 13, 2022):

Officers of the Gibbs Society of Biological Thermodynamics shall be elected at the annual meeting of the Board of Directors during the annual conference. A President-Elect shall be chosen at each annual meeting. At the close of the annual meeting, the President-Elect will become the current President and the current President will become the Past-President. The Secretary and Treasurer shall serve renewable 3-5-year terms. The Treasurer is authorized to invest society funds in checking accounts and interest-bearing accounts such as money market accounts, savings accounts and/or certificates of deposit. The Treasurer will inform the Board of Directors prior to transfers of Society funds. The Treasurer shall provide the Board of Directors with a financial report at least twice a year, 1) at the annual meeting of the Board of Directors and 2) after all expenses and income from the meeting have been resolved. Two members of the Board of Directors in addition to the Treasurer will be authorized to have access to the bank account.

Bylaw 3 (established February 22, 2017; edited May 16, 2022):

The Board of Directors shall provide oversight and serve as advisors to the current organizers of the Gibbs Conference on Biothermodynamics. It is the responsibility of the Board of Directors to provide documentation on how to organize a Gibbs Conference to the conference organizers. This includes a summary of financial reports from the treasurer; a list of email addresses from the secretary; and a check list of organizational tasks that will be revised in consultation with the organizers of the immediate past meeting.

Bylaw 4 (established May 13, 2022):

All donations to the Gibbs Society of Biological Thermodynamics, regardless of source, will be unrestricted, will be deposited in the general fund, and will be used at the discretion of the Board of Directors to support and improve the conference and other activities of the Gibbs Society of Biological Thermodynamics. All donors will be informed of this before their gifts are accepted, and donations will be acknowledged in the program book and on the website for the conference.

These Bylaws were discussed by email exchanges and approved by vote during a zoom meeting of the Board of Directors:

Madeline Shea, Long-Term Board Member*
Vince LiCata, Long-Term Board Member*
Aaron Lucius, Past-President
Jim Horn, President
Sarah Bondos, Past-President
Andrew Herr, Secretary
Karen Lewis, Treasurer

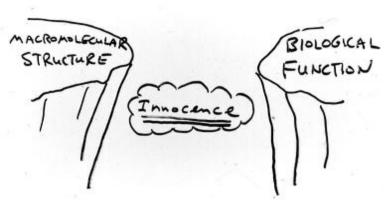
These bylaws replace those established June 3, 2022.

^{*}Following approval of these Bylaws on January 20, 2023, these "Long-Term Board Members" became "Members-at-Large".

15th Annual Gary K. Ackers Lecture in Biothermodynamics 2023 Lecturer – Karen Fleming, 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 Washington University in St. Louis 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.

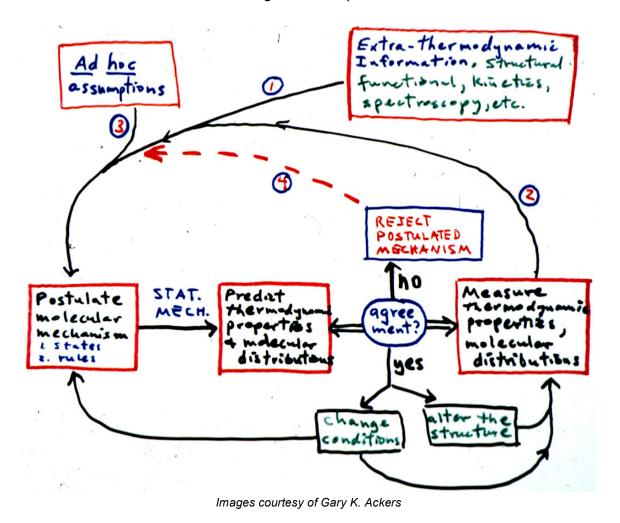


Gary was a pioneer in the development of methods and application of principles of equilibrium thermodynamics to the study of linkage in complex macromolecular assemblies. Studies from his laboratory on the energetics of self-association and ligand binding in human hemoglobin proved unequivocally that 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

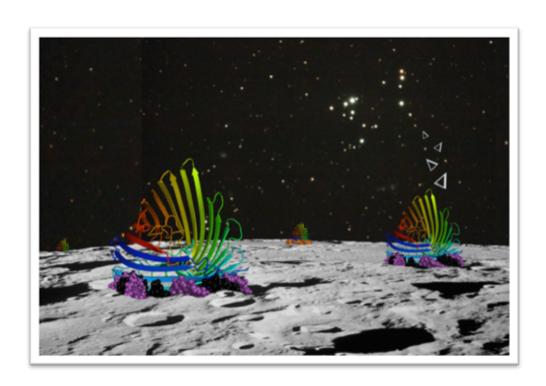
methods for measuring free energies of repressor-operator interactions. The footprinting assay remains an effective tool for measuring the extremely tight binding constants that are often encountered in site-specific interactions between proteins and nucleic acids. Those studies paved the way for similar 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.

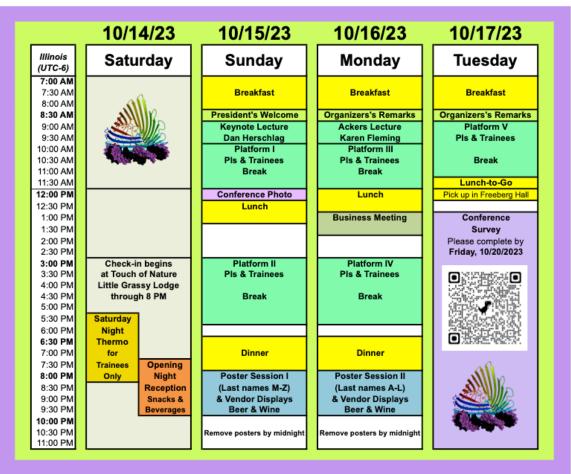




Schedule of Events October 14-17, 2023



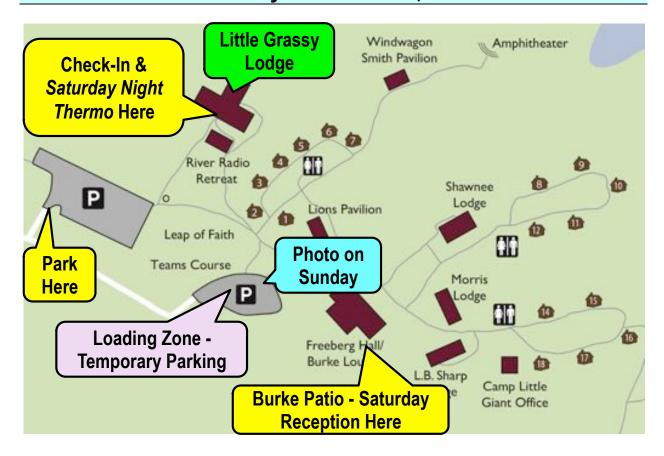
Δ Gibbs₃₇



Time listed in leftmost column indicates approximate starting time for event in that block.

Prime Time Thermo

Saturday ♦ October 14, 2023



3:00 – 8:00 pm Check-in at Little Grassy Lodge for those staying on-site.

Everyone - Collect name badge & program book in Little Grassy Lodge.

If you need to park in the Loading Zone near Freeberg to unload luggage or displays, please move cars **before noon on Sunday** to accommodate setup of the conference photo that will be held there.

Attractions in Southern Illinois (https://southernillinoistourism.org) include Giant City State Park orchards with apple cider, doughnuts, apple butter & pumpkins (Rendleman Orchards & Market and Flamm Orchards are popular), and plenty of small town charm in Makanda.

