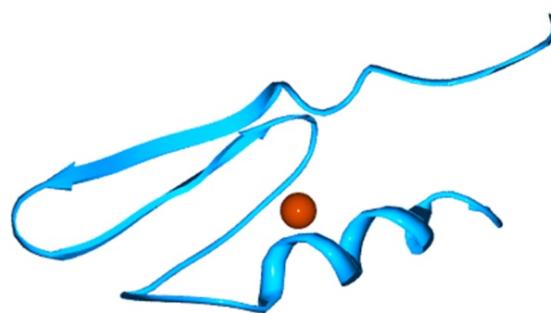
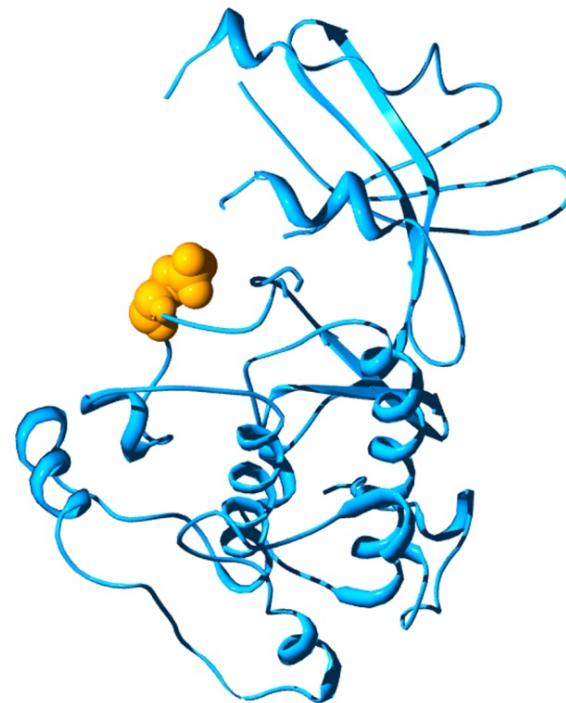


EXAMPLES OF FUNCTIONAL SITES

A. Zinc-binding site



B. Phosphorylation site



C. DNA-binding site



MOLECULAR CONSEQUENCES (ON P53)

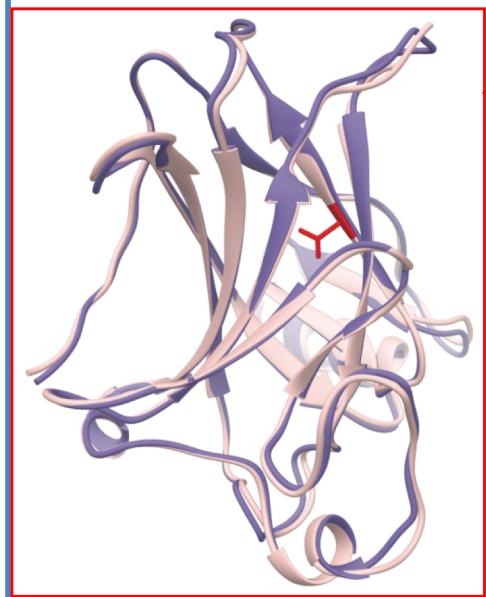


R175H: Metal-binding



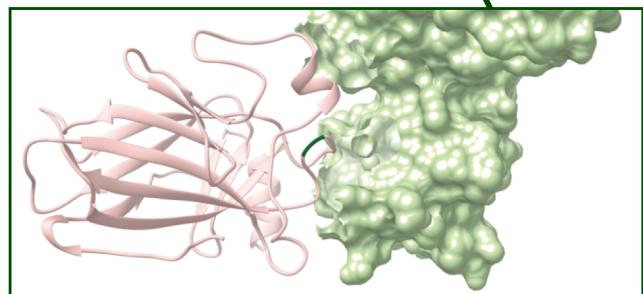
K120R: Acetylation

V143A: Stability

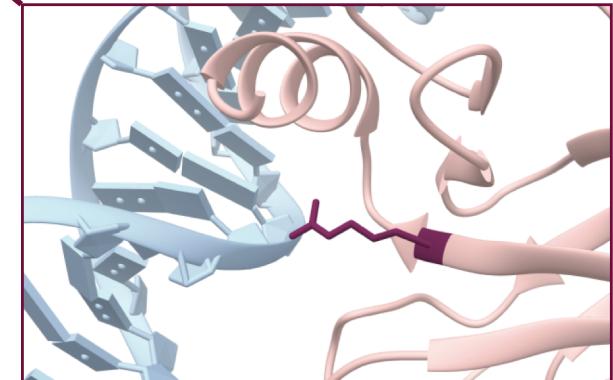


p53 – tumor suppressor protein

G245S: Protein-binding

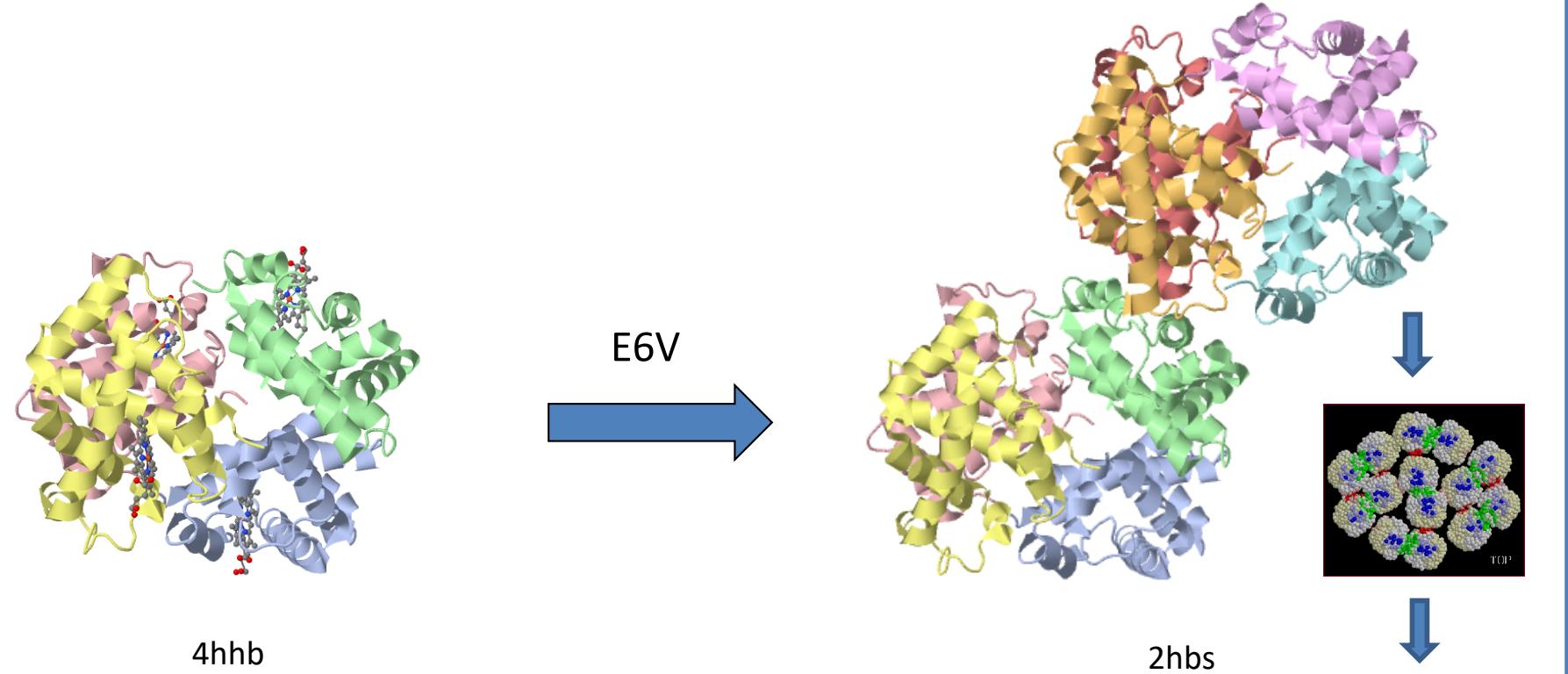


R273H: DNA-binding



PDB structures: 2ybg, 2j1w, 1ycs and 1tup

SICKLE CELL DISEASE: GAIN OF FUNCTION



Sickle Cell Disease:

Autosomal recessive disorder

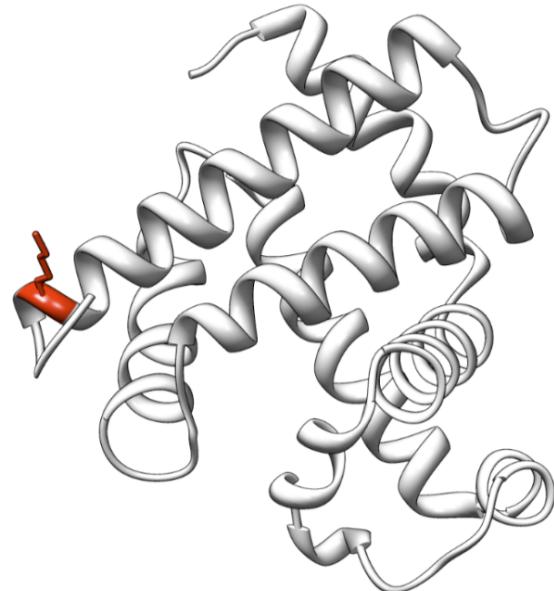
E6V in HBB causes interaction w/ F85 and L88

Formation of amyloid fibrils

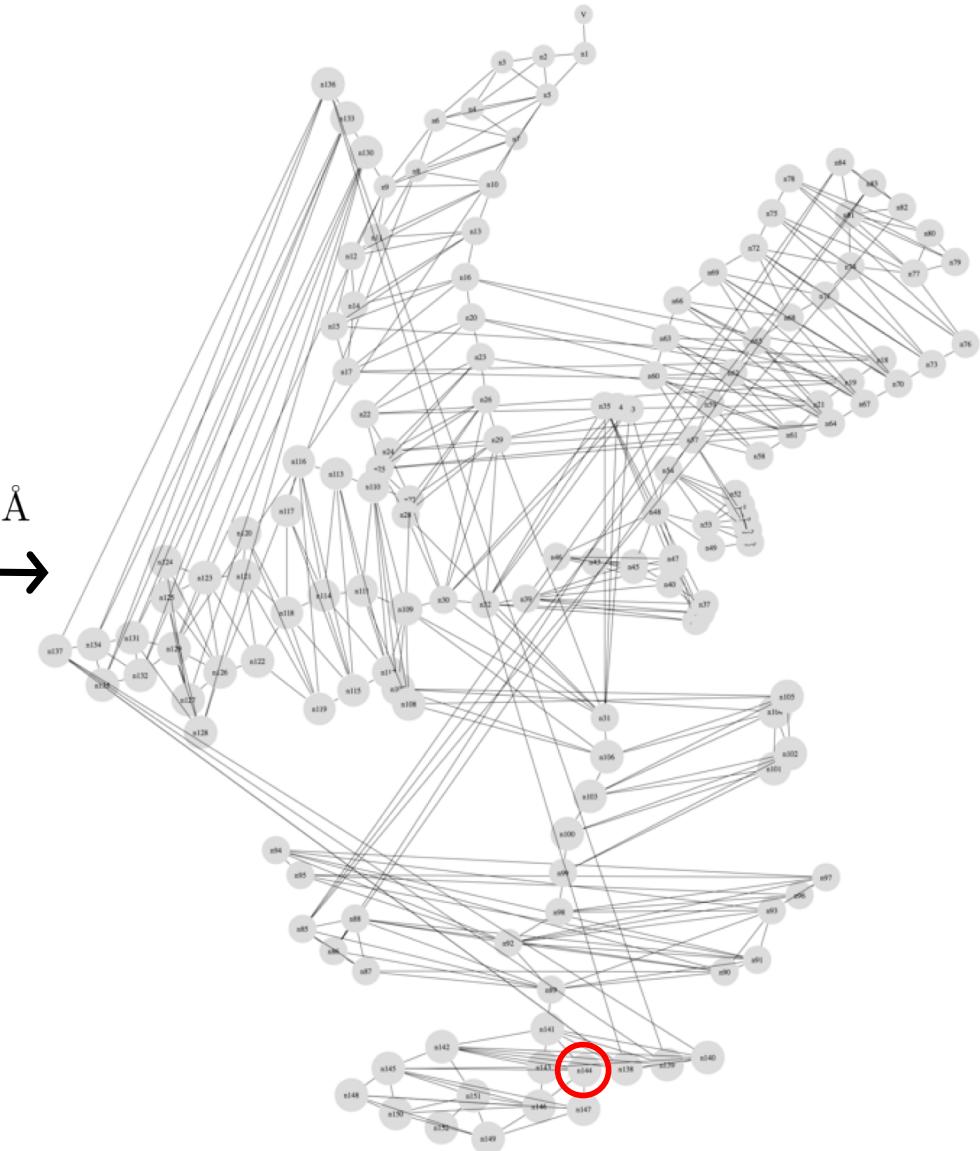
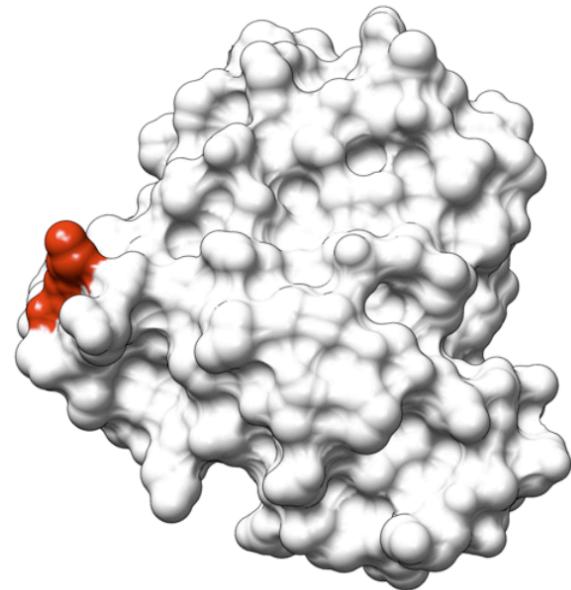
Abnormally shaped red blood cells, leads to sickle cell anemia

