a Find a selection record

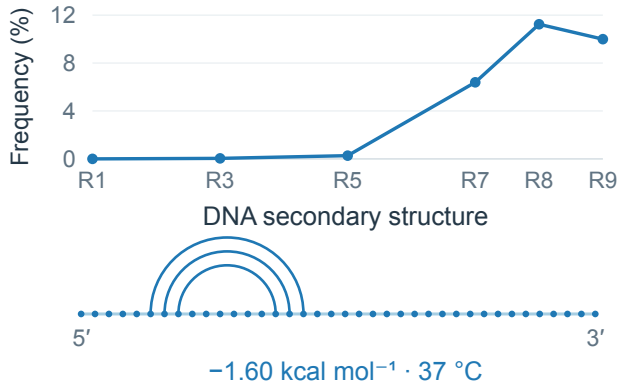
SAA

TARGET	UNIPROT
SAA	P0DJI8 ↗

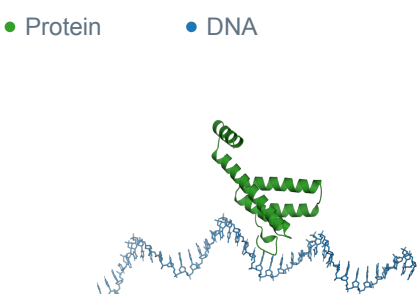
Human serum amyloid A

38-nt DNA selection candidate

b Follow selection and folding



c Inspect a predicted complex



Boltz-2 · ipTM 0.411

d Examine the interface



Minimum 2.42 Å