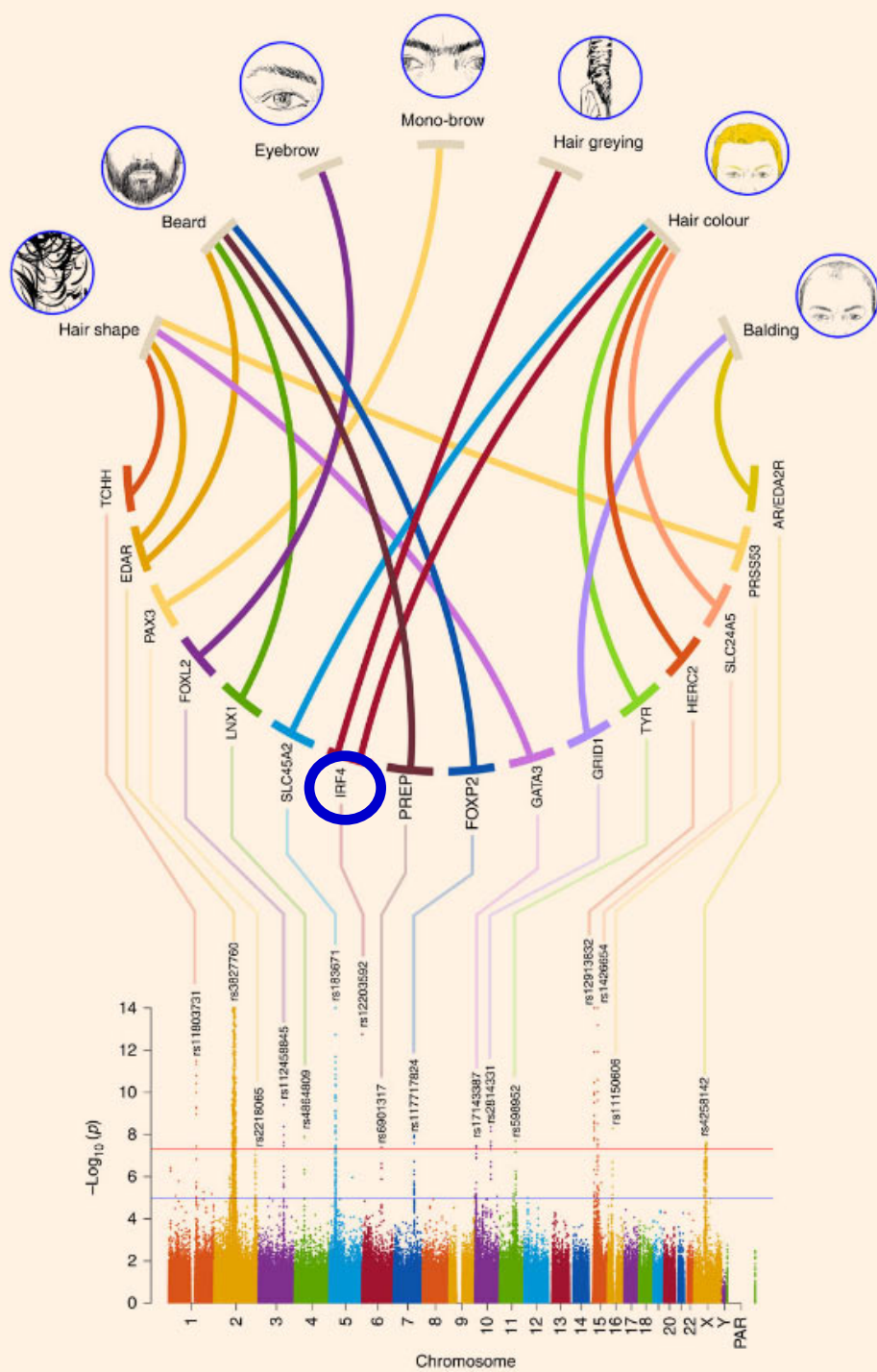
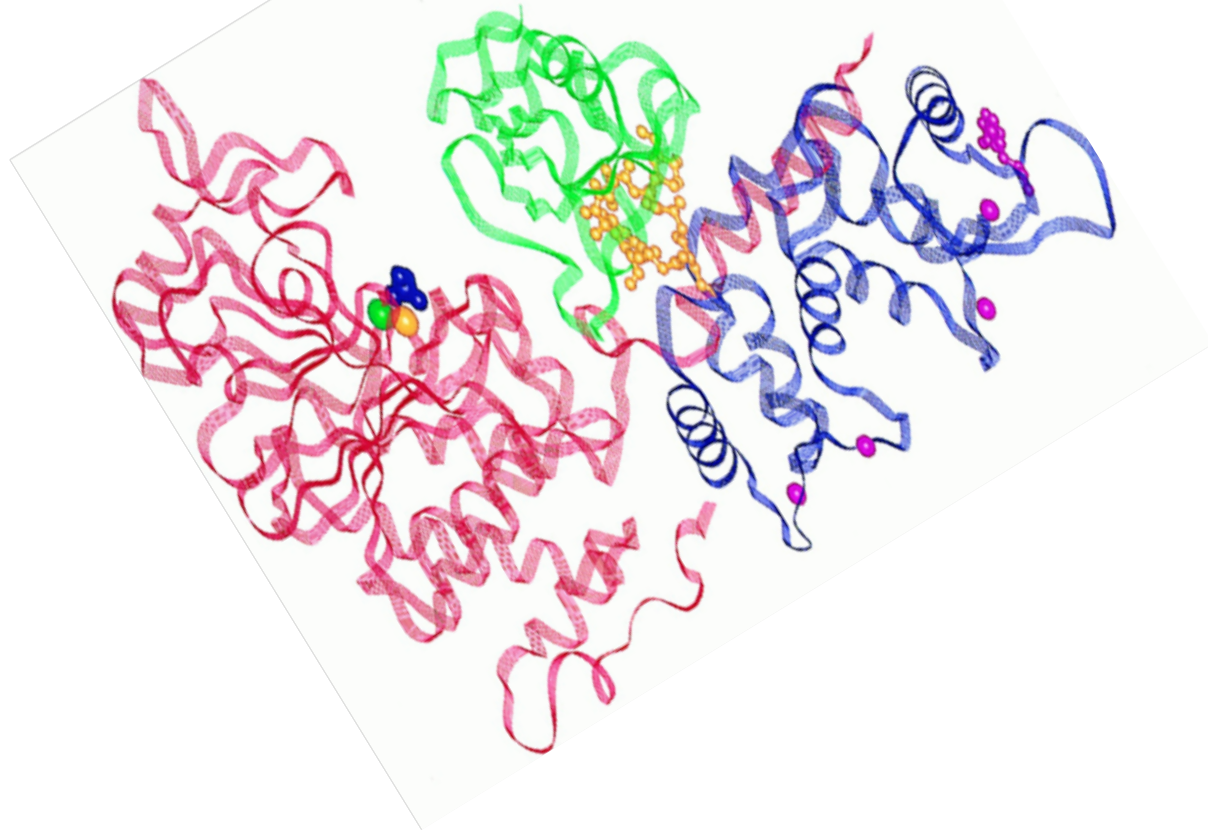


Which process or biomolecule would you study with a chemical probe if you had one in hand?

