

NSLS-II and research into combating COVID-19 at BNL

J.P. Hill, Director, NSLS-II

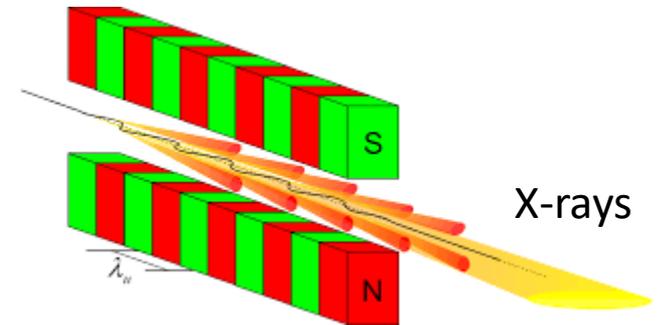
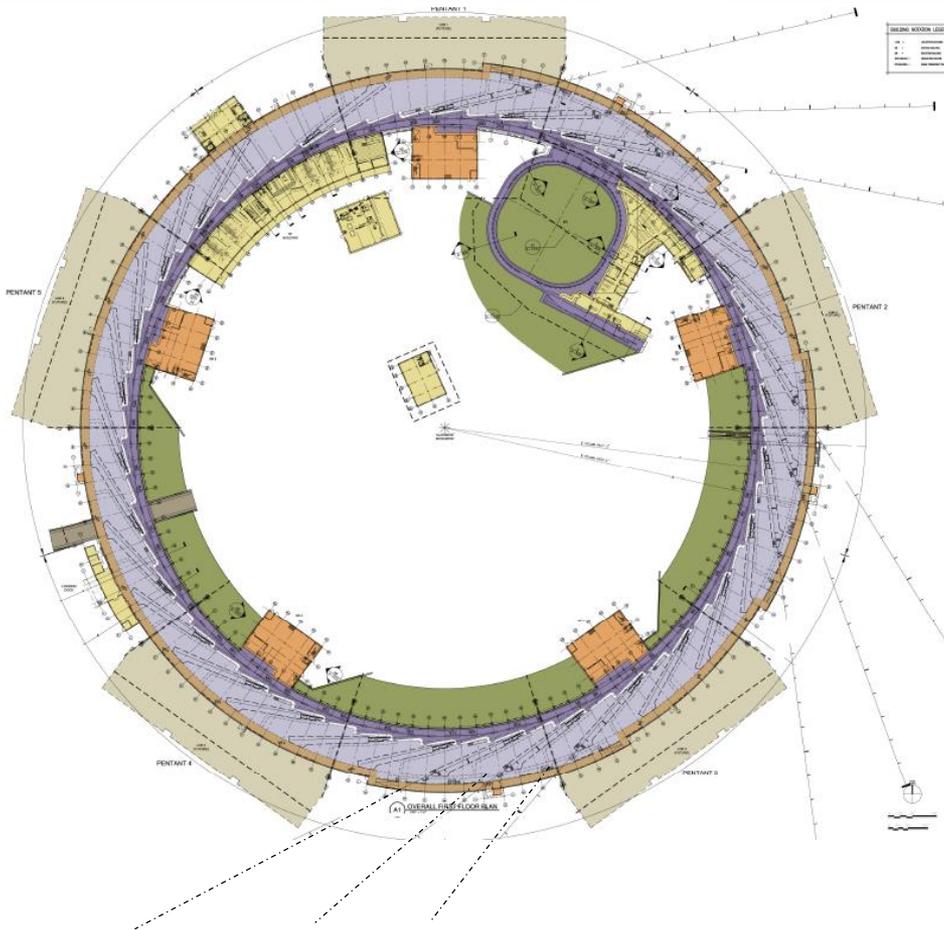
CAC May 15th 2020

NSLS-II – A World-Leading Microscope

- NSLS-II is a state-of-the-art light source covering far infrared to hard X-rays
- Offers best in-the-world characterization capabilities
- Compatible with devices, real-world materials and “operando” conditions
- ~1 km around, can fit Yankee Stadium inside the ring



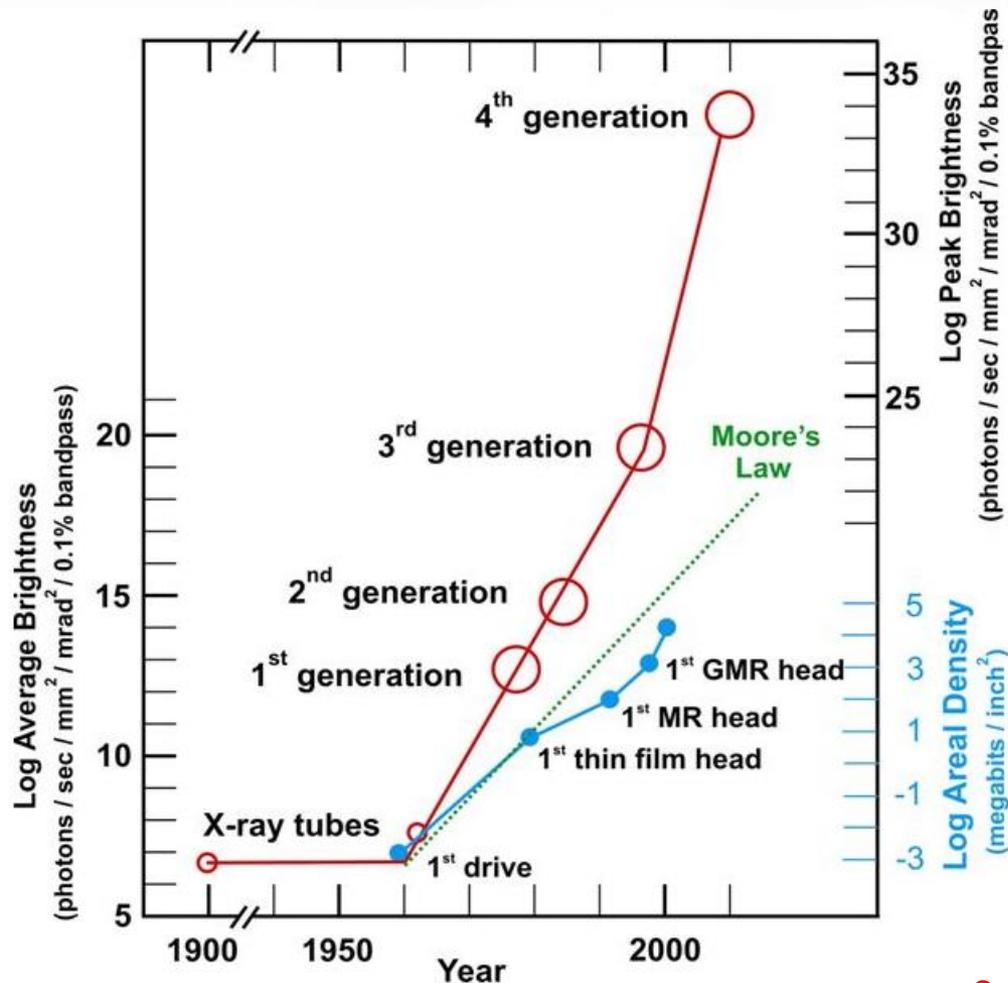
NSLS-II: Best in class from far-IR to hard x-ray



User Facility

- Capacity for ~ 60 beamlines
- 1800 Users in FY19
- Proposal access. Free if intend to publish.
- Proprietary fee (\$430/hr)

X-ray sources brightness vs time and Moore's Law



Improvements in x-ray sources have outpaced even Moore's Law in microelectronics.

