



## wwPDB EM Map Validation Summary Report ⓘ

Dec 9, 2020 – 11:40 pm GMT

EMDB ID : EMD-6914  
Title : Serial cryotomogram of a metaphase budding yeast cell.  
Authors : , Ng.C.; , Deng.L.; , Chen.C.; , Lim.H.; , Shi.J.; , Surana.U.; , Gan.L.  
Deposited on : 2018-02-22  
Resolution : Not provided

This is a wwPDB EM Map Validation Summary 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	tomography	Depositor
Imposed symmetry	Not Provided	Depositor
Number of images used	61	Depositor
Resolution determination method	Not provided	Depositor
CTF correction method	Phase flipping done in Etomo 4.10	Depositor
Microscope	FEI TITAN KRIOS	Depositor
Voltage (kV)	300	Depositor
Electron dose ( $e^-/\text{\AA}^2$ )	1.6	Depositor
Minimum defocus (nm)	10.0	Depositor
Maximum defocus (nm)	Not provided	Depositor
Magnification	15678.	Depositor
Image detector	FEI FALCON II (4k x 4k)	Depositor