



Full wwPDB EM Map Validation Report ⓘ

Dec 9, 2020 – 01:59 pm GMT

EMDB ID : EMD-9392
Title : Bacteriophage L virion
Authors : , Parent.KN.; , Schrad.JR.
Deposited on : 2019-01-04
Resolution : 4.20 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

A user guide is available at

<https://www.wwpdb.org/validation/2017/EMMapValidationReportHelp>

with specific help available everywhere you see the ⓘ symbol.

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	singleParticle	Depositor
Imposed symmetry	POINT, I	Depositor
Number of images used	7879	Depositor
Resolution determination method	FSC 0.143 CUT-OFF	Depositor
CTF correction method	estimated manually	Depositor
Microscope	FEI TITAN KRIOS	Depositor
Voltage (kV)	300	Depositor
Electron dose ($e^-/\text{\AA}^2$)	27.0	Depositor
Minimum defocus (nm)	0.35	Depositor
Maximum defocus (nm)	2.49	Depositor
Magnification	Not provided	Depositor
Image detector	DIRECT ELECTRON DE-20 (5k x 3k)	Depositor