



Full wwPDB EM Map Validation Report ⓘ

Dec 9, 2020 – 12:43 pm GMT

EMDB ID : EMD-5255
Title : Seeing the Portal in Herpes Simplex Virus type I B-capsids. Map1: Icosahedral reconstruction of the HSV-1 B-capsid Map2: symmetry free (c1) reconstruction of the HSV-1 B-capsid Map3: c12 symmetrized portal density extracted from symmetry free (c1) reconstruction of the HSV-1 B-capsid Map4: c1 portal density extracted from symmetry free (c1) reconstruction of the HSV-1 B-capsid
Authors : , Rochat.RH.; , Liu.X.; , Murata.K.; , Nagayama.K.; , Rixon.FJ.; , Chiu.W.
Deposited on : 2010-12-15
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	6300	Depositor
Resolution determination method	FSC 0.5 CUT-OFF	Depositor
CTF correction method	No Correction	Depositor
Microscope	JEOL 2200FS	Depositor
Voltage (kV)	200	Depositor
Electron dose ($e^-/\text{\AA}^2$)	20	Depositor
Minimum defocus (nm)	0.0	Depositor
Maximum defocus (nm)	0.2	Depositor
Magnification	56000.0	Depositor
Image detector	GENERIC TVIPS (4k x 4k)	Depositor