

Broadly neutralizing antibodies and a rational approach to HIV vaccine design

Dennis Burton

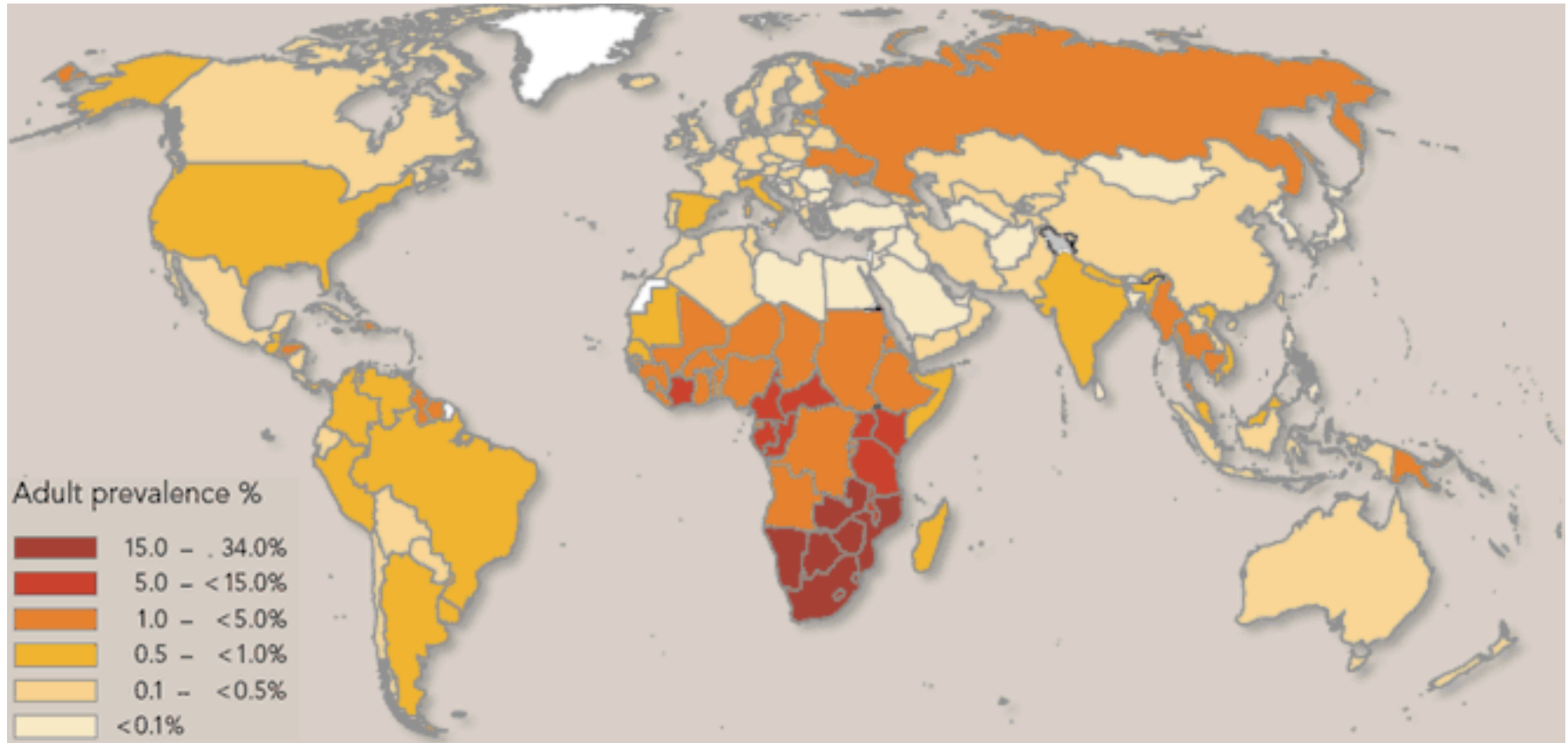
**Dept of Immunology & Microbial Science and
IAVI Neutralizing Antibody Center & Center
for HIV/AIDS Vaccine Immunology and
Immunogen Discovery
(CHAVI-ID), The Scripps Research Institute**

Ragon Institute of MGH, Harvard and MIT

**AIDS, Science & Society, UCSD, La Jolla,
Nov 20th 2014**

A global view of HIV infection

33 million people living with HIV, 2009



AIDS compromises human development ...

- Reversing the spread of AIDS is one of the eight U.N. Millennium Development Goals
- The AIDS pandemic undermines:
 - Poverty reduction
 - Improvements in child and maternal health
 - Improvements in nutrition
 - Gains in basic education
 - Control of other infectious diseases

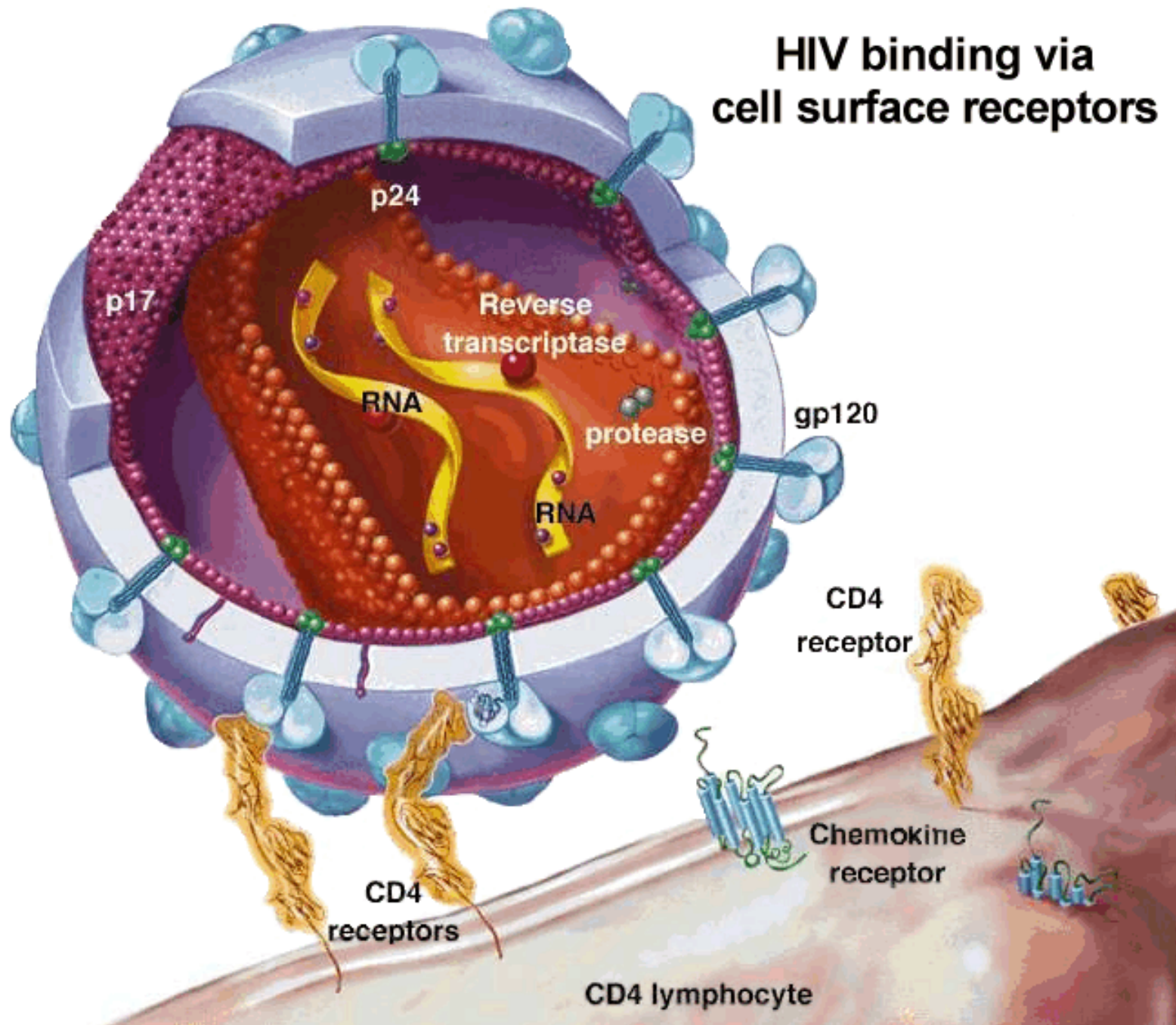


The power of vaccines

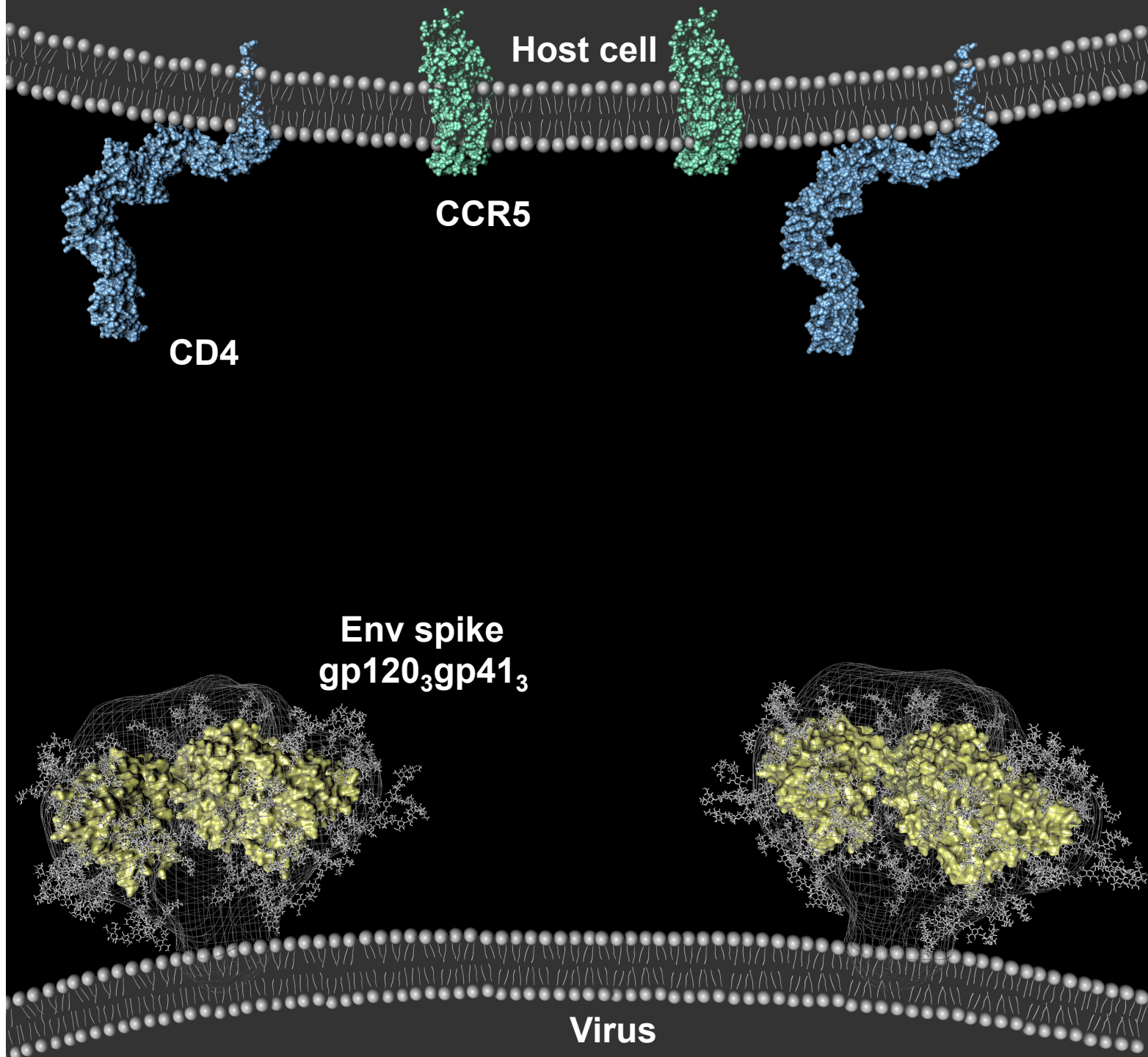
- **Approximately 300m individuals died of smallpox in the 20th Century.**
- **Following the eradication program based on vaccination, the last person died of smallpox on 9/11/1978.**

Oldstone, MBA “Viruses, Plagues & History”, 1998, Oxford University Press.

HIV binding via cell surface receptors



The players in HIV entry.