Manifestation of disease vastly different over patients

PREDICTION OF FUNCTIONAL RESIDUES FROM STRUCTURE

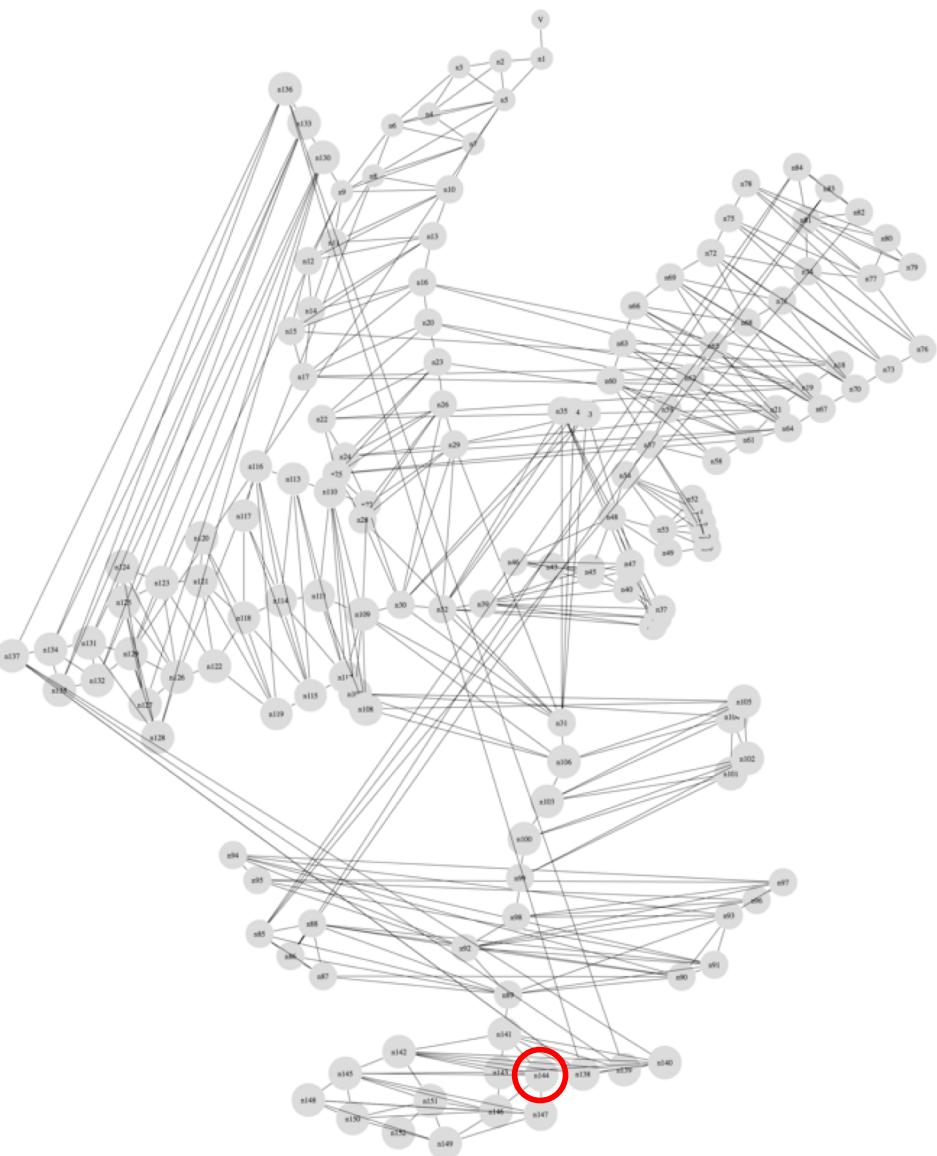
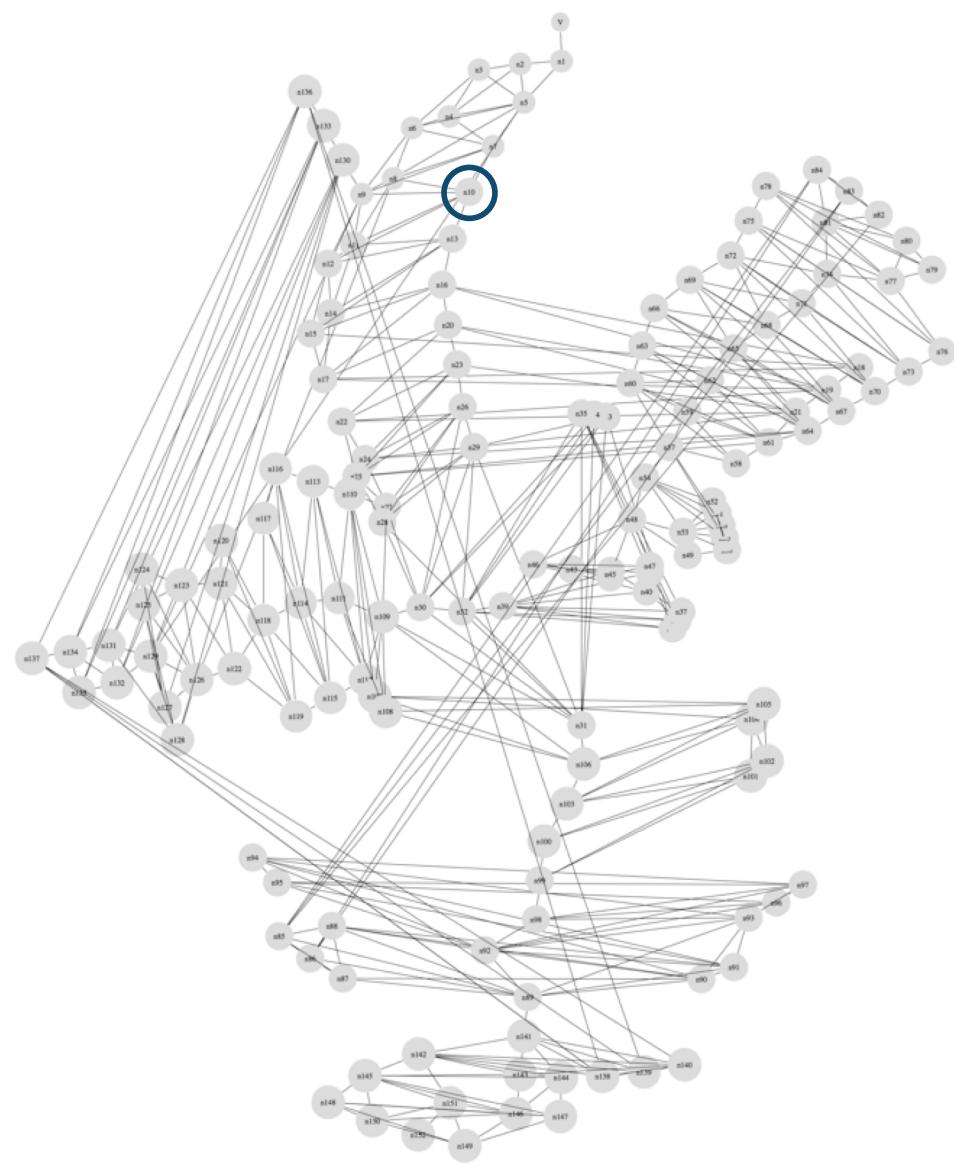


$$C_{\alpha} - C_{\alpha} \leq 6\text{\AA}$$
A black arrow pointing from the ribbon diagram towards the network graph below.

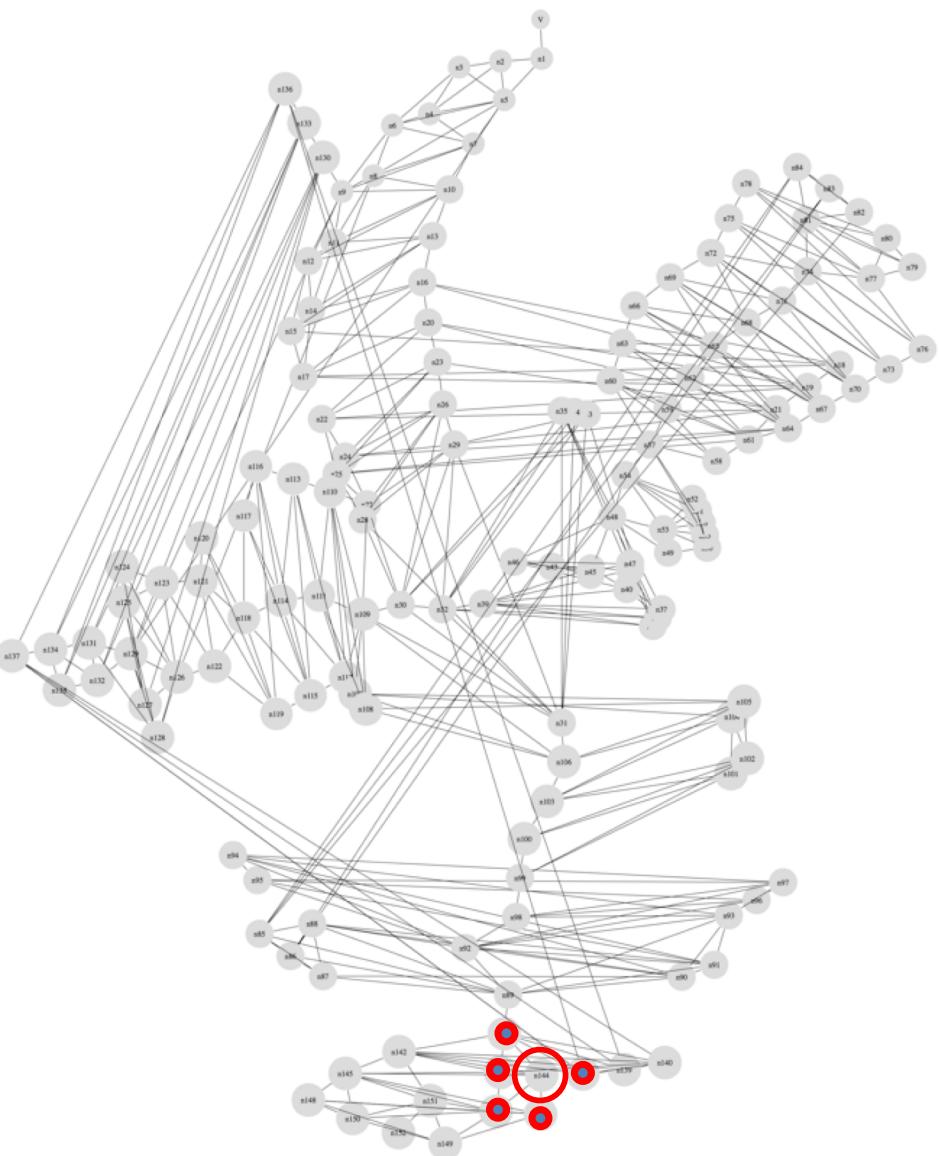
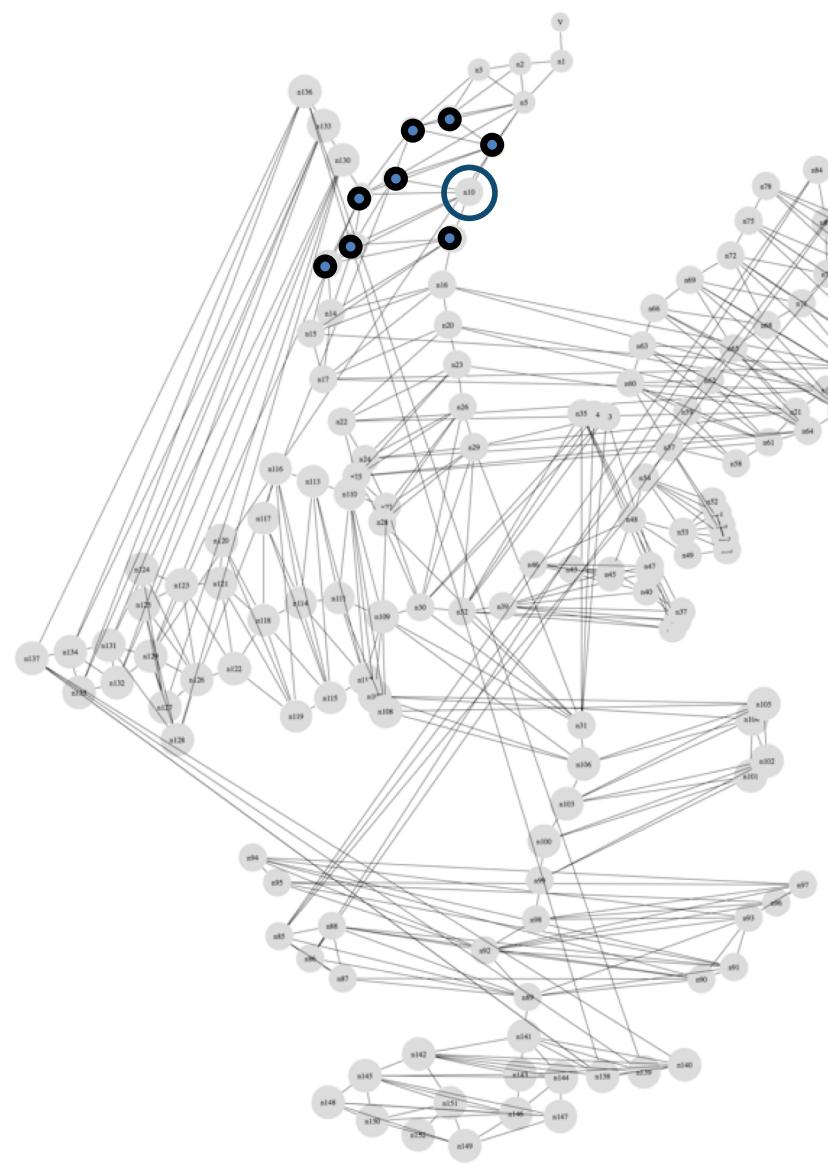


