

## wwPDB EM Validation Summary Report (i)

### Mar 31, 2021 - 09:57 am BST

EMDB ID	) :	EMD-1582		
Title	e :	Three-dimensional structure of the 70S E. coli ribosome in its native 3D orga-		
		nization in polysomes.		
Authors	3 :	Brandt, F.; Elcock, A.H.; Etchells, S.A.; Ortiz, J.O.; Hartl, F.U.; Baumeister,		
		W.		
Deposited or	ı :	2008-11-07		
Resolution	Resolution : 30.00 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/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:

# 1 Experimental information (i)

Property	Value	Source
EM reconstruction method	SUBTOMOGRAM AVERAGING	Depositor
Imposed symmetry	Not Provided	
Number of subtomograms used	Not provided	
Resolution determination method	FSC 0.5 CUT-OFF	Depositor
CTF correction method	Not provided	
Microscope	FEI/PHILIPS CM200FEG	Depositor
Voltage (kV)	160	Depositor
Electron dose $(e^-/\text{Å}^2)$	50	Depositor
Minimum defocus (nm)	3.0	Depositor
Maximum defocus (nm)	4.0	Depositor
Magnification	53960.0	Depositor
Image detector	GENERIC TVIPS (4k x 4k)	Depositor
Maximum map value	0.000	Depositor
Minimum map value	-0.000	Depositor
Average map value	-0.000	Depositor
Map value standard deviation	0.000	Depositor
Recommended contour level	9.2e-09	Depositor
Map size (Å)	716.8, 716.8, 716.8	wwPDB
Map dimensions	128, 128, 128	wwPDB
Map angles (°)	90, 90, 90	wwPDB
Pixel spacing (Å)	5.6, 5.6, 5.6	Depositor



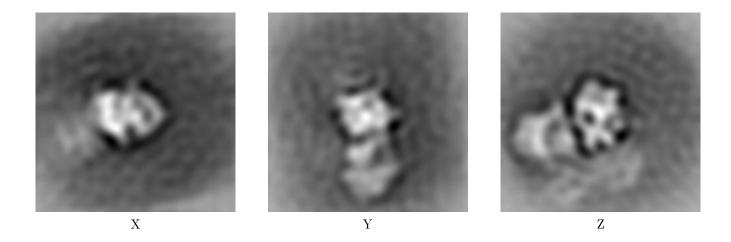
## 2 Map visualisation (i)

This section contains visualisations of the EMDB entry EMD-1582. These allow visual inspection of the internal detail of the map and identification of artifacts.

No raw map or half-maps were deposited for this entry and therefore no images, graphs, etc. pertaining to the raw map can be shown.

## 2.1 Orthogonal projections (i)

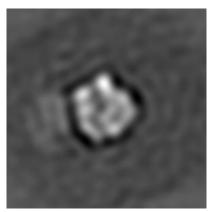
#### 2.1.1 Primary map



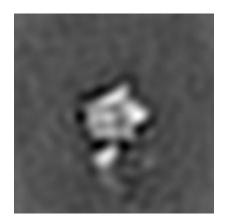
The images above show the map projected in three orthogonal directions.

## 2.2 Central slices (i)

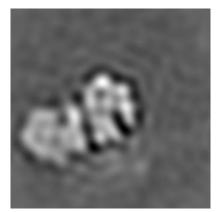
### 2.2.1 Primary map



X Index: 64



Y Index: 64



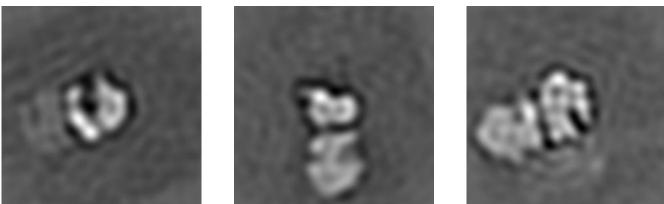
Z Index: 64



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

### 2.3 Largest variance slices (i)

#### 2.3.1 Primary map



X Index: 69

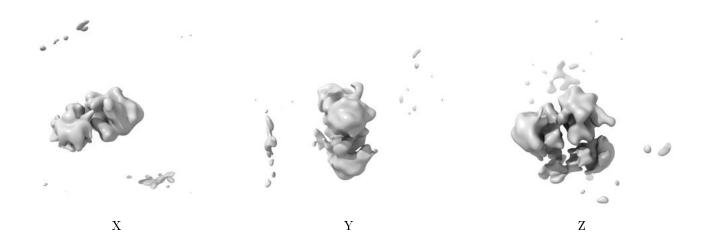
Y Index: 49



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

## 2.4 Orthogonal surface views (i)

#### 2.4.1 Primary map



The images above show the 3D surface view of the map at the recommended contour level 9.2e-09. These images, in conjunction with the slice images, may facilitate assessment of whether an appropriate contour level has been provided.



