

**Summary of integrative structure determination of Dimer structure of the solute carrier
SLC26Dg (PDB ID: 8ZZV, PDB-Dev ID: PDBDEV_0000031)**

1. Model Composition	
Entry composition	- SLC26Dg: Chain A (379 residues) - SLC26Dg: Chain B (379 residues)
Datasets used for modeling	- Experimental model, PDB ID: 5DA0 - Other, File: 10.5281/zenodo.2638061
2. Representation	
Resolution	Atomic
Number of rigid bodies, flexible units	4, 2
Rigid bodies	- A: 1-333, 338-379 - B: -
Flexible units	- A: 334-337 - B: 334-337
Structural coverage (rigid bodies)	100%
3. Restraints	
Physical principles	Information about physical principles was not provided
Experimental data	
4. Validation	
Number of ensembles	0
Number of models in ensembles	Not applicable
Number of deposited models	1
Model precision (uncertainty of models)	Model precision can not be calculated with one structure
Data quality	Data quality has not been assessed
Model quality: assessment of atomic segments	Model-1: Clashscore = 0.86, Number of Ramachandran outliers = 10, Number of sidechain outliers = 17
Model quality: assessment of excluded volume	Not applicable
Fit to data used for modeling	Fit of model to information used to compute it has not been determined

<i>Fit to data used for validation</i>	Fit of model to information not used to compute it has not been determined
5. Methodology and Software	
1. <i>Method</i>	?
<i>Name</i>	?
<i>Number of computed models</i>	?
<i>Software</i>	BioEn (version Not available)