5:30 – 8:15 Saturday Night Thermo Event for trainees only – full schedule on next page

7:30 – 10:00 pm	Opening Reception on Burke Patio
	Light refreshments, beer, wine, and soft drinks will be served.
	Participants are expected to make dinner arrangements
	independently.

8:30 PM	Little Grassy Lodge "Friends Room"
	Meeting of Presentation Logistics Committee and
	Moderators of Platform Sessions to review AV setup.

Saturday Night Thermo ♦ October 14, 2023

Saturday Night Thermo - Event for trainees only

Moderators and Co-Organizers

Govinda Hancock, Fanning Lab, Loyola University Chicago Kacey Mersch, Lohman Lab, Washington University in St. Louis Kristen Young, Fanning Lab, Loyola University Chicago

Faculty Co-Organizers

Ana Maria Soto, Towson University Vince J. LiCata, Louisiana State University

5:30 pm	Dinner for trainees who registered in advance	
	Little Grassy Lodge "Friends Room"	

6:00 – 7:00 pm Flash Talks in Little Grassy Lodge "Friends Room" Open to trainees only. No registration required.

Speakers - please work with Moderators to test your presentation in advance.

- High-resolution models of conformational dynamics in DNA recognition by the master transcription factor PU.1 Tyler N. Vernon, Poon Lab, Georgia State University
- Disruptions of DNA Base Pair Cooperativity Probed with Time-Resolved Infrared Spectroscopy, Brennan Ashwood, Tokmakoff Lab, The University of Chicago
- 3. Insights into How Chaperone Net Charge and Sequence Patterning Influence Nucleic Acid Folding,

Gabrielle M. Perkins, Holmstrom Lab, University of Kansas

- 4. Molecular basis of ligand-dependent Nurr1-RXRalpha activation, **Xiaoyu Yu**, Kojetin Lab, Vanderbilt University
- The Role of Structural and Dynamic Rearrangements in the Functional Regulation of a Classically Allosteric Protein,
 Darex J. Vera-Rodríguez, Lee Lab, University of North Carolina at Chapel Hill
- 6. Engineering a Biomimetic Protein Corona for Predictive Nanoparticle Behavior **Tanveer Shaikh**, Fitzkee Lab, Mississippi State University
- Excluded Volume and Weak Interactions in Crowded Solutions
 Modulate Conformations and RNA Binding of an Intrinsically Disordered Tail
 Madison Stringer, Soranno Lab, Washington University in St Louis

7:00 – 7:15 pm Refreshment Break

7:15 – 8:15 pm Career Panel in Little Grassy Lodge "Friends Room" Session open to all trainees but not Pls. Sponsored by the Serge N. Timasheff Scholarship Donors

8:15 pm Adjourn to join ongoing reception for all participants.

Sunday Morning ♦ October 15, 2023

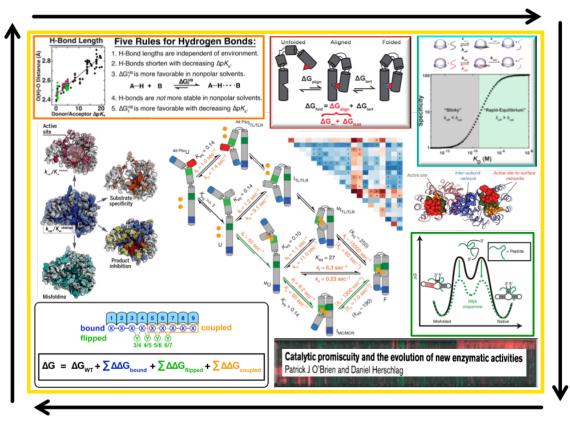
♦ Posters in Poster Session I may be mounted on Sunday morning. Use assigned number.

7:00 – 8:20 am **Breakfast in Freeberg & Burke & Surrounding Area** Individuals who requested special meals in advance because of dietary restrictions should follow signage in Freeberg to separate serving area, and ask staff for assistance.

Platform Session I

8:15 am	Speakers - Please connect your laptop in advance of the session. AV support provided by Yafan Yu, Piepenbrink Lab, Univ. of Nebraska
8:30 – 8:40 am	President's Welcome James R. Horn, Northern Illinois University
8:40 – 8:50 am	Organizers' Remarks Tonya Zeczycki, Brody School of Medicine, East Carolina University Krishna Mallela, Skaggs School of Pharmacy, Univ. of Colorado
Moderator	Pierce O'Neil, Swint-Kruse Lab, KUMC

woderator	Pierce O Neil, Swillt-Kluse Lab, Kolvic	
37 th Annual (37 th Annual Gibbs Conference Keynote Lecture	
8:50 – 9:00 am	Speaker Introduction by Tonya Zeczycki Brody School of Medicine, East Carolina University	
9:00 – 9:50 am	From Structure–Function to Ensemble–Function: A new paradigm for quantitative understanding of protein function Dan Herschlag, Stanford University	
9:50 – 10:00 am	Questions for Keynote Speaker	



Platform Session I - continued

10:00 – 10:30 am	Break – Refreshments in River Radio Retreat
	Please move cars in parking lot near Freeberg before noon to accommodate setup of the conference photo.
10:30 – 10:45 am 10:45 – 10:50 am	Identification of a covert evolutionary pathway between two protein folds Devlina Chakravarty, Porter Lab, NLM NCBI NIH Questions for speaker
10:50 – 11:15 am	The Genetic Landscape of a Metabolic Interaction
	Kimberly Reynolds, Univ. of Texas Southwestern Medical Center
11:15 – 11:20 am	Questions for speaker
11:20 – 11:35 am	Teasing More out of ITC Using System-Specific Bayesian Inference Douglas Walker, Oregon State University
11:35 – 11:40 am	Questions for speaker
11:40 – 11:50 am	General Discussion for Session
12:00 pm	Conference Photo near Freeberg Hall Photographer Karen Fleming Gather on hill near parking lot by Freeberg Hall Please move cars in parking lot before noon to accommodate setup of the conference photo.

12:15 pm	Lunch in Freeberg & Burke & Surrounding Tents
	Individuals who identified dietary restrictions should follow signage
	in Freeberg to separate serving area, and ask staff for assistance.

1:00 – 2:45 pm Free Time until Afternoon Session
Look near the check-in counter in Little Grassy Lodge for information about possible organized events. Information about local parks and attractions is also available near the entrance to Little Grassy Lodge.

Attractions in Southern Illinois (https://southernillinoistourism.org) include Giant City State Park, orchards with apple cider, doughnuts, apple butter & pumpkins (Rendleman Orchards & Market and Flamm Orchards are popular), and plenty of small town charm in Makanda.