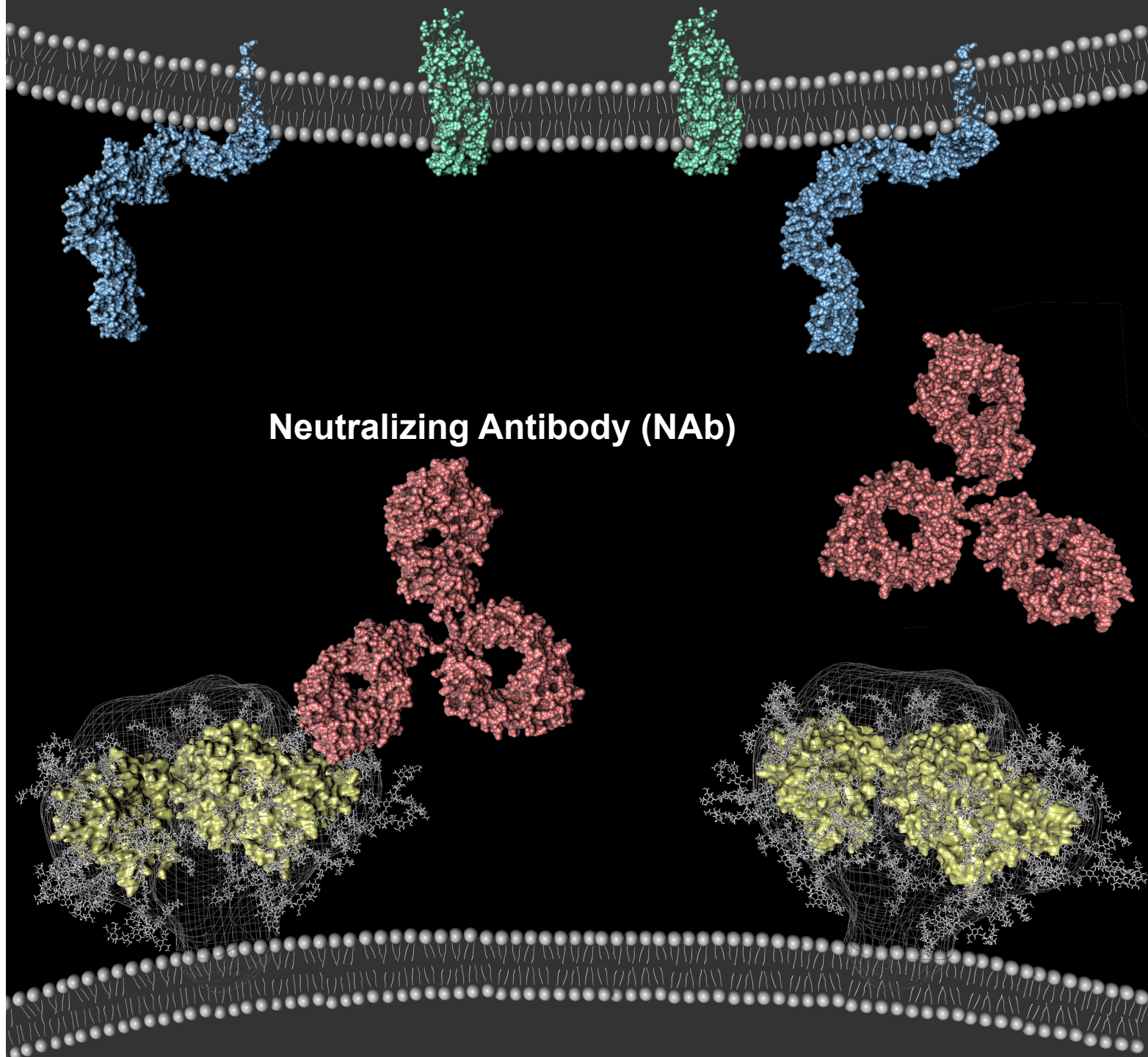


**NAbs block
viral entry.**

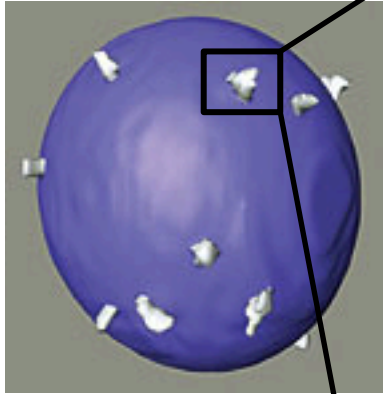
**Binding to
Env spike is
necessary &
sufficient for
neutralizn.**

**Vaccine
problem is
to induce
Abs that
bind to Env
spikes.**

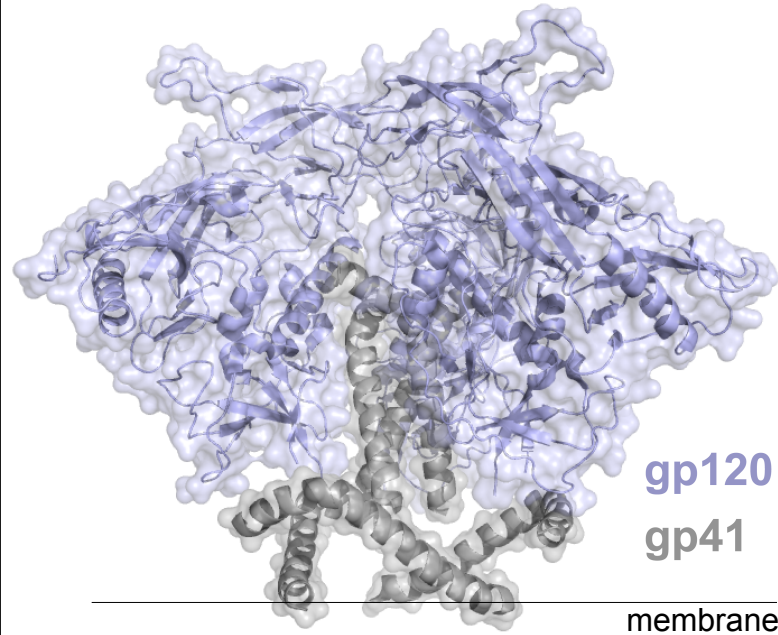
**HIV
variability
dictates
need for
broadly
NAbs**



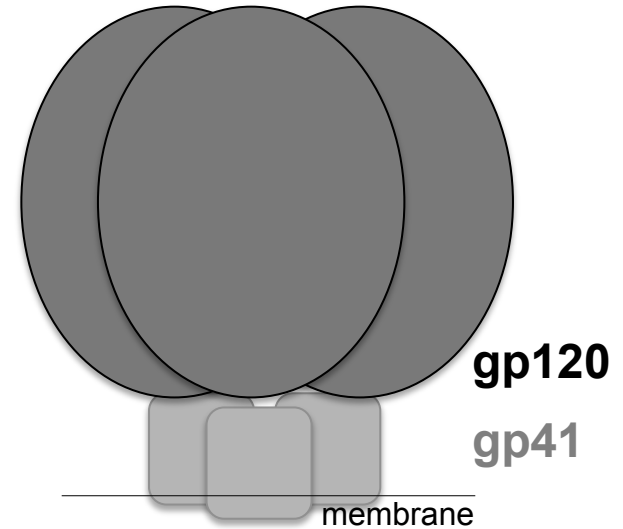
The HIV Env spike is a metastable trimer of heterodimers



HIV



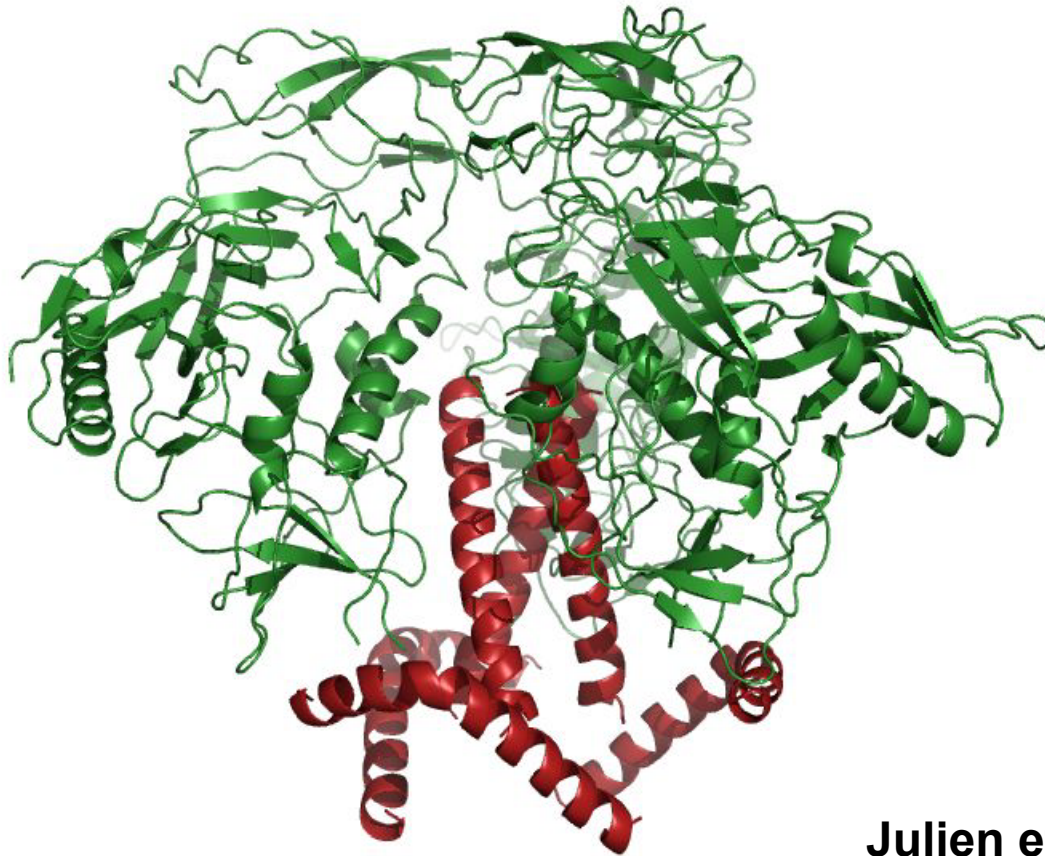
**Structural representation
(BG505 SOSIP.664)**



