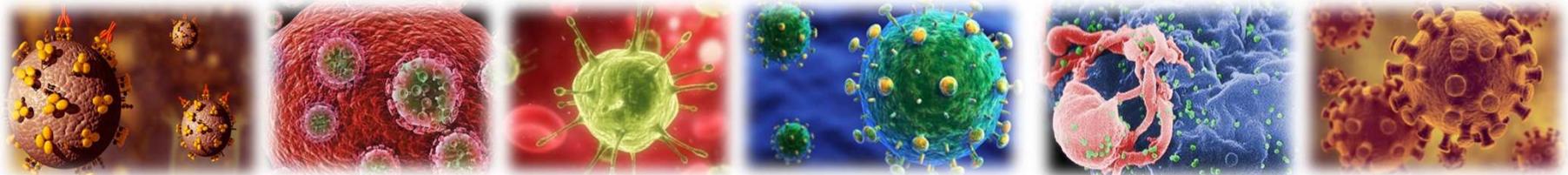


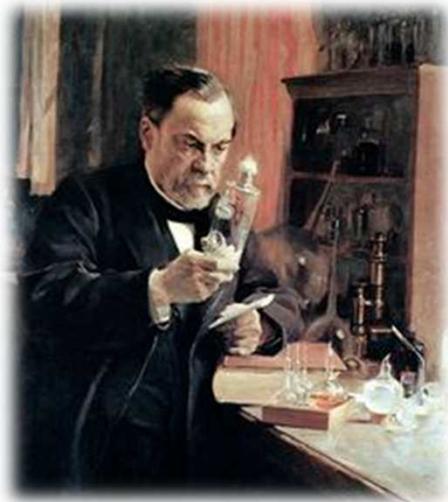
From Traditional to New-Generation Vaccines: Can We Overcome the Impossible?

Anan Jongkaewwattana, Ph.D

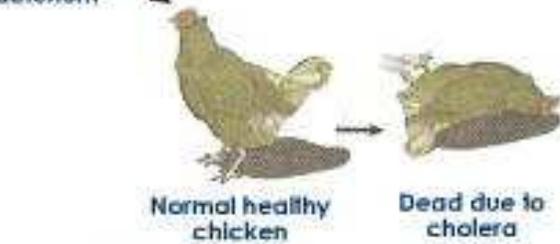
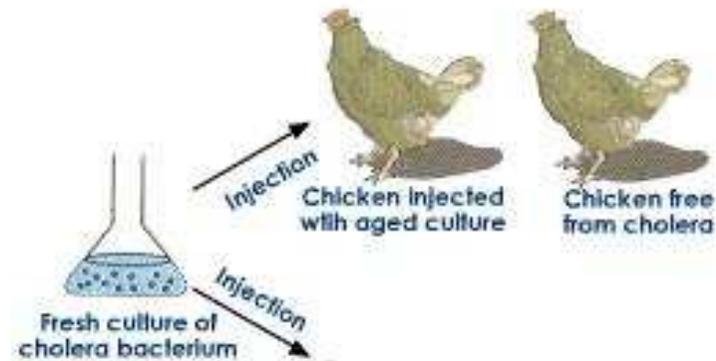
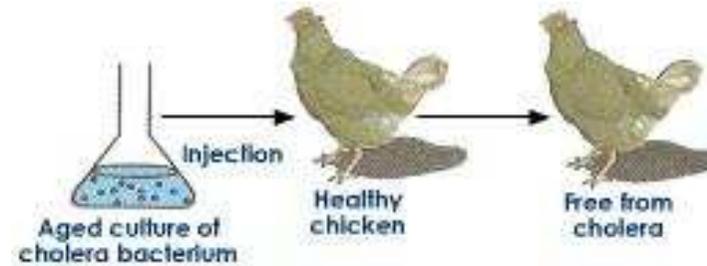
**Virology and Antibody Technology Research Unit
BIOTEC, NSTDA**



Isolate – Inactivate - Inject

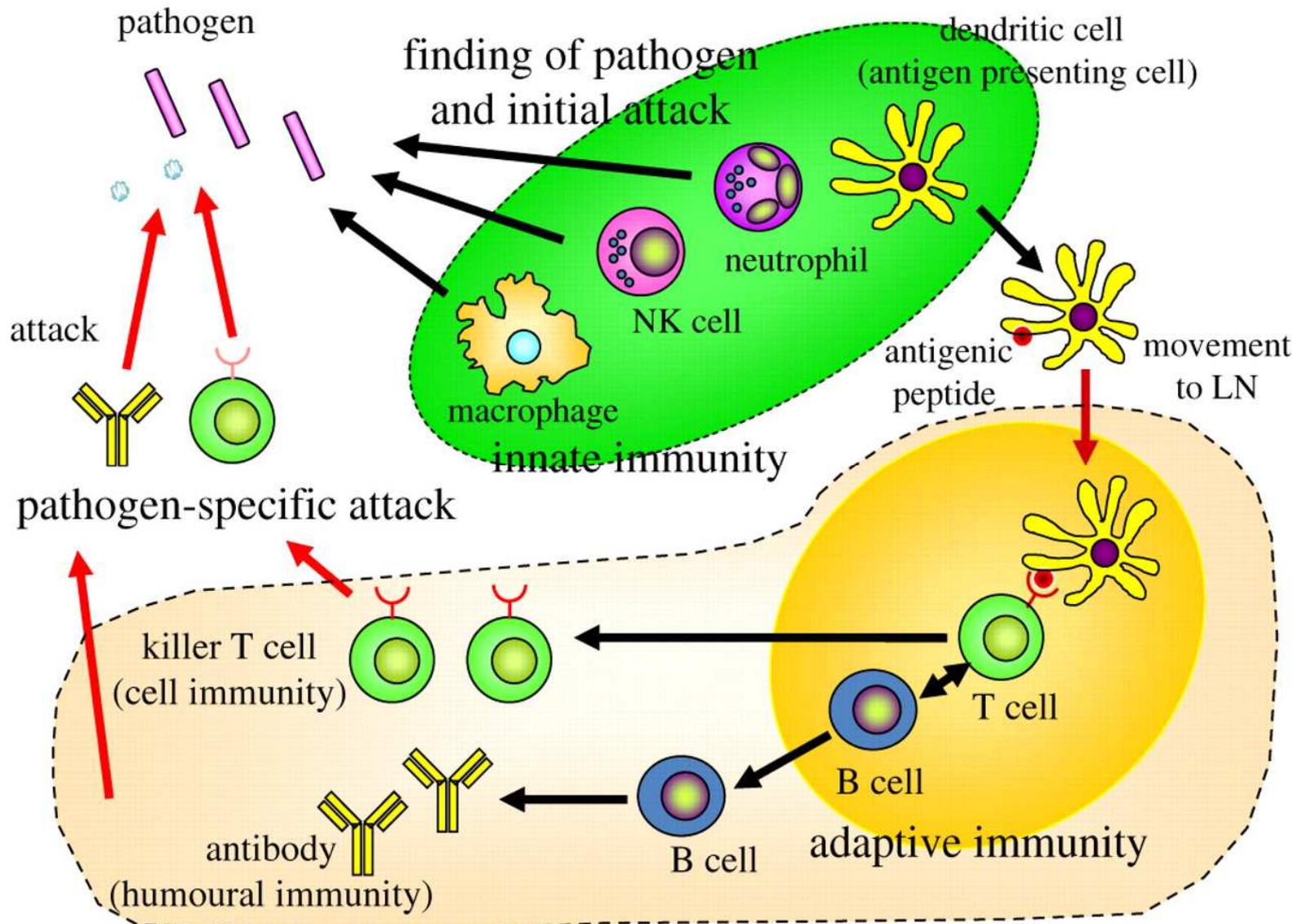


Louis Pasteur (1811-1895)