# Genome Wide Association Study: Hair-related phenotypes

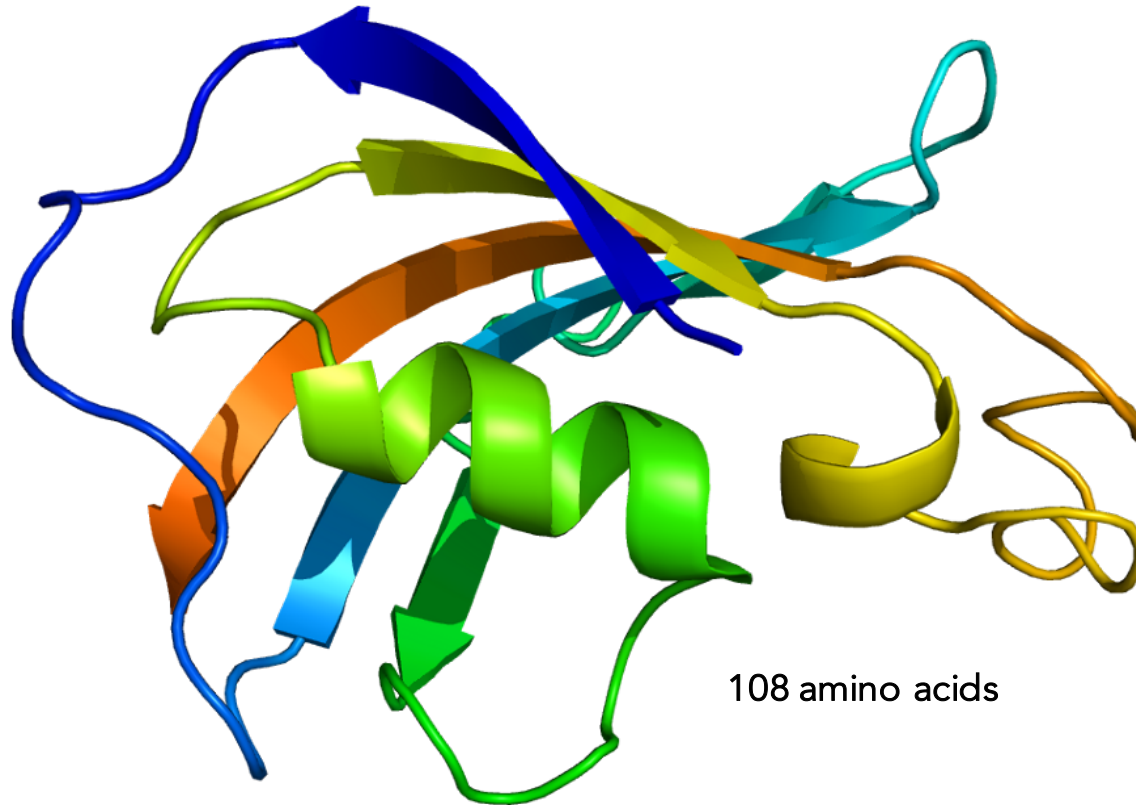


# For the love of proteins: FKBP12 and immunophilins



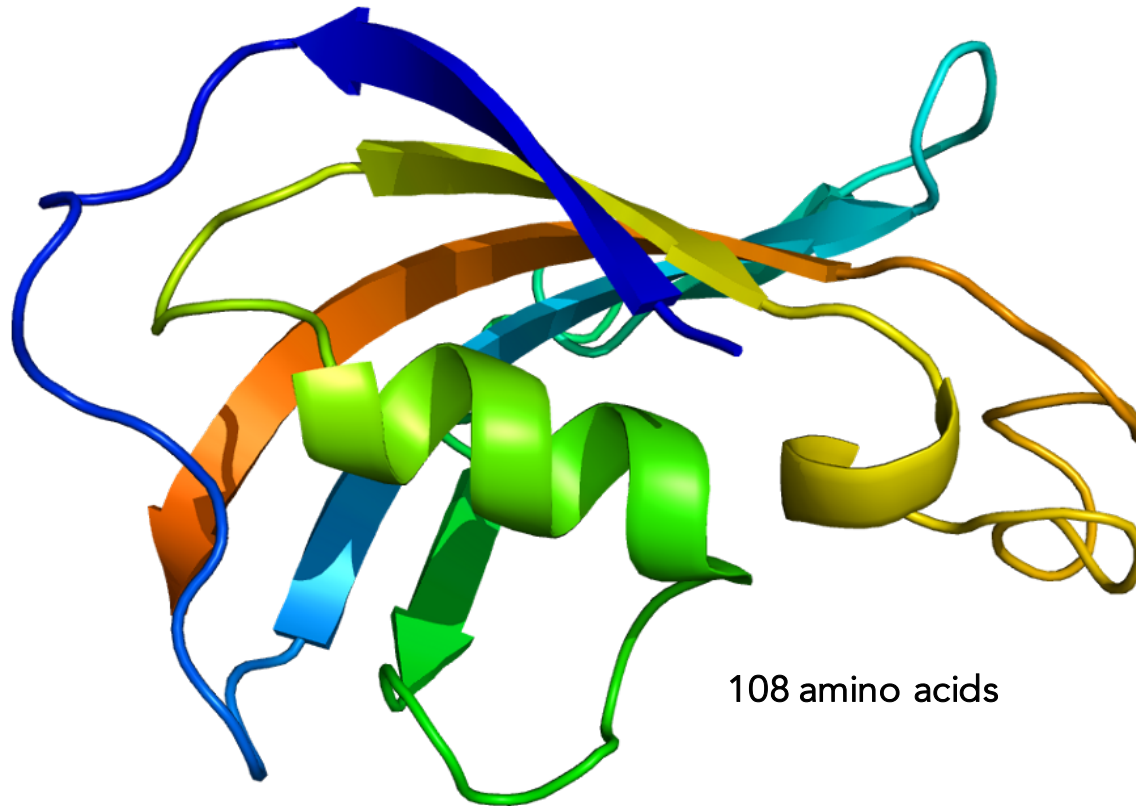
February 16, 2017

Our target protein: **FKBP12**



**FK-506** Binding Protein that is **12** kilodaltons

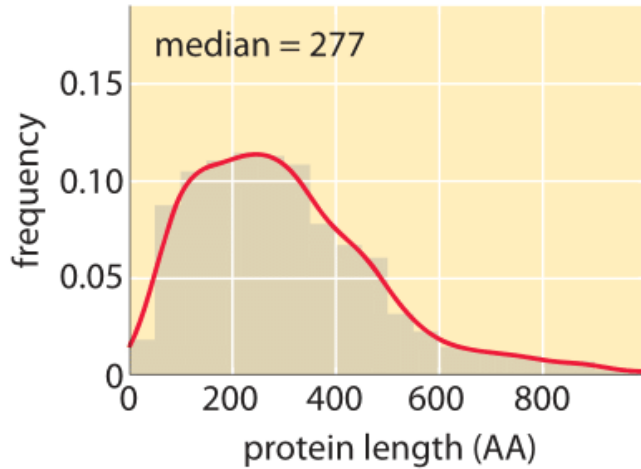
Our target protein: **FKBP12**



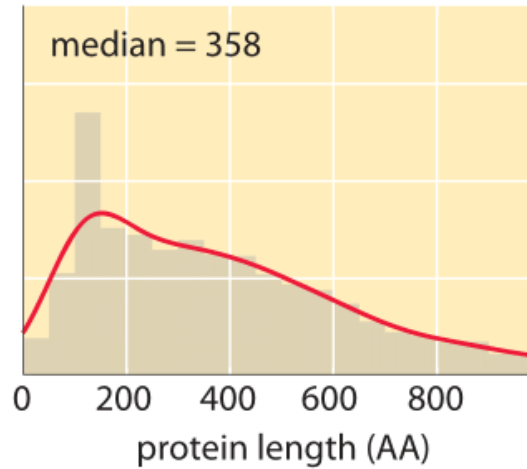
**FK-506** Binding Protein that is **12 kilodaltons**

# How big is the typical protein?

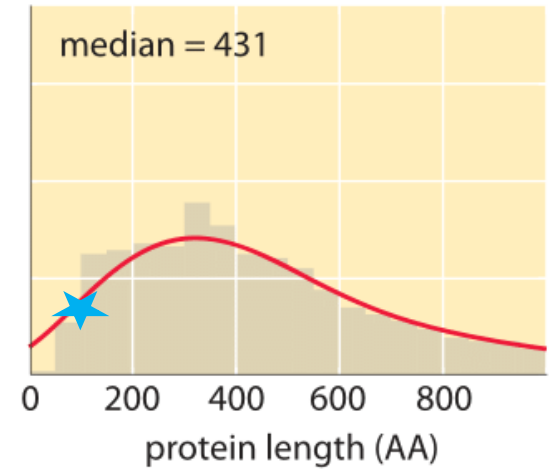
*E. coli* [N=4,303]



budding yeast [N=6,723]



human HeLa [N=22,257]



fkbp12  
12 kDa

insulin  
5.8 kDa

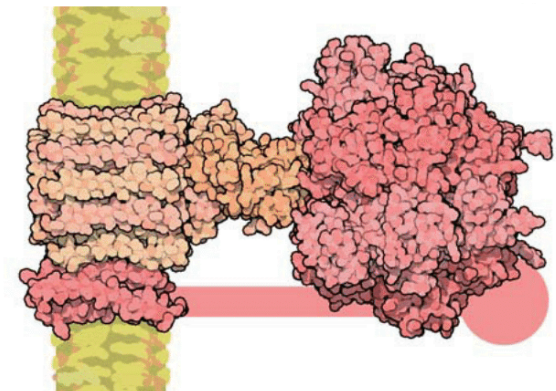
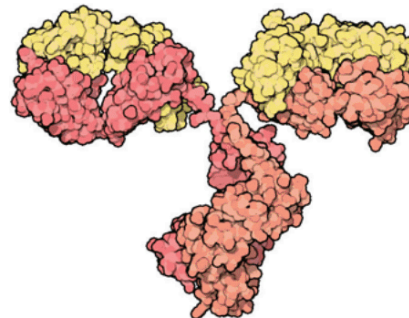
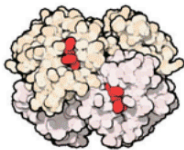
trypsin  
23.3 kDa

