

# **Full wwPDB Integrative Structure Validation Report**

## August 20, 2019 -- 05:25 PM

PDB ID	PDBDEV00000005
Molecule Name	Serum Albumin Domain A Structure
Title	Serum Albumin Domain Structures in Human Blood Serum by Mass Spectrometry and Computational E
Authors	Belsom A;Schneider M;Fischer L;Brock O;Rappsilber J

#### The following softwares were used in the production of this report:

Integrative Modeling Package: Version XX
Molprobity: Version XX
Phenix: Version XX
Integrative Modeling Validation Package: Version XX

### 1. Overall quality at a glance

### 2. Entry composition

There are 1 unique types of models in this entry. The entry contains 1 chains.

Molecule ID	Molecule Name	Chain ID	Total Residues
1	HSA_A	Α	197

There are 2 software packages reported in this entry.

ID	Software Name	Software Version	Software Classification
1	Rosetta MBS	None	Model Building
2	EPC-map	None	Contact Predictor

There are 2 unique datasets used to build the model(s) in this entry.

ID	Dataset Type	Database Name	Data Access Code
1	CX-MS data	PRIDE	PXD001692
2	unspecified	Not Listed	None

3. Data quality
4. Model quality
4.1 Too-close contacts
4.2 Torsion angles
4.2.1 Protein backbone
4.2.2 Protein sidechains
5. Fit of model and data
6. Uncertaintiy of model