

Alex Sodt

BS Chem/Physics Univ. Washington 2000

2002 Q-Chem software developer

PhD 2007 Martin Head-Gordon (electronic structure)

2007-9 post-doc with Teresa Head-Gordon (biophysics)

2010-2015 post-doc with Richard Pastor@NIH (lipid biophysics)

Currently Principal Investigator NICHD/NIH

Eunice Kennedy Schriver National Institute of Child Health and Human Development



A QUANTUM LEAP INTO THE FUTURE OF CHEMISTRY

Q-Chem Employment Opportunities:

Research Associate in Computational Chemistry:

Q-Chem, Inc., is seeking a scientist for a research associate position at its main office in Pleasanton, CA. The successful candidate should have a Ph.D. in theoretical chemistry or a closely-related field, with significant programming experience (C++, Fortran). Duties will include implementation of novel features, algorithm development, and code maintenance. Previous experience as a developer of a quantum chemistry software package is highly desirable. Candidates must be legally authorized to work for a US company and willing to relocate to California. The initial contract will be for one year, renewable upon mutual consent.

Please send a CV to jobs@q-chem.com. Review of applications will commence on Jan. 25, 2016, and continue until the position is filled.

Q-Chem Inc., 6601 Owens Drive, Suite 105, Pleasanton, CA 94588

Internship Opportunity for Graduate Students and Postdocs:

Q-Chem is announcing an internship program for graduate students and postdocs. The goal of the program is to accelerate the development of new capabilities and core infrastructure; to provide training to new generation of Q-Chem developers; and to help academic developers integrate new features into Q-Chem.

Two types of internships will be supported:

1. **Targeted internships aimed at specific developments initiated by Q-Chem, Inc.** In this case, Q-Chem will provide travel support (domestic airfare and ground transportation) and a stipend of \$2,500 per month (\$625 per week), to be used to cover local accommodation costs.
2. **Internships initiated by academic developers.** Q-Chem will provide 50% of travel support (domestic airfare and ground transportation) and a stipend of \$1250 per month towards local expenses.

Home

How to Buy

Features

Technical Support

Workshops/Webinars

Instructional Materials

Workshops/Webinars

About Us

Contact Us

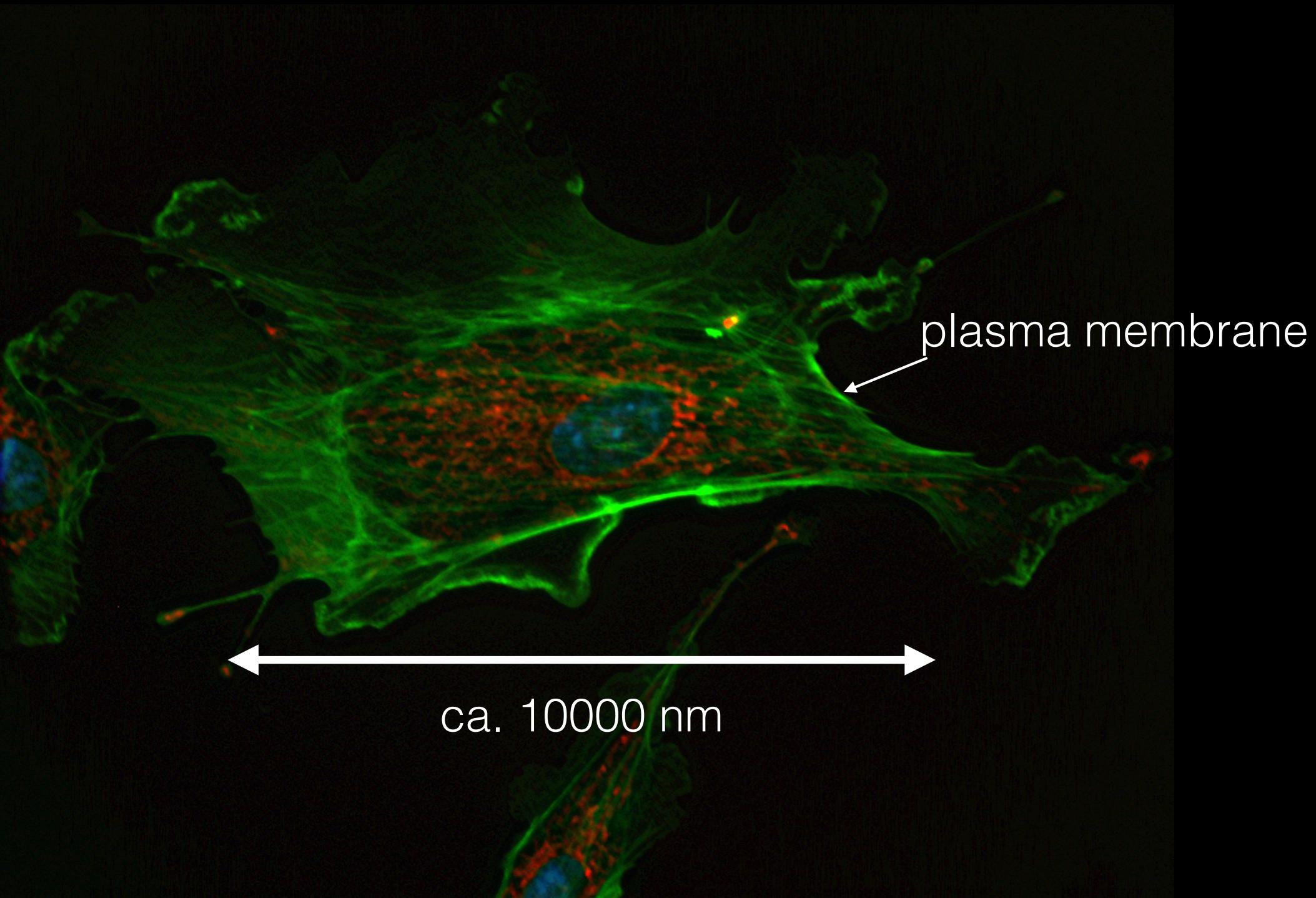
Free Evaluation

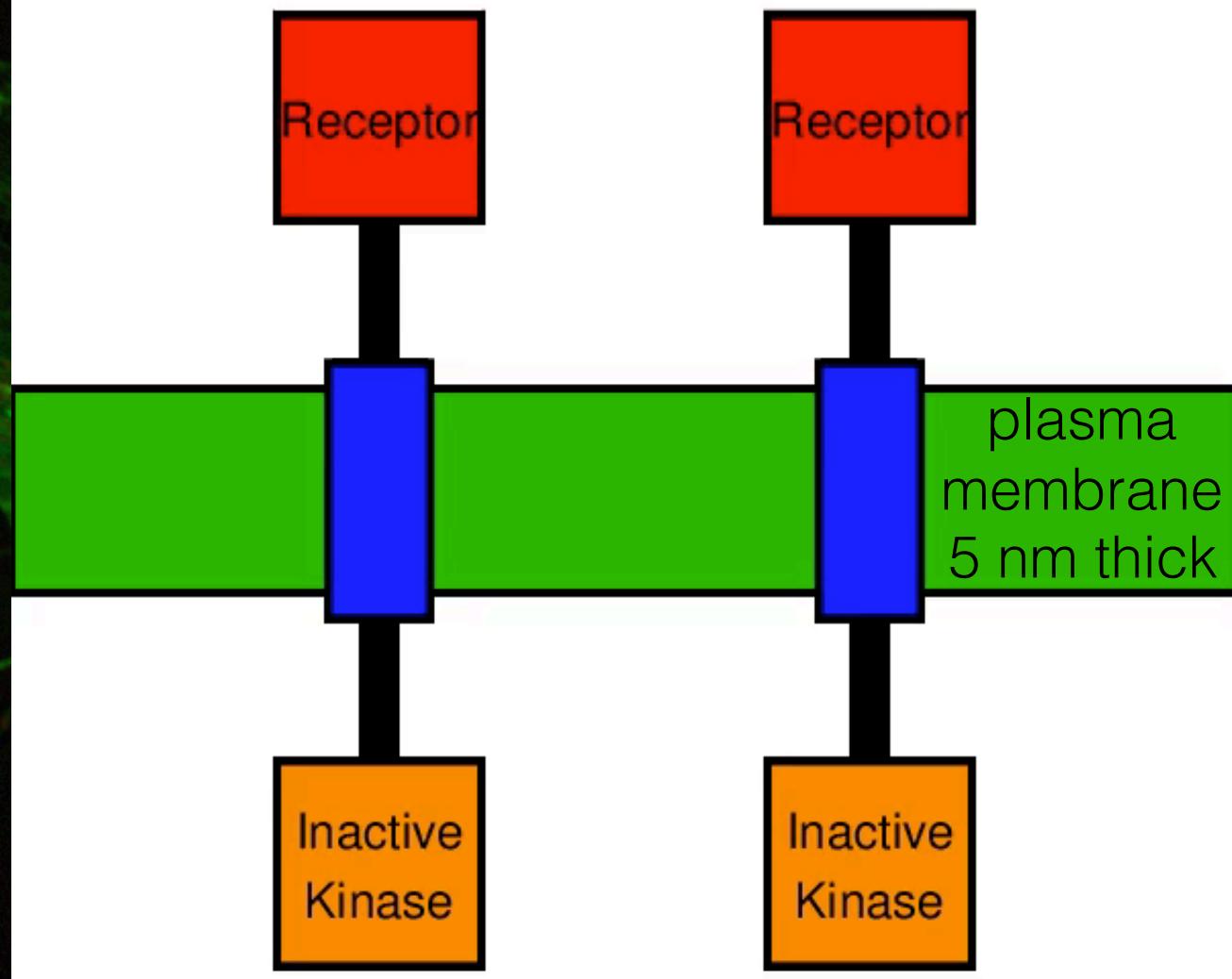
Find us on
Facebook

Q-Chem Forum:

iOpenShell

Center for computational
studies of electronic structure
and dynamics

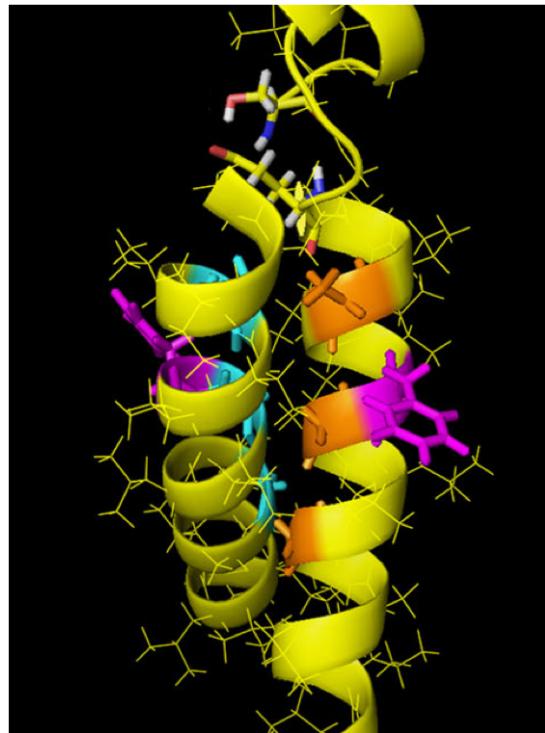




at membrane

ms)

