



Full wwPDB EM Map Validation Report ⓘ

Dec 9, 2020 – 11:48 am GMT

EMDB ID : EMD-30003
Title : Cryo-EM map of CRF1 receptor bound to CRF and Gs protein
Authors : , Liang.YL.; , Belousoff.MJ.; , Zhao.P.; , Danev.R.; , Sexton.PM.; , Wootten.D.
Deposited on : 2020-02-10
Resolution : 2.70 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 ⓘ symbol.

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

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

1 Experimental information

Property	Value	Source
EM reconstruction method	singleParticle	Depositor
Imposed symmetry	Not Provided	Depositor
Number of images used	515871	Depositor
Resolution determination method	FSC 0.143 CUT-OFF	Depositor
CTF correction method	Not provided	Depositor
Microscope	FEI TITAN KRIOS	Depositor
Voltage (kV)	300	Depositor
Electron dose ($e^-/\text{\AA}^2$)	64.0	Depositor
Minimum defocus (nm)	0.4	Depositor
Maximum defocus (nm)	1.6	Depositor
Magnification	105000.0	Depositor
Image detector	GATAN K3 BIOQUANTUM (6k x 4k)	Depositor