	Final Map	Final Map Deconvolved
Masked Resolution (Å) at FSC 0.143	3.9	-
Unmasked Resolution (Å) at FSC 0.143	4.2	-
	Coordinate Refinement	
Map to Model Masked Resolution (Å) at FSC 0.5	6.2	4.6
	Model Composition	
Total Atoms	33,508	
Hydrogens	16,644	
Ligands	0	
	Model Geometry (RMSD)	
Bond Lengths (Å)	0.020	
Bond Angles (°)	1.765	
	Ramachandran Plot	
Favored (%)	97.41	
Allowed (%)	2.40	
Outliers (%)	0.19	
	Validation	
MolProbity Score	0.84	
Clash Score	0.69	
Rotamer Outliers (%)	0	

Table 1 | CryoEM Map Resolutions, Model Refinement, and Model Statistics