



## Full wwPDB EM Map Validation Report ⓘ

Dec 9, 2020 – 01:40 pm GMT

EMDB ID : EMD-9040  
Title : Analysis of bacteriophage phiX174 tomograms using the locked self-rotation function  
Authors : , Fokine.A.; , Sun.Y.; , Khare.B.; , Rossmann.MG.  
Deposited on : 2018-08-13  
Resolution : Not provided

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.

---

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	tomography	Depositor
Imposed symmetry	Not Provided	Depositor
Number of images used	41	Depositor
Resolution determination method	Not provided	Depositor
CTF correction method	Not provided	Depositor
Microscope	FEI TITAN KRIOS	Depositor
Voltage (kV)	300	Depositor
Electron dose ( $e^-/\text{\AA}^2$ )	1.73	Depositor
Minimum defocus (nm)	3.0	Depositor
Maximum defocus (nm)	Not provided	Depositor
Magnification	11000.0	Depositor
Image detector	GATAN K2 SUMMIT (4k x 4k)	Depositor