hemoglobin  
64.5 kDa

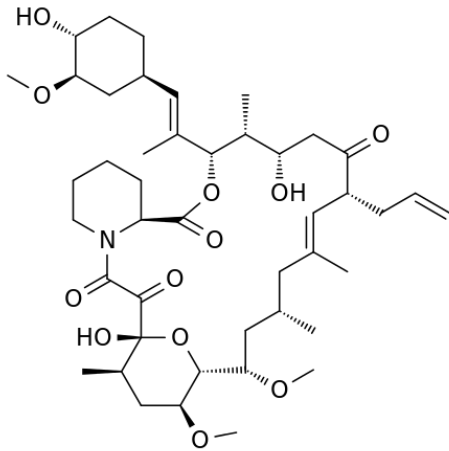
hexokinase  
102 kDa

immunoglobulin G  
150 kDa

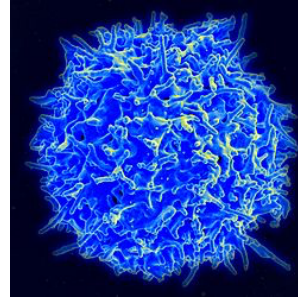
ATP synthase complex  
>500 kDa



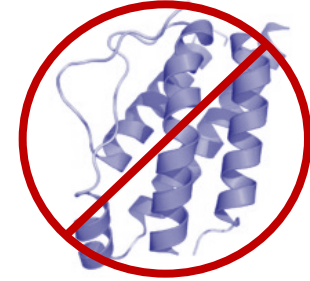
# FK-506 is an immunosuppressant drug



**FK-506 (Tacrolimus)**  
inhibits the development of  
T-cells for immune responses



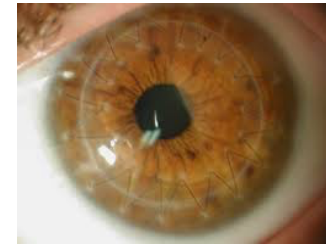
blocks T-cell proliferation



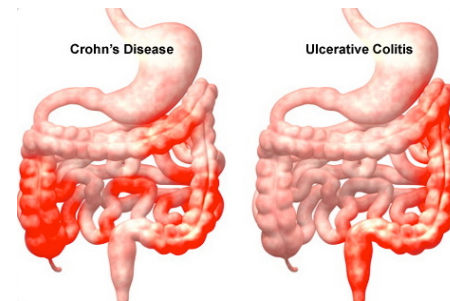
inhibits secretion of  
interleukin-2



eczema and other  
skin conditions

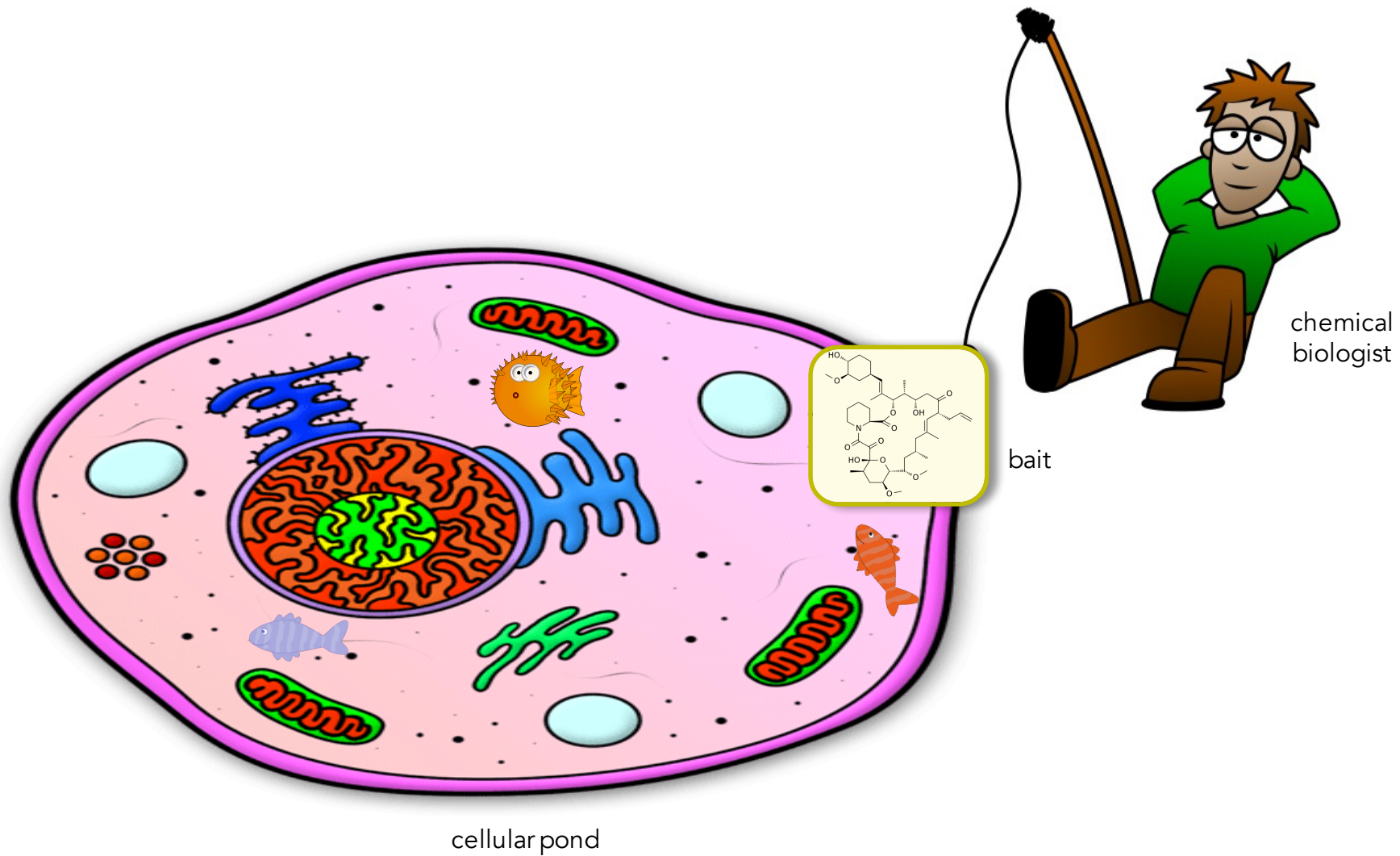


transplant rejection



inflammatory bowel disorders

# Fishing for the target of FK506

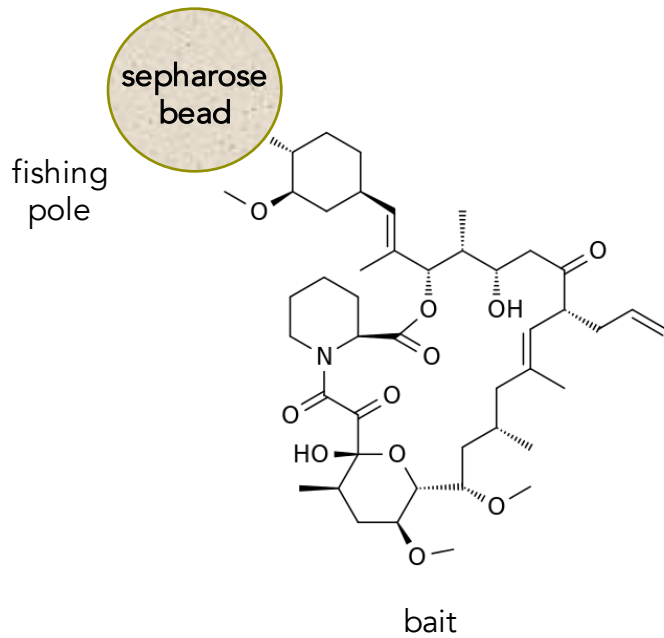




# Fishing for the target of FK506

## affinity chromatography

couple 'bait' to matrix



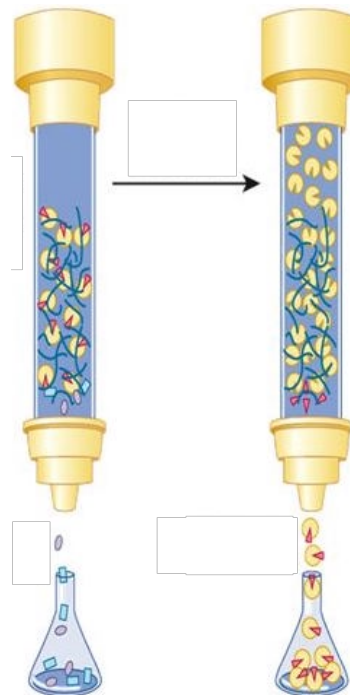
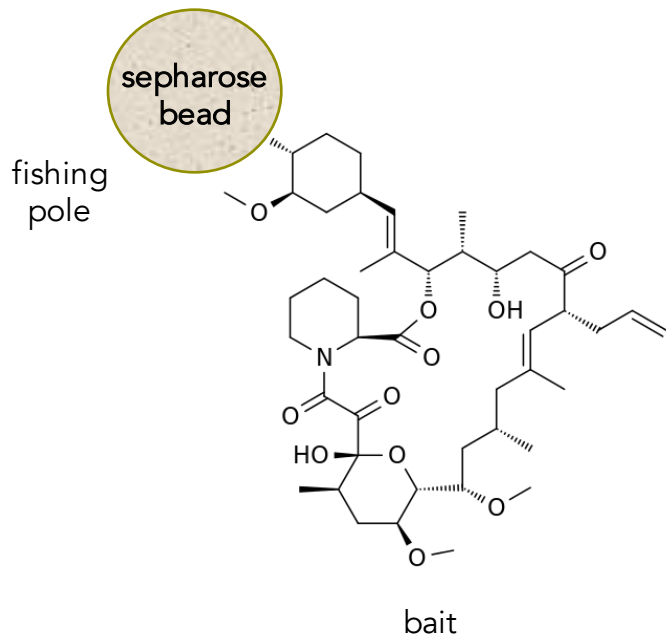
# Fishing for the target of FK506

## affinity chromatography

couple 'bait' to matrix

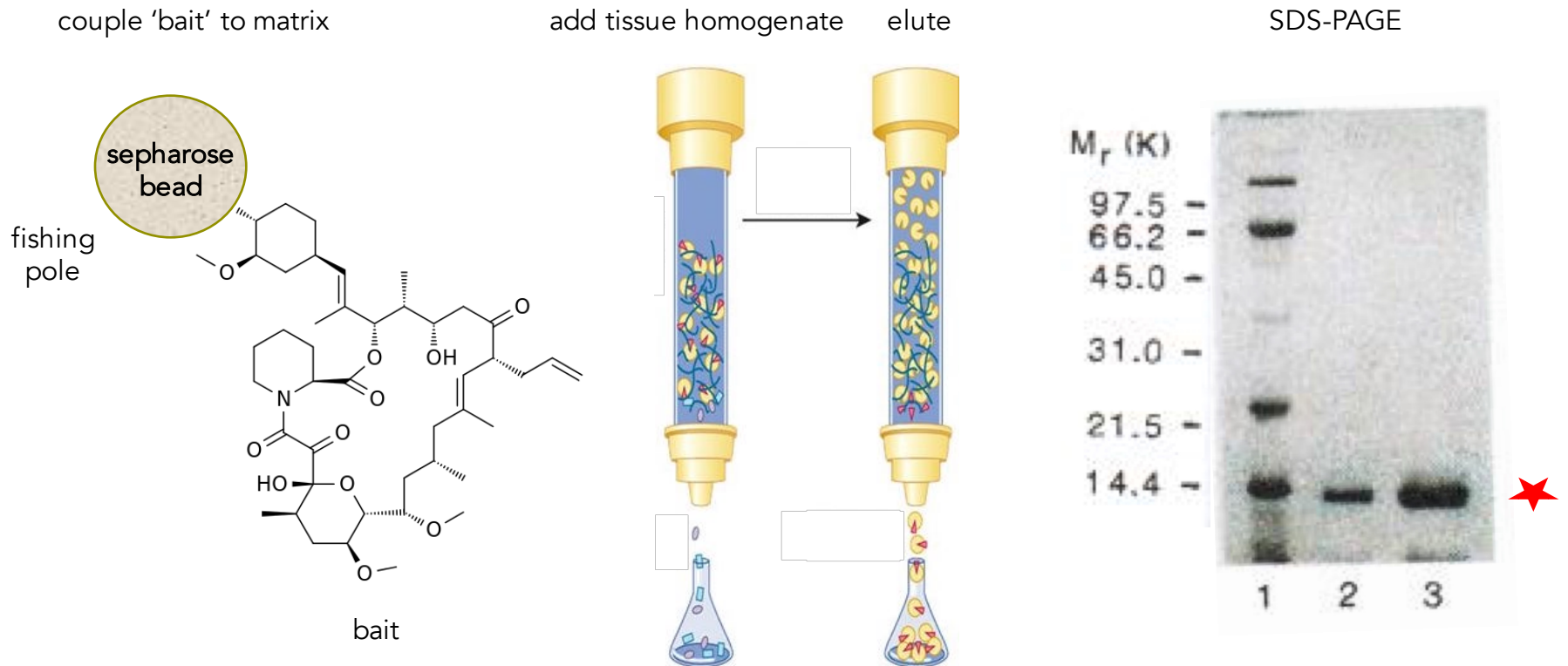
add tissue homogenate

elute



# Fishing for the target of FK506

## affinity chromatography



Harding *et al.* Nature 341, 758-760 (1989)  
Siekierka *et al.* Nature 341, 755-757 (1989)