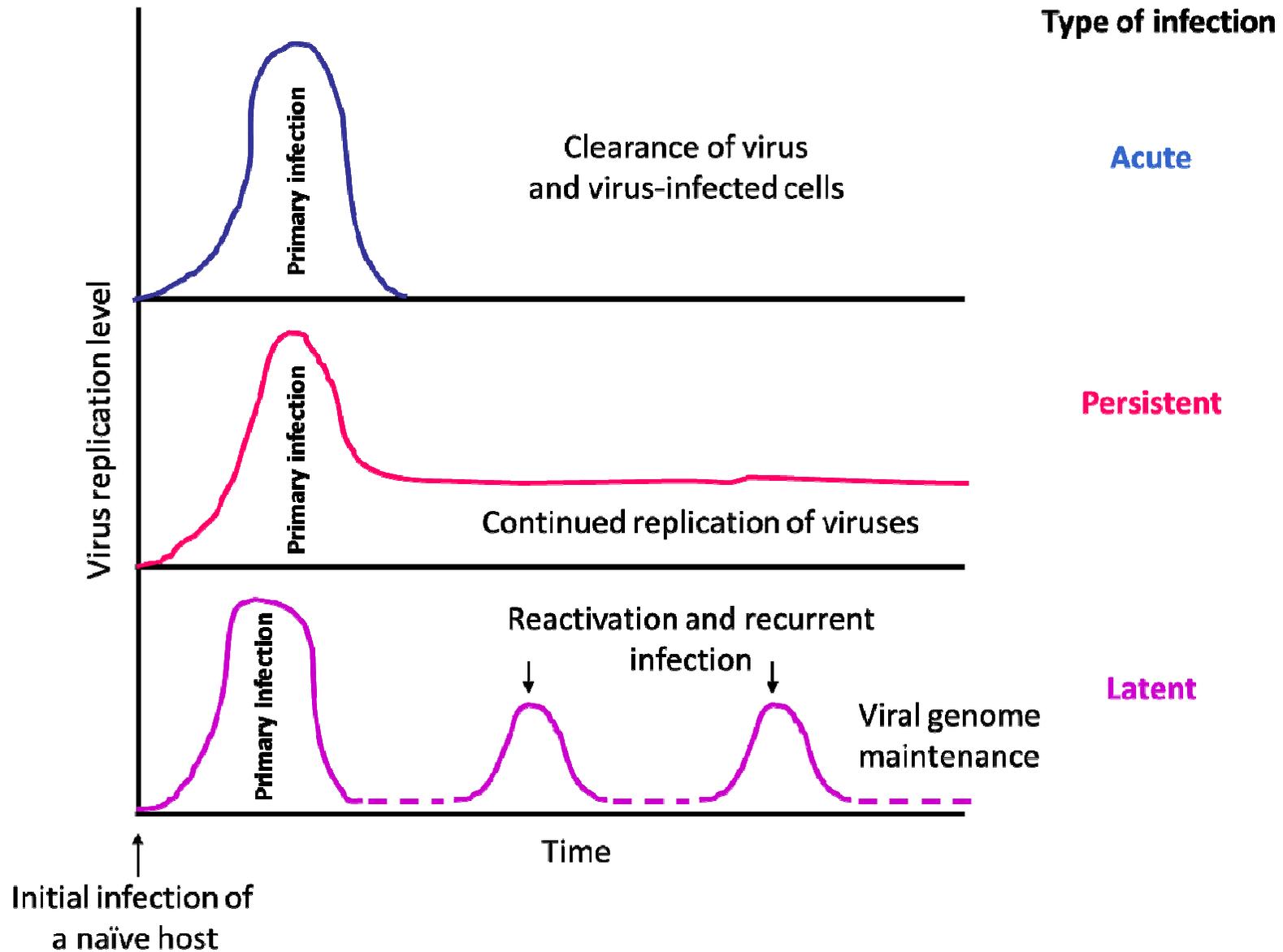


The classic experiment of Pasteur with chicken (fowl) cholera

Innate and Adaptive immunity: The Concept



Outcomes of virus infection can vary



Class I Pathogen

Infects narrow age range

Host exhibits spontaneous recovery

Host generates long lasting protective immunity

Priming with wild-type or attenuated pathogen induces protection

Genetically stable with limited antigenic variation

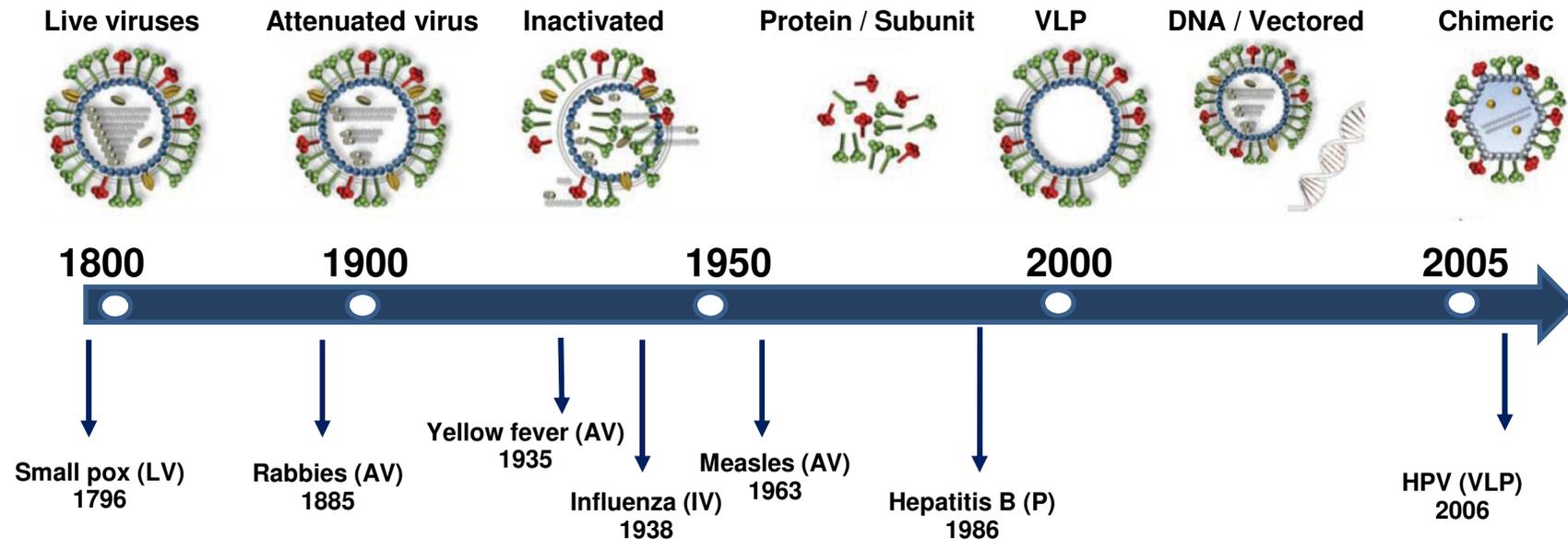
Immune responses are directed to multiple epitopes

