

# ***NANUQ™***

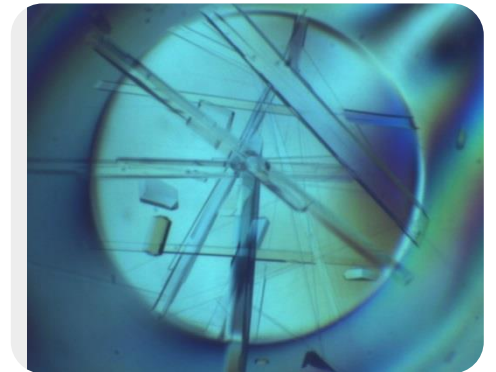
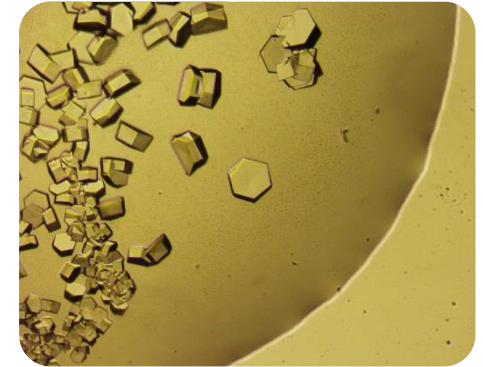
**Advanced Cryocooling Technology  
for Biomolecular Cryocrystallography**

**MiTeGen, LLC  
Ithaca, NY**



# Crystallography requires good crystals - *and that they stay good to the end of data collection*

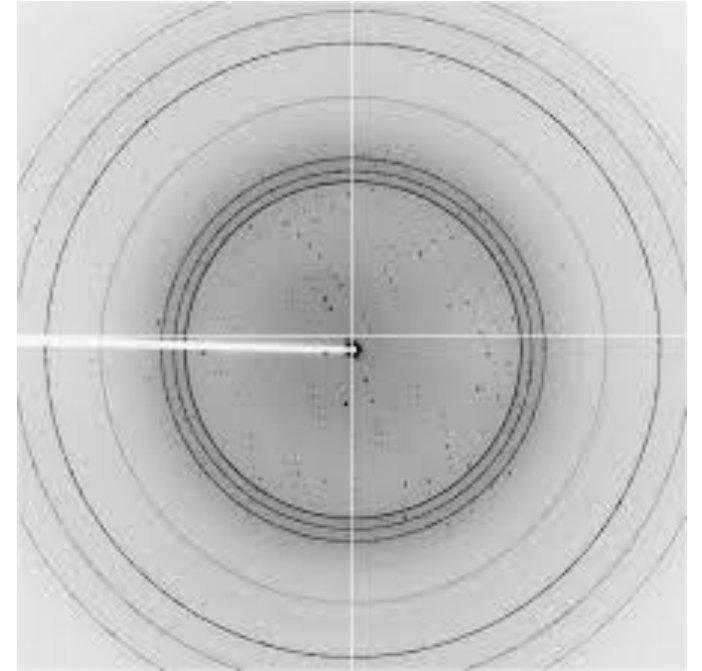
- Getting good crystals requires  
Time + Effort + Resources (\$)
- Once you have good crystals, don't **damage** or **waste** them.
- Use protocols that **maximize data quality**, **maximize throughput**, **minimize crystal-to-crystal variability**, and **minimize** risks of crystal **contamination**, **damage** and **loss**.



➡ Use MiTeGen's **NANUQ™** automated cryocooler!

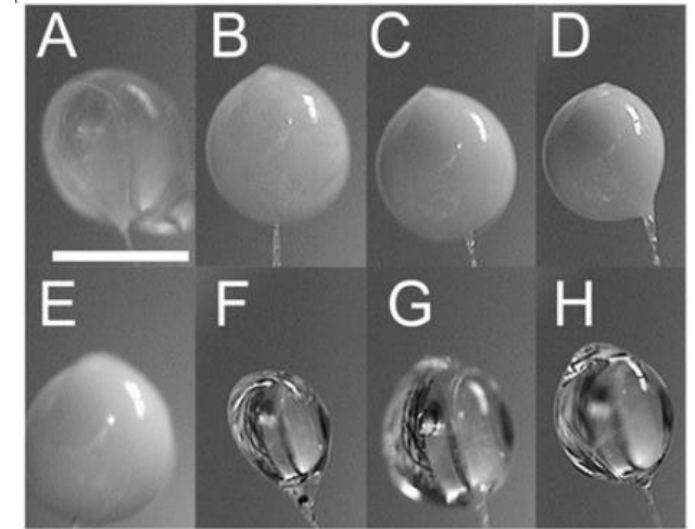
# Challenges in Biomolecular CryoCrystallography

- Cooling **damages crystals**, especially when ice forms.
- **Ice forms** in **inside** and **outside** crystals.
- **Ice** accumulates in  $\text{LN}_2$  and **contaminates crystals**.
- **20% of PDB deposited data sets** - and a much larger fraction of all data sets - **show structure factor errors due to ice**.
- High-value targets **with large solvent contents** (e.g., membrane proteins) and **large solvent cavities** (e.g., large complexes) are the most challenging to cool successfully.



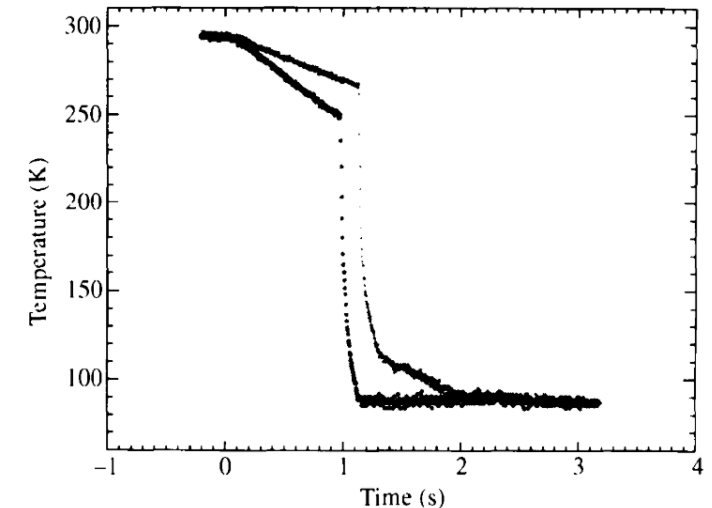
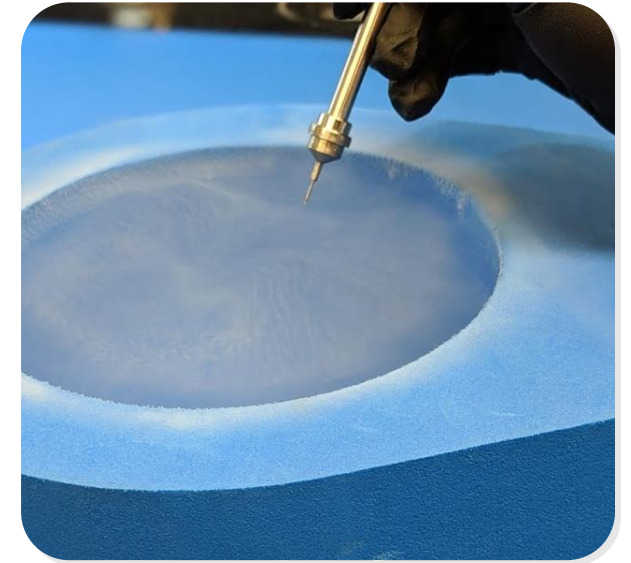
# You have Ice? Why not just use more cryoprotectant?

- **Cryoprotectant soaks** reduce ice, but can **damage** or **dissolve** crystals.
- Cryoprotectants can **modify protein conformation**.
- **Cryoprotectants reduce electron density contrast** and **resolution**.
- Cryoprotectants complicate electron density in solvent cavities.
- **Cryoprotectants can displace** or be mistaken for **ligands** (e.g., in HT fragment screening)



# Challenges in cooling by hand plunging in LN<sub>2</sub>

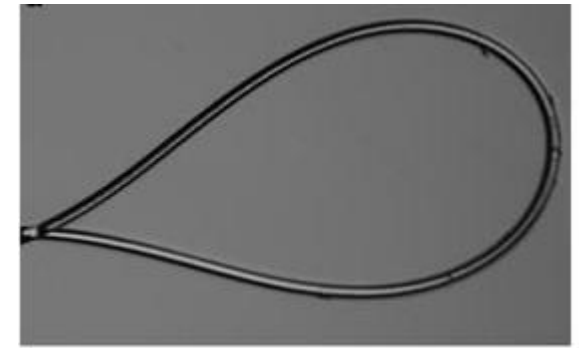
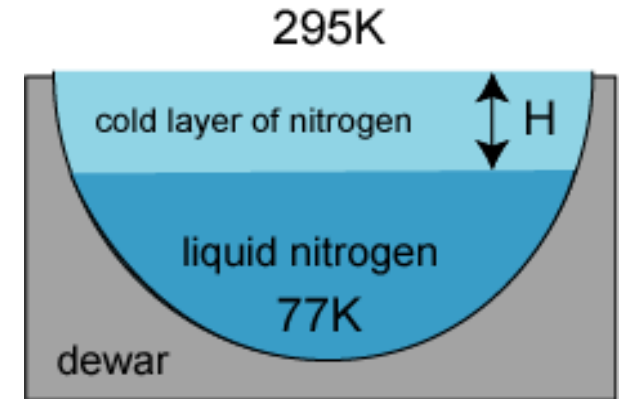
- Cooling **damage** depends on **cooling rate** and **cryoprotectant concentration**.
- **Cooling rates** in hand plunging are **modest** (100-1000 K/s) , so **cryoprotectants** are often needed **to eliminate ice**.
- **Cooling times** in hand plunging are **0.1 to 1 s**, **too long to kinetically capture** the biological temperature **structural ensemble**.



