



Full wwPDB EM Map Validation Report i

Dec 9, 2020 – 12:27 pm GMT

EMDB ID : EMD-5048
Title : CLIC2-RyR1 interaction and structural characterization by cryo-electron microscopy
Authors : , Meng.X.; , Wang.G.; , Viero.C.; , Wang.Q.; , Mi.W.; , Su.X.; , Wagenknecht.T.; , Williams.A.; , Liu.Z.; , Yin.C-C.
Deposited on : 2009-01-20
Resolution : 25.00 Å(reported)

This is a Full wwPDB EM Map Validation 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	Not Provided	Depositor
Number of images used	15889	Depositor
Resolution determination method	FSC 0.5 CUT-OFF	Depositor
CTF correction method	Each particle	Depositor
Microscope	FEI TECNAI F20	Depositor
Voltage (kV)	200	Depositor
Electron dose ($e^-/\text{\AA}^2$)	10	Depositor
Minimum defocus (nm)	1.5	Depositor
Maximum defocus (nm)	4.5	Depositor
Magnification	50760.0	Depositor
Image detector	KODAK SO-163 FILM	Depositor