

Protein stabilisation: Design, Experiments and Assessment (ProteSta)

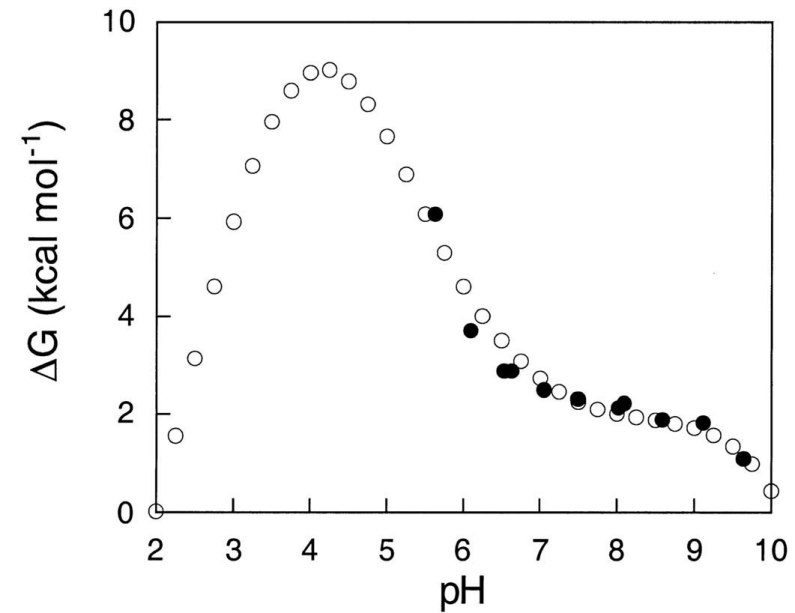
ALS1 (BIFI-LACRIMA) 3rd-7th July 2023, Zaragoza, Spain

- ▶ Protein stabilization by formulation: some rational
- ▶ Examples
- ▶ Protein stabilization by mutagenesis: some rational
- ▶ Examples
- ▶ Three-state proteins

► Protein stabilization by formulation: some rational

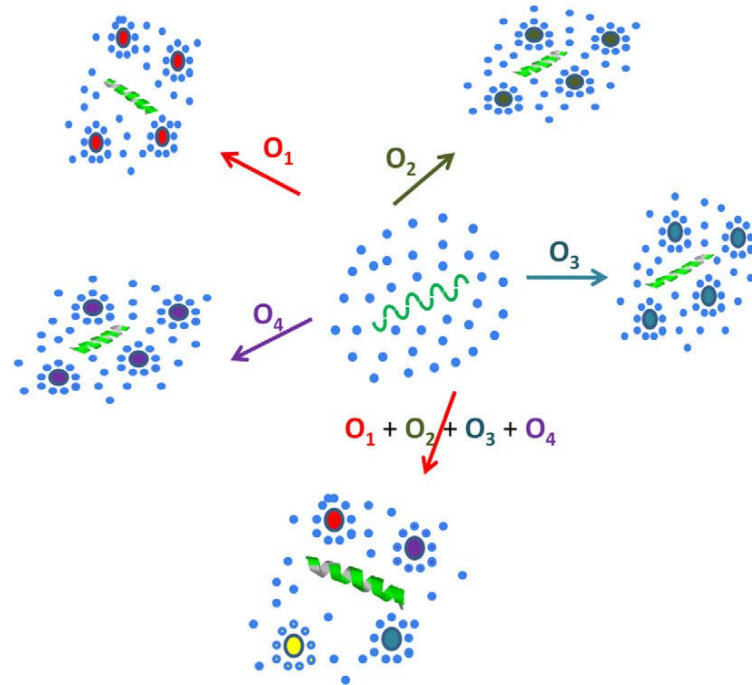
Some chemicals you may considered

- Buffers
- Salts (NaCl)
- Amino acids (histidine, glycine, arginine)
- Sugars (sucrose, manitol, sorbitol)
- Polymers (dextran, PEG)
- Specific binders to the native state: ligand binding



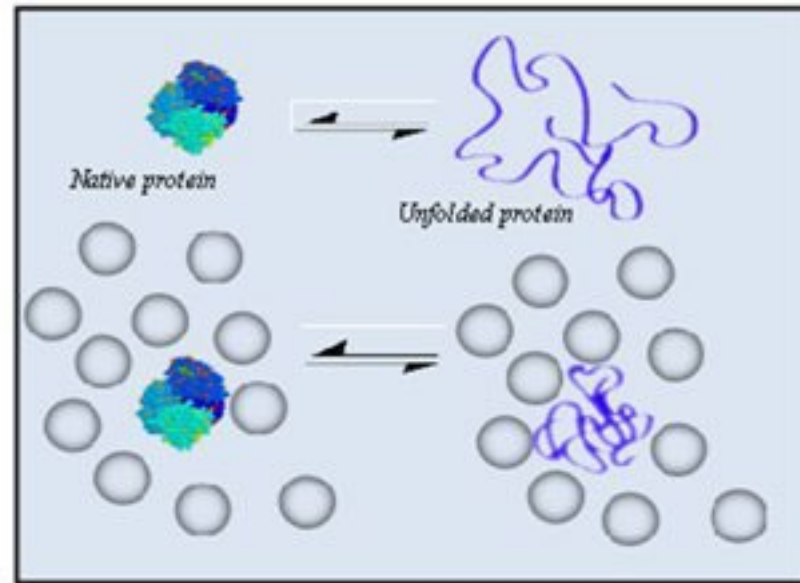
Salt-induced stabilization of apoflavodoxin at neutral pH is mediated through cation-specific effects. Maldonado et al. Protein Science 11: 1260-1273

Protein stabilization by reducing water availability



Role of naturally occurring osmolytes in protein folding and stability
R. Kumar. Archives of Biochemistry and Biophysics 491 (2009) 1–6

Protein stabilization by crowding the solution



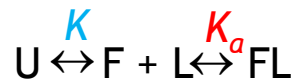
► Protein stabilization by formulation: some rational

Protein stabilization by ligand binding



$$K = \frac{[F]}{[U]}$$

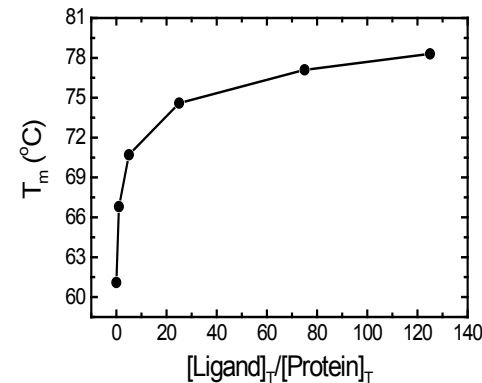
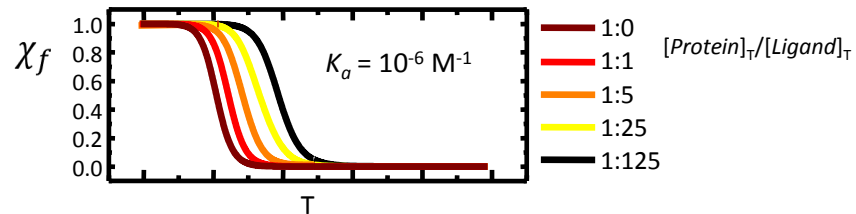
$$\Delta G = -RT \ln K$$



$$K_{lig} = \frac{[F] + [FL]}{[U]}$$

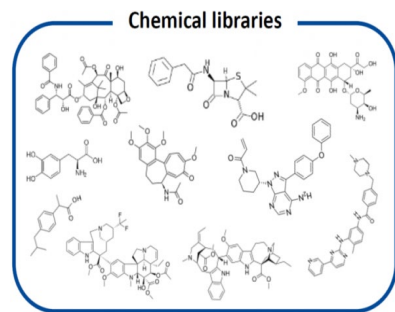
$$K_a = \frac{[FL]}{[F] \times [L]}$$

$$\Delta G_{lig} = \Delta G - RT \ln(1 + K_a \times [L])$$



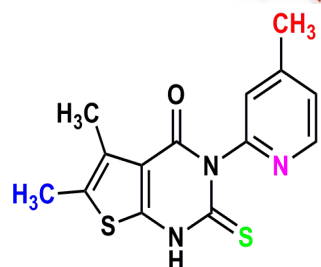
Protein stabilization by ligand binding: application in drug discovery

