



# Preliminary Full wwPDB EM Validation Report ⓘ

Oct 3, 2022 – 08:20 AM EDT

Deposition ID : D\_1000268928

**This wwPDB validation report is NOT for manuscript review**

This is a Preliminary Full wwPDB EM Validation Report.

This report is produced by the wwPDB Deposition System during initial deposition but before annotation of the structure.

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/EMTomogramValidationReportHelp>  
with specific help available everywhere you see the ⓘ symbol.

The types of validation reports are described at

<http://www.wwpdb.org/validation/2017/FAQs#types>.

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

EMDB validation analysis : 0.0.1.dev43  
Validation Pipeline (wwPDB-VP) : 2.31.2

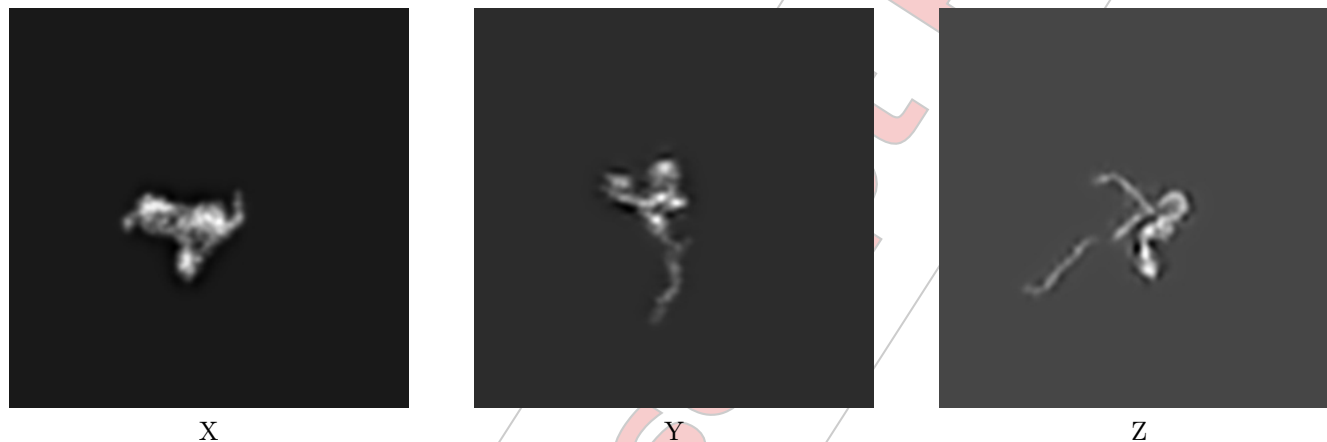
# 1 Experimental information ⓘ

| Property                             | Value                                   | Source    |
|--------------------------------------|---|-----------|
| EM reconstruction method             | TOMOGRAPHY                              | Depositor |
| Imposed symmetry                     | POINT, Not provided                     |           |
| Number of tilted images used         | Not provided                            |           |
| Resolution determination method      | Not provided                            |           |
| CTF correction method                | PHASE FLIPPING AND AMPLITUDE CORRECTION | Depositor |
| Microscope                           | FEI TITAN KRIOS                         | Depositor |
| Voltage (kV)                         | 300                                     | Depositor |
| Electron dose ( $e^-/\text{\AA}^2$ ) | 5.4                                     | Depositor |
| Minimum defocus (nm)                 | 2500                                    | Depositor |
| Maximum defocus (nm)                 | 3500                                    | Depositor |
| Magnification                        | Not provided                            |           |
| Image detector                       | GATAN K2 QUANTUM (4k x 4k)              | Depositor |
| Maximum voxel value                  | 1.283                                   | Depositor |
| Minimum voxel value                  | -0.326                                  | Depositor |
| Average voxel value                  | 0.001                                   | Depositor |
| Voxel value standard deviation       | 0.016                                   | Depositor |
| Recommended contour level            | 0.07                                    | Depositor |
| Tomogram size (Å)                    | 1480.0, 1480.0, 1480.0                  | wwPDB     |
| Tomogram dimensions                  | 200, 200, 200                           | wwPDB     |
| Tomogram angles (°)                  | 90.0, 90.0, 90.0                        | wwPDB     |
| Grid spacing (Å)                     | 7.4, 7.4, 7.4                           | Depositor |

