



## wwPDB EM Validation Summary Report ⓘ

Feb 19, 2025 – 12:12 pm GMT

EMDB ID : EMD-17134  
Title : 2.9Angstrom 50S ribosome focused-refined map in chloramphenicol-treated Mycoplasma pneumoniae cells  
Authors : Xue, L.; Spahn, C.; Schacherl, M.; Mahamid, J.  
Deposited on : 2023-04-17  
Resolution : 2.90 Å(reported)

This is a wwPDB EM Validation Summary Report for a publicly released PDB entry.

We welcome your comments at [validation@mail.wwpdb.org](mailto: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	SUBTOMOGRAM AVERAGING	Depositor
Imposed symmetry	POINT, C1	Depositor
Number of subtomograms used	30774	Depositor
Resolution determination method	FSC 0.143 CUT-OFF	Depositor
CTF correction method	Not provided	
Microscope	FEI TITAN KRIOS	Depositor
Voltage (kV)	300	Depositor
Electron dose ( $e^-/\text{\AA}^2$ )	3.34	Depositor
Minimum defocus (nm)	1.0	Depositor
Maximum defocus (nm)	3.25	Depositor
Magnification	64000.0	Depositor
Image detector	GATAN K3 BIOQUANTUM (6k x 4k)	Depositor