This has driven advances in the x-ray field that are as revolutionary as seen in the field of computing

Credit: J. Stohr

NSLS-II capabilities

Exquisite resolution and sensitivity, and the expertise to use them.

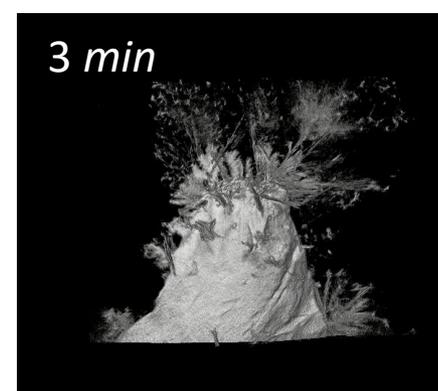
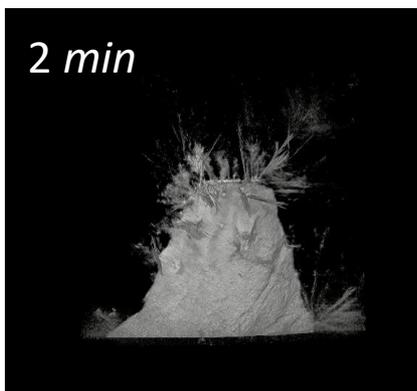
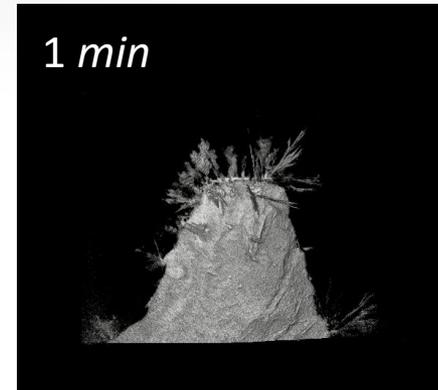
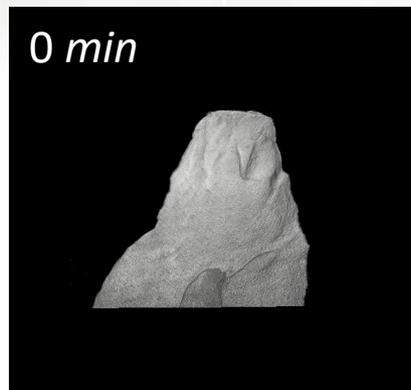
Some examples of the work we do:

- 1) **Energy storage:** Chemistry in real time on the nm length scale
 - Can we prevent your cell phone batteries dying?
 - Can we store wind-generated electricity?
- 2) **Materials synthesis:** Novel nano-functional materials
 - Can we build better solar cells?
 - Can we design new ways to deliver medicines?
- 3) **Environmental Science:** understanding toxins moving through plants
 - How can we protect the food web?
- 4) **Microelectronics:** Legacy electronics and state-of-the-art
 - Can we ensure our electronics perform as we expect?
- 5) **Magnetism:** Spintronics
 - What materials are beyond Moore's law?

Energy Storage

Full field imaging

- 30 nm spatial resolution over 100 micron field of view
- 3D images in seconds
- x10 faster than any similar instrument in the world
- Allows first 3D movies of dynamic processes



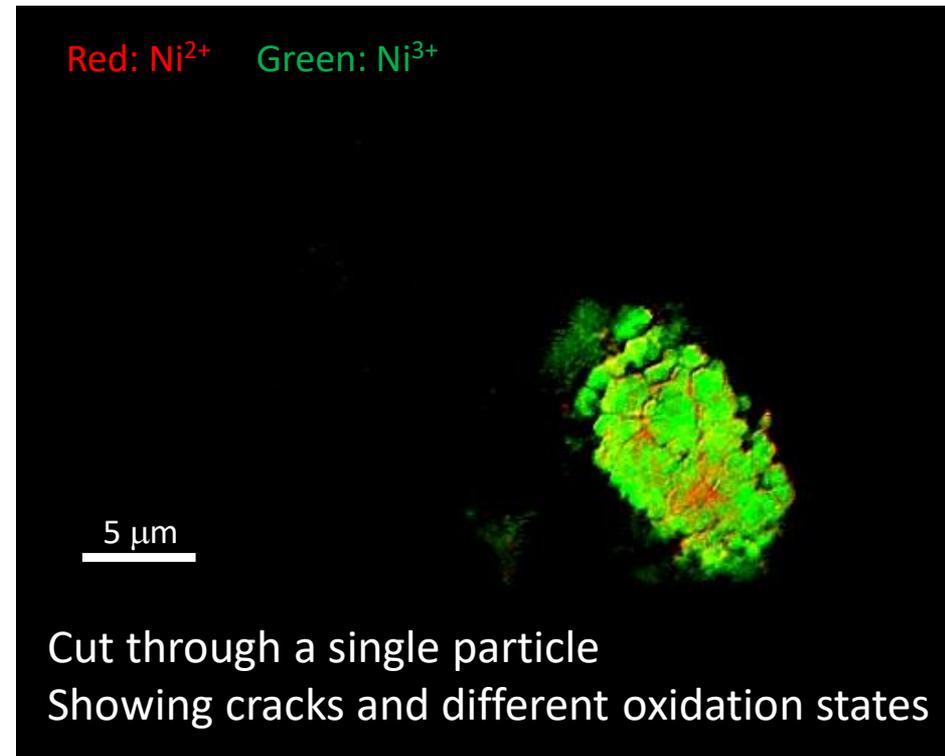
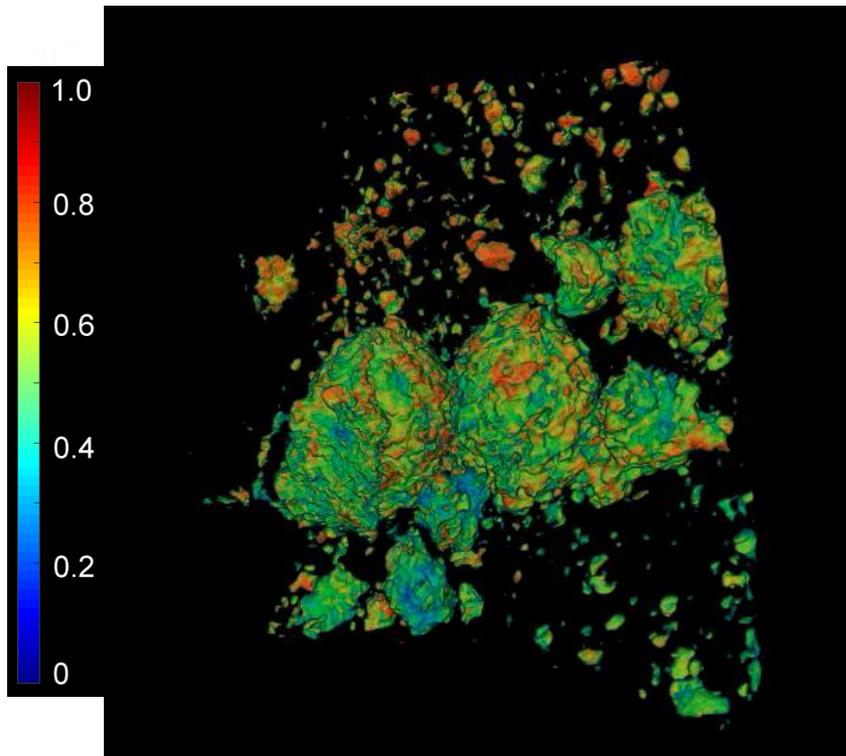
Appl. Phys. Lett (2018)