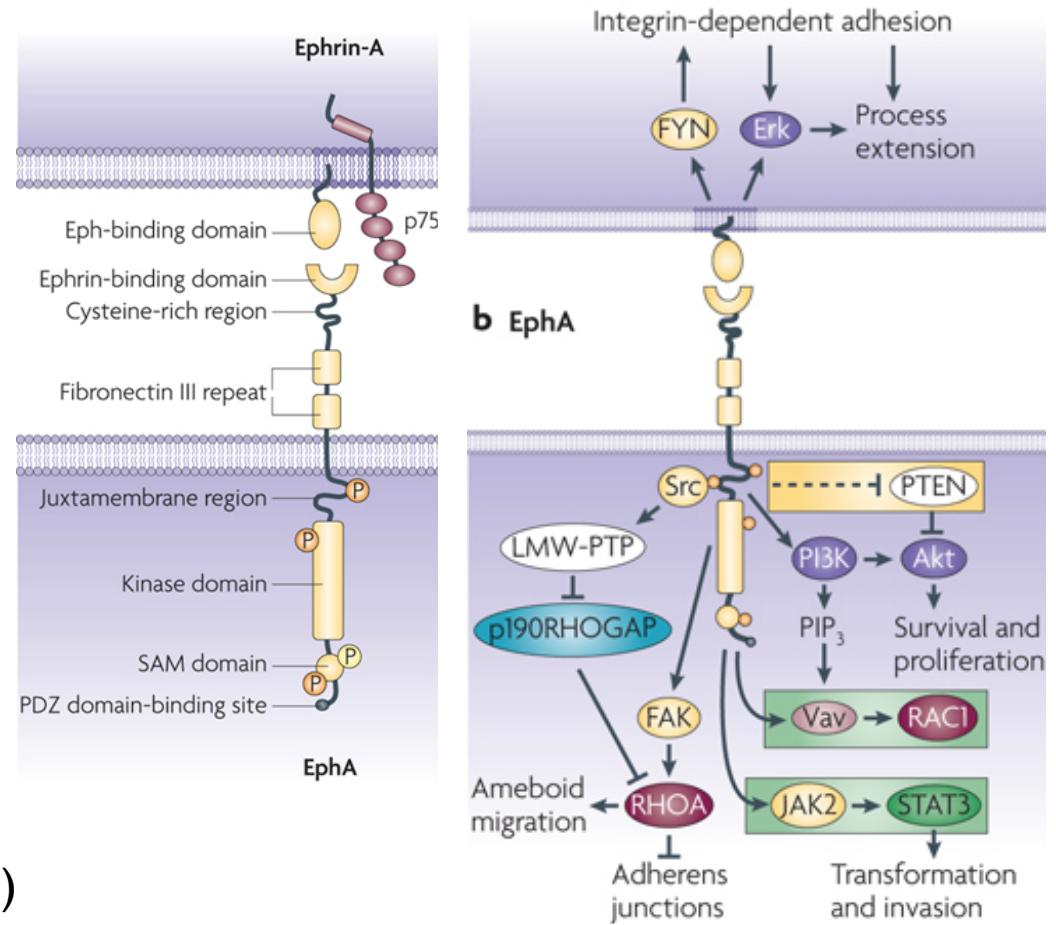
Simple system,
carefully controlled:
transmembrane region,
small bilayer patch



EphA1, Zhang et al, Proteins 81 365-376 (2013)

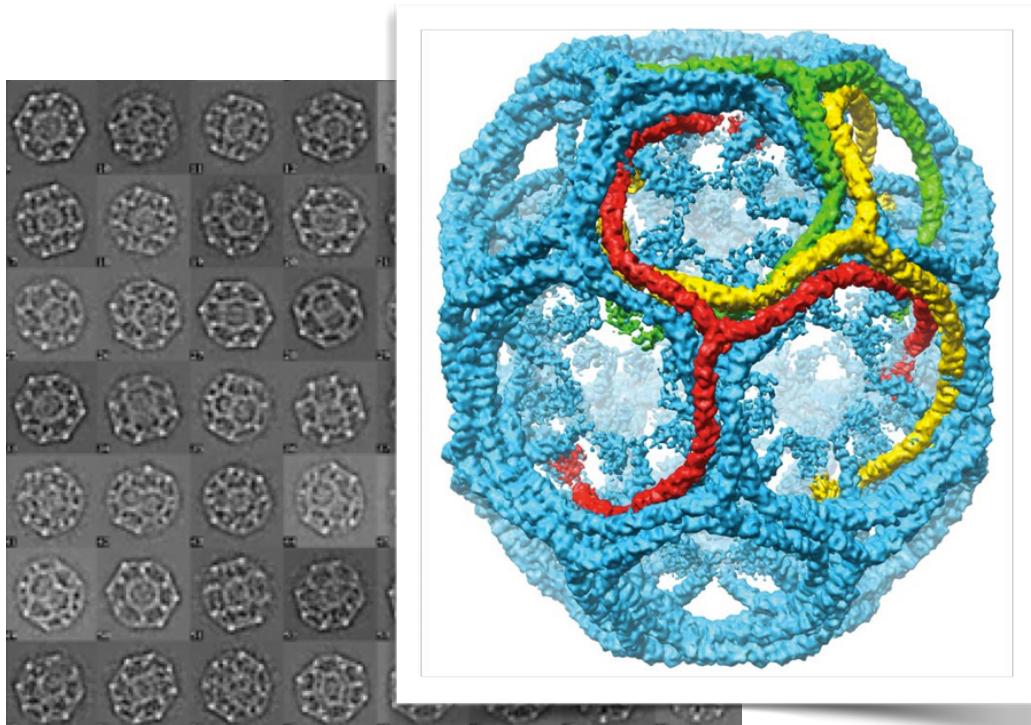
Bocharaov et al, J. Biol. Chem., 283, 29385 (2008)

Would like to do full system,
in a realistic environment



Pasquale, Nat. Rev. Cancer, 10 165 (2010)

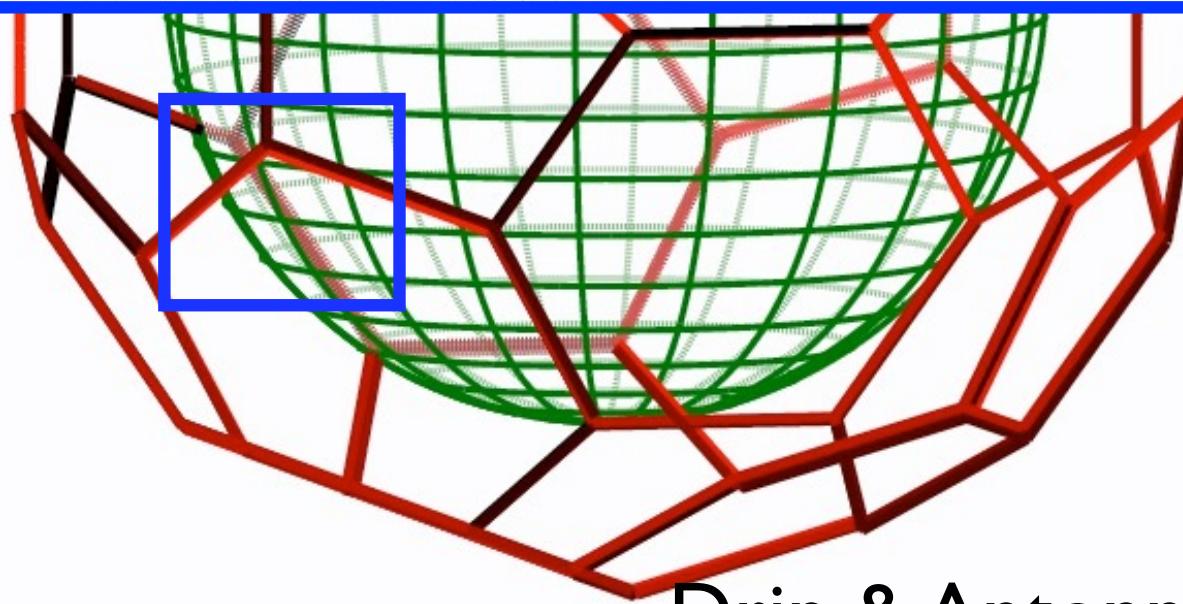
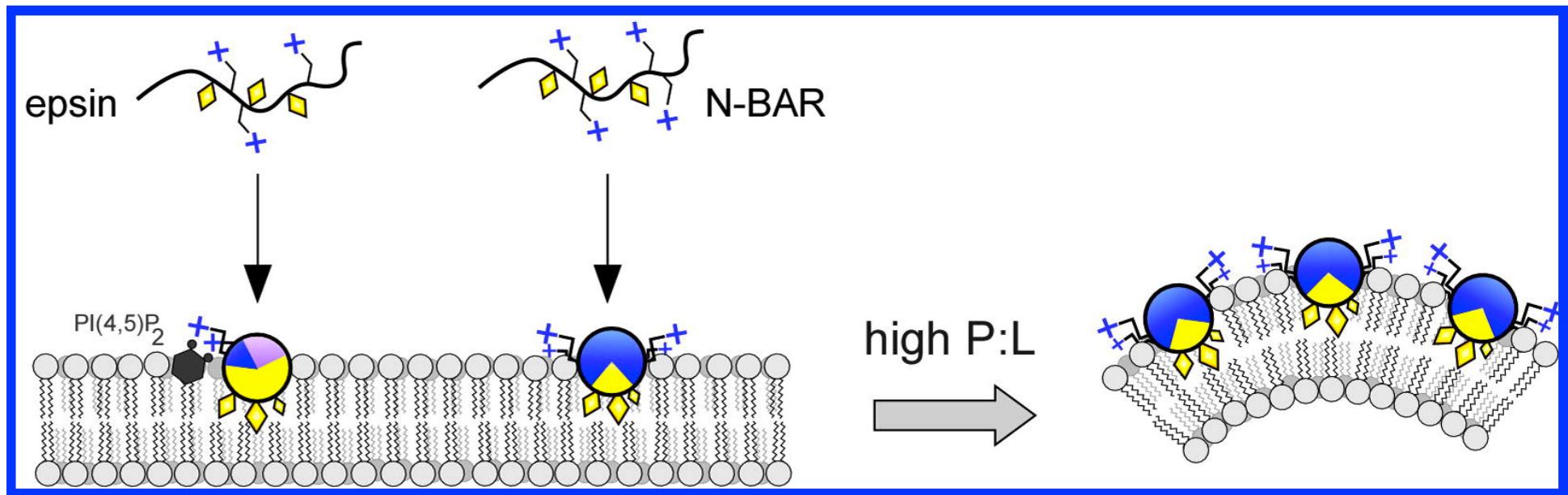
Transport of receptors from the membrane into the cell



