



Full wwPDB EM Map Validation Report ⓘ

Dec 9, 2020 – 12:38 pm GMT

EMDB ID : EMD-5462
Title : Cryo-electron tomography of a trimeric soluble HIV Env construct, gp140
SOSIP, in complex with Fab 17b
Authors : , Tran.EEH.; , Borgnia.MJ.; , Kuybeda.O.; , Schauder.DM.; , Bartesaghi.A.; ,
Frank.GA.; , Sapiro.G.; , Milne.JLS.; , Subramaniam.S.
Deposited on : 2012-08-06
Resolution : 8.80 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	subtomogramAveraging	Depositor
Imposed symmetry	Not Provided	Depositor
Number of images used	39001	Depositor
Resolution determination method	FSC 0.143 CUT-OFF	Depositor
CTF correction method	Not provided	Depositor
Microscope	FEI TITAN KRIOS	Depositor
Voltage (kV)	80	Depositor
Electron dose ($e^-/\text{\AA}^2$)	20	Depositor
Minimum defocus (nm)	1.5	Depositor
Maximum defocus (nm)	5.0	Depositor
Magnification	138888.0	Depositor
Image detector	GATAN ULTRASCAN 4000 (4k x 4k)	Depositor