



wwPDB EM Map Validation Summary Report ⓘ

Dec 7, 2020 – 04:19 pm GMT

EMDB ID : EMD-0589
Title : Cryo-EM structure of the human TFIIH core complex: Map classified for presence of the p62 BSD2 domain.
Authors : Greber, B.J.; Toso, D.; Fang, J.; Nogales, E.
Deposited on : 2019-02-19
Resolution : 3.90 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 | SINGLE PARTICLE | Depositor |
| Imposed symmetry | POINT, C1 | Depositor |
| Number of particles used | 92200 | Depositor |
| Resolution determination method | FSC 0.143 CUT-OFF | Depositor |
| CTF correction method | CTF correction during 3D reconstruction in RELION 3. | Depositor |
| Microscope | FEI TITAN KRIOS | Depositor |
| Voltage (kV) | 300 | Depositor |
| Electron dose ($e^-/\text{\AA}^2$) | 50.0 | Depositor |
| Minimum defocus (nm) | 0.5 | Depositor |
| Maximum defocus (nm) | 0.7 | Depositor |
| Magnification | 43478.0 | Depositor |
| Image detector | GATAN K2 SUMMIT (4k x 4k) | Depositor |