Table S3 Electron microscopy and model building information, Related to Figure 2

Data collection	
EM equipment	FEI Titan Krios
Acceleration voltage	300 kV
Detector	FEI Falcon III
Pixel size	1.16 Å
Electron dose	40 e-/ Å ²
Defocus range	0.8 – 4 μm
Reconstruction	
Software	RELION
Number of Particles (in high-resolution structure)	308,163
Accuracy of rotation	1.0 °
Accuracy of translation	0.7 pixel
Final resolution	3.4 Å
Model building	

Software COOT

Refinement

Software PHENIX

Map sharpening B-factor -178.0

Model composition

Protein residues 8183
RNA nucleotides 275
GTP 1

IP6	1
Validation	
RMS deviation	
Bond length	0.007
Bond angle	1.252
Ramachandran plot statistics	
Preferred	84.63
Allowed	15.03
Outlier	0.33
Clash score	5.81
MolProbity score	1.98