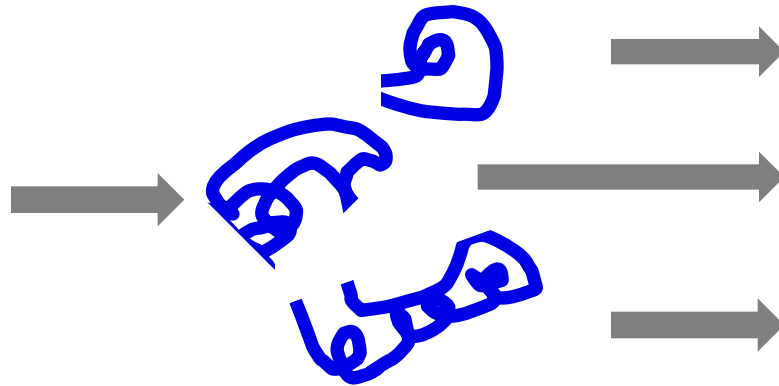
# Fishing for the target of FK506

determining the protein sequence

purified FKBP protein



FKBP proteolytic fragments



N-terminal sequencing  
methods

IHRTPATS

RGRTHAT

TYTLTWN

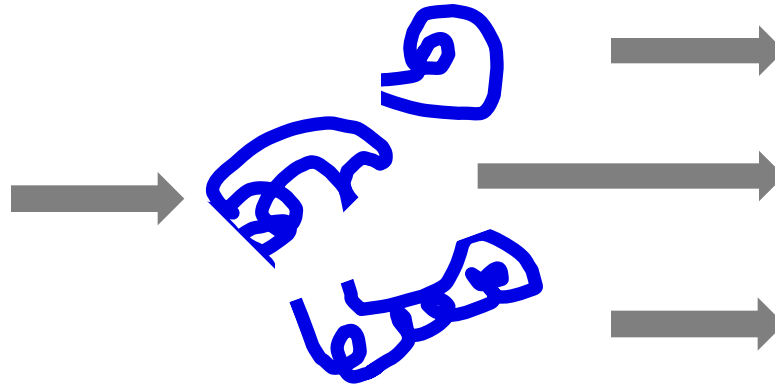
# Fishing for the target of FK506

## determining the protein sequence

purified FKBP protein



FKBP proteolytic fragments

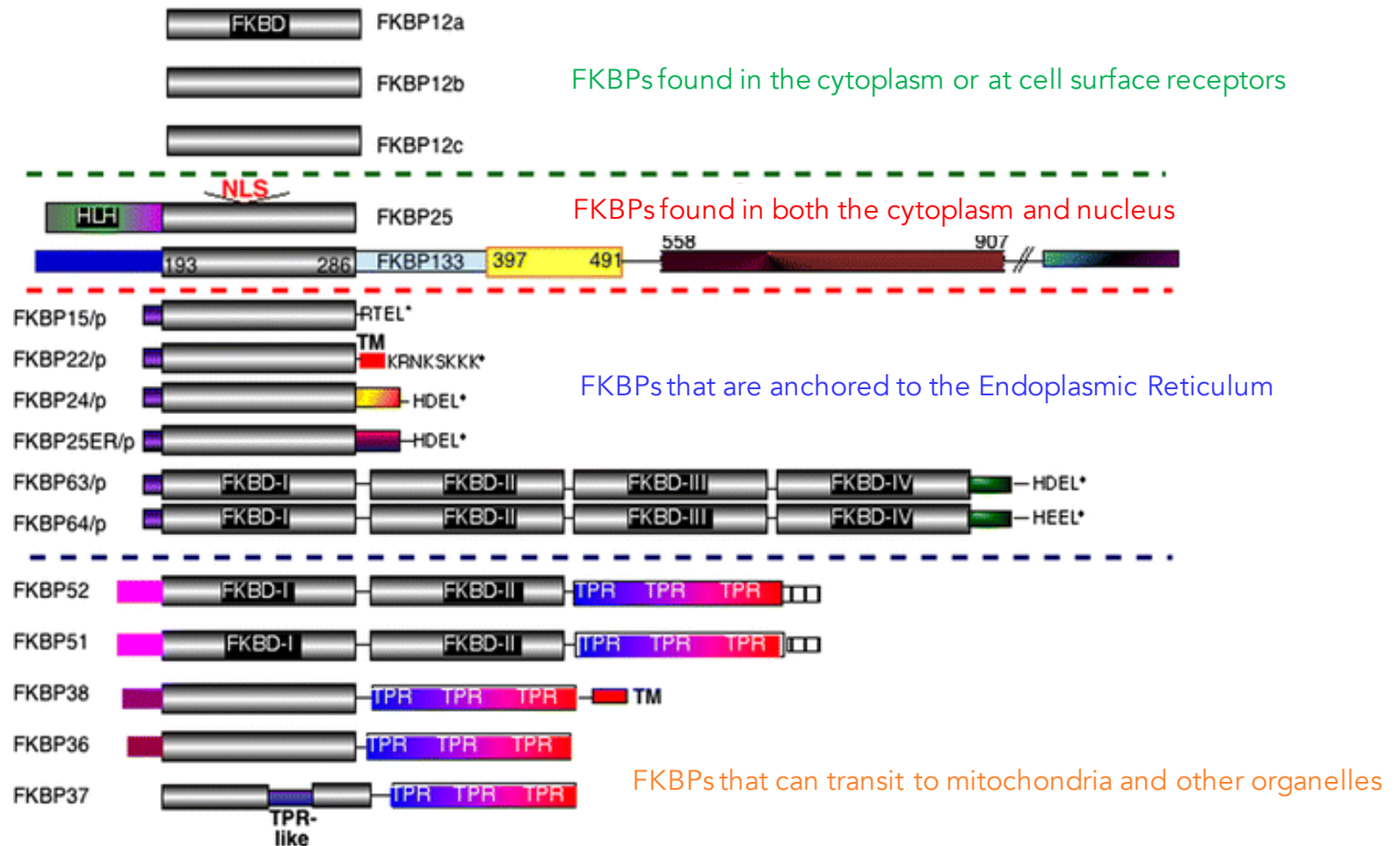


108 amino acid peptide chain  
(11.8 kilodaltons)

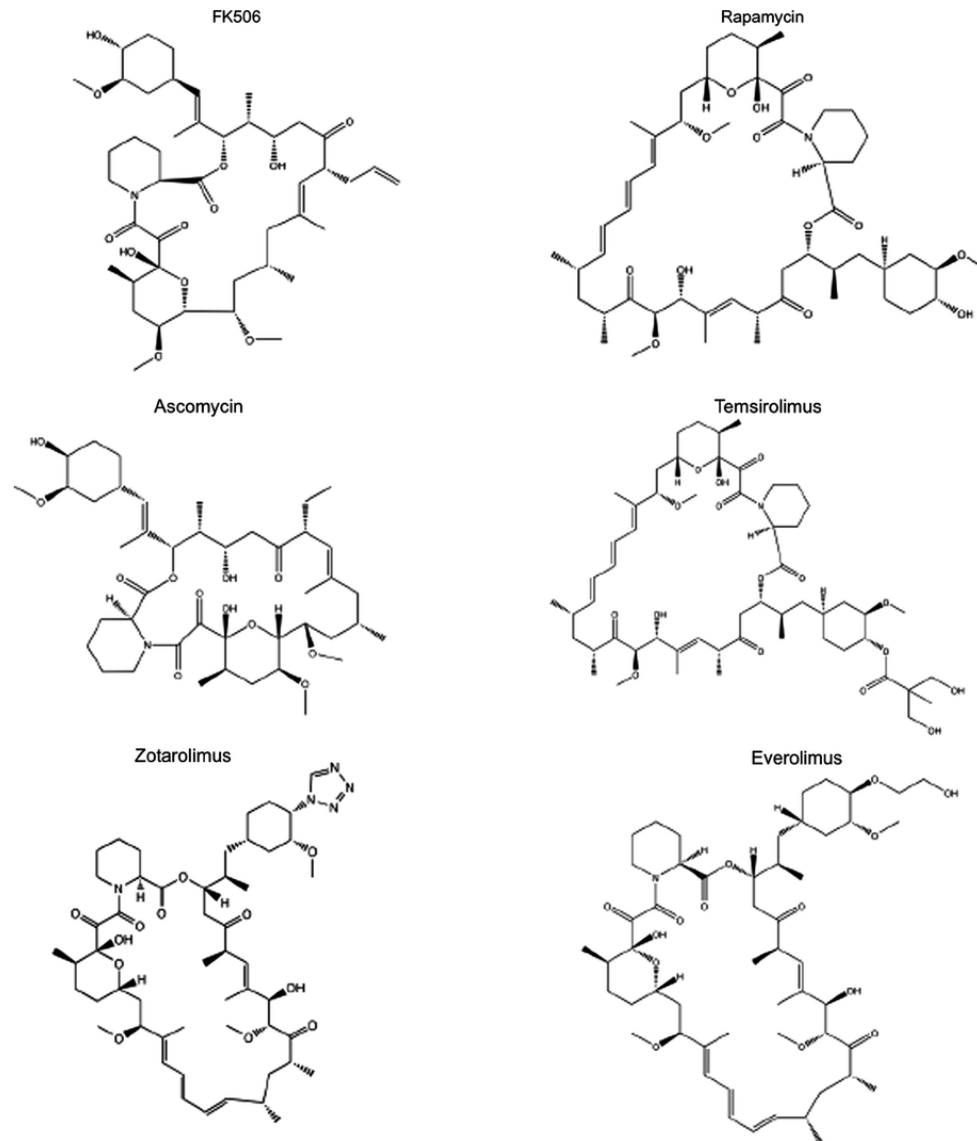
MGVQVETISPGDG  
RTFPKRGQTCVVH  
YTGMLDGGKKFDS  
SRDRNKPFKFVLG  
KQEVIRGWEEGVA  
QMSVGQRAKLTISP  
DYAYGATGHPGIIP  
PNATLIFDVELLKE

# FKBPs are everywhere!

*FKBPs encoded in the human genome*



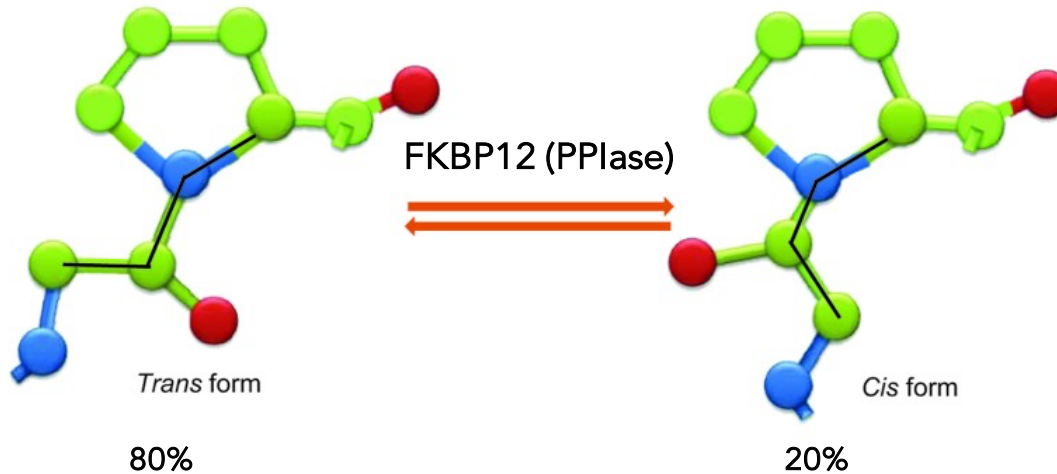
# Several drugs bind to FKBP12 with high affinity



anti-immune effects  
anti-tumor effects

# FKBP12 is a peptidyl-prolyl isomerase

Proline is highly constrained  
-  
Side chain is bonded to its secondary amine Nitrogen

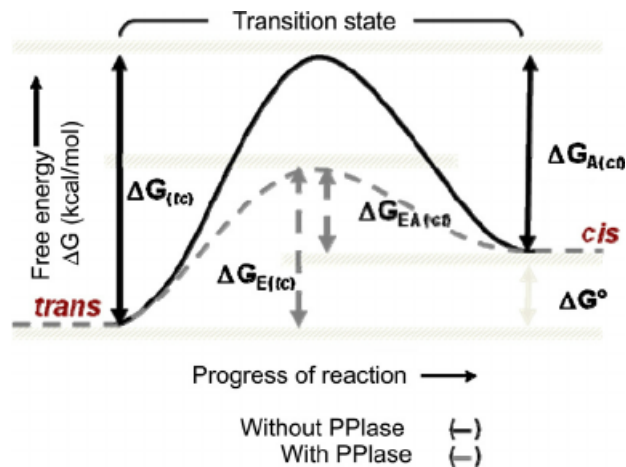
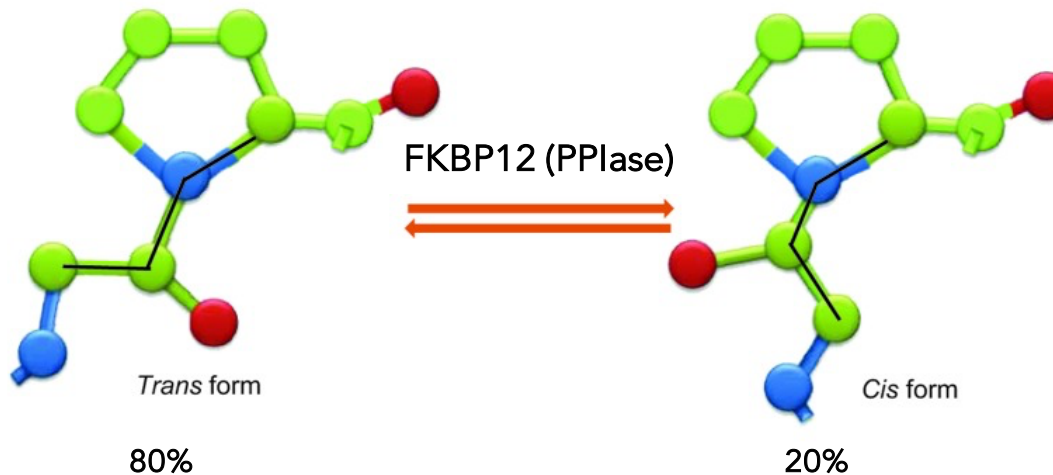




