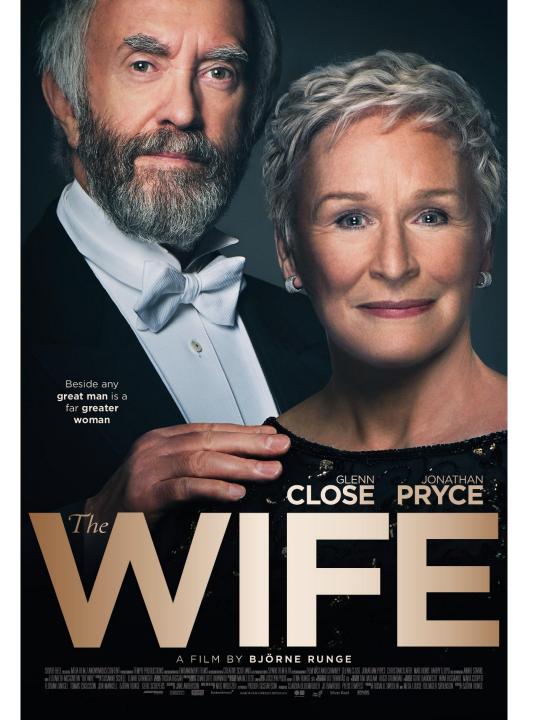
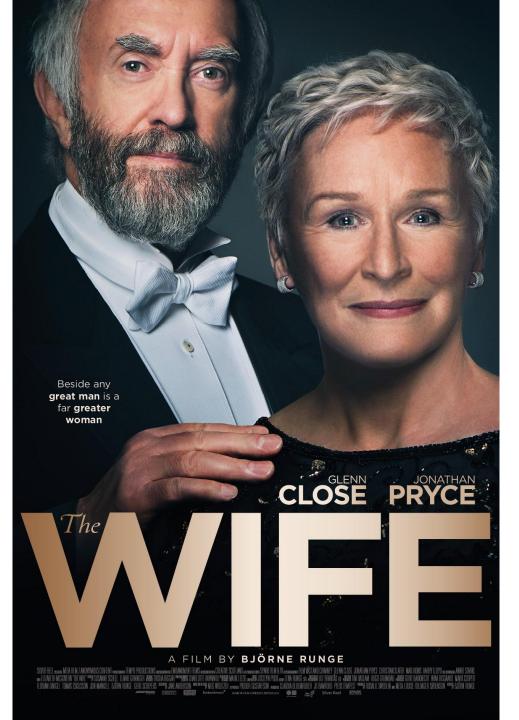
Phage Display: Simple Evolution in a Petri Dish





George P. Smith
Division of Biological Sciences
University of Missouri
Nobel Prize Lecture in Chemistry
December 8, 2018

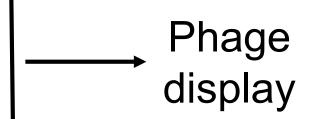




Margie did not invent phage display

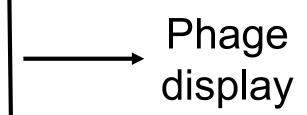
My science community

- Molecular biologists
- Evolutionary biologists
- Immunologists
- Protein chemists
- Phage biologists
- Mathematicians
- Philosophers of science
- Bayesian statisticians