*T.-Y. Teng & K. Moffat, 1998*

# Challenges in cooling by hand plunging in LN<sub>2</sub>

- **Cooling rates** are **highly irreproducible** (by a factor up to **100 ×**) due to cooling in **cold gas** above LN<sub>2</sub> and to variable plunge speeds.
- Cooling rate variability is a **major cause of crystal nonisomorphism**.
- Why does one crystal diffract better than the next? Was it how it was grown, how it was cooled, or ... ?
- **Crystals** are **lost** during hand plunging and during manual loading into pucks. **Loops** sent to synchrotrons often arrive with **no crystals**.



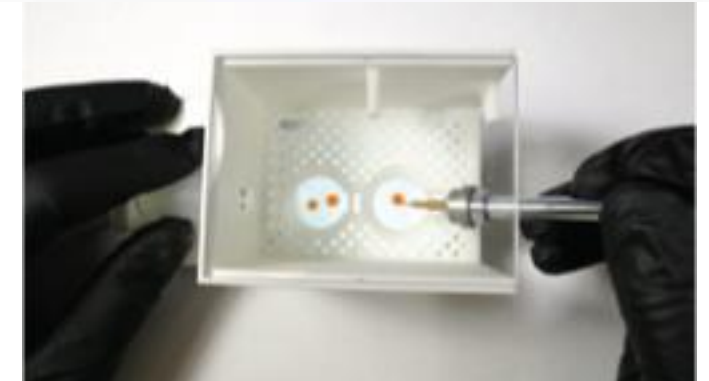
# What Would an Optimal Cryocrystallography Workflow Look Like?

1. Mount as-grown crystals and check diffraction at **room T** using MiTeGen's MicroRT™ system. Triage bad crystals.
2. Harvest crystals **in a humidity-controlled environment**.





# Watershed™ Humidity Control System

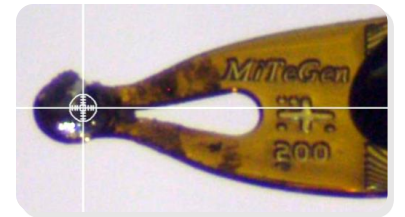


- Prevent drops from drying out while harvesting
- Perform controlled crystal dehydration to improve diffraction



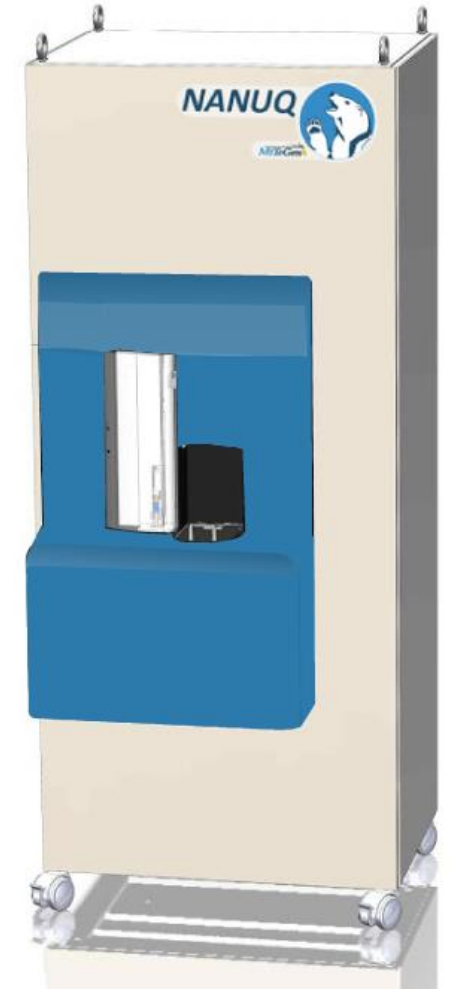
# What Would an Optimal Cryocrystallography Workflow Look Like?

1. Mount as-grown crystals and check diffraction at **room T** using MiTeGen's MicroRT™ system. Triage bad crystals.
2. Harvest crystals in a **humidity-controlled environment**.
3. Do a quick swipe through oil or a 10% cryo solution. **Do not soak in penetrating cryoprotectants.**
4. **Remove excess liquid** from around the crystal (and use MiTeGen's MicroLoops or MicroMounts).
5. **Cool as fast as possible in ice-free LN2.**
6. Store in an **ice-free** environment until data collection.

