



## Full wwPDB EM Map Validation Report ⓘ

Dec 9, 2020 – 01:12 pm GMT

EMDB ID : EMD-7486  
Title : The tether and tether head complex and I1 dynein complex region extracted from the cryo-electron tomography and subtomographic average (1885 axone-mal repeats) of isolated Chlamydomonas fap44 mutant cilia  
Authors : , Fu.G.; , Wang.Q.; , Phan.N.; , Urbanska.P.; , Joachimiak.E.; , Lin.J.; , Wloga.D.; , Nicastro.D.  
Deposited on : 2018-03-01  
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	POINT, C1	Depositor
Number of images used	1885	Depositor
Resolution determination method	FSC 0.5 CUT-OFF	Depositor
CTF correction method	Not provided	Depositor
Microscope	FEI TECNAI F30	Depositor
Voltage (kV)	300	Depositor
Electron dose ( $e^-/\text{\AA}^2$ )	1.5	Depositor
Minimum defocus (nm)	6.0	Depositor
Maximum defocus (nm)	8.0	Depositor
Magnification	13500.	Depositor
Image detector	GATAN ULTRASCAN 1000 (2k x 2k)	Depositor