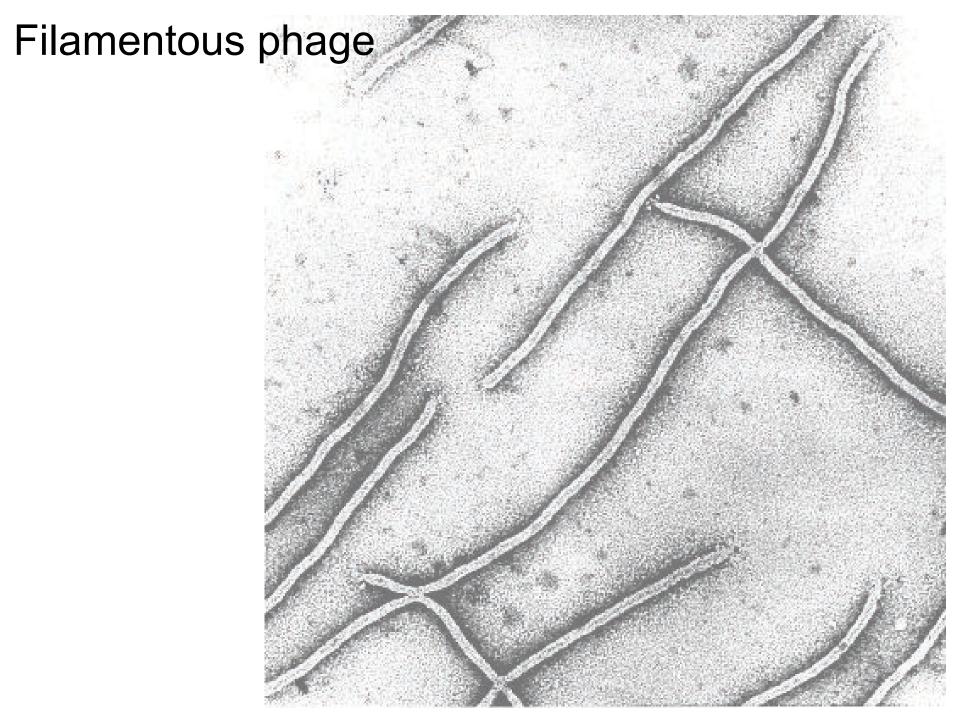


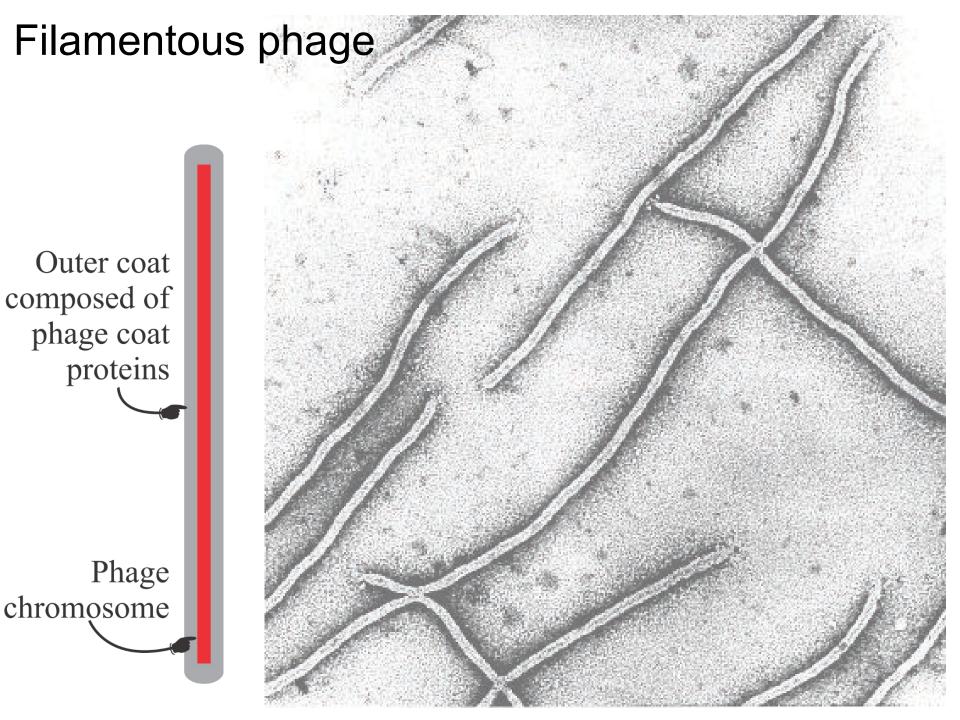
Our science community

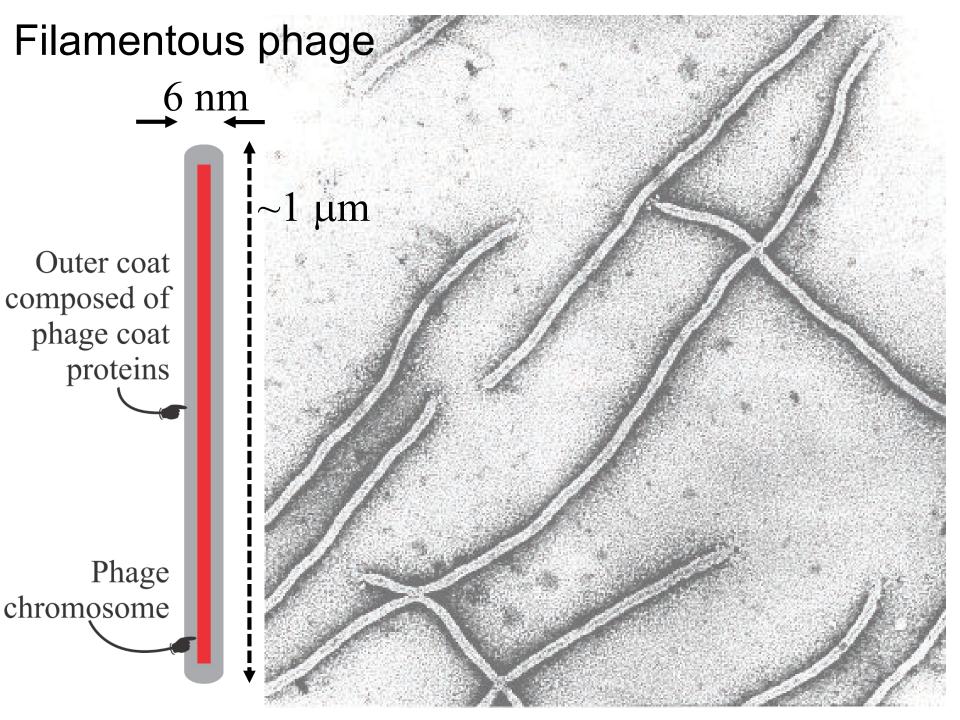