204 M known chemicals

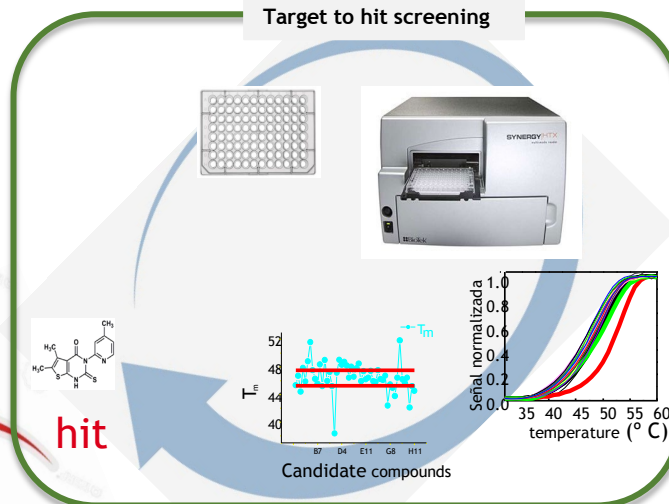


+
Target

Hit to lead medicinal chemistry



Target to hit screening



Testing in animal models



Improving and more preclinical



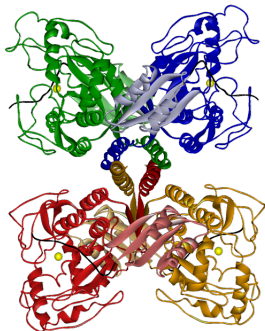
Laboratorio Avanzado de Cribado e Interacciones Moleculares de Aragón



Year	Target protein	Type of compound	Application	Patent
2008	PAH	pharmacological chaperons	Phnylketonuria (PKU)	X
2009	Fld	antimicrobials	<i>Helicobacter pylori</i>	X
2009	Stem cells	apoptotic cell death specific of ESC or EF	Cell therapy	
2012	Aβ(1-42)	aggregation inhibitors	Alzheimer disease	X
2013	NS3 protease	antivirals	Hepatitis C	
2016	hIAPP	aggregation inhibitors	Type II Diabetes	
2016	PEPCK-C	Enzyme inhibitors	Isoform phenotyping	
2017	IDP	antitumoral	Pancreatic cancer	
2017	FNR	antimicrobial	<i>Xanthomonas citri</i>	
2018	α-syn	aggregation inhibitors	Parkinson disease	X
2018	FAD synthetase	antimicrobials	<i>C. ammoniagenes</i> <i>M. Tuberculosis</i> <i>S. pneumoniae</i>	
2019	AIF	apoptosis regulation	Apoptosis regulation	

Phenylketonuria

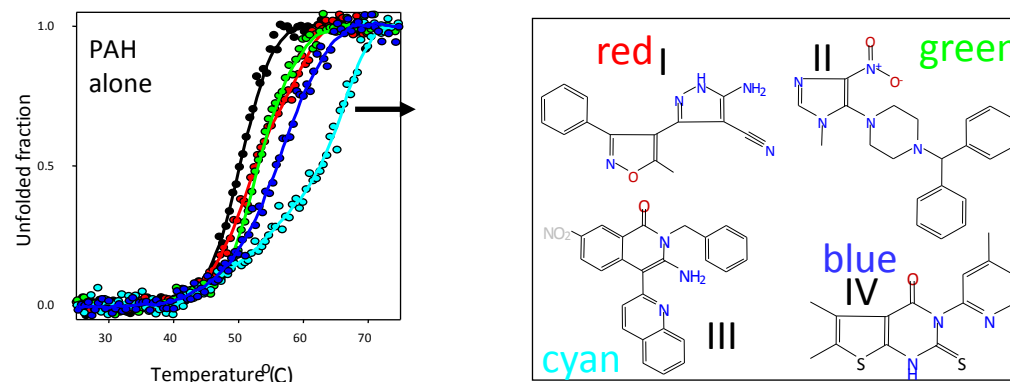
(1:15000 in Europe)



Destabilizing mutations in PAH
2208 variants known
(as of May 22, 2023)

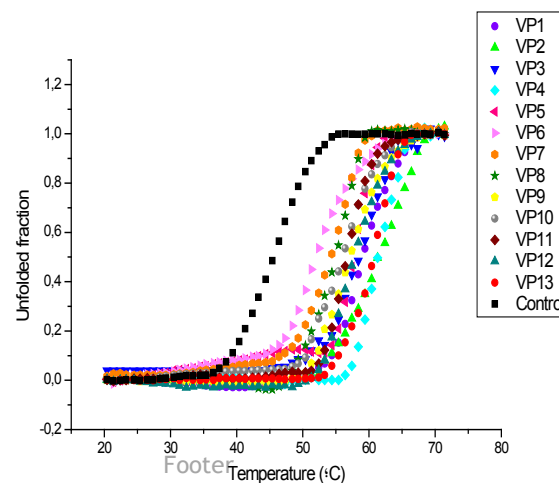
Pharmacological chaperons

1500 compounds screened → 4 stabilizing compounds found



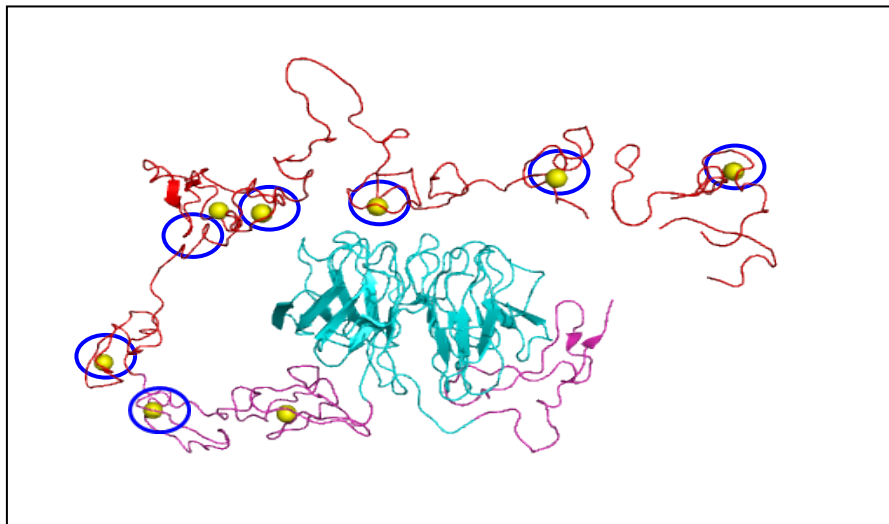
Identification of pharmacological chaperones as potential therapeutic agents to treat phenylketonuria
Pey et al. 2008. Journal of clinical investigation 118: 2858-2867.

10000 compounds screened → more stabilizing compounds found

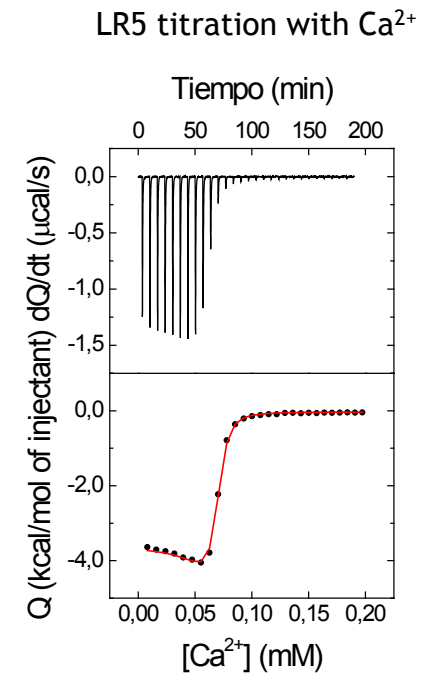
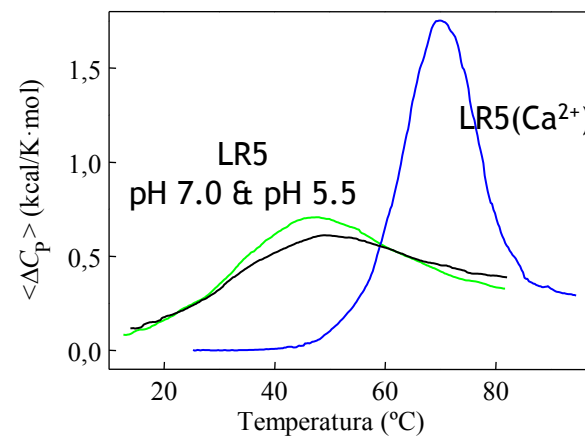