# FKBP12 is a peptidyl-prolyl isomerase

Proline is highly constrained

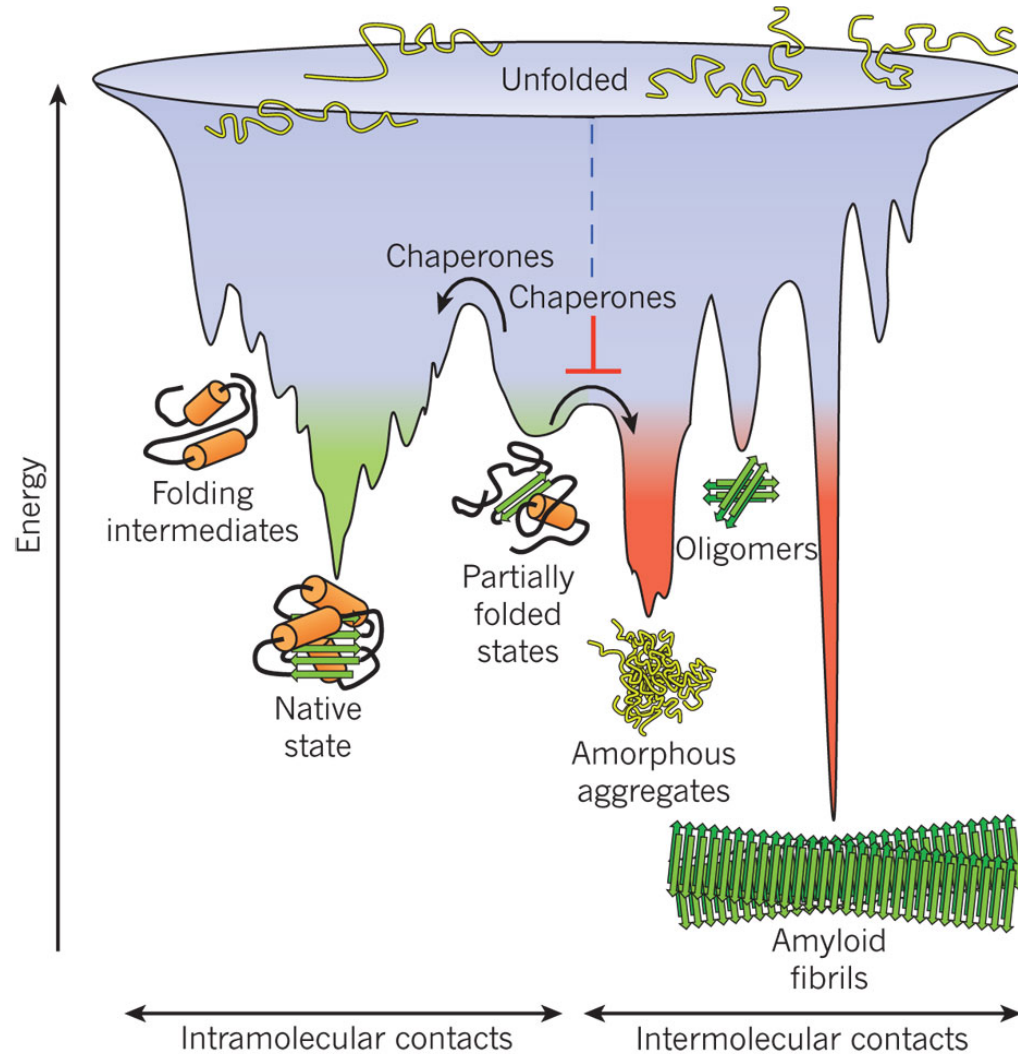
Side chain is bonded to its secondary amine Nitrogen



$E_a \sim 20$  kcal/mol

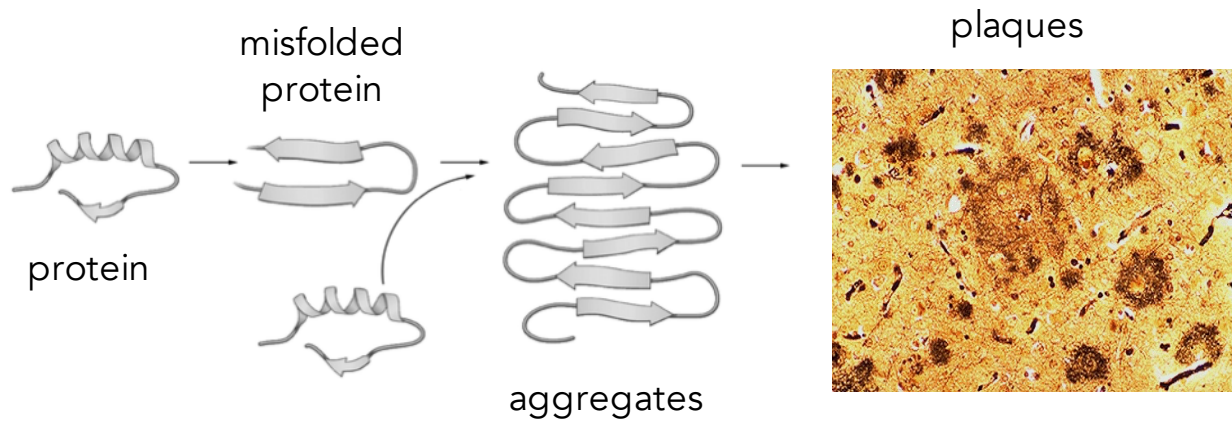
Enzyme catalyzes rate acceleration of  $10^6$  fold over non-enzymatic cis-trans isomerization

# FKBP12 is a molecular chaperone

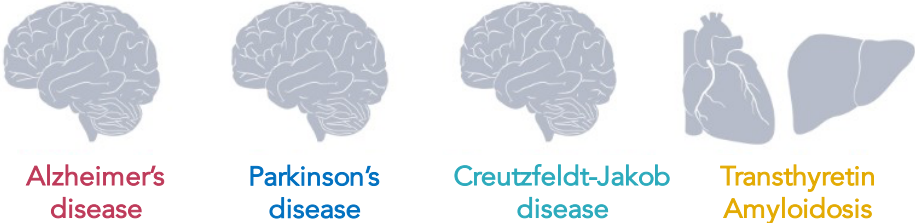
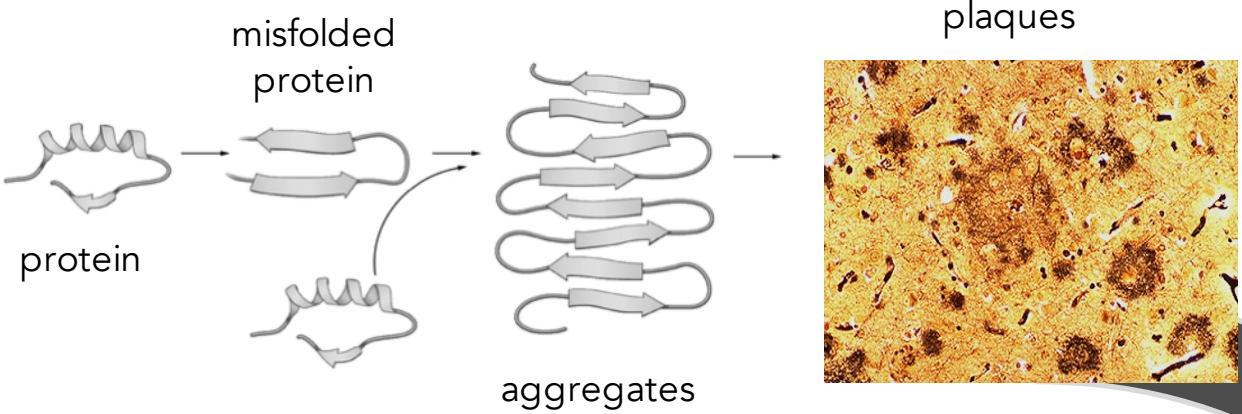




