



Full wwPDB EM Map Validation Report ⓘ

Dec 9, 2020 – 12:38 pm GMT

EMDB ID : EMD-5641
Title : cryo-EM structure of CCT5 complex with 1mM ATP/AlFx (symmetry-free)
Authors : , Sergeeva.OA.; , Chen.B.; , Haase-Pettingell.C.; , Ludtke.SJ.; , Chiu.W.; , King.JA.
Deposited on : 2013-04-14
Resolution : Not provided

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	2974	Depositor
Resolution determination method	Not provided	Depositor
CTF correction method	each frame	Depositor
Microscope	JEOL 2010F	Depositor
Voltage (kV)	200	Depositor
Electron dose ($e^-/\text{\AA}^2$)	20	Depositor
Minimum defocus (nm)	1.5	Depositor
Maximum defocus (nm)	3.5	Depositor
Magnification	71361.0	Depositor
Image detector	GENERIC GATAN (4k x 4k)	Depositor