electron microscopy

Fotin et al, Nature 432 573 (2004)

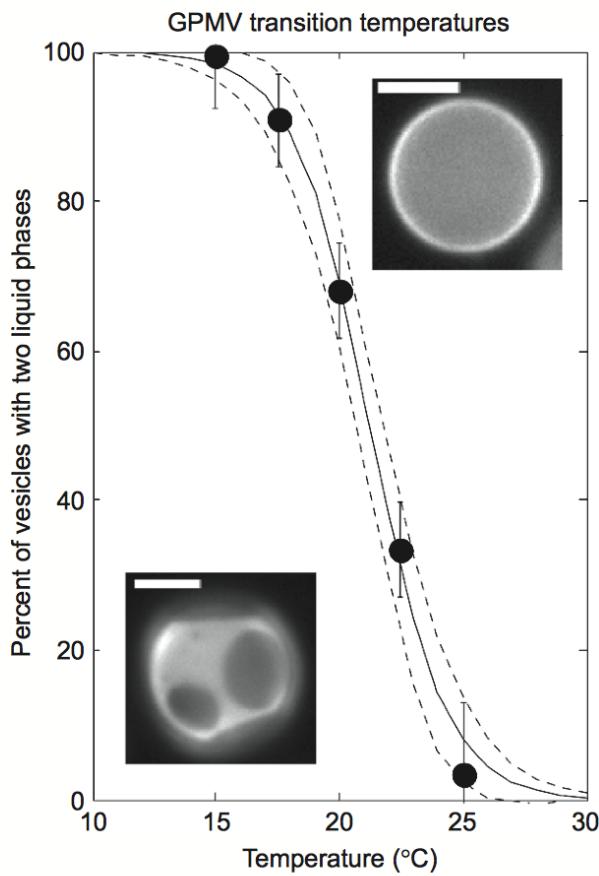
Protein-lipid curvature coupling



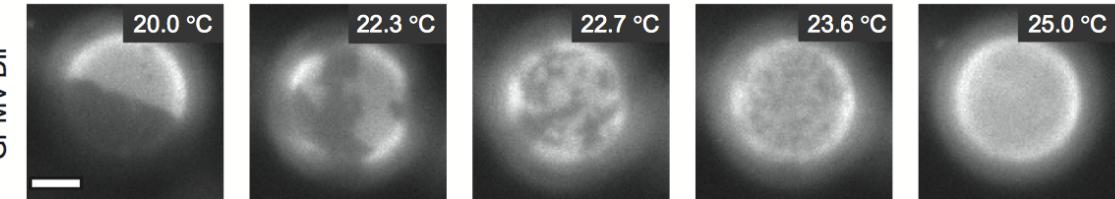
Drin & Antony FEBS Letters
584 1840 (2010)