Sunday Afternoon ♦ October 15, 2023

Platform Session II

2:45 pm	Speakers - Please connect your laptop in advance of the session. AV support provided by Alexandra Lucas , Mallela Lab, Univ. of Colorado Anschutz Medical Campus
Moderator	Taylor Devlin, Fleming Lab, Johns Hopkins
3:00 – 3:25 pm	Interactions of a riboswitch RNA with fluorogenic probes for multiplexed RNA detection in live cells with the Riboglow-FLIM sensor Esther Braselmann, Georgetown University
3:25 – 3:30 pm	Questions for speaker
3:30 – 3:45 pm	Thermodynamics of Selective DNA Binding by Cre Recombinase Jonathan Montgomery, Foster Lab, Ohio State University Questions for speaker
3:45 – 3:50 pm	·
3:50 – 4:05 pm	Evaluating the role of membrane localization on protein dimerization using MD simulations Adip Jhaveri, Johnson Lab, Johns Hopkins University
4:05 – 4:10 pm	Questions for speaker
4:10 – 4:40 pm	Break – Refreshments in River Radio Retreat
4:40 – 5:05 pm 5:05 – 5:10 pm	Intrinsically Disordered Regions Promote Protein Refoldability, Facilitating Retrieval from Biomolecular Condensates Stephen Fried, Johns Hopkins University Questions for speaker
·	·
5:10 – 5:25 pm	Control of G Protein-Coupled Receptor Function via Membrane Interacting Intrinsically Disordered C-Terminal Domains Chiara Mancinelli, Levitz & Eliezer Lab, Weill Cornell Medicine Questions for speaker
5:25 – 5:30 pm	·
5:30 – 5:55 pm	Intrinsically disordered regions that drive phase separation form a robustly distinct protein class Steven Whitten, Texas State University
5:55 – 6:00 pm	Questions for speaker
6:00 – 6:10 pm	General Discussion
6:30 pm	Dinner in Freeberg & Burke & Surrounding Area

Sunday Evening ♦ October 15, 2023

Poster Session I (N-Z)

8:00 – 10:00 pm Posters on display in Freeberg Hall and Burke Lounge

Instructions & assigned numbers are listed after the schedule of talks.

Displays by Sponsors in Freeberg Hall (upper level)
Please set up displays near beer, wine, and soft drinks

Monday Morning ◆ October 16, 2023

- ♦ Posters in Poster Session II may be mounted on Monday morning. Use assigned number.
- ♦ A white board will be available in Little Grassy Lodge, near the check-in window.

 If you can offer someone a ride, please write your name, the time when you are leaving Touch of Nature, number of people you can accommodate and which airport you are using.

7:00 – 8:20 am Breakfast in Freeberg & Burke & Surrounding Area

Platform Session III

8:15 am **Speakers - Please connect your laptop in advance of the session.**

AV support provided by Yingzi Xia, Fried Lab, Johns Hopkins Univ.

Moderator Arianna Lacen, Lee Lab, Univ Alabama at Birmingham

8:30 - 8:40 am Announcements - Tonya Zeczycki & Krishna Mallela, Organizers

15th Annual Gary K. Ackers Lecture in Biothermodynamics

8:40 – 8:50 am Speaker Introduction

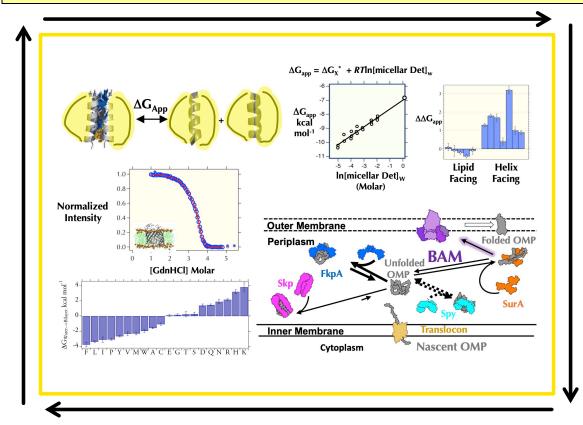
Janice Robertson, Washington University in St. Louis

8:50 – 9:40 am Visualizing the Periplasmic Chaperone Network Interactions

with Unfolded Clients

Karen Fleming, Johns Hopkins University

9:40 – 9:50 am Questions for Ackers Lecturer



Platform Session III - continued

9:50 – 10:20 am	Break – Refreshments in River Radio Retreat
10:20 – 10:35 am 10:35 – 10:40 am	An Intrinsically Disordered Region of the FACT Subunit, Spt16, Regulates Chromatin Dynamics in Controlling the Assembly of Transcription Factors at the Promoter for Transcription Initiation Priyanka Barman, Bhaumik Lab, Southern Illinois Univ. School of Medicine Questions for speaker
10:40 – 11:05 am 11:05 – 11:10 am	Probing molecular interactions in liquid condensed phases by solution NMR spectroscopy Nicholas Fawzi, Brown University Questions for speaker
11:10 – 11:25 am 11:25 – 11:30 am	Phase separation by SARS-CoV-2 N-protein is disrupted outside of a narrow range of nucleic acid stoichiometry Patrick Laughlin, Zlotnick Lab, Indiana University Questions for speaker
11:30 – 11:40 pm	General Discussion
11:40 – 12:00 pm	Announcements – Tonya Zeczycki & Krishna Mallela, Organizers Introduction of Attending Commercial Sponsors Reminder of Meeting of Conference Volunteers & Past Organizers

1:00 – 2:45 pm Free Time until Afternoon Session Look near the check-in counter in Little Grassy Lodge for information about possible organized events. Information about local parks and attractions is also available near the entrance to Little Grassy Lodge.

Attractions in Southern Illinois (https://southernillinoistourism.org) include Giant City State Park, orchards with apple cider, doughnuts, apple butter & pumpkins (Rendleman Orchards & Market and Flamm Orchards are popular), and plenty of small town charm in Makanda.

1:15 – 2:15 pm	Business Meeting of Past Organizers and Committee Pls		
Little Grassy Lodge - Friends Room			
	Note - Area will be unavailable to other attendees during this time.		

Monday Afternoon ♦ October 16, 2023

Platform Session IV

2:45 pm	Speakers - Please connect your laptop in advance of the session. AV support provided by Xinzhe Ren, Fried Lab, Johns Hopkins Univ.
Moderator	Vaibhav Upadhyay, Mallela Lab, Univ. of Colorado Anschutz Medical Campus
3:00 – 3:25 pm	PDZ domains: Model systems for studying allostery, dynamics, and binding specificity Ernesto J. Fuentes, University of Iowa
3:25 – 3:30 pm	Questions for speaker
3:30 – 3:45 pm	Protein design enables interrogation of thermodynamic signal transduction mechanisms Katherine Hatstat, DeGrado Lab, Univ. of California San Francisco
3:45 – 3:50 pm	Questions for speaker
3:50 – 4:05 pm	Energy-Preserving Variational Auto-Encoding and Decoding of Atomistic Protein Structure
4:05 – 4:10 pm	Joseph DePaolo-Boisvert, Minh Lab, Illinois Inst. of Technology Questions for speaker
4:10 – 4:40 pm	Break – Refreshments in River Radio Retreat
	bleak - Kellesiillelits III Kivel Kadio Ketleat
4:40 – 5:05 pm	The Mechanistic States of ABC Importers Heather Pinkett, Northwestern University
	The Mechanistic States of ABC Importers
4:40 – 5:05 pm	The Mechanistic States of ABC Importers Heather Pinkett, Northwestern University
4:40 – 5:05 pm 5:05 – 5:10 pm	The Mechanistic States of ABC Importers Heather Pinkett, Northwestern University Questions for speaker Thermodynamic Stability as a Determinant of the pK _a values of Ionizable Residues in Hydrophobic Environments in Proteins
4:40 – 5:05 pm 5:05 – 5:10 pm 5:10 – 5:25 pm	The Mechanistic States of ABC Importers Heather Pinkett, Northwestern University Questions for speaker Thermodynamic Stability as a Determinant of the pK _a values of Ionizable Residues in Hydrophobic Environments in Proteins Miranda Hurst, García-Moreno Lab, Johns Hopkins University
4:40 – 5:05 pm 5:05 – 5:10 pm 5:10 – 5:25 pm 5:25 – 5:30 pm	The Mechanistic States of ABC Importers Heather Pinkett, Northwestern University Questions for speaker Thermodynamic Stability as a Determinant of the pK _a values of Ionizable Residues in Hydrophobic Environments in Proteins Miranda Hurst, García-Moreno Lab, Johns Hopkins University Questions for speaker Force-based Detection of Transient Topoisomerase IA Dynamics
4:40 – 5:05 pm 5:05 – 5:10 pm 5:10 – 5:25 pm 5:25 – 5:30 pm 5:30 – 5:55 pm	The Mechanistic States of ABC Importers Heather Pinkett, Northwestern University Questions for speaker Thermodynamic Stability as a Determinant of the pK _a values of Ionizable Residues in Hydrophobic Environments in Proteins Miranda Hurst, García-Moreno Lab, Johns Hopkins University Questions for speaker Force-based Detection of Transient Topoisomerase IA Dynamics Maria Mills, University of Missouri
4:40 - 5:05 pm 5:05 - 5:10 pm 5:10 - 5:25 pm 5:25 - 5:30 pm 5:30 - 5:55 pm 5:55 - 6:00 pm	The Mechanistic States of ABC Importers Heather Pinkett, Northwestern University Questions for speaker Thermodynamic Stability as a Determinant of the pK _a values of Ionizable Residues in Hydrophobic Environments in Proteins Miranda Hurst, García-Moreno Lab, Johns Hopkins University Questions for speaker Force-based Detection of Transient Topoisomerase IA Dynamics Maria Mills, University of Missouri Questions for speaker
4:40 - 5:05 pm 5:05 - 5:10 pm 5:10 - 5:25 pm 5:25 - 5:30 pm 5:30 - 5:55 pm 5:55 - 6:00 pm 6:00 - 6:10 pm	The Mechanistic States of ABC Importers Heather Pinkett, Northwestern University Questions for speaker Thermodynamic Stability as a Determinant of the pK _a values of Ionizable Residues in Hydrophobic Environments in Proteins Miranda Hurst, García-Moreno Lab, Johns Hopkins University Questions for speaker Force-based Detection of Transient Topoisomerase IA Dynamics Maria Mills, University of Missouri Questions for speaker General Discussion Announcements - Officers for 2023-24 & Organizers for Gibbs 38