Tracking 3D silver nano-dendritic growth in real time,
under *in-situ* chemical reaction conditions: $\text{Cu} + \text{AgNO}_3$
 $\rightarrow \text{Ag} + \text{Cu}(\text{NO}_3)_2$

Energy Storage

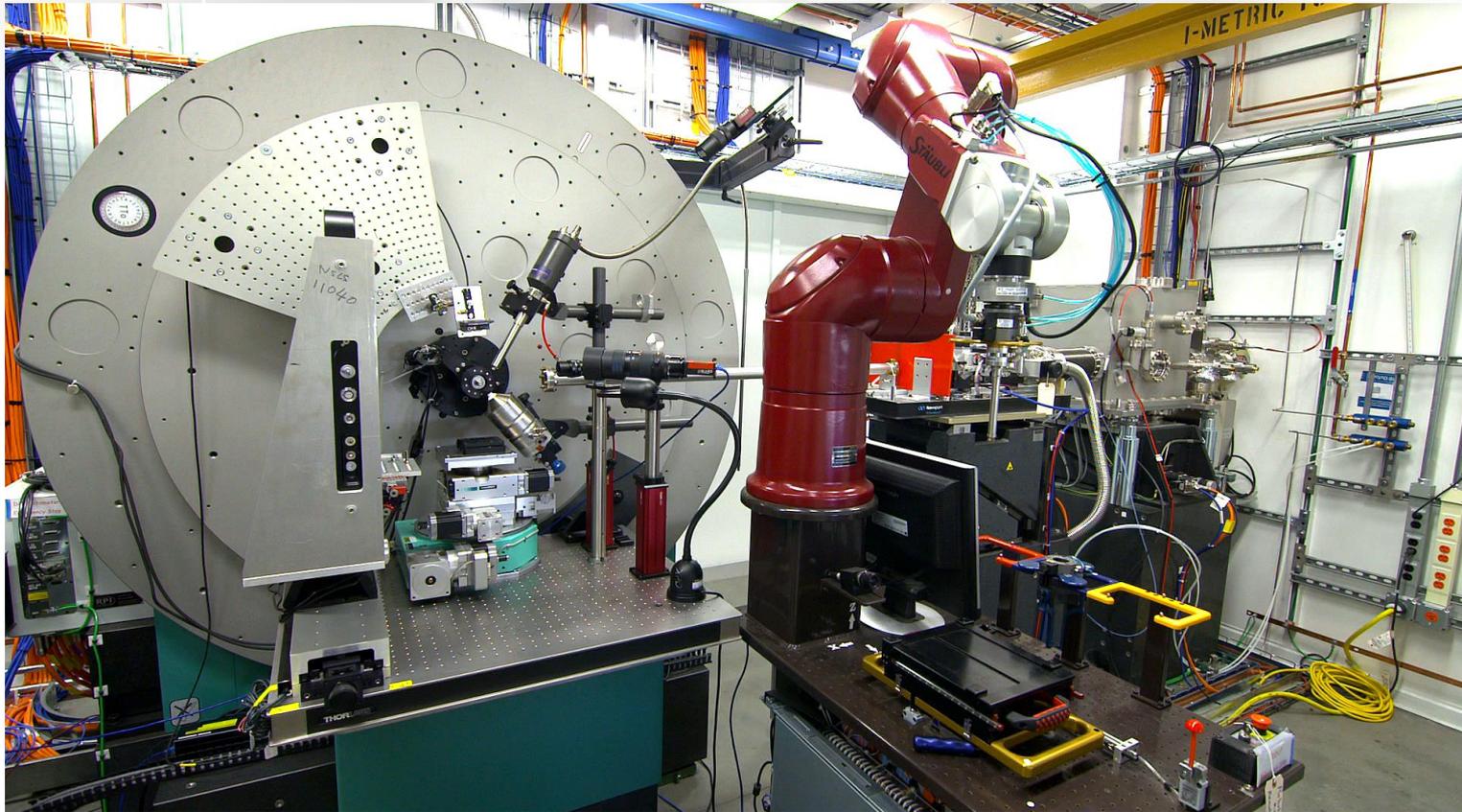
Brightness of x-ray beam allows very fast imaging with nanoscale resolution in 3D

3D rendition of Ni^{3+} concentration



Materials Synthesis

Measurements of atomic structure

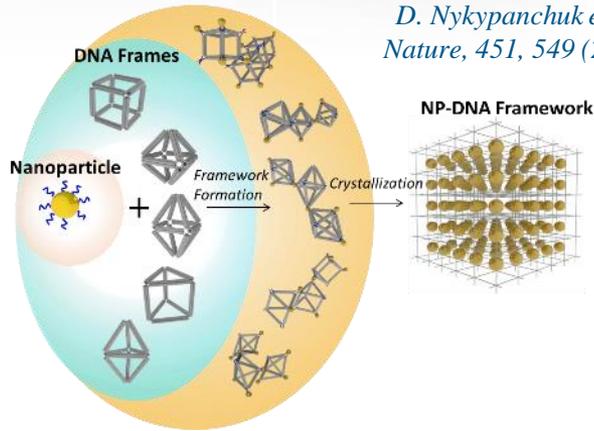


- Precise measurements of atomic structure *in-situ*
- Synthesis, batteries, catalysts, nanoparticles,.....
- Robot to process large numbers of samples quickly

