



Full wwPDB EM Map Validation Report ⓘ

Dec 9, 2020 – 01:24 pm GMT

EMDB ID : EMD-8066
Title : HKU1 spike with attached foldon domain and wild-type furin-cleavage site
Authors : , Kirchdoerfer.RN.; , Cottrell.CA.; , Wang.N.; , Pallesen.J.; , Yassine.HM.; ,
Turner.HL.; , Corbett.KS.; , Graham.BS.; , McLellan.JS.; , Ward.AB.
Deposited on : 2016-02-02
Resolution : 20.00 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	POINT, C3	Depositor
Number of images used	10039	Depositor
Resolution determination method	FSC 0.5 CUT-OFF	Depositor
CTF correction method	Not provided	Depositor
Microscope	FEI TECNAI SPIRIT	Depositor
Voltage (kV)	120	Depositor
Electron dose ($e^-/\text{\AA}^2$)	28.0	Depositor
Minimum defocus (nm)	1.0	Depositor
Maximum defocus (nm)	1.5	Depositor
Magnification	52000.	Depositor
Image detector	TVIPS TEMCAM-F416 (4k x 4k)	Depositor