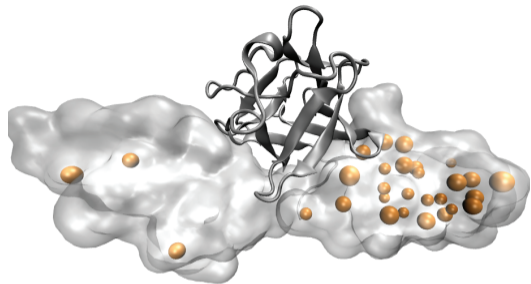


A.

- Hinge residues
- Buried residues
- Protein-protein interaction site
- Post-translational modifications
- HGMD site (w/o annotation overlap)
- HGMD site (w/annotation overlap)

B.

residue 150

residue 350