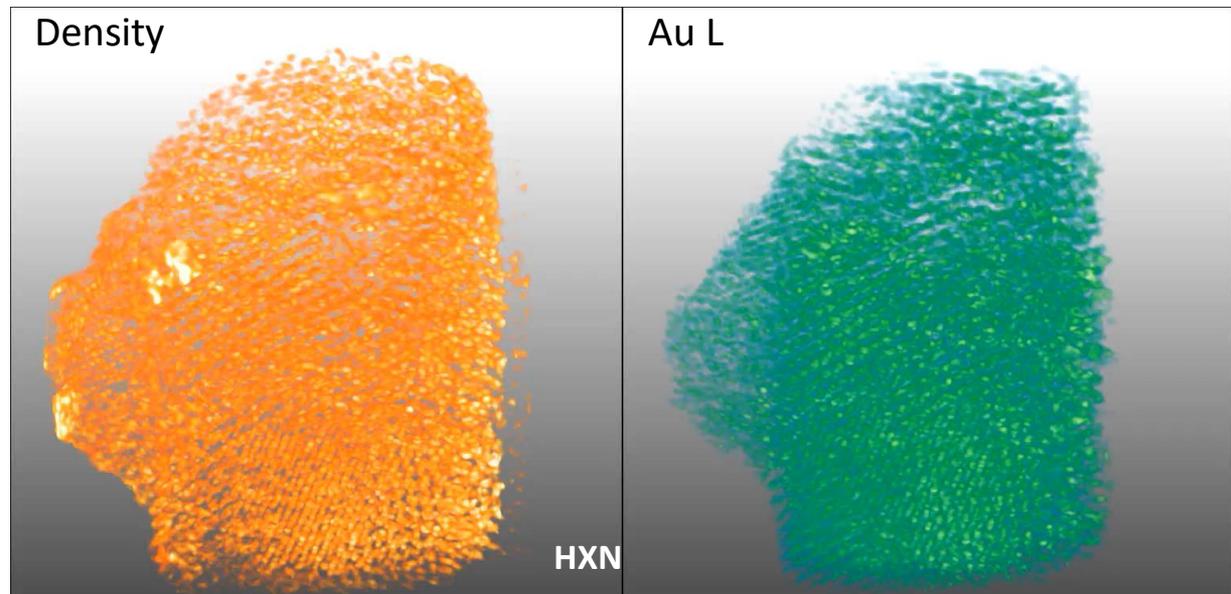
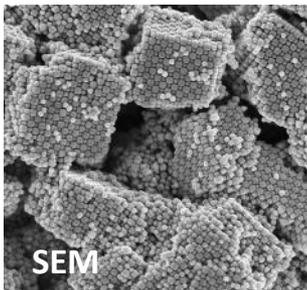
DNA guided assembly

O. Gang (CFN), H. Yan (NSLS-II)

*D. Nykypanchuk et al,
Nature, 451, 549 (2008)*



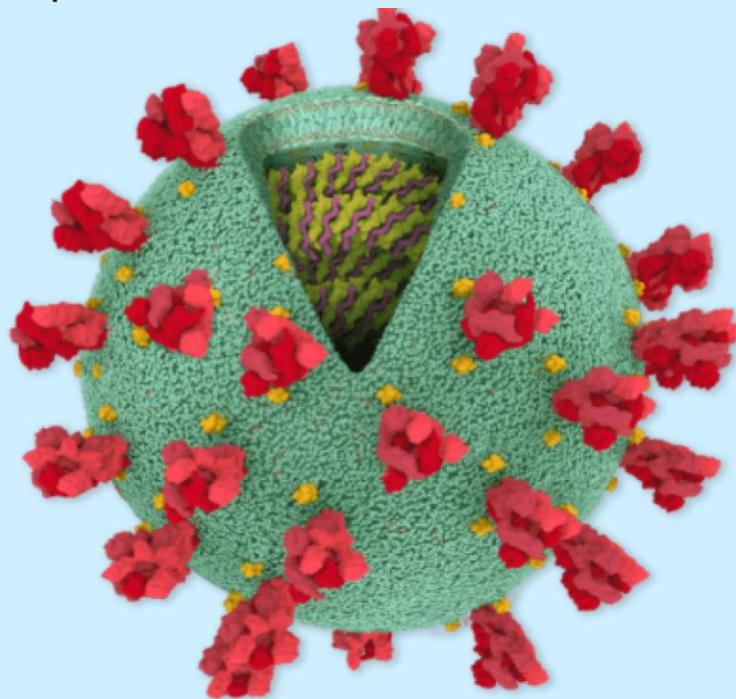
- DNA-guided 3D assembly process offers the potential to synthesize entirely new materials



