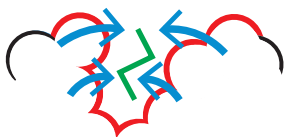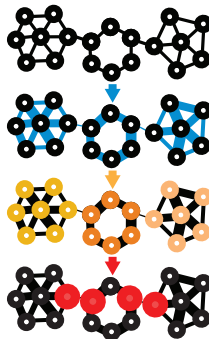predicting **allosteric residues** with **dynamics**

surface

interior

sequence-based analyses

across species

amongst humans