



Full wwPDB EM Map Validation Report ⓘ

Dec 9, 2020 – 01:44 pm GMT

EMDB ID : EMD-8716
Title : Cryo-EM reconstruction of B41 SOSIP.664 in complex with fragment antigen binding of PGV04
Authors : , Pallesen.J.; , Ozorowski.G.; , de.Val.N.; , Ward.AB.
Deposited on : 2017-04-28
Resolution : 7.40 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	9475	Depositor
Resolution determination method	FSC 0.143 CUT-OFF	Depositor
CTF correction method	Not provided	Depositor
Microscope	FEI TECNAI F20	Depositor
Voltage (kV)	200	Depositor
Electron dose ($e^-/\text{\AA}^2$)	34.0	Depositor
Minimum defocus (nm)	1.0	Depositor
Maximum defocus (nm)	4.5	Depositor
Magnification	41322.	Depositor
Image detector	GATAN K2 SUMMIT (4k x 4k)	Depositor