



## wwPDB EM Map Validation Summary Report ⓘ

Dec 9, 2020 – 12:29 pm GMT

EMDB ID : EMD-5220  
Title : The architecture of the DNA polymerase-PCNA-DNA ternary complex  
Authors : , Mayanagi.K.; , Nishida.H.; , Kiyonari.S.; , Saito.M.; , Kohda.D.; , Ishino.Y.;  
          , Shirai.T.; , Morikawa.K.  
Deposited on : 2010-07-23  
Resolution : 19.00 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	singleParticle	Depositor
Imposed symmetry	Not Provided	Depositor
Number of images used	18897	Depositor
Resolution determination method	FSC 0.5 CUT-OFF	Depositor
CTF correction method	Not provided	Depositor
Microscope	JEOL 1010	Depositor
Voltage (kV)	100	Depositor
Electron dose ( $e^-/\text{\AA}^2$ )	Not provided	Depositor
Minimum defocus (nm)	Not provided	Depositor
Maximum defocus (nm)	Not provided	Depositor
Magnification	400000.0	Depositor
Image detector	GENERIC GATAN	Depositor