Ex: Measles, mumps, rubella, diphtheria, Canine distemper, rabies, poliovirus

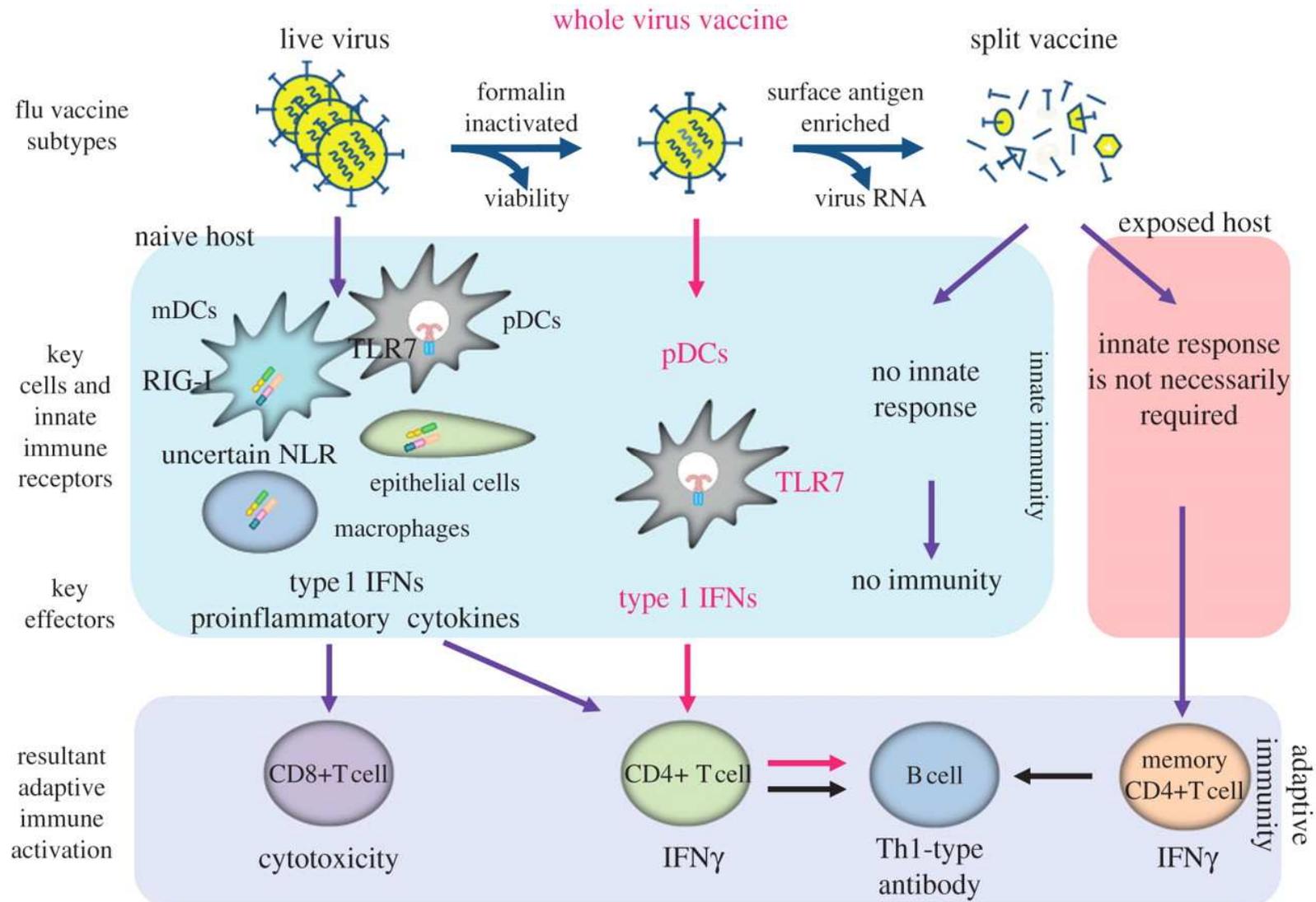
Technological milestones

Experimental Sciences 18-19 th Century Inoculation	Tissue Culture – 1930s Eggs, Animal cell culture	Recombinant DNA – 1980s Sequencing Cloning PCR Gene Delivery	Synthetic Biology – 2000s DNA and Peptide Synthesis
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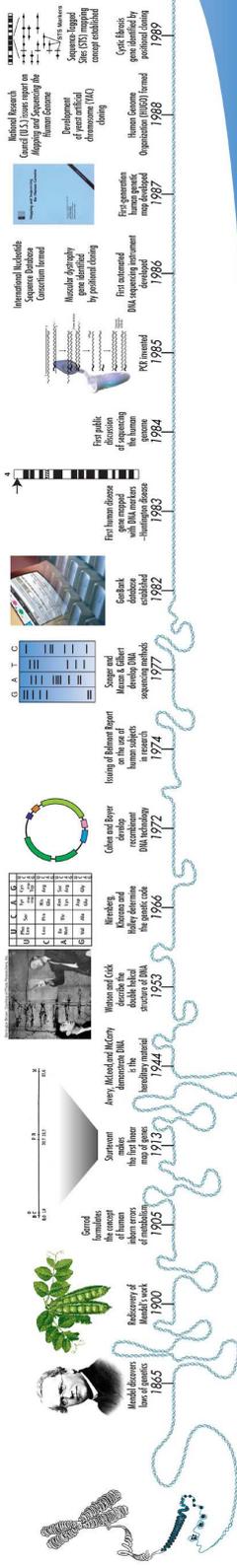
Vaccine types



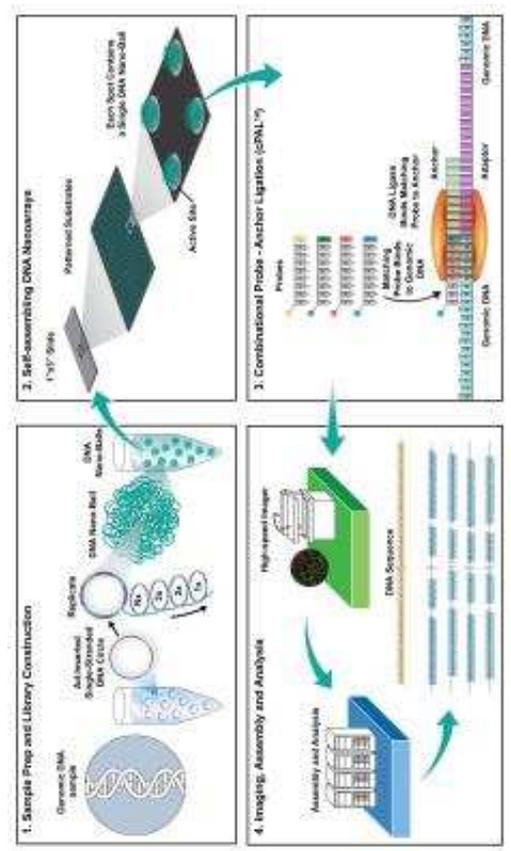
Each vaccine type has its pros and cons



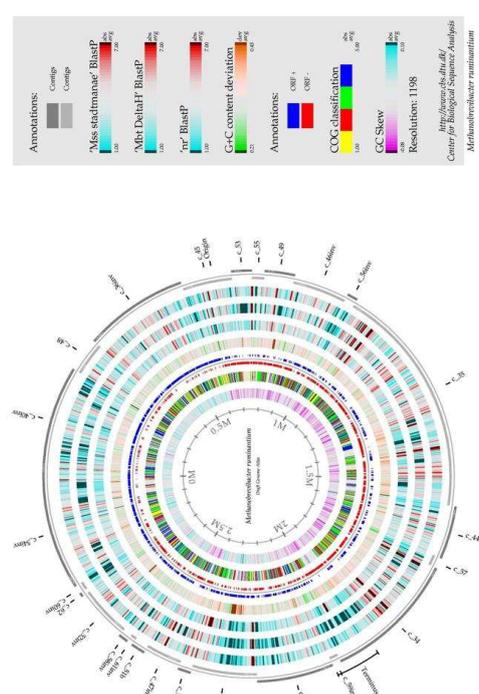
HGP



1990 Human Genome Project (HGP) launched in the U.S. Ethical, legal and social implications (ELSI) programs founded at NIH and DOE.	1991 First U.S. Genome Center established.	1992 Second-generation human genetic map developed.	1993 New five-year plan for the HGP in the U.S. published.	1994 HGP's human genetic mapping goal achieved.	1995 HGP's human physical mapping goal achieved.	1996 First human gene map established.	1997 DOE forms Joint Genome Institute (JGI). MORF becomes NHGRI.	1998 Incorporation of 30,000 genes into human genome map. New five-year plan for the HGP in the U.S. published.	1999 Full-scale human sequencing begins.	2000 Draft version of human genome sequence completed. President Clinton and Prime Minister Blair support free access to genome information.	2001 Draft version of human genome sequence published.	2002 Draft version of mouse genome sequence completed and published.	2003 Finished version of human genome sequence completed.
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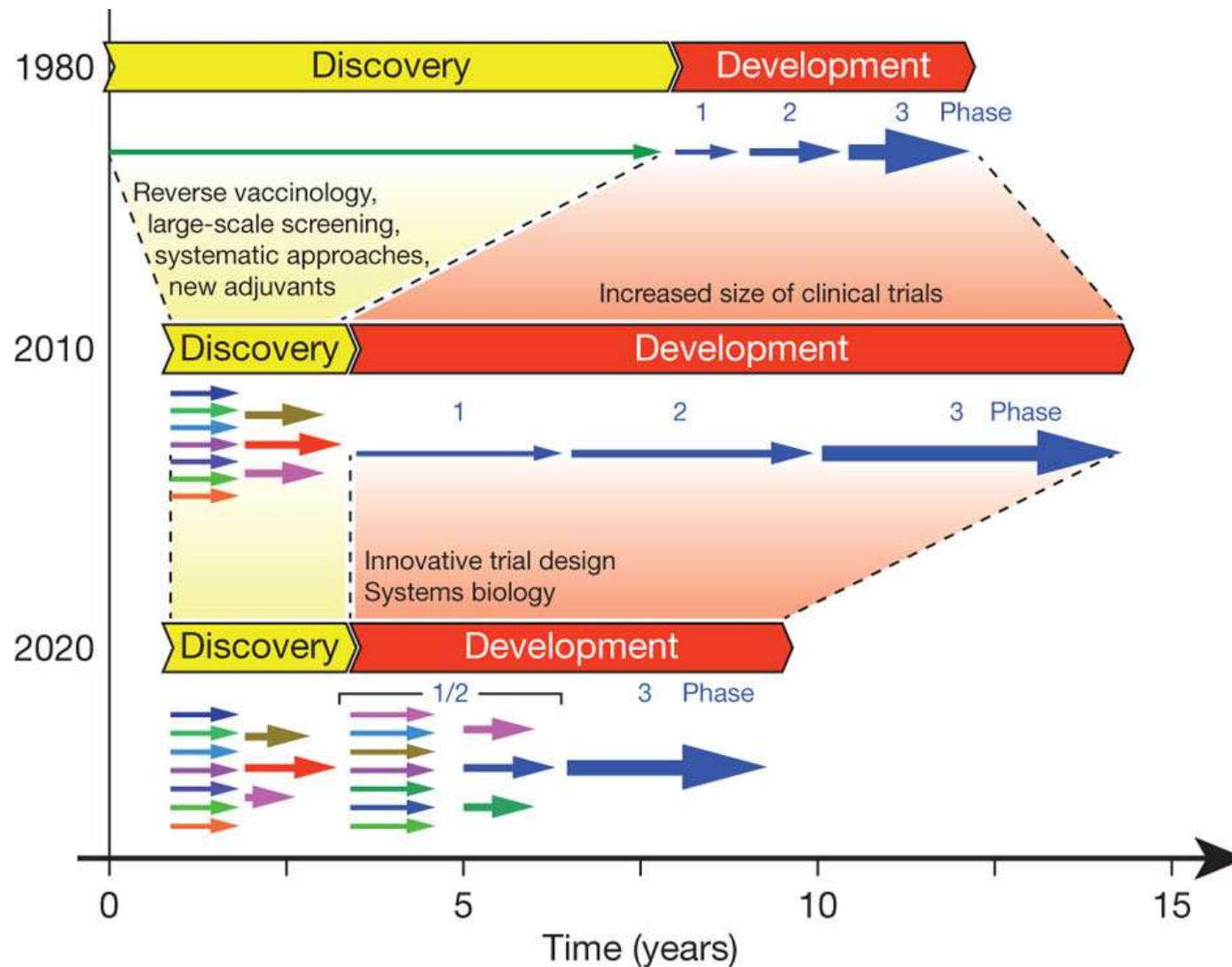


