



Full wwPDB EM Validation Report ⓘ

Mar 3, 2025 – 12:28 pm GMT

EMDB ID : EMD-52523
Title : CryoEM structure of F-ENA fibers on the spores of *Bacillus thuringiensis* serovar *kurstaki*
Authors : Sleutel, M.; Remaut, H.
Deposited on : 2025-01-13
Resolution : 3.23 Å(reported)

This is a Full wwPDB EM 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 types of validation reports are described at

<http://www.wwpdb.org/validation/2017/FAQs#types>.

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) : 2.41

1 Experimental information

Property	Value	Source
EM reconstruction method	HELICAL	Depositor
Imposed symmetry	HELICAL, twist=42.73°, rise=33.54 Å, axial sym=C3	Depositor
Number of segments used	196540	Depositor
Resolution determination method	FSC 0.143 CUT-OFF	Depositor
CTF correction method	Not provided	
Microscope	JEOL CRYO ARM 300	Depositor
Voltage (kV)	300	Depositor
Electron dose ($e^-/\text{\AA}^2$)	60.0	Depositor
Minimum defocus (nm)	1.0	Depositor
Maximum defocus (nm)	2.0	Depositor
Magnification	Not provided	
Image detector	GATAN K3 (6k x 4k)	Depositor