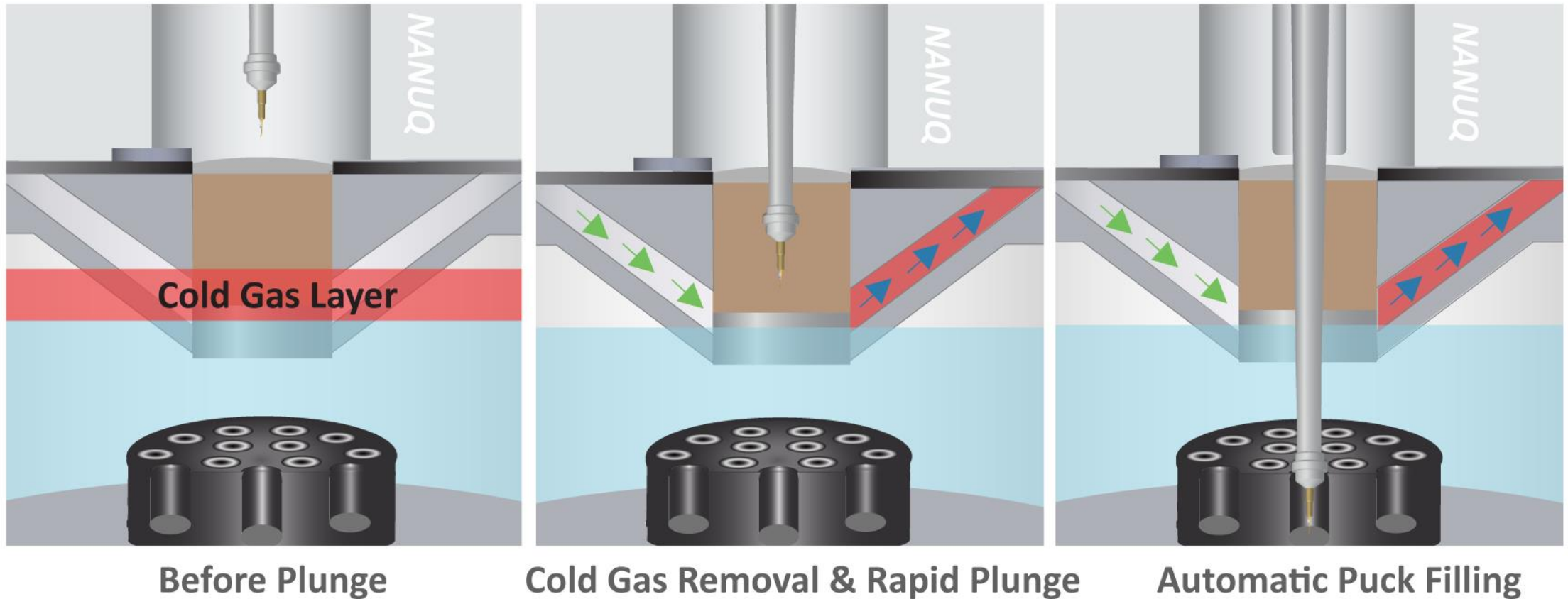


# NANUQ™:

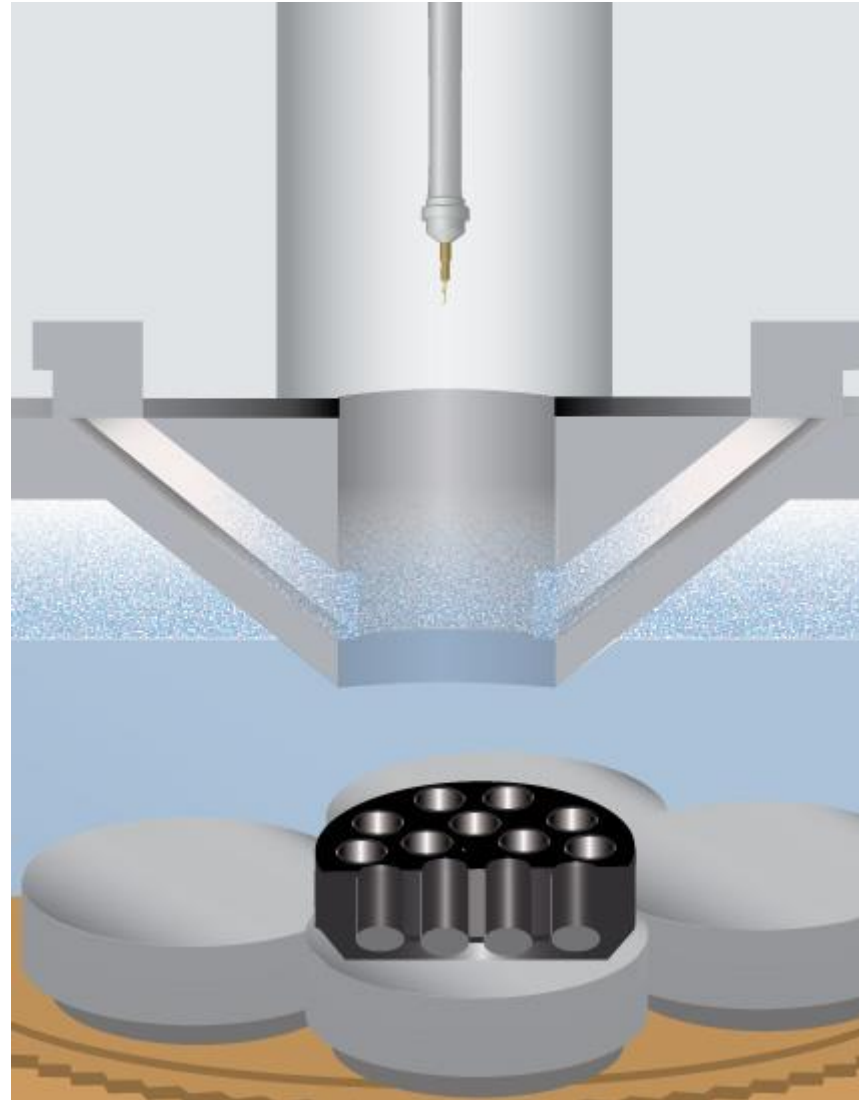
- Eliminates ice formation and ice contamination
- Minimizes sample-to-sample variability
- Provides complete control of every plunge – ultrafast cooling, slow cooling, or anything in between
- Standardizes cooling for every crystal and every user
- Automates critical steps between crystal harvesting and X-ray data collection



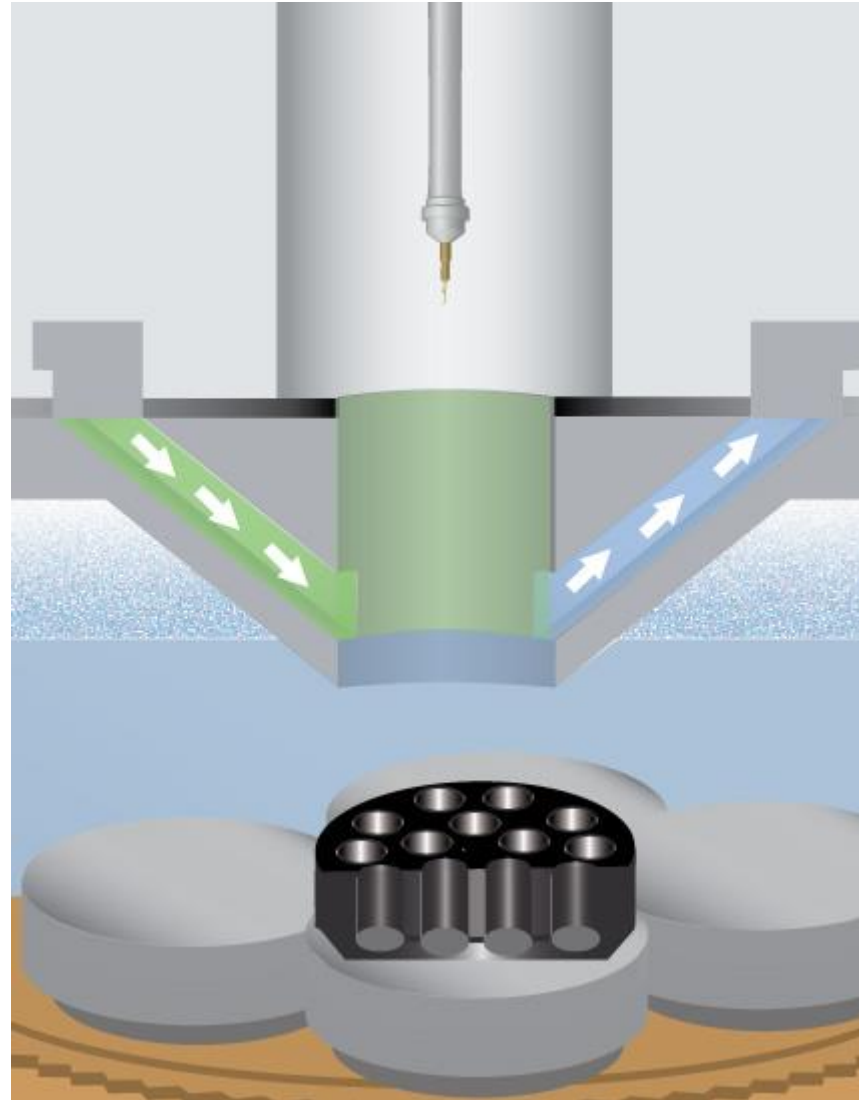
# NANUQ™: Key Design Features and Patented Technology



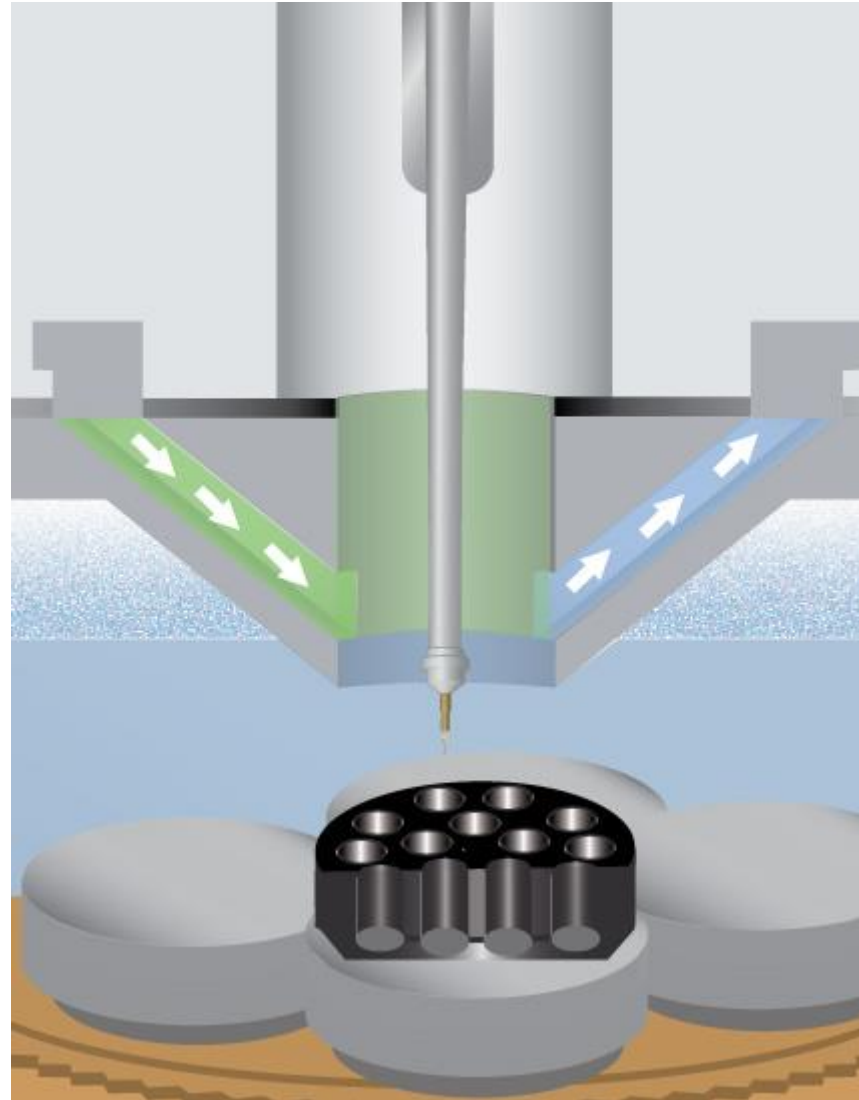
# *NANUQ™*: Hyperquenching (Fast Cooling) Plunge Sequence



