



wwPDB EM Validation Summary Report ⓘ

Dec 5, 2020 – 11:32 PM EST

EMDB ID : EMD-1001
Title : Real space refinement of acto-myosin structures from sectioned muscle.
Authors : Chen, L.F.; Blanc, E.; Chapman, M.S.; Taylor, K.A.
Deposited on : 2002-06-18
Resolution : 40.00 Å(reported)

This is a wwPDB EM 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/EMTomogramValidationReportHelp>

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 : 0.0.0.dev61
Validation Pipeline (wwPDB-VP) : 2.15.1

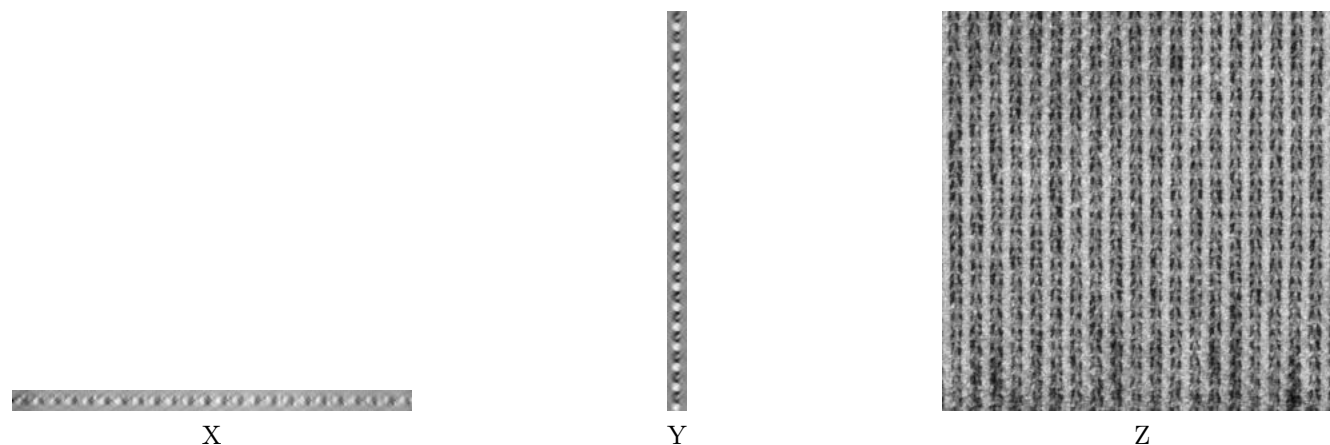
1 Experimental information

| Property | Value | Source |
|--------------------------------------|---------------------------|-----------|
| EM reconstruction method | TOMOGRAPHY | Depositor |
| Imposed symmetry | Not Provided | |
| Number of tilted images used | 30 | Depositor |
| Resolution determination method | OTHER | Depositor |
| CTF correction method | none | Depositor |
| Microscope | FEI/PHILIPS EM400 | Depositor |
| Voltage (kV) | 100 | Depositor |
| Electron dose ($e^-/\text{\AA}^2$) | Not provided | |
| Minimum defocus (nm) | Not provided | |
| Maximum defocus (nm) | Not provided | |
| Magnification | 17000.0 | Depositor |
| Image detector | Not provided | |
| Maximum voxel value | 366.680 | Depositor |
| Minimum voxel value | -417.992 | Depositor |
| Average voxel value | 1.860 | Depositor |
| Voxel value standard deviation | 47.792 | Depositor |
| Recommended contour level | 81.2 | Depositor |
| Tomogram size (\AA) | 9280, 9280, 464 | wwPDB |
| Tomogram dimensions | 600, 600, 30 | wwPDB |
| Tomogram angles ($^\circ$) | 90, 90, 90 | wwPDB |
| Grid spacing (\AA) | 15.4667, 15.4667, 15.4667 | Depositor |

