



## wwPDB EM Map Validation Summary Report ⓘ

Dec 9, 2020 – 01:03 pm GMT

EMDB ID : EMD-6950  
Title : Subtomogram average of a nucleosome with long linker DNA from a Volta phase contrast cryotomogram of a cryo-FIB lamella of an interphase HeLa cell  
Authors : , Cai.S.; , Bock.D.; , Pilhofer.M.; , Gan.L.  
Deposited on : 2018-04-28  
Resolution : 21.00 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](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.

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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	subtomogramAveraging	Depositor
Imposed symmetry	3D CRYSTAL, $a=193.2$ Å, $b=193.2$ Å, $c=193.2$ Å, $\alpha=90^\circ$ , $\beta=90^\circ$ , $\gamma=90^\circ$ , space group=P1	Depositor
Number of images used	693	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 ( $e^-/\text{\AA}^2$ )	2.0	Depositor
Minimum defocus (nm)	0.0	Depositor
Maximum defocus (nm)	Not provided	Depositor
Magnification	14493.0	Depositor
Image detector	GATAN K2 SUMMIT (4k x 4k)	Depositor