# *NANUQ™*: Hyperquenching Plunge Sequence

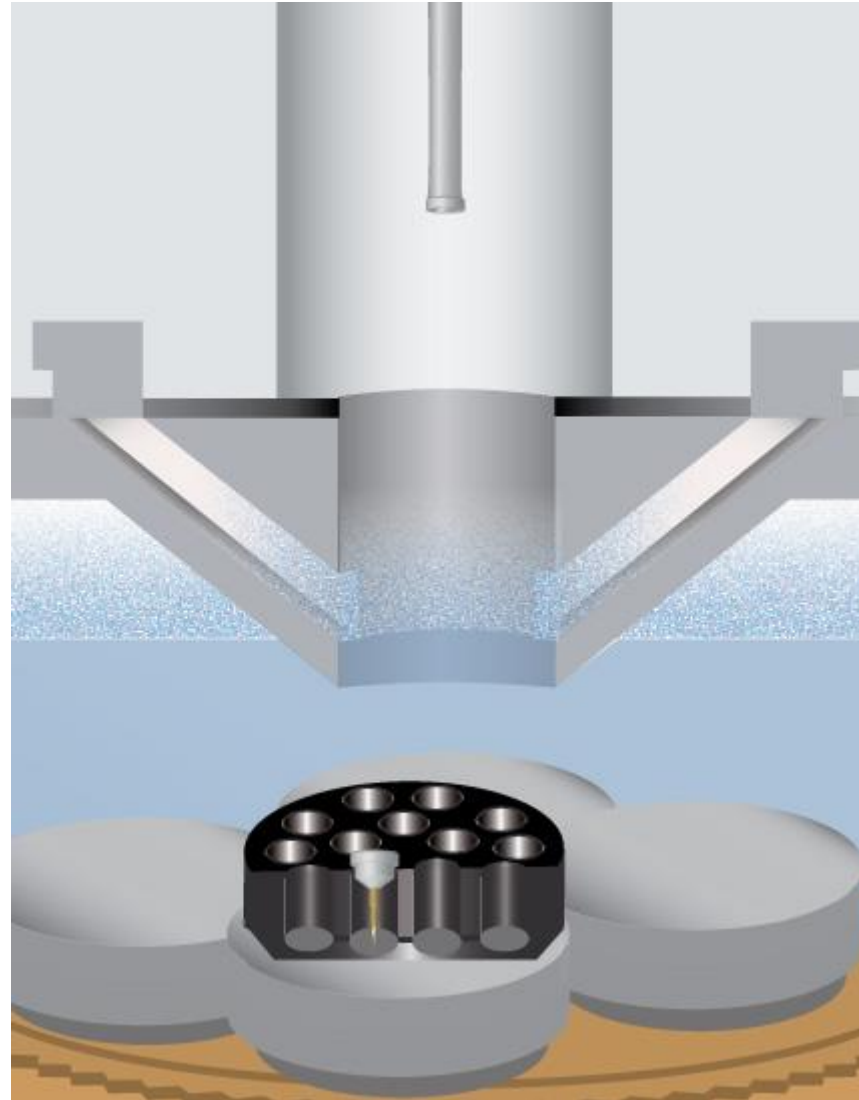


# NANUQ™: Hyperquenching Plunge Sequence





# *NANUQ™*: Hyperquenching Plunge Sequence

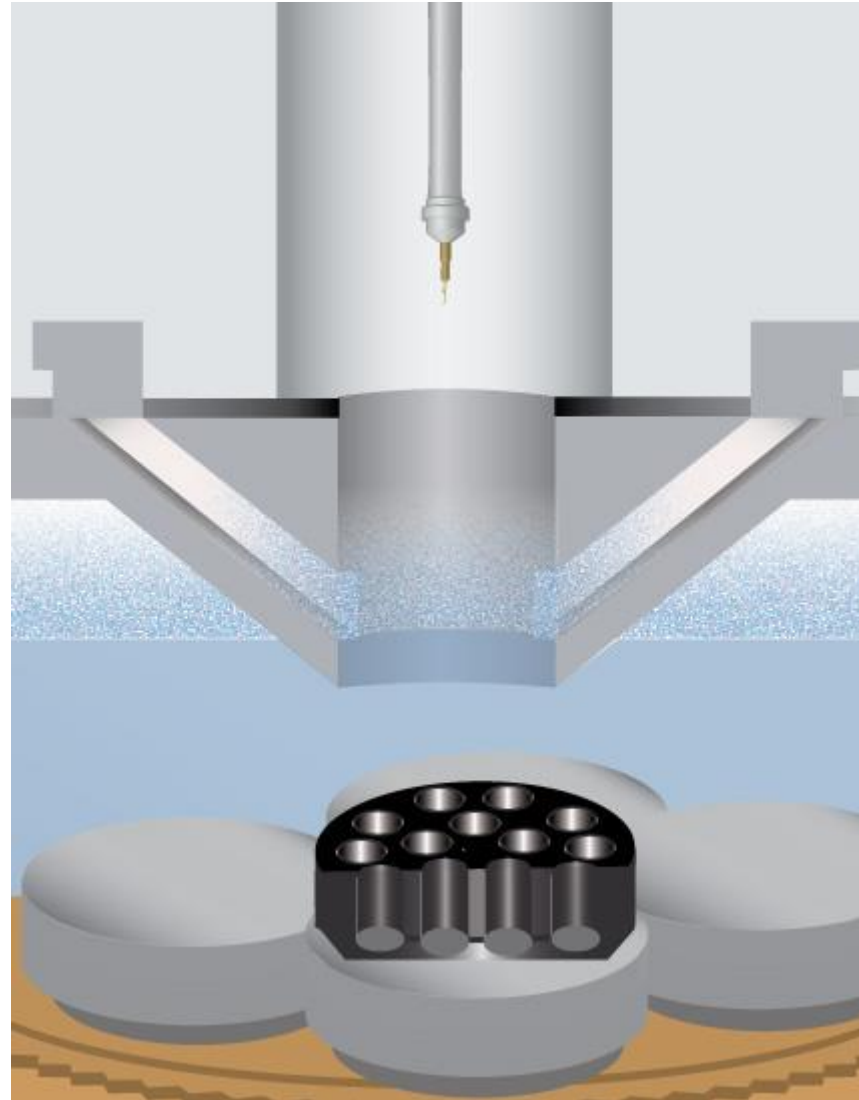


# *NANUQ*<sup>™</sup>: Slow- and Variable-Rate Cooling

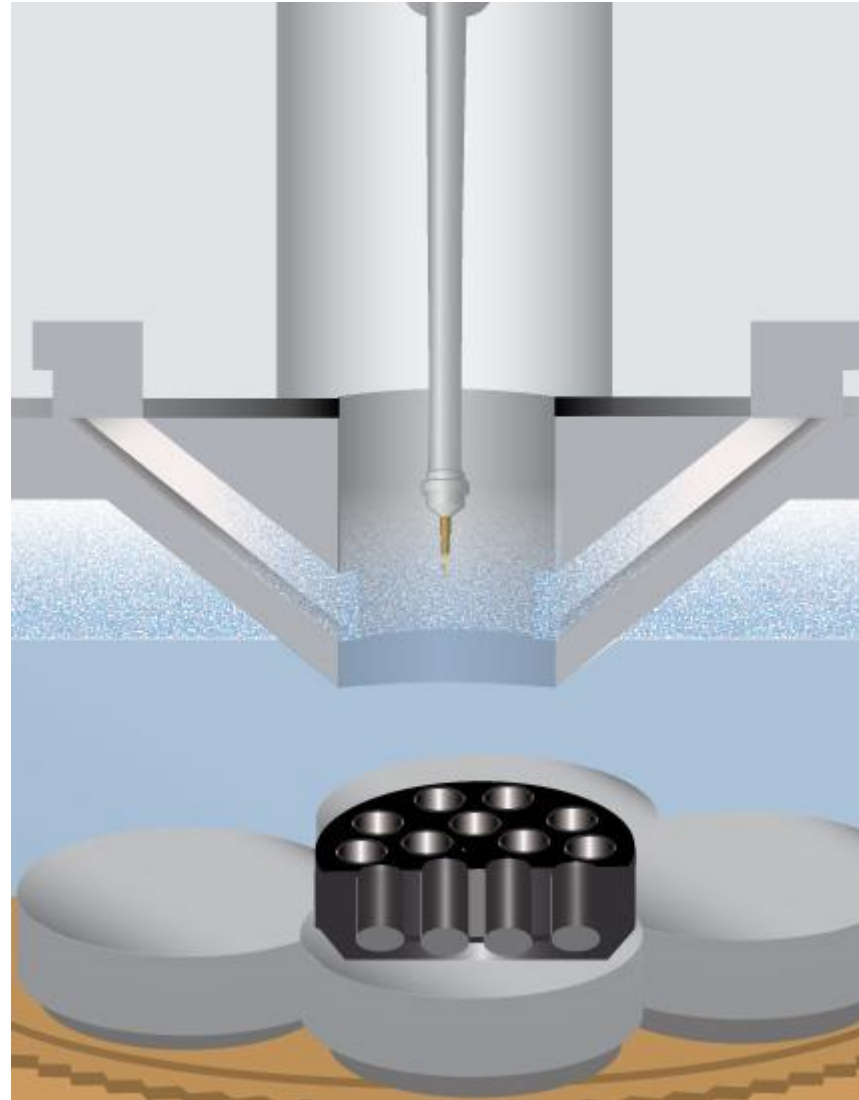
**The fastest cooling may not always give the best results.**

- **Large crystals** can develop internal thermal gradients during fast cooling that can lead to crystal damage.
- Flows of internal solvent during cooling can increase disorder. Allowing more time for this flow can reduce disorder.
- The protein and lattice may undergo favorable relaxations during slow cooling that improve overall crystal order.