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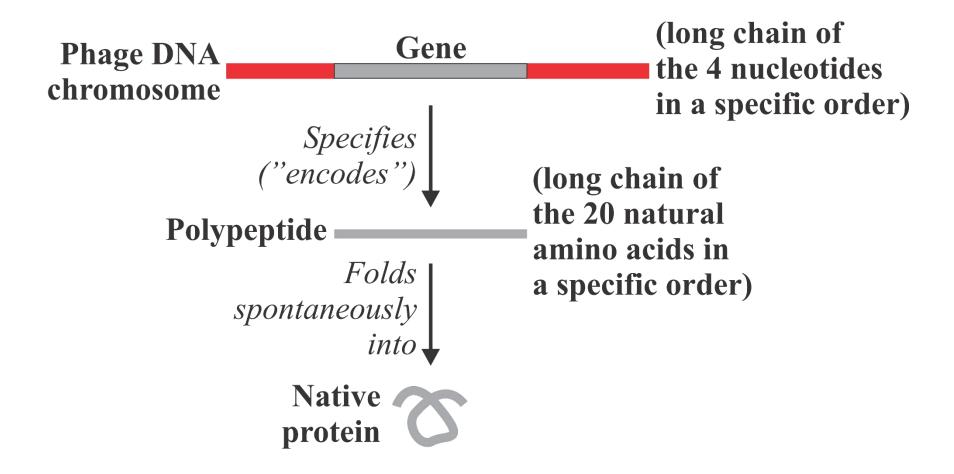
Phage = virus that infects bacteria