Myoglobin 1.4A X-ray PDB: 2jho 153 residues

PROTEIN CONTACT GRAPH



PROTEIN CONTACT GRAPH



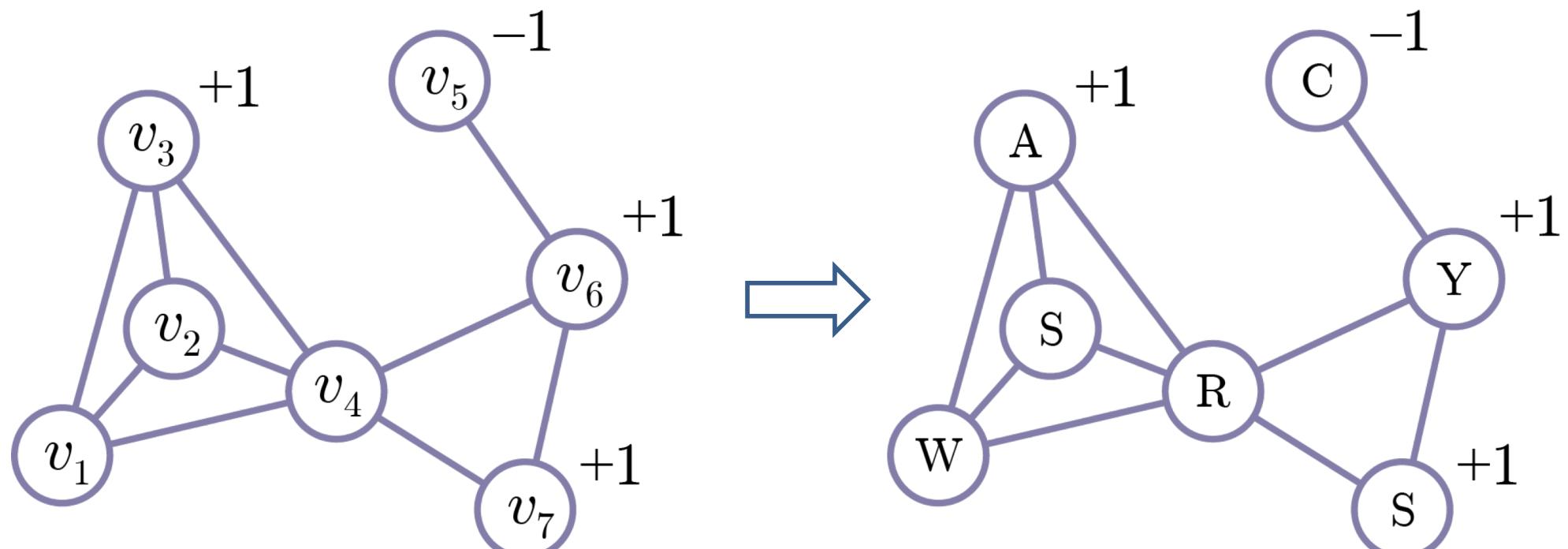
NOTATION AND PROBLEM FORMULATION

$$G = (V, E)$$

$$t : V \rightarrow \{-1, +1\}$$

$$\Sigma = \{A, C, \dots, W, Y\}$$

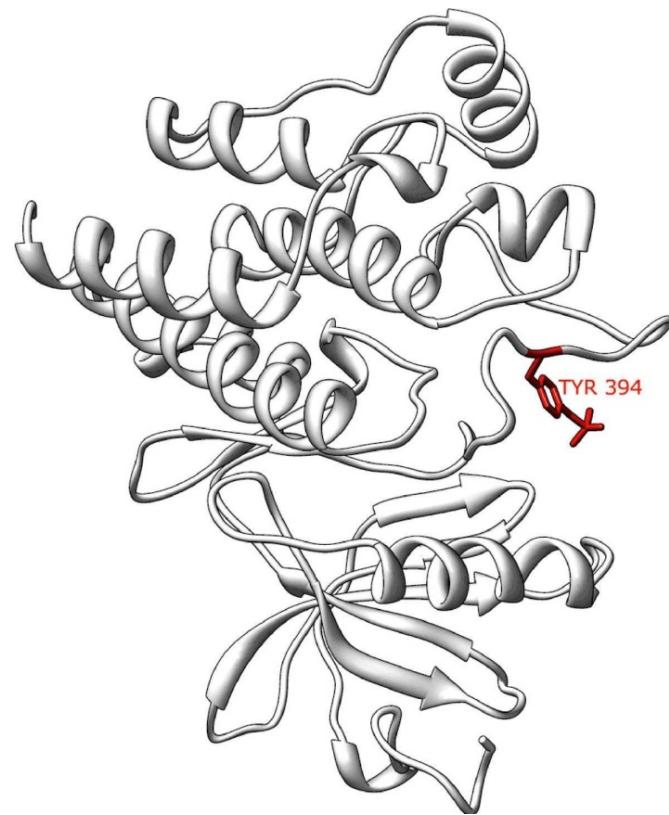
$$f : V \rightarrow \Sigma$$



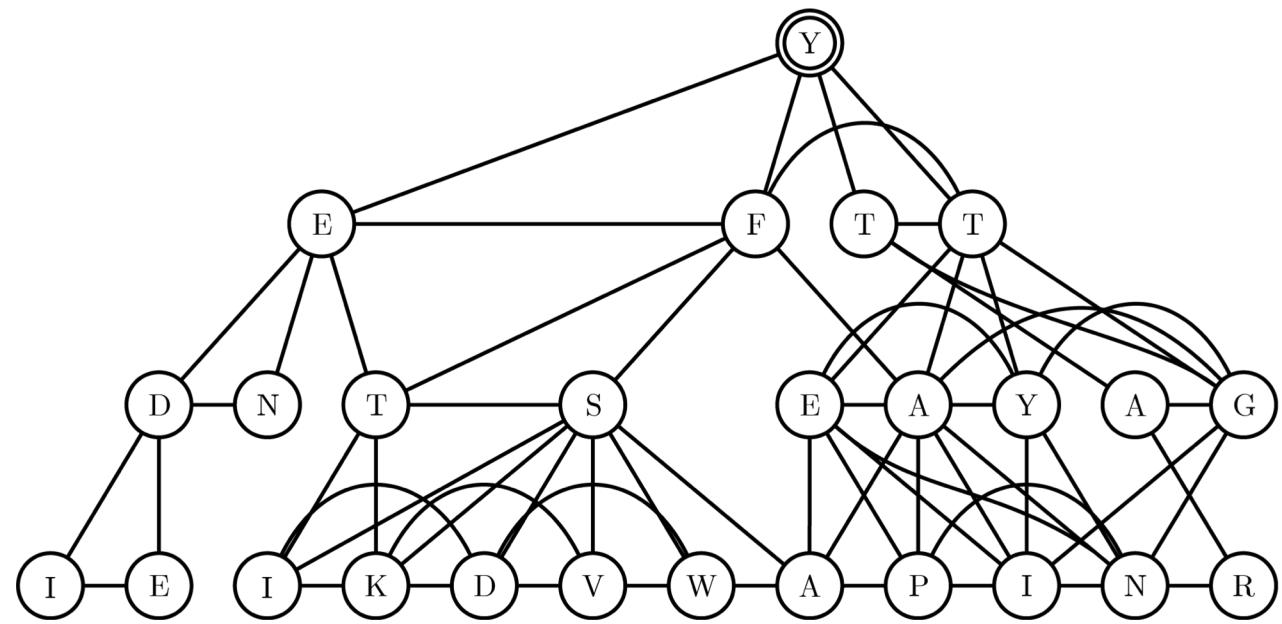
Objective: predict class label for each unlabeled vertex

e.g. $\Pr(t(v_4) = +1 | D) = 0.9$

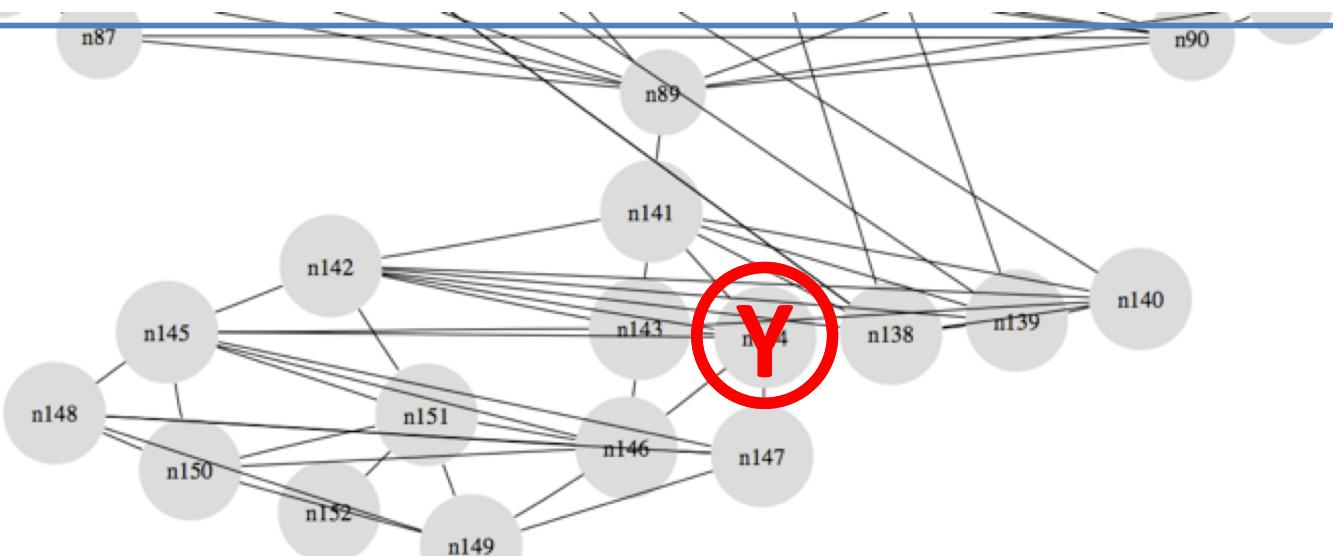
RESIDUE NEIGHBORHOOD



Y394 of human lymphocyte kinase



