



Full wwPDB EM Map Validation Report ⓘ

Dec 9, 2020 – 12:51 pm GMT

EMDB ID : EMD-6232
Title : Negative stain single particle electron microscopy of Marburg Virus Glycoproteins bound to antibodies from a human survivor
Authors : , Flyak.A.; , Ilinykh.PA.; , Murin.CD.; , Garron.T.; , Shen.X.; , Fusco.ML.; , Hashiguchi.T.; , Bornholdt.ZA.; , Slaughter.C.; , Sapparapu.G.; , Ksiazek.TG.; , Ward.AB.; , Sapphire.EO.; , Bukreyev.A.; , Crowe.JE.
Deposited on : 2015-01-06
Resolution : 27.00 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

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	Not Provided	Depositor
Number of images used	978	Depositor
Resolution determination method	OTHER	Depositor
CTF correction method	Not provided	Depositor
Microscope	FEI TECNAI SPIRIT	Depositor
Voltage (kV)	120	Depositor
Electron dose ($e^-/\text{\AA}^2$)	Not provided	Depositor
Minimum defocus (nm)	1.0	Depositor
Maximum defocus (nm)	1.0	Depositor
Magnification	52000.0	Depositor
Image detector	OTHER	Depositor