



Full wwPDB EM Map Validation Report ⓘ

Dec 9, 2020 – 01:53 pm GMT

EMDB ID : EMD-9059
Title : CryoEM structure of human enterovirus D68 A-particle (pH 6.0, room temperature, 30 sec)
Authors : , Liu.Y.; , Rossmann.MG.
Deposited on : 2018-08-27
Resolution : 3.46 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, 1 | Depositor |
| Number of images used | 14000 | Depositor |
| Resolution determination method | FSC 0.143 CUT-OFF | Depositor |
| CTF correction method | On-the-fly CTF correction during 2D alignment and 3D reconstruction | Depositor |
| Microscope | FEI TITAN KRIOS | Depositor |
| Voltage (kV) | 300 | Depositor |
| Electron dose ($e^-/\text{\AA}^2$) | 28.0 | Depositor |
| Minimum defocus (nm) | 1.1 | Depositor |
| Maximum defocus (nm) | 5.4 | Depositor |
| Magnification | 81000. | Depositor |
| Image detector | GATAN K2 SUMMIT (4k x 4k) | Depositor |