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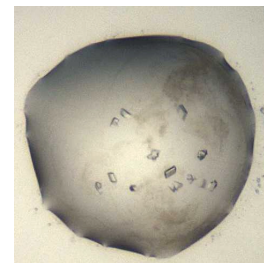
🌐 Crystallisation of Enterovirus D68 3C protease



Forked from [Crystallisation of Enterovirus D68 3C protease](#)

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ASAP Discovery



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Protocol status: Working

We use this protocol and it's working

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Disclaimer

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Abstract

The development of effective broad-spectrum antivirals forms an important part of preparing for future pandemics. A cause for concern is the currently emerging pathogen Enterovirus D68 (EV-D68) which primarily spreads through respiratory routes causing mostly mild to severe respiratory illness but, in severe cases, acute flaccid myelitis. The 3C protease of EV-D68 is a potential target for the development of antiviral drugs due to its essential role in the viral life cycle and high sequence conservation. This protocol was used to grow D68 3C ProB crystals that were applied high-throughput crystallographic follow up compound screening on D68 3C.

Materials


SwissCI 3 lens crystallization plates <https://swissci.com/product/3-lens-crystallisation-plate/> **Codes:**

Midi: UVXPO-3LENS 3W96T-PS 3W96T-UVP

[M] 1 Molarity (M) Tris adjusted to  7.8 with NaOH, Molecular Dimensions, Catalog # MD2-027-PH 7.8

[M] 1 Molarity (M) Ammonium acetate, Molecular Dimensions, Catalog # MD2-002-PH

50% w/v PEG 3350, Molecular Dimensions, Catalog # MD2-250-9

Purified D683C protein ([M] 35 mg/mL) in [M] 10 millimolar (mM) HEPES,  7.5 , [M] 0.5 Molarity (M) NaCl, 5% glycerol, [M] 0.5 millimolar (mM) TCEP



Safety warnings

⚠ Follow all handling warning for the chemicals used in the crystallisation screen composition.

Enterovirus D68 3C protease expression and purification

- 1 **The protein used for crystallisation was expressed and purified using the following protocol.**

Protocol



NAME

Enterovirus D68 3C protease large scale purification protocol

CREATED BY

Korvus Wang

PREVIEW

Equipment needed

- 2 **Formulatrix Rock Imager** (or incubator of choice)
SPT mosquito

Equipment

Mosquito HV

NAME

High Volume 16-Channel Robotic Liquid Handler

TYPE

SPT LabTech

BRAND

3097-01057

SKU

<https://www.sptlabtech.com/products/liquid-handling/mosquito-hv/>^{LINK}

P100 8 multi-channel pipette

SwissCI 3 lens plate

Crystallisation experiment

1d

- 3 **Prepare seed stock:**




**Protocol**

NAME



Diamond XChem Seeding Protocol

CREATED BY

Peter Marples

PREVIEW1: 1 000 000 dilution  Sample seeds**4 Protein and buffer requirements:** 14.4 µL  35 mg/mL  Sample 2.88 mL Crystallisation screen 7.2 µL seeds, dilution 1:1 000 000**5 ccCrystallisation screen composition:** 0.1 Molarity (M) Tris  7.8 0.2 Molarity (M) Ammonium acetate

26% w/v PEG 3350


Stock solutions used: 1 Molarity (M) Tris adjusted to  7.8 with NaOH 1 Molarity (M) Ammonium acetate


50% w/v PEG 3350

Note

The crystallisation screen can be stored in a duran bottle or aliquoted into 96 deep well block for easy dispensing into SwissCI 3 lens plates.


For long term storage keep the crystallisation screen in the fridge at 4°C.

6 Dispense  30 µL Crystallisation screen into SwissCI 3 lens plate reservoir wells using a 100 µl multi-channel pipette.

Dispense  50 nL  35 mg/mL  Sample to each lens using the SPT mosquito.



Dispense  100 nL Crystallisation screen to each lens using the SPT mosquito.

Dispense  25 nL Seeds to each lens using the SPT mosquito.

Drop ratio: 2:4:1

Final drop volume: 175 nl

7 Incubate at  20 °C for  24:00:00 h in Formulatrix Rock Imager.

1d

Imaging Schedule: The first images are taken after 12 h and the imaging schedule follows a Fibonacci sequence of days for further collections.

8

Expected result

Crystals typically appear after 24 hours and reach their maximum size after ~24 h with some precipitation often remaining.