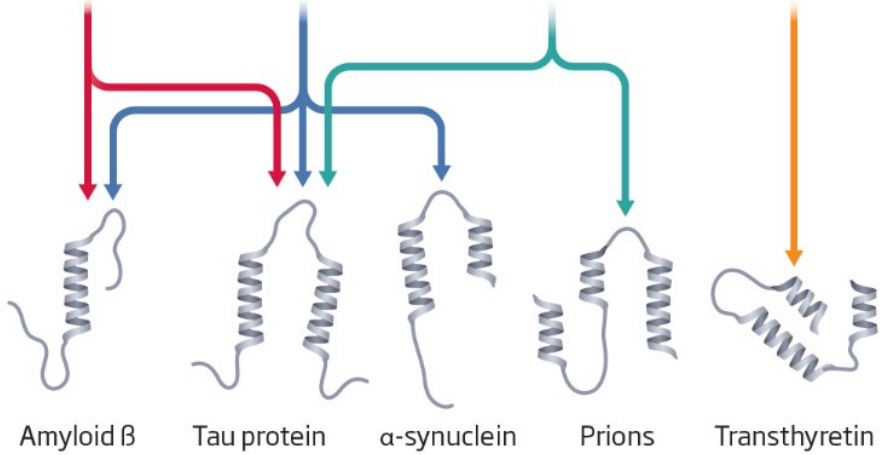
# FKBPs in diseases of protein folding



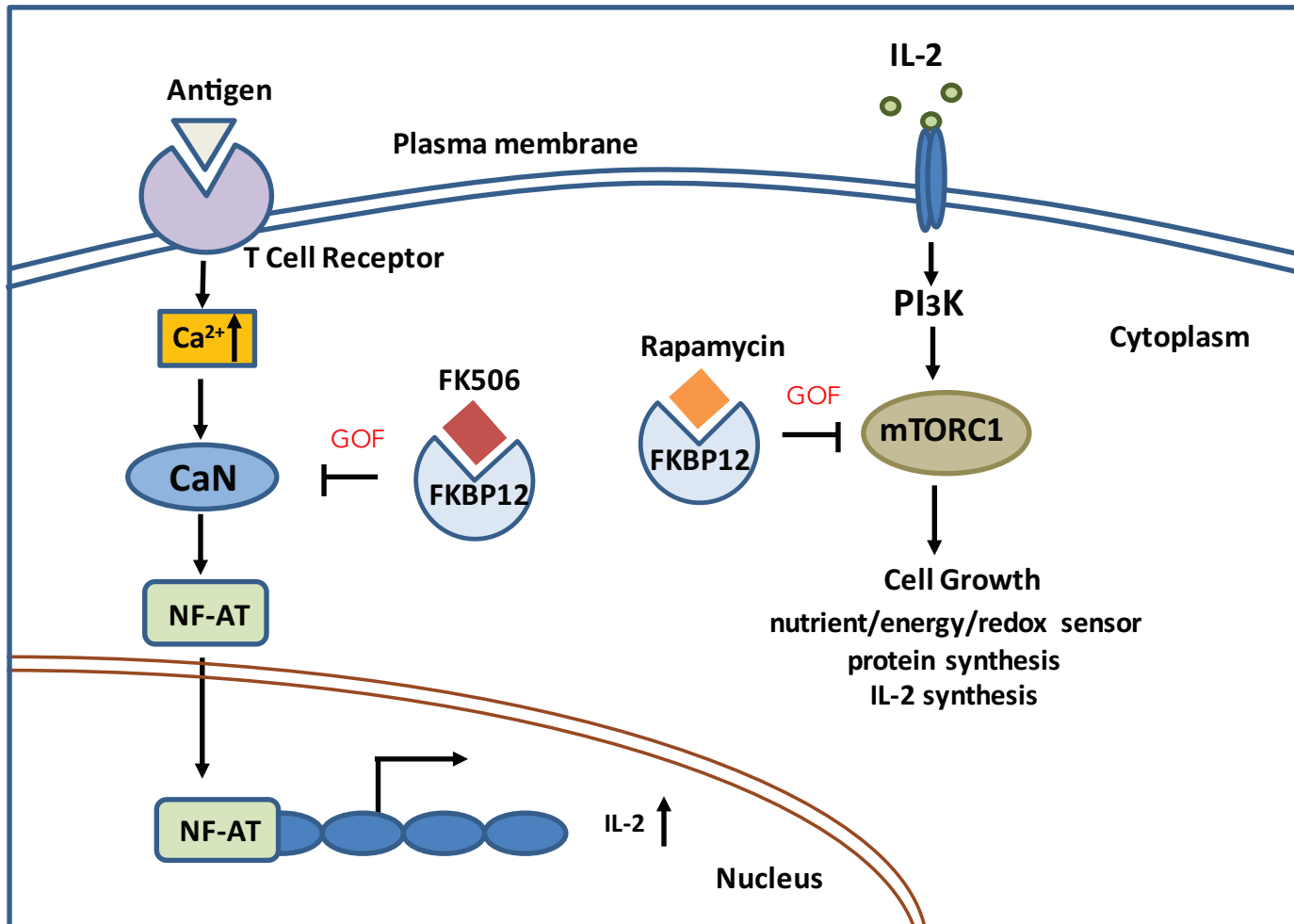
# FKBPs in diseases of protein folding



higher FKBP52 = lower Tau aggregation

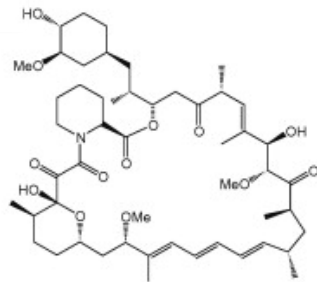


# FKBP12 'gains a function' to inhibit T-cell activity

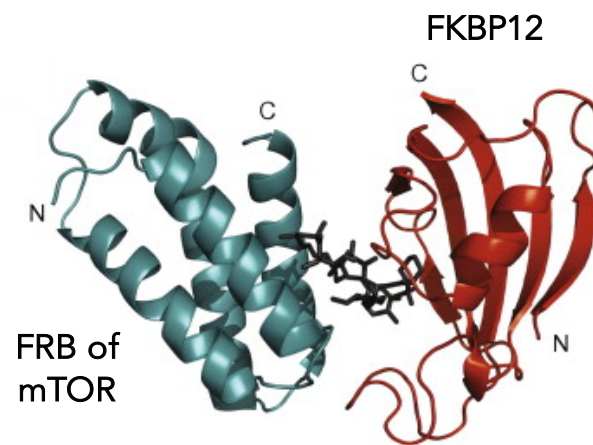


# FKBP12 'ternary' complexes

Rapamycin and mTOR



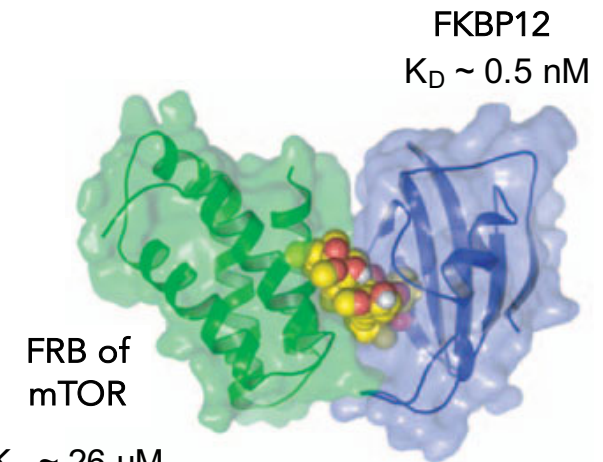
rapamycin



FRB of  
mTOR

FKBP12

$K_D \sim 26 \mu\text{M}$



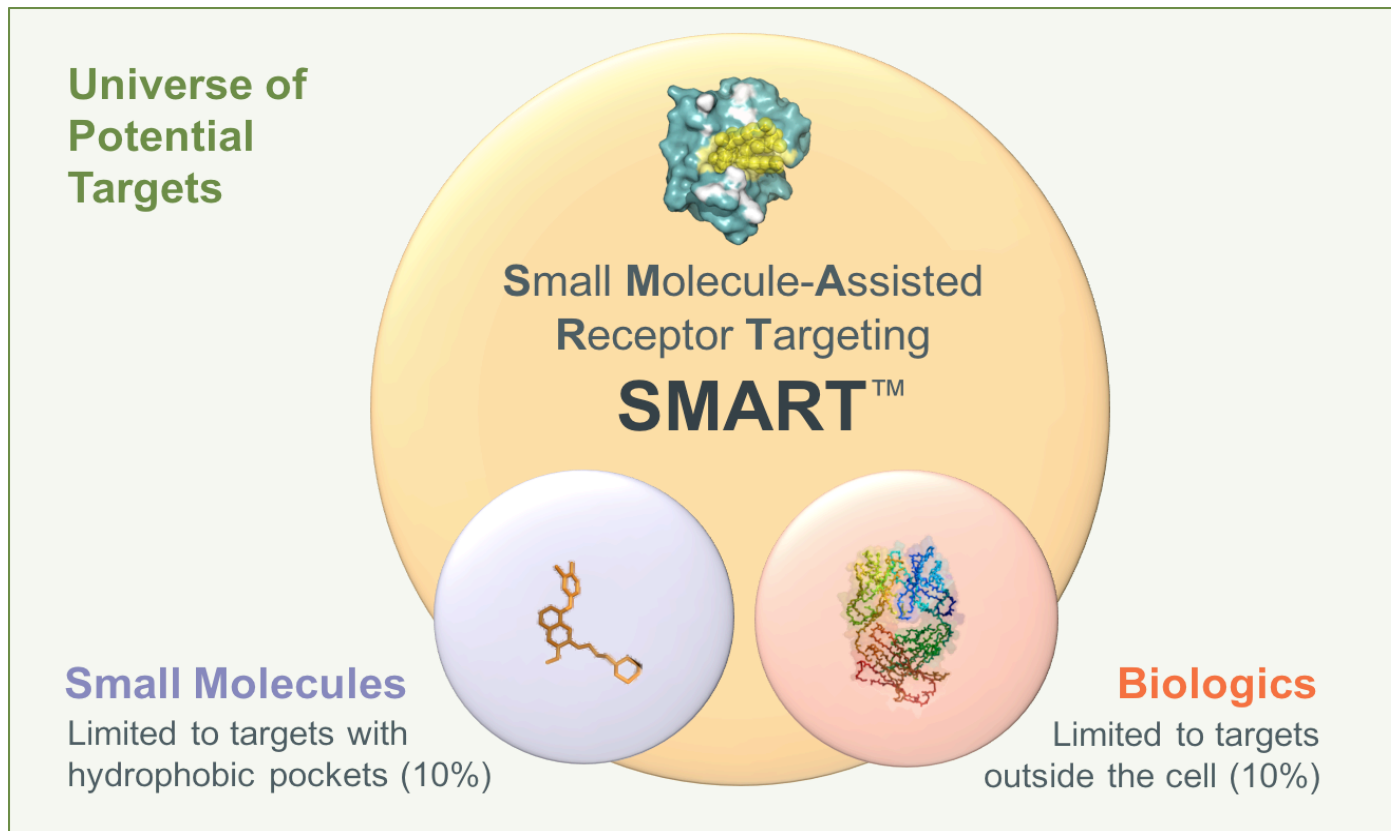
FKBP12  
 $K_D \sim 0.5 \text{ nM}$

$K_D \sim 12 \text{ nM}$

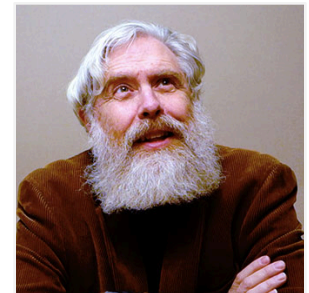
Choi *et al.* Science, 273, 239-42 (1996)  
Banaszynski, Liu, and Wandless, J. Am. Chem Soc., 127, 4815-21 (2005)

PDB entry:  
1NSG

# Drugging the 'undruggable' through GOF



Greg Verdine, Harvard



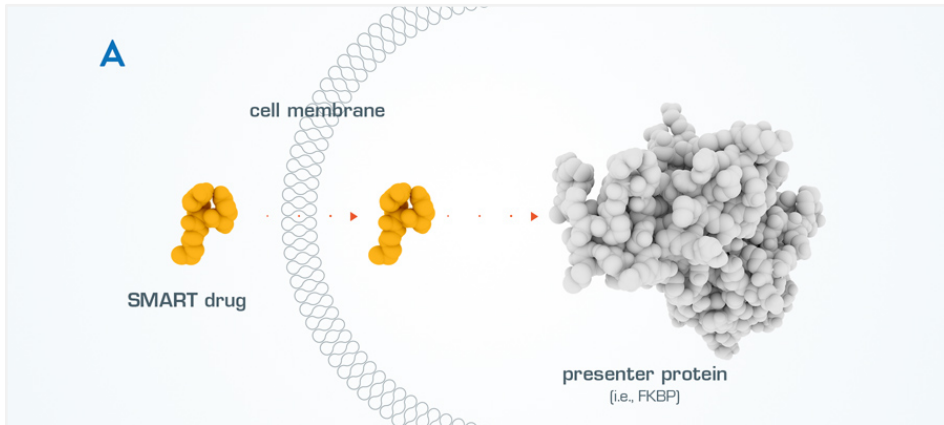
George Church, Harvard



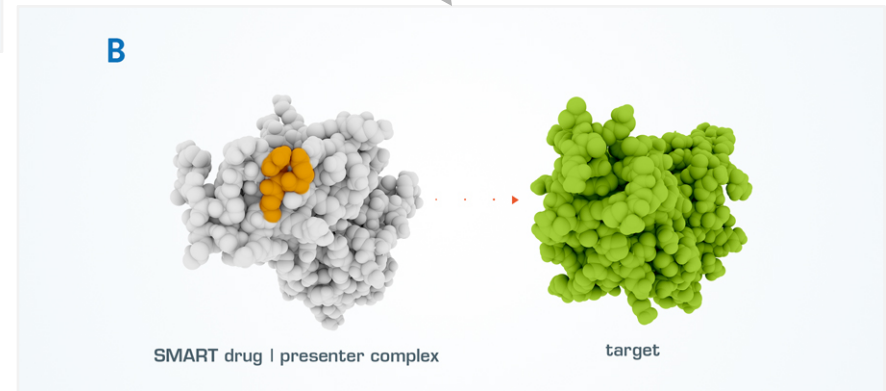
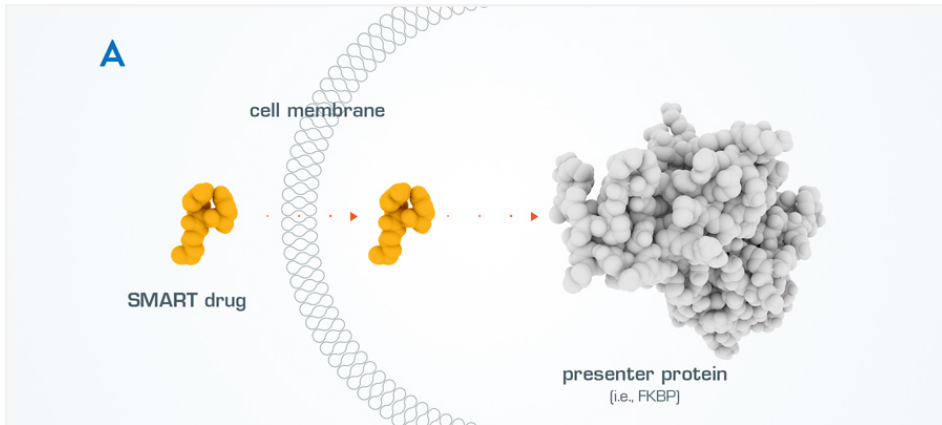
Jim Wells, UCSF



