



wwPDB EM Map Validation Summary Report ⓘ

Dec 9, 2020 – 12:47 pm GMT

EMDB ID : EMD-6103
Title : Cryo-EM Structure of the 80S-RAC complex from yeast
Authors : , Zhang.YX.; , Ma.CY.; , Yuan.Y.; , Zhu.J.; , Li.NN.; , Chen.C.; , Wu.S.; ,
Yu.L.; , Lei.JL.; , Gao.N.
Deposited on : 2014-09-28
Resolution : 10.20 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

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	89801	Depositor
Resolution determination method	OTHER	Depositor
CTF correction method	defocus grouping	Depositor
Microscope	FEI TECNAI F20	Depositor
Voltage (kV)	200	Depositor
Electron dose ($e^-/\text{\AA}^2$)	20	Depositor
Minimum defocus (nm)	Not provided	Depositor
Maximum defocus (nm)	Not provided	Depositor
Magnification	Not provided	Depositor
Image detector	GATAN ULTRASCAN 4000 (4k x 4k)	Depositor