



Full wwPDB EM Map Validation Report ⓘ

Dec 9, 2020 – 12:35 pm GMT

EMDB ID : EMD-5487
Title : Rotavirus VP6 protein
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Veesler.D.; , Pan.J.; , Harrison.SC.; , Potter.CS.; , Carragher.B.; , Grigori-
eff.N.
Deposited on : 2012-08-25
Resolution : 4.90 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	807	Depositor
Resolution determination method	FSC 0.143 CUT-OFF	Depositor
CTF correction method	Each particle	Depositor
Microscope	FEI TECNAI F20	Depositor
Voltage (kV)	200	Depositor
Electron dose ($e^-/\text{\AA}^2$)	32	Depositor
Minimum defocus (nm)	1.0	Depositor
Maximum defocus (nm)	2.4	Depositor
Magnification	42135.0	Depositor
Image detector	DIRECT ELECTRON DE-12 (4k x 3k)	Depositor