Modeling goals: lateral organization

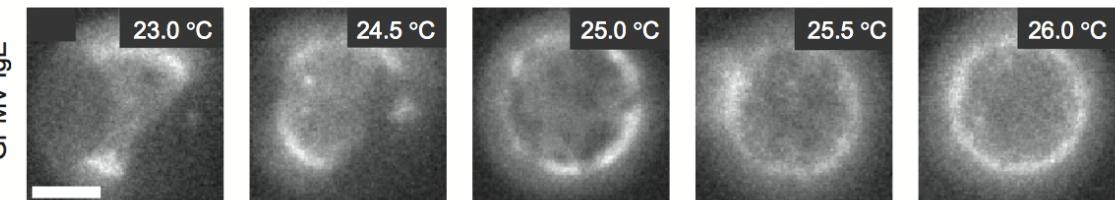
a



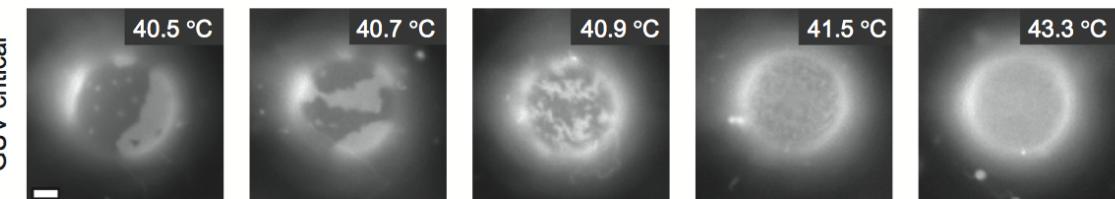
b



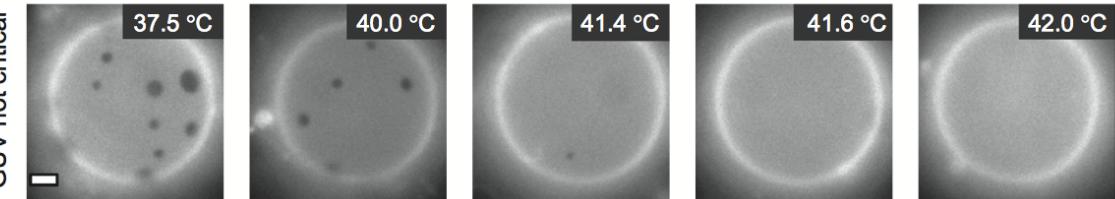
c



d



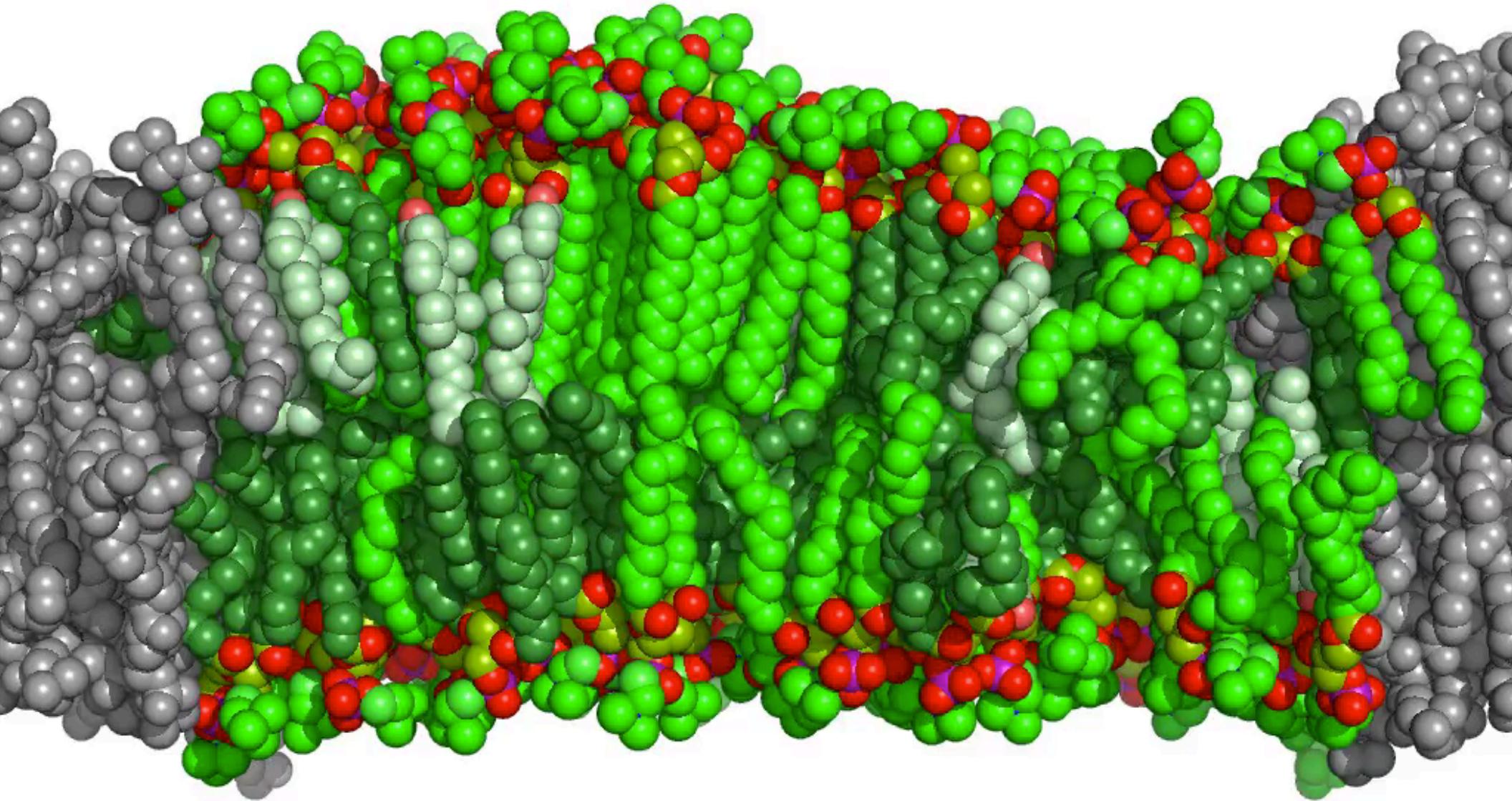
e

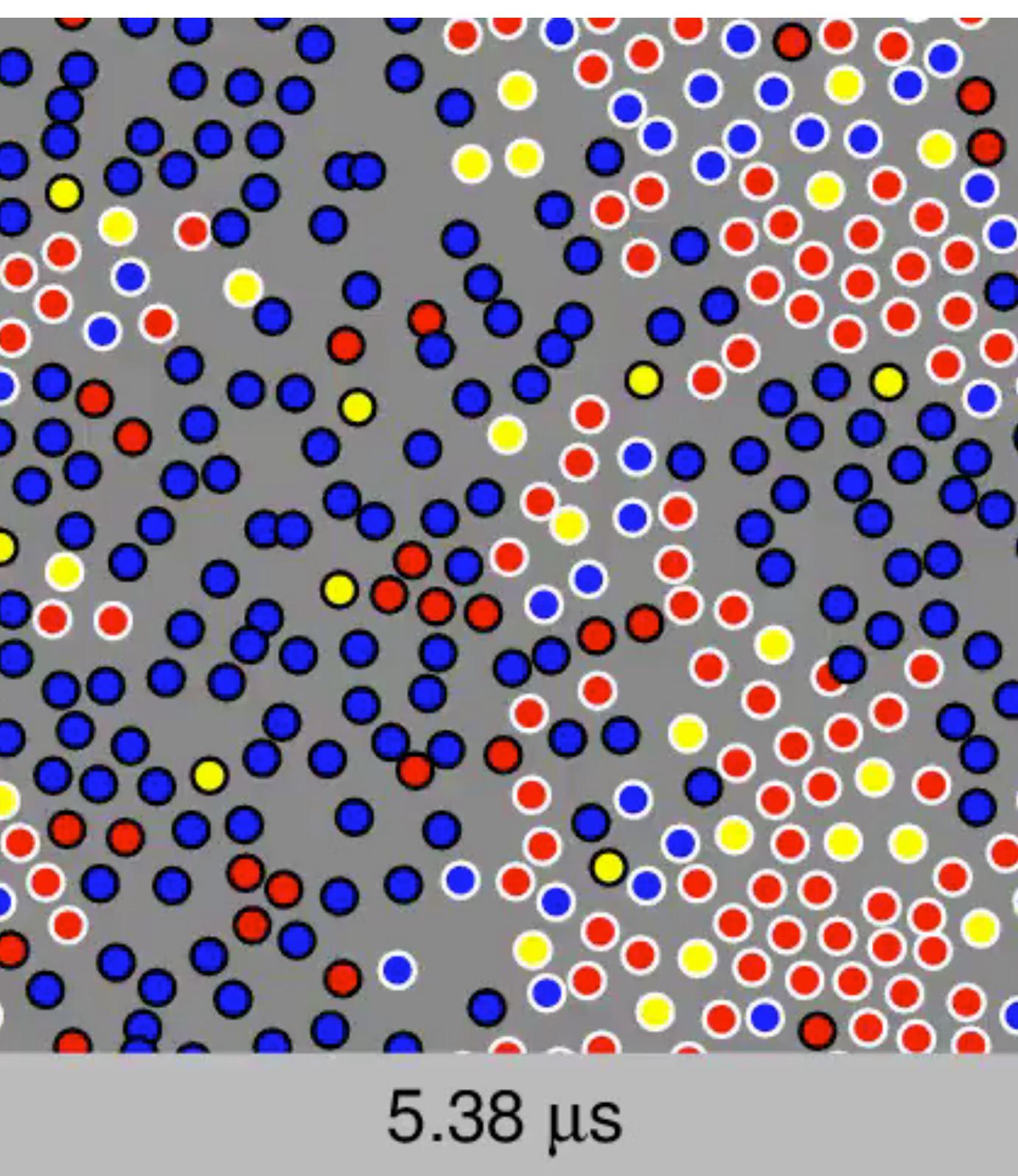


Veatch et al, ACS Chemical Biol, 3 2008

Liquid
ordered

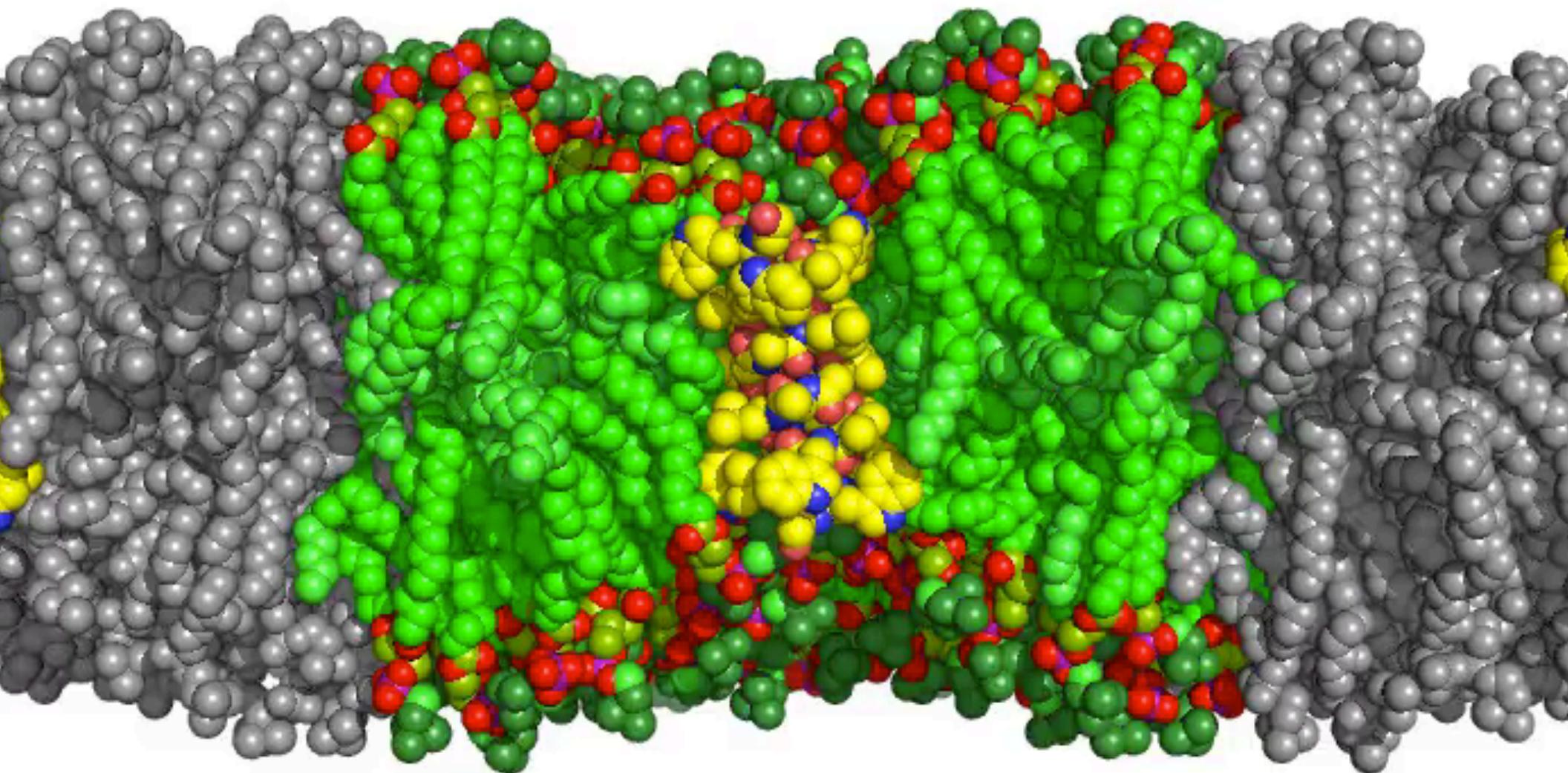
Liquid
disordered





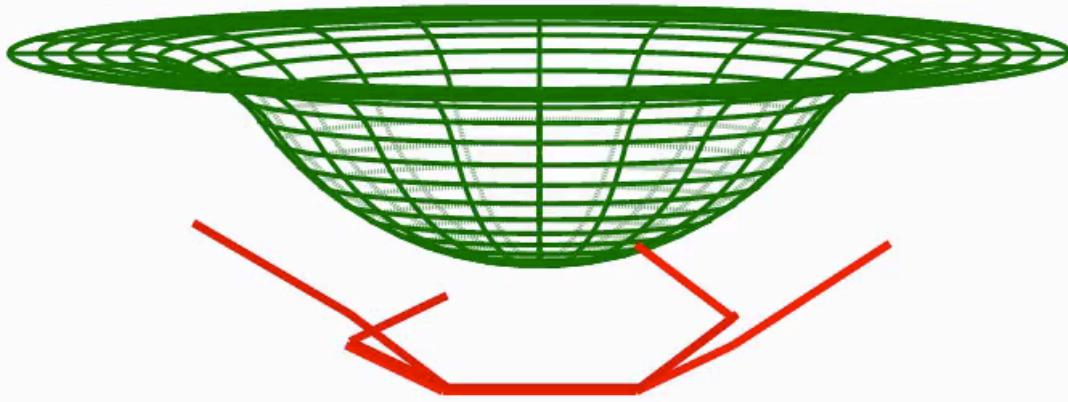
Unexpected
computational
technique:
Automated
“structure-free”
identification of
domains
by a hidden Markov
model

