



wwPDB EM Map Validation Summary Report ⓘ

Dec 9, 2020 – 01:31 pm GMT

EMDB ID : EMD-8360
Title : Asymmetric reconstruction of bacteriophage MS2
Authors : , Nazarov.SU.; , Guerrero-Ferreira.RC.; , Zhong.Q.; , Kohn.T.; , Leiman.PG.
Deposited on : 2016-08-30
Resolution : 10.50 Å(reported)

This is a wwPDB EM Map Validation Summary 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, C1	Depositor
Number of images used	29719	Depositor
Resolution determination method	FSC 0.143 CUT-OFF	Depositor
CTF correction method	Not provided	Depositor
Microscope	FEI TECNAI F20	Depositor
Voltage (kV)	200	Depositor
Electron dose ($e^-/\text{\AA}^2$)	25.0	Depositor
Minimum defocus (nm)	0.8	Depositor
Maximum defocus (nm)	3.0	Depositor
Magnification	Not provided	Depositor
Image detector	FEI EAGLE (4k x 4k)	Depositor