Complete Genomics' Sequencing Process

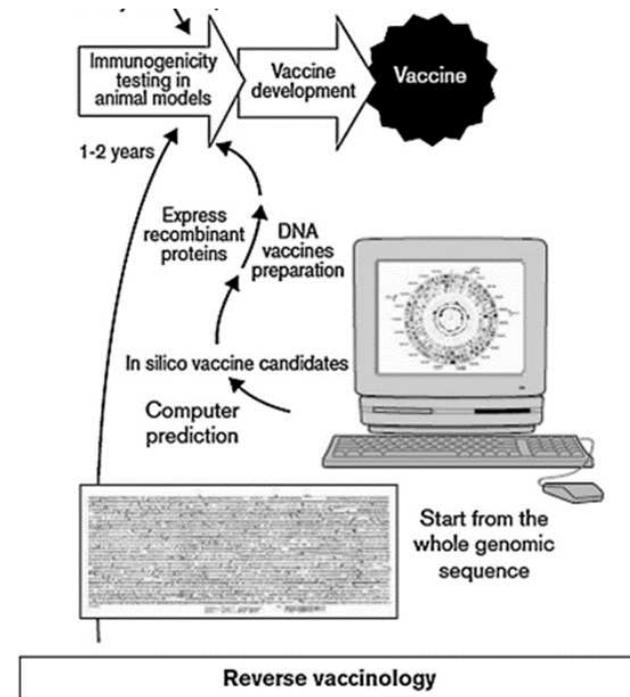
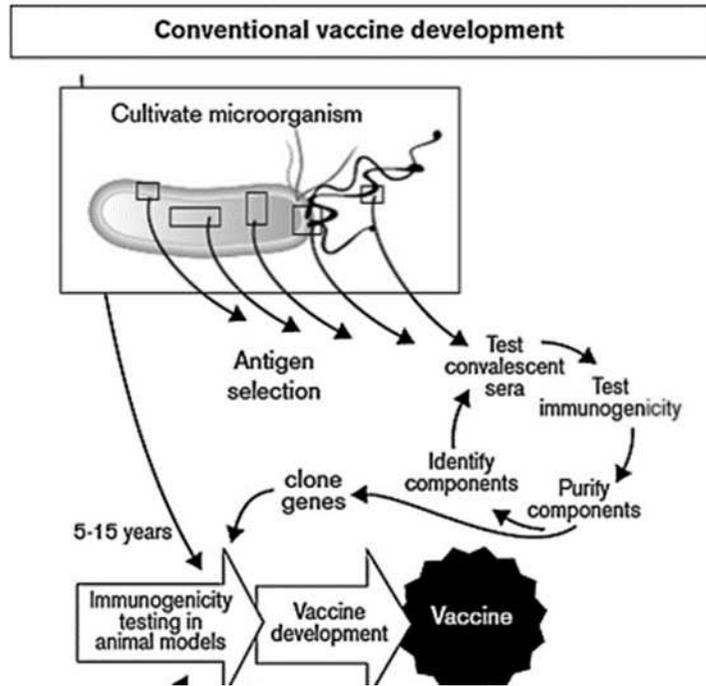
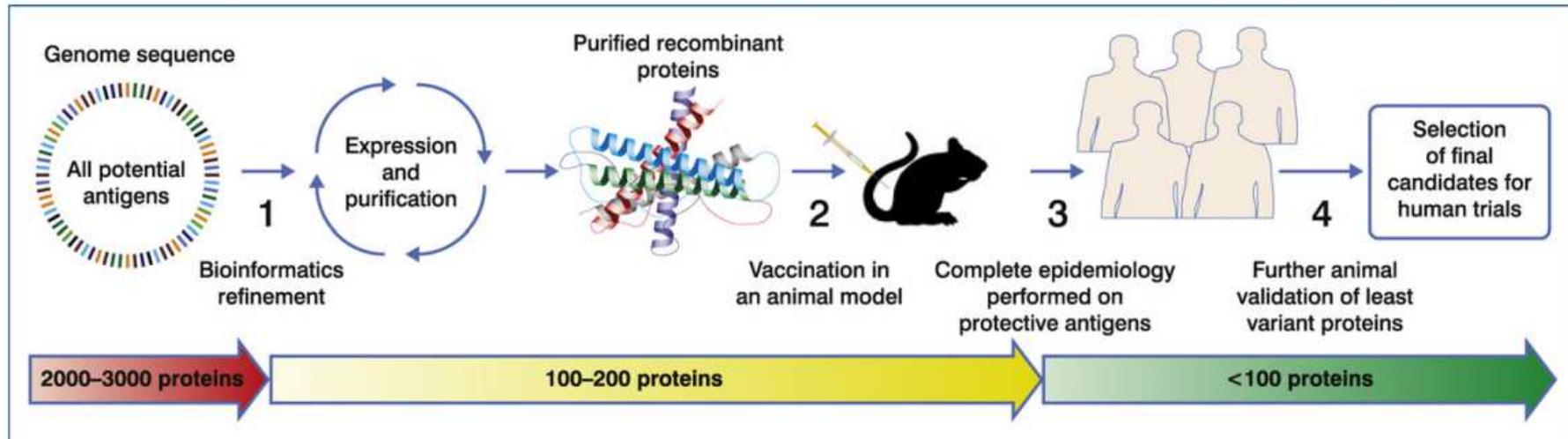


