



Full wwPDB EM Map Validation Report ⓘ

Dec 9, 2020 – 01:20 pm GMT

EMDB ID : EMD-7861
Title : Full length HIV-1 Envelope Glycoprotein clone PC64M4C054 in complex with
PGT151 Fab and PCT64-13C Fab
Authors : , Berndsen.ZT.; , Rantalainen.K.; , Ward.AB.
Deposited on : 2018-05-09
Resolution : 30.00 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 | POINT, C1 | Depositor |
| Number of images used | 2395 | Depositor |
| Resolution determination method | FSC 0.5 CUT-OFF | Depositor |
| CTF correction method | Not provided | Depositor |
| Microscope | FEI TECNAI SPIRIT | Depositor |
| Voltage (kV) | 120 | Depositor |
| Electron dose ($e^-/\text{\AA}^2$) | 25.0 | Depositor |
| Minimum defocus (nm) | Not provided | Depositor |
| Maximum defocus (nm) | Not provided | Depositor |
| Magnification | Not provided | Depositor |
| Image detector | TVIPS TEMCAM-F416 (4k x 4k) | Depositor |