SARS-COV-2 research at NSLS-II

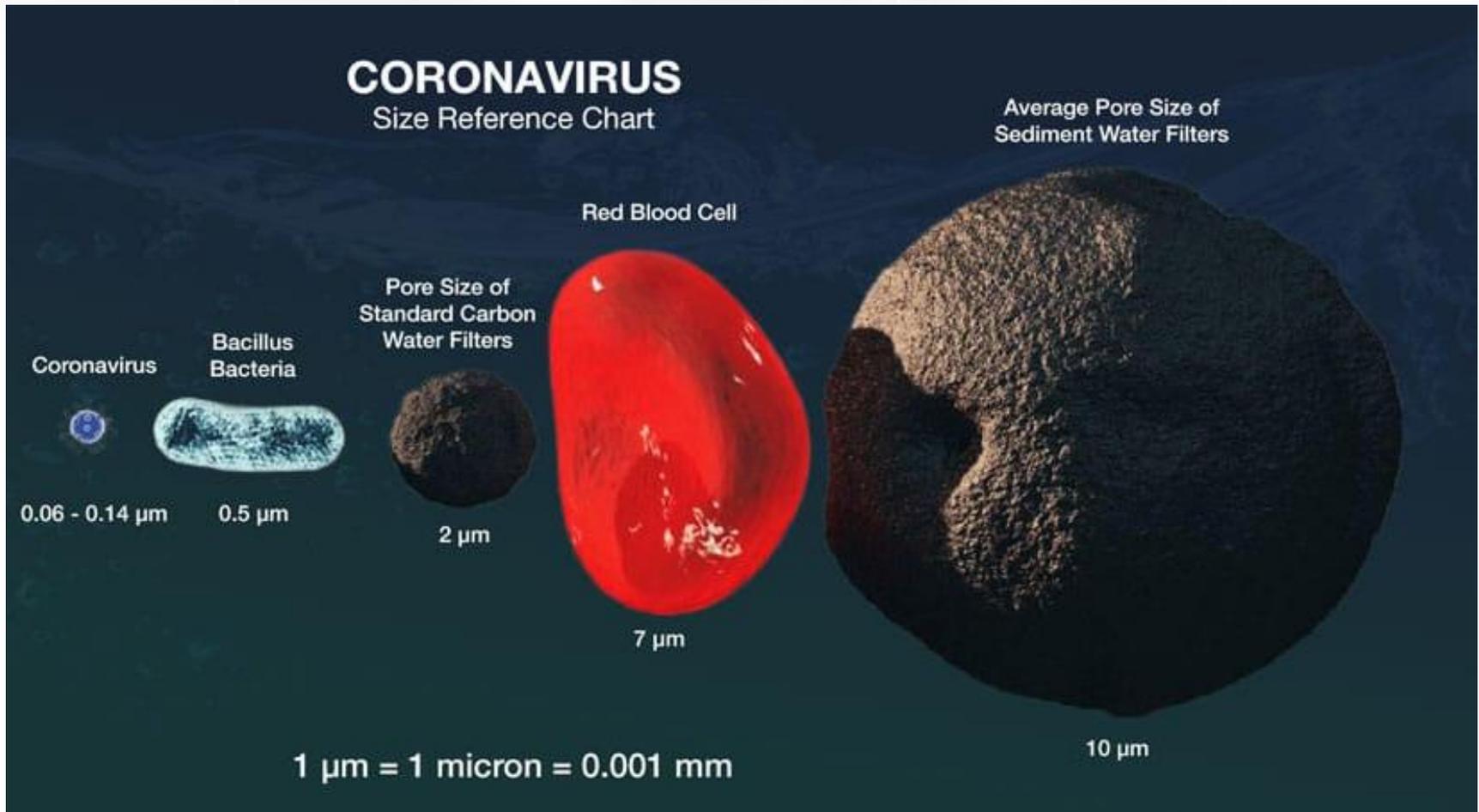
SARS-COV-2

NY Times April 3rd

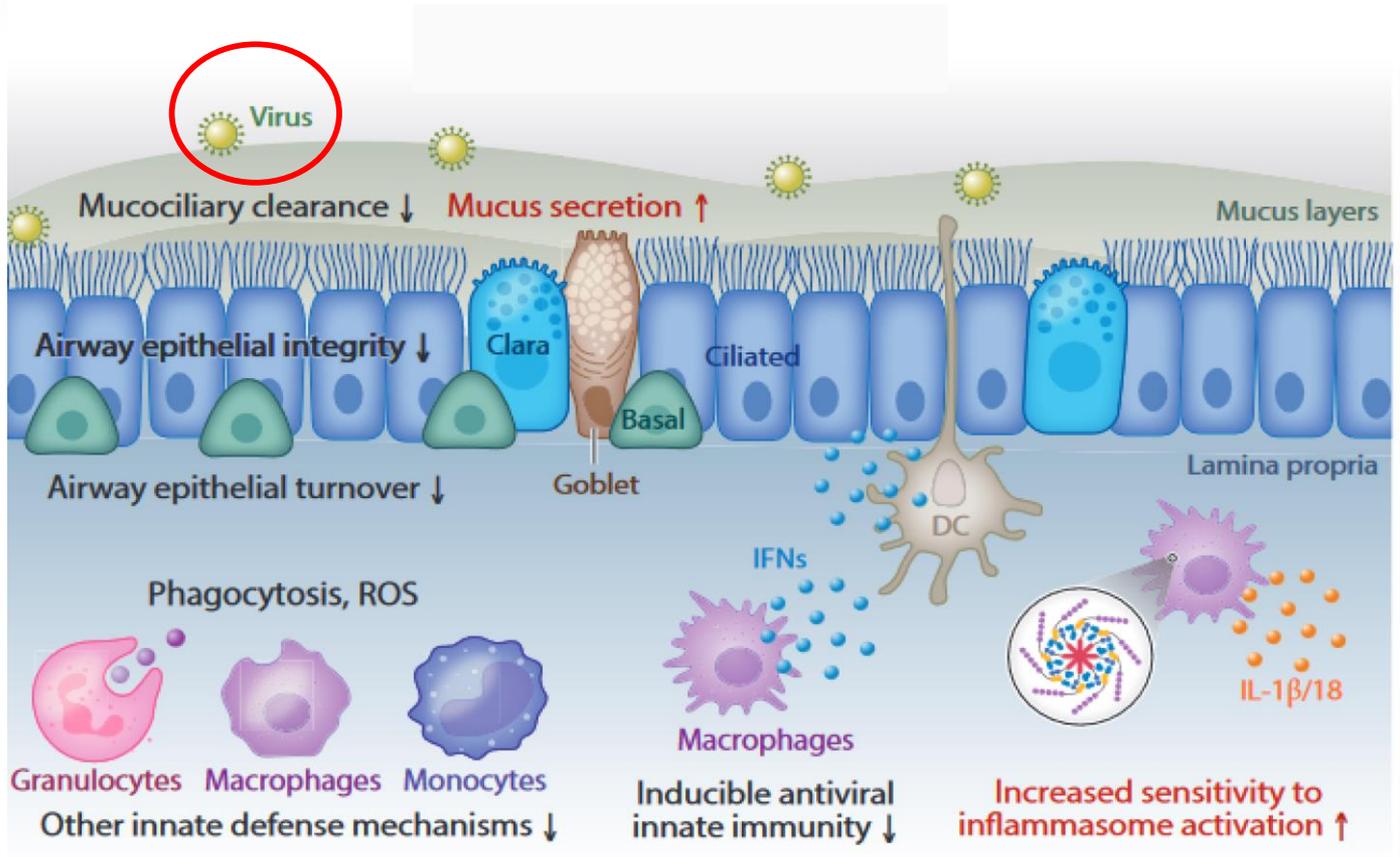


A virus is “simply a piece of bad news wrapped up in protein,” the biologists Jean and Peter Medawar wrote in 1977.

Size of virus particle



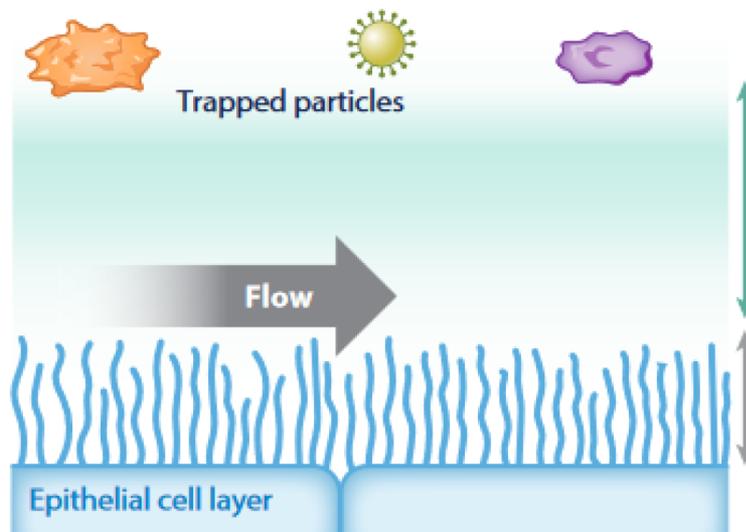
Respiratory Tract



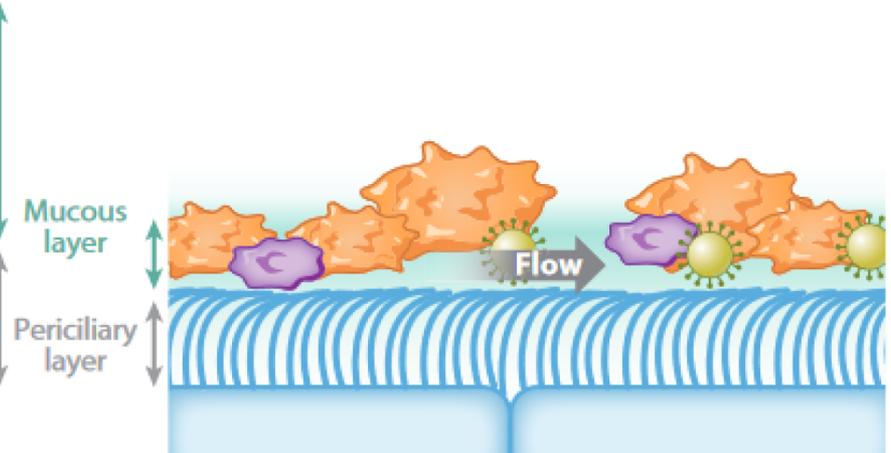
Moriyama et al, Ann Rev Virol, 16 March 2020

Primary Defense

a Humid breathing air (hydrated)



b Dry breathing air (dehydrated)