Monday Evening ◆ October 16, 2023

Poster Session II (A-M)

8:00 – 10:00 pm Posters on display in Freeberg Hall and Burke Lounge

Instructions & assigned numbers are listed after the schedule of talks.

Displays by Sponsors in Freeberg Hall (upper level)
Please set up displays near beer, wine, and soft drinks

Tuesday Morning ◆ October 17, 2023



When you leave Touch of Nature, please leave your room key at the counter in Little Grassy Lodge.



7:00 - 8:20 am

Breakfast in Freeberg & Burke & Surrounding Area

Platform Session V

	8:25 am	Speakers - Please connect your laptop in advance of the sessio AV support provided by Pritam Chakraborty , Bhaumik Lab, SIUC		
	8:40 – 8:45 am	Announcements – Tonya Zeczycki & Krishna Mallela, Organizers		
	Moderator	Mishghan Humayun, Durie Lab, University of Missouri		
	8:45 – 9:10 am	RNA-mediated ribonucleoprotein assembly controls TDP-43 nuclear retention Yuna Ayala, St. Louis University School of Medicine		
	9:10 – 9:15 am	Questions for speaker		
	9:15 – 9:30 am 9:30 – 9:35 am	Homomeric Protein Assembly <i>In Vivo</i> McKenze Moss, P. Clark Lab, University of Notre Dame Questions for speaker		
	9:35 – 10:00 am	·		
	9:35 – 10:00 am	Exploiting structural features and thermodynamics to develop a library of reagents coupling folding status to protein quality control outcomes Rick Page, Miami University		
	10:00 – 10:05 am	Questions for speaker		
	10:05 – 10:20 am	Short Break – Refreshments in River Radio Retreat		
	10:20 – 10:35 am	Xeno interactions between MHC-I proteins and molecular chaperones enable ligand exchange		
		on a broad repertoire of HLA allotypes Yi Sun, Sgourakis Lab, University of Pennsylvania		
	10:35 – 10:40 am	on a broad repertoire of HLA allotypes		
	10:35 – 10:40 am 10:40 – 11:05 am	on a broad repertoire of HLA allotypes Yi Sun, Sgourakis Lab, University of Pennsylvania		
		on a broad repertoire of HLA allotypes Yi Sun, Sgourakis Lab, University of Pennsylvania Questions for speaker Some Assembly Required: Multiple Modes of Self-association for a Staphylococcal Biofilm Protein		
	10:40 – 11:05 am 11:05 – 11:10 am 11:10 – 11:25 am	on a broad repertoire of HLA allotypes Yi Sun, Sgourakis Lab, University of Pennsylvania Questions for speaker Some Assembly Required: Multiple Modes of Self-association for a Staphylococcal Biofilm Protein Andrew Herr, Cincinnati Children's Hospital Questions for speaker Controlled Disorder: Phosphorylation Alters the Ensemble Structures of the Protein SRSF1 Talia Fargason, Zhang Lab, University of Alabama at Birmingham		
	10:40 – 11:05 am 11:05 – 11:10 am 11:10 – 11:25 am 11:25 – 11:30 am	on a broad repertoire of HLA allotypes Yi Sun, Sgourakis Lab, University of Pennsylvania Questions for speaker Some Assembly Required: Multiple Modes of Self-association for a Staphylococcal Biofilm Protein Andrew Herr, Cincinnati Children's Hospital Questions for speaker Controlled Disorder: Phosphorylation Alters the Ensemble Structures of the Protein SRSF1 Talia Fargason, Zhang Lab, University of Alabama at Birmingham Questions for speaker		
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	10:40 - 11:05 am 11:05 - 11:10 am 11:10 - 11:25 am 11:25 - 11:30 am 11:30 - 11:35 am 11:35 am	on a broad repertoire of HLA allotypes Yi Sun, Sgourakis Lab, University of Pennsylvania Questions for speaker Some Assembly Required: Multiple Modes of Self-association for a Staphylococcal Biofilm Protein Andrew Herr, Cincinnati Children's Hospital Questions for speaker Controlled Disorder: Phosphorylation Alters the Ensemble Structures of the Protein SRSF1 Talia Fargason, Zhang Lab, University of Alabama at Birmingham Questions for speaker		



Please complete the Gibbs₃₇ Post-Conference Survey

by Friday, October 20, 2023.

You may reach it using the QR code to the right or this link. https://forms.gle/DaWdDpPUX51Y7R3p7



Equilibrium binding checklist

		Equilibrium Binding Checklist			
Binding partner 1 (BP1):					
Binding partner 2 (BP2):					
Method:					
CONDITIONS:	Temperature	: Buffer & pH:			
	Salt(s):				
	Other:				
A. Required:					
☐ 1. Vary incubation	on time to test fo	or equilibration.			
Time range):	Number of time points:			
		BP2 concentration(s):			
Time-indep	endence across	s the entire binding curve?	Y D N D		
☐ 1.1. Alternative a	approach: meas	sure $k_{ m off}$.			
<i>k</i> _{off} :		Calculated equilibration time (5 half-lives):			
☐ 2. Vary the cond	entration of bot	h binding partners.			
Concentrat	ion range of 'tra	ace' binding partner:			
K_{D}^{app} indep	endent of trace	binder concentration?	Y D N D		
Concentrat	ion range show	ing invariant $K_{ extsf{D}}^{ ext{app}}$:			
Binding eq	uation used:	☐ hyperbolic ☐ quadratic			
Binding curves	shown?		Y D N D		
Systematic devi	ations from the	binding curve?	Y DND		
K D ^{app} :		(upper limit if dependent on trace binder concentration)			
B. Recommended:					
□ 1. Test <i>K</i> _D by an	independent ar	pproach.			
Alternative	e approach:				
$K_{ extsf{D}}^{ extsf{app}}$ from	alternative appr	roach:			
☐ 2. Determine the	e fraction of act	ive protein by titration.			
K _D correct	ed for active pro	otein fraction?	Y N		
Fraction o	f active protein:				
Comments:			1		

Checklist shared with permission from Keynote Speaker Dan Herschlag

How to measure and evaluate binding affinities

Jarmoskaite, AlSadhan, Vaidyanathan & Herschlag (2020) eLife 9:e57264

(https://doi.org/10.7554/eLife.57264)



Poster Information & Assigned Boards



Poster Sessions - General Information

Dates and Locations

Posters will be presented during evening sessions on Sunday and Monday in both Burke Lounge and Freeberg Dining Hall. Each poster session will start at **8:00 pm**. Free beverages will be served.

