



wwPDB EM Map Validation Summary Report (i)

Dec 9, 2020 – 11:21 am GMT

EMDB ID : EMD-21313

Title : Cryo-EM structure of a substrate-engaged Bam complex with alternative cysteine-cysteine crosslink

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Deposited on : 2020-02-03

Resolution : 6.50 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 (i) symbol.

The following versions of software and data (see [references \(1\)](#)) were used in the production of this report:

EMDB validation analysis : FAILED

Validation Pipeline (wwPDB-VP) : 2.13

1 Experimental information i

Property	Value	Source
EM reconstruction method	singleParticle	Depositor
Imposed symmetry	POINT, C1	Depositor
Number of images used	233064	Depositor
Resolution determination method	FSC 0.143 CUT-OFF	Depositor
CTF correction method	Not provided	Depositor
Microscope	FEI TITAN KRIOS	Depositor
Voltage (kV)	300	Depositor
Electron dose ($e^-/\text{\AA}^2$)	55.0	Depositor
Minimum defocus (nm)	1.1	Depositor
Maximum defocus (nm)	3.2	Depositor
Magnification	58717.0	Depositor
Image detector	GATAN K3 BIOQUANTUM (6k x 4k)	Depositor