Material model of membrane deformation around a protein



Membrane
deformation
energetics:

lipid composition:
link of
molecular property
to
model energy



Molecular hypothesis

sphingomyelin
hydrogen bonding determines
membrane material properties

design simulation

Thermo. prop. vs.
concentration

analysis

visualization/theory

interpret results

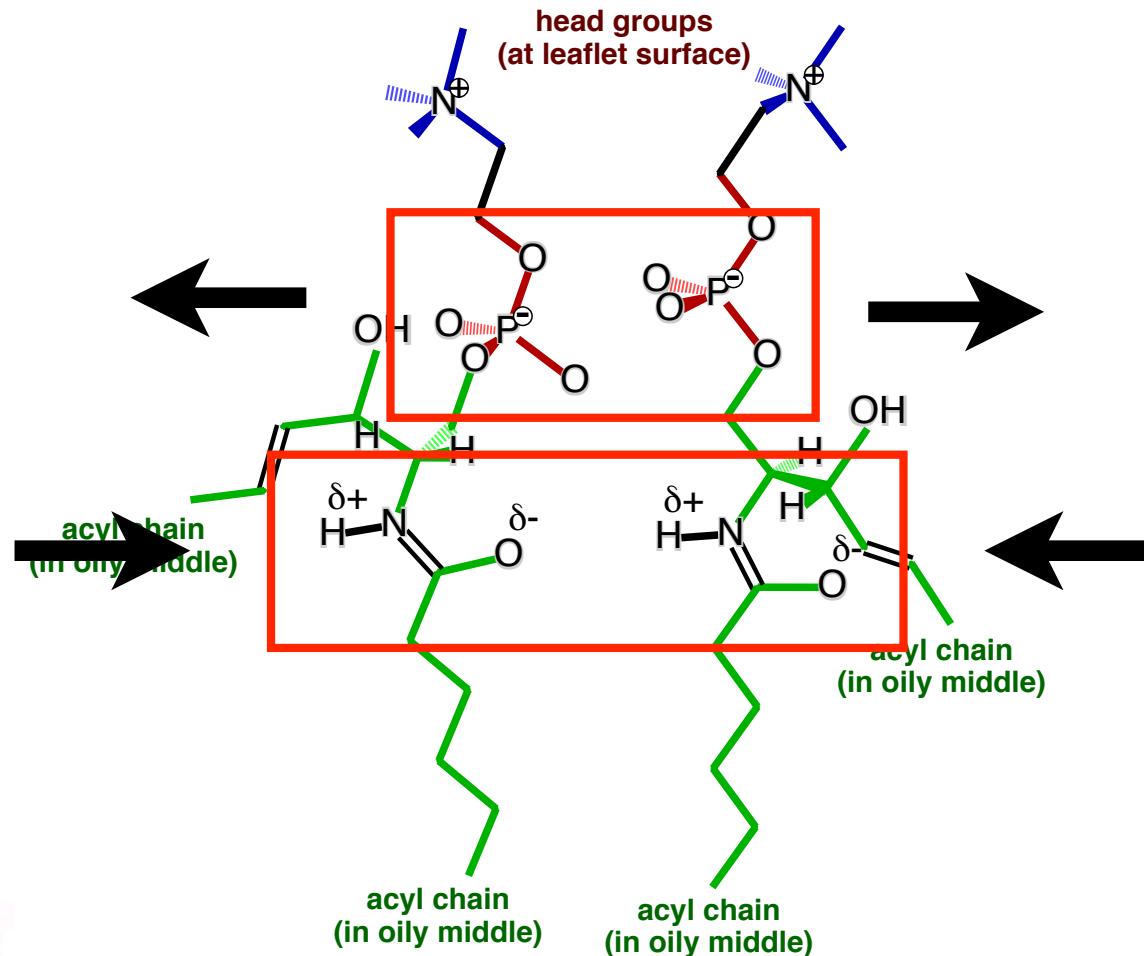
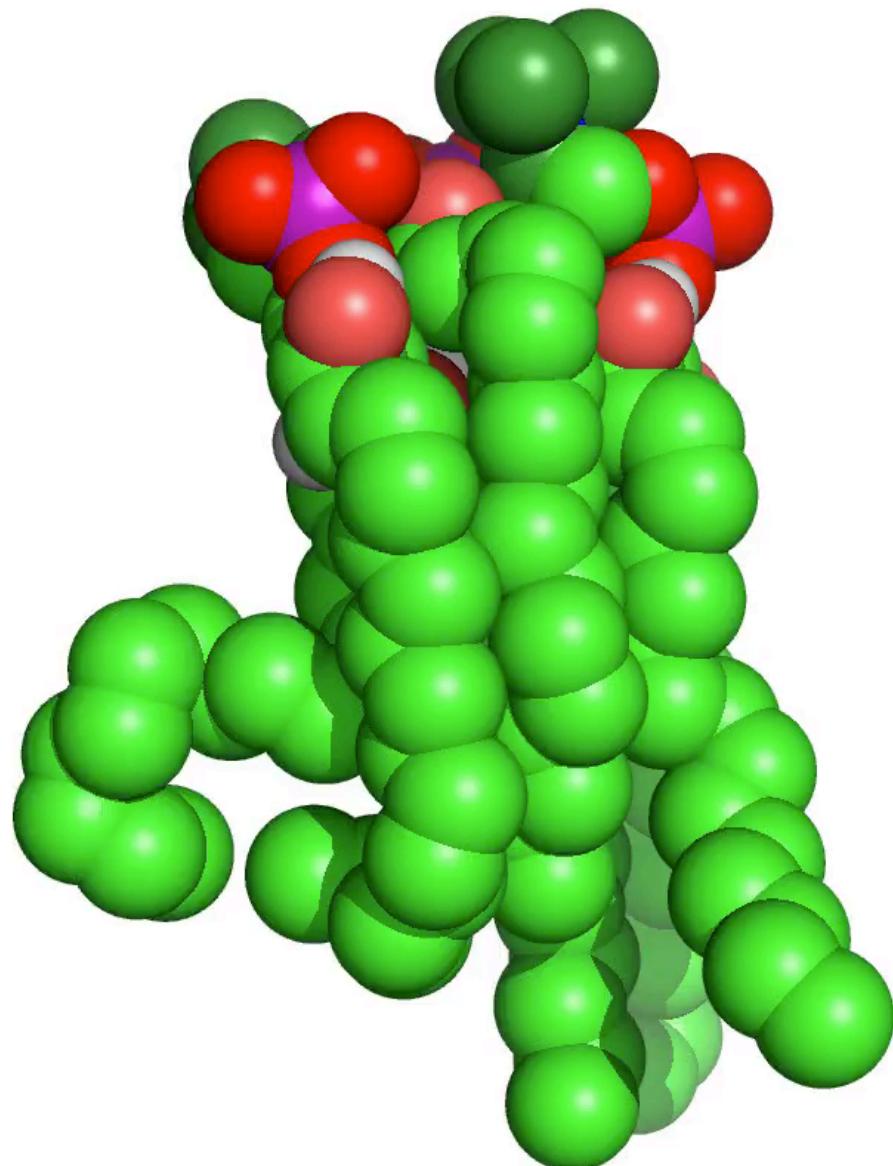
Develop model

validate

NMR experiment



Sphingomyelin



Large positive spontaneous curvature

Molecular hypothesis

design simulation

analysis

interpret results

validate

sphingomyelin
hydrogen bonding determines
membrane material properties

Thermo. prop. vs.
concentration

visualization/theory

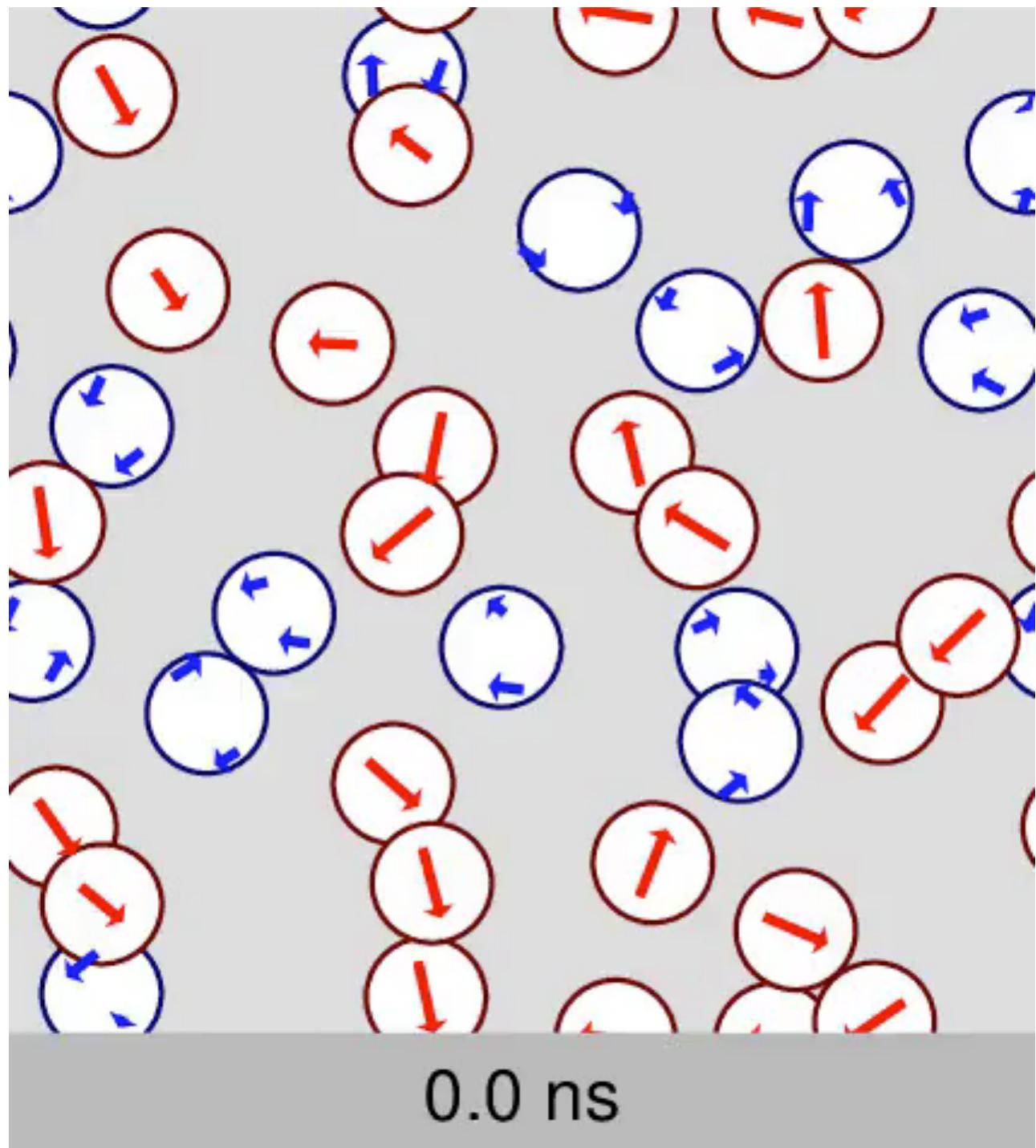
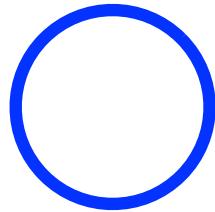
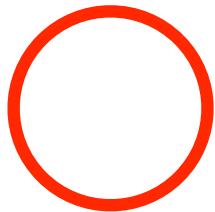
Develop model