Protein stabilization by ligand binding: role in cellular mechanisms

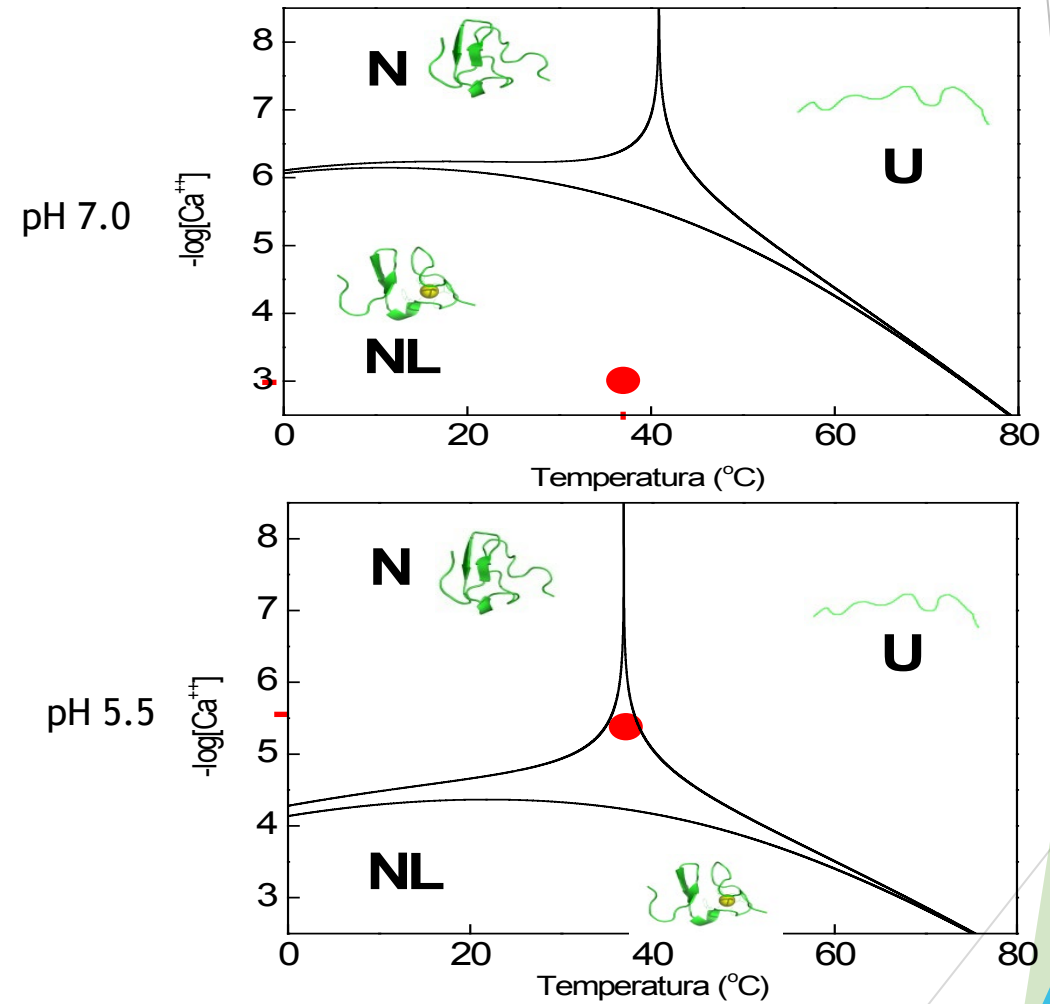
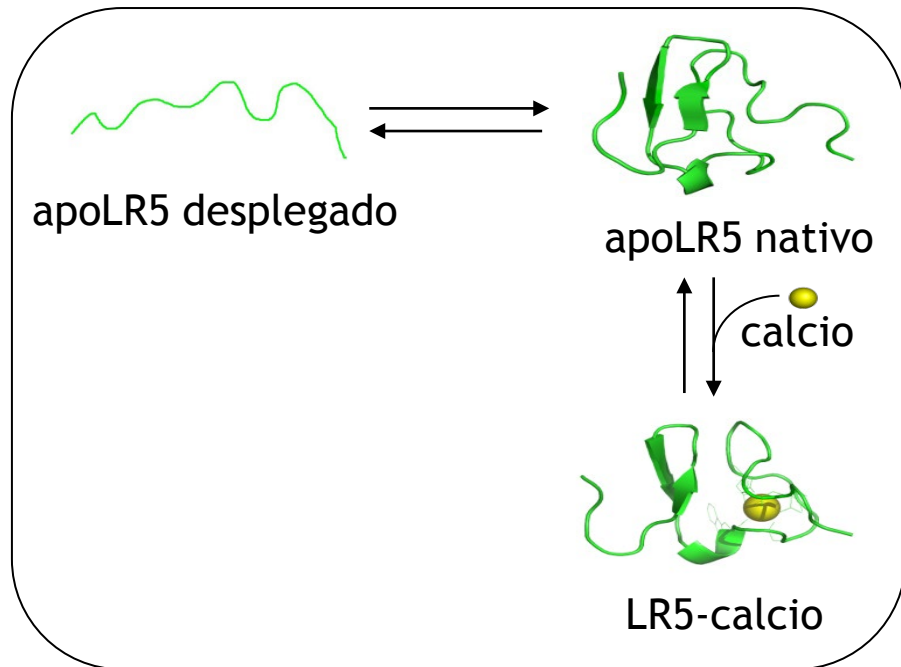
LDL delivery by LDL receptor



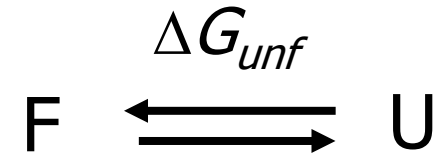
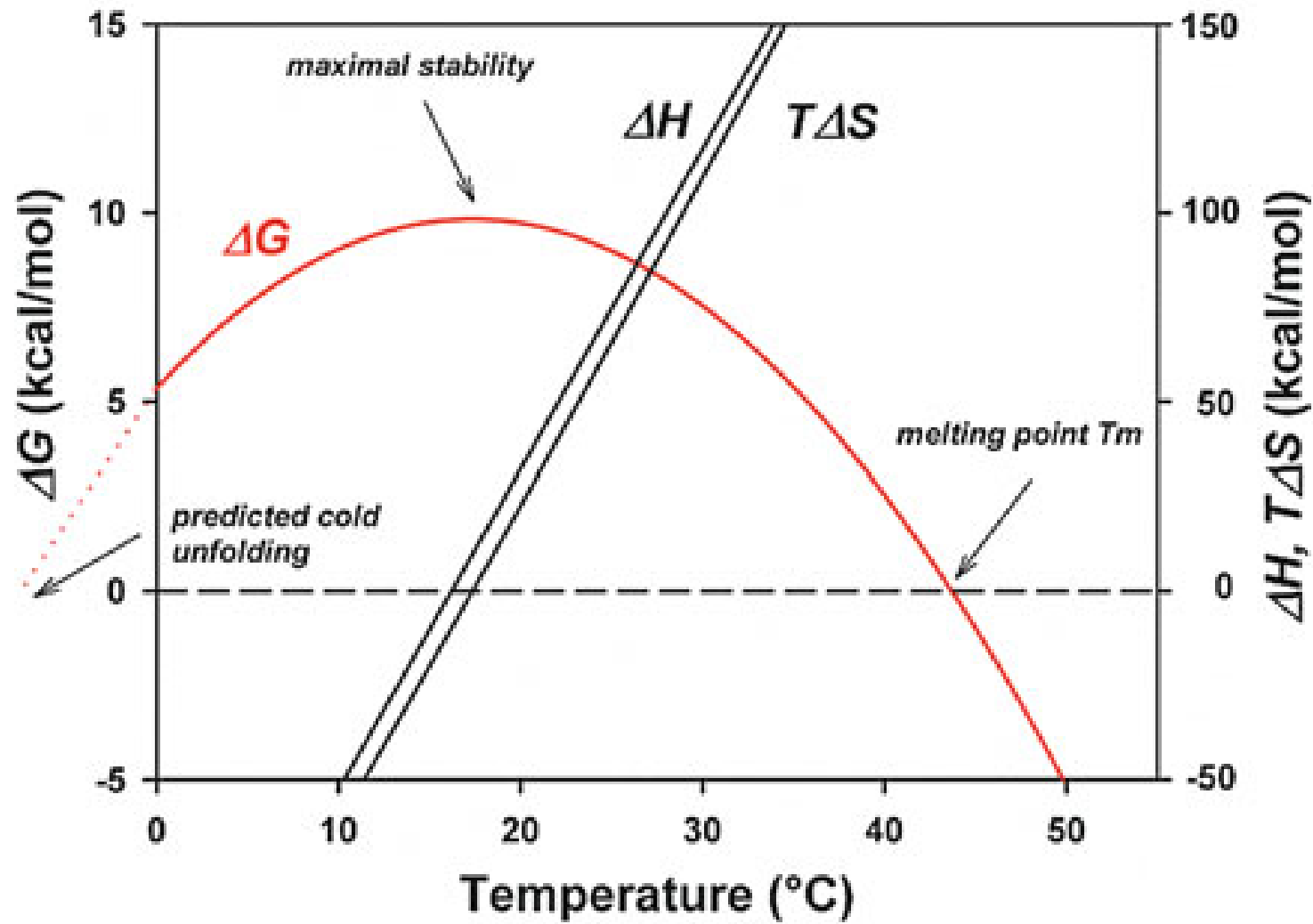
Arias-Moreno X, Velazquez-Campoy A, Rodríguez JC, Pocoví M, Sancho J. Mechanism of low density lipoprotein (LDL) release in the endosome: implications of the stability and Ca^{2+} affinity of the fifth binding module of the LDL receptor. J Biol Chem. 2008 Aug 15;283(33):22670-9.