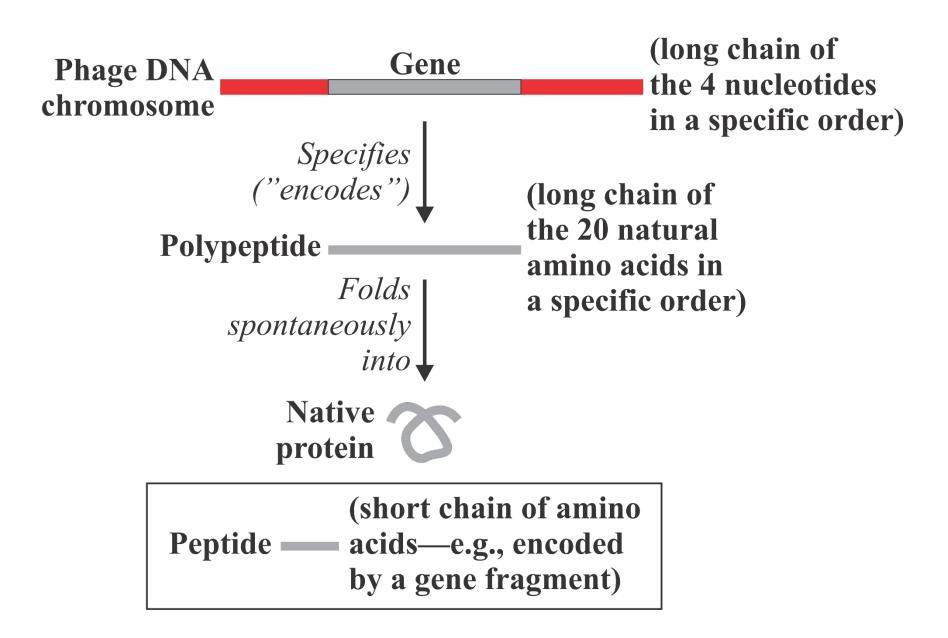




Crash course in molecular biology

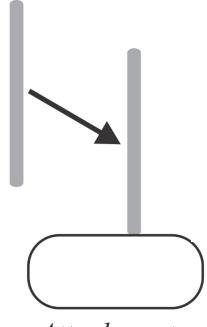


Crash course in molecular biology

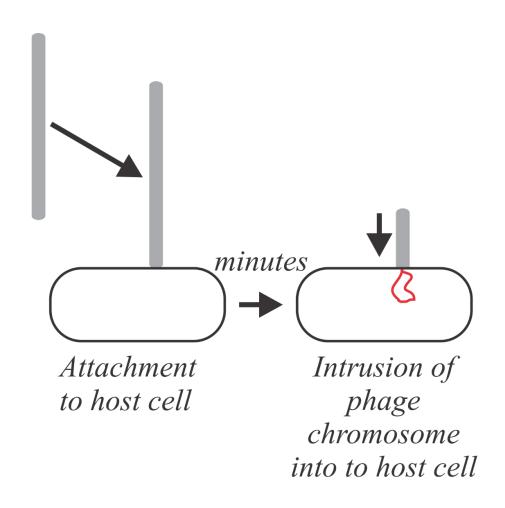


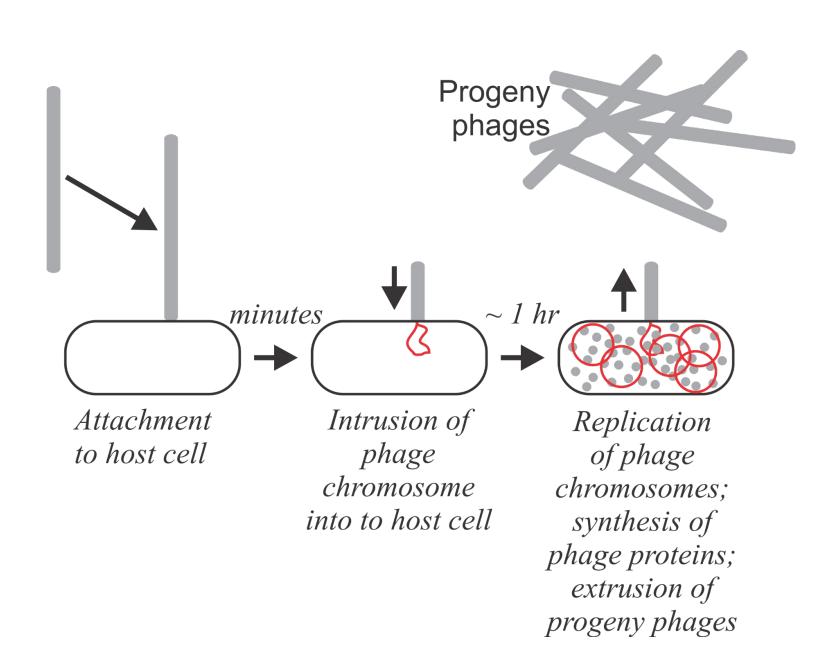
Filamentous phage infection cycle (simplified)

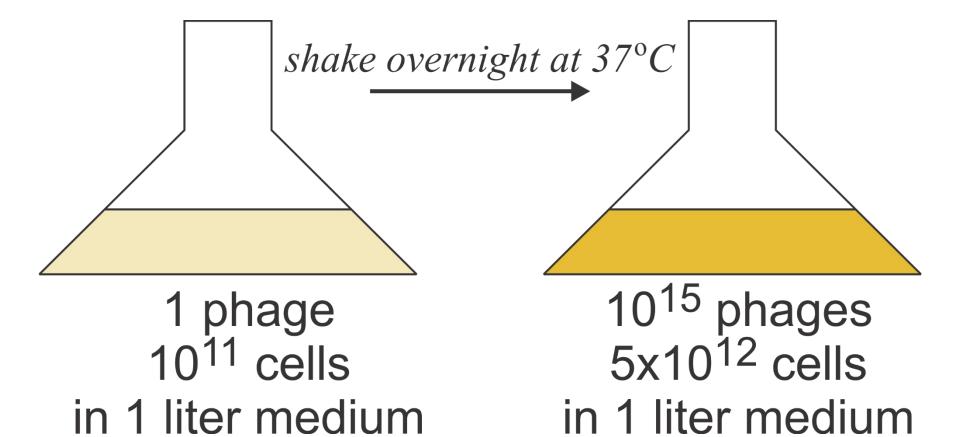
Bacterial cell



Attachment to host cell









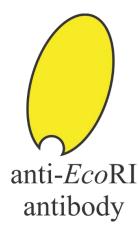
Bob Webster, Department of Biochemistry, Duke University (now retired to North Carolina coast)

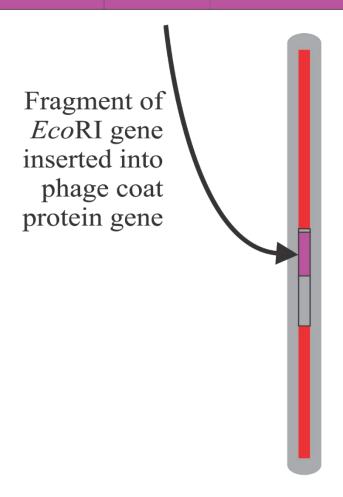
This coat protein gene encodes a coat protein that's partly exposed at one tip of the phage



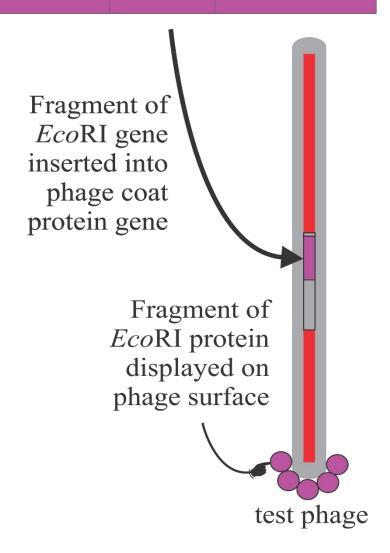
Paul Modrich Nobel Prize in Chemistry, 2015

