



## Full wwPDB EM Map Validation Report ⓘ

Dec 9, 2020 – 12:41 pm GMT

EMDB ID : EMD-5745  
Title : Zernike Phase Contrast electron cryo-tomography of cyanophage Syn5 assembly intermediates  
Authors : , Dai.W.; , Fu.C.; , Raytcheva.D.; , Flanagan.J.; , Khant.HA.; , Liu.X.; , Rochat.RH.; , Haase-Pettingell.C.; , Piret.J.; , Ludtke.SJ.; , Nagayama.K.; , Schmid.MF.; , King.JA.; , Chiu.W.  
Deposited on : 2013-08-13  
Resolution : 47.00 Å(reported)

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	137	Depositor
Resolution determination method	FSC 0.143 CUT-OFF	Depositor
CTF correction method	Not provided	Depositor
Microscope	JEOL 2200FS	Depositor
Voltage (kV)	200	Depositor
Electron dose ( $e^-/\text{\AA}^2$ )	1	Depositor
Minimum defocus (nm)	0.0	Depositor
Maximum defocus (nm)	1.0	Depositor
Magnification	25000.0	Depositor
Image detector	GATAN ULTRASCAN 4000 (4k x 4k)	Depositor