



Full wwPDB EM Map Validation Report ⓘ

Dec 7, 2020 – 03:44 pm GMT

EMDB ID : EMD-0354
Title : CryoEM structure of Leviviridae PP7 coat protein dimer capsid with a loop insertion and a C-terminal extension (PP7-aloop-PP7-150loop)
Authors : Liangjun, Z.; Kopylov, M.; Carragher, B.; Potter, C.S.; Finn, M.G.
Deposited on : 2018-11-27
Resolution : 3.43 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	SINGLE PARTICLE	Depositor
Imposed symmetry	POINT, 1	Depositor
Number of particles used	3066	Depositor
Resolution determination method	FSC 0.143 CUT-OFF	Depositor
CTF correction method	Not provided	Depositor
Microscope	FEI TITAN KRIOS	Depositor
Voltage (kV)	300	Depositor
Electron dose ($e^-/\text{\AA}^2$)	35.0	Depositor
Minimum defocus (nm)	-0.0007	Depositor
Maximum defocus (nm)	0.002	Depositor
Magnification	130000.	Depositor
Image detector	GATAN K2 SUMMIT (4k x 4k)	Depositor