



## Full wwPDB EM Map Validation Report ⓘ

Dec 9, 2020 – 12:38 pm GMT

EMDB ID : EMD-5683  
Title : Molecular structure of the native influenza hemagglutinin H1 on virions  
Authors : , Harris.AK.; , Meyerson.JR.; , Matsuoka.Y.; , Kuybeda.O.; , Moran.A.; , Bliss.D.; , Das.SR.; , Yewdell.JW.; , Sapiro.G.; , Subbarao.K.; , Subramaniam.S.  
Deposited on : 2013-06-03  
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](mailto: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.

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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	subtomogramAveraging	Depositor
Imposed symmetry	Not Provided	Depositor
Number of images used	416	Depositor
Resolution determination method	Not provided	Depositor
CTF correction method	Not provided	Depositor
Microscope	FEI POLARA 300	Depositor
Voltage (kV)	200	Depositor
Electron dose ( $e^-/\text{\AA}^2$ )	100	Depositor
Minimum defocus (nm)	2.5	Depositor
Maximum defocus (nm)	2.5	Depositor
Magnification	34000.0	Depositor
Image detector	GENERIC CCD	Depositor