## 2 Tomogram visualisation ⓘ

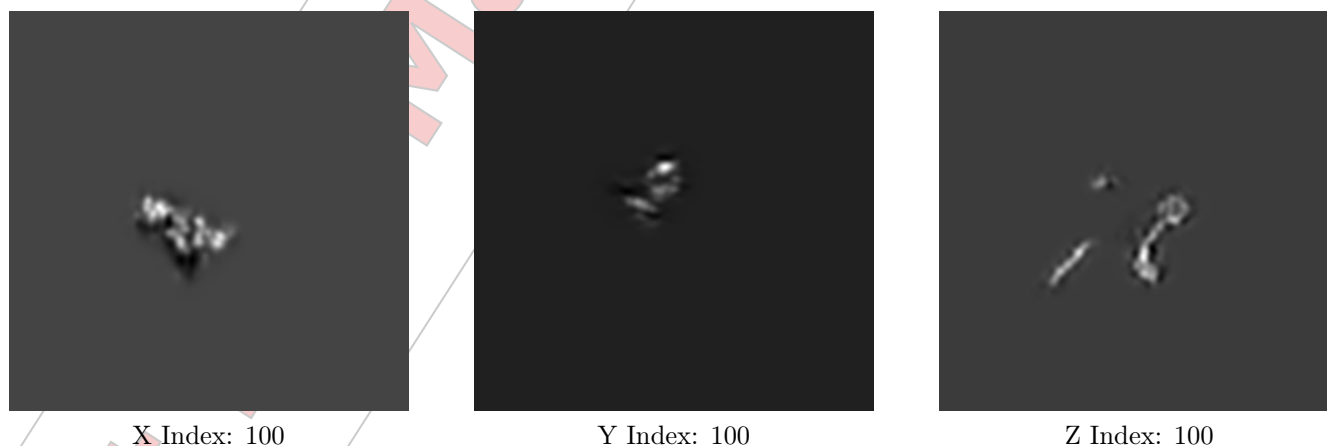
This section contains visualisations of the EMDB entry D\_1000268928. These allow visual inspection of the internal detail of the tomogram and identification of artifacts.

### 2.1 Orthogonal projections ⓘ



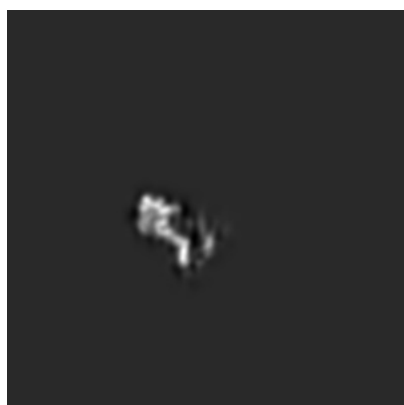
The images above show the tomogram projected in three orthogonal directions.

### 2.2 Central slices ⓘ



The images above show central slices of the tomogram in three orthogonal directions.

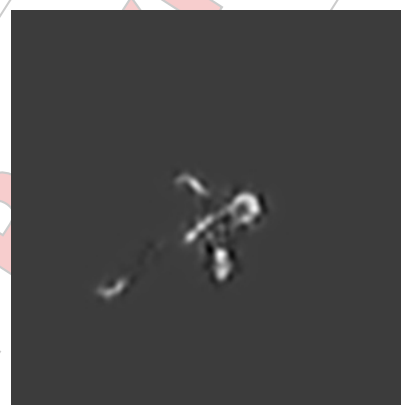
## 2.3 Largest variance slices [i](#)



X Index: 103



Y Index: 88



Z Index: 93

The images above show the largest variance slices of the tomogram in three orthogonal directions.