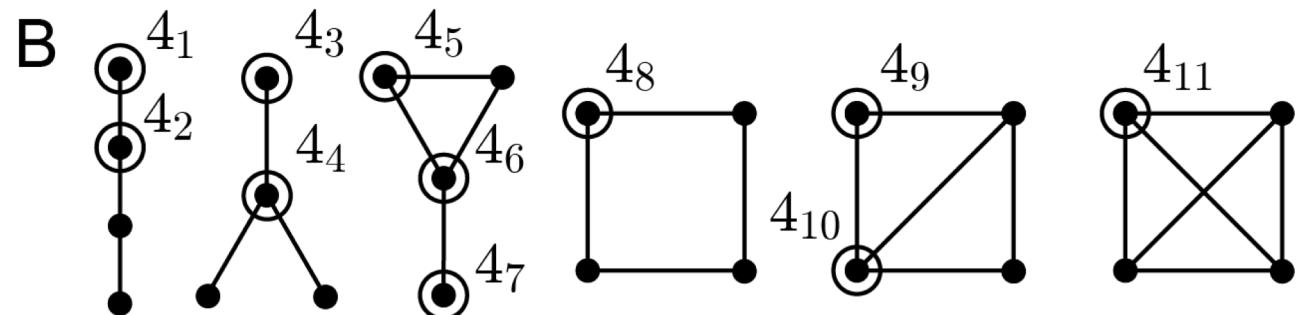
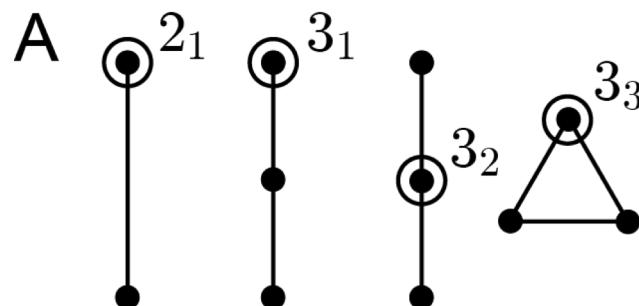
Depth-3 graph neighborhood for Y394



Idea:

1. Count small graphs rooted at the vertex of interest
 - i. *Graphlets*
2. Create a similarity measure between the counts for two vertices
 - i. *Kernel function*
3. Use kernel functions for supervised classification
 - i. *Support vector machines*

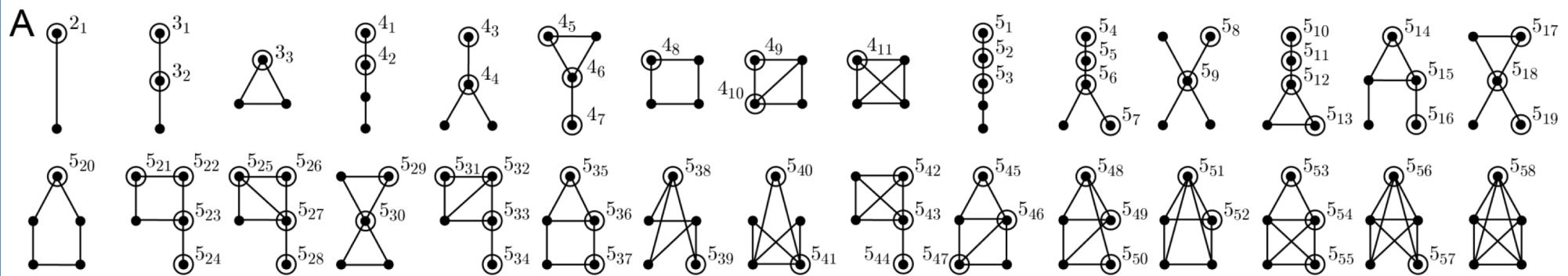
⇒ Graphlet Kernel



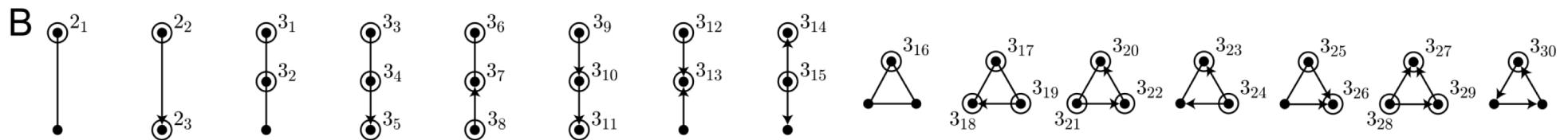
Graphlets: simple small connected rooted graphs.

