



## Full wwPDB EM Map Validation Report ⓘ

Dec 9, 2020 – 12:26 pm GMT

EMDB ID : EMD-5026  
Title : Cryo-negative stain structure of the yeast transcription factor TFIID at 20 Å resolution  
Authors : , Papai.G.; , Tripathi.MK.; , Ruhlmann.C.; , Crucifix.C.; , Jennings.JL.; , Link.AJ.; , Weil.PA.; , Schultz.P.  
Deposited on : 2008-08-20  
Resolution : 22.00 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	singleParticle	Depositor
Imposed symmetry	Not Provided	Depositor
Number of images used	10205	Depositor
Resolution determination method	FSC 0.5 CUT-OFF	Depositor
CTF correction method	phase flipping on each particle from one micrograph	Depositor
Microscope	FEI TECNAI F20	Depositor
Voltage (kV)	200	Depositor
Electron dose ( $e^-/\text{\AA}^2$ )	15	Depositor
Minimum defocus (nm)	0.137	Depositor
Maximum defocus (nm)	1.96	Depositor
Magnification	40080.0	Depositor
Image detector	KODAK SO-163 FILM	Depositor