Cartoon illustration

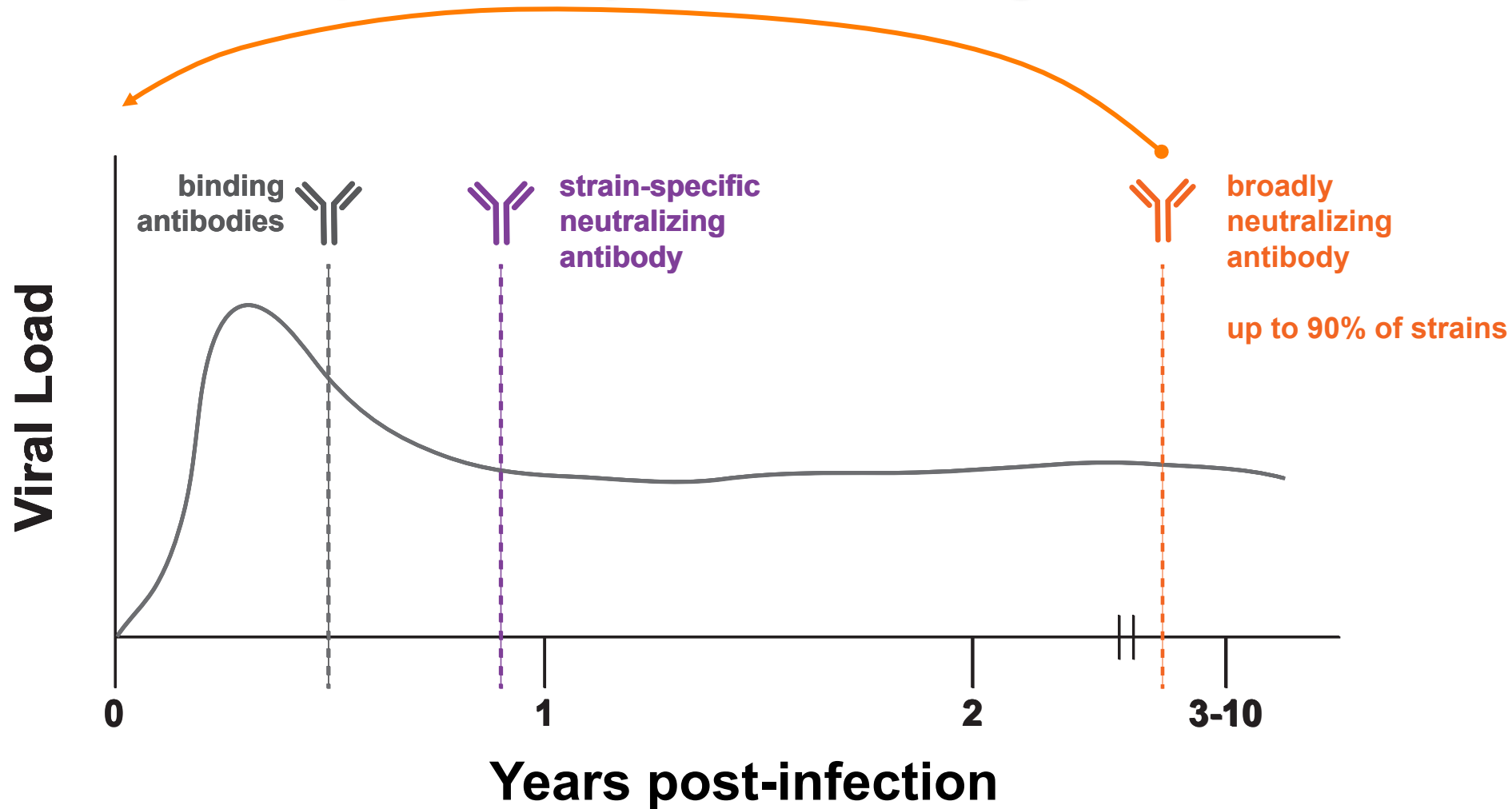
HIV Env Trimer

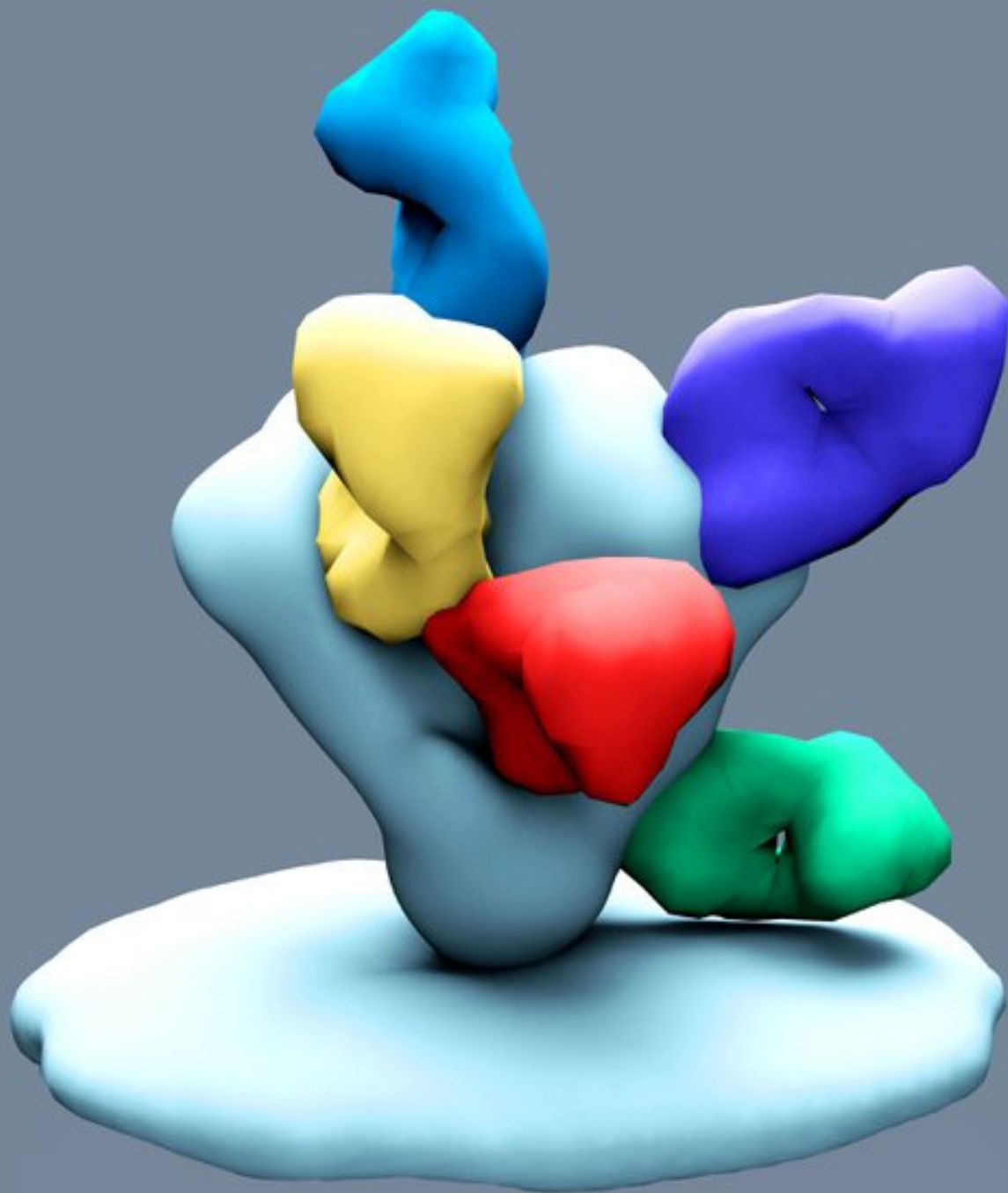
Jardine, Menis & Schief



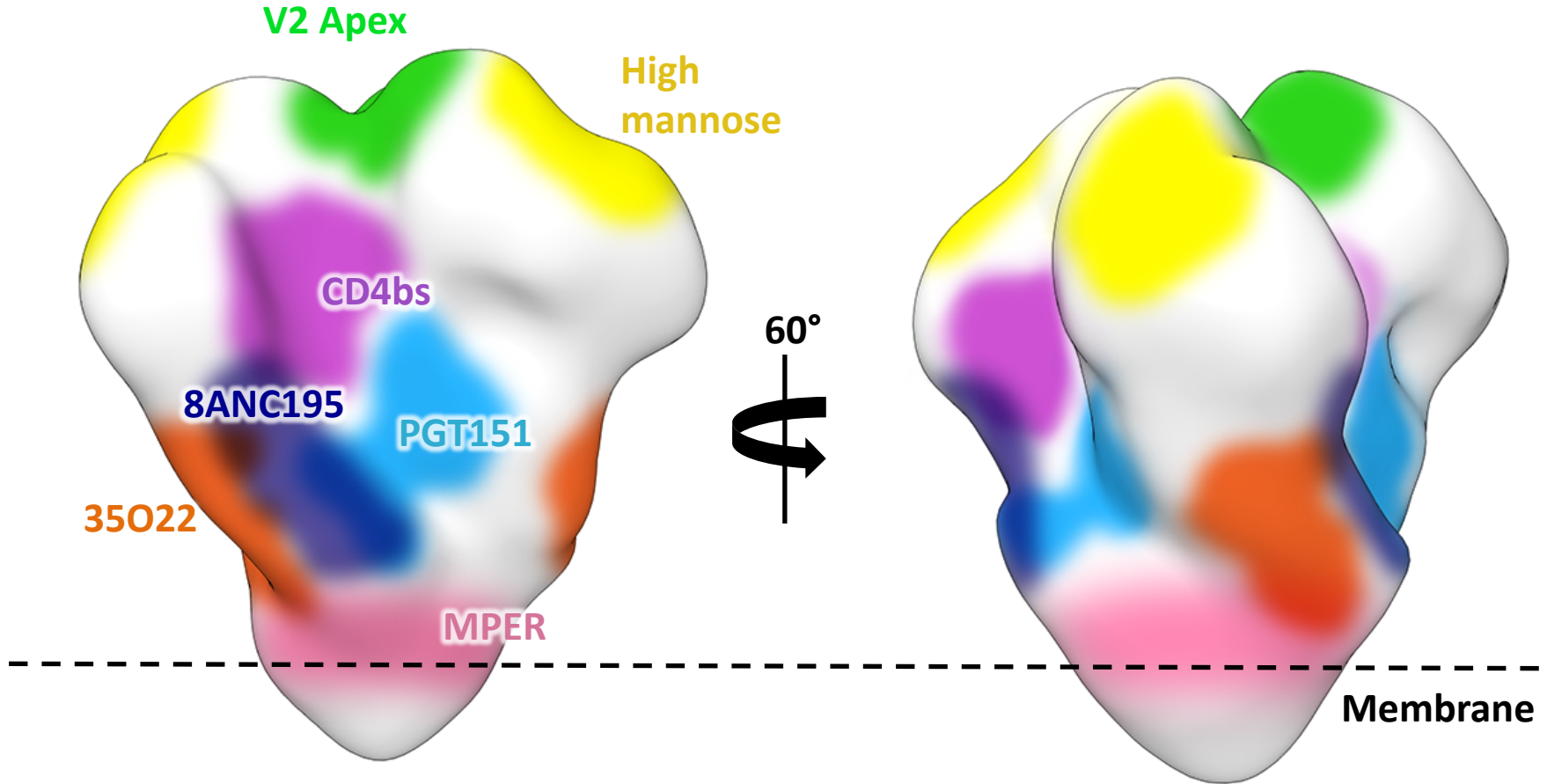
Julien et al, 2013, Science

Despite the evasion tactics by HIV Env, up to 20% of chronically-infected individuals develop broadly neutralizing antibodies

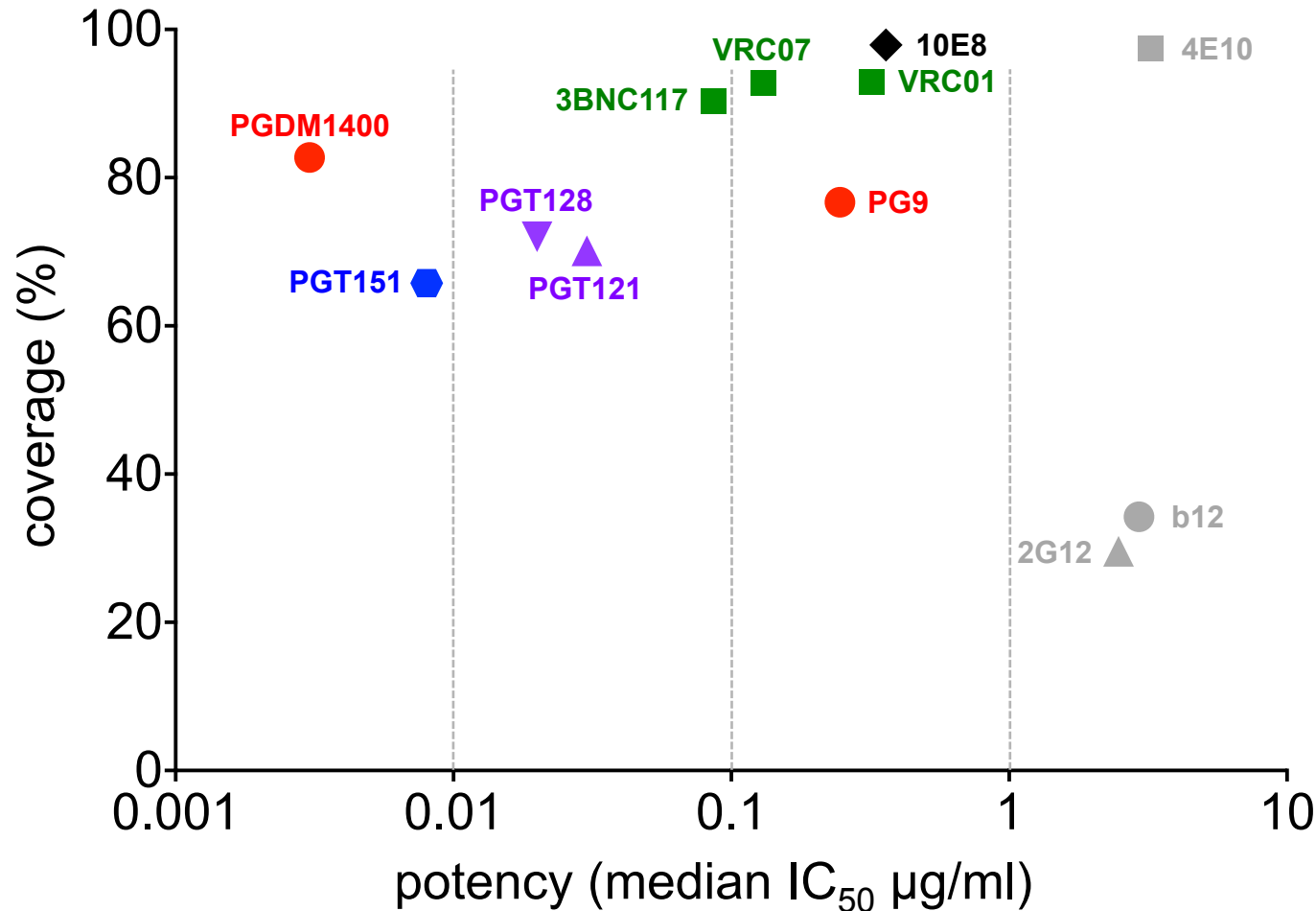




Broadly neutralizing MAbs define Env sites of vulnerability



Breadth and potency of prototype HIV bnAbs



Critical: better donors & HuMAb isolation methods

Example of broad serum neutralization by an IAVI Protocol G “elite neutralizer”

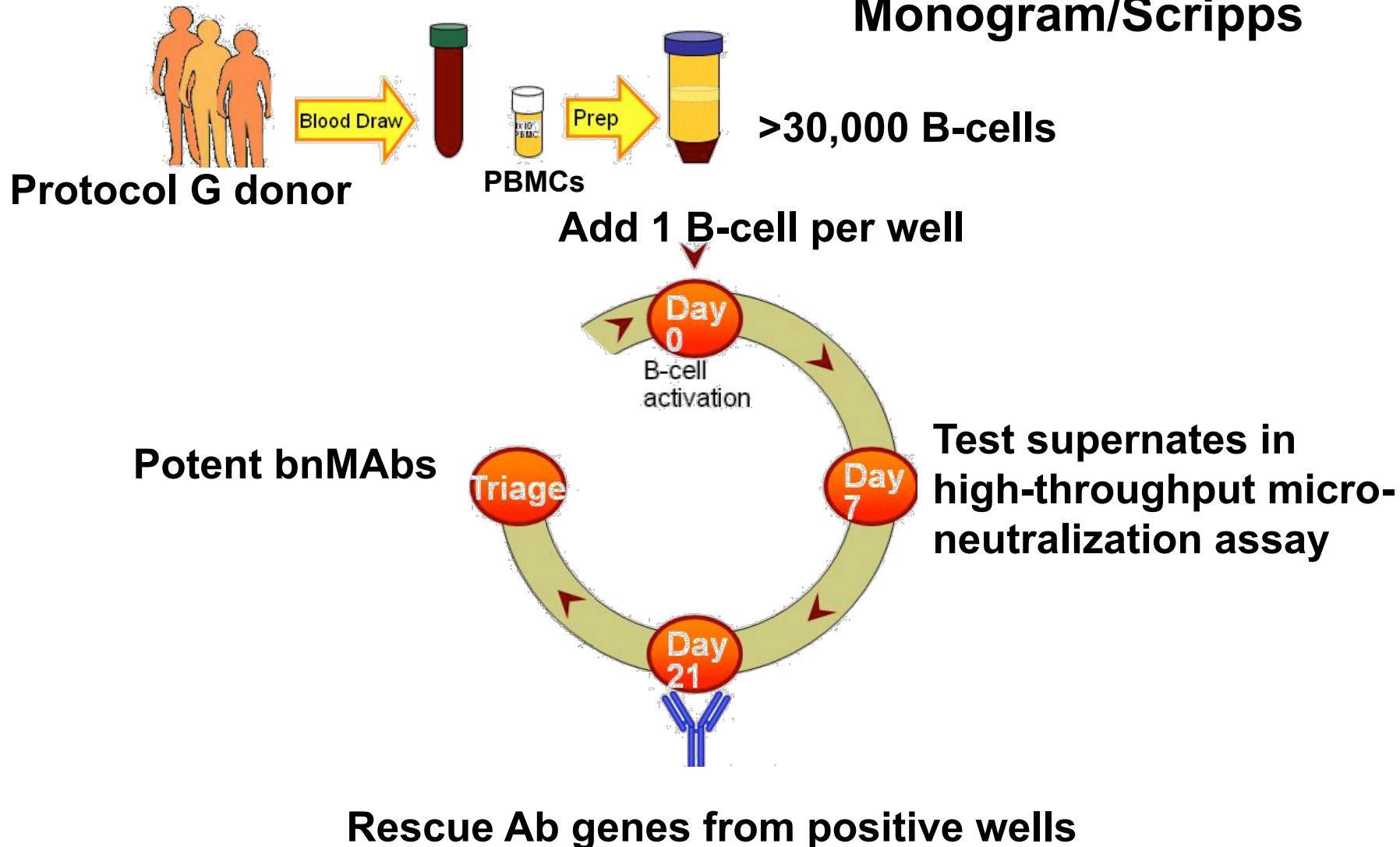
	Clade A	Clade B		Clade C		CRF01_AE
Virus	94UG103	92BR020	JRCSF	IAVI C22	93IN905	92TH021
Serum IC ₅₀	900	900	2700	2700	2700	2700