GRAPHLETS

Undirected:



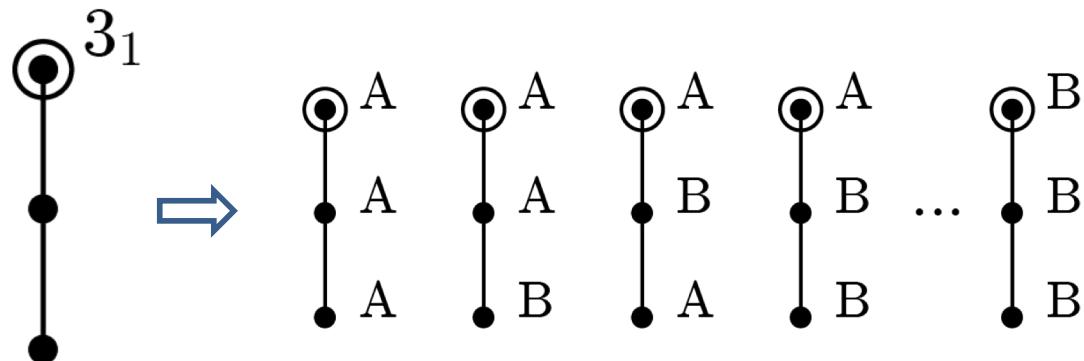
Directed:



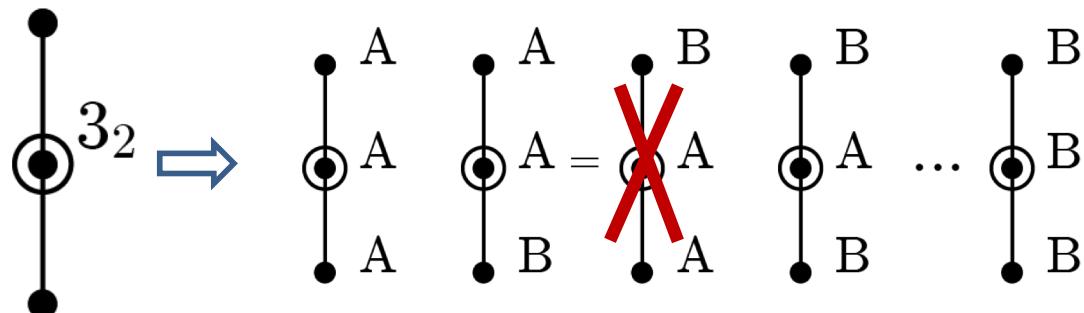
Graphlets: simple small (typically of order 5 or less) connected rooted graphs.

LABELED GRAPHLETS

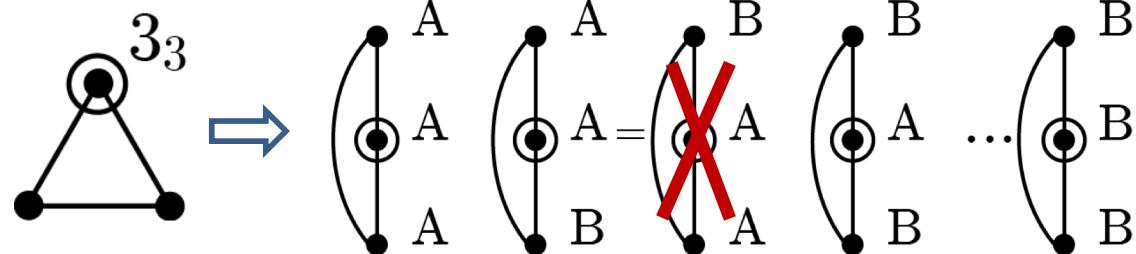
$$\kappa(n, \Sigma) = 20$$



$$|\Sigma|^n = 2^3 = 8$$



$$|\Sigma| \cdot \left(\frac{|\Sigma|+1}{|\Sigma|-1}\right) = 6$$

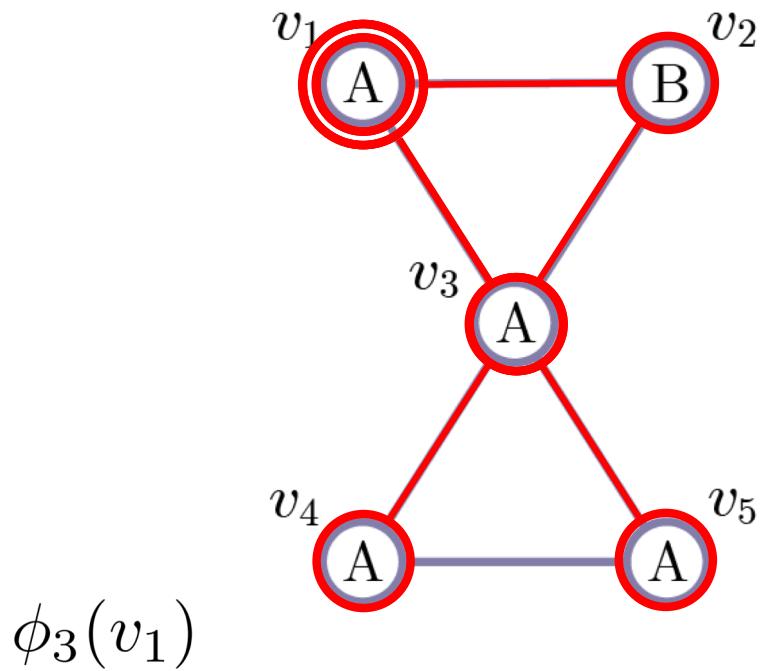


same symmetry class



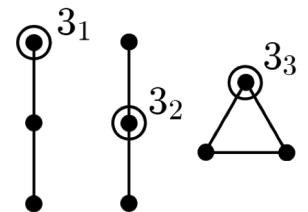
$$|\Sigma| \cdot \left(\frac{|\Sigma|+1}{|\Sigma|-1}\right) = 6$$

EXAMPLE



$$V = \{v_1, v_2, v_3, v_4, v_5\} \quad \Sigma = \{\text{A, B}\}$$

$$\begin{aligned} f : V &\rightarrow \Sigma \\ f(v_1) &= \text{A} \\ f(v_2) &= \text{B} \\ &\vdots \end{aligned}$$



AAA AAB ABA ABB BAA BAB BBA BBB AAA AAB ABB BAA BAB BBB AAA AAB ABB BAA BAB BBB

2																	1			
---	--	--	--	--	--	--	--	--	--	--	--	--	--	--	--	--	---	--	--	--

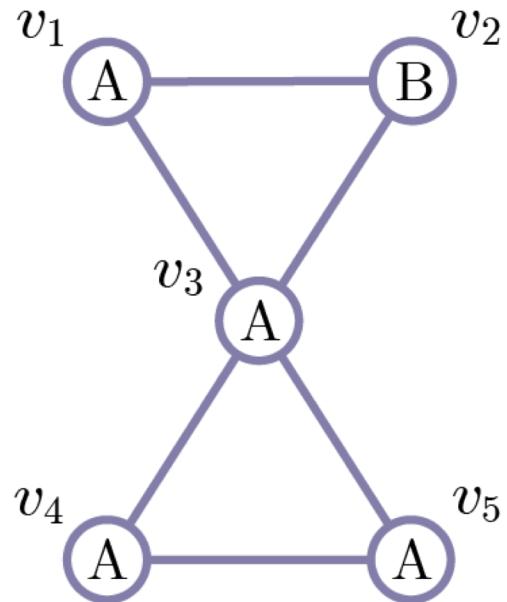
←———— 3₁ —————>———— 3₂ —————>———— 3₃ —————>