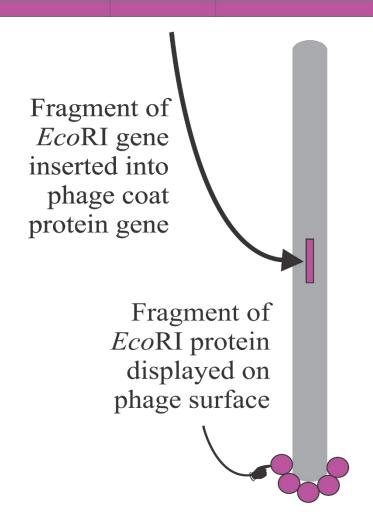
EcoRI gene

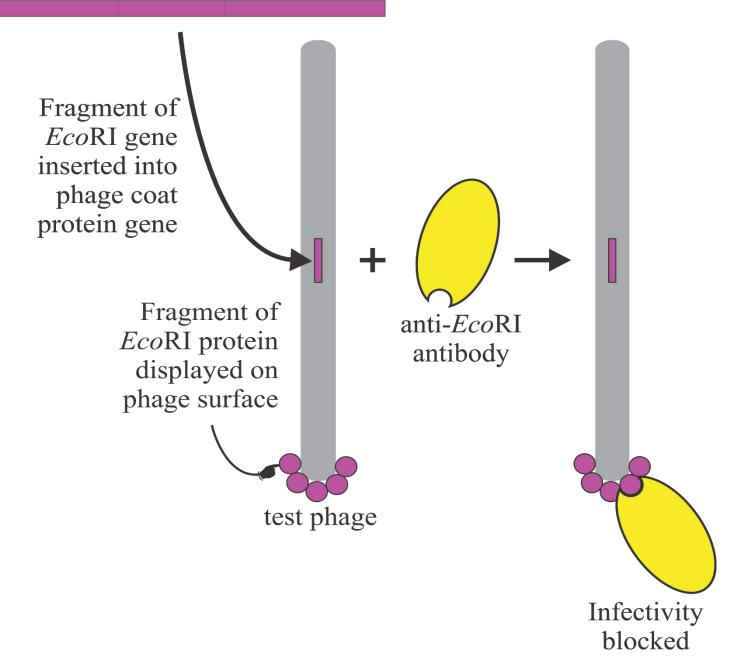




test phage





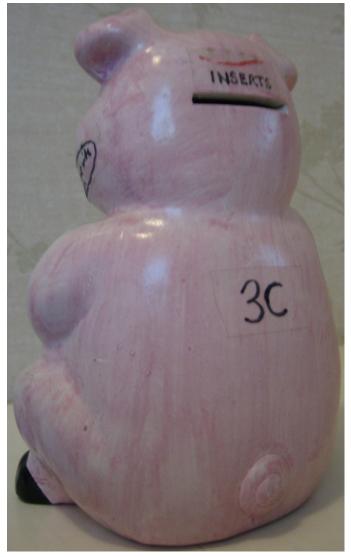


Infectivity

blocked

Why is that interesting? Please wait...





Steve developed a practical phage display vector (after some false starts like pIG3C) and affinity selection as grad student, 1985-1988.

Darn good colleague Jamie Scott

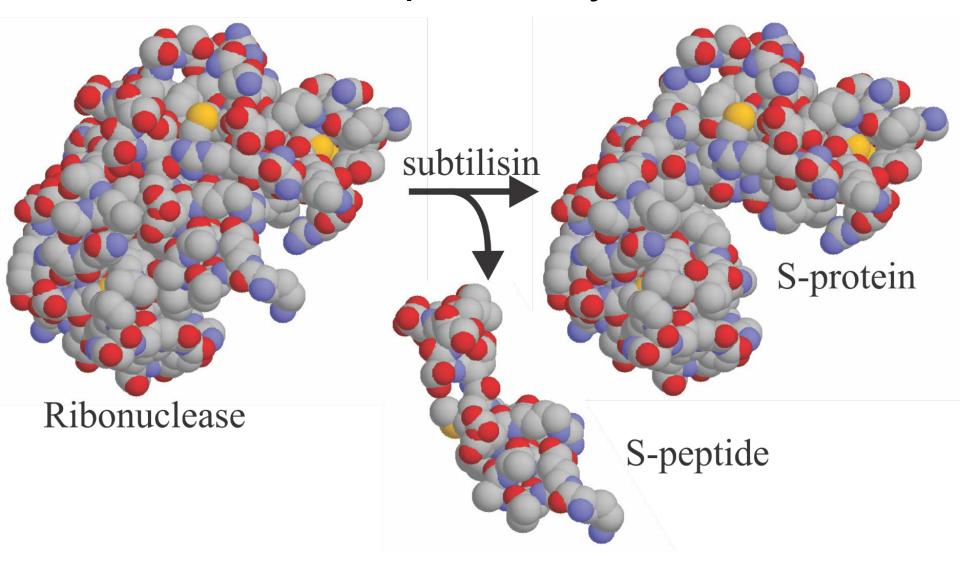
Jamie Scott first demonstrated affinity selection of peptides from large random peptide libraries as postdoc, 1988-1991.



