



wwPDB EM Map Validation Summary Report ⓘ

Dec 9, 2020 – 01:05 pm GMT

EMDB ID : EMD-7005
Title : Negative stain reconstruction of the peroxisomal AAA-ATPase Pex1/Pex6 complex associated with substrate Pex15
Authors : , Chowdhury.S.; , Gardner.BM.; , Castanzo.DT.; , Stjepanovic.G.; , Stefely.MS.; , Hurley.JH.; , Martin.A.; , Lander.GC.
Deposited on : 2017-09-01
Resolution : 23.20 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	POINT, C1	Depositor
Number of images used	14678	Depositor
Resolution determination method	FSC 0.143 CUT-OFF	Depositor
CTF correction method	EMAN1 was used for phase flipping whole micrograph before particle extraction.	Depositor
Microscope	FEI TECNAI SPIRIT	Depositor
Voltage (kV)	120	Depositor
Electron dose ($e^-/\text{\AA}^2$)	20.0	Depositor
Minimum defocus (nm)	0.5	Depositor
Maximum defocus (nm)	1.5	Depositor
Magnification	52000.0	Depositor
Image detector	TVIPS TEMCAM-F416 (4k x 4k)	Depositor