# *NANUQ™*: Slow / Variable-Rate Cooling Plunge Sequence

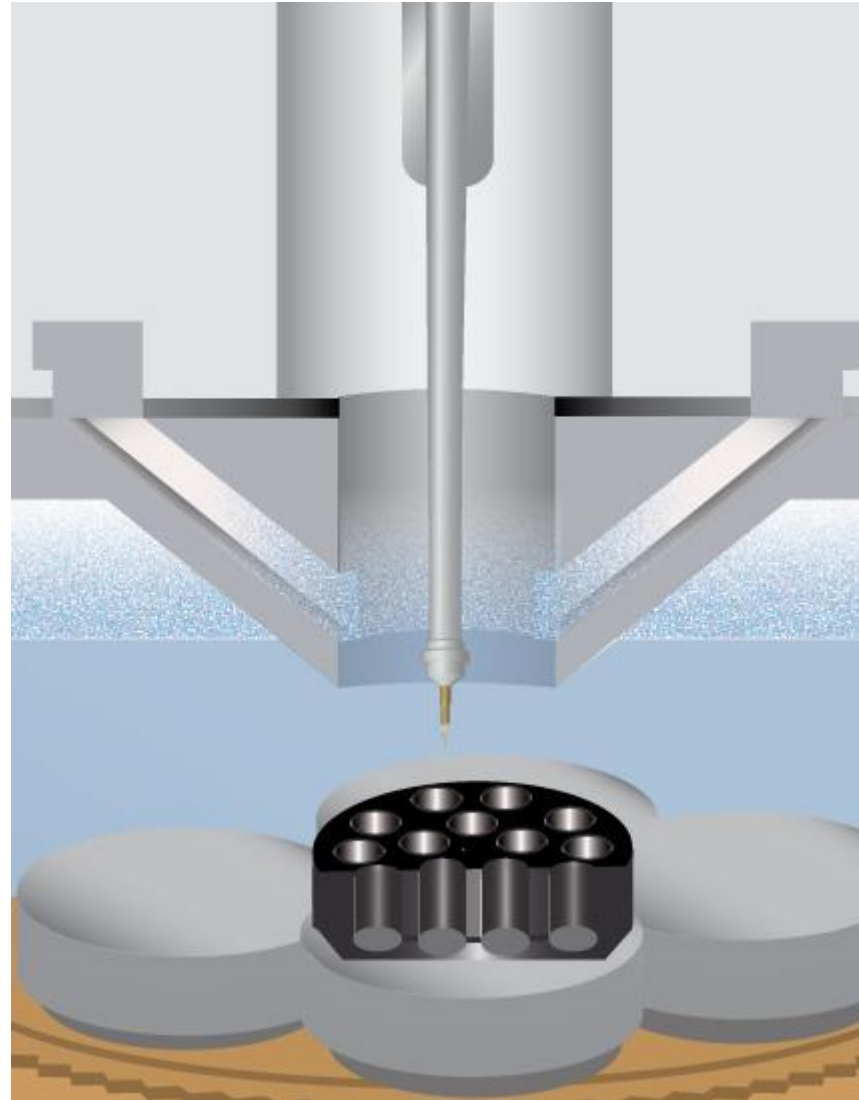


# *NANUQ™*: Slow / Variable-Rate Cooling Plunge Sequence

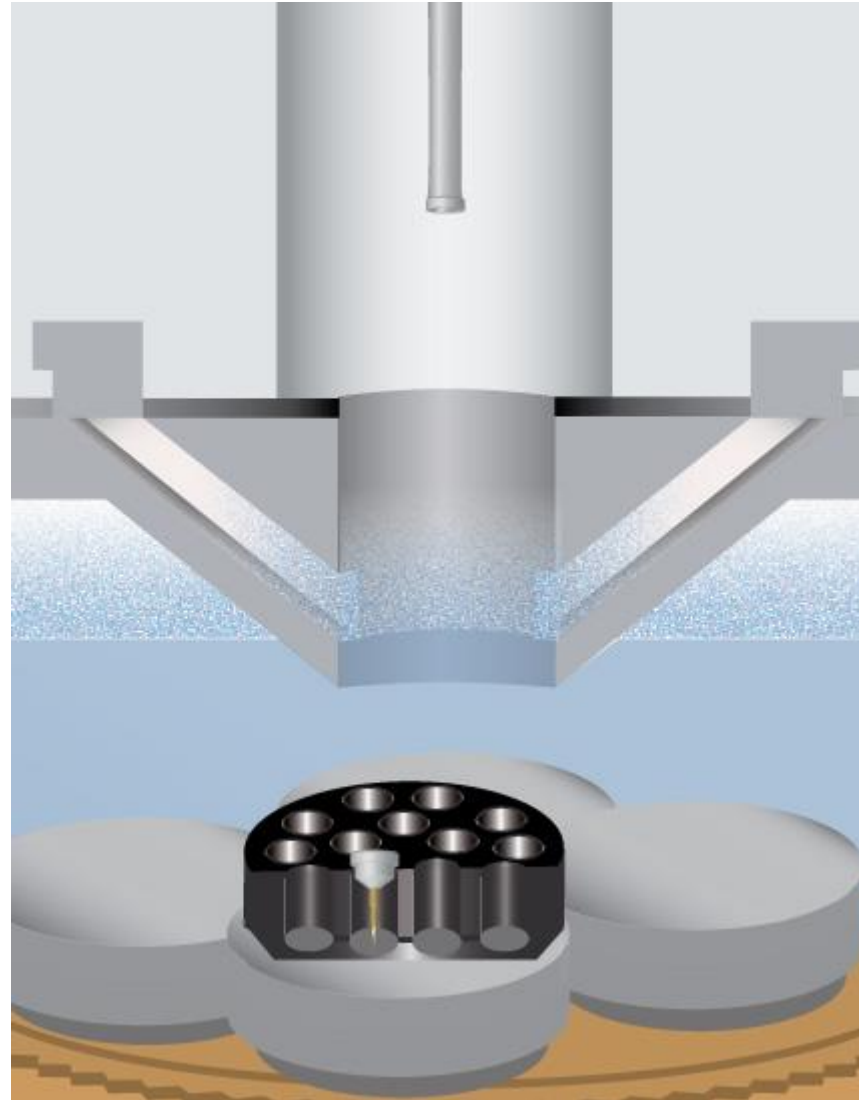


Cooling rate  
determined by  
sample transit time  
through cold gas,  
i.e., by plunge speed

# *NANUQ™*: Slow / Variable-Rate Cooling Plunge Sequence



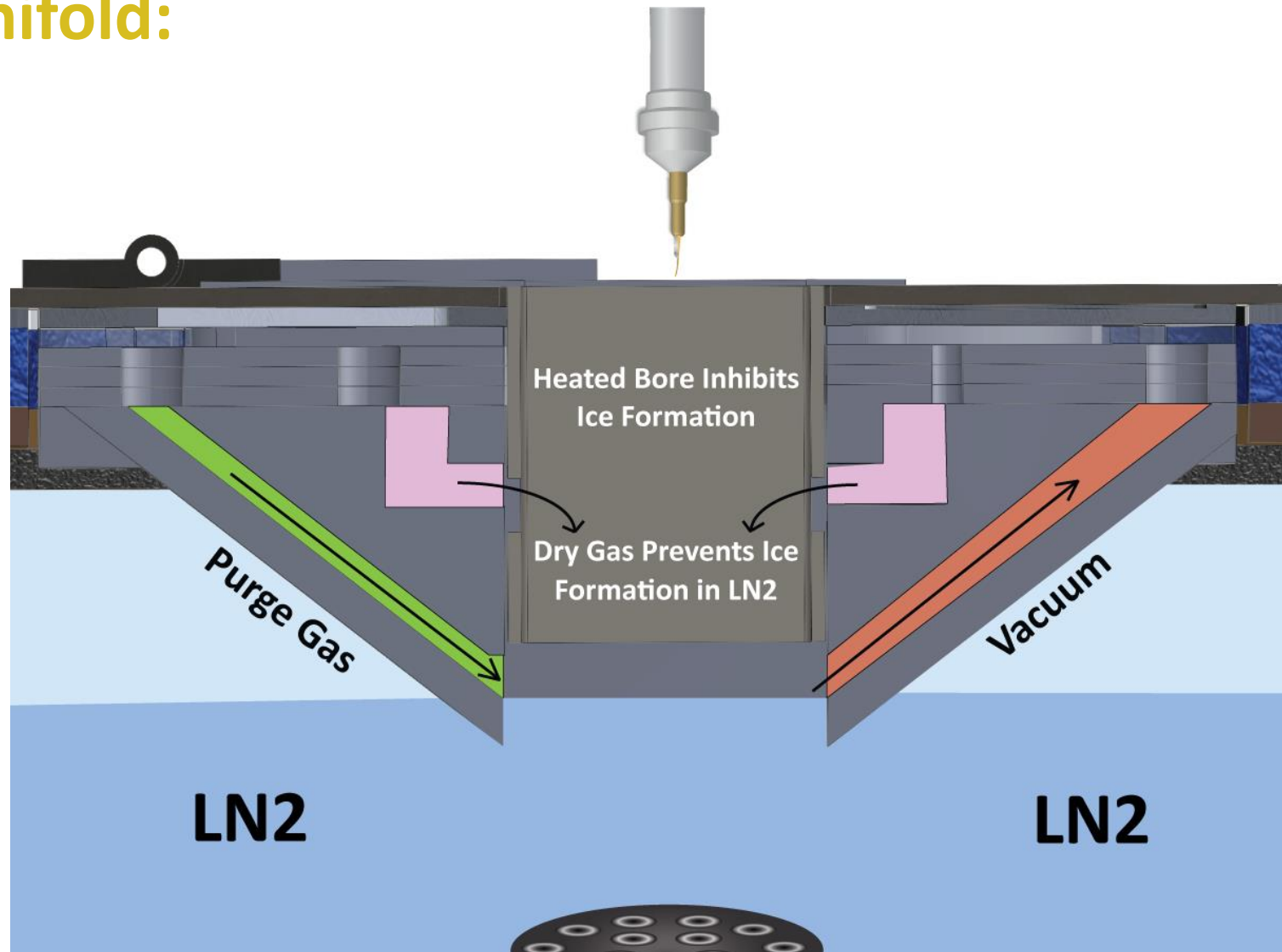
# *NANUQ™*: Slow / Variable-Rate Cooling Plunge Sequence