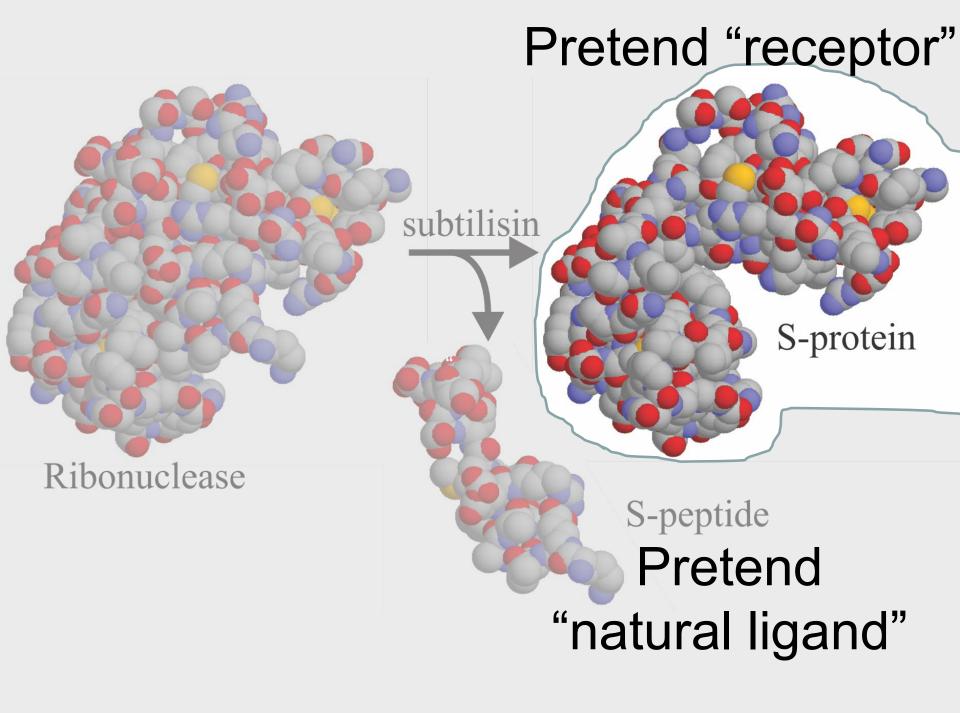


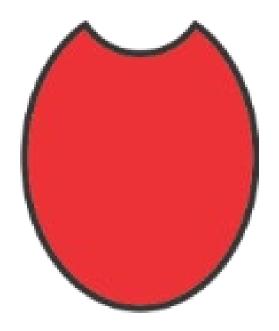
Robert Davis came to the lab as chief manager and technician in the summer of 1989. We calculate that he sequenced a million DNA bases using old-fashioned radioactive technology.

The S-protein system



Fred Richards at Yale, late 50's; brought to our lab by John Ladbury and David Schultz



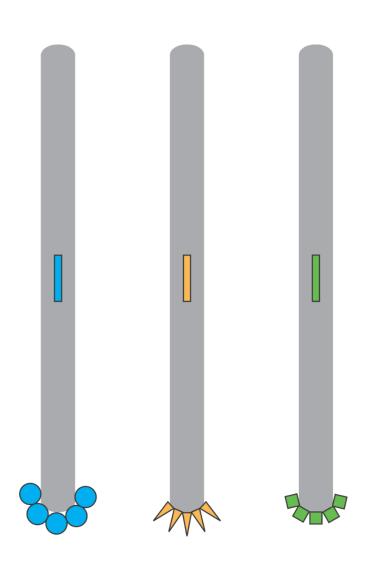


S-protein "receptor"

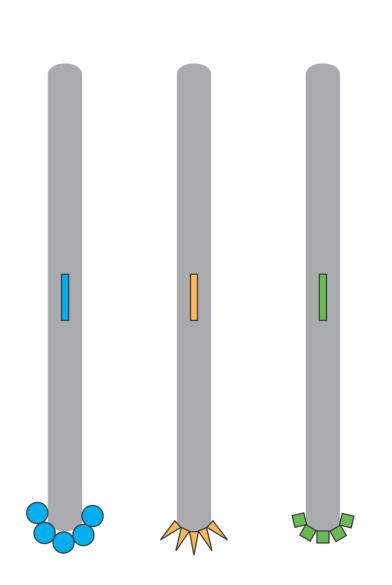


Jinan Yu, now a researcher at Hainan University, China.

