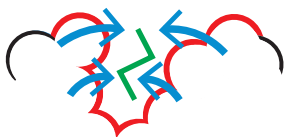
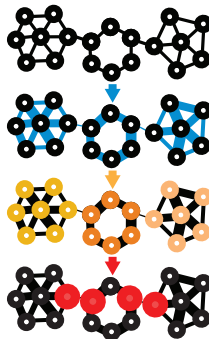


predicting **allosteric residues** with **dynamics**

surface



interior



sequence-based analyses

across species

YQDLKPENLAIDQQGYDQLWALGQ
 YRDLRPQNLLIEQQGEIQVWALGV
 YAALRPQNLHIFQMGYIQAWVLGV
 YDDLAPPNDLTVQOGNAQDWALGV

amongst humans

