



Full wwPDB Integrative Structure Validation Report

November 07, 2019 -- 03:08 PM

<i>PDB ID</i>	<i>PDBDEV00000022</i>
Molecule Name	Structural dynamics of the E6AP/UBE3A-E6-p53 enzyme-substrate complex
Title	Structural dynamics of the E6AP/UBE3A-E6-p53 enzyme-substrate complex
Authors	Carolyn Sailer;Fabian Offensperger;Alexandra Julier;Kai-Michael Kammer;Ryan Walker-Gray;Matthew G. Gold;Martin Scheffner;Florian Stengel

The following software were used in the production of this report:

Molprobit : Version 4.4
Integrative Modeling Validation Package : Version 1.0

1. Overall quality at a glance

2. Ensemble information

This entry consists of 1 distinct ensemble.

<i>Ensemble number</i>	<i>Ensemble name</i>	<i>Model ID</i>	<i>Number of models</i>	<i>Clustering method</i>	<i>Clustering feature</i>	<i>Cluster precision</i>
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1	E6 main cluster	1	500	Other	RMSD	None
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3. Model composition

3.1 Summary

This entry consists of 1 unique models, with 2 subunits in each model. A total of 4 datasets or restraints was used to build this entry. Each model is represented by 2 rigid bodies and 0 flexible or non-rigid units.

3.2 Entry composition

There is 1 unique type of model in this entry. This model is titled E6AP E6/None respectively.

<i>Model ID</i>	<i>Subunit number</i>	<i>Subunit ID</i>	<i>Subunit name</i>	<i>Chain ID</i>	<i>Total residues</i>
1	1	1	E6AP HECT Domain	A	350
1	2	2	E6	B	143

3.3 Datasets used for modeling

There are 4 unique datasets used to build the models in this entry.

<i>ID</i>	<i>Dataset type</i>	<i>Database name</i>	<i>Data access code</i>
1	CX-MS data	Not listed	None
2	Comparative model	Not listed	None
3	Experimental model	PDB	1C4Z
4	Experimental model	PDB	4XR8

4. Representation

This entry has only one representation and includes 2 rigid bodies and 0 flexible units.

<i>Chain ID</i>	<i>Rigid bodies</i>	<i>Non-rigid segments</i>
A	497-846:Comparative model/None.	-
B	1-143:Comparative model/None.	-

5. Methodology and software

<i>Step number</i>	<i>Protocol ID</i>	<i>Method name</i>	<i>Method type</i>	<i>Number of computed models</i>	<i>Multi state modeling</i>	<i>Multi scale modeling</i>
1	1	IMP	MC based Bayesian sampling using crosslinks	720000	False	False

There is 1 software package reported in this entry.

<i>ID</i>	<i>Software name</i>	<i>Software version</i>	<i>Software classification</i>	<i>Software location</i>
1	Integrative Modeling Platform (IMP)	git checkout 2018/01/08 (commit 5eb8151c651256d50bbcd847932bc913df94090c)	integrative model building	https://integrativemodeling.org

6. Data quality

7. Model quality

8. Fit of model to data used for modeling

9. Fit of model to data not used for modeling

10. Uncertainty of model