NMR experiment

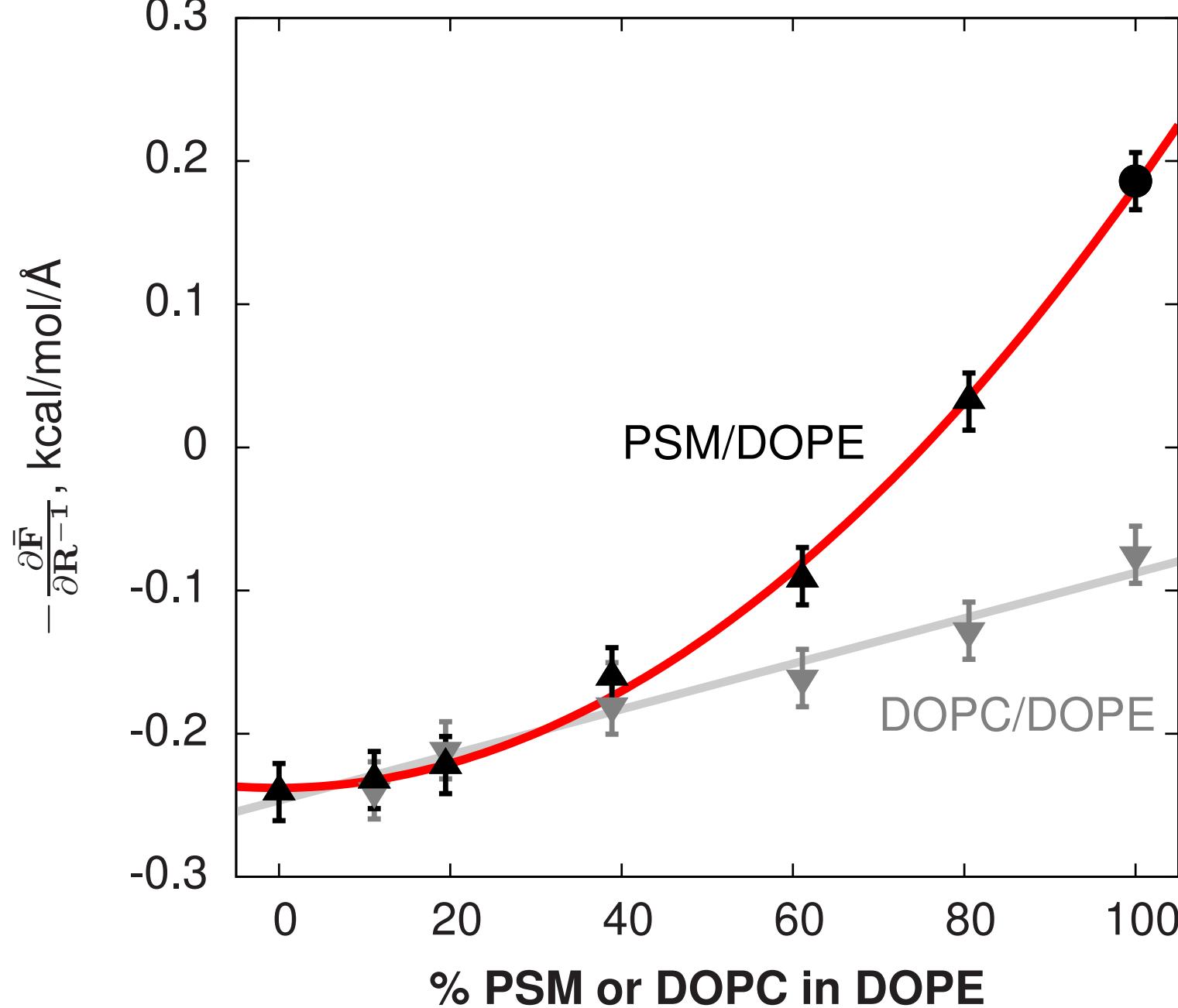


PSM

PE



Curvature preference vs. concentration



Molecular hypothesis

design simulation

analysis

refine/communicate
results

validate

sphingomyelin
hydrogen bonding determines
membrane material properties

Thermo. prop. vs.
concentration

visualization/theory

Develop model

NMR experiment



Model Development

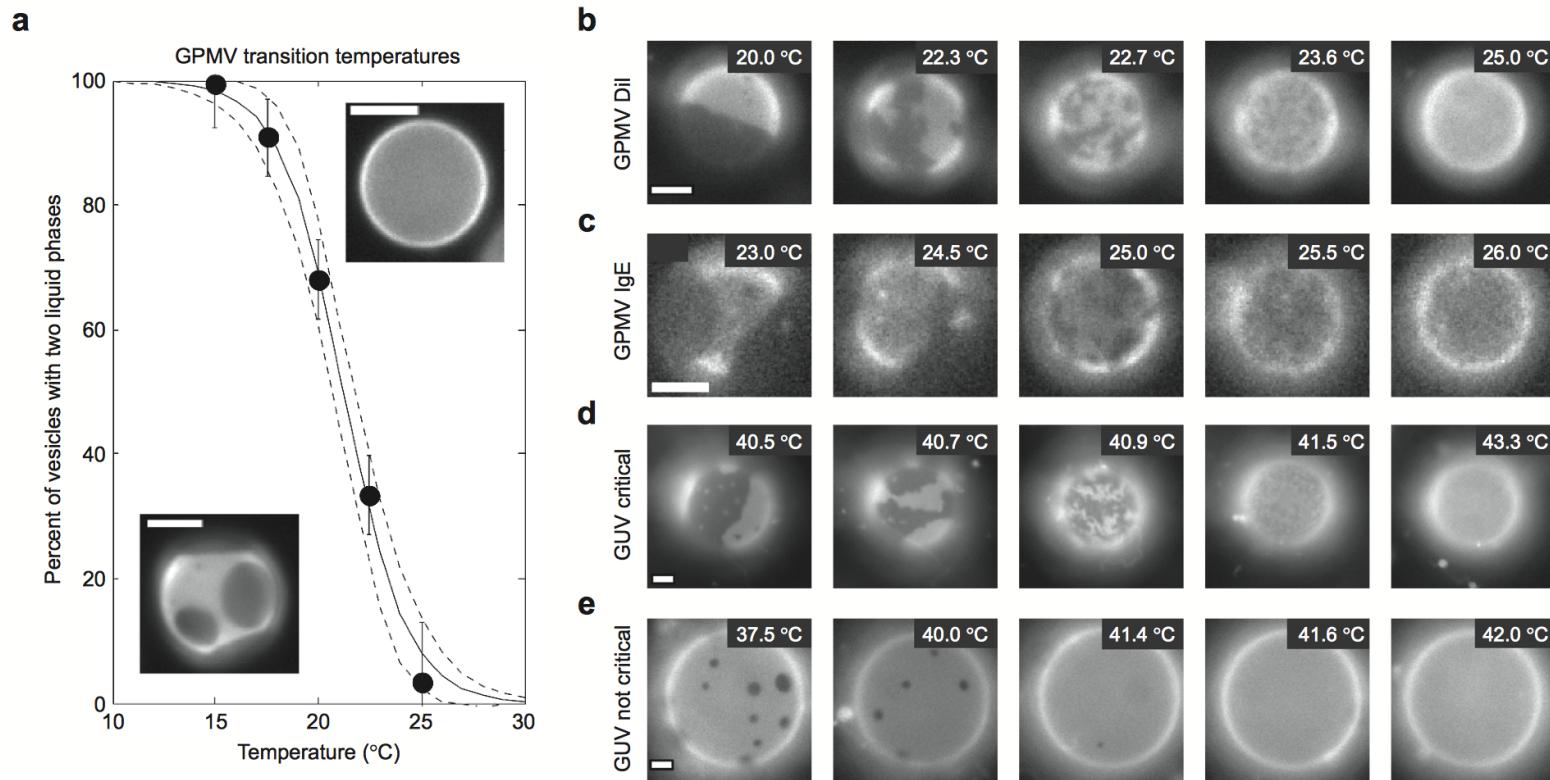
Caveolae (Enriched in sphingomyelin)



Parton/Simons Nature Reviews, 2007

Model Development

sphingomyelin: favors ordered domains



Veatch et al,
ACS Chemical Biol, **3** 2008

Mid-term goal:

Develop open-source membrane model
incorporating my theory/results (curvature, organization)
pluggable into systems biology/cell models

I will need:
creative coders with strong math background,
optimally: know how to develop with a group!

