

**Summary of integrative structure determination of Integrative structure of Apo-GAFab
(PDB ID: 9A0L, PDB-Dev ID: PDBDEV_00000057)**

1. Model Composition	
Entry composition	GAFab: chain(s) A, B (399 residues)
Datasets used for modeling	<ul style="list-style-type: none"> - Crosslinking-MS data, PRIDE: PXD020817 - Experimental model, PDB: 6X88
2. Representation	
Number of representations	1
Scale	Atomic
Number of rigid and flexible segments	0, 2
3. Restraints	
Physical principles	Information about physical principles was not provided
Experimental data	<ul style="list-style-type: none"> - 1 unique CrossLinkRestraint: sulfo-SDA, 55 crosslinks - 1 unique CrossLinkRestraint: BS3, 22 crosslinks - 1 unique CrossLinkRestraint: EDC, 7 crosslinks
4. Validation	
Number of ensembles	0
Number of models in ensembles	Not applicable
Number of deposited models	1
Model precision (uncertainty of models)	Not available
Data quality	Data quality has not been assessed
Model quality: assessment of atomic segments	<ul style="list-style-type: none"> - Clashscore: 0.00 - Ramachandran outliers: 42 - Sidechain outliers: 30
Fit to data used for modeling	Satisfaction of crosslinks: 98.81%
Fit to data used for validation	Fit of model to information not used to compute it has not been determined
5. Methodology and Software	
1. Name	None

Software

- [Integrative Modeling Platform \(IMP\)](#) (version Not available)
- [Modeller](#) (version Not available)