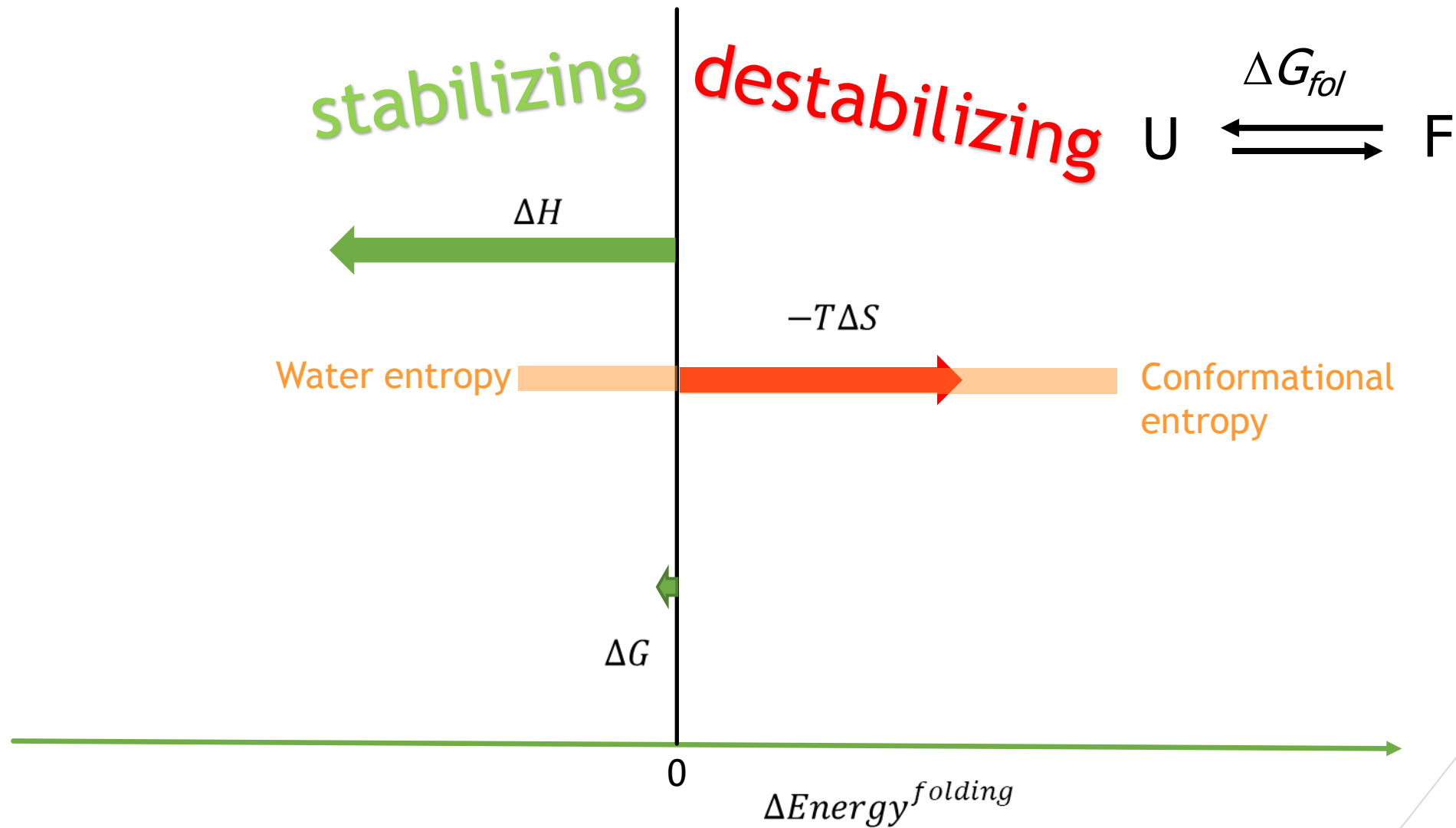
► Examples



► Protein stabilization by mutagenesis: some rational

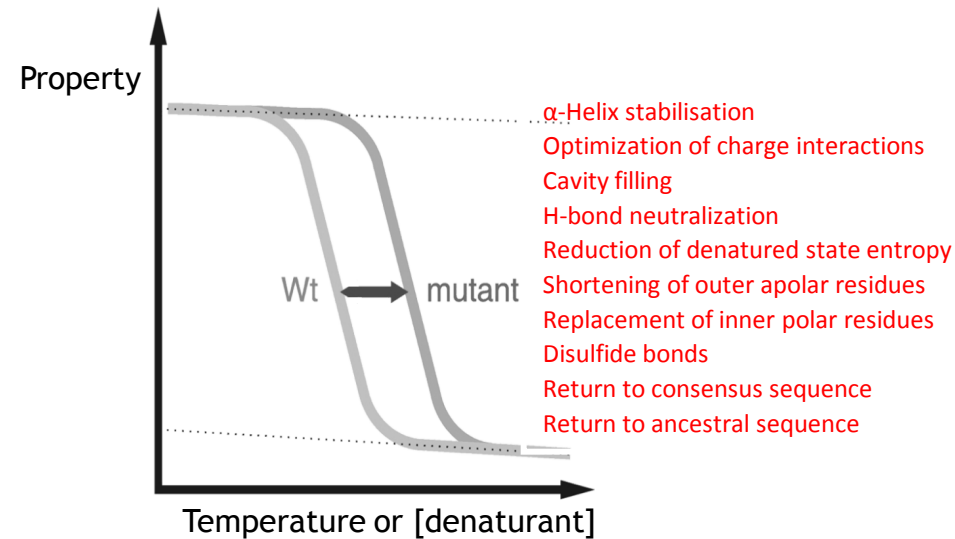
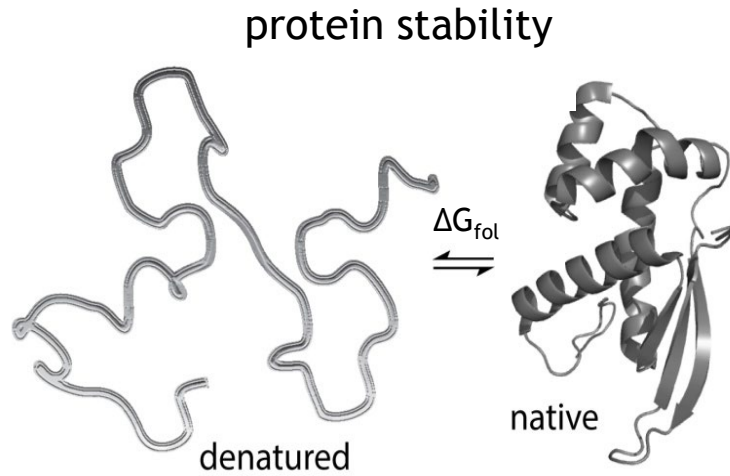


► Protein stabilization by mutagenesis: some rational



► Protein stabilization by mutagenesis: some rational

To get inspiration you may think of ΔH , ΔS or evolution

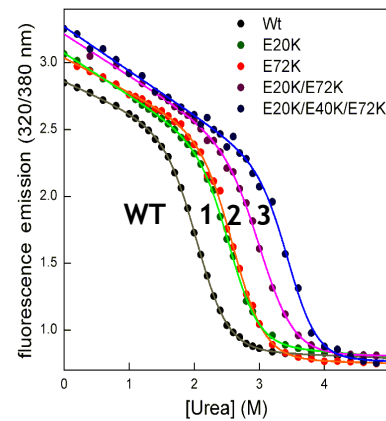
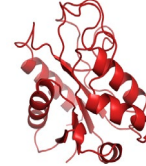


$$\Delta\Delta G_{\text{mut-wt}} = \Delta G_{\text{mut}} - \Delta G_{\text{wt}} > 0.5 \text{ kcal/mol}$$

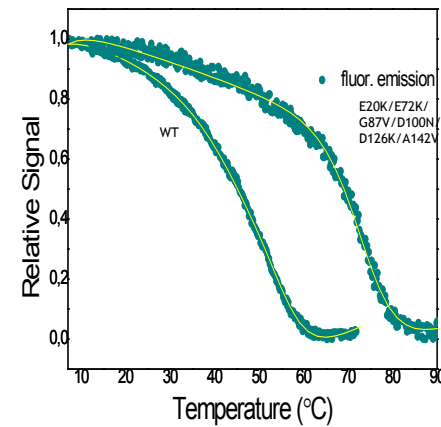
- Random substitution: 2.5 % stabilizing mutations
- Directed evolution; deep sequencing
- **Designed substitutions**
- **AI??**