Center for Biological Sequence Analysis
Mikhael Reiterman

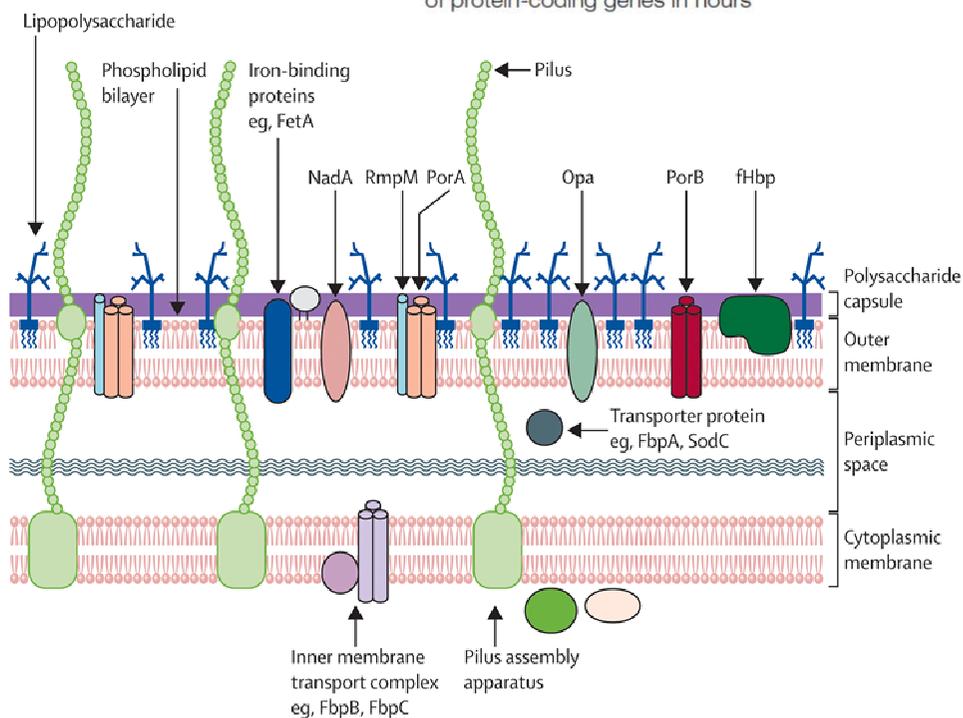
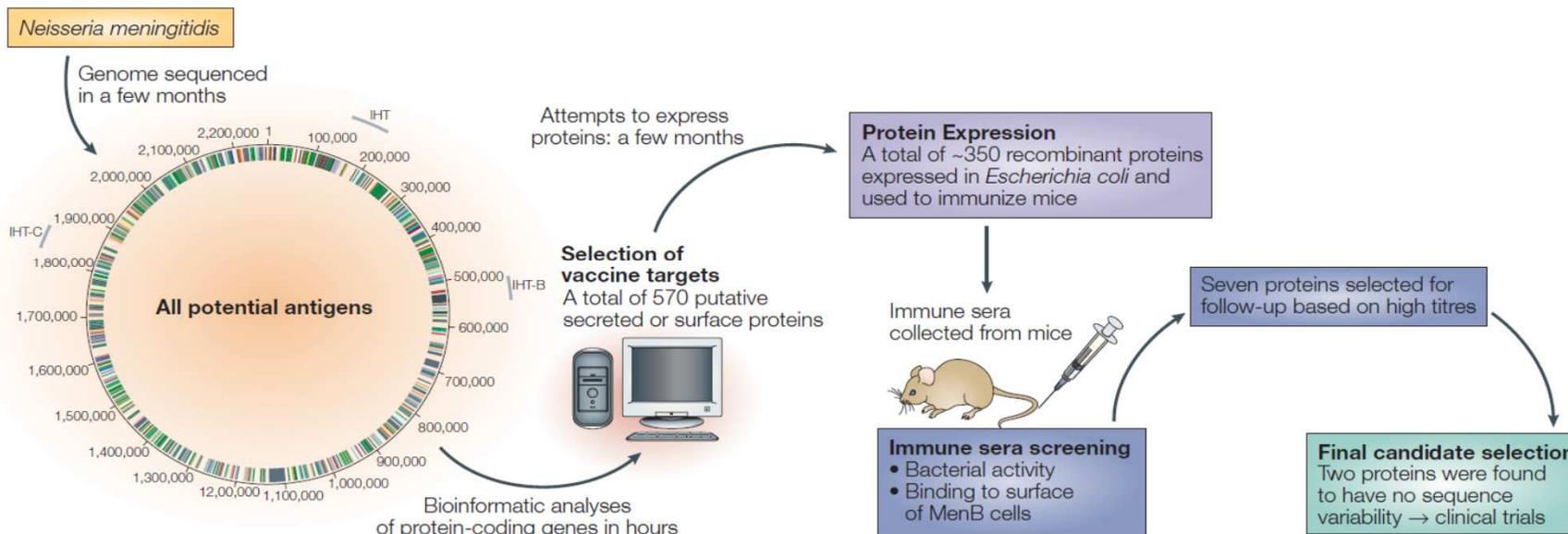
Evolution of vaccine development during the last 30 years



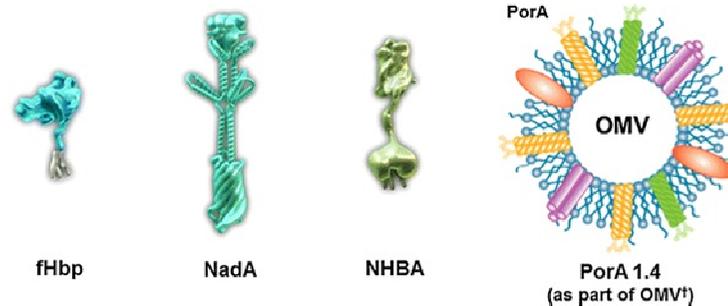
Reverse Vaccinology: The Concept



Reverse Vaccinology: The Application

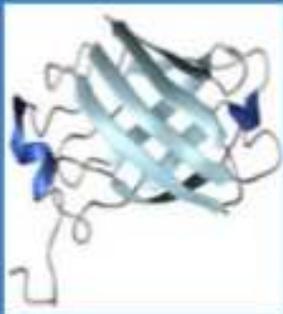
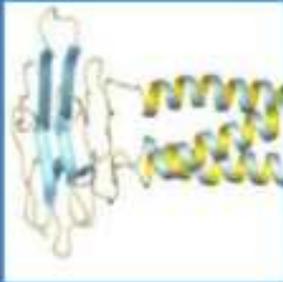
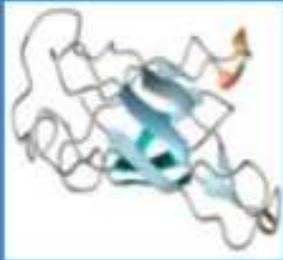


4CMenB:



Novartis investigational MenB 4CMenB vaccine (Bexsero[®])

- Bexsero (previously known as 4CMenB or rMenB+OMV) contains 4 main antigens.
- Three recombinant proteins discovered by genome mining/reverse vaccinology combined with OMV from the New Zealand outbreak strain (NZ 98/254).

			
rHBP-1	NadA	NHBA	PorA (presented as part of an OMV)

Published March 28, 2016

JEM

Perspective

Reverse vaccinology 2.0: Human immunology instructs vaccine antigen design

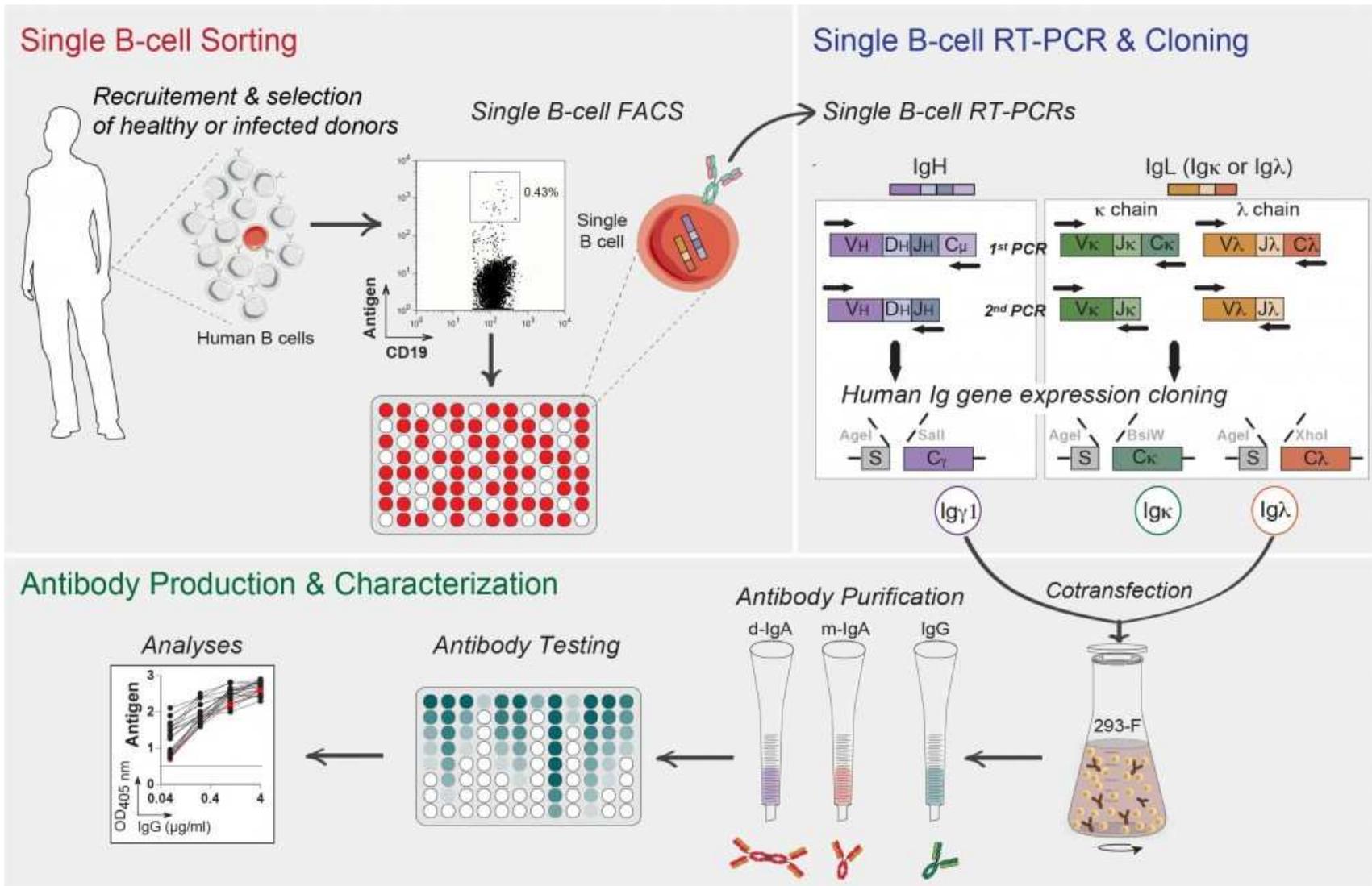
Rino Rappuoli, Matthew J. Bottomley, Ugo D'Oro, Oretta Finco, and Ennio De Gregorio

