Table 3 The backbone confirmation of models found from rampage serve

Analysis	Protein structure	Domain structure
Number of residues in favoured region	517 ( 91.8%)	159 ( 88.3%)
Number of residues in allowed region	34 ( 6.0%)	15 ( 8.3%)
Number of residues in outlier region	12 ( 2.1%)	6 ( 3.3%)