



wwPDB EM Map Validation Summary Report ⓘ

Dec 9, 2020 – 12:54 pm GMT

EMDB ID : EMD-6426
Title : Negative-stain electron microscopy of a full-length transmembrane receptor tyrosine kinase (PDGFR) in complex with PDGFB
Authors : , Chen.PH.; , Unger.VM.; , He.X.
Deposited on : 2015-08-12
Resolution : 27.00 Å(reported)

This is a wwPDB EM Map Validation Summary 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	4234	Depositor
Resolution determination method	OTHER	Depositor
CTF correction method	Not provided	Depositor
Microscope	OTHER	Depositor
Voltage (kV)	120	Depositor
Electron dose ($e^-/\text{\AA}^2$)	Not provided	Depositor
Minimum defocus (nm)	0.5	Depositor
Maximum defocus (nm)	2.0	Depositor
Magnification	30000.0	Depositor
Image detector	GENERIC CCD	Depositor