Session I - Sunday evening "N-Z"

Posters whose abstracts have first authors with last names starting with N-Z.

Session II - Monday evening "A-M"

Posters whose abstracts have first authors with last names starting with A-M.

Poster Numbers

Board #1 will be available for posting job openings and training program information.

Boards from #2 on are numbered according to the **last name of the first author**, regardless of who is presenting the poster.

Presenters are asked to place their poster on the board that corresponds to the numeral in their poster listing for Sunday (S2, S3, etc.) or Monday (M2, M3, etc.). That listing follows these instructions.

Poster Sizes & Boards

The poster boards come from a variety of sources. They are variable in size and made of materials ranging from foamcore boards to repurposed cubicle dividers.

Width: Making posters ≤ 48 inches wide is recommended.

Mounting: Pushpins will be provided, but some boards are very dense. You may wish to bring your own Velcro (hook and loop) buttons for mounting your poster.

Times

Posters may be mounted in the morning for viewing during the day,

Posters should be taken down by midnight of the assigned session day.

Confidentiality – No photos or recording without permission

Please remember that the content of all presentations (both talks and posters) for this meeting are confidential material and may contain unpublished results. Abstracts will not be posted online. Please ask permission from authors before taking photos of posters or poster material. Please do not record talks, posters, or any presented material unless a speaker has given you express permission.















Poster Session I ♦ Sunday, October 15, 2023

Posters numbered alphabetically by first author's last name (N-Z) (Posters S1 – S60)

- S1 Open Post for Educational Programs & Employment Opportunities
- S2 Poly(ADP-Ribose) has Cation and Length Dependent Structures. Tong (George)
 Wang, Kush Coshic, Mohsen Badiee, Aleksei Aksimentiev, Anthony K. L. Leung, and
 Lois Pollack, Cornell University
- S3 On the Mechanism of Dimer Activation of SF1 DNA Helicases. Binh Nguyen, John Hsieh, and Timothy M. Lohman, Washington University
- S4 Engineering Multidirectional pH-Dependent Antibody Interactions Through the Introduction of Ionizable Residues Within a VHH Homodimer Interface. Praise Oguntokun, Peter Gungel, Emily Camposeo, Tosha Laughlin, Hyeyoung Eom, Aaron Manz, Kevin Beck, Kylie Zawisza, Moly Eilbes, and James R. Horn, Northern Illinois University
- Mechanism of Protein Stabilization by Sugars in Crowded Solutions. Gil I. Olgenblum and Daniel Harries, The Hebrew University
- Rheostatic Contributions to Protein Stability Can Obscure a Position's Functional Role. Pierce T. O'Neil, Braelyn M. Page, Liskin Swint-Kruse, and Aron Fenton, University of Kansas Medical Center
- S7 Unfolded Outer Membrane Protein (uOMP) Modeling gives Insight into Protein Folding. Andrea L. Ori and Karen G. Fleming, Johns Hopkins University
- Structural Insights into Binding of Polyglutamylated Tetrahydrofolate by Soybean SHMT8. Luckio Owuocha and Lesa Beamer, University of Missouri
- S9 Illumination of the U1 Complex Recruitment to the 5' Splice Site in Alternative Splicing through Characterization of U1-70K SRSF1 Interaction. Trenton Paul, Jamal Shariq, Ethan Ekpenyong, Talia Fargason, Zihan Zhang, Ivon De Silva, and Jun Zhang, University of Alabama at Birmingham
- Pioneering the Details: A Multi-Method Collaboration to Gain Insight into Structure and Thermal Stability of Lipid Nanoparticles (LNPs) for mRNA Delivery. <u>Jérémie Parot</u>, Sven Even Borgos, Alicja Molska, Natalia Markova, and Martin Textor, Applied Photophysics Ltd.
- S11 Insights into How Chaperone Net Charge and Sequence Pattering Influence Nucleic Acid Folding. Gabrielle M. Perkins, Parnian Arafi, and Erik D. Holmstrom, University of Kansas
- Functional Site Distant Mutations of hGMPK Alter Dynamics and Kinetics. Andrea R. Poole, Mark Vincent C. dela Cerna, and T. Michael Sabo, University of Louisville
- S13 Thermodynamic Study of Substrate Binding with PrnA from *Burkholderia ambifaria*. KD Power, AE Rankins, SL Stokes, and JP Emerson, Mississippi State University
- S14 Cryo-EM of Human Factor Va and Factor Va-APC Complex. Suhaila Rahman and Enrico Di Cera, Saint Louis University School of Medicine
- S15 Effect of Myopalladin Ig3 Domain Cardiomyopathy Mutations on Actin-Binding and Bundling. Asha Rankoth Arachchige, Julie Tran, Alia Michaelis, and Moriah Beck, Wichita State University

- S16 Effect of N-linked Glycosylation and Disulfide Bonds on Yeast ER Proteome Refoldability. Xinzhe Ren, Siyuan Zhou, and Stephen D. Fried, Johns Hopkins University
- Studying the Effects of Transcription Factor IIS on RNA Polymerase II. Ryan Requijo, David Schneider, and Aaron Lucius, University of Alabama at Birmingham
- **S18** Thermodynamics of DNA Junctions with dU's. Alex Rohe, Irine Khutsisvili, and Luis A. Marky, University of Nebraska Medical Center
- Precision Engineering of Biological Function with Large-Scale Measurements and Machine Learning. David Ross, Drew S. Tack, Peter D. Tonner, Abe Pressman, Nathan D. Olson, Sasha F. Levy, Eugenia F. Romantseva, Nina Alperovich, and Olga Vasilyeva, National Institute of Standards and Technology
- S20 ¹⁵N Chemical Shift Anisotropy of Proteins: Reassessment through ¹⁵N Relaxation at Ultra-High Magnetic Fields. <u>T. Michael Sabo</u>, John O. Trent, David A. Case, Christian Griesinger, and Nasrollah Rezaei-Ghaleh, University of Louisville
- S21 Key Structural role of a Conserved *cis*-Proline Revealed by the P285S Variant of Soybean Serine Hydroxymethyltansferase 8. Vindya Samarakoon, Luckio F. Owuocha, Jamie Hammond, Melissa G. Mitchum, and Lesa J. Beamer, University of Missouri
- **S22 Mixed, Non-Classical Behavior in a Classic Allosteric Protein.** Paul J. Sapienza, Jeffrey P. Bonin, Dinusha Jinasena, Kelin Li, Henry Dieckhaus, Konstantin Popov, Jeff Aubé, and <u>Andrew L. Lee</u>, University of North Carolina, Chapel Hill
- S23 Riboglow-FLIM: Unveiling Subcellular RNA Localization through Multiplexing with Fluorescence Lifetime Imaging Microscopy. Nadia Sarfraz and Esther Braselmann, Georgetown University
- **S24** Structural Characterization of Avian Secretory Antibodies. Rebecca Schneider and Beth Stadtmueller, University of Illinois Urbana-Champaign
- S25 Using HDX-MS to Probe the Dynamics of Helix Unfolding and the Denatured State of GlpG. Abigail L. Schroetera, Saba Kanwalb, Heedeok Hongb, and Tobin R. Sosnick, University of Chicago
- S26 Ubiquitin-Proteasome System Regulation of a Chromatin Remodeling Factor FACT and its Interactome in Controlling Gene Expression With Link to Cancer. Rwik Sen, Amala Kaja, Priyanka Barman, Sukesh R. Bhaumik, Southern Illinois University School of Medicine
- **S27** Mapping Conformational Landscapes of Intrinsically Disordered Proteins. Hossain Shadman, Jesse D. Ziebarth, and Yongmei Wang, The University of Memphis
- S28 L. Monocytogenes Infection Activates PML Antibacterial Response via Interactions with Host Splicing Machinery. Luke Shafik and Esther Braslemann, Georgetown University
- S29 Engineering a Biomimetic Protein Corona for Predictive Nanoparticle Behavior.