# NANUQ™: Key Design Features and Patented Technology

## Plunge manifold:



# NANUQ™: Key Design Features and Patented Technology

## Plunge manifold:

- **Controls plunge environment** to remove or define cold gas layers above the  $\text{LN}_2$  and provide a controlled temperature plunge path.
- **Eliminate cold gas** for fastest cooling.
- **Trap cold gas** for slow / variable rate cooling.
- Eliminates waves on the  $\text{LN}_2$  surface for fastest cooling.
- Isolates the  $\text{LN}_2$  and cold surfaces from ambient air, eliminating frosting.
- Computer-controlled dry gas flows, vacuum, and heaters **minimize  $\text{LN}_2$  consumption**.

# NANUQ™: Key Design Features and Patented Technology



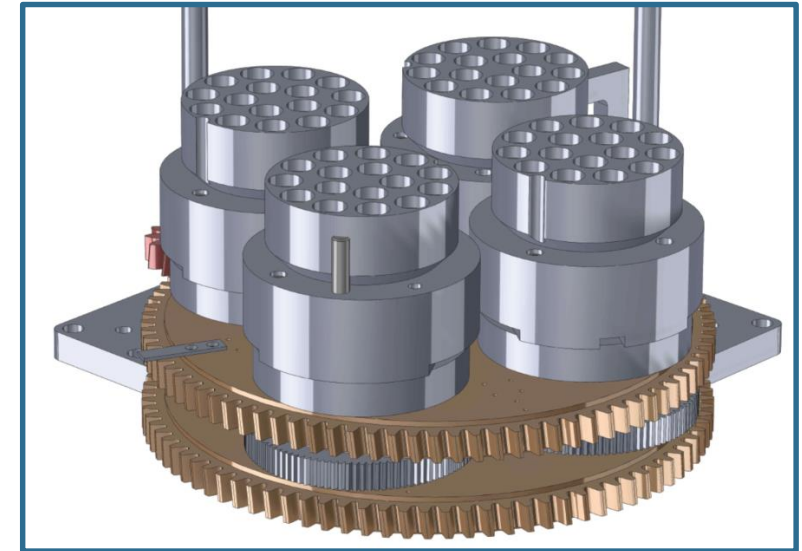
## High speed sample translation stage:

- Programmable sample plunge speeds from 3 m/s down to  $< 0.01$  m/s.
- Remove cold gas and plunge fast for fastest cooling.
- Trap cold gas and plunge slow for slower cooling.

# NANUQ™: Key Design Features and Patented Technology

## Automated sample puck loading:

- Carousel holds **4 Uni-Pucks** and 64 samples in  $\text{LN}_2$ .
- Pucks are positioned in a programmable sequence to capture cold samples and organize them for optimal synchrotron data collection.
- Pucks are easily loaded into *NANUQ* and then transferred to a dry shipper or storage Dewar.
- All **manual handling** of cold samples is **eliminated**.



# NANUQ™: Key Design Features and Patented Technology

## High capacity LN<sub>2</sub> storage:

- Large LN<sub>2</sub> capacity and efficient insulation allow continuous operation for **>12 hours** in high-throughput applications **without frosting**.



# NANUQ™: Key Design Features and Patented Technology

## Automated, computer-controlled operation:

- Automatically or manually trigger sample plunge
- Time from crystal harvest to fully cryocooled is < 4 seconds
- Sample cycle time < 15 secs; cool 4 samples/minute
- Minimizes time for crystal dehydration, maximizes throughput





ApplicationFile

HomeLN2 LevelPucksOperateMonitor

READY TO HARVEST  
AND LOAD SAMPLE

Running

START  
SEQUENCE

Auto plunge on door close

Plunge on command

Door

Wand In

	PUCK #	WELL #
Last filled	1	1
CURRENT	1	2
Next in Queue	1	2

Plunger Home

LN2 Level

Hatch

N2 pressure

Vacuum


Puck Ready

COMMs

SYSTEM READY

COMMs

NANUQ



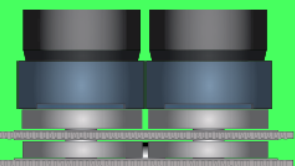
EMERGENCY STOP

When ready to proceed harvest sample insert Wand in holder close Door and press "Push to plunge"

