



wwPDB EM Map Validation Summary Report i

Dec 9, 2020 – 01:31 pm GMT

EMDB ID : EMD-8466
Title : Human Myeloma IgG4 Structure
Authors : , Ryazantsev.S.; , Tischenko.V.; , Zavyalov.V.
Deposited on : 2016-11-01
Resolution : 32.00 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

A user guide is available at

<https://www.wwpdb.org/validation/2017/EMMapValidationReportHelp>

with specific help available everywhere you see the i symbol.

The following versions of software and data (see [references](#) ①) were used in the production of this report:

EMDB validation analysis : FAILED
Validation Pipeline (wwPDB-VP) : 2.13

1 Experimental information i

| Property | Value | Source |
|--------------------------------------|---|-----------|
| EM reconstruction method | singleParticle | Depositor |
| Imposed symmetry | POINT, C1 | Depositor |
| Number of images used | 1000 | Depositor |
| Resolution determination method | FSC 0.5 CUT-OFF | Depositor |
| CTF correction method | Images were close to focus, no CTF correction implemented | Depositor |
| Microscope | JEOL 100CX | Depositor |
| Voltage (kV) | 80 | Depositor |
| Electron dose ($e^-/\text{\AA}^2$) | 100.0 | Depositor |
| Minimum defocus (nm) | Not provided | Depositor |
| Maximum defocus (nm) | Not provided | Depositor |
| Magnification | 80000.0 | Depositor |
| Image detector | KODAK 4489 FILM | Depositor |