 <u>Tanveer Shaikh</u>, Dhanush Amarasekara, Kenneth Hulugalla, Railey Mayatt, Thomas Werfel, and Nicholas C. Fitzkee, Mississippi State University
- S30 Reflections of a Half Century of Research on DNA Biothermodynamics. Richard D. Sheardy, Texas Woman's University
- **S31** *In Vivo* Nearest Neighbor Parameters for RNA. Jacob P. Sieg, Elizabeth A. Jolley, Melanie J. Huot, Paul Babitzke, and Philip C. Bevilacqua, Dept of Chemistry, Penn State University
- Optimizing Non-Consensus SPOP Binding Motifs Influences the Cooperativity
 Observed in Multivalent Binding. Scott A. Showalter, Nolan Jacob, and Emery T.
 Usher, Penn State University

- S33 Investigating the Mechanism of IgA Assembly in Avian Species. <u>Asta Simonovic</u>, Rebecca Schneider, and Beth Stadtmueller, University of Illinois Urbana-Champaign
- **S34** Prodomain Characterization of ADAM10/17. Conner E. Slone and Tom C. Seegar, University of Cincinnati College of Medicine
- Insights into the Isoenergetic Monomeric Structures of the Hepatitis C Virus 3'X RNA. Parker D. Sperstad and Erik D. Holmstrom, University of Kansas
- S36 Interrogating Protein Features that Allow Functional Tuneability from Single Amino Acid Substitutions. Shwetha Sreenivasan, Anastasiia Sivchenko, Aron Fenton, and Liskin Swint-Kruse, University of Kansas Medical Center
- S37 Assessing Lifetime-Differences of Riboglow-FLIM for Multiplexed RNA Imaging. Zachary Stickelman, Nadia Sarfraz, and Esther Braselmann, Georgetown University
- The Impact of Crowding on the SARS-CoV-2 Nucleocapsid Protein. Madison
 Stringer, Jasmine Cubuk, J. Jeremias Incicco, Debjit Roy, Melissa D. Stuchell-Brereton, and Andrea Soranno, Washington University in St. Louis
- Single-Molecule Spectroscopy of Apolipoprotein E Reveals a Complex Conformational Ensemble. Melissa D. Stuchell-Brereton, Upasana L. Mallimadugula, Justin J. Miller, Jasmine Cubuk, Debjit Roy, Anshuman Jaysingh, Carl Frieden, Greg Bowman, and Andrea Soranno, Washington University in St. Louis
- S40 Mapping by Oxidation the Binding of Small and Large Ligands on Proteins. Yan Sun, Damian Houde, Roxana Iacob, Robert Swift, Jason Baird, Michael Holliday, Sergei Khrapunov, Subray Hegde, Simone Sidoli, and Michael Brenowitz, Albert Einstein College of Medicine
- S41 Comparing the Stability of Riboswitches from Different Organisms. <u>Victoria E.</u> Suntich and Ana Maria Soto, Towson University
- S42 G-Quadruplex Stabilization by TMPyP4 Monitored Through Introduction of Complimentary DNA. <u>Drew Symasek</u>, Alan Gunter, and Hui-Ting Lee, University of Alabama at Birmingham
- Structure-Based Engineering of Secretory Immunoglobulins that Neutralize Influenza Virus. Emma Thames, Sonya Kumar Bharathkar, and Beth Stadtmueller, University of Illinois, Urbana-Champaign
- S44 Dystrophin Interacts Differentially with Dystrobrevin Isoforms. Vaibhav Upadhyay, Sudipta Panja, Shashikant Ray, and Krishna M. G. Mallela, University of Colorado-Anschutz
- S45 All-Atom Simulations of Phosphorylated Intrinsically Disordered Proteins. Emery T. Usher, Martin J. Fossat, and Alex S. Holehouse, Washington University in St. Louis
- S46 Hepatitis B Virus Capsid Demonstrates Negative Cooperativity When Bound by Capsid Assembly Modulators. Caleb J. Valkner, Angela Patterson, Balasubramanian Venkatakrishnan, Samson Francis, and Adam Zlotnick, Indiana University
- S47 The Role of Structural and Dynamic Rearrangements in the Functional Regulation of a Classically Allosteric Protein. <u>Darex J. Vera-Rodríguez</u>, Paul J. Sapienza, and Andrew L. Lee, University of North Carolina at Chapel Hill
- S48 High-Resolution Models of Conformational Dynamics in DNA Recognition by the master Transcription Factor PU.1. Tyler N. Vernon, J. Ross Terrell, Markus W. Germann, W. David Wilson, and Gregory M. K. Poon, Georgia State University
- S49 Global vs. Local Specialization: An Examination of the Divergence in Overall vs.
 Clustered Amino Acid Enrichments in Extremophilic DNA Polymerase I Homologs.
 Elena Voisin, Larissa Cortes Morales, Alyce Fields, and Allyn Schoeffler, Loyola
 University New Orleans

- S50 Coarse-Grained Modelling of RNA: Capturing the Dynamics of Sugar Pucker Conformational Transitions. Yiheng Wu, Riccardo Alessandri, Aria Coraor, Xiangda Peng, Tobin Sosnick, and Juan de Pablo, University of Chicago
- **S51** Exploring Misfolded Proteins with Crosslinking Mass Spectrometry. Yingzi Xia and Stephen Fried, Johns Hopkins University
- Systematic Study on Effects of DnaKJ and Trigger Factor on Co-translational Folding in *E. coli*. Divya Yadav, Idil Demiralp, Mark Fakler, and Stephen D. Fried, Johns Hopkins University
- S53 Bringing AUC into the cGMP Space to Meet FDA/EMA Regulations. Alexander E. Yarawsky, Erik Gough, Valeria Zai-Rose, Natalya Figueroa, Hazel Cunningham, John Burgner, Michael DeLion, and Lake N. Paul, BioAnalysis, LLC
- A Novel Lasofoxifene/Elacestrant Hybrid Antiestrogen Illuminates Unique
 Relationship Between Ligand Binding Pose and Anti-Tumoral Action in Y537S
 ESR1 Breast Cancer Xenografts. Kristen Young, Govinda Hancock, Steve Kregel, and
 Sean Fanning, Loyola University Chicago
- **Molecular Basis of Nurr1-RXRα Activation.** <u>Xiaoyu Yu</u>, Jinsai Shang, and Douglas J. Kojetin, Vanderbilt University
- S56 The Structural Basis for DNA-uptake by *Acinetobacter*. Yafan Yu and Kurt Piepenbrink, University of Nebraska-Lincoln
- Structural Characterization of the Type IV Secretion System of Legionella pneumophila. Mishghan Zehra, Wing-Cheung Lai, and Clarissa Durie, University of Missouri- Columbia
- **The Central Role of RS in Spliceosome Assembly.** <u>Jun Zhang</u>, University of Alabama at Birmingham
- S59 Charges in Hydrophobic Environments in Proteins: A Study of Arg Ion-Pairs.