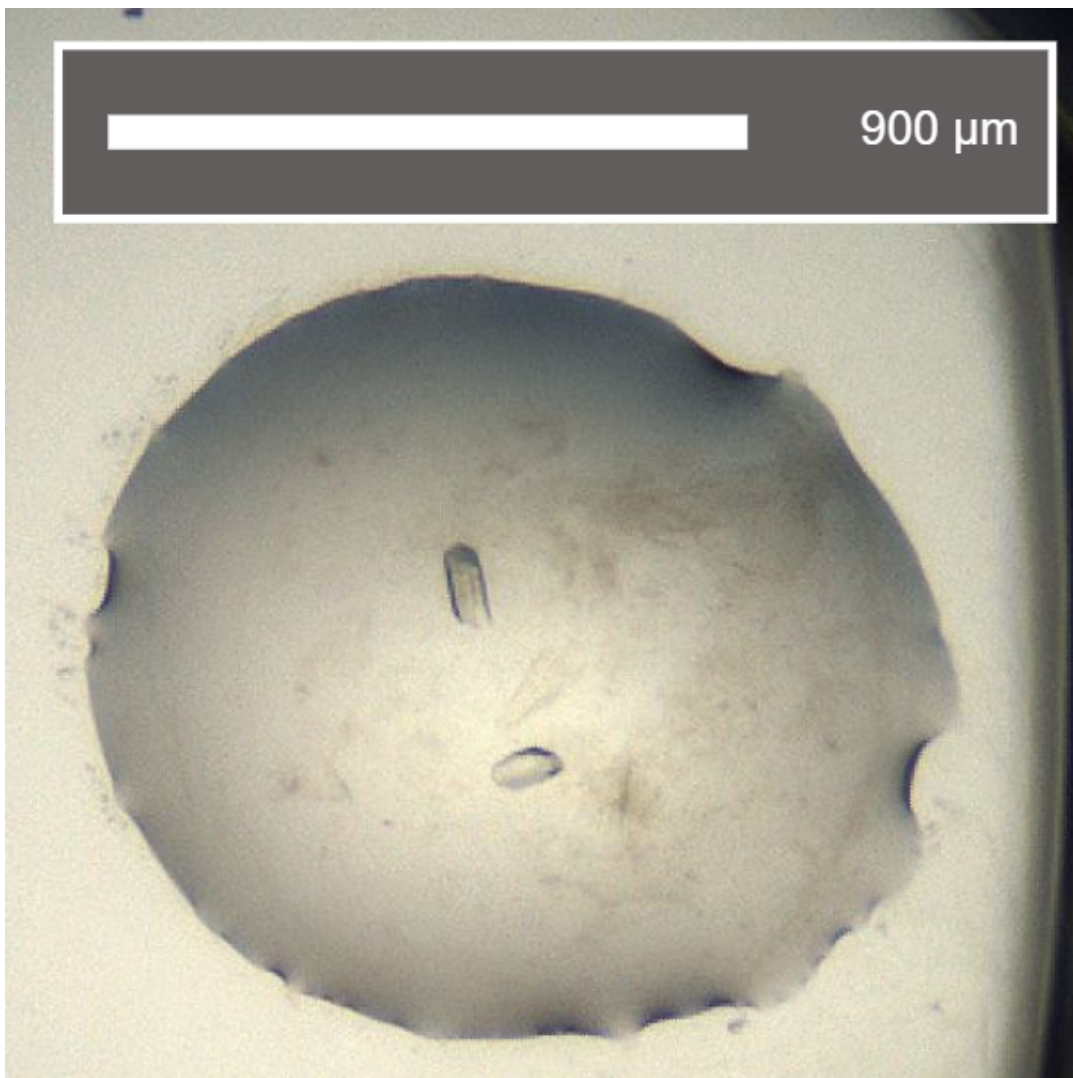
Morphology: small shards.

Size: ~40 μm in length and ~40 μm in width, depth of the crystals is ~20 μm , giving a glass shard appearance

Average resolution: 1.5 \AA

Space group: $P2_1$

Unit cell: 39.7 \AA , 105 \AA , 43.5 \AA
90.00°, 110.00°, 90.00°



An example of a drop containing D68 3C protease crystals.



Data collection at Synchrotron

9 Diamond Light Source
Unattended Data Collection (UDC)
Data Collection Temperature: 100K
Detector: DECTRIS EIGER2 X 9M
Beamline: I04-1
Wavelength: 0.9212 Å
Resolution (Å): 1.62
Beam Size (µm): 60 X 50
Number of images: 3600
Oscillation: 0.10°
Exposure (s): 0.0020
Transmission (%): 100
Flux (ph/s): 9.50e+11

Protocol references

Crystallographic fragment screen of Enterovirus D68 3C protease and iterative design of lead-like compounds using structure-guided expansions, <https://doi.org/10.1101/2024.04.29.591650>