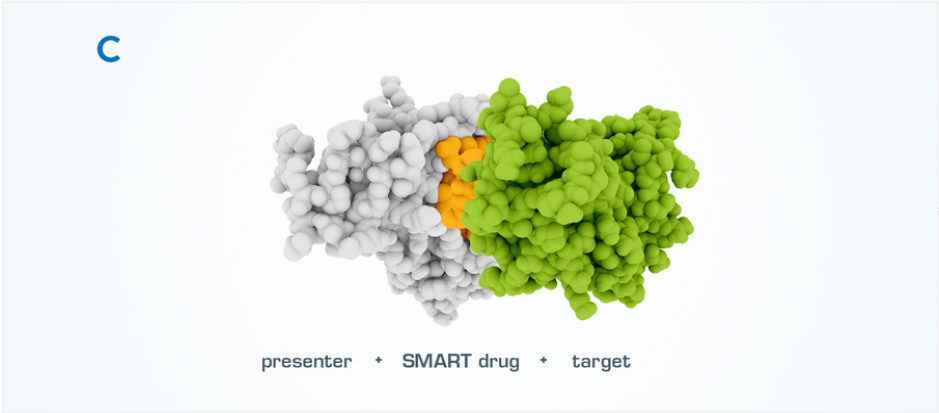
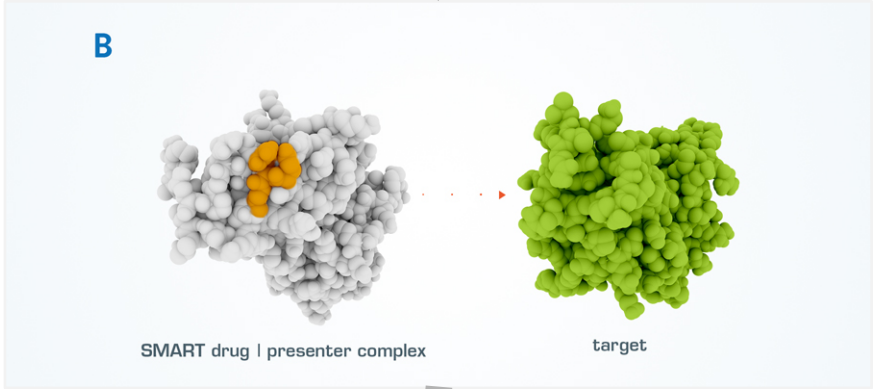
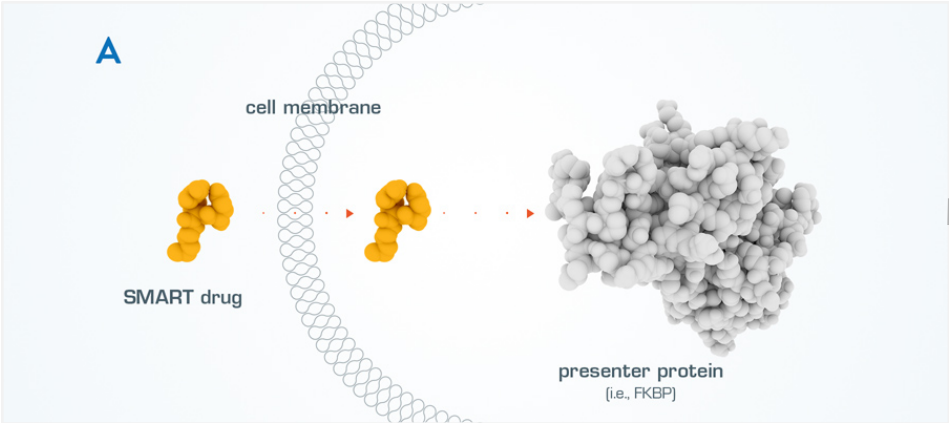
# SMART™ – small molecule-assisted receptor targeting



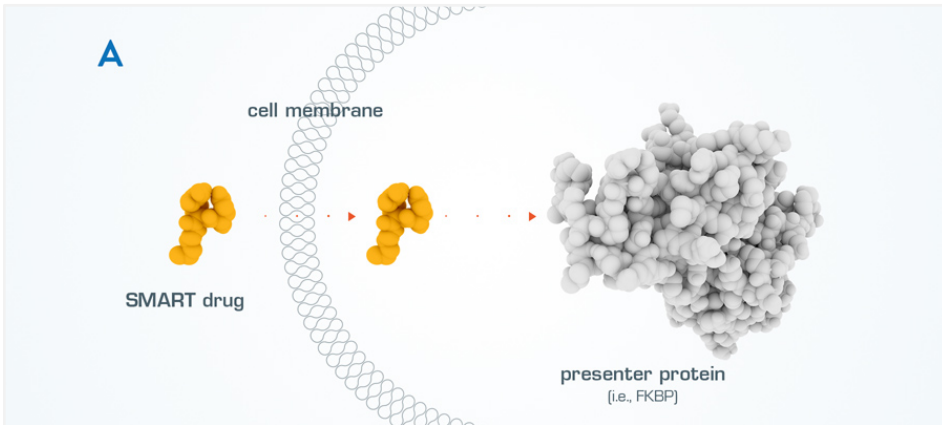
# SMART™ – small molecule-assisted receptor targeting



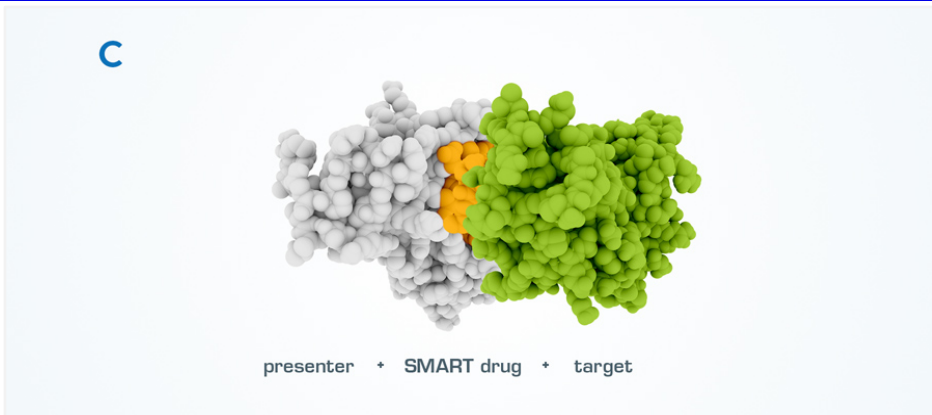
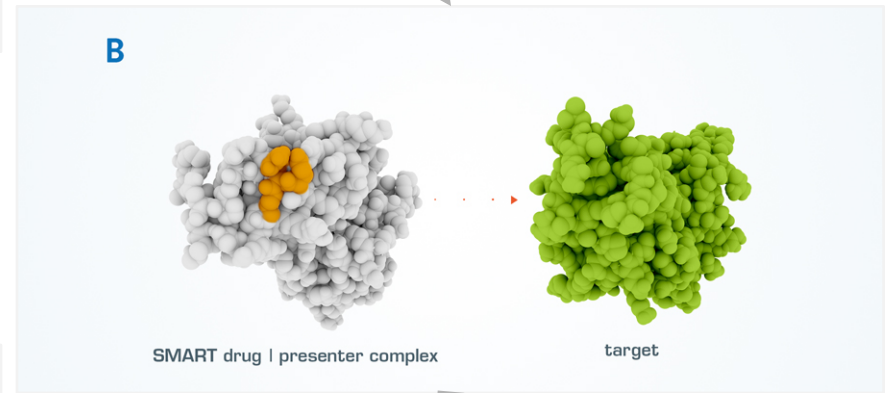
# SMART™ – small molecule-assisted receptor targeting



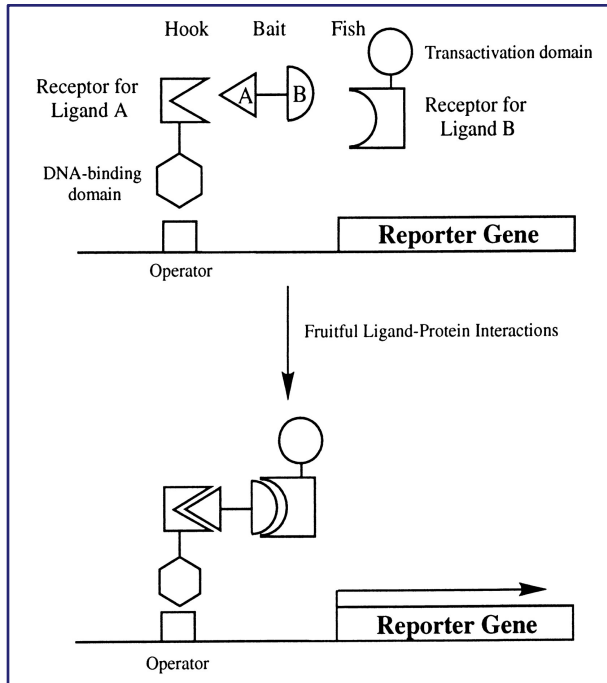
# SMART™ – small molecule-assisted receptor targeting



Novel molecules that you find during this class may serve as new starting points for this concept, providing new molecular interfaces with FKBP12 that can be used to engage a new proteins through design or screening



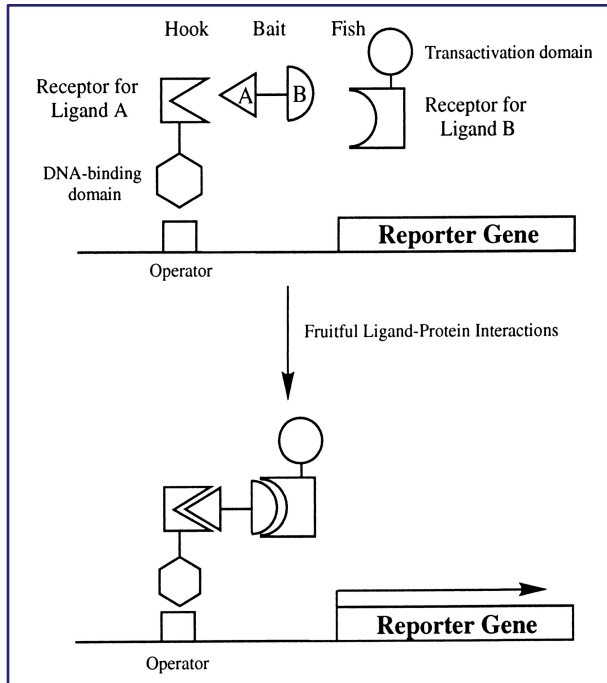
# FKBP12 as a tool for biological engineering - preview



