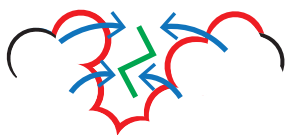
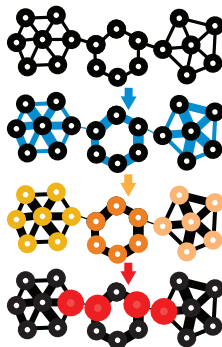


predicting allosteric residues

surface



interior



sequence-based analyses

across species

amongst humans

Y	Q	D	L	K	P	E	N	L	A	I	D	Q	Q	G	Y	D	Q	L	W	A	L	G	Q
Y	R	D	L	R	P	Q	N	L	L	I	E	Q	Q	G	E	I	Q	V	W	A	L	G	V
Y	A	A	L	R	P	Q	N	L	H	I	F	Q	M	G	Y	I	Q	A	W	V	L	G	V
Y	D	D	L	A	P	P	N	D	L	T	V	Q	Q	G	N	A	Q	D	W	A	L	G	V