At target level

Displacer Mode  
NORMAL

LN2 level 4.98 mm



13:47:22:67 - Thumb Homed

13:47:25:26 - Puck moving to position

13:47:30:44 - Puck in position

auto scroll

Well Fill Color Key

Well/puck not loaded

Well not selected

Well selected as slow plunge

Well selected as fast plunge

Well filled

Error filling well

Puck 1

Puck 2

Puck 4

Puck 3

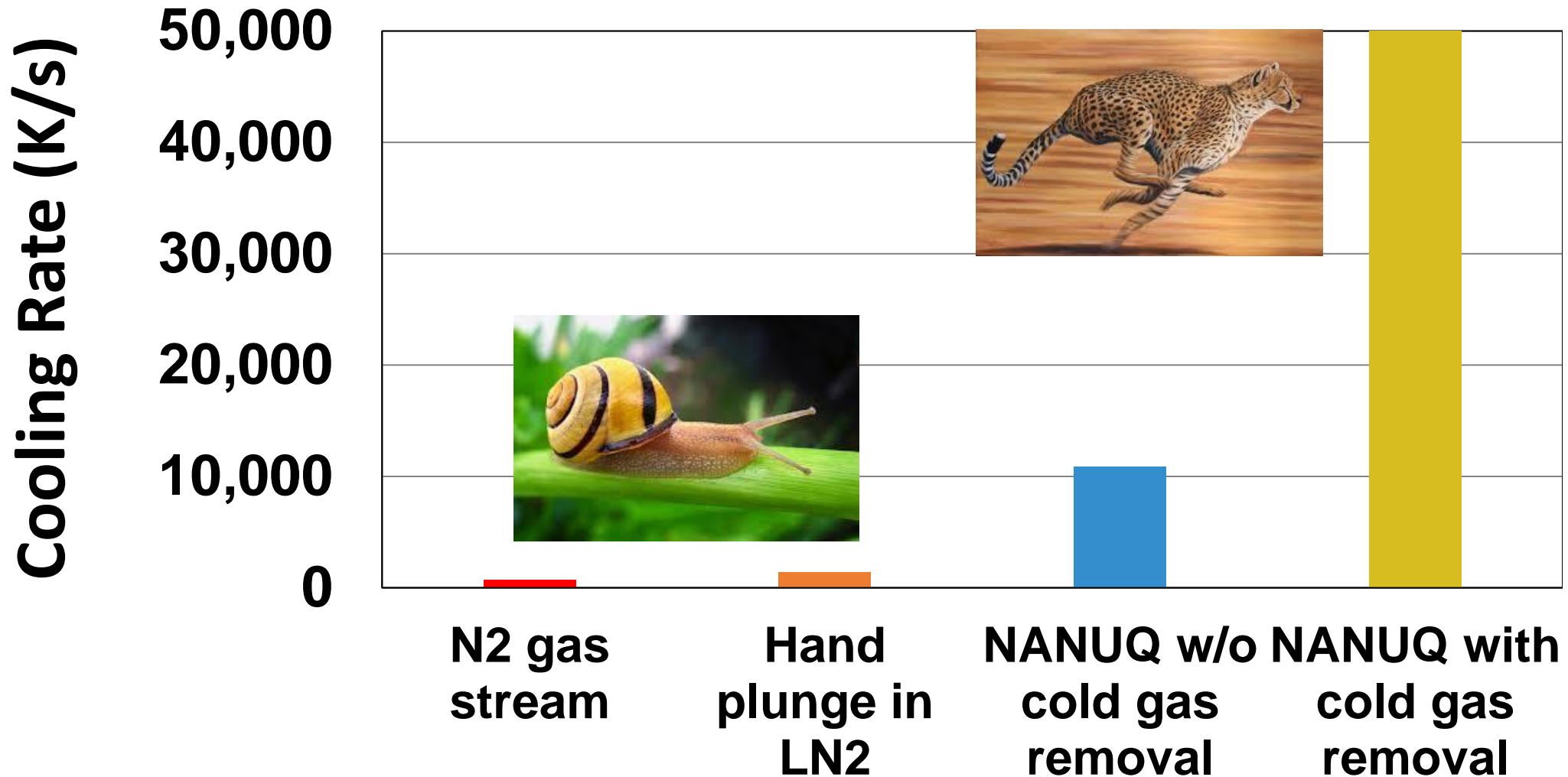
scroll position  
33

MiTeGen

27

# ***NANUQ™* : Key Advantages**

**Maximize cooling rates using LN<sub>2</sub>:**



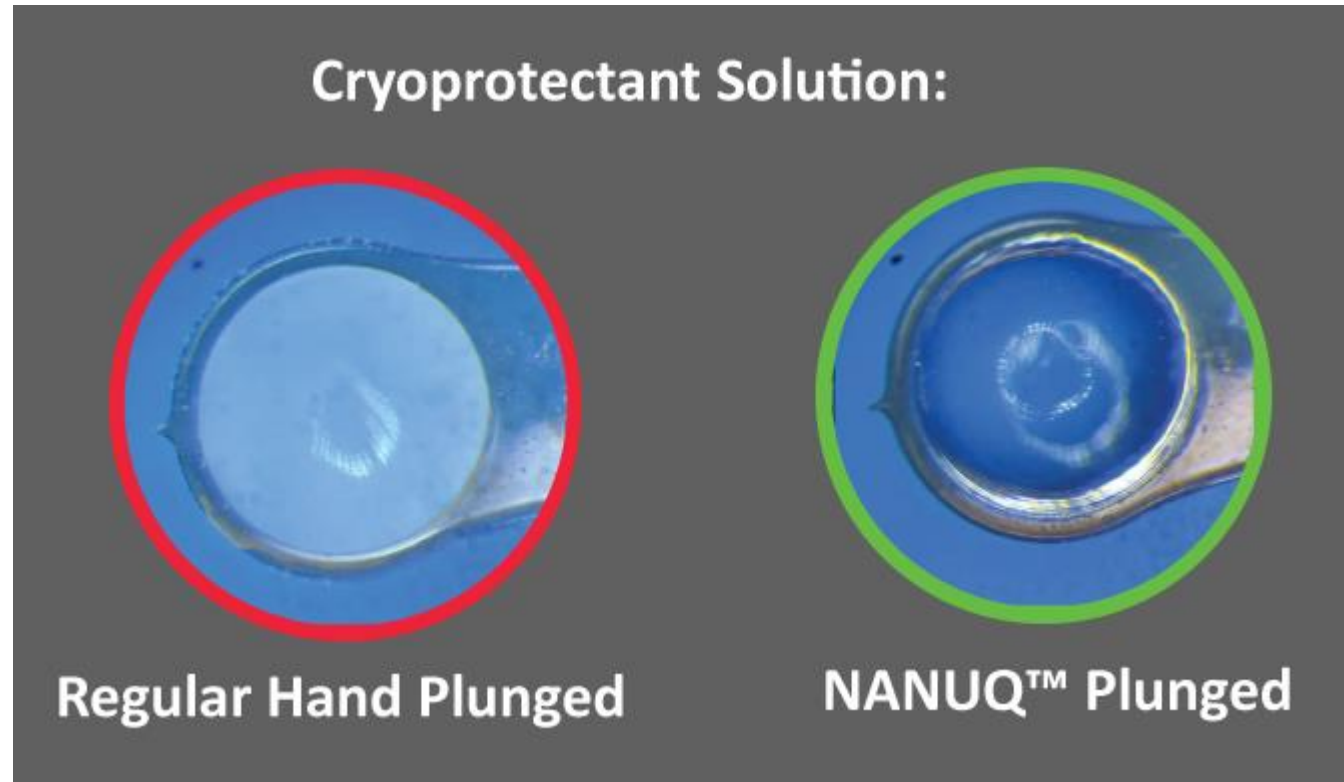
# *NANUQ™* : Key Advantages

## Maximize cooling rates using LN<sub>2</sub>:

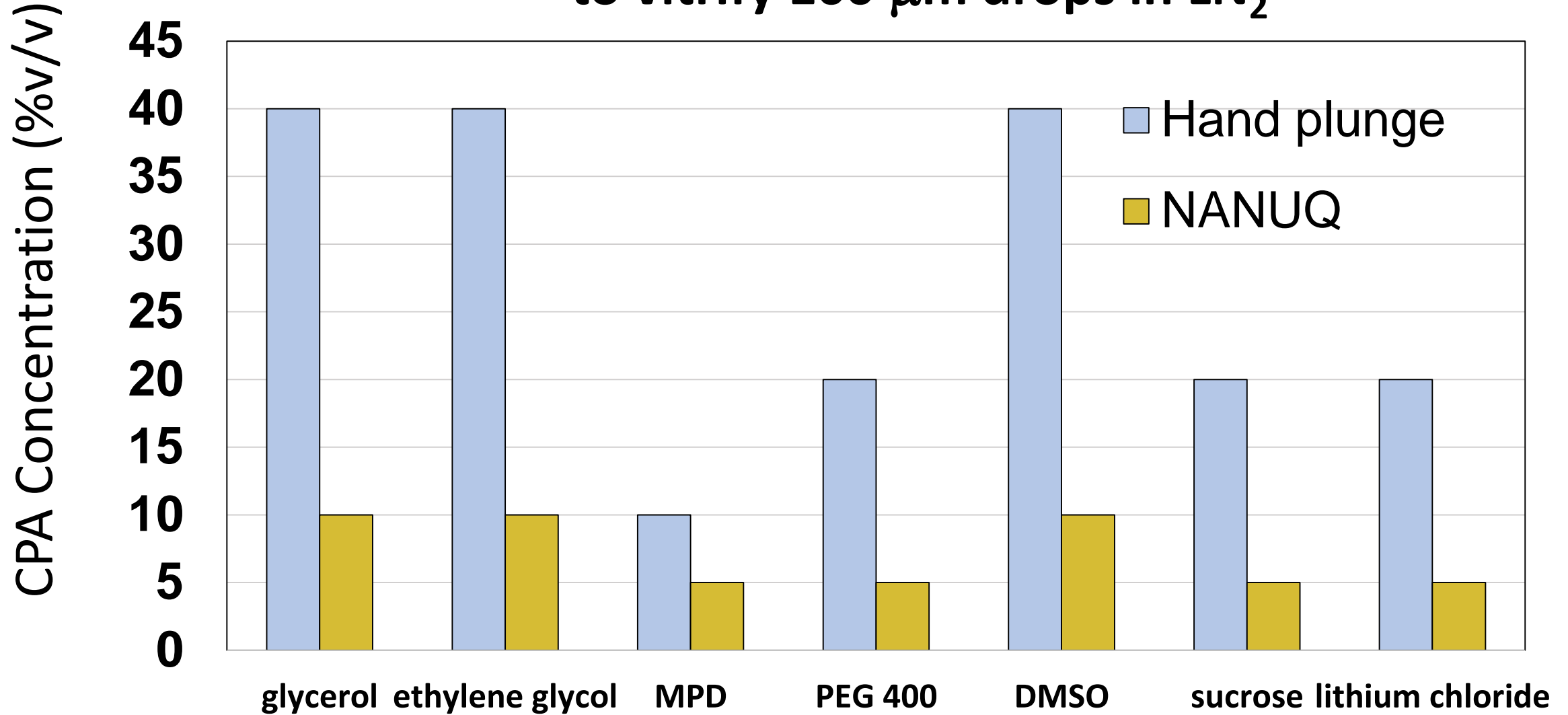
- Cooling rates to **>50,000 K/s**.
- **Cooling times < 2 ms**  $\Rightarrow$  **better kinetic capture** of room temperature structure & ligand binding configuration.
- Can also cool at **intermediate (100 K/s)** and **slow (down to 1 K/s)** rates to **simulate gas stream cooling** and for quasi-equilibrium cooling, which may give better diffraction outcomes (e.g., for large crystals).
- **Highly reproducible cooling**  $\Rightarrow$  science-based **optimization of cooling and diffraction**.