Random peptide library

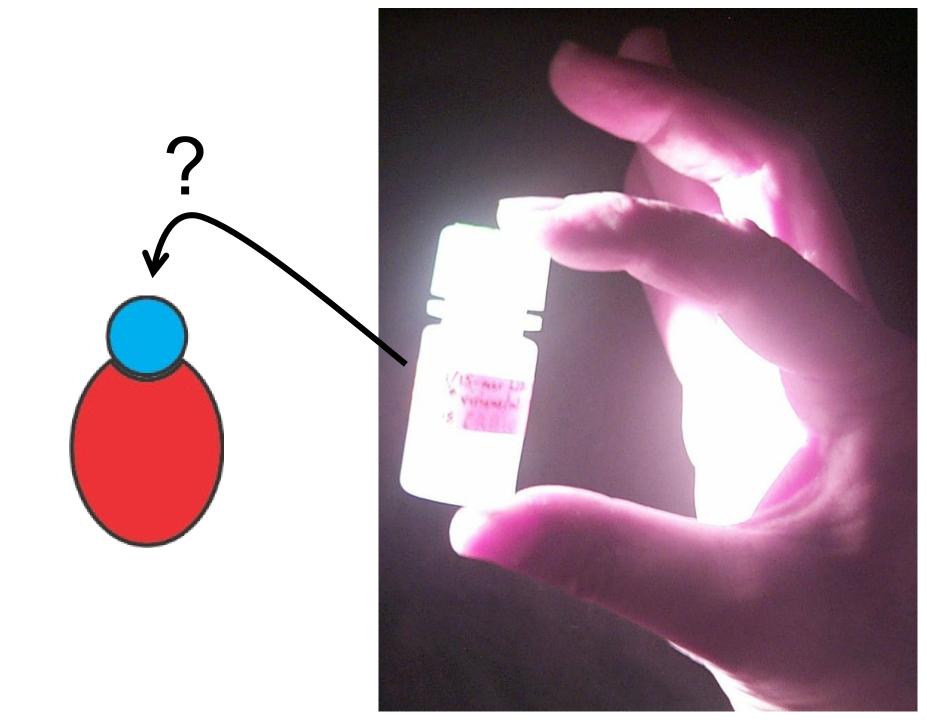


Random peptide library

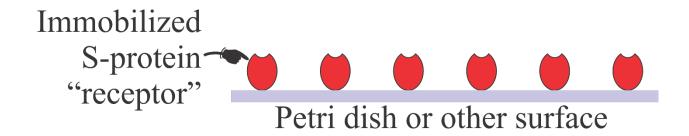


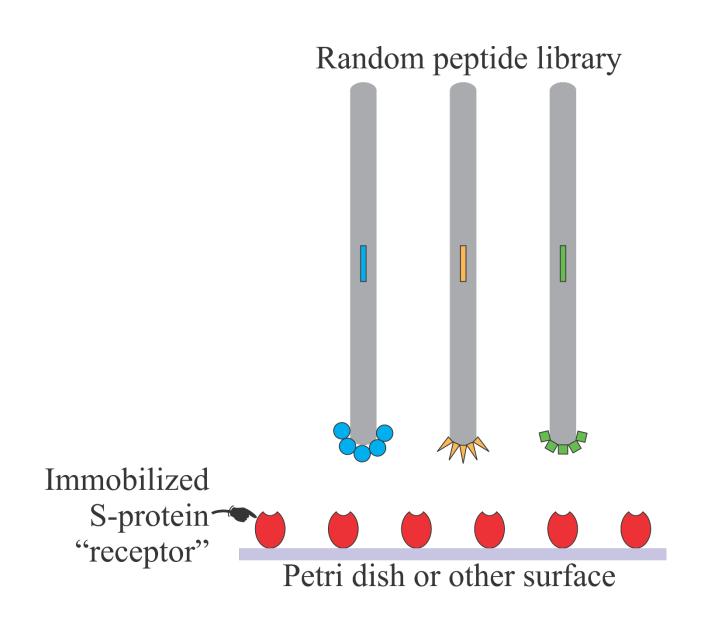


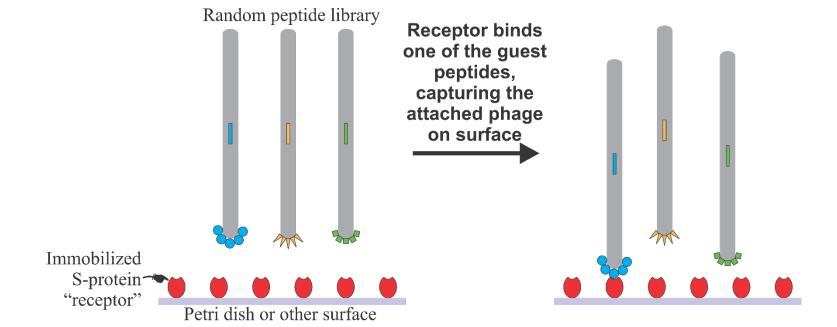
10¹⁵ phages representing 250 million phage clones, each clone displaying a different 15-amino acid guest peptide. [T. Nishi et al., FEBS Letters 399, 237–240 (1996)]

