



wwPDB EM Map Validation Summary Report ⓘ

Dec 9, 2020 – 01:26 pm GMT

EMDB ID : EMD-8239
Title : EM map of the intact yeast Elongator complex
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Deposited on : 2016-06-08
Resolution : 25.20 Å(reported)

This is a wwPDB EM Map Validation Summary 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	Not Provided	Depositor
Number of images used	8190	Depositor
Resolution determination method	FSC 0.143 CUT-OFF	Depositor
CTF correction method	CTF parameters were calculated on entire micrographs prior to any data processing using CTFFIND within SPIDER. Phase-flip correction was done in SPIDER.	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	49000.	Depositor
Image detector	FEI EAGLE (4k x 4k)	Depositor