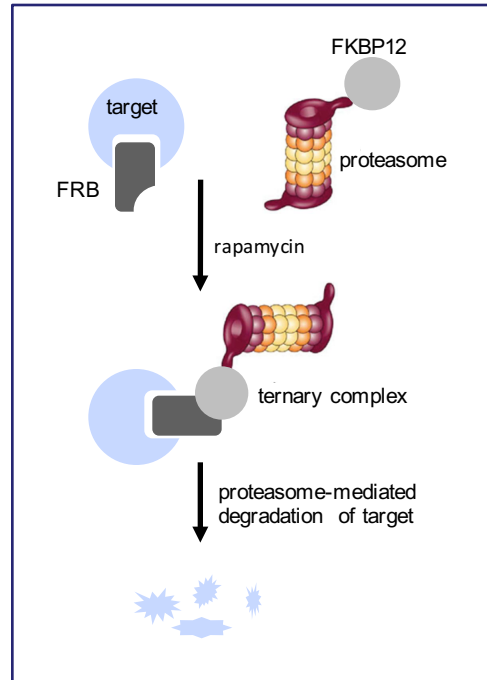
controlling  
transcription

proximity induction strategies

# FKBP12 as a tool for biological engineering - preview



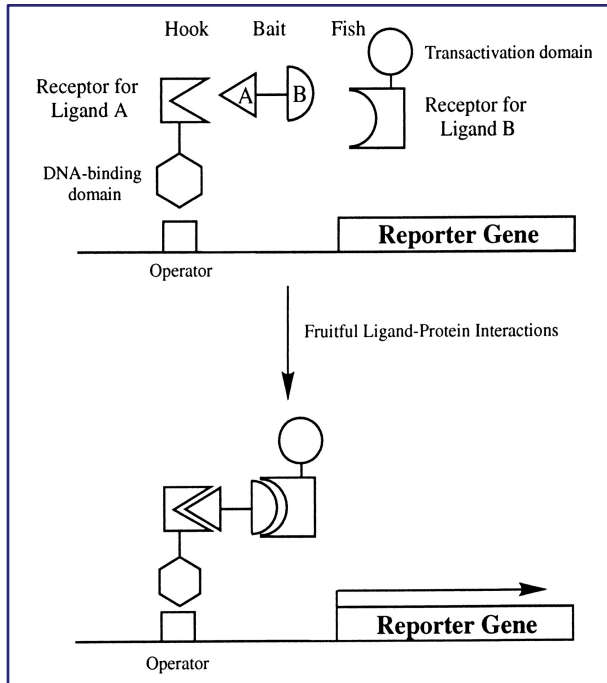
controlling  
transcription



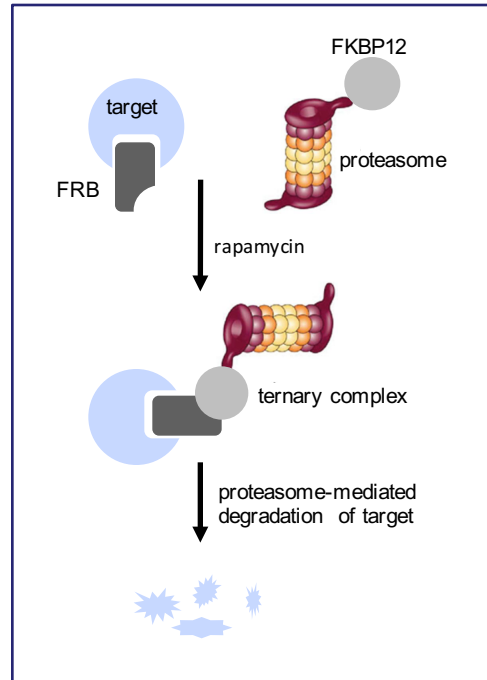
inducing  
protein degradation

proximity induction strategies

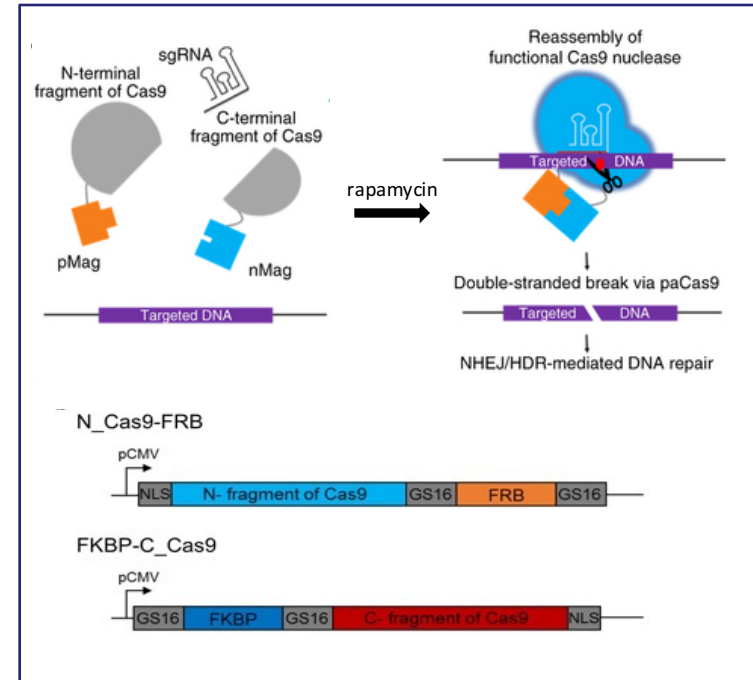
# FKBP12 as a tool for biological engineering - preview



controlling transcription



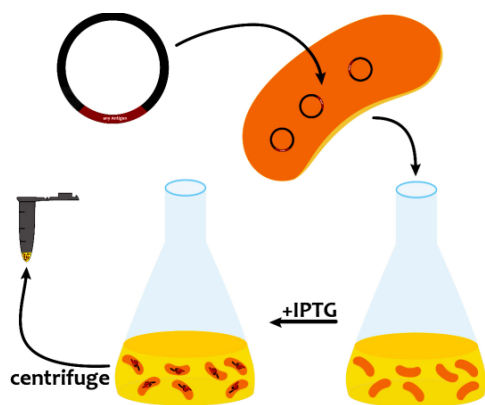
inducing protein degradation



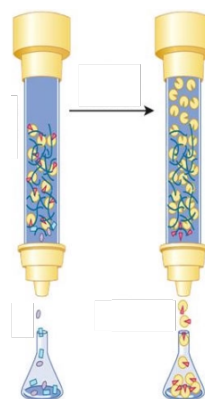
induced genome editing

proximity induction strategies

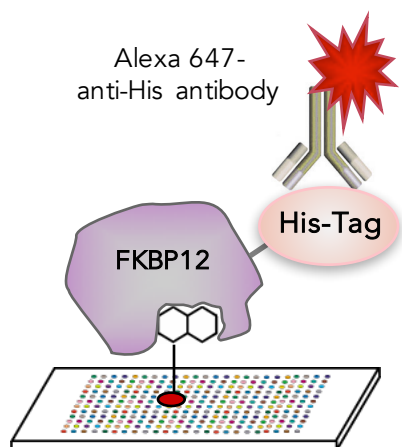
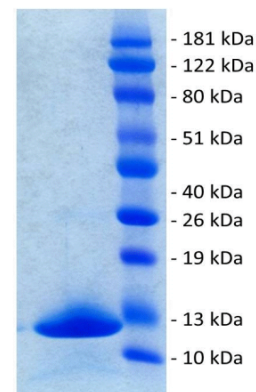
# Reminder - our path to probe discovery



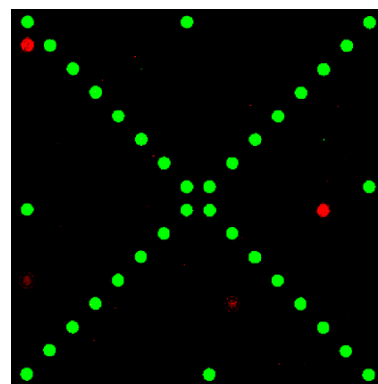
**overexpress FKBP12**  
lab day 1



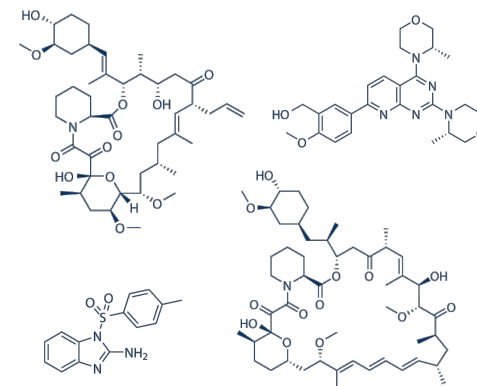
**purify and characterize FKBP12**  
lab days 2 and 3



**SMM screen**  
lab day 4



**scan images and analyze data**  
lab days 5 and 6



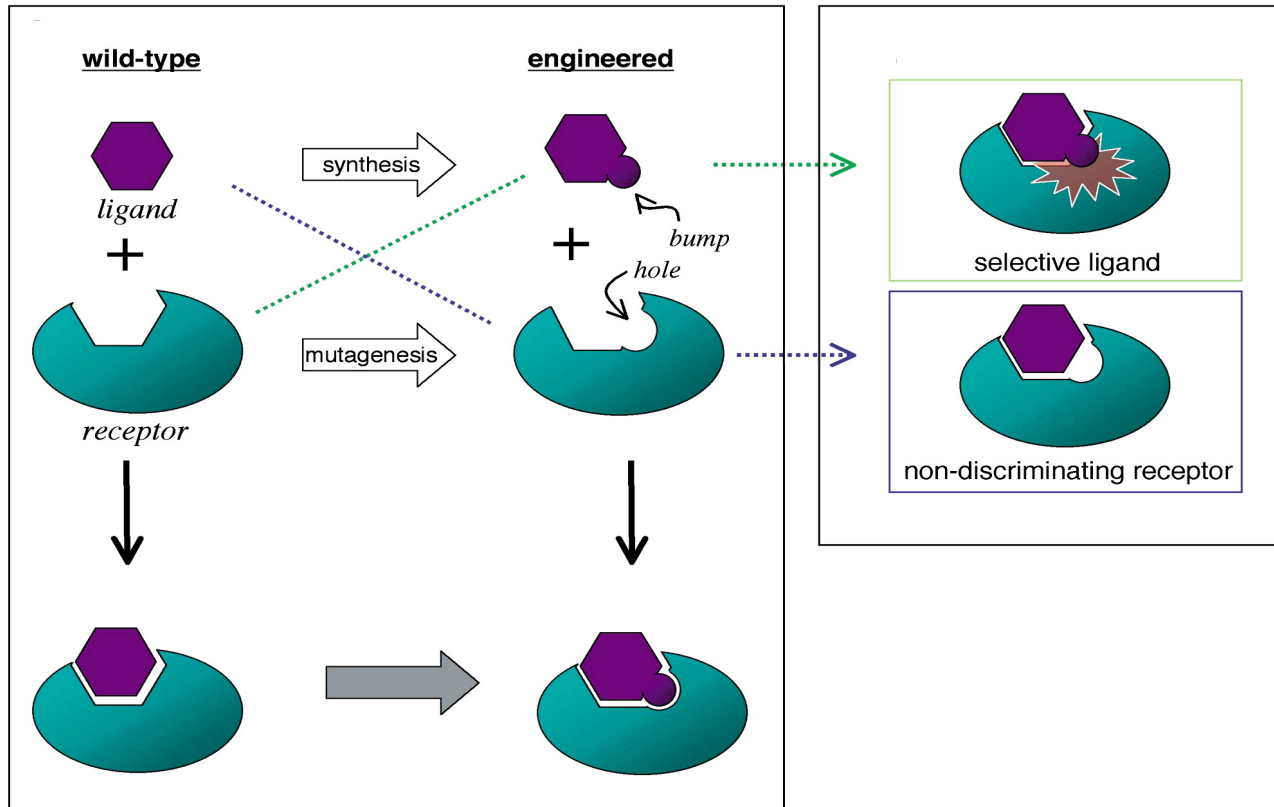
**compare hit lists for teams**  
lab day 7



# Our path to probe discovery - lectures

2/14/17	Lecture 1	Intro to chemical biology: small molecules, probes, and screens
2/16/17	Lecture 2	For the love of proteins: FKBP12 and immunophilins
2/21/17	No Lecture	
2/23/17	Lecture 3	Small-molecule microarrays
2/28/17	Lecture 4	Analyzing SMM data sets ( <a href="#">Shelby Doyle</a> )
3/2/17	Lecture 5	Engineering fun with chemical probes
3/7/17	Lecture 6	Wrap up discussion: suggestions for how to report your findings

# FKBP12 as a tool for biological engineering - preview



orthogonal receptor ligand pairing  
aka 'bump-hole'