2 Tomogram visualisation [i](#)

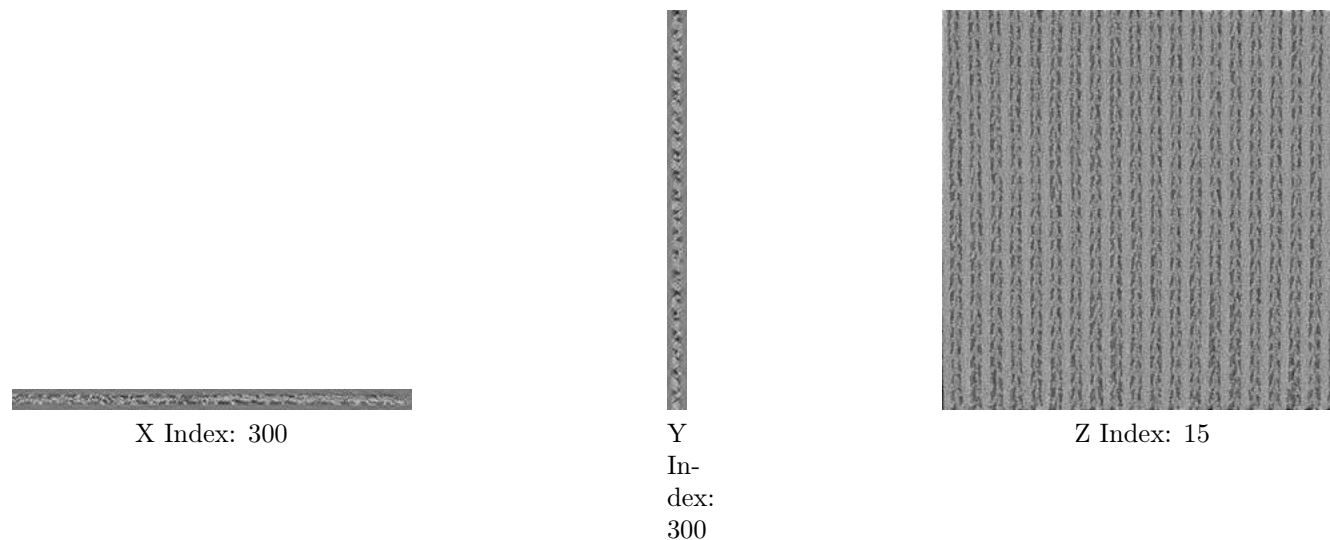
This section contains visualisations of the EMDB entry EMD-1001. These allow visual inspection of the internal detail of the tomogram and identification of artifacts.

2.1 Orthogonal projections [i](#)



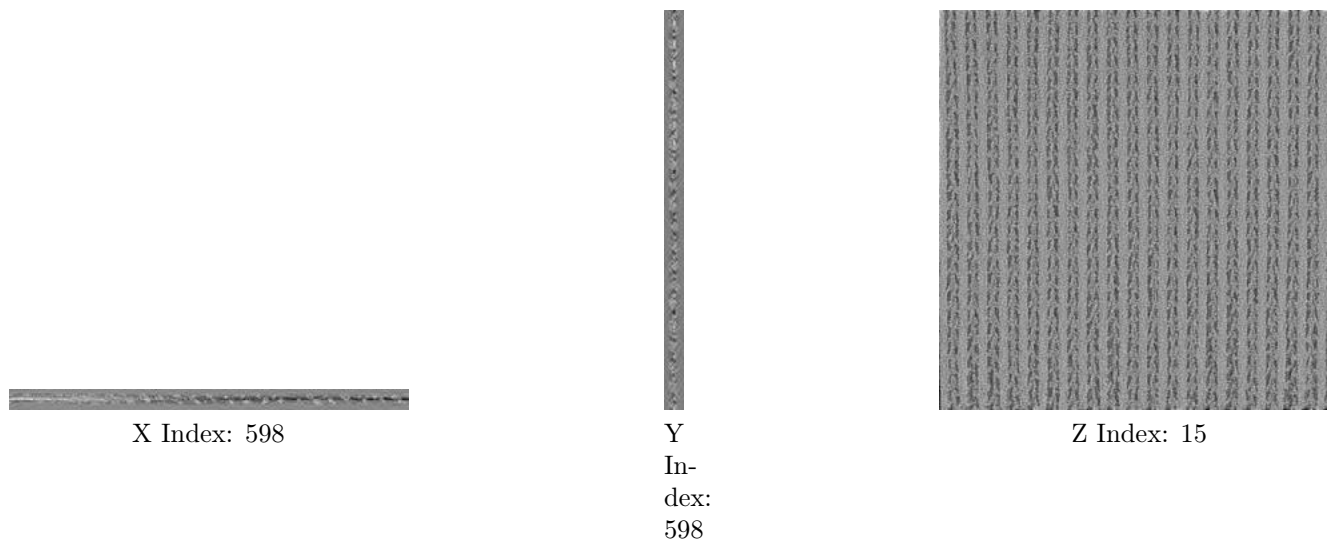
The images above show the tomogram projected in three orthogonal directions.