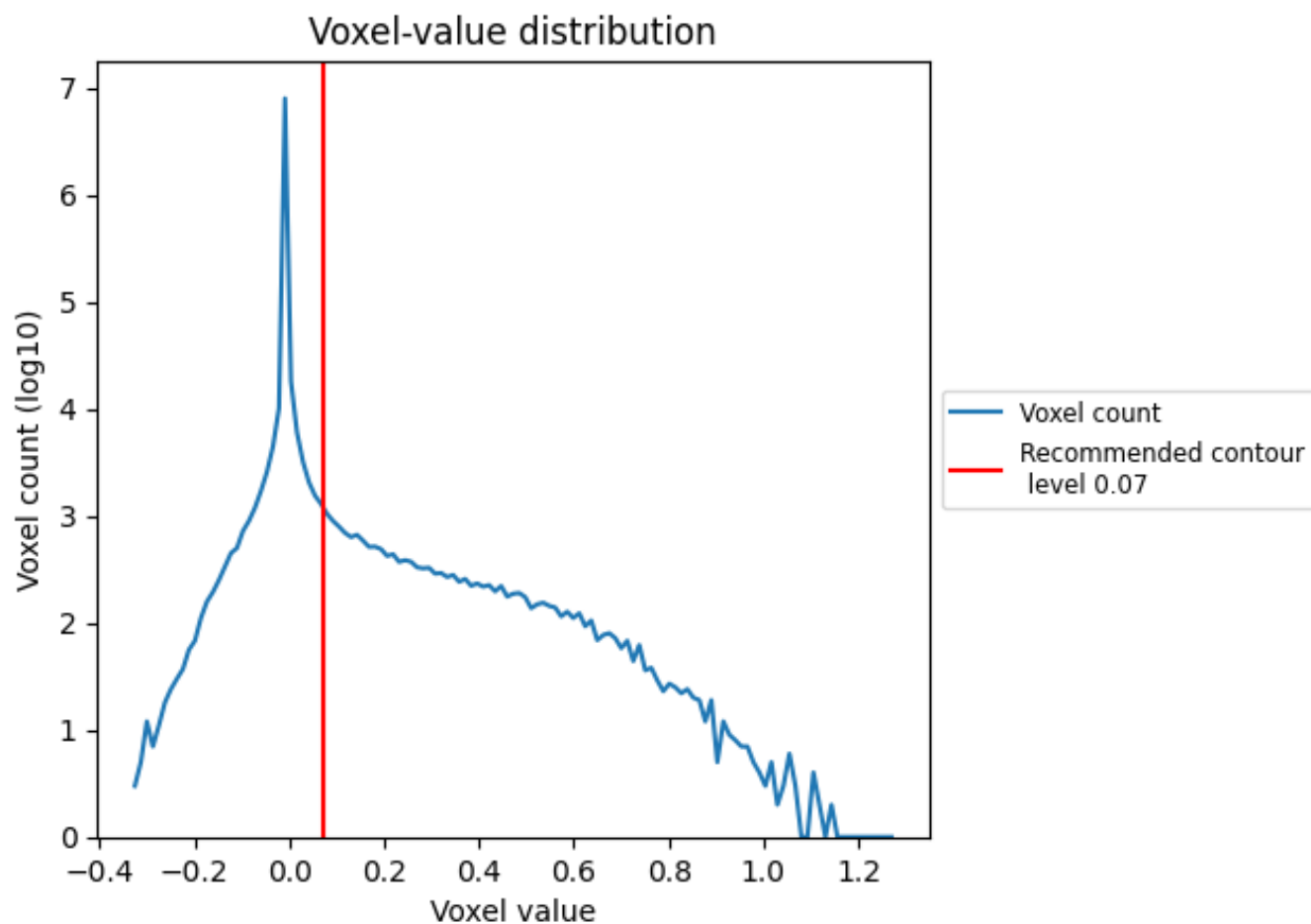
## 2.4 Mask visualisation [i](#)

This section was not generated. No masks/segmentation were deposited.

### 3 Tomogram analysis [i](#)

This section contains the results of statistical analysis of the tomogram.

#### 3.1 Voxel-value distribution [i](#)



The voxel-value distribution is plotted in 128 intervals along the x-axis. The y-axis is logarithmic.