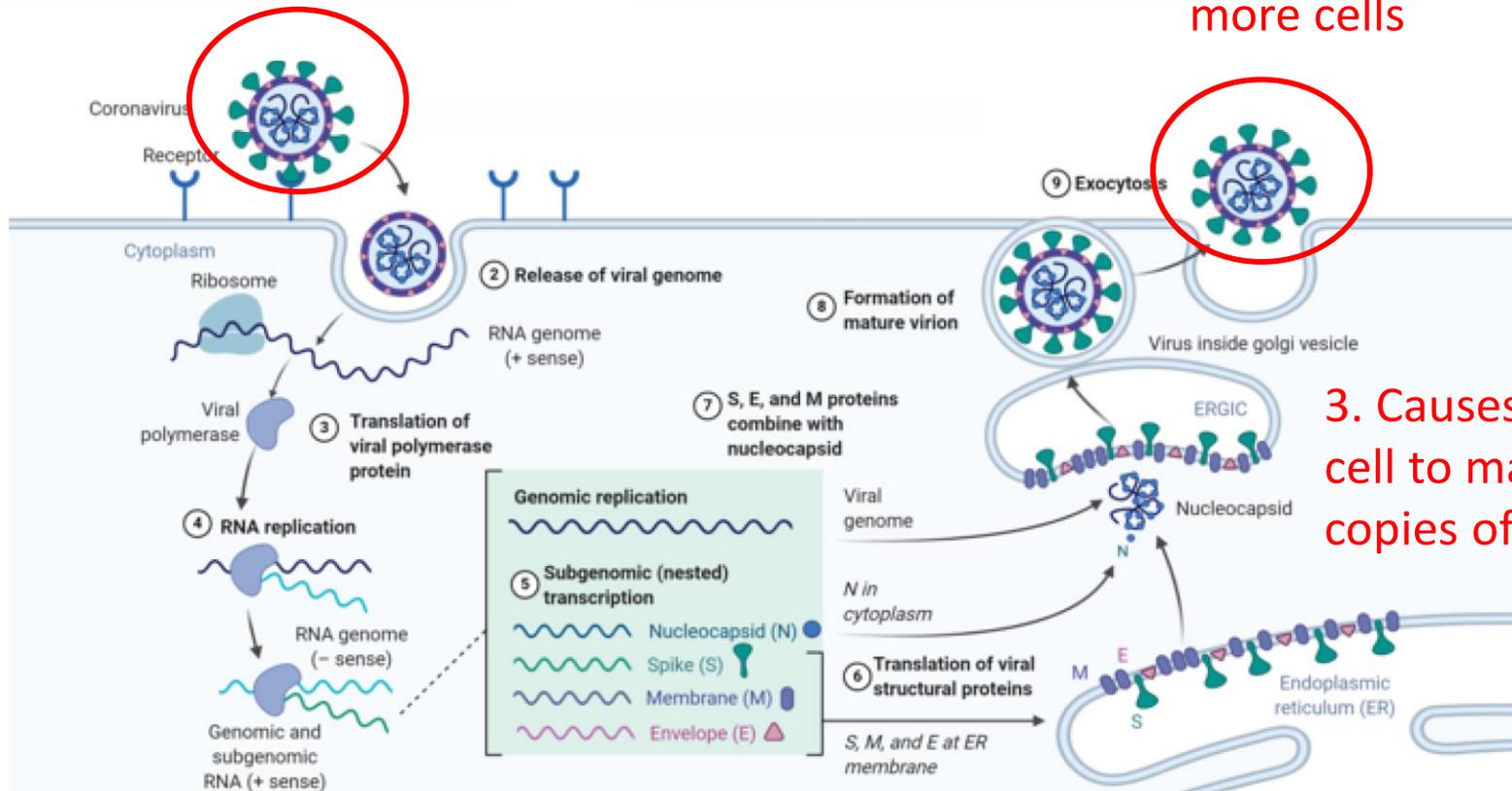
Moriyama et al, Ann Rev Virol, 16 March 2020

Keep hydrated. Use room humidifier

How the virus infects you

1. Virus binds to a “receptor” on your cell

4. Ejects those copies into the body to infect more cells



3. Causes your cell to make more copies of the virus

2. Releases its genome inside your cell

Protein structure

We need to understand the precise shape of the key proteins involved in the replication of the virus in order to interrupt it

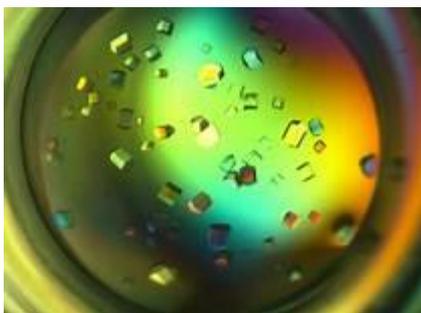
A lot has been learned since Jan 2020:



