

NCCAT Session Report

GUP2: chameleon access

GUP2 application

Project ID: NCCAT-GUP2-KG220701

Project name: Structural basis for nuclear export of HIV-1RNA Mediated by Rev

Response Element

Primary user name: Keerthu Gottipati

eRA Commons user name: kegottip

Institution: University of Texas Medical Branch

Submission date: 07/01/2022

Sample information

1) Name/Title of macromolecule of interest: †RNA_RRE237

2) Molecular weight: 100 kDa

3) Storage buffer: 50 mM HEPES

4) Ligands/Binding partners in sample (if applicable):

5) Standard storage temperature (in °C): -80C

6) Storage time (max time at storage temperature and/or at RT):

7) Highest soluble concentration tested (mg/mL): 10mg/ml

8) Issue the sample is facing: ice thickness and particle density

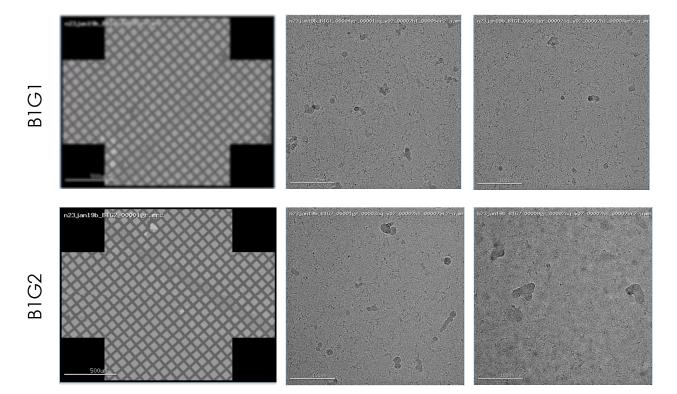
9) Additional information:

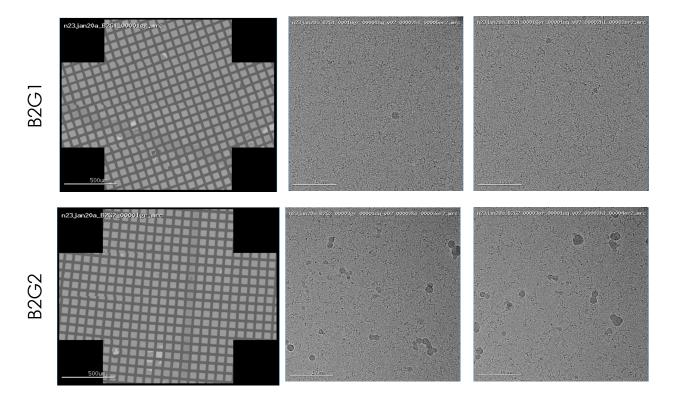




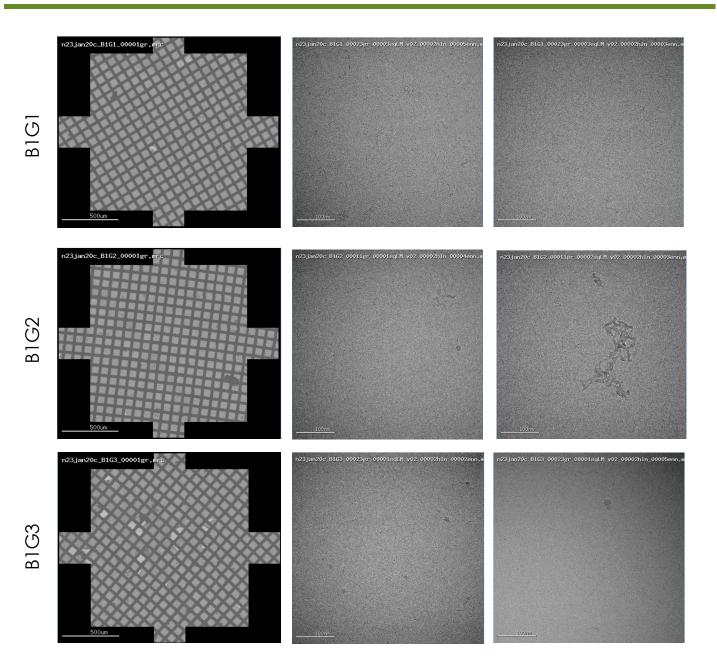
chameleon session log

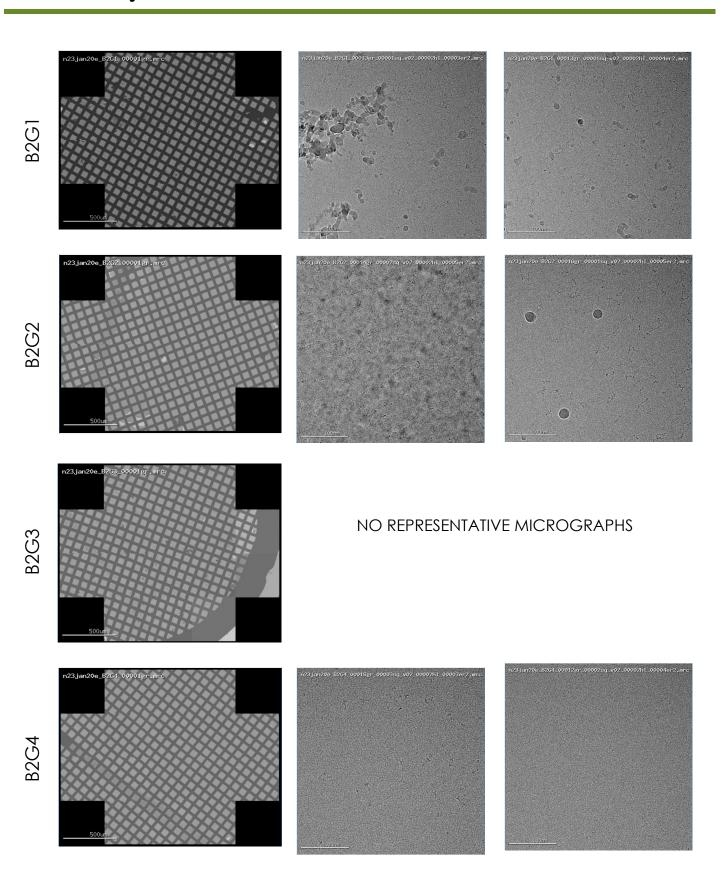
Session	Grid	Notes
n23jan19b Microscope: Krios 7 Camera: Falcon 4 Dose: 47.32 e ⁻ /Å Mag: 130000 Pixel Size: 0.959 Å Number of images: 553 Concentration: 10mg/ml	BIGI	Acceptable overwicked Small collection done on this grid
	B1G2	Acceptable underwicked
n23jan20a Microscope: Krios 7 Camera: Falcon 4 Dose: 47.32 e ⁻ /Å Mag: 130000 Pixel Size: 0.959 Å Number of images: 8 Concentration: 6.66mg/ml	B2G1	Acceptable overwicked
	B2G2	Acceptable overwicked
n23jan20c Microscope: Glacios 1 Camera: Falcon 3 Dose: 51.68 e ⁻ /Å Mag: 120000 Pixel Size: 1.204 Å Number of images: 25 Concentration: 5mg/ml	B1G1	Acceptable overwicked with gradient
	B1G2	Acceptable overwicked
	B1G3	Acceptable overwicked
n23jan20e Microscope: Krios 7 Camera: Falcon 4 Dose: 47.32 e ⁻ /Å Mag: 130000 Pixel Size: 0.959 Å Number of images: 32 Concentration: BOX 2- 5mg/ml & BOX 3 – 6.66mg/ml	B2G1	Acceptable underwicked with gradient Not enough squares on stripe cannot use for collection
	B2G2	Acceptable overwicked
	B2G3	Acceptable overwicked with gradient Grid cannot be used for collection
	B2G4	Acceptable overwicked
	B3G1	Acceptable overwicked with gradient Not enough squares on stripe cannot use for collection
	B3G2	Acceptable overwicked





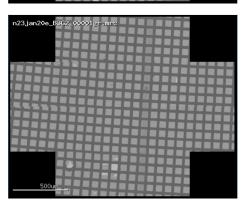
Session: n23jan20c – Glacios 1 – NCCAT-GUP2-KG220701 BOX 012023-1

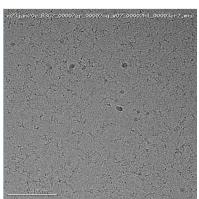


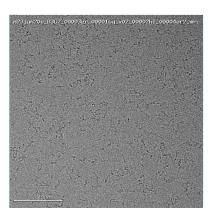


h23jan20e_B301_00001gr_mrc

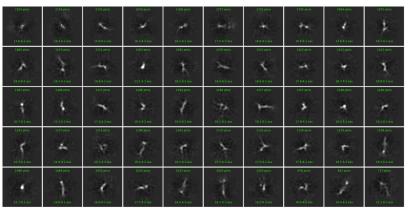
NO REPRESENTATIVE MICROGRAPHS







Data Processing: n23jan18b – Krios 7 – NCCAT-GUP2-KG220701 BOX 011923-1

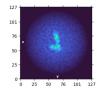


cryoSPARC v4.12 cryoSPARC live 2D classes: 635,902 particles

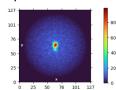
2nd Round 2D classes: 220,976 particles 3rd Round 2D classes: 210,708 particles Classes selected for Ab-initio: 144,487 particles

This image are the 2D classes for the Refinement:
67,629 particles

Ab-initio Class 000 (47.39%)



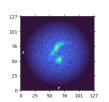


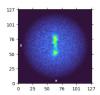


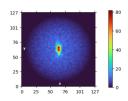




Ab-initio Class 001 (52.61%)



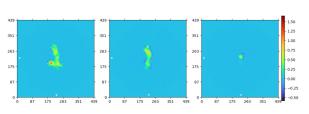








Homogenous Refinement of Ab-initio class 000









Acknowledgements

Some of this work was performed at the National Center for CryoEM Access and Training (NCCAT) and the Simons Electron Microscopy

Center located at the New York Structural Biology Center, supported by the NIH Common Fund Transformative High Resolution Cryo Electron Microscopy

program (U24 GM229539), and by grants from the Simons Foundation (SF349247) and NY S tate Assembly.