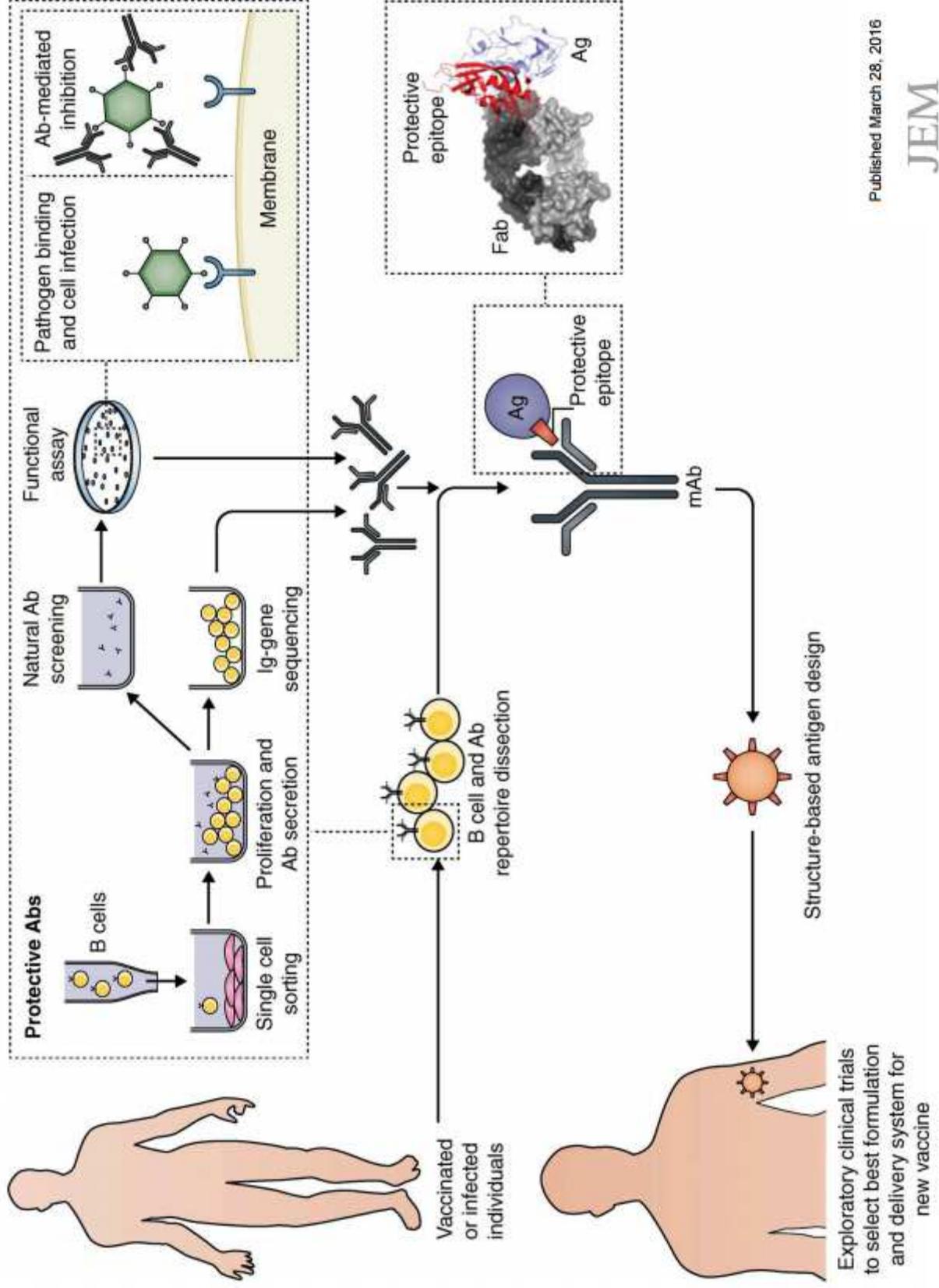
GlaxoSmithKline Vaccines S.r.l., 53100 Siena, Italy

Traditionally, vaccines have been developed by cultivating infectious agents and isolating the inactivated whole pathogen or some of its purified components. 20 years ago, reverse vaccinology enabled vaccine discovery and design based on information deriving from the sequence of microbial genomes rather than via the growth of pathogens. Today, the high throughput discovery of protective human antibodies, sequencing of the B cell repertoire, and the increasing structural characterization of protective antigens and epitopes provide the molecular and mechanistic understanding to drive the discovery of novel vaccines that were previously impossible. We are entering a "reverse vaccinology 2.0" era.



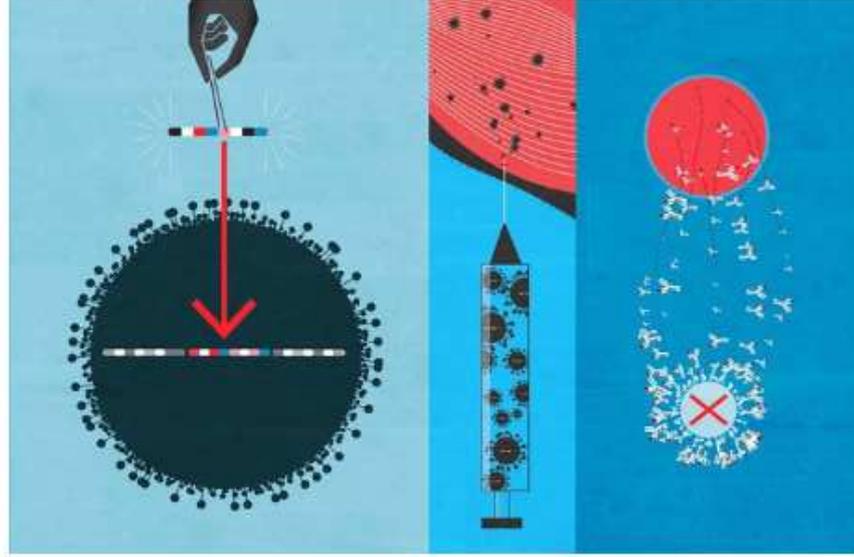
Accelerating Antibody Discovery by B cell cloning