The stabilizing effect tends to be accumulative

Anabaena PCC 7119 flavodoxin

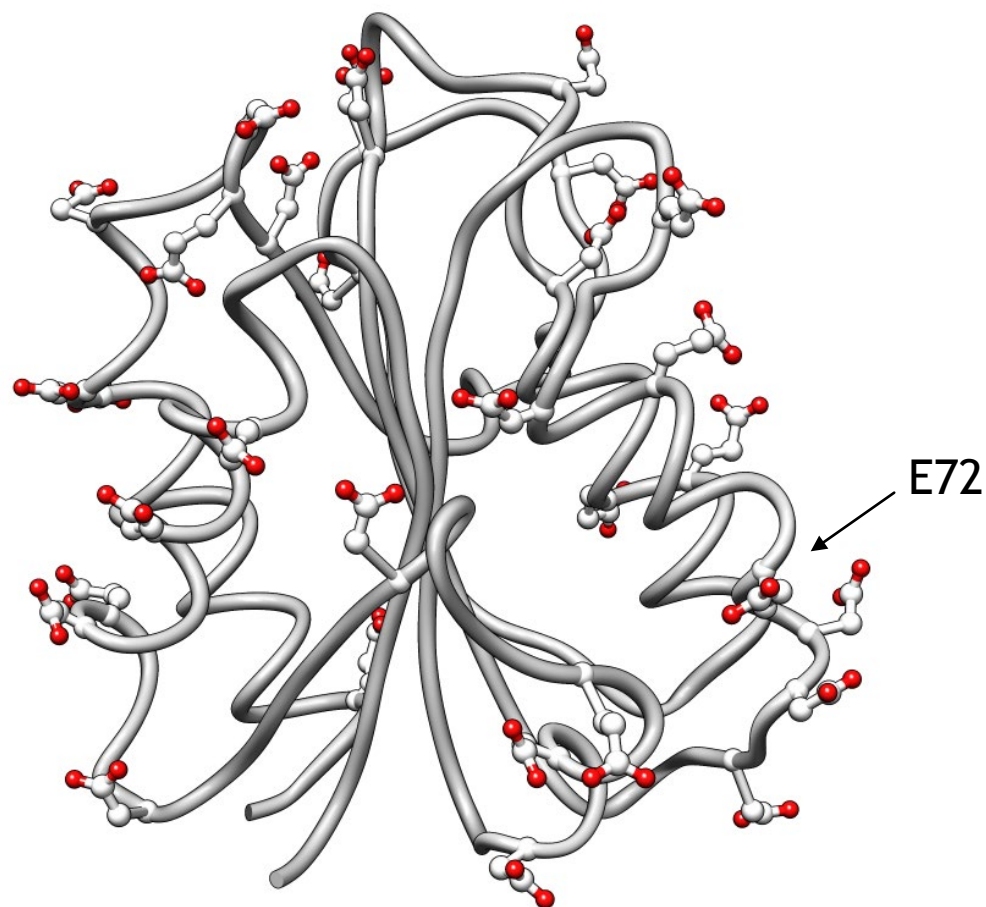


Bueno et al.
J. Mol. Biol. (2006) 358:701-712

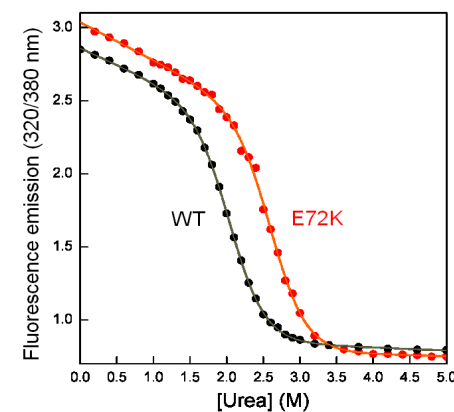
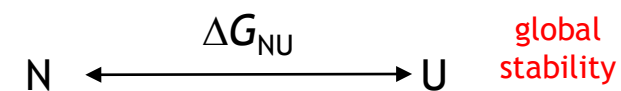


Lamazares et al.,
Sci Rep. 5:9129 (2015)

► Three-state proteins



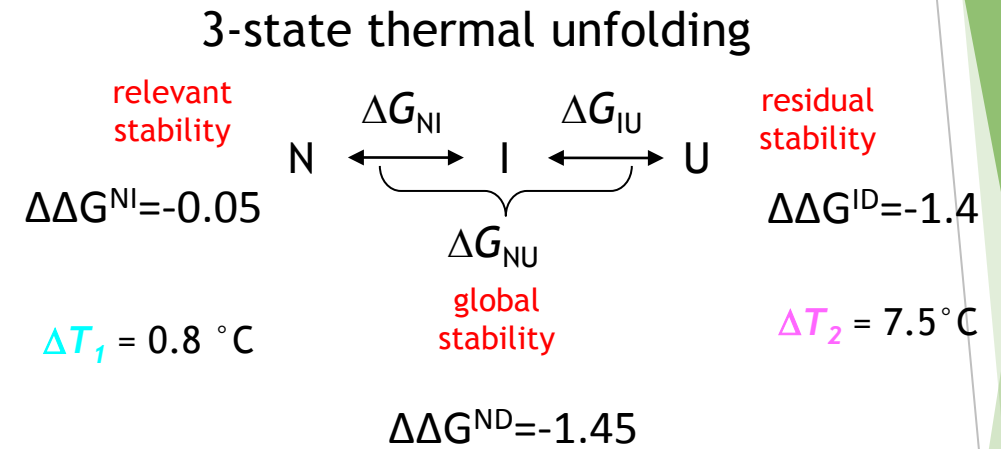
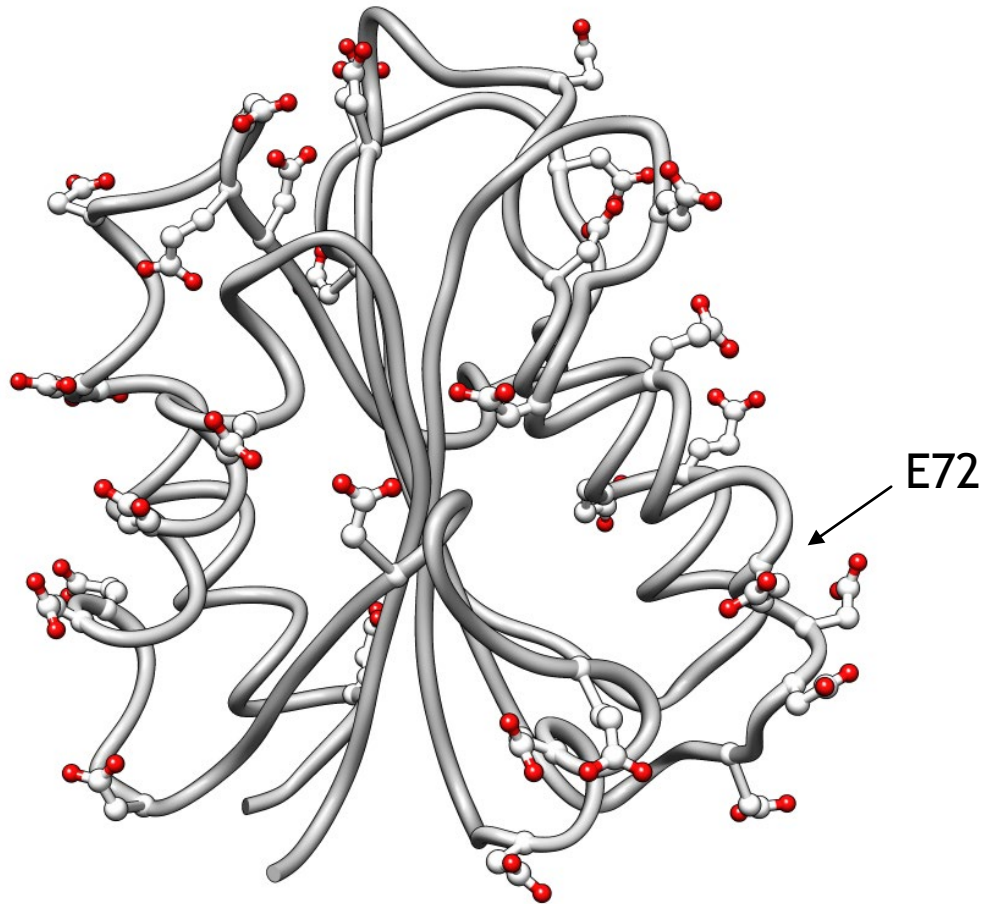
2-state chemical unfolding



