



## Full wwPDB EM Map Validation Report ⓘ

Dec 7, 2020 – 04:26 pm GMT

EMDB ID : EMD-0654  
Title : Structural basis of Dot1L stimulation by histone H2B lysine 120 ubiquitination.  
5.2A reconstruction of Dot1L on H2BK120Ub nucleosome  
Authors : Valencia-Sanchez, M.I.; De Ioannes, P.; Wang, M.; Vasilyev, N.; Chen, R.;  
Nudler, E.; Armache, J.-P.; Armache, K.-J.  
Deposited on : 2019-03-09  
Resolution : 5.20 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](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	SINGLE PARTICLE	Depositor
Imposed symmetry	Not Provided	Depositor
Number of particles used	85029	Depositor
Resolution determination method	FSC 0.143 CUT-OFF	Depositor
CTF correction method	Not provided	Depositor
Microscope	FEI TALOS ARCTICA	Depositor
Voltage (kV)	200	Depositor
Electron dose ( $e^-/\text{\AA}^2$ )	41.0	Depositor
Minimum defocus (nm)	Not provided	Depositor
Maximum defocus (nm)	Not provided	Depositor
Magnification	Not provided	Depositor
Image detector	GATAN K2 SUMMIT (4k x 4k)	Depositor