



Full wwPDB EM Map Validation Report ⓘ

Dec 9, 2020 – 12:55 pm GMT

EMDB ID : EMD-6318
Title : Three-dimensional structure of AcrAB and MacA-TolC alpha-hybrid dimer complex
Authors : , Kim.J-S.; , Jeong.H.; , Song.S.; , Kim.H-Y.; , Lee.K.; , Hyun.J.; , Ha.N-C.
Deposited on : 2015-04-15
Resolution : 24.00 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	1563	Depositor
Resolution determination method	OTHER	Depositor
CTF correction method	CTF correction to each particle	Depositor
Microscope	FEI TECNAI SPIRIT	Depositor
Voltage (kV)	120	Depositor
Electron dose ($e^-/\text{\AA}^2$)	Not provided	Depositor
Minimum defocus (nm)	0.7	Depositor
Maximum defocus (nm)	1.2	Depositor
Magnification	66667.0	Depositor
Image detector	GATAN ULTRASCAN 4000 (4k x 4k)	Depositor