



## Full wwPDB EM Map Validation Report ⓘ

Dec 9, 2020 – 12:29 pm GMT

EMDB ID : EMD-5217  
Title : Visualizing the structural changes of bacteriophage epsilon15 and its Salmonella host during infection  
Authors : , Chang.JT.; , Schmid.MF.; , Haase-Pettingell.C.; , Weigele.PR.; , King.JA.; , Chiu.W.  
Deposited on : 2010-07-22  
Resolution : Not provided

This is a Full wwPDB EM Map Validation Report for a publicly released PDB entry.

We welcome your comments at [validation@mail.wwpdb.org](mailto:validation@mail.wwpdb.org)

A user guide is available at

<https://www.wwpdb.org/validation/2017/EMMapValidationReportHelp>  
with specific help available everywhere you see the ⓘ symbol.

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The following versions of software and data (see [references ⓘ](#)) were used in the production of this report:

EMDB validation analysis : **FAILED**  
Validation Pipeline (wwPDB-VP) : 2.13

# 1 Experimental information

Property	Value	Source
EM reconstruction method	subtomogramAveraging	Depositor
Imposed symmetry	Not Provided	Depositor
Number of images used	Not provided	Depositor
Resolution determination method	Not provided	Depositor
CTF correction method	Not provided	Depositor
Microscope	JEOL 3200FSC	Depositor
Voltage (kV)	300	Depositor
Electron dose ( $e^-/\text{\AA}^2$ )	65	Depositor
Minimum defocus (nm)	6.0	Depositor
Maximum defocus (nm)	9.0	Depositor
Magnification	20000.0	Depositor
Image detector	GENERIC CCD	Depositor