



Full wwPDB EM Map Validation Report ⓘ

Dec 9, 2020 – 12:41 pm GMT

EMDB ID : EMD-5341
Title : Chemically controlled assembly of 1, 2 and 3-dimensional protein arrays
Authors : , Brodin.JD.; , Ambroggio.XI.; , Tang.C.; , Parent.KN.; , Baker.TS.; , Tezcan.FA.
Deposited on : 2011-09-06
Resolution : 10.00 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	helical	Depositor
Imposed symmetry	HELICAL, twist=5.3°, rise=25.7 Å, axial sym=C9	Depositor
Number of images used	Not provided	Depositor
Resolution determination method	OTHER	Depositor
CTF correction method	ROBEM and IHRSR	Depositor
Microscope	FEI POLARA 300	Depositor
Voltage (kV)	200	Depositor
Electron dose ($e^-/\text{\AA}^2$)	15	Depositor
Minimum defocus (nm)	1.75	Depositor
Maximum defocus (nm)	3.675	Depositor
Magnification	52027.0	Depositor
Image detector	GENERIC GATAN (4k x 4k)	Depositor