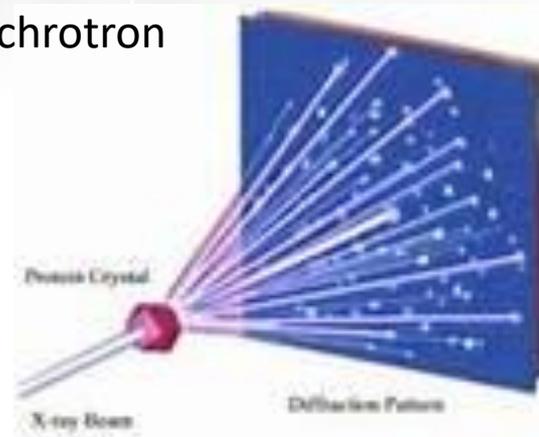
The spike protein structure

How to determine the atomic structure

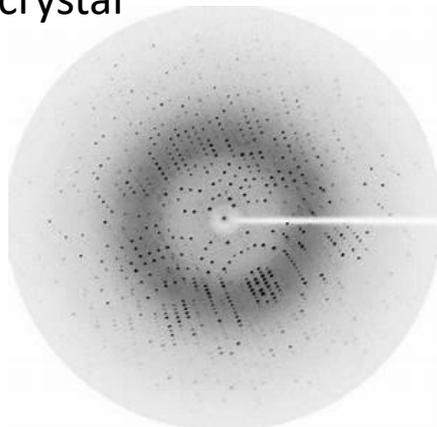
1. Protein crystals



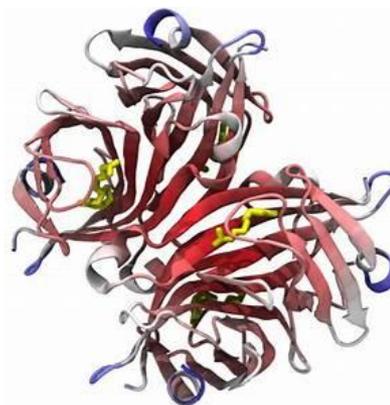
2. Synchrotron x-rays



3. X-ray diffraction pattern from crystal



4. Atomically precise protein structure

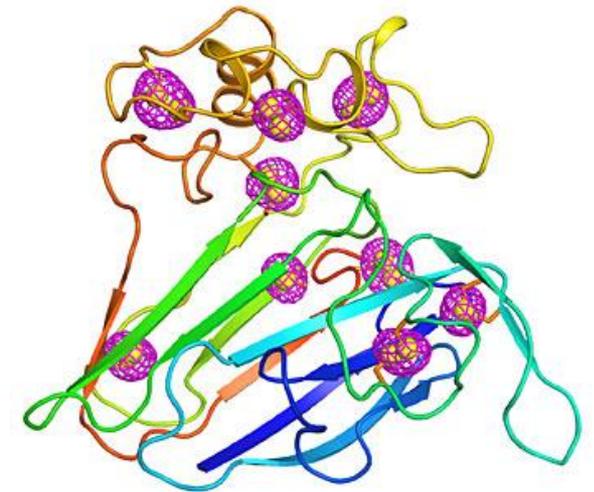
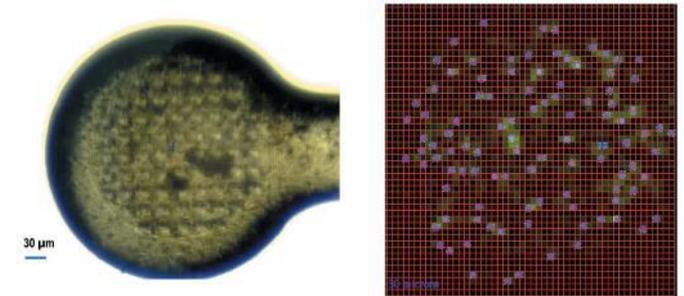


96% of all drugs approved by the FDA in the last 15 years used synchrotron x-rays.

How does NSLS-II help?

- X-ray scattering from protein crystals gives structure to atomic precision
- Growing large enough single crystals is the bottleneck in determining structure
- Automation and intense x-ray beams at NSLS-II allow the study of smallest crystals in the world
- Down to ~ 1 micron ($1/100^{\text{th}}$ of the width of a human hair)
- This greatly speeds up solving the structure and searching for new drugs

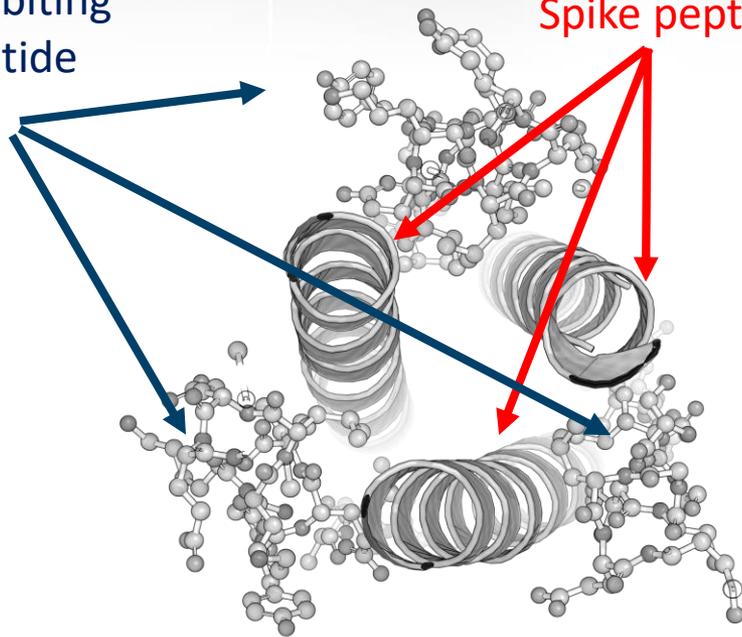
Guo et al. *IUCr J.* 6, Part 4, July 2019.



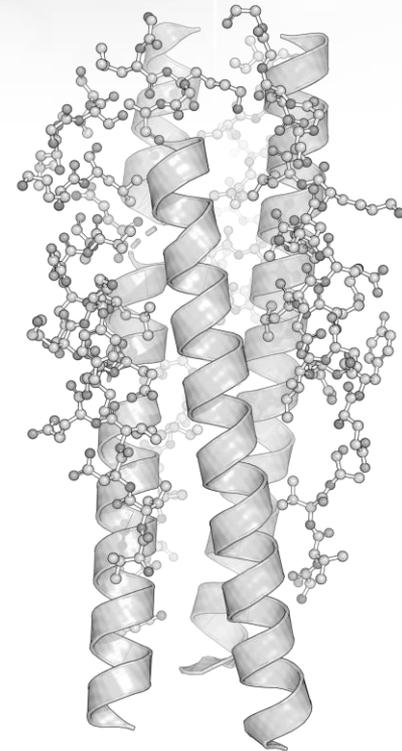
Some NSLS-II results:

Inhibiting peptide

Spike peptide



Top view



Side view

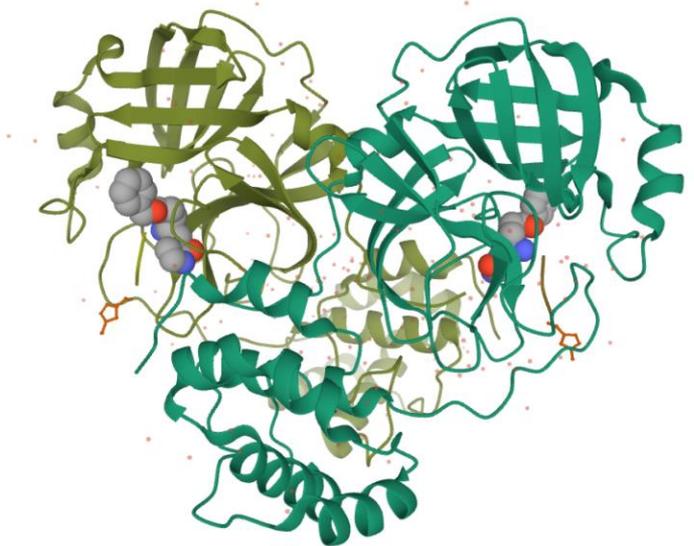
Goal is to find something that binds with the spike protein and prevents it fusing with our cell membrane

Peptide = piece of a protein

Inhibiting viral proteins

Another approach at NSLS-II is to try and disrupt the viral proteins once they are inside your cell

The main protease from COVID-19, with ligand bound



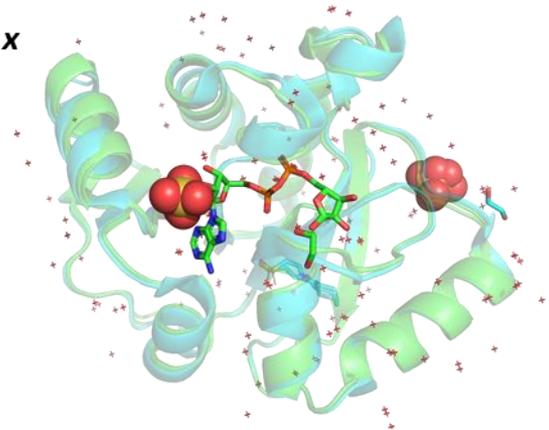
Is it possible to repurpose existing (FDA approved) drugs to inhibit this proteins activity?

We have grown crystals of this and are presently trying various drugs to see if we can determine how and where they bind

How is it safe?

