

**Summary of integrative structure determination of NMR Structure of  
Sa1\_V90T at 5 Degrees Celsius (PDB ID: 9A20, PDB-Dev ID:  
PDBDEV\_00000132)**

<b>1. Model Composition</b>	
<a href="#">Entry composition</a>	3-alpha NMR structure of Sa1_V90T at 5 degrees Celsius: chain(s) A (95 residues)
<a href="#">Datasets used for modeling</a>	- NMR data, BMRB: <a href="#">51338</a>
<b>2. Representation</b>	
<a href="#">Number of representations</a>	1
<a href="#">Scale</a>	Atomic
Number of <a href="#">rigid</a> and <a href="#">flexible</a> segments	0, 1
<b>3. Restraints</b>	
<a href="#">Physical principles</a>	Information about physical principles was not provided
<a href="#">Experimental data</a>	
<b>4. Validation</b>	
<a href="#">Number of ensembles</a>	0
<a href="#">Number of models in ensembles</a>	Not applicable
<a href="#">Number of deposited models</a>	10
<a href="#">Model precision (uncertainty of models)</a>	Not available
<a href="#">Data quality</a>	Data quality has not been assessed
<a href="#">Model quality: assessment of atomic segments</a>	<ul style="list-style-type: none"> <li>- Clashscore: 0.00-7.12</li> <li>- Ramachandran outliers: 0-1</li> <li>- Sidechain outliers: 0-2</li> </ul>
<a href="#">Fit to data used for modeling</a>	Fit of model to information used to compute it has not been determined
<a href="#">Fit to data used for validation</a>	Fit of model to information not used to compute it has not been determined
<b>5. Methodology and Software</b>	
1. <a href="#">Name</a>	None
<a href="#">Software</a>	<a href="#">CS-Rosetta</a> (version Not available)