IC₅₀ = serum dilution giving 50% neutralization of virus isolate in Monogram pseudovirus assay

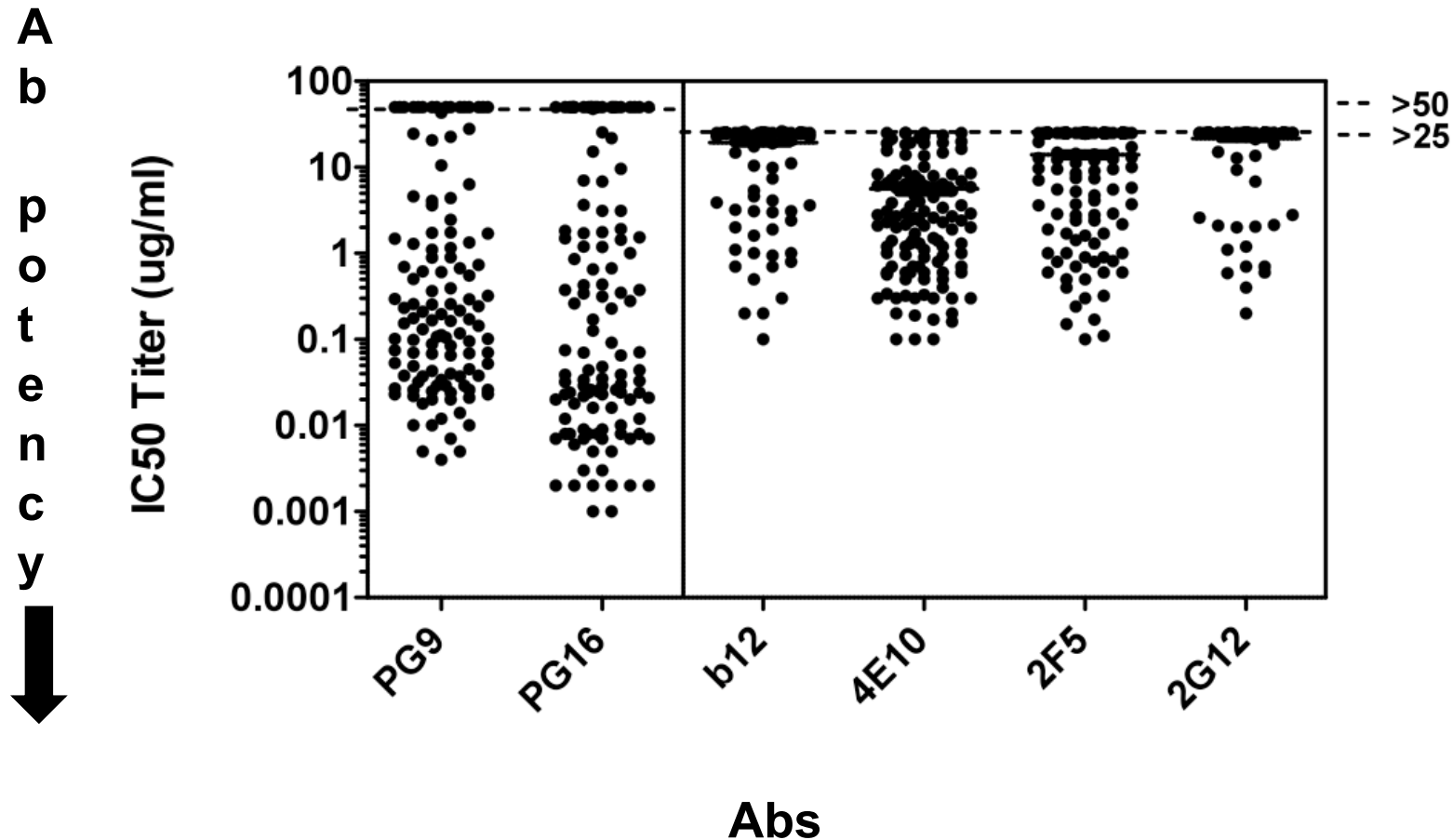
(Simek et al, J Virol, 2009)

Isolation of bnMAbs from Protocol G donors

IAVI/Theraclone/
Monogram/Scripps



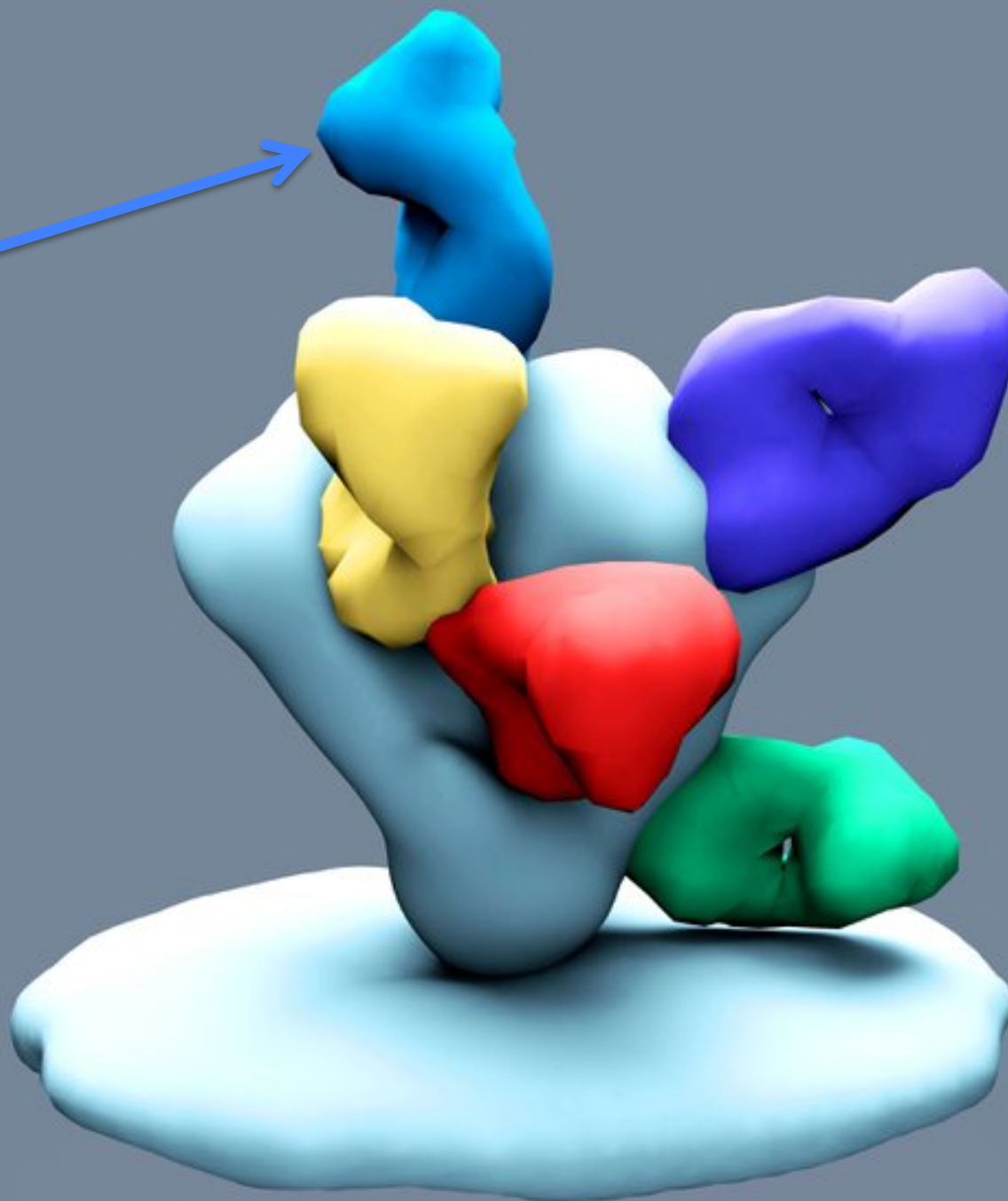
Broad and potent MAbs PG9 and PG16 isolated by direct neutralization screening approach (Walker et al, Science, 2009)



Each dot represents one virus (118 viruses)

Mike Seaman

PGDM1400



PGT121

PGT121 and PGDM1400 neutralization together on 106-virus panel

	PGT121	PGDM1400	PGT121+PGDM1400
Median IC50, µg/ml	0.015	0.005	0.007
% breadth	63	80	98

Sok, van Gils et al, PNAS, in press

PGT 121 effectively protects against high-dose mucosal SHIV challenge in macaques

Challenge SHIV	Challenge route	Complete protection Lowest dose (mg/kg)	50% animals protected dose (mg/kg)	Collabr
162P3	IVAG	1	~0.2	Watkins
162P3	IVAG	2	~0.4	Barouch
AD8E0	IR	1	-	Martin
DH12-V3AD8	IR	0.2	-	Martin
B founder	IR	10	-	Lifson

Potent bnMAbs have therapeutic effects in the SHIV/macaque model

nature

November 2013
Volume 503
Issue 7475

Therapeutic efficacy of potent neutralizing HIV-1-specific monoclonal antibodies in SHIV-infected rhesus monkeys

Dan H. Barouch, James B. Whitney, Brian Moldt, Florian Klein, Thiago Y. Oliveira, Jinyan Liu, Kathryn E. Stephenson, Hui-Wen Chang, Karthik Shekhar, Sanjana Gupta, Joseph P. Nkolola, Michael S. Seaman, Kaitlin M. Smith, Erica N. Borducchi, Crystal Cabral, Jeffrey Y. Smith, Stephen Blackmore, Srisowmya Sanisetty, James R. Perry, Matthew Beck, Mark G. Lewis, William Rinaldi, Arup K. Chakraborty, Pascal Poignard, Michel C. Nussenzweig and Dennis R. Burton

nature

November 2013
Volume 503
Issue 7475

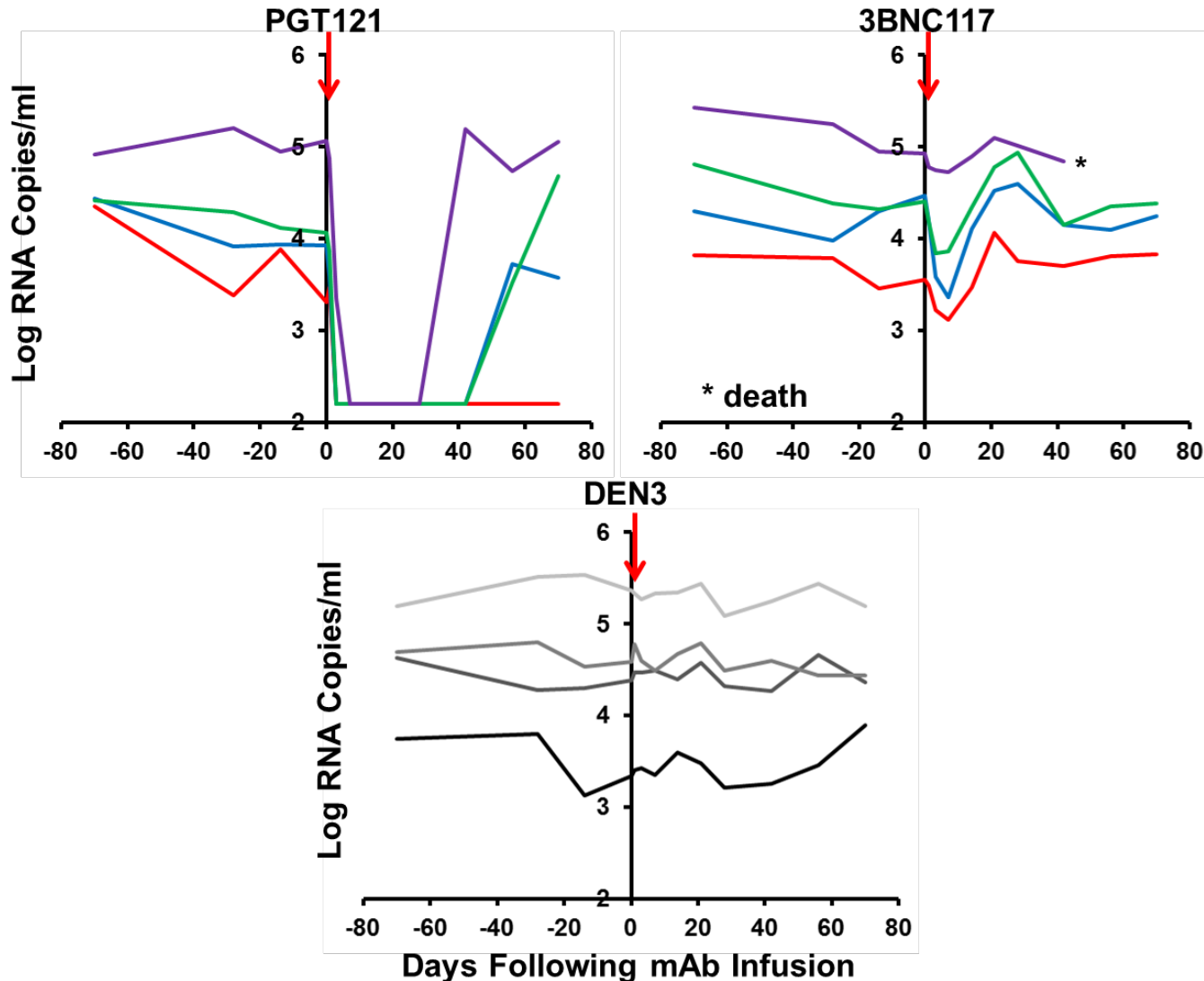
Antibody-mediated immunotherapy of macaques chronically infected with SHIV suppresses viraemia