 Yipeng Zhang, Jasmine A. Forbes, Jaime L. Sorenson, Aaron C. Robinson, Jamie L. Schlessman, and Bertrand Garcia-Moreno E., Johns Hopkins University
- S60 Characterization of U2AF35 and Interaction between U2AF35 and SRSF1. Zihan Zhang, Ivon De Silva, Talia Fargason, and Jun Zhang, University of Alabama at Birmingham
- S61 Elucidating the effect of disease-causing mutations on the structure-function of the CT domain of dystrophin. Shashikant Ray*, Vaibhav Upadhyay, Sudipta Panja, Jeffrey Kearns, and Krishna M.G. Mallela















Poster Session II ♦ Monday, October 16, 2023

Posters numbered alphabetically by first author's last name (A to M) (Posters M1 – M61)

- M1 Open Post for Educational Programs & Employment Opportunities
- M2 Investigating the Role of Small Basic Protein in the Formation of Staphylococcal Biofilms. P. Ethan Adkins, Alexander Yarawsky, and Andrew Herr, Cincinnati Children's Hospital Medical Center
- M3 The Novel Interaction of a DNA Quadruplex with a Gemini Surfactant. Derek Aguilar, Helen Nembaware, Alexys Ginegaw, Alaina Jordan, Payton Justice, Rafaela Dominguez, and Richard D. Sheardy, Texas Woman's University
- M4 Conformational Disorder in Regulation of Enzyme and Nanoparticle Catalysts.
 Yeongseo An, Dan Burns, Rochelle Dotas, Timothy Egner, Balabadra Khatiwada, Trang
 Nguyen, Jeffrey Purslow, Sergey Sedinkin, Aayushi Singh, and Vincenzo Venditti, Iowa
 State University
- M5 Negative Cooperativity Creates a Frustrated System in *E. Coli* Adenylate Kinase.

 <u>Anna Andrick</u>, Miranda Russo, Sarah Brantley, Ananya Majumdar, and Vincent J Hilser,
 Johns Hopkins University
- M6 Disruptions of DNA Base-Pair Cooperativity Probed with Time-Resolved Infrared Spectroscopy. Brennan Ashwood, Michael S Jones, Andrew L. Ferguson, and Andrei Tokmakoff, University of Chicago
- M7 Sequential Mixing Stopped Flow Method to Investigate the Kinetic Mechanisms of Protein Translocation Catalyzed by Clpb. <u>Jaskamaljot Kaur Banwait</u>, and Aaron L. Lucius, University of Alabama at Birmingham
- M8 Determining Residue Interactions Leading to Variability in Interfacial Free Energy in a Repeat Protein System. Soumya Behera, Mark Petersen, Sai Harshitha Dakoor, and Doug Barrick, Johns Hopkins University
- M9 Conformational Entropy and Hydration in Molecular Recognition by Proteins.
 Anthony Bishop, Jose A. Caro, Taylor R. Cole, Hayley Hobley, Glorise Montalvo-Torres,
 Weimin Tan, and <u>Josh Wand</u>, Texas A&M University
- M10 Characterizing a *Mycobacterium Tuberculosis* Riboswitch Candidate. Swadha V. Bhatt, Emily Paris, and Ana Maria Soto, Towson University
- M11 Interactions Between Bacterial Amyloids and α-Synuclein Leads to Prolonged Innate Immune Responses. Whitney G. Bond and Tonya N. Zeczycki, Brody School of Medicine at East Carolina University
- **M12** Exploring the Druggability of Proteins. Adrian Calderon and Jannette Carey, Princeton University
- M13 A Dynamic Protein that is not Allosteric. Jannette Carey, Princeton University
- **M14** RNAP Fidelity Using High-Throughput Kinetics. Zachariah I. Carter, William O'Brien, and Andrew F. Gardner, New England Biolabs, Inc.
- M15 The Staphylococcus aureus Srrb Sensor Histidine Kinase Binds Dicarboxylic Acids Through an Extracellular Cache Domain. Delaney Catania, and Nicholas Hammons, University of Iowa
- M16 A Thermodynamic Analysis of CLC Transporter Dimerization in Lipid Bilayers.
 Rahul Chadda, Taeho Lee, Robyn Mahoney-Kruszka, Elizabeth G. Kelley, Nathan
 Bernhardt, Priyanka Sandal, and <u>Janice L. Robertson</u>, Washington University in St.
 Louis

- M17 Neural Network Interpretation Suggests Sequence Determinants of a Protein Domain that Interconverts between Alpha Helix and Beta Sheet Folds. Ethan Chen, Joseph Schafer, Joseph Thole, and Lauren Porter, National Institutes of Health
- M18 Subtelomeric Sequence of TERRA Regulates the Level of Gene Expression. <u>Tanvir Ahmed Chowdhury</u>, and Hui-Ting Lee, University of Alabama at Birmingham
- M19 Assembly Thermodynamics of a Bivalent Heterotrimer in the Notch Signaling Pathway. Cyril Cook, Kristen Ramsey, and Douglas Barrick, Johns Hopkins University
- M20 μ-Opioid Receptor Structural Conformations that Lead to Different Signaling Pathways. <u>David Cooper</u>, Joseph DePaolo-Boisvert, Stanley Nicholson, Barien Gad, and David Minh, Illinois Institute of Technology
- M21 The Properties of Elastic Protein Materials are Directed by the Properties of their Disordered Monomeric Precursors. Giovanni M. Crump, Jonathan M. Preston, and Ronald L. Koder, The City College of New York
- M22 Single-Molecule Spectroscopy of the SARS-CoV-2 Nucleocapsid Protein. <u>Jasmine Cubuk</u>, J. Jeremias Incicco, Debjit Roy, Melissa D. Stuchell-Brereton, Kathleen B. Hall, and Andrea Soranno, Washington University in St. Louis
- M23 A Novel Thiol-Dependent Inhibitory Modification of the Zinc Regulatory
 Transcription Factor ADCR, AJ. Cutright, EL. Matthews, T. Shaikh, KD. Power, JA.
 Thornton, NC. Fitzkee, SL. Stokes, and JP. Emerson, Mississippi State University
- **M24** LLPS of Human Glucocorticoid Receptor. Margaret Daugherty, Will Gerash, Nate Kesti, Devlin Swanson, and David Bain, Colorado College
- M25 An Exploration of How G-Quadruplex Binding of SRSF1 Expands its RNA-Binding Capability, N.I.U. De Silva, T. Fargason, N.E Lehman, H-T. Lee, and J. Zhang, University of Alabama at Birmingham
- **M26** FkpA Binds Outer Membrane Proteins as a Heterogeneous Ensemble of Conformations. <u>Taylor A. Devlin</u>, Dagan C. Marx, Michaela A. Roskopf, and Karen G. Fleming, Johns Hopkins University
- M27 Demystifying Dynamics: How you can Use Force Spectroscopy to Help Understand the Dynamics and Functions of Proteins, Nucleic Acids, and Phase Separated Droplets. Matthew Dilsaver, Trey Simpson, and Andrea Candelli, LUMICKS
- M28 Quantifying What We Can't See: Protein Degradation in the Cell. <u>Jacob Diehl</u> and Patricia Clark, University of Notre Dame
- M29 Engineering Allosteric Sites Across the Surface of a Model Enzyme. <u>Jerry C. Dinan,</u> James W. McCormick, and Kimberly A. Reynolds, University of Texas Southwestern Medical Center
- M30 The Role of Protein-Protein Interactions in Fluc Dimerization Stability and Function. Melanie Ernst, Jennifer Chen, and Janice L. Robertson, Washington University in St. Louis
- M31 Single-Molecule Spectroscopy Reveals Residue-Specific Impacts on Domain Folding Of Apolipoprotein E. <u>Klaudio Fatmiri</u>, Melissa D. Stuchell-Brereton, and Andrea Soranno, Washington University in St. Louis
- M32 E. coli RecBCD Nuclease Domain Regulates Helicase Activity but not Single Stranded DNA Translocation Activity. Nicole Fazio, Kacey N. Mersch, Linxuan Hao, and Timothy M. Lohman, Washington University
- M33 Improving Anti-Breast Cancer Activities by Optimizing Torsion Angle Energies.