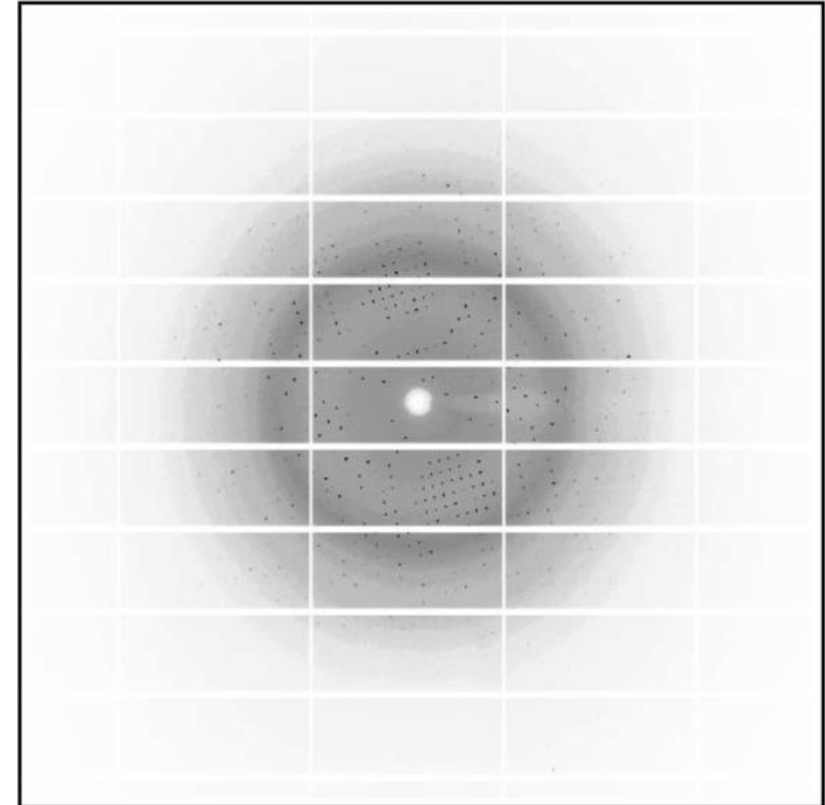
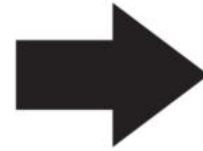
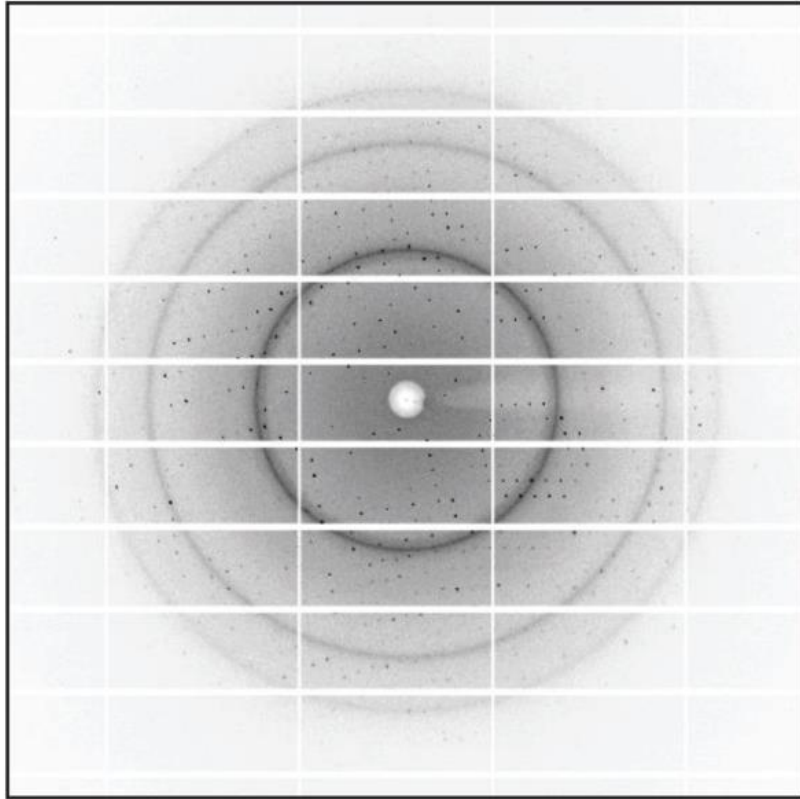
# *NANUQ™*: Key Advantages

**Minimize cryoprotectant concentrations:**



## Minimum CPA concentration (%v/v) to vitrify 100 $\mu\text{m}$ drops in $\text{LN}_2$





**Hand plunged (LN2)**  
**Sample: CA IX-mimic Crystal**  
**Cryoprotectant: 20% sucrose**  
**Result: Significant ice rings**

**Cryocooled on NANUQ™ (LN2)**  
**Sample: CA IX-mimic Crystal**  
**Cryoprotectant: none**  
**Result: No ice rings**



# *NANUQ™*: Key Advantages

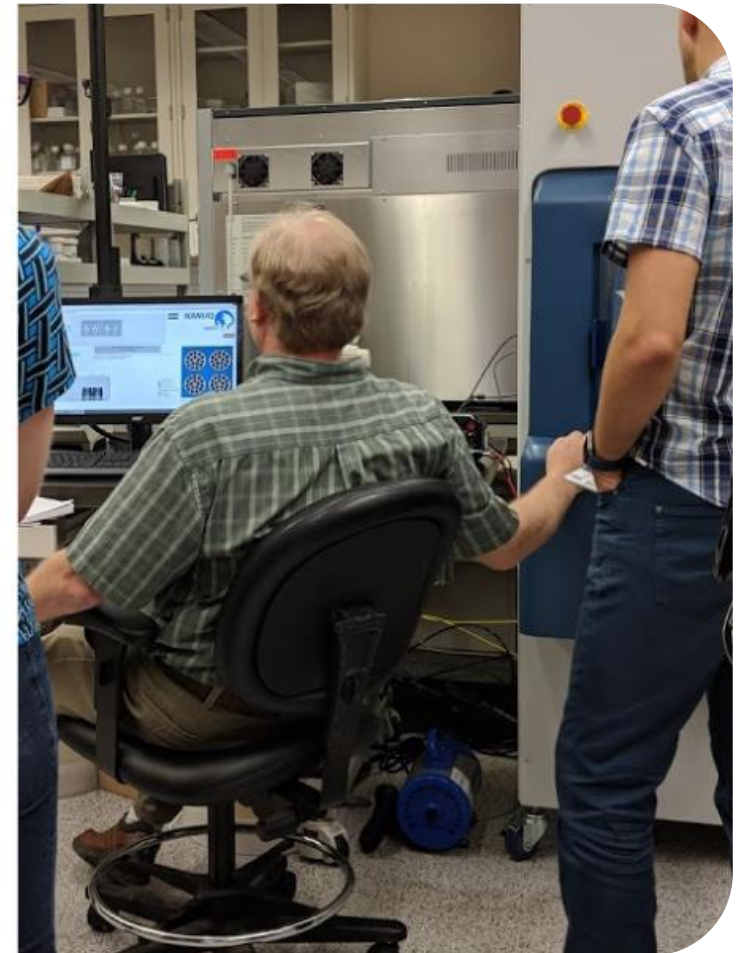
## Use less (no) cryoprotectant, simplify cryoprotection:

- **No penetrating cryoprotectants required** to prevent internal ice. Only add for contraction matching.
- **No cryo soaks necessary** - just quick swipes through a dilute cryo solution / oil to protect / remove external solvent.
- No crystal damage or dissolving by cryoprotectants.
- **Ideal for ligand/fragment screening:** no ligand displacement by or confusion with cryoprotectants.

## **NANUQ™: Key Advantages**

- Cool **immediately** after harvesting, with **complete confidence**.
- **No ice** from **cooling**, **LN<sub>2</sub>** or **frosting** during puck loading.
- **No lost crystals** due to manual plunging / loading accidents.
- Easily **sequence and track** samples in pucks for optimized data collection.
- Get **everyone** in your lab using the **same reliable protocols**.
- Provides the **reproducibility** and **flexibility** to rationally optimize your cooling protocols.

# How Can You Get the Technology?



**NANUQ™ shipped and installed at a customer location.**

We can help with instrumentation proposal writing and justification statements.

# Access the Technology: *NANUQ*<sup>™</sup> Facility in Ithaca, NY Open to Users



# NANUQ™ Access Facility

## Cool your crystals with NANUQ™

### Getting Your Crystals to MiTeGen:

- Bring your crystals via car, or ship them to us.
- Come in advance and set up plates. We can monitor them.
- Send your solutions + plates. We can set them up manually & monitor them.

### Cooling Your Crystals:

- Do it yourself with our support.
- Have our expert staff harvest, cool, & ship them anywhere.



# NANUQ™ Access Facility

## Cool your crystals with NANUQ™

### Available equipment:

- Vibration-free Crystallization Incubator
- Zeiss V8 & V20 Microscopes with imaging
- Watershed™ Humidity Controlled Harvesting
- Worthington HC34 & HC35 Refrigerators
- Worthington CX100 Dry shippers
- Harvesting loops, tools, materials....



# NANUQ™ Access Facility

## How to Apply for Access Time

1. Visit our NAF Website:  
- [www.mitegen.com/NAF](http://www.mitegen.com/NAF)
2. Fill out the Request Trial form
3. Our team will be in touch to get the process started!

We look forward to having you see the difference NANUQ™ can make!



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## NANUQ™ Access Facility

UTILIZE ADVANCED CRYOCOOLING TO IMPROVE RESULT

Introductory trial experiments to give researchers access to the impact of advanced cooling techniques.

Try preparing your samples with NANUQ™ cryocooling technology to:

- Prevent damaging ice formation
- Improve sample diffraction quality
- Reduce or eliminate cryoprotection
- Guarantee reproducible crystal-to-crystal cooling

- Visit the facility and cool them yourself, or
- Ship us samples for us to cool, or
- Have us set up trays and perform the cryocooling

**REQUEST TRIAL INFO**

Name \*

Email \*

SUBMIT





# *Thank you!*

**- From Everyone at MiTeGen!**

**Contact:**

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