

# Full wwPDB Integrative Structure Validation Report

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PDB ID	PDBDEV00000027
Molecule Name	A metastable contact and structural disorder in the estrogen receptor transactivation domain
Title	A metastable contact and structural disorder in the estrogen receptor transactivation domain
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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	Estrogen receptor	A	184

There is 1 software package reported in this entry.

ID	Software Name	Software Version	Software Classification
1	iSPOT	None	model building

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

ID	Dataset Type	Database Name	Data Access Code
1	SAS data	SASBDB	SASDEE2
2	unspecified	Not Listed	None

### **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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