Masashi Shingai, Yoshiaki Nishimura, Florian Klein, Hugo Mouquet, Olivia K. Donau, Ronald Plishka, Alicia Buckler-White, Michael Seaman, Michael Piatak Jr, Jeffrey D. Lifson, Dimitar S. Dimitrov, Michel C. Nussenzweig, and Malcolm A. Martin

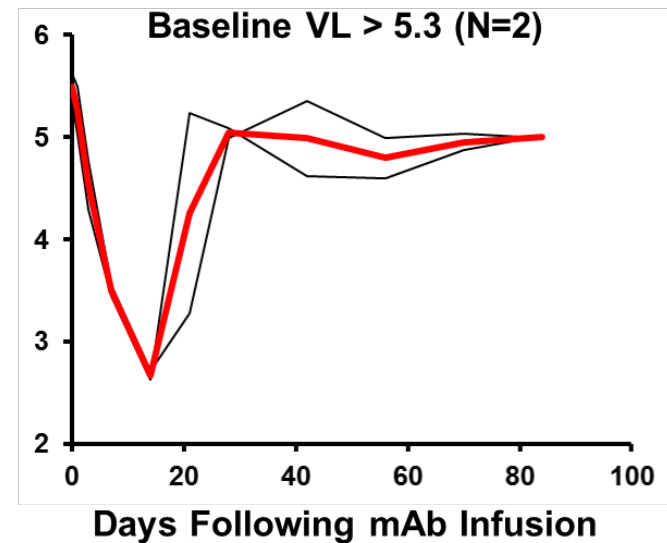
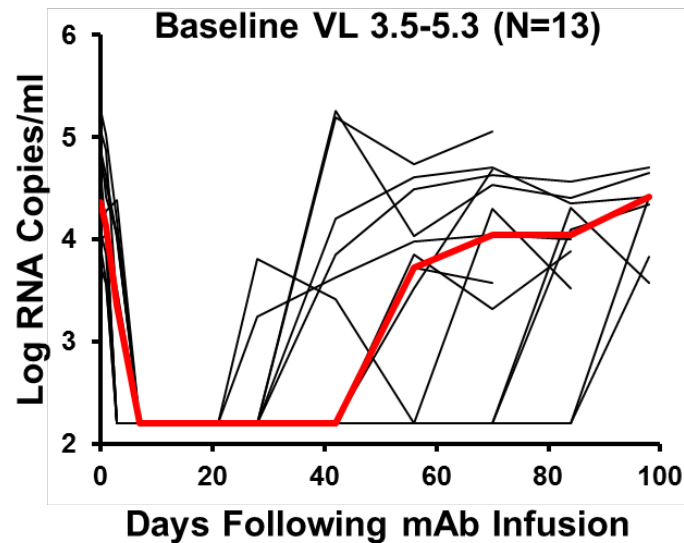
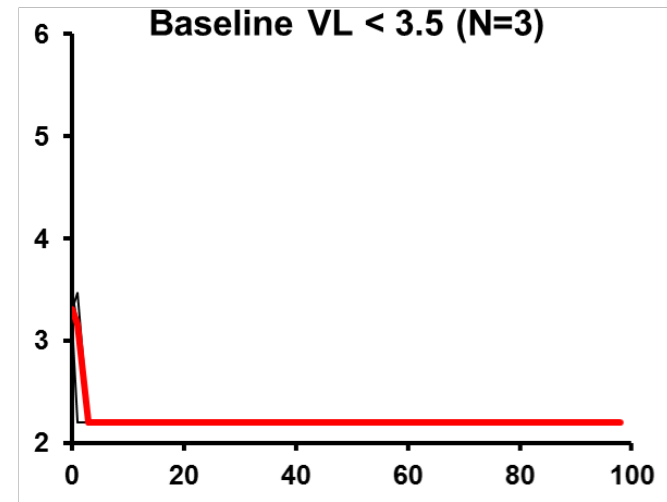
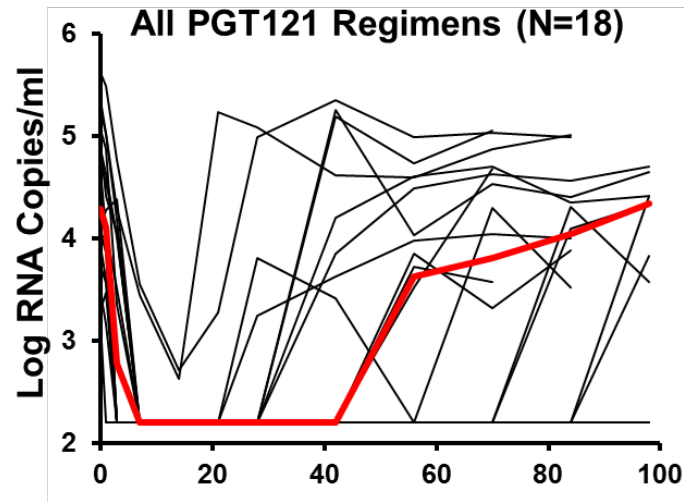
Therapeutic efficacy of PGT121 monotherapy

- **12 monkeys infected with SHIV-SF162P3 for 9 months**
- **Log setpoint viral loads 3.3 - 5.4**
- **Single IV infusion on day 0**
 - **PGT121 (10 mg/kg; N=4)**
 - **3BNC117 (10 mg/kg; N=4)**
 - **DEN3 isotype control (10 mg/kg; N=4)**

Rapid virologic control following a single infusion of PGT121 alone



Summary of therapeutic efficacy of PGT121 or PGT121-containing mAb cocktails



Perhaps the single largest problem in HIV vaccine design based on Abs is the nature of broadly neutralizing epitopes.

-these epitopes exist in a sea of glycans and variable protein residues, presenting major challenges to recognition by and elicitation of bnAbs.

How do Abs in natural HIV infection deal with glycans and variability?

October 2005
Volume 102
Number 42

PNAS

Proceedings of the National Academy of Sciences of the United States of America

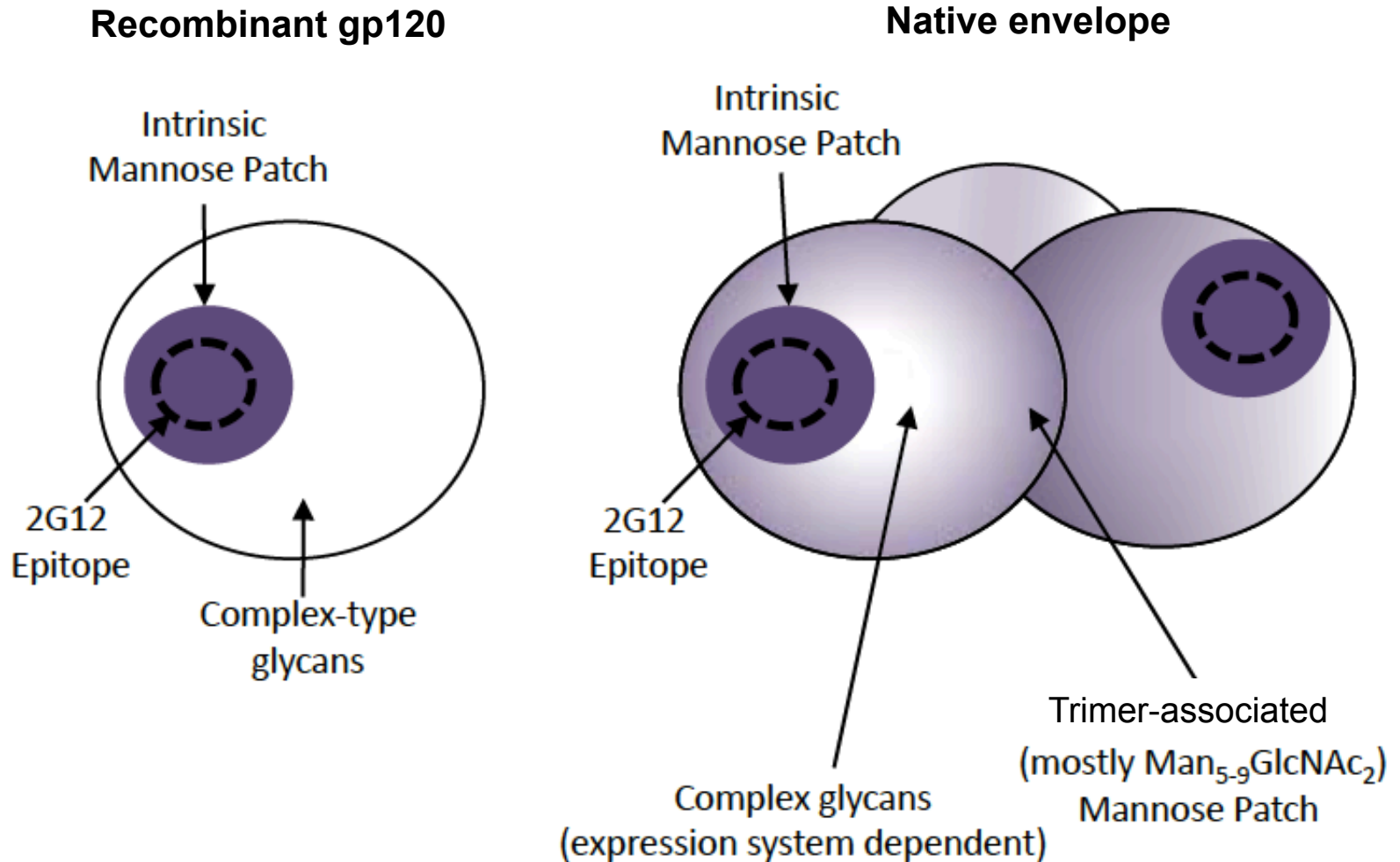
www.pnas.org

Antibody vs. HIV in a clash of evolutionary titans

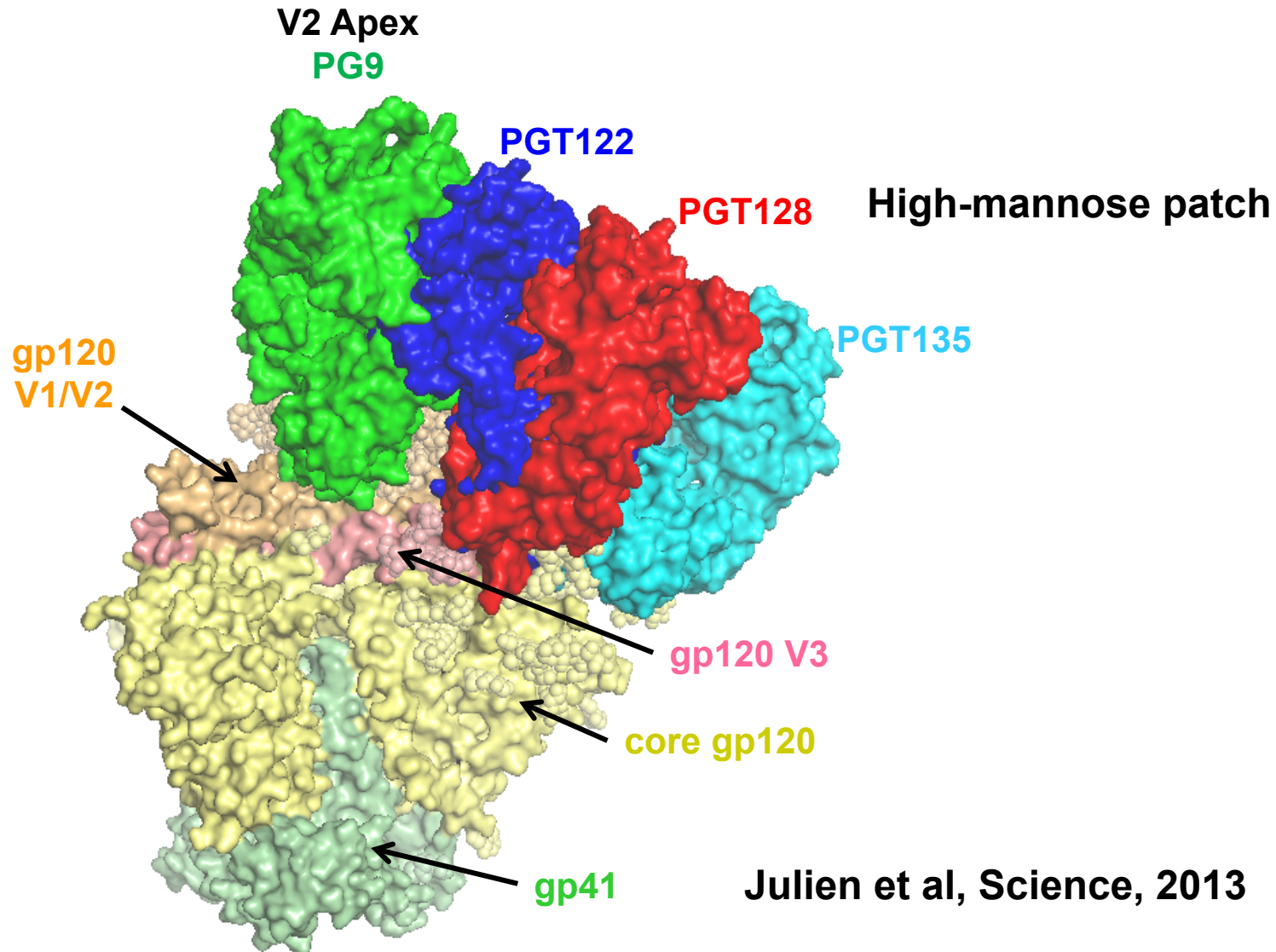
Dennis R. Burton, Robyn L. Stanfield, and
Ian A. Wilson

