



## Full wwPDB EM Map Validation Report ⓘ

Dec 9, 2020 – 12:32 pm GMT

EMDB ID : EMD-5435  
Title : dATP-inhibited class Ia ribonucleotide reductase from E. coli: alpha4beta4 open conformation 5  
Authors : , Zimanyi.CM.; , Ando.N.; , Brignole.E.; , Asturias.FJ.; , Stubbe.J.; , Drennan.CL.  
Deposited on : 2012-06-13  
Resolution : 33.40 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.

---

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	344	Depositor
Resolution determination method	FSC 0.5 CUT-OFF	Depositor
CTF correction method	each particle	Depositor
Microscope	FEI TECNAI F20	Depositor
Voltage (kV)	120	Depositor
Electron dose ( $e^-/\text{\AA}^2$ )	25	Depositor
Minimum defocus (nm)	0.275	Depositor
Maximum defocus (nm)	1.801	Depositor
Magnification	50000.0	Depositor
Image detector	GATAN ULTRASCAN 4000 (4k x 4k)	Depositor