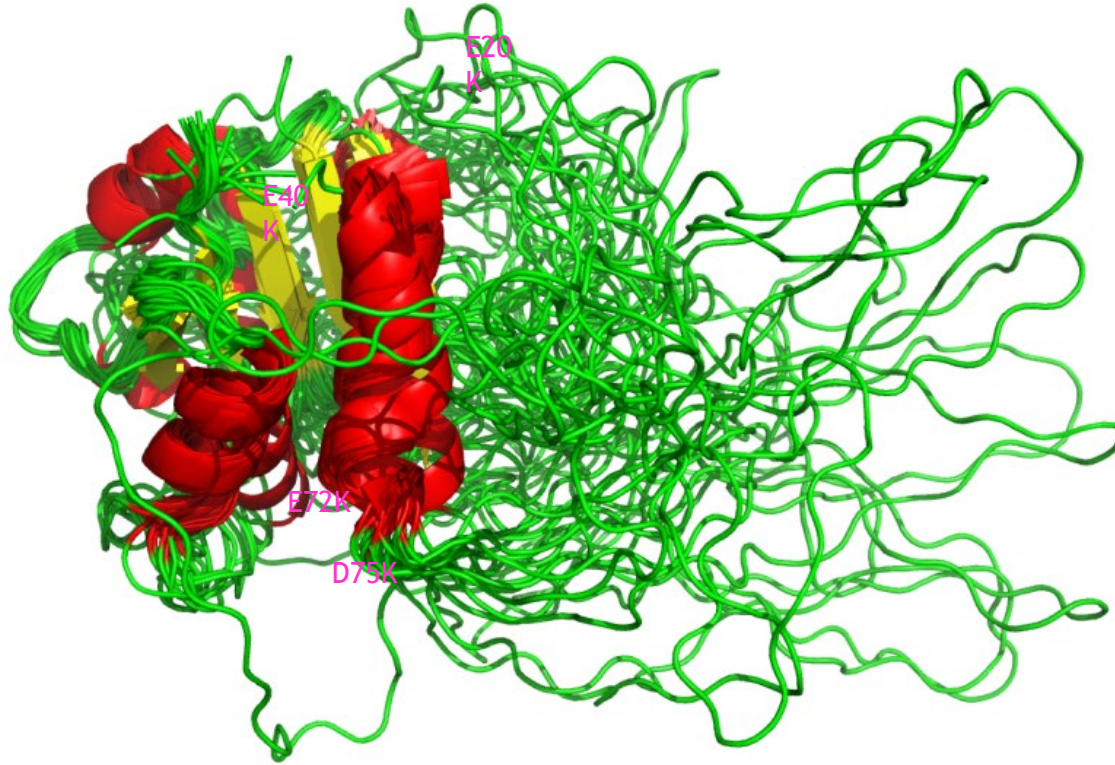
$$\Delta\Delta G_{NU} = -1.4 \text{ Kcal mol}^{-1}$$

Campos et al., J. Mol. Biol (2004) 344:223-237

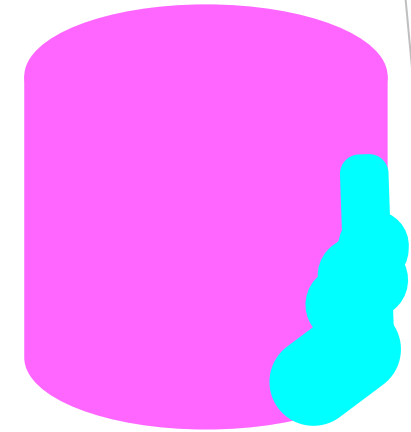
► Three-state proteins



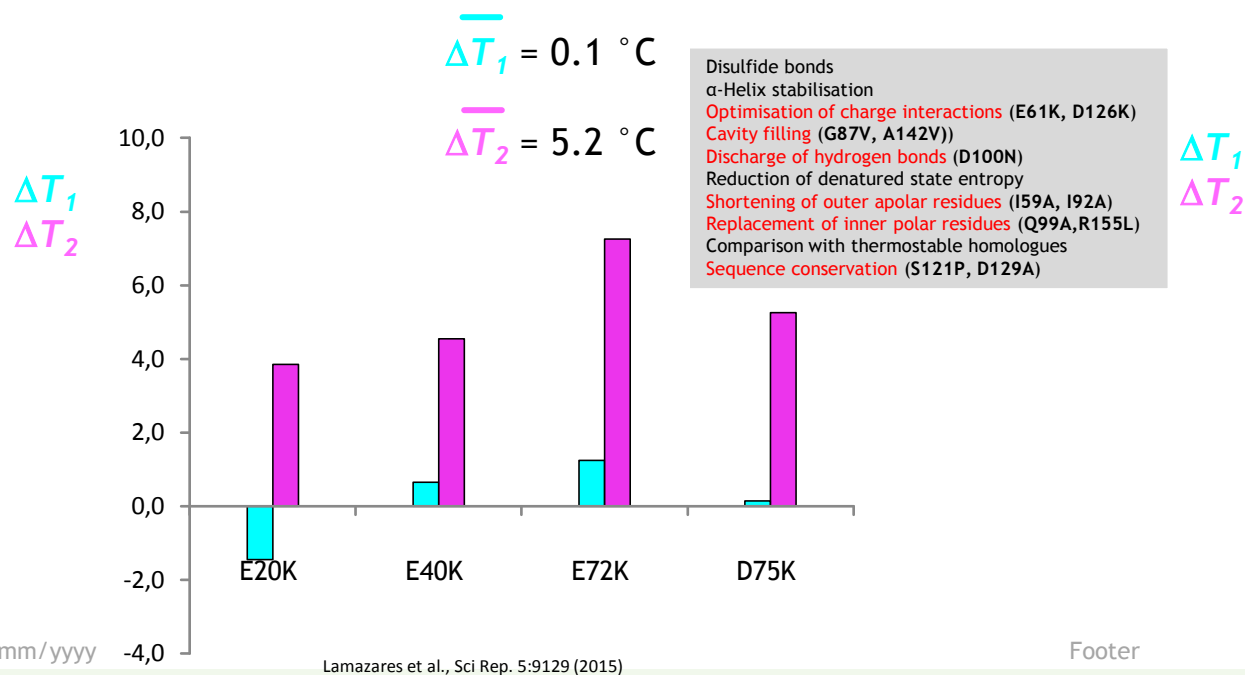
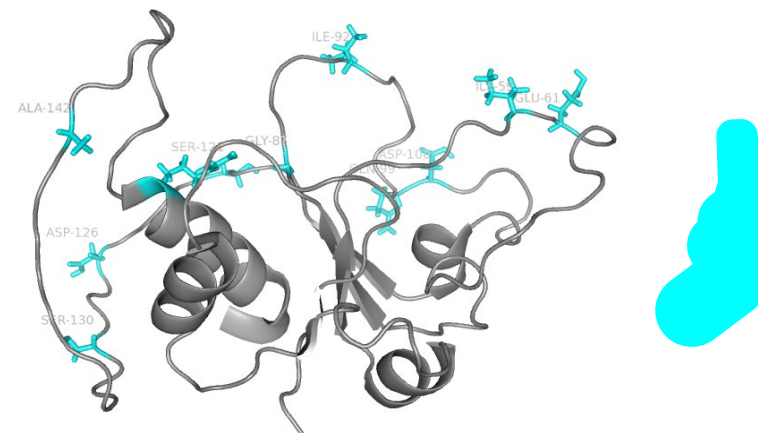
► Three-state proteins



Ayuso-Tejedor et al., J. Mol. Biol. 400:922-934 (2010)