Molecular hypothesis

design simulation

analysis

interpret results

validate

sphingomyelin
hydrogen bonding determines
membrane material properties

Thermo. prop. vs.
concentration

visualization/theory

Develop model

NMR experiment



Faculty Search

Was not given an interview at any of the research universities
in the MD/DC area with multiple rounds of application:

Johns Hopkins,

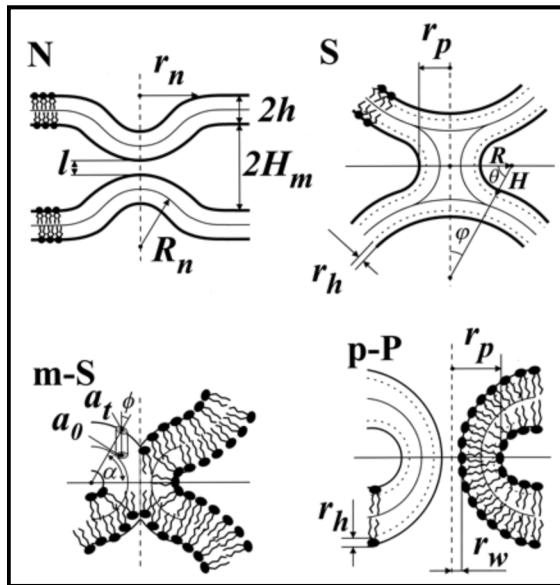
U Maryland (main, Baltimore County),

American

U Delaware

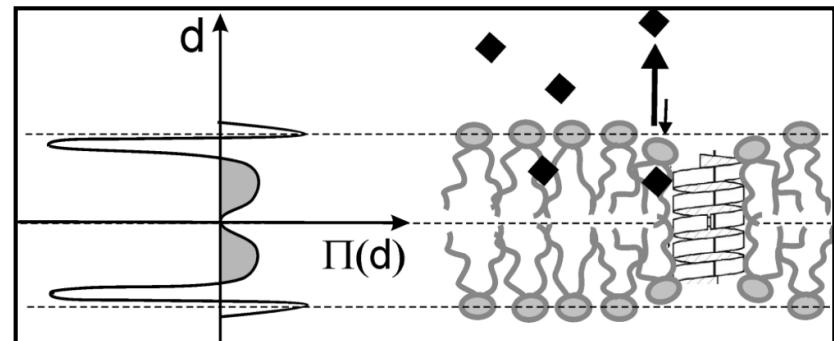
Even after I got a K award (a baby R01)
I was still rejected.

Where I was hired: My division at the NICHD includes:

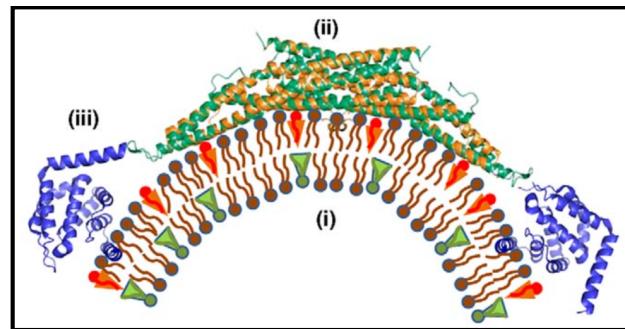


Joshua Zimmerberg
Program Head,
Physical Biology
viral membrane fusion

all experimentalists



Sergey Bezrukov
Chief, Section on Molecular Transport
membrane regulation of transport



Leonid Chernomordik
Chief, Section on Membrane Biology
membrane remodeling in biology

Director



Dr. Francis Collins

previously led
humane genome project

my building!
(schd for demolition
2013)



~1200 Investigators
10% strong computational
interest

Another American with Ebola will be flown to
NIH in Bethesda

A Print 99 Save for Later Reading List



Alberto Cavigli is beginning the multi-step process of identifying antibodies against Ebola in samples from vaccinated volunteers at the National Institutes of Health in Bethesda, Md., on Feb. 4. (Cliff Owen/AP)

By Julie Zauzmer March 12, 2015 Follow @JulieZauzmer

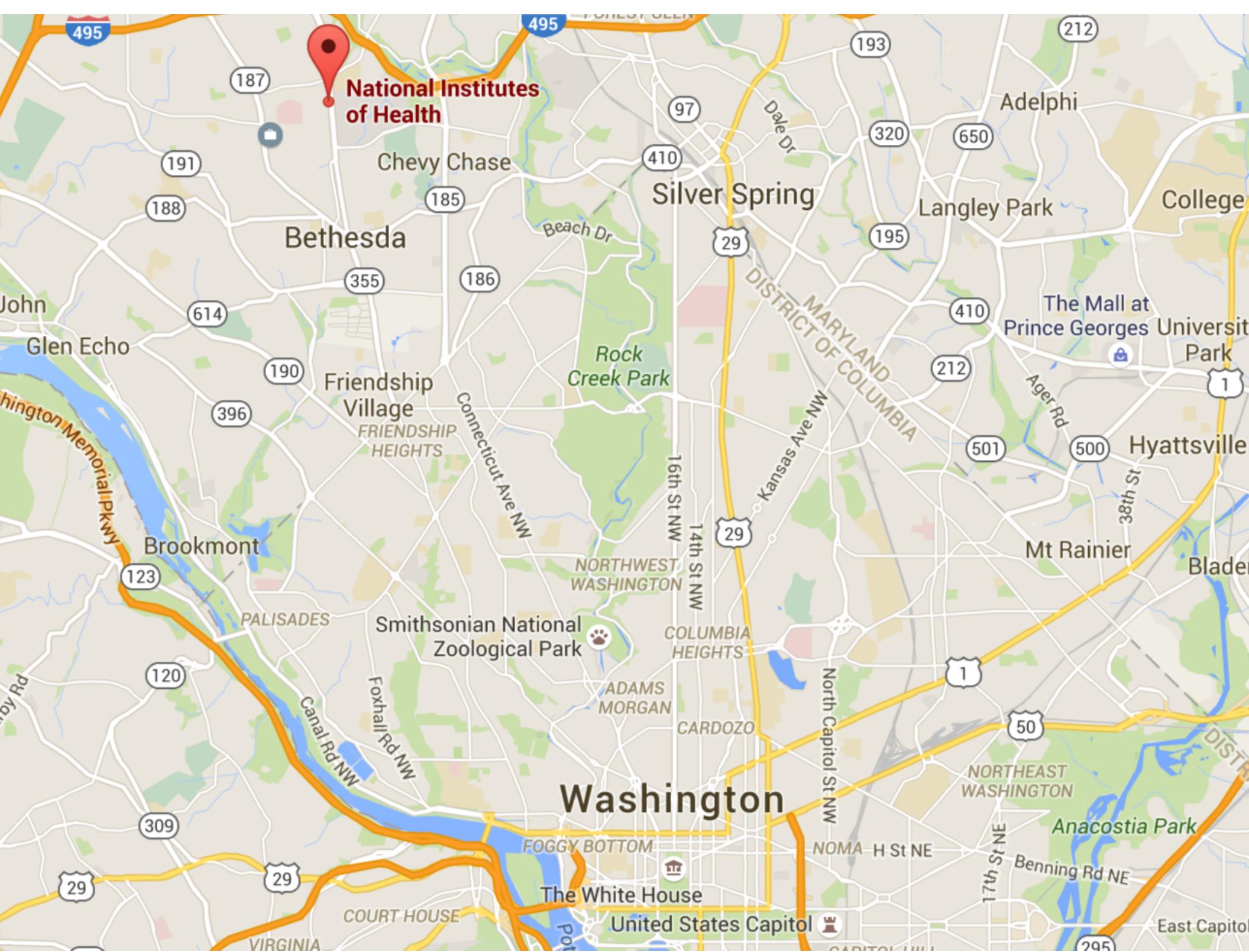
An American health-care worker has come down with Ebola and will be transported to the National Institutes of Health in Bethesda for treatment, NIH announced in a news release on Thursday.

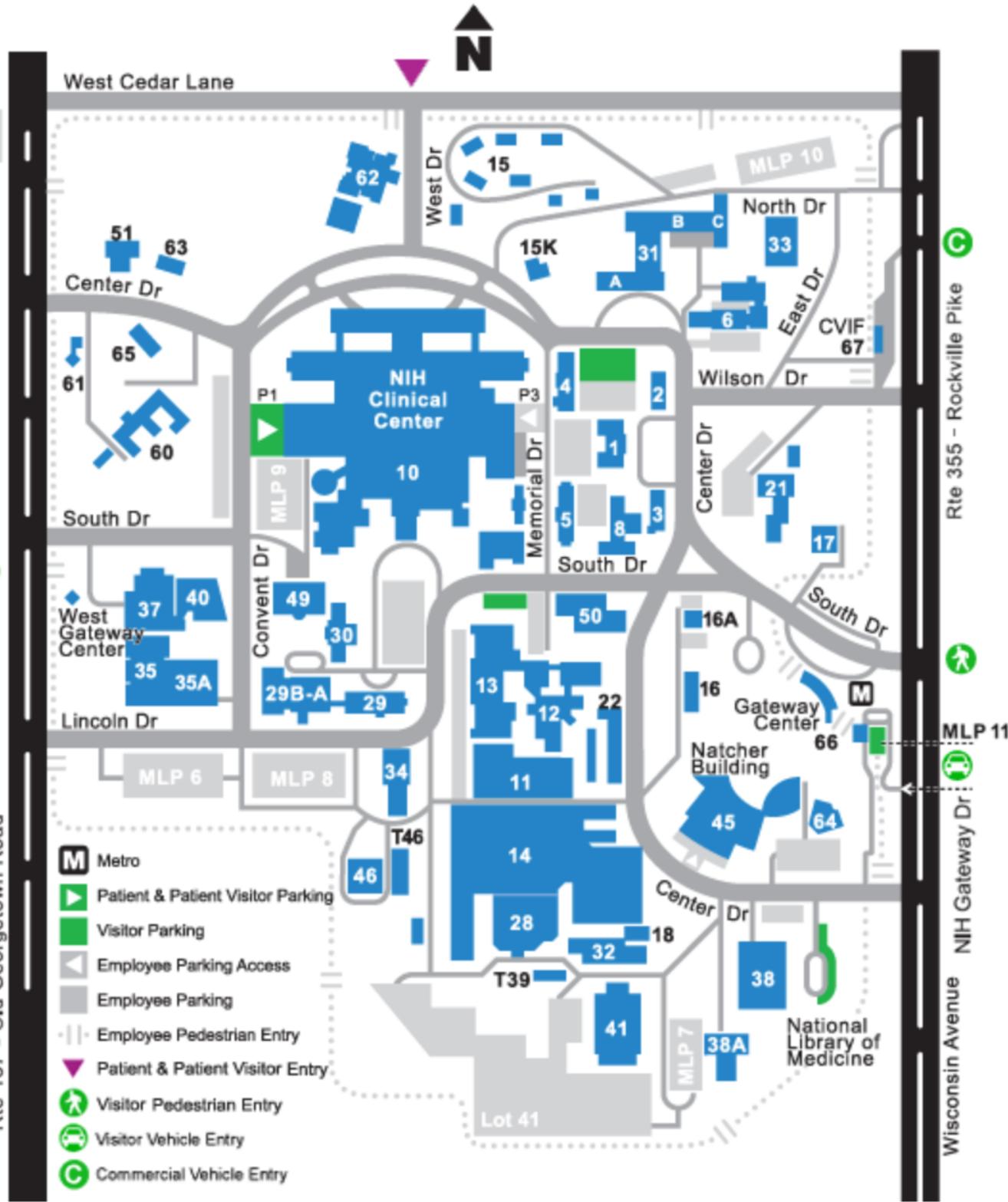
nextday
BLINDS

Spring is in the air.
Quality is in the details.