2.2 Central slices [i](#)



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

2.3 Largest variance slices [i](#)



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

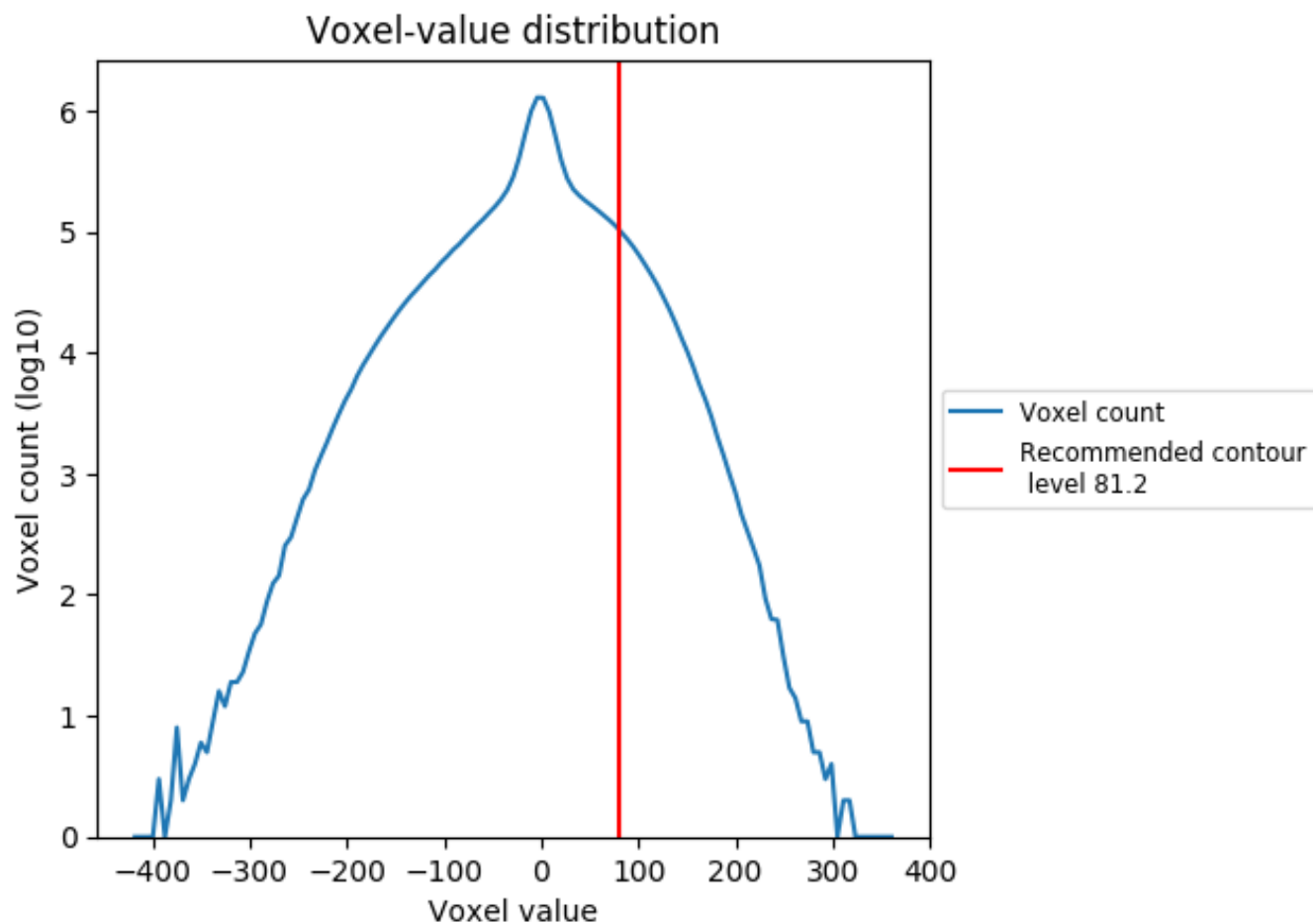
2.4 Mask visualisation [i](#)

This section was not generated.

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.