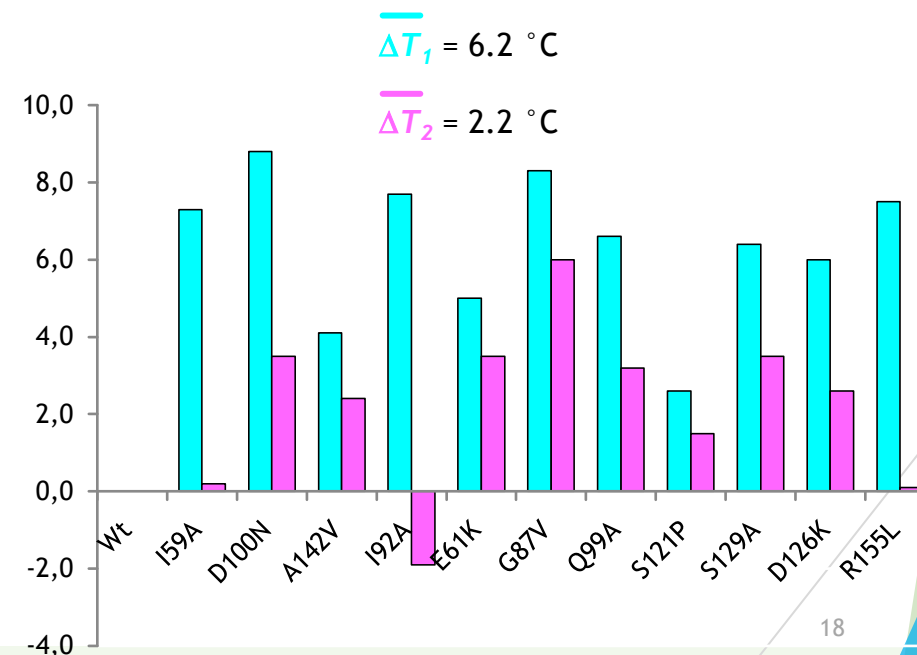


► Three-state proteins

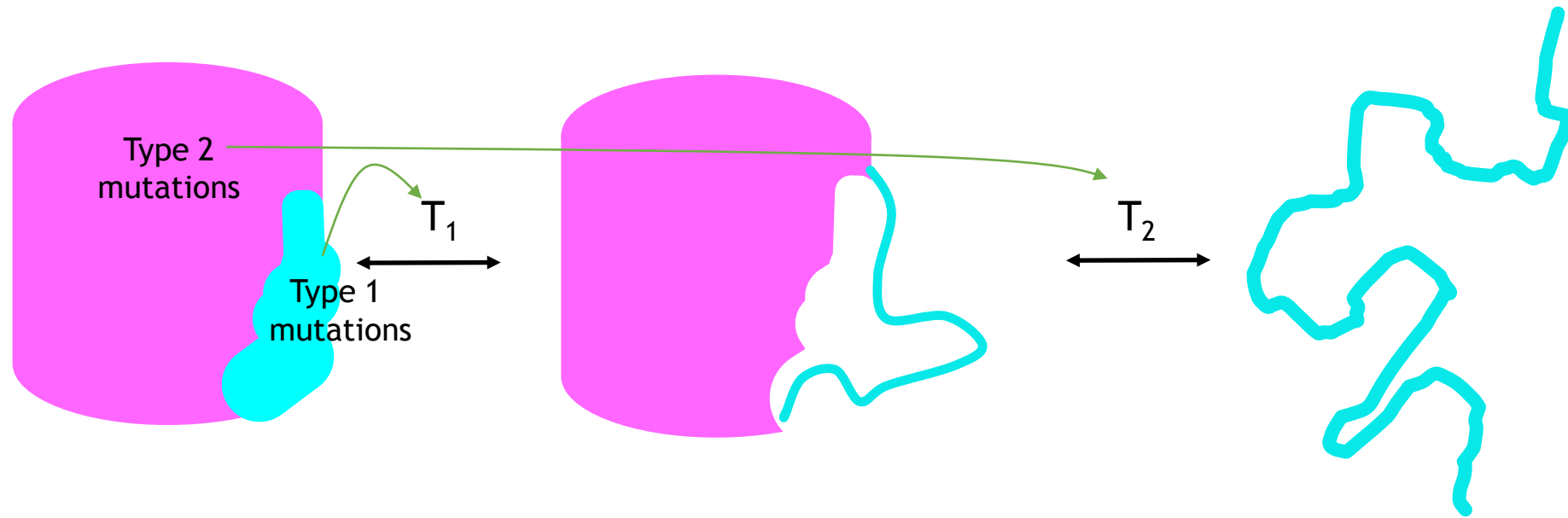


Lamazares et al., Sci Rep. 5:9129 (2015)

Footer



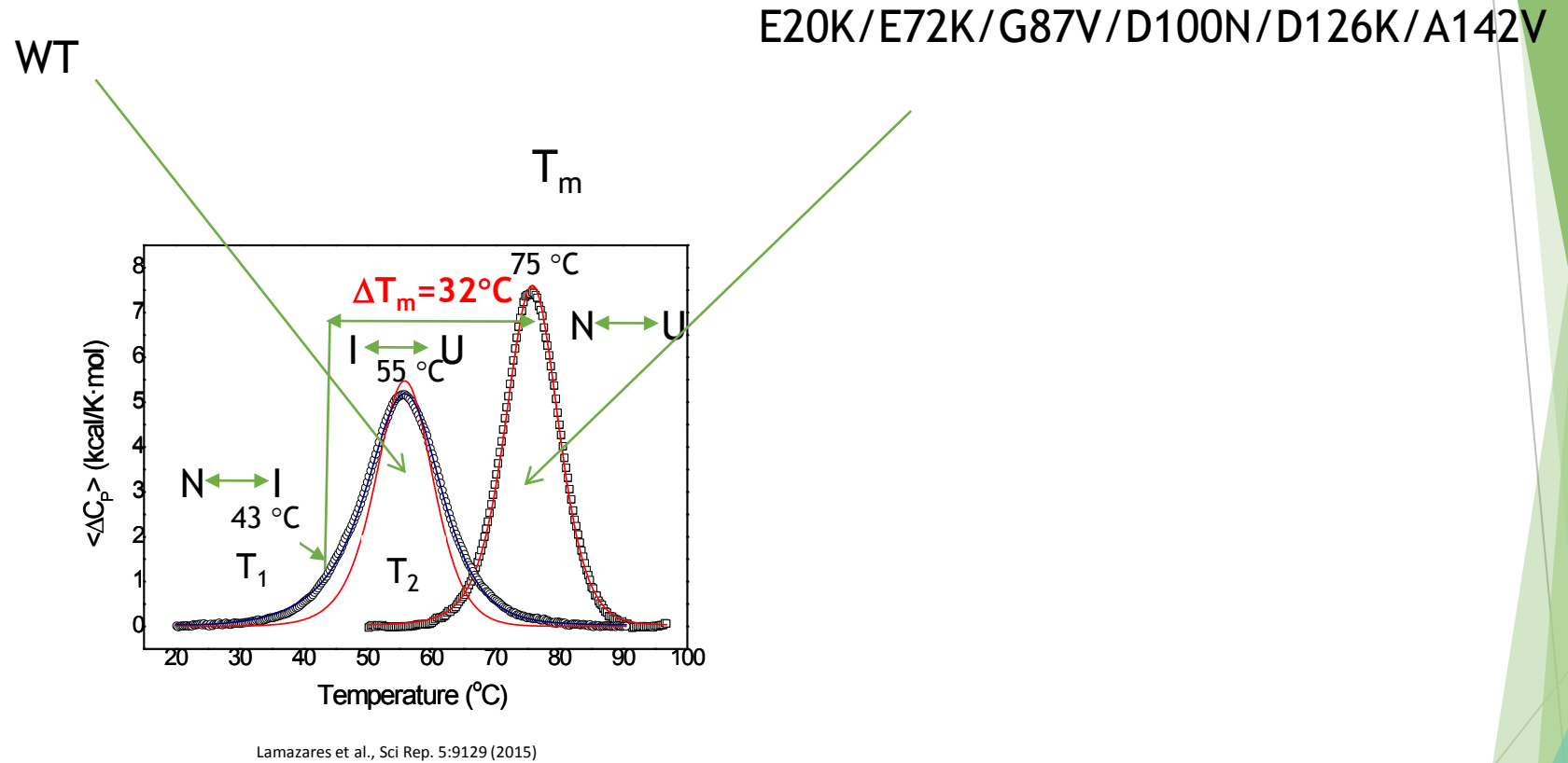
► Three-state proteins



Type 1 mutations: located in the weakest region or at the interface with the rest, they increase the relevant stability

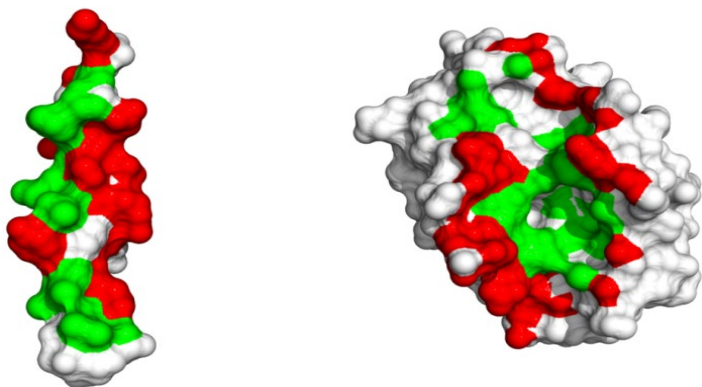
Type 2 mutations: they only increase the residual stability of the intermediate

► Three-state proteins

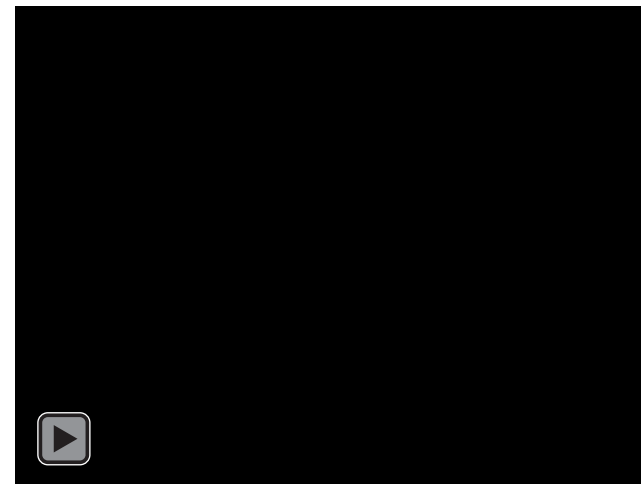


► Three-state proteins

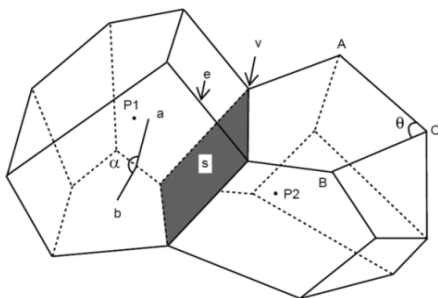
Polarity ratio



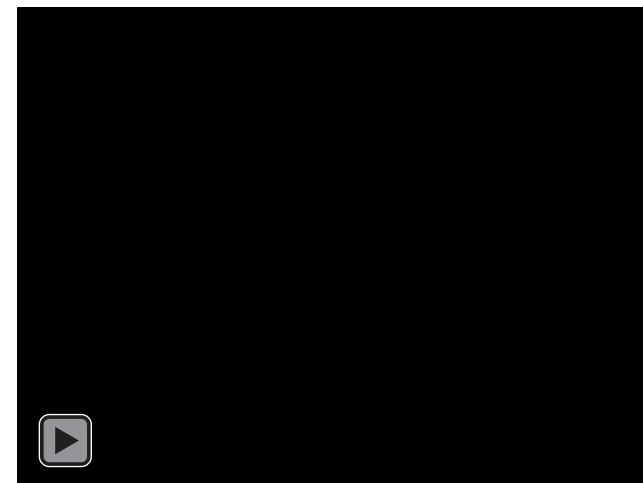
$$PR_{intraface} = \frac{\sum_{i=1}^m SASA_{(polar)_i}}{\sum_{j=1}^n SASA_{(hydrophobic)_j}}$$



