

# Full wwPDB Integrative Structure Validation Report

September 06, 2019 -- 02:06 PM

PDB ID	PDBDEV00000007
Molecule Name	Serum Albumin Domain C Structure
Title	Serum Albumin Domain Structures in Human Blood Serum by Mass Spectrometry and Computational E
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**The following softwares were used in the production of this report:**

*Integrative Modeling Package : Version XX*  
*Molprobit : Version XX*  
*Phenix : Version XX*  
*Integrative Modeling Validation Package : Version XX*

## 1. Overall quality at a glance

## 2. Entry composition

There is 1 unique type of model in this entry. Model has 1 chains respectively.

Molecule ID	Molecule Name	Chain ID	Total Residues
1	HSA_C	A	192

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 in this entry.

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

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### **3. Data quality**

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### **4. Model quality**

#### **4.1 Too-close contacts**

#### **4.2 Torsion angles**

##### **4.2.1 Protein backbone**

##### **4.2.2 Protein sidechains**

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### **5. Fit of model and data**

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### **6. Uncertainty of model**

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