1. We are NOT looking at the virus itself. We are looking at pieces of some of the proteins
2. These pieces cannot infect a cell
3. The pieces are synthesized in a lab, knowing the gene sequence of the viral protein. They do NOT come from the virus. There is no chance of a virus particle being included in the synthesis
4. Think of them like “photocopies”. They are inert
5. The crystals are kept at liquid nitrogen temperatures and tracked in and out of the lab. A robot handles them at the beamline

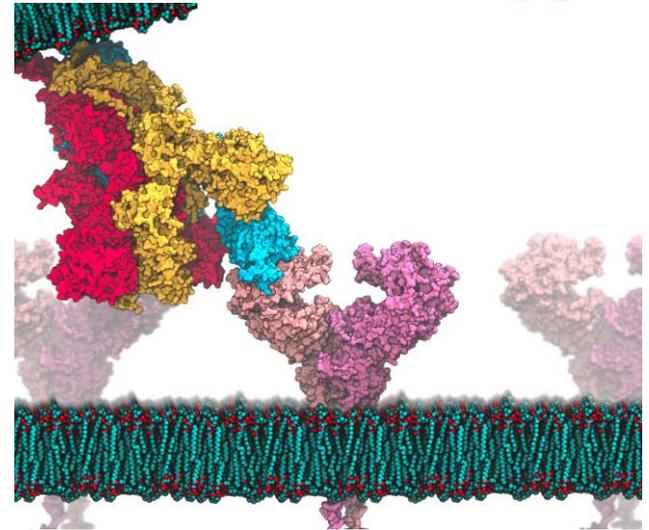
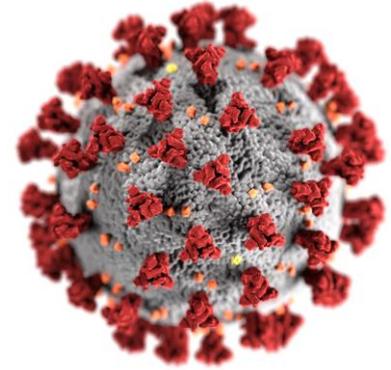
ADRP x
2



Computational Research

Computer-based Drug Discovery

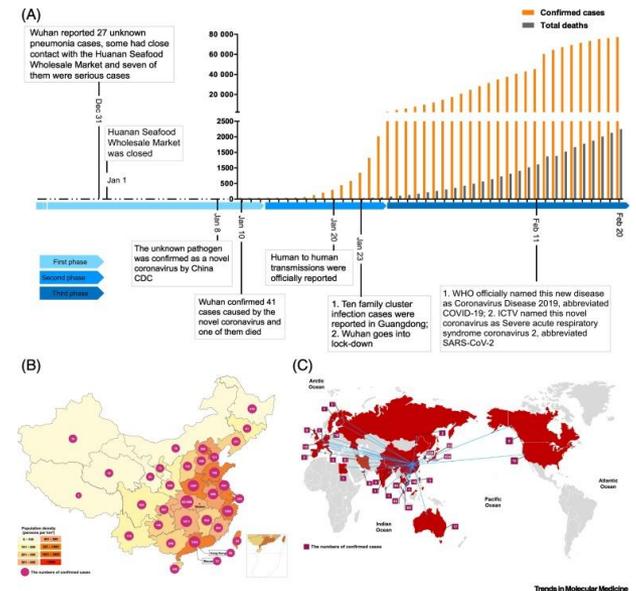
- There are 68 'Pockets' on the spikes where a potential vaccine/drug could 'bind' to stop the virus from connecting
- 4 billion drug compounds that could potentially 'bind' to those pockets
- Gives **68 x 4 billion** possibilities!
- BNL and partners are using computers to test out all possible options and assess which ones are most likely to succeed.
- The most likely drug compounds can then be tested through experiments and drug trials.



Work funded by DOE's National Virtual Biotechnology Laboratory

Epidemiology: Predicting the spread of infection

- Goal is to predict:
 - How many will get infected? How many will need hospital treatment? What measures will be effective?
- These questions need to be answered for a given region:
 - What protective measures are in place and when? Are people adhering to restrictions? How many people travel through the region? What is the general health of the population?
- Running these complex models is very time consuming and each model has to be run many, many times to account for different scenarios and changes.
- Using Artificial Intelligence, BNL and its partners are creating faster models that would allow us to study more scenarios quicker - this will help us to gain more confidence in the results

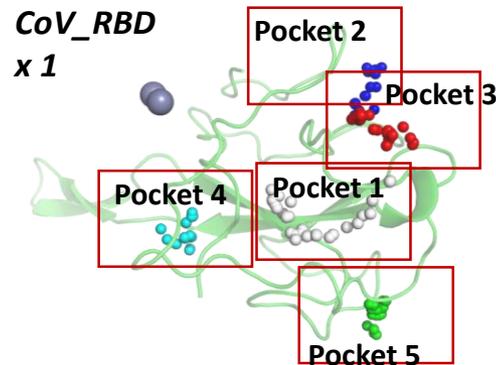
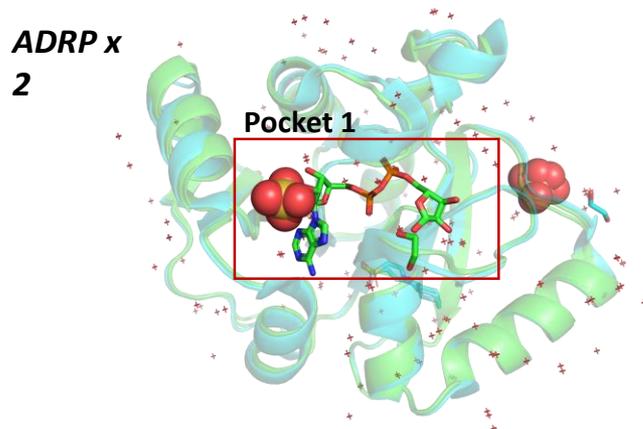
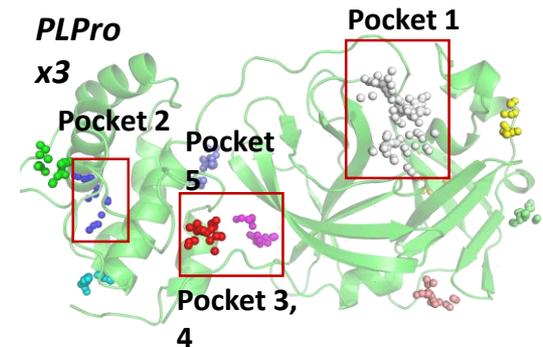
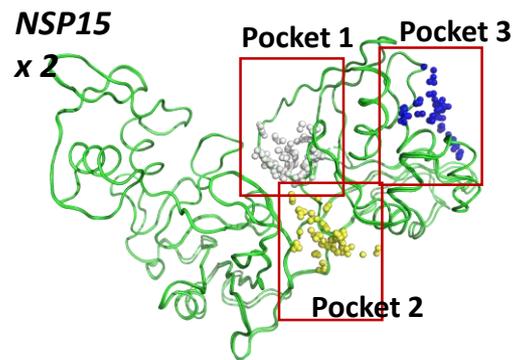