 Emma Fink, David Zak, Govinda Hancock, Kristen Young, and Sean W. Fanning, Loyola University Chicago

- M34 Enzyme Kinetic Model for the Coronavirus Main Protease Including Dimerization and Ligand Binding. Barien Gad, Van Ngoc Thuy La, Lulu Kang, David D. L. Minh, Illinois Institute of Technology in Chicago
- M35 Exploring Interfaces Linked to Antigen Binding in the Creation of pH-Dependent Fab/Antigen Interactions. Peter Gungel, Emily Camposeo, Praise Oguntokun, Tosha Laughlin, Hyeyoung Eom, and James R. Horn, Northern Illinois University
- M36 Identity and Distribution of Histone H3 Tail Basic Residues Determine NCP Phase Separation Propensity. Erin F. Hammonds and Emma A. Morrison, Medical College of Wisconsin
- M37 PA0012 is a Cyclic di-GMP Effector that Biases *P. aeruginosa* Type IV Pili Twitching Motility. Nicholas A. Hammons, Kaitlin D. Yarrington, Christopher Ptak, Devin J. Hendrix, Dominique H. Limoli, and Ernesto J. Fuentes, University of Iowa
- M38 An Unconventional Estrogen Receptor Conformation Induces SUMO Expression in Breast Cancer Cells. Govinda Hancock, Kristen Young, and Sean W. Fanning, Loyola University Chicago
- M39 Parameter Dependence of the Solubility Limit for DiSodium Phosphate. Caleb Huang and B. Montgomery Pettitt, University of Texas Medical Branch
- M40 Determining the Mechanism of Translocation of ClpA while Unfolding Polypeptide Substrates. Liana Islam and Aaron L. Lucius, University of Alabama at Birmingham
- M41 Can an Intrinsically Disordered Protein Qualify as a Fold Switch or Shape Shifter?

 Bhavika T. Kaparthi, Alexandra Klinger, Corie Ralston, Sayan Gupta, Line Kristensen, and Sarah E. Bondos, Texas A&M University
- M42 Argonaute 2 Binding Affinity to Determine Repression of mi-mRNA Pairs. Christina Karadiakos and Alan Chen, University at Albany, State University of New York
- M43 A Host-Guest System for Quantifying Thermodynamic Parameters in the Nanoparticle Protein Corona. Chathuri S. Kariyawasam, Abdullahi T. Aborode, Thayane Lopes de Sousa, and Nicholas C. Fitzkee, Mississippi State University
- M44 Computational Protein Design as a Tool for Probing Relative Energetic Contributions of Core Packing and Metal Coordination. Dylan Klein and Vikas Nanda, Rutgers University
- M45 Hydroxyl Radical Footprinting Yields Ensemble-averaged Atomic Level
 Conformational Descriptors of Intrinsically Disordered Proteins. Alexandra Klinger,
 Decipher Hydro
- M46 PAD Specificity to Histone Tail Arginines. Alex Kowalczyk and Emma Morrison, Medical College of Wisconsin
- M47 An Allosteric Redox Switch in Domain V of β2-glycoprotein I Controls Membrane Binding and Anti-Domain I Autoantibody Recognition. Suresh Kumar, Mathivanan Chinnaraj, William Planer, Xiaobing Zuo, Paolo Macor, Francesco Tedesco, John Wulf II, Katherine Basore, Nicola Pozzi, and Edward A. Doisy, Saint Louis University School of Medicine
- M48 Using smFRET to Investigate Structural Dynamics Between G Quadruplexes and T-Loops in vitro. Arianna Lacen and Hui-Ting Lee, University of Alabama at Birmingham
- M49 Integrated Use of Biophysics to Improve Data and Decision Quality in Drug Discovery. Orthogonal use of GCI, DSF, ITC and DLS. May Poh Lai, Natalia Markova, Edward FitzGerald, and Alastair Davy, Malvern Panalytical
- **M50** Coupled Binding and Folding of Ribonuclease S. Yumin Lee, Yiheng Wu, Brennan Ashwood, Balamurugan Dhayalan, Isabelle Gagnon, Tobin R. Sosnick, and Andrei Tokmakoff, University of Chicago

- **M51** Folding of Prestin's Anion-Binding Site and the Mechanism of Outer Hair Cell Electromotility. Xiaoxuan Lin, Patrick Haller, Navid Bavi, Nabil Faruk, Eduardo Perozo, and Tobin R. Sosnick, University of Chicago
- M52 Recognition of Recombination Intermediates by the Mismatch Repair Protein Msh2-Msh6. Zane Lombardo, Tai Lon Tan, Jesse Pellman, Bold Boldbayar, Amy Du, and Ishita Mukerji, Wesleyan University
- M53 Direct Prediction of Intrinsically Disordered Protein Conformational Properties from Sequence. <u>Jeffrey M. Lotthammer</u>, Garrett M. Ginell, Daniel Griffith, Ryan J. Emenecker, Alex S. Holehouse, Washington University
- M54 Examining the Impact of NTD Mutations on RBD Conformational Dynamics and ACE2 Receptor Binding Affinity of the SARS-CoV-2 Spike Protein. Alexandra Lucas, Vaibhav Upadhyay, Casey Patrick, Shashikant Ray, and Krishna M.G. Mallela, University of Colorado-Anschutz
- M55 Lipid Dependent Thermodynamics of GPCR Function. Ed Lyman, University of Delaware
- M56 Establishing the PahZ1 peptidase scaffold. <u>Justin D. Marsee</u>, Joshua Couch, Brittany Williams, Yordanos Williams, Connor Coughran, Michael Mercante, and Justin M. Miller, Middle Tennessee State University
- M57 Regulation of GPCR Function by its Disordered C-Terminal Domain. <u>Dagan Marx</u>, Joon Lee, Alexa Strauss, David Eliezer, and Joshua Levitz, Weill Cornell Medicine
- M58 Thermodynamically Characterized Conformational Changes of Zinc Dependent SczA. EL. Matthews, AJ. Cutright, SL. Stokes, and JP. Emerson, Mississippi State University
- M59 Protein Supercharging-induced Phase Change Enhances Optical Nanosensor Functionality. James J McCann, Mijin Kim, Ewelina Randall, Chen Chen, Zvi Yaari, Yu Chen, Daniel Heller, and Ronald Koder, City College of New York
- **M60** High-throughput Analysis of Twister Ribozymes. <u>Lauren N. McKinley</u>, McCauley O. Meyer, and Philip C. Bevilacqua, Penn State University
- M61 Sequence and Structural Features that Govern Disorder and Secretion of Autotransporter Proteins. Cedrick D. Mukinay, Michael C. Baxa, Tobin R. Sosnick, and Patricia L. Clark, University of Notre Dame

















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