**Anti-glycan/protein bnMAbs
(V2/apex and high mannose patch)**

Model for native envelope glycosylation

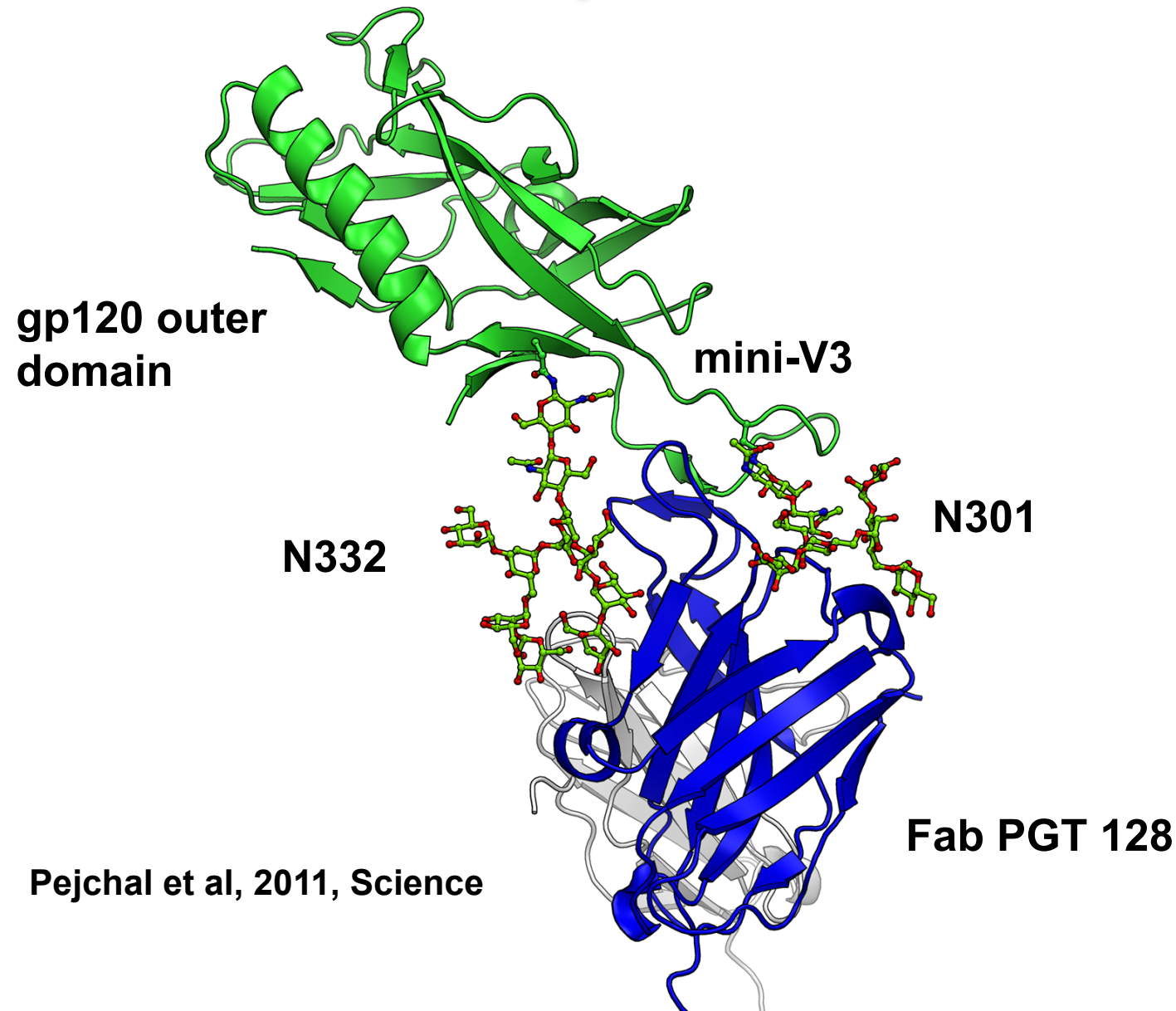


Glycan-dependent bnMAbs bound to SOSIP Env gp140 trimer



Julien et al, Science, 2013

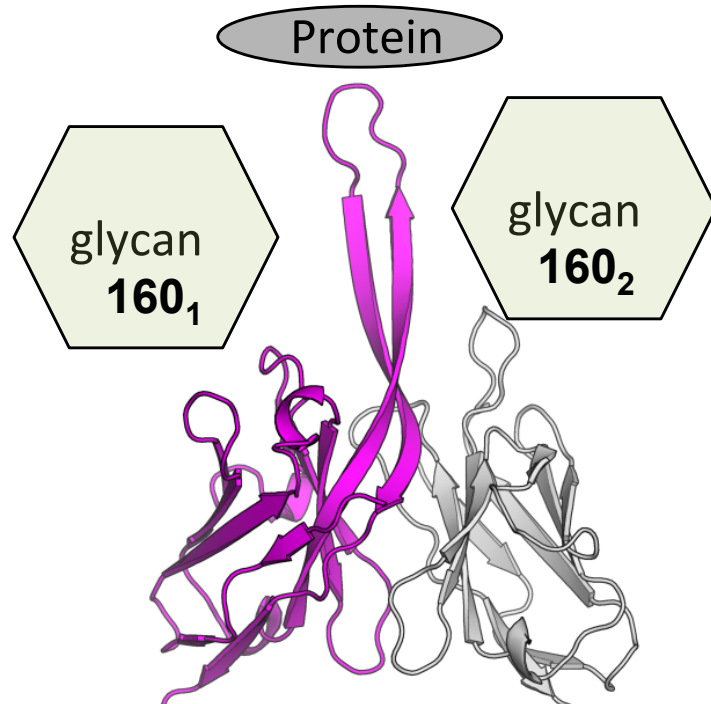
PGT 128 penetrates the glycan shield and accesses the protein surface underneath



Pejchal et al, 2011, Science

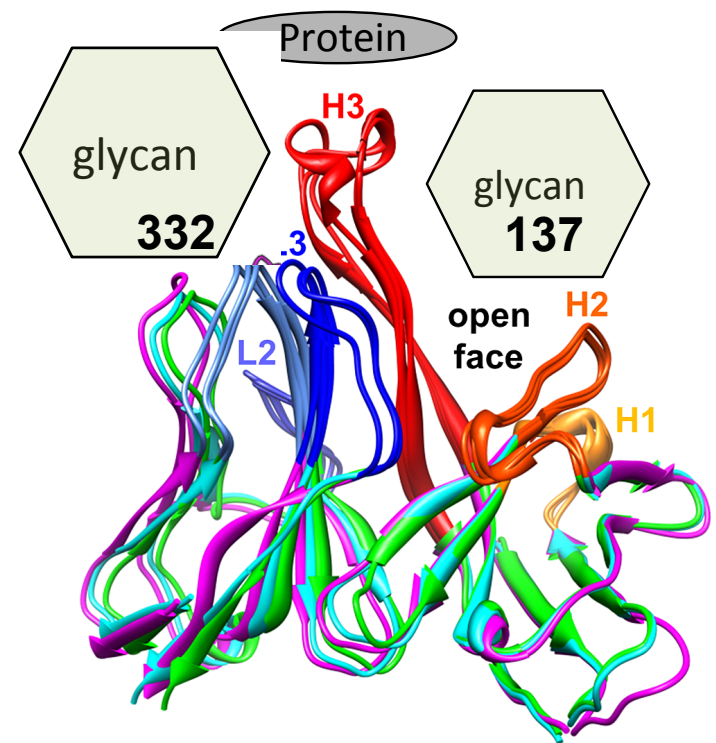
V2 apex and high-mannose patch bnAb recognition

V2 apex



PGT145

High mannose patch

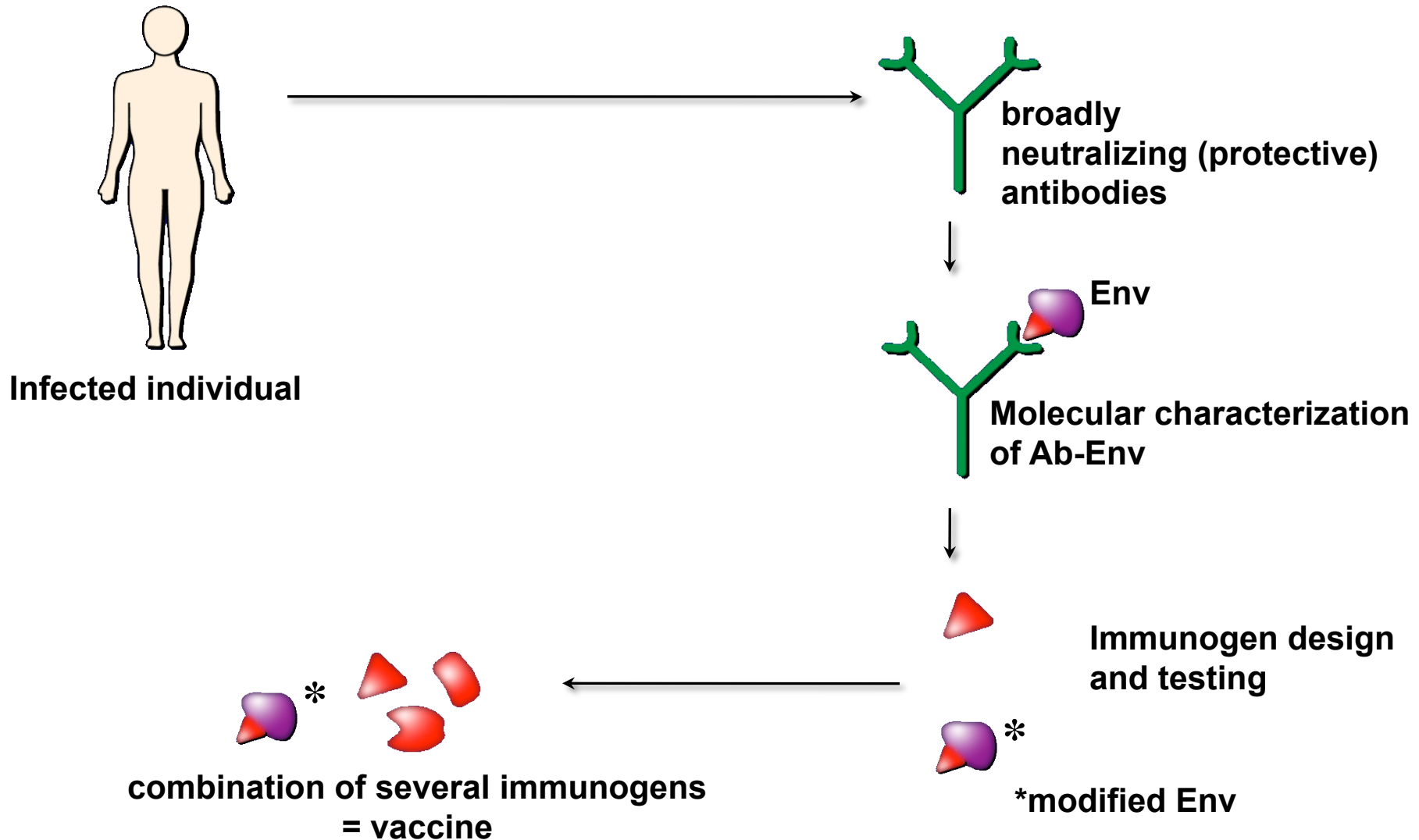


PGT121

PGT124

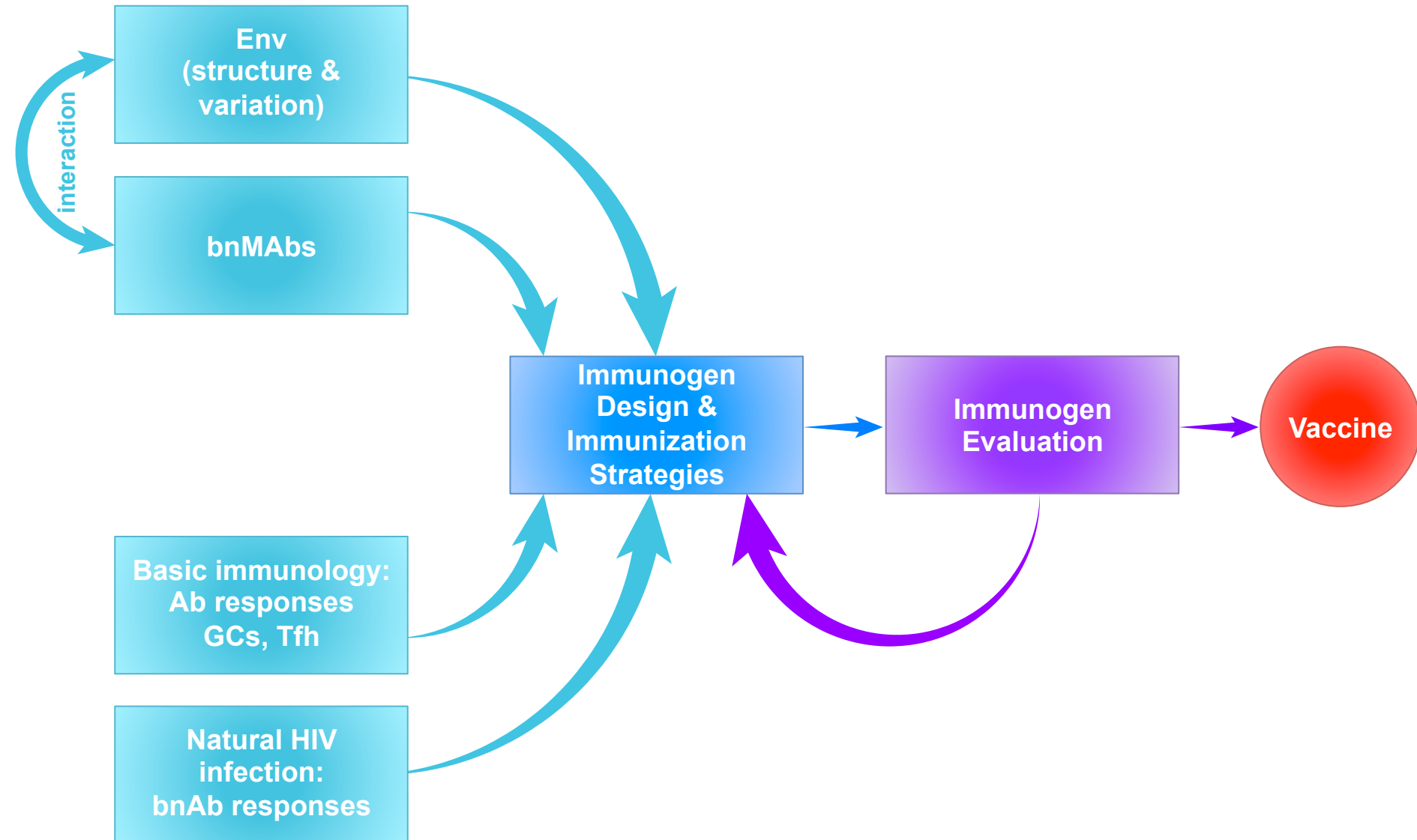
How do we move from data on bnAb-Env interactions to immunogen design?