1	1															1			
---	---	--	--	--	--	--	--	--	--	--	--	--	--	--	--	---	--	--	--

$\phi_3(v_5)$

$$k_3(v_1, v_5) = \langle \phi_3(v_1), \phi_3(v_5) \rangle = 2$$

GRAPHLET KERNELS



$$k_n(u, v) = \langle \phi_n(u), \phi_n(v) \rangle$$

where

$$\phi_n(v) = (\varphi_{n_1}(v), \varphi_{n_2}(v), \dots, \varphi_{n_{\kappa(n, \Sigma)}}(v))$$

$$k(u, v) = \sum_{n=1}^N k_n(u, v)$$

Graphlet kernel

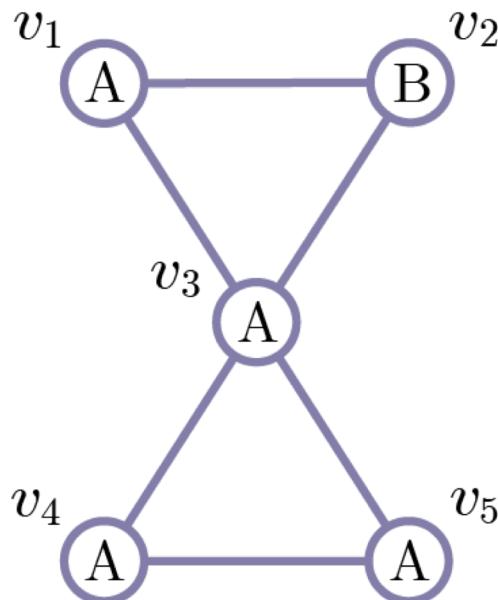
$$k'(u, v) = \frac{k(u, v)}{\sqrt{k(u, u) k(v, v)}}$$

Normalized graphlet kernel

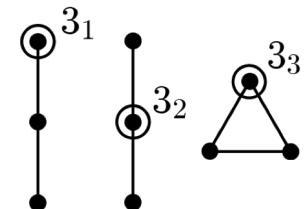
EDIT DISTANCE GRAPHLET KERNELS



Jose Lugo-Martinez



IDEA: Allow approximate matching,
i.e. allow mismatch in labels or edges



AAA AAB ABA ABB BAA BAB BBA BBB AAA AAB ABB BAA BAB BBB AAA AAB ABB BAA BAB BBB

2	2	2		2									1	1	1		1	
---	---	---	--	---	--	--	--	--	--	--	--	--	---	---	---	--	---	--

←———— 3₁ —————>———— 3₂ —————>———— 3₃ —————>

2	2	1	1	1	1								1	1		1	
---	---	---	---	---	---	--	--	--	--	--	--	--	---	---	--	---	--

$$k_{(3,1)}^l(v_1, v_5) = \left\langle \phi_{(3,1)}^l(v_1), \phi_{(3,1)}^l(v_5) \right\rangle = 14$$

RESULTS: PHOSPHORYLATION

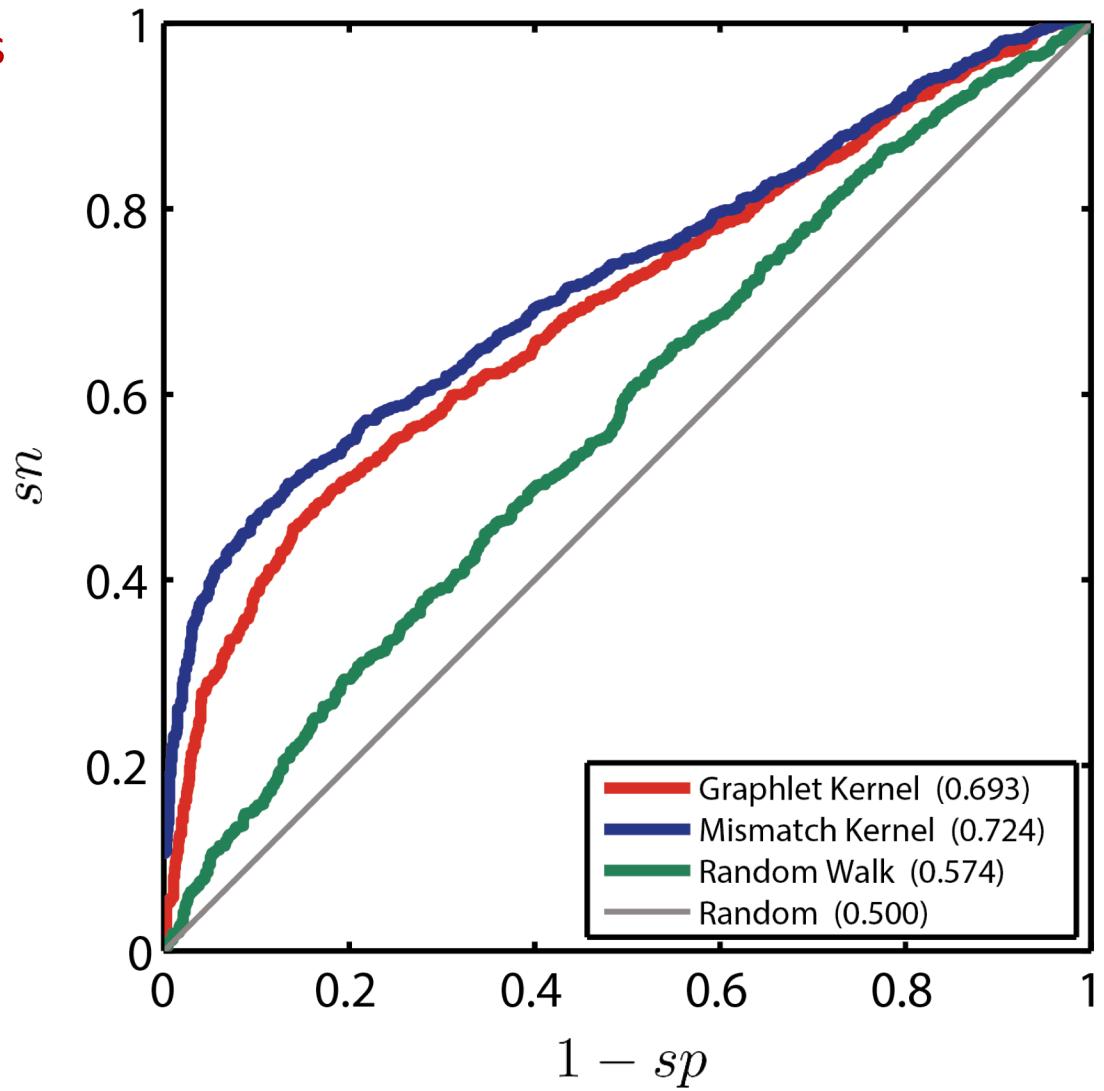
Task: predict phosphorylation sites
from protein structures

Data:

