



Full wwPDB EM Map Validation Report ⓘ

Dec 9, 2020 – 01:27 pm GMT

EMDB ID : EMD-8291
Title : EM map of the Elp123 subcomplex of yeast Elongator
Authors : , Setiaputra.D.; , Cheng.DTH.; , Hansen.JM.; , Yip.CK.
Deposited on : 2016-07-16
Resolution : 24.10 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

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	16362	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