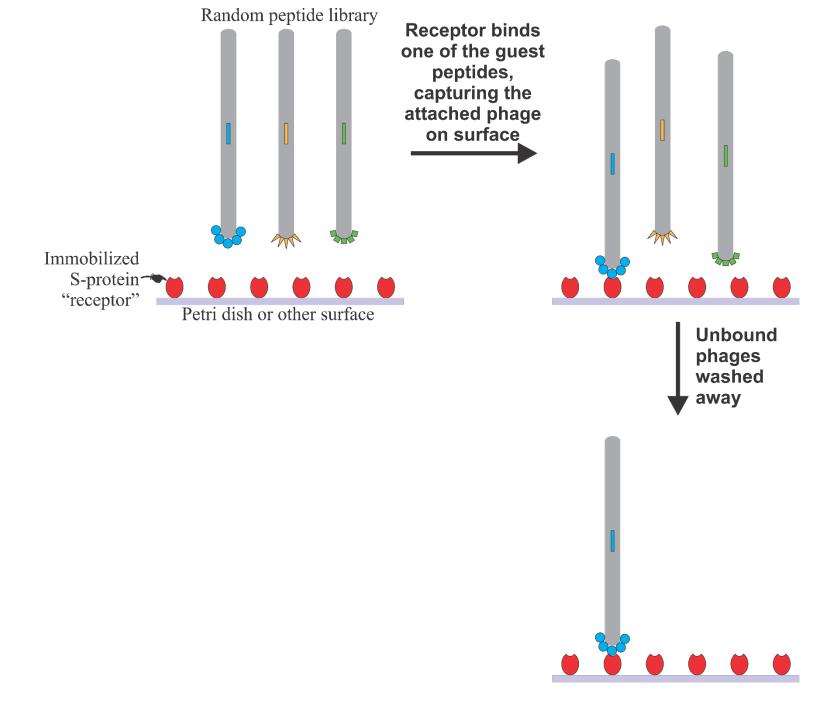


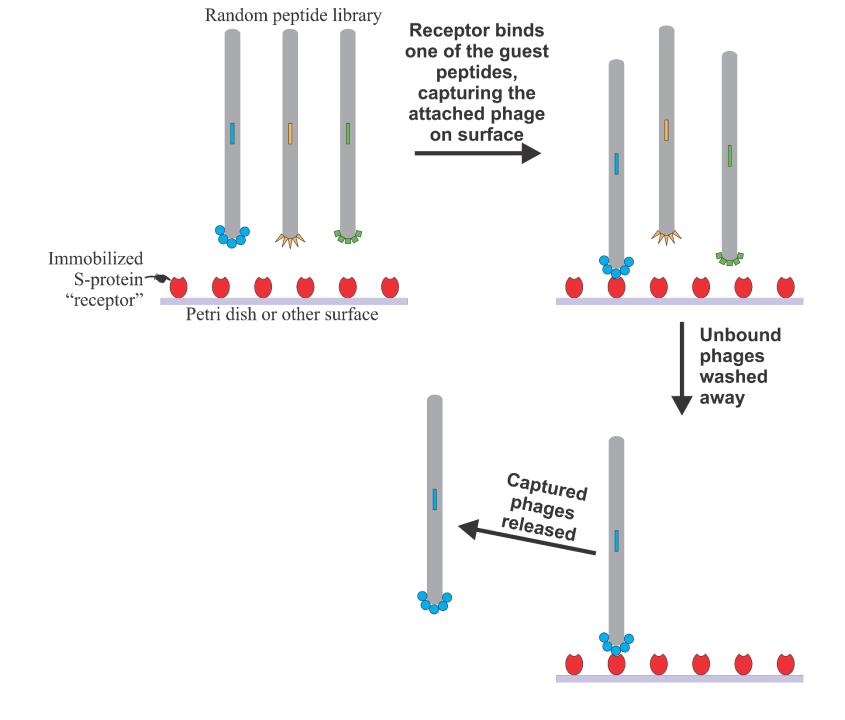
Affinity selection

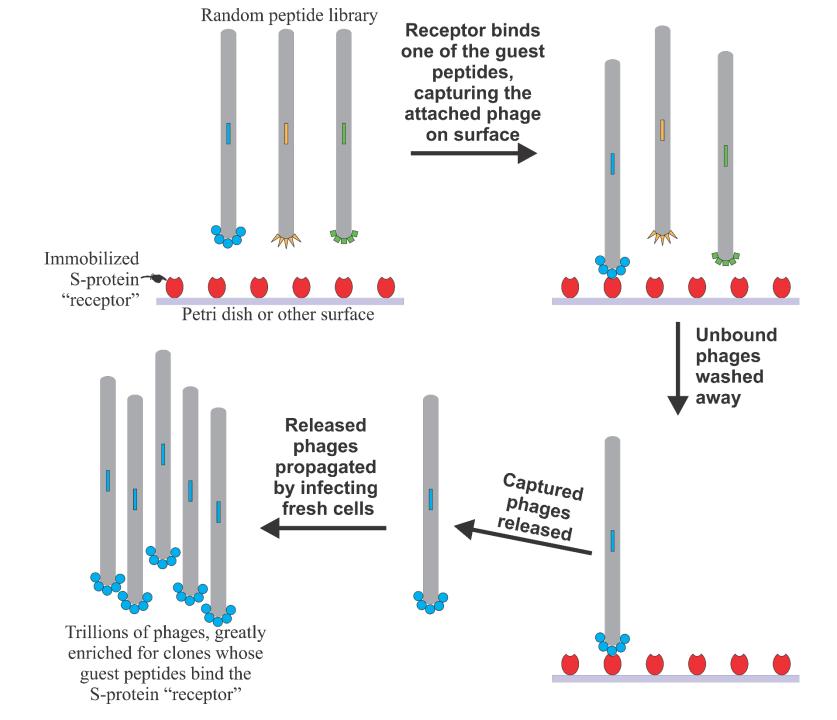












Dominant sequence among selected peptides

Selected peptide NRAWSEFLWQHLAPV

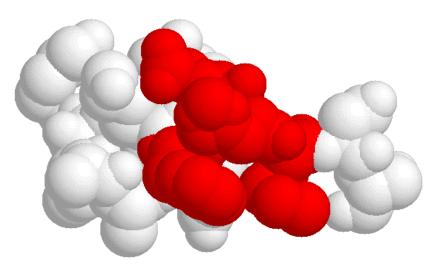
(one-letter abbreviations for amino acids)

Dominant sequence among selected peptides aligns with S-peptide "natural ligand"

Selected peptide NRAWSEFLWQHLAPV

S-peptide KETAAAKFERQHMDSSTSAA

Buried amino acids



Artificial evolution in the petri dish

Evolution in the living world

- Diversification
- Natural selection
- Adaptation

Affinity-selection from random peptide libraries

- Construction of library
- Affinity selection
- Peptide with desired activity