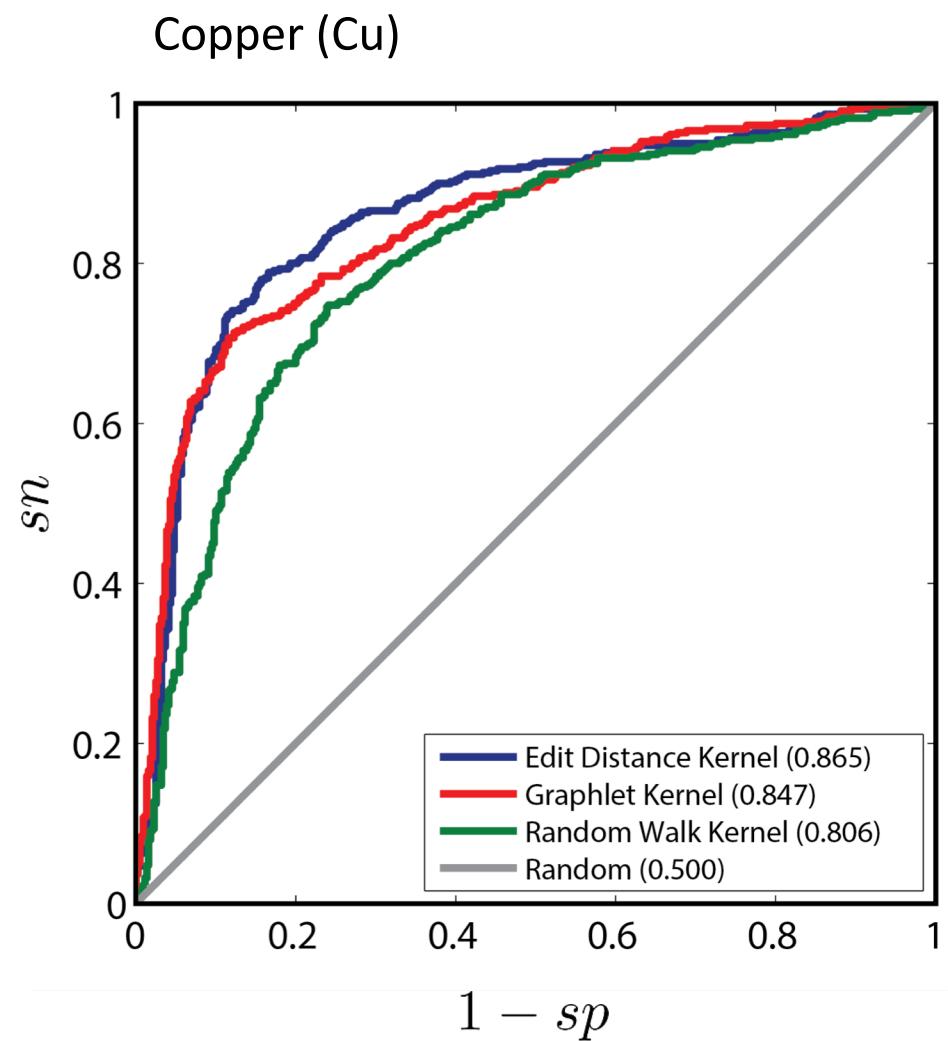
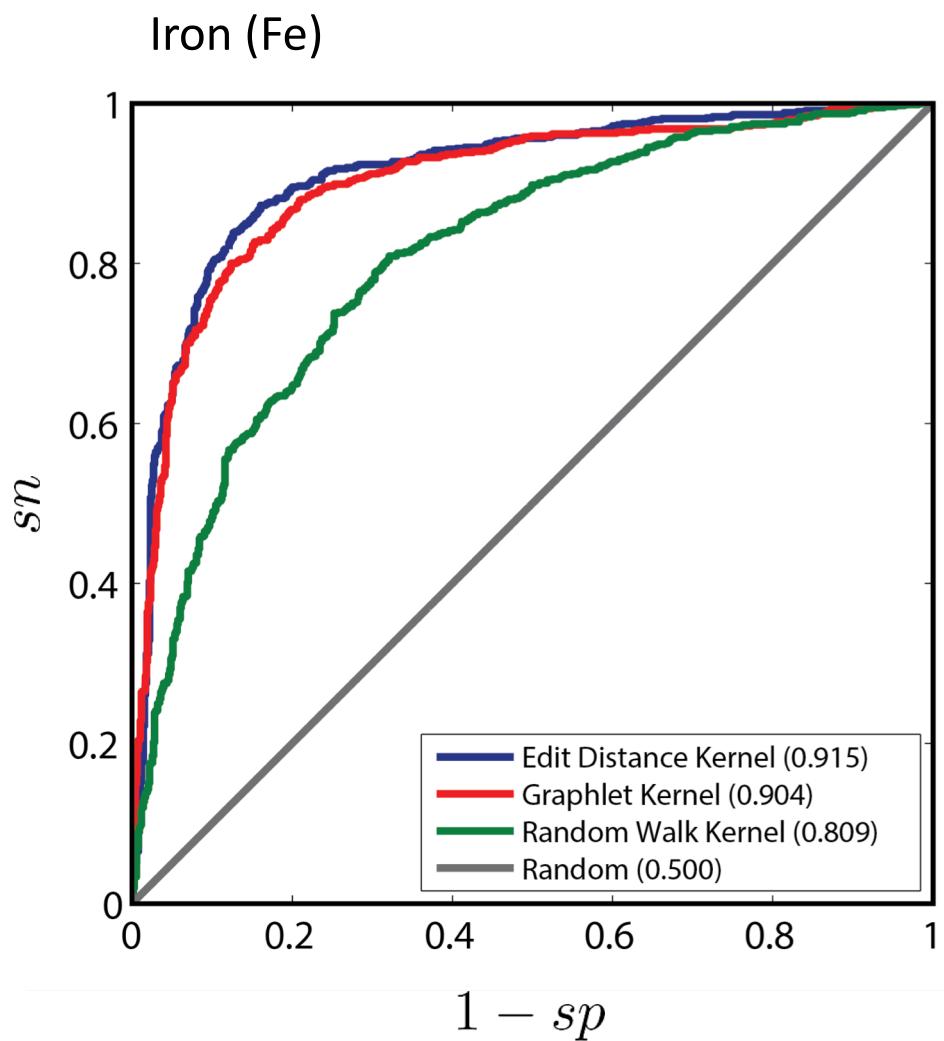
$$n_+ = 1157$$

$$n_- = 1157$$

$$|\Sigma| = 40$$

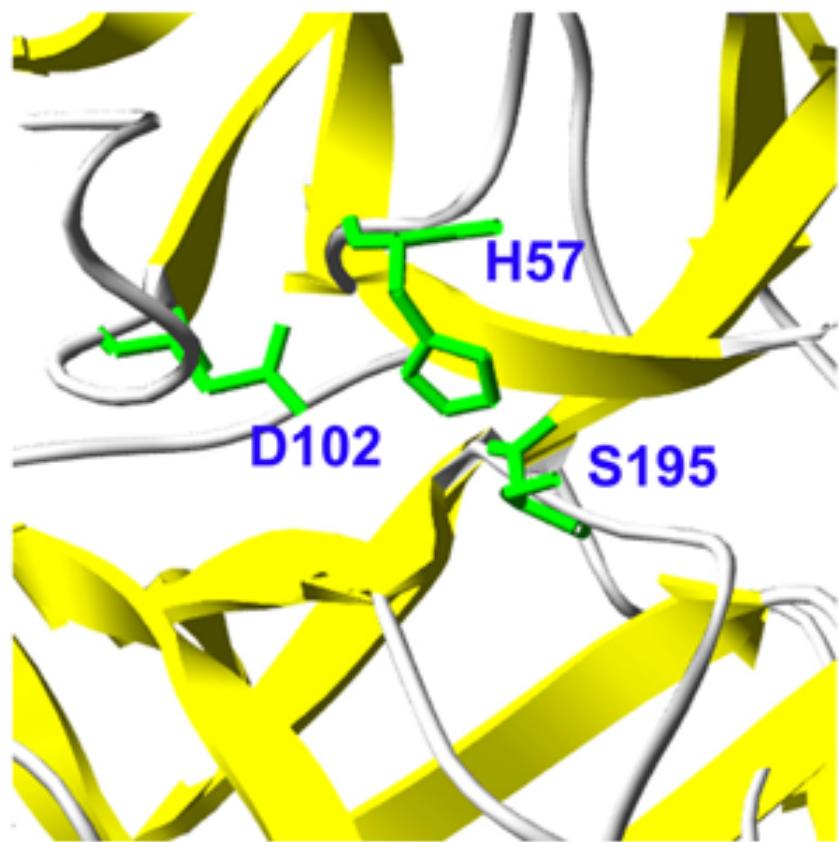


RESULTS: METAL BINDING RESIDUES



LOSS OF CATALYTIC ACTIVITY CAUSES DISEASE

Catalytic triad of F9 (1frn)



- F9 is activated in response to injury of the blood vessel and has a key role in blood clot formation;
- Mutations in F9 give rise to the X-linked recessive disorder, hemophilia B;
- H57R leads to the loss of catalytic activity.

GAIN OF CATALYTIC ACTIVITY CAUSES DISEASE

- a member of the proteinase K subfamily of subtilases that reduces the number of LDL receptors in liver through a posttranscriptional mechanism.
- D374Y leads to a 10-fold increase in catalytic activity that causes hypercholesterolemia.

Catalytic residues of PCSK9 (2qtw)