Protection Without a Vaccine

By CARL ZIMMER MARCH 9, 2015

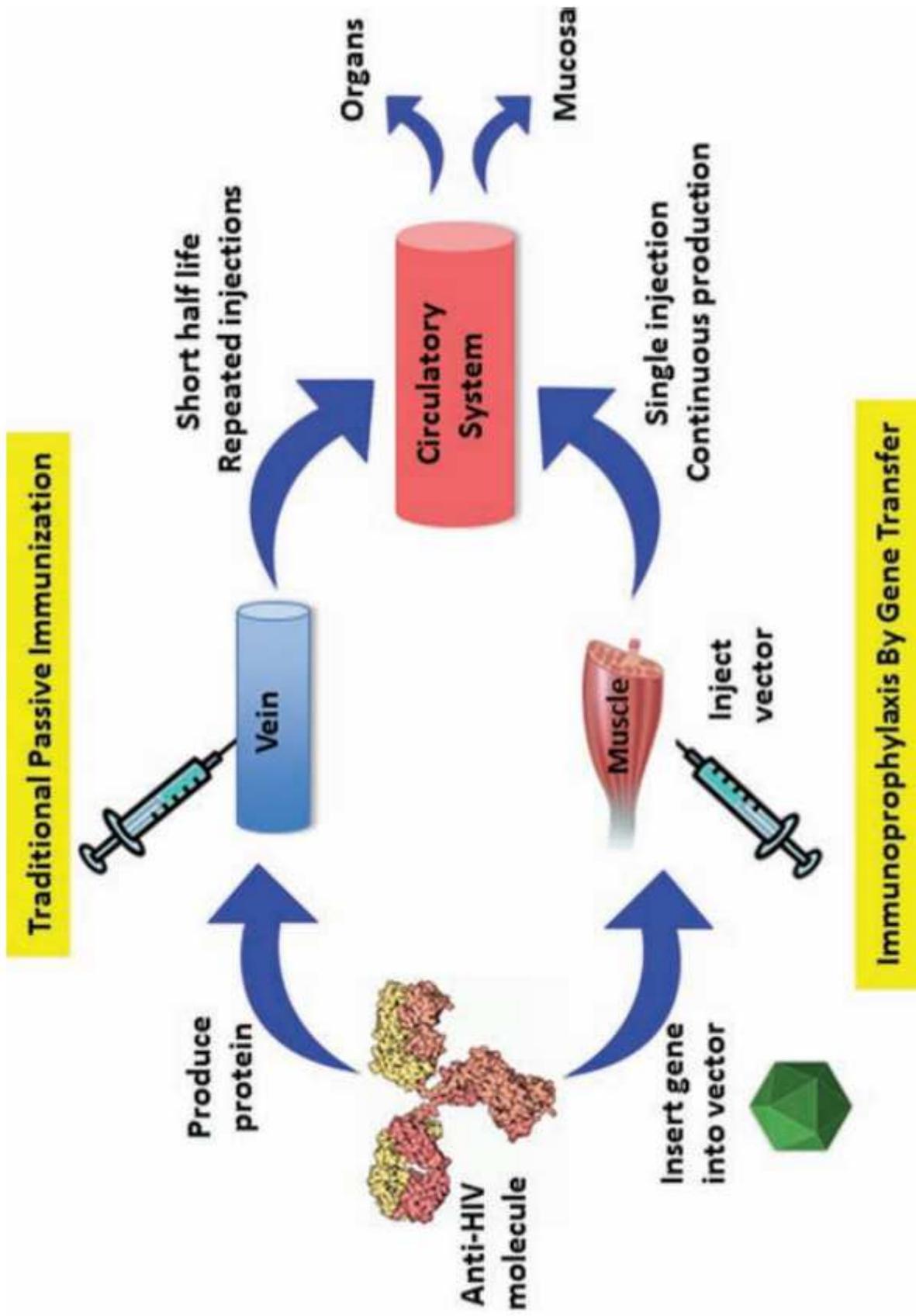


John Hersey

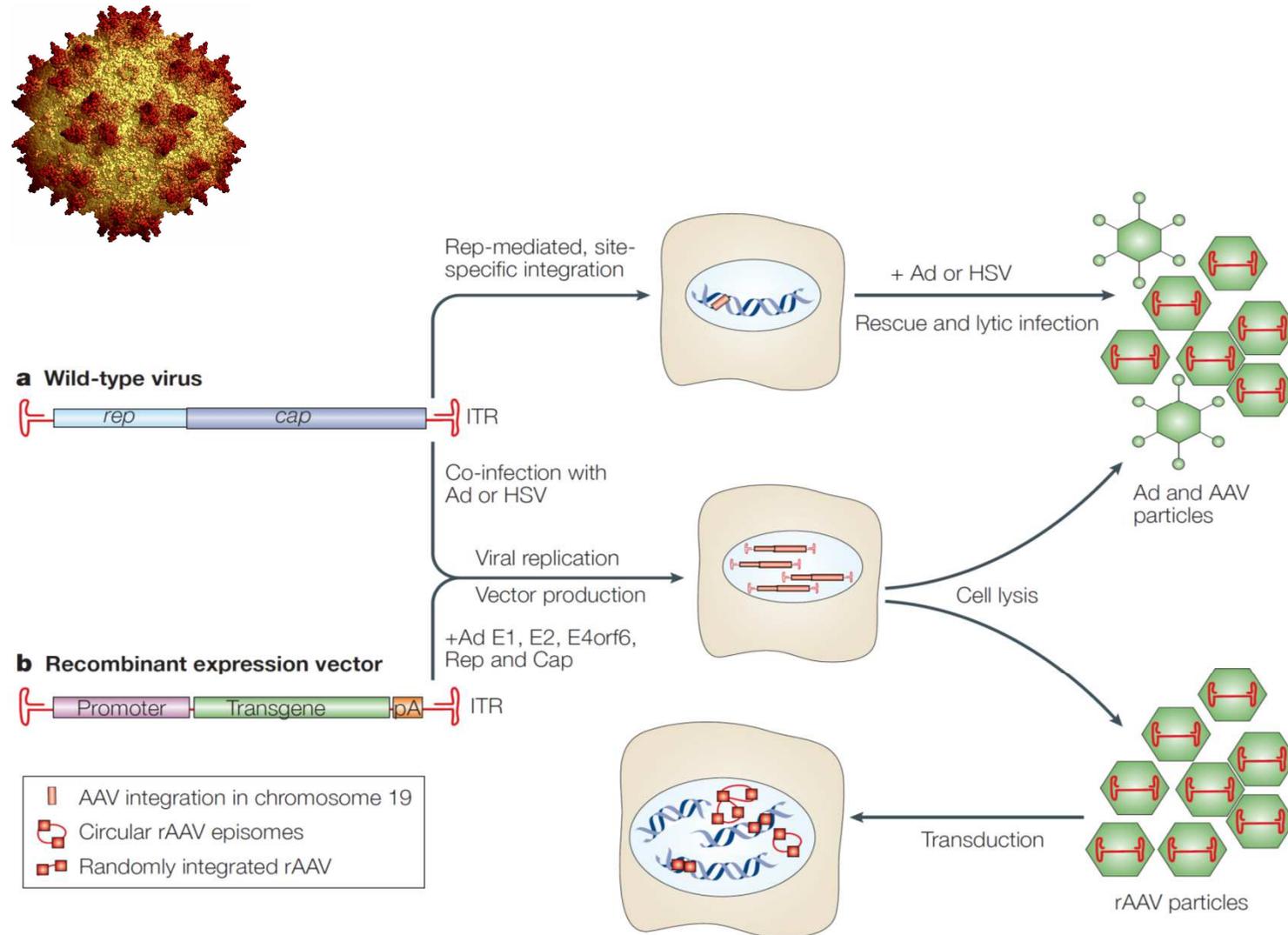
Last month, a team of scientists announced what could prove to be an enormous step forward in the fight against H.I.V.

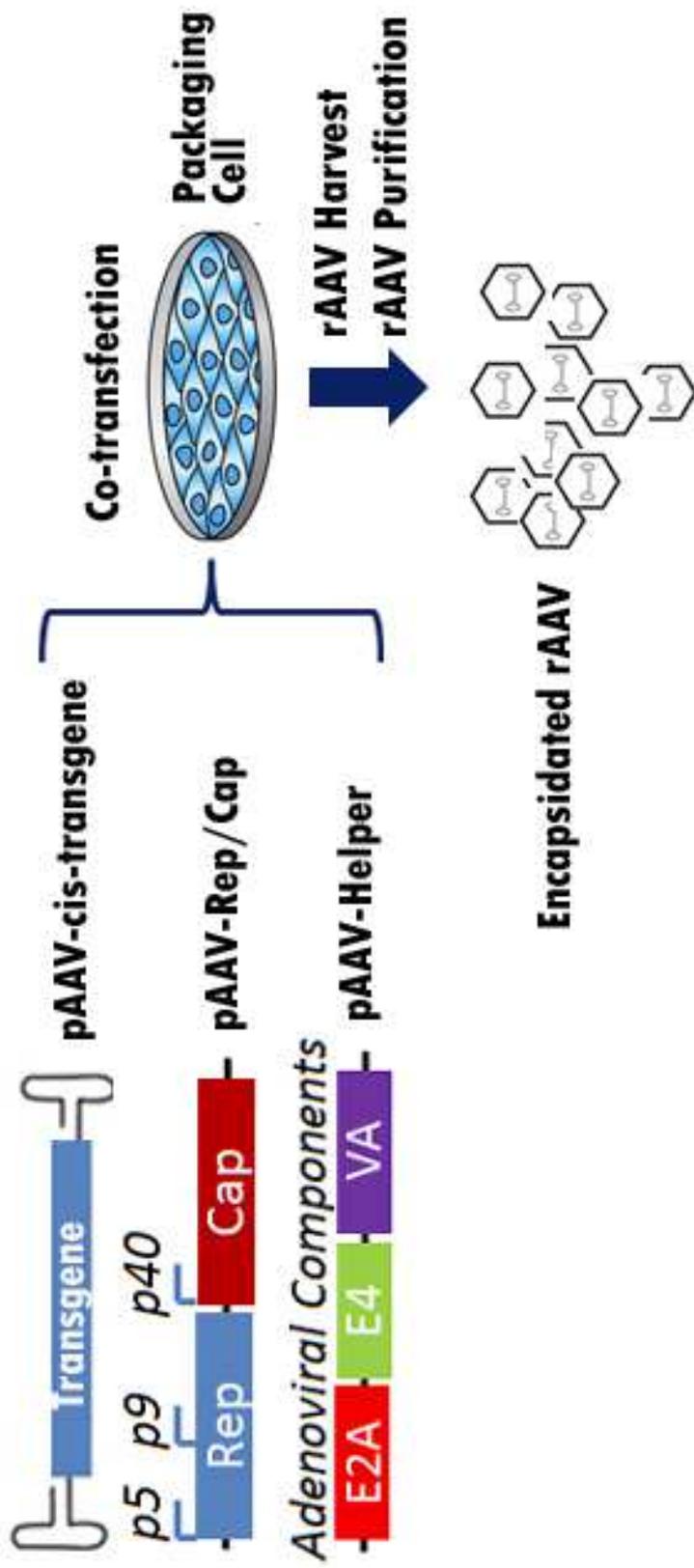
Scientists at Scripps Research Institute said they had developed an artificial antibody that, once in the blood, grabbed hold of the virus and inactivated it. The molecule can eliminate H.I.V. from infected monkeys and protect them from future infections.

