



## Full wwPDB EM Map Validation Report ⓘ

Dec 9, 2020 – 12:49 pm GMT

EMDB ID : EMD-6080  
Title : Cryo electron tomography of infective bacteriophage T4  
Authors : , Hu.B.; , Margolin.W.; , Molineux.IJ.; , Liu.J.  
Deposited on : 2014-09-01  
Resolution : 40.00 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](mailto: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.

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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             | subtomogramAveraging        | Depositor |
| Imposed symmetry                     | Not Provided                | Depositor |
| Number of images used                | 1985                        | Depositor |
| Resolution determination method      | OTHER                       | Depositor |
| CTF correction method                | Not provided                | Depositor |
| Microscope                           | FEI POLARA 300              | Depositor |
| Voltage (kV)                         | 300                         | Depositor |
| Electron dose ( $e^-/\text{\AA}^2$ ) | 100                         | Depositor |
| Minimum defocus (nm)                 | 4.0                         | Depositor |
| Maximum defocus (nm)                 | 6.0                         | Depositor |
| Magnification                        | 23000.0                     | Depositor |
| Image detector                       | TVIPS TEMCAM-F415 (4k x 4k) | Depositor |