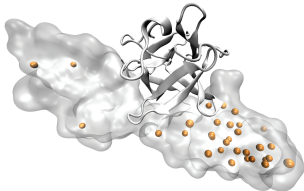


A

- Predicted allosteric (surface)
- Predicted allosteric (interior)
- Hinge residues
- Buried residues
- Protein-protein interaction site
- Post-translational modifications
- HGMD site (w/o annot. overlap)
- HGMD site (w/annot. overlap)

B