Packing density



$$\rho_{intraface} = \frac{\sum_{i=1}^N V_i^{\circ}}{\sum_{i=1}^N V_i}$$



► Three-state proteins

Old LIPS web

(only precalculated proteins) monomers)

ProteinLIPS

Prediction of locally unstable segments of proteins

ProteinLIPS scans protein structures, computes packing densities and surface polarities, and predicts the locally unstable segments which are likely to be the first parts of the protein to unfold, the last parts to fold or to be involved in protein dynamics. You can query using PDB codes or keywords, e.g. if interested in a family of proteins.

Submission form

Please, fill in all fields. Only ASCII printable characters are allowed.

Input type (*)

☒ Simple Search ☐ Complex Search

Search (*)

All

Examples: 1BWV, HYDROLASE, LYSOZYME, HOMO SAPIENS, CHICKEN

SUBMIT

Angarica VE, Sancho J. Protein dynamics governed by interfaces of high polarity and low packing density. PLoS One. 2012;7(10):e48212. doi: 10.1371/journal.pone.0048212. Epub 2012 Oct 26.

dd/mm/yyyy

<http://webapps.bifi.es/lips>

Footer



► Three-state proteins

New LIPS web (any monomer or oligomer). Under construction.

LIPS test

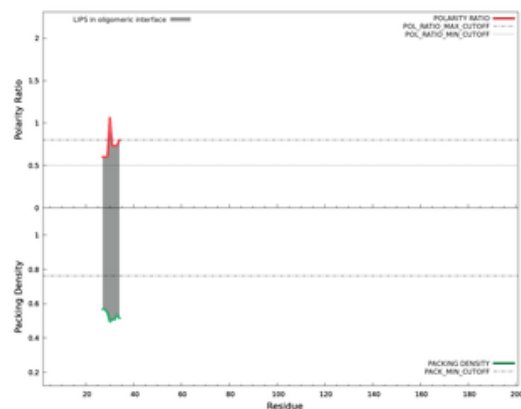
PDB: [7W6W](#)

Estructura: 2-homo

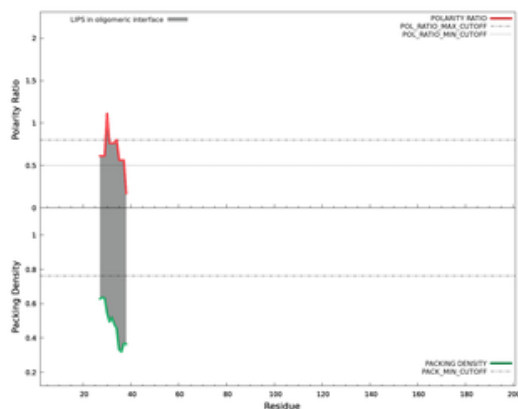
Cadenas elegidas: AB



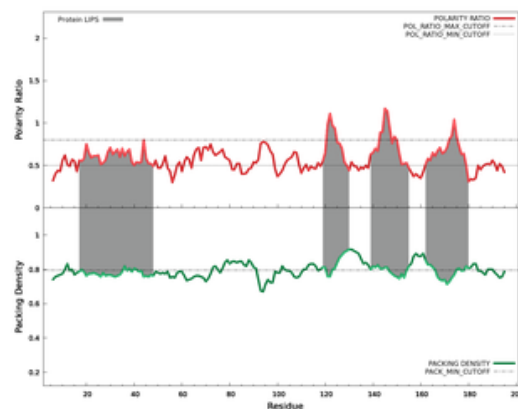
Preprocesado del PDB: Downloaded PDB from RCSB. No important issues found in the PDB analysed! Any minor issue resting in the PDB file will be translated to the generated LIPS profiles.



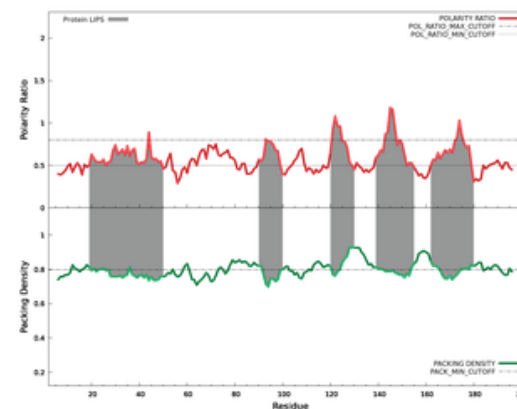
7w6w_dim_A.png



7w6w_dim_B.png



7w6w_mon_A.png



7w6w_mon_B.png

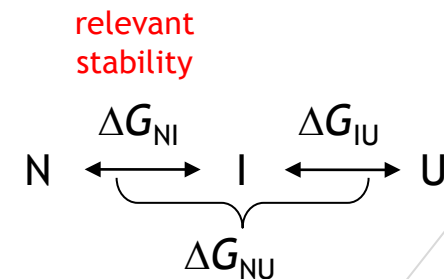
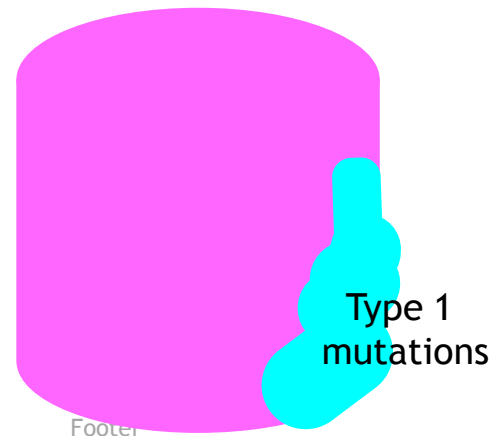
	40	60	80	100	120	140	160	180
dimer_A	DQQTMEIHHDKHNTYVTKLNAAVEGTDLESKSEIEIVANLDSVPENIQTAVRNNGGGHLNHSLEWELLTPNSEEGTGVVDKIKEQNGSLDAFKEEFANQAAARFGSGWAWLVNDGKLEIVTTPNQDNPLTEGKTPILGLDVWEHAYYWKYQNKRPDIYISAFWNVVNW							
dimer_B	DQQTMEIHHDKHNTYVTKLNAAVEGTDLESKSEIEIVANLDSVPENIQTAVRNNGGGHLNHSLEWELLTPNSEEGTGVVDKIKEQNGSLDAFKEEFANQAAARFGSGWAWLVNDGKLEIVTTPNQDNPLTEGKTPILGLDVWEHAYYWKYQNKRPDIYISAFWNVVNW							
monomer_A	DQQTMEIHHDKHNTYVTKLNAAVEGTDLESKSEIEIVANLDSVPENIQTAVRNNGGGHLNHSLEWELLTPNSEEGTGVVDKIKEQNGSLDAFKEEFANQAAARFGSGWAWLVNDGKLEIVTTPNQDNPLTEGKTPILGLDVWEHAYYWKYQNKRPDIYISAFWNVVNW							
monomer_B	DQQTMEIHHDKHNTYVTKLNAAVEGTDLESKSEIEIVANLDSVPENIQTAVRNNGGGHLNHSLEWELLTPNSEEGTGVVDKIKEQNGSLDAFKEEFANQAAARFGSGWAWLVNDGKLEIVTTPNQDNPLTEGKTPILGLDVWEHAYYWKYQNKRPDIYISAFWNVVNW							

TO ENGINEER A MORE STABLE PROTEIN YOU WOULD LIKE:

1- To get a list of promising stabilizing mutations

and, unless the protein is 2-state

2-To identify the likely unstable regions of the protein



- ▶ Protein stabilization by formulation: some rational
- ▶ Examples
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- ▶ Examples
- ▶ Three-state proteins

dd/mm/yyyy

Footer

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