But this treatment is not a vaccine, not in any ordinary sense. By delivering synthetic genes into the muscles of the monkeys, the scientists are essentially re-engineering the animals to resist disease. Researchers are testing this novel approach not just against H.I.V., but also Ebola, malaria, influenza and hepatitis.

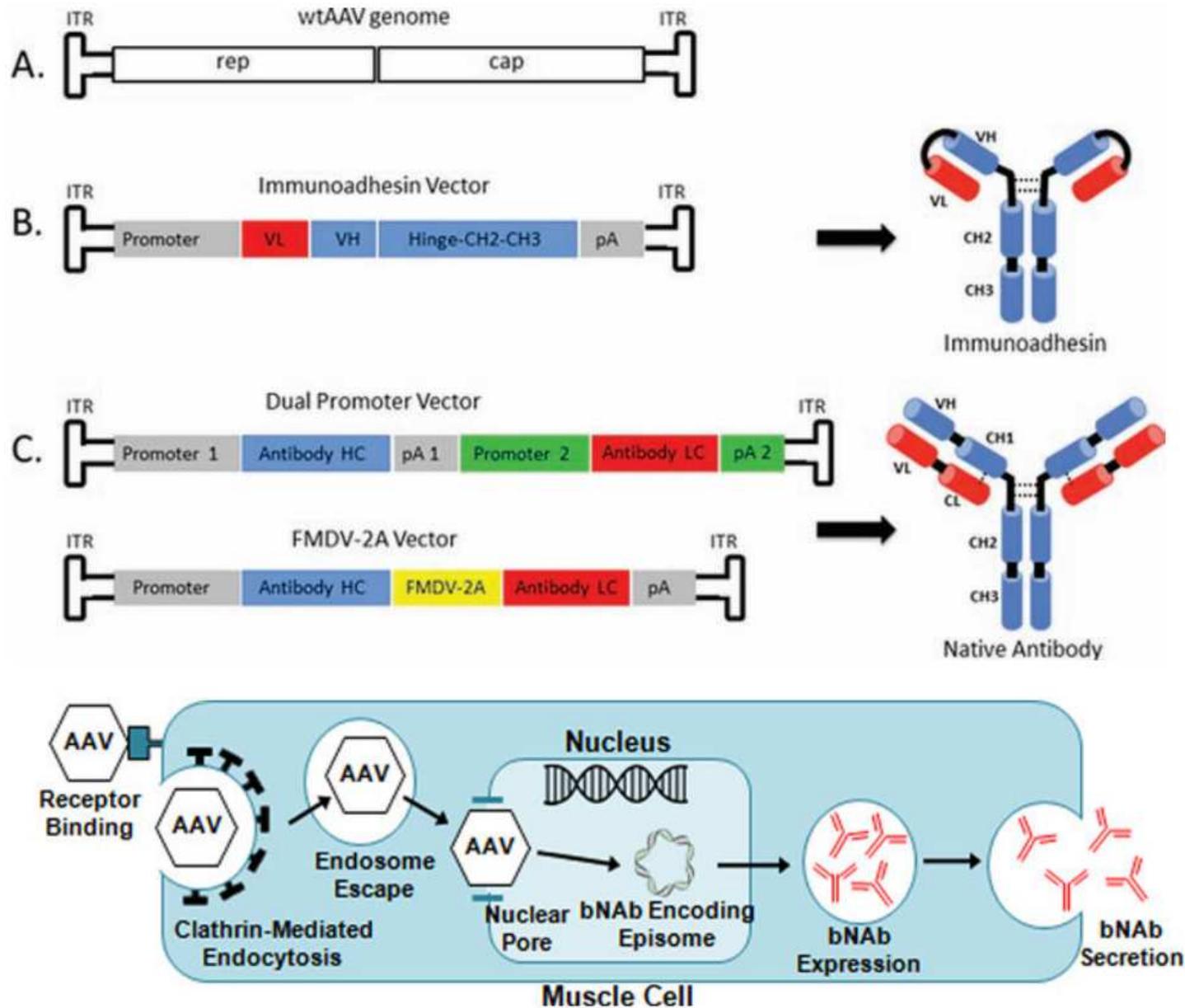


rAAV-based Technology





Engineering rAAV to deliver Nab *in vivo*



Generation of Neutralizing Activity against Human Immunodeficiency Virus Type 1 in Serum by Antibody Gene Transfer

Anne D. Lewis,¹ Ruju Chen,¹ David C. Montefiori,² Philip R. Johnson,^{1,3,4*} and K. Reed Clark^{1,3,4†}

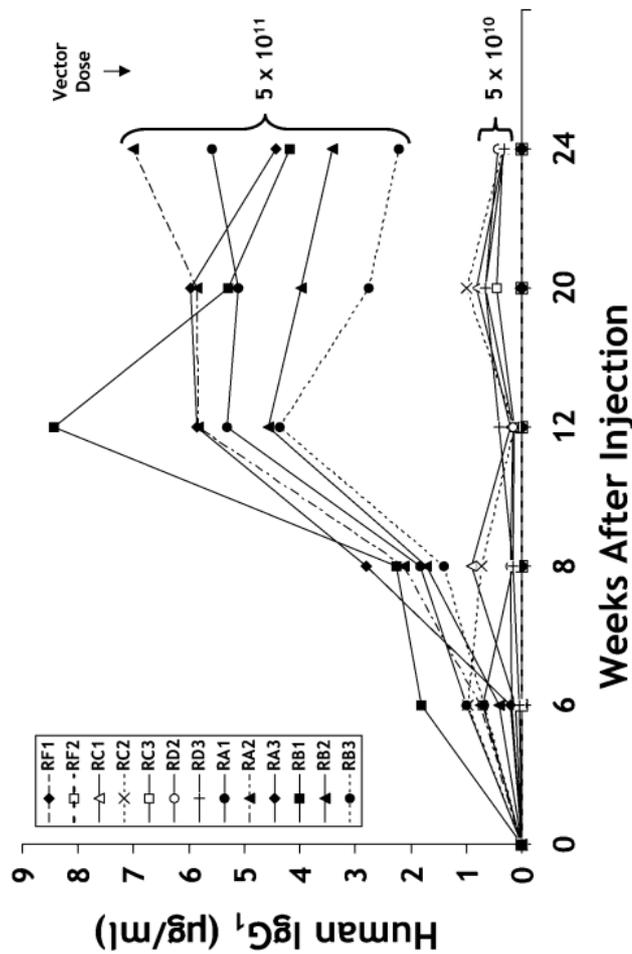


TABLE 4. Persistence of vector DNA in mouse muscle

Vector (dose) ^a	Mouse	Avg no. of genome copies/nucleus ^b
PBS	RF3	0.003
	RF4	0.012
rAAV/IgG1b12 (5×10^{10})	RC1	9.0
	RC2	5.5
	RC3	0.5
	RD2	1.2
	RD3	4.5
rAAV/IgG1b12 (5×10^{11})	RA1	42.5
	RA2	3.2
	RA3	20.0
	RB1	27.1
	RB2	0.4
	RB3	19.3

^a Dose is measured as the DRP (see Materials and Methods).

^b Values represent the average number of rAAV genomes per nucleus observed in the quadriceps muscles after rAAV injection. A total of 60 ng of muscle DNA (10,000 nucleus equivalents) was analyzed by quantitative Taqman PCR by using the CMV primer-probe set (see Materials and Methods). All samples were harvested 24 weeks after injection.



BIOTEC
a member of NSTDA

เนคเทค
NSTDA

Virology and Cell Technology Laboratory

**National Center for Genetic Engineering
and Biotechnology (BIOTEC), NSTDA**

Email: anan.jon@biotec.or.th



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NATIONAL CENTER FOR GENETIC ENGINEERING AND BIOTECHNOLOGY

Thank you