## 2.5 Mask visualisation (i)

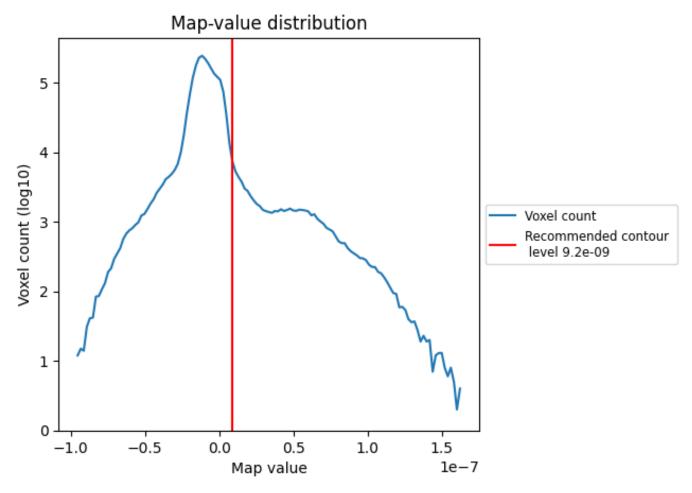
This section was not generated. No masks/segmentation were deposited.



## 3 Map analysis (i)

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

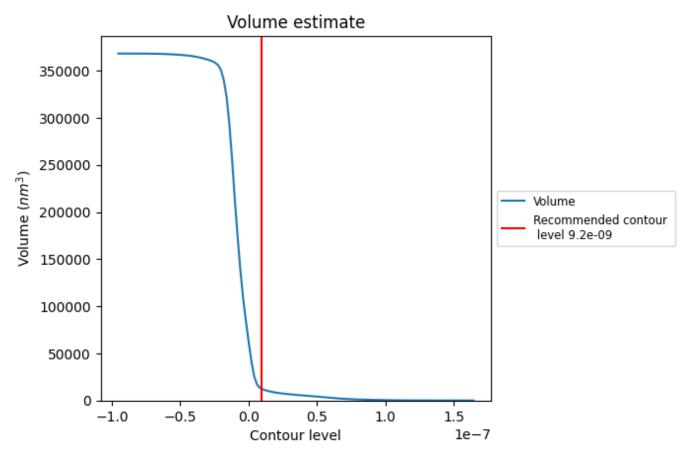
## 3.1 Map-value distribution (i)



The map-value distribution is plotted in 128 intervals along the x-axis. The y-axis is logarithmic. A spike in this graph at zero usually indicates that the volume has been masked.



### 3.2 Volume estimate (i)

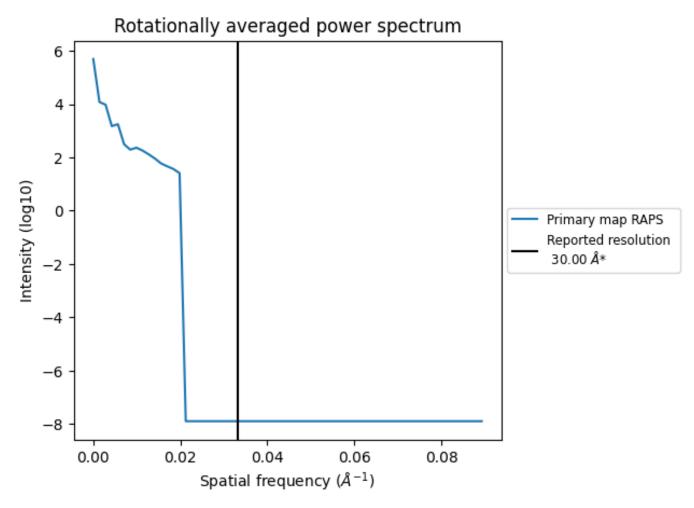


The volume at the recommended contour level is  $12656 \text{ nm}^3$ ; this corresponds to an approximate mass of 11433 kDa.

The volume estimate graph shows how the enclosed volume varies with the contour level. The recommended contour level is shown as a vertical line and the intersection between the line and the curve gives the volume of the enclosed surface at the given level.



## 3.3 Rotationally averaged power spectrum (i)



\*Reported resolution corresponds to spatial frequency of 0.033  ${\rm \AA}^{-1}$ 



# 4 Fourier-Shell correlation (i)

This section was not generated. No FSC curve or half-maps provided.