Reverse engineering a viral vaccine



(adapted from Burton, Nat. Rev. Immunol., 2:706, 2002)

Rational Approach to HIV Vaccine Design



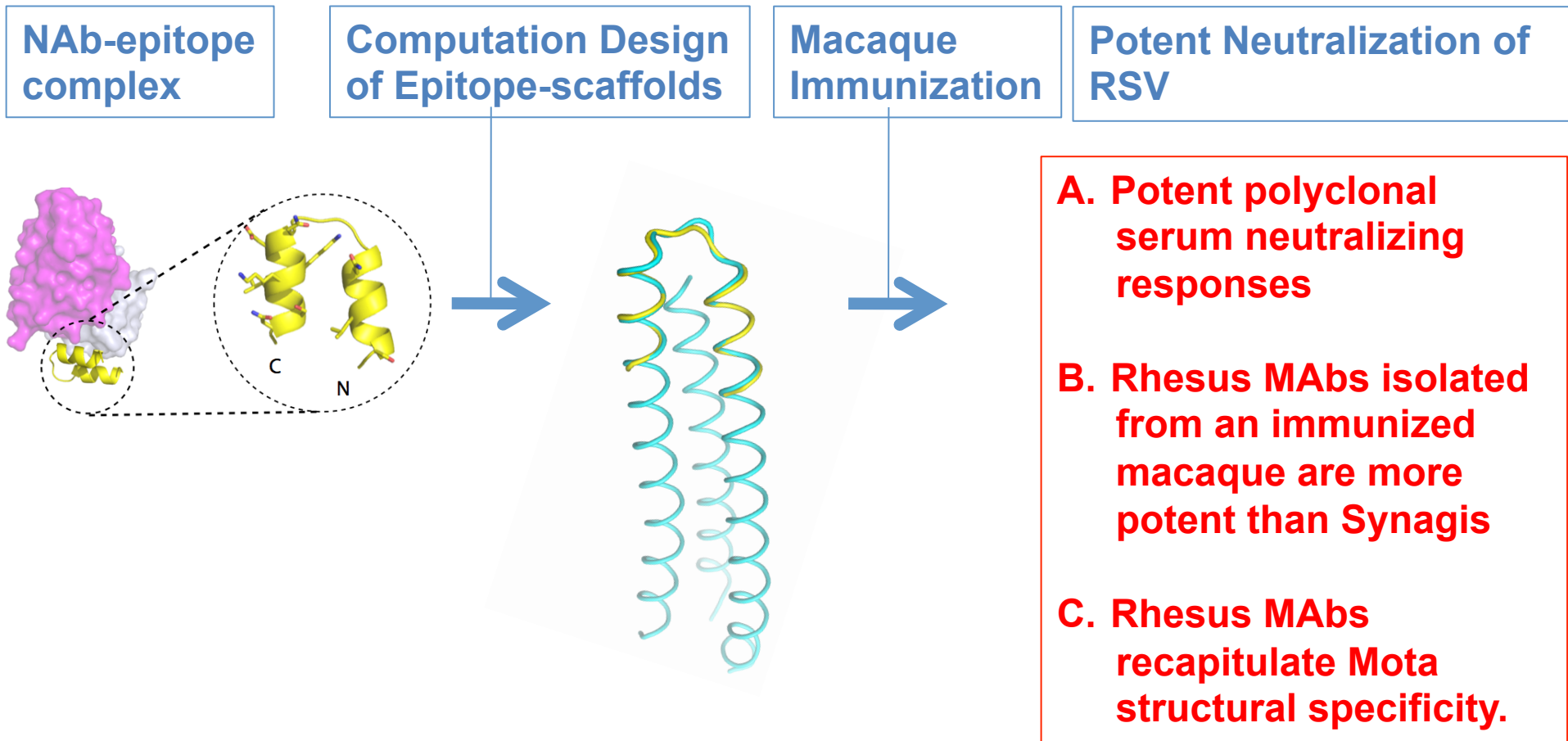
Structure-based design of a fusion glycoprotein vaccine for respiratory syncytial virus

Jason S. McLellan, Man Chen, M. Gordon Joyce, Mallika Sastry, Guillaume B. E. Stewart-Jones, Yongping Yang, Baoshan Zhang, Lei Chen, Sanjay Srivatsan, Anqi Zheng, Tongqing Zhou, Kevin W. Graepel, Azad Kumar, Syed Moin, Jeffrey C. Boyington, Gwo-Yu Chuang, Cinque Soto, Ulrich Baxa, Arjen Q. Bakker, Hergen Spits, Tim Beaumont, Zizheng Zheng, Ningshao Xia, Sung-Youl Ko, John-Paul Todd, Srinivas Rao, Barney S. Graham, and Peter D. Kwong

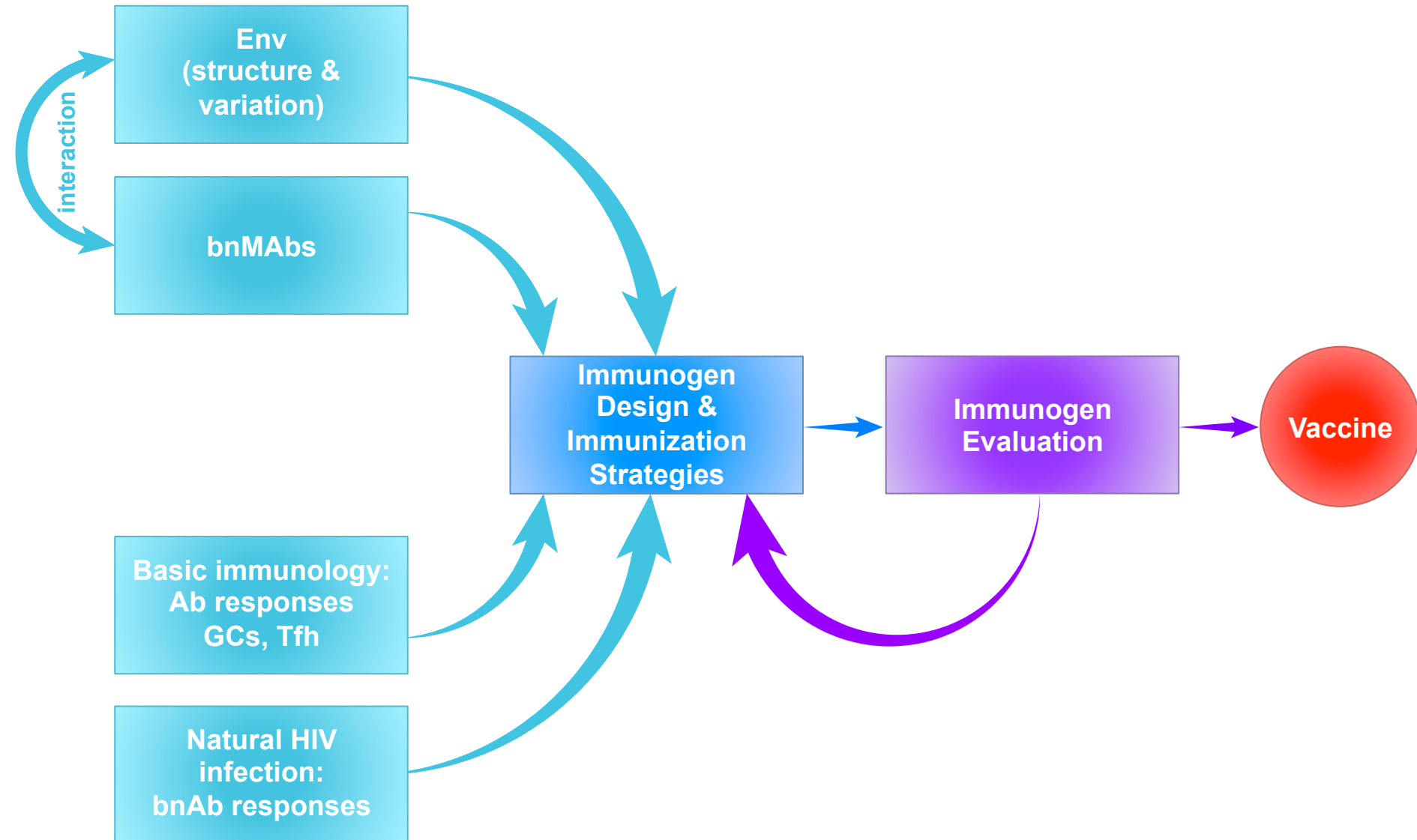
Proof of principle for epitope-focused vaccine design

Bruno E. Correia, John T. Bates, Rebecca J. Loomis, Gretchen Baneyx, Chris Carrico, Joseph G. Jardine, Peter Rupert, Colin Correnti, Oleksandr Kalyuzhniy, Vinayak Vittal, Mary J. Connell, Eric Stevens, Alexandria Schroeter, Man Chen, Skye MacPherson, Andreia M. Serra, Yumiko Adachi, Margaret A. Holmes, Yuxing Li, Rachel E. Klevit, Barney S. Graham, Richard T. Wyatt, David Baker, Roland K. Strong, James E. Crowe Jr, Philip R. Johnson and William R. Schief

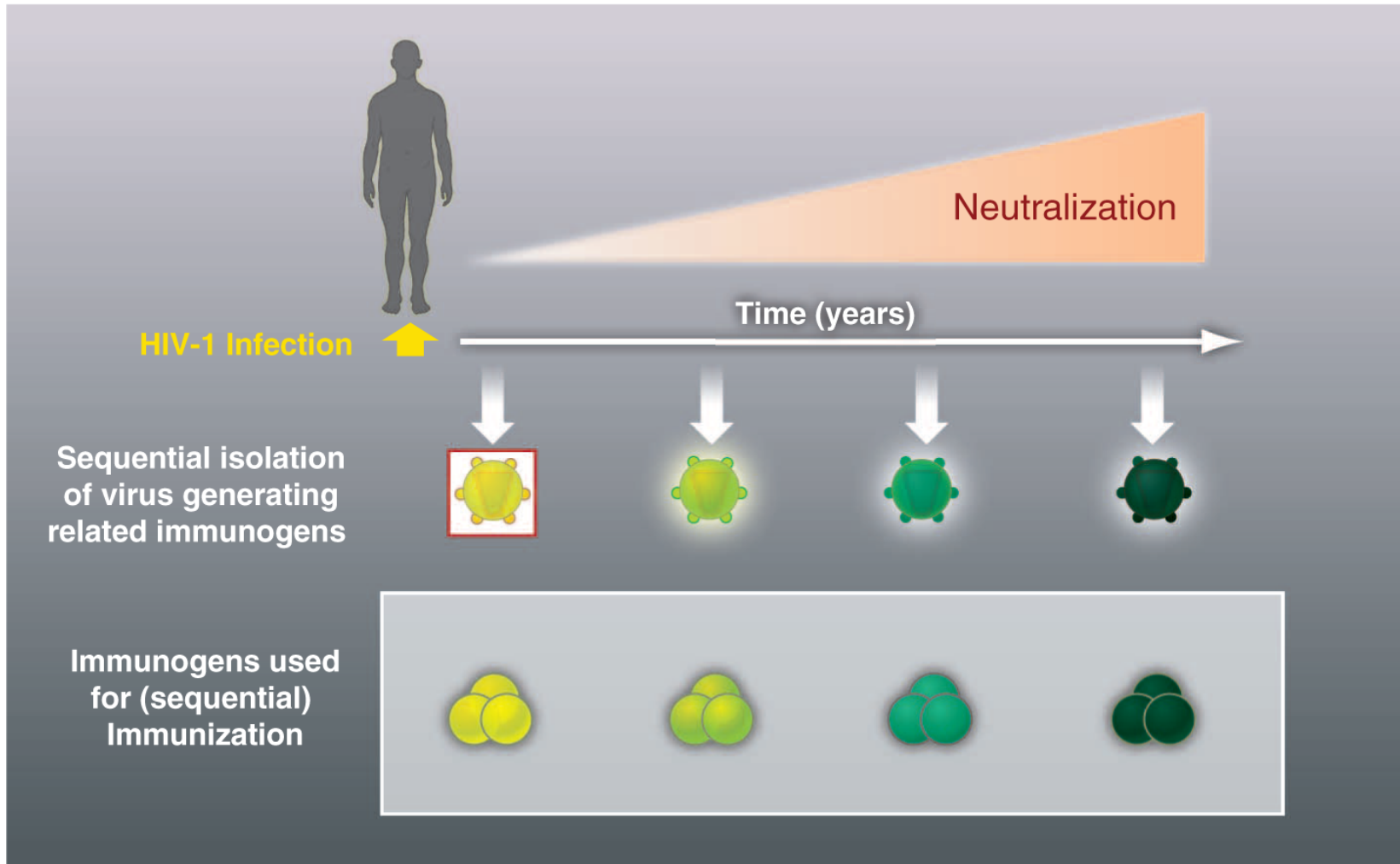
Proof of concept for epitope-focused vaccine design: epitope-scaffolds induce potent neutralization of RSV in NHPs



Rational Approach to HIV Vaccine Design

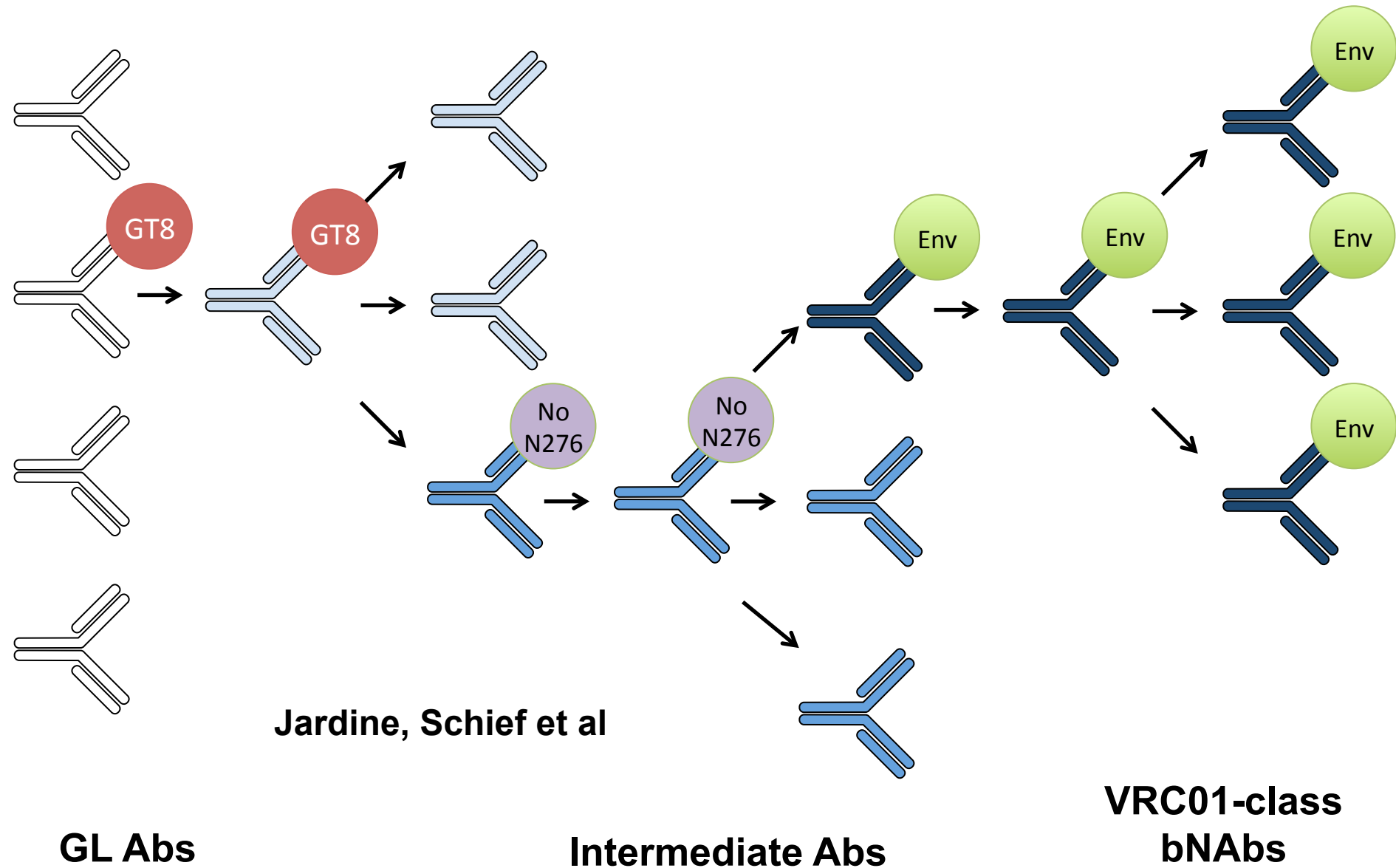


Vaccine Design based on studying Co-Evolution of Antibody and Virus



Klein et al, Science, 2013

The Reductionist Approach: VRC01-like CD4bs Abs-one iteration



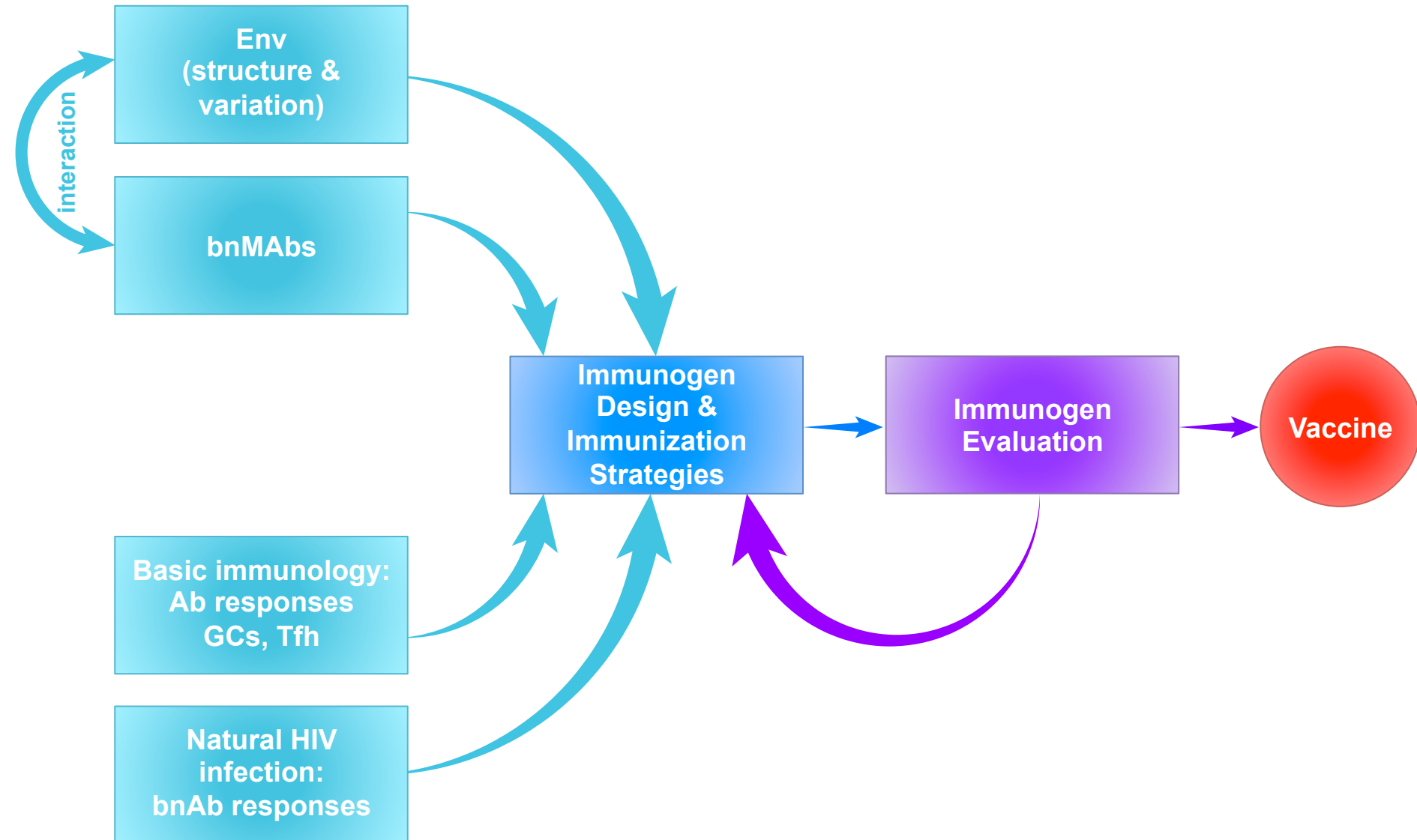


Rational HIV immunogen design to target specific germline B cell receptors.

Jardine J, Julien JP, Menis S, Ota T, Kalyuzhniy O, McGuire A, Sok D, Huang PS, MacPherson S, Jones M, Nieuwma T, Mathison J, Baker D, Ward AB, Burton DR, Stamatatos L, Nemazee D, Wilson IA, Schief WR.

Science. 2013 340:711-6

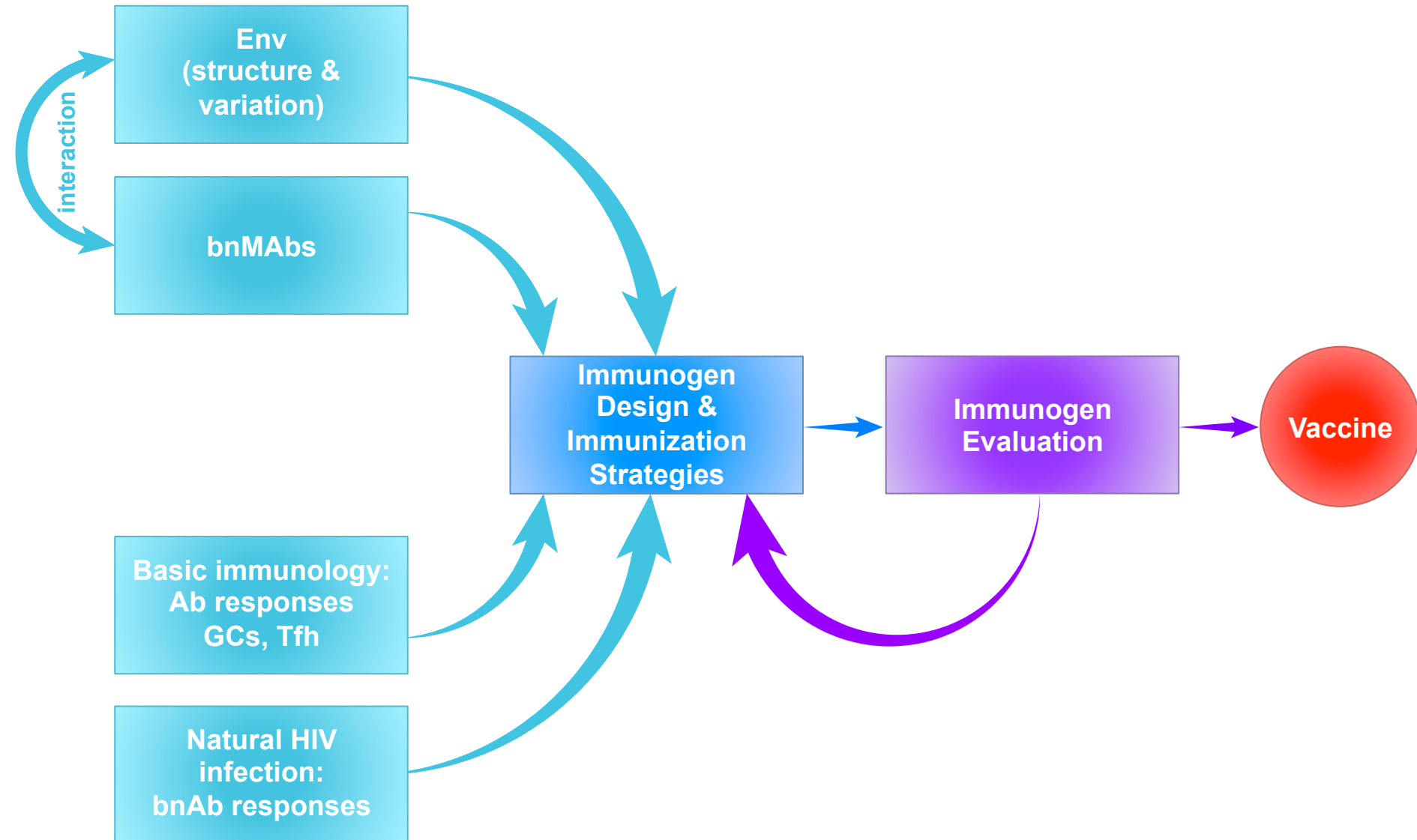
Rational Approach to HIV Vaccine Design



HIV immunogen evaluation

- **Iteration!**
- **Prefer humans! Animal models-macaques, mice expressing human Ab repertoires, germline (GL)-bnAb knockin mice. In vitro models.**
- **Detailed analysis of Ab responses-Next Generation Sequencing, isolation of MAbs, innate signatures, Tfh cells.**

Rational Approach to HIV Vaccine Design



Collaborators

CHAVI-ID

Rafi Ahmed
Dan Barouch
Dennis Burton
Max Crispin
Shane Crotty
Adam Godzik
Julie McElrath
Michel Nussenzweig
Bali Pulendran
Bill Schief
Guido Silvestri
Bruce Walker
Andrew Ward
Ian Wilson
Rich Wyatt
Jiang Zhu

Scripps

David Nemazee
Chi-Huey Wong
Jim Paulson

IAVI/Scripps

Pascal Poignard

VRC

John Mascola
Peter Kwong

Duke CHAVI-ID

Bart Haynes
Bette Korber
Robin Shattock

IAVI

Wayne Koff
Clinical sites
Members of the
Neutralizing
Antibody
Consortium
Protocol G & C
scientists &
donors

Monogram

Theraclone

Ragon

Bruce Walker
Galit Alter
Dan Barouch
Arup Chakra-
-borty

Cornell

John Moore

Miami

David Watkins



National Institute of
Allergy and
Infectious Diseases



*Scripps Center for HIV/AIDS Vaccine
Immunology & Immunogen Discovery*



International AIDS
Vaccine Initiative



The Collaboration for AIDS Vaccine Discovery

BILL & MELINDA
GATES foundation



Global HIV Vaccine
Enterprise



Ragon Institute
of MGH, MIT and Harvard