SCHEDULE APPOINTMENT >

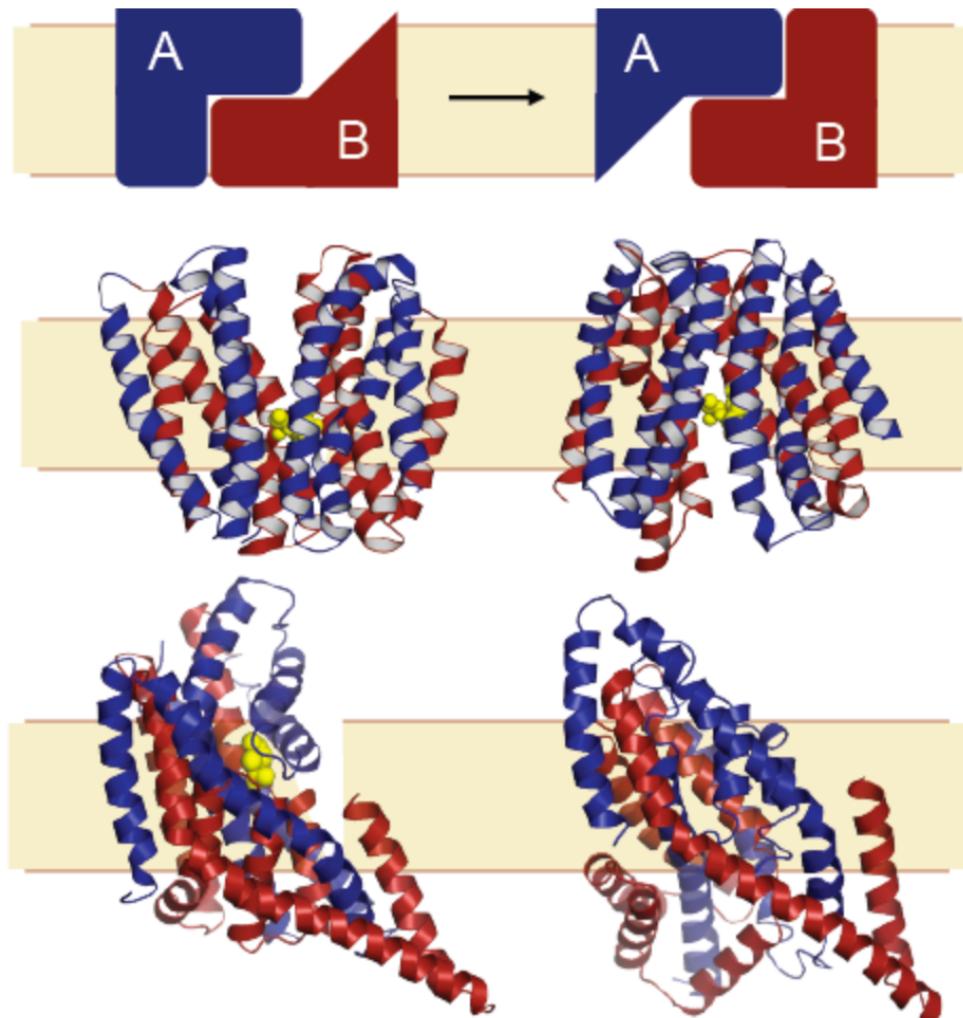
Most Read





Lucy Forrest

The Pseudo-Symmetry of Transporter Structural Changes



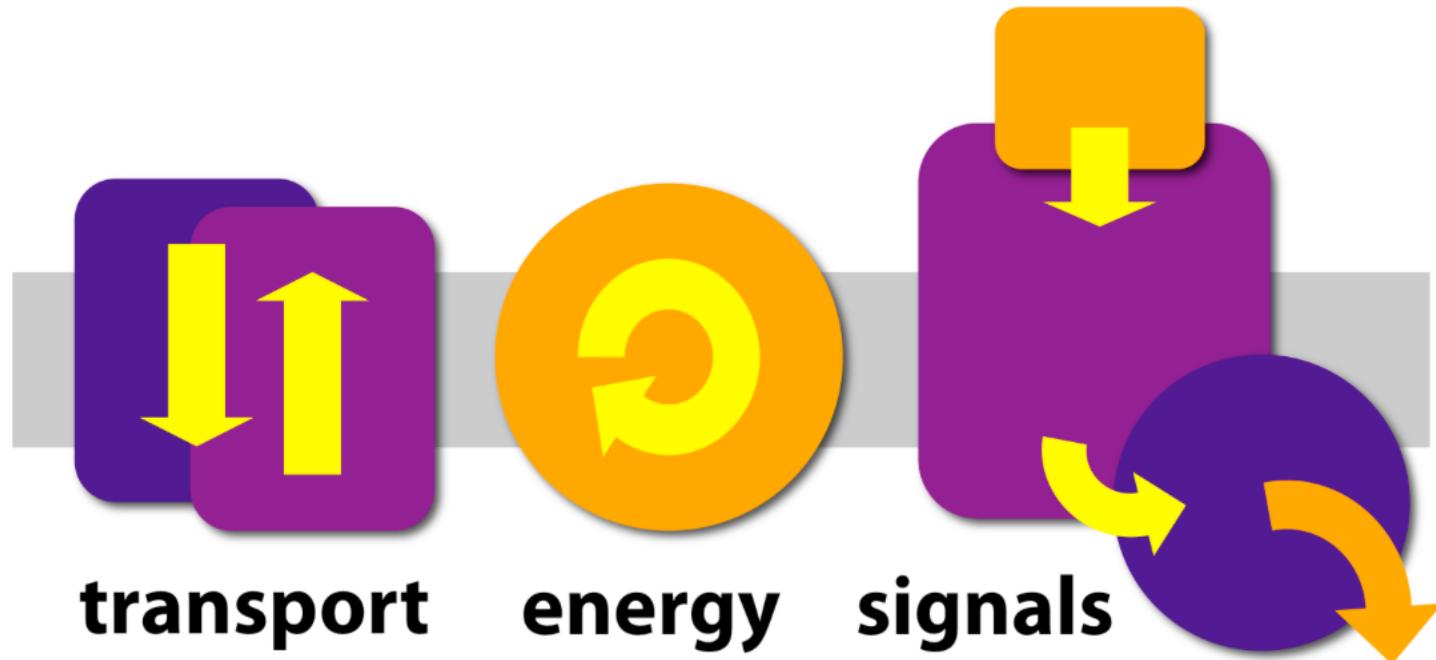
Relating structure/
sequence
to mechanism

AlignMe
software

[http://
www.bioinfo.mpg.de/
AlignMe/faq.html](http://www.bioinfo.mpg.de/AlignMe/faq.html)

<https://neuroscience.nih.gov/ninds/Faculty/Profile/lucy-forrest.aspx>

Jose Faraldo-Gomez



Specific channel function
free energy methodology

<http://www.faraldolab.org>

Robert Best

- Balance of secondary structure in forcefields.
- Modeling protein folding kinetics using reduced diffusion to capture off-path dynamics.
- Protein folding mechanisms

Part of chemical physics lab including
Attila Szabo
Ad Bax (NMR)

<http://spin.niddk.nih.gov/best/home.html>

Computation at the NIH



Partition	FreeNds	FreeCPUs	Per-Node Resources			
			Cores	CPUs	Mem	Disk
norm*	0/310	2542/7754	16	32	124g	800g
unlimited	0/14	0/234	16	32	124g	800g
niddk	26/86	1168/2736	16	32	124g	800g
ibfdr	39/191	1248/6112	16	32	61g	800g
ibqdr	2/102	64/3264	16	32	30g	400g
ibqdr	38/99	1216/3168	16	32	61g	400g
gpu	12/14	164/168	12	12	45g	600g
gpu	0/20	96/608	16	32	124g	800g
gpu	0/4	16/112	16	32	124g	800g

Biowulf



ca. 5000 cores
for Pastor/Brooks labs

<http://www.lobos.nih.gov>

Opportunities now

<https://www.training.nih.gov/programs/sip>

Summer internships: high school/undergrad/grad

EDUCATION LEVEL COMPLETED	MONTHLY STIPEND
HIGH SCHOOL	Before Graduation
	\$1,740
UNDERGRADUATE	After Graduation
	\$1,940
	After 1 year
GRADUATE	After 2 years
	\$2,040
	After 3 years or >
GRADUATE	\$2,140
	\$2,240
	Less than 1 year
	\$2,340
	After 1 year
	\$2,660
	After 2 years
	\$2,960
	After 3 years or >
	\$3,160

Opportunities now

https://www.training.nih.gov/programs/postbac_irta

Application is probably equivalent to grad school.
You must convince an NIH PI to offer you a job,
probably with a convincing personal appeal.

POSTBACCALAUREATE IRTA STIPEND RANGES FOR ALL AREAS -
Discretionary for Montana (RML-NIAID)

YEARS OF EXPERIENCE	INITIAL STIPEND	SECOND YEAR
0-1 year	\$27,700	\$29,700
1-2 years	\$29,700	\$31,000

Anxieties during a long job search

Could I work on a problem I was not obsessed with?

Had I become dependent on my obsession to do things others couldn't do?

Could I work in a group?

I've developed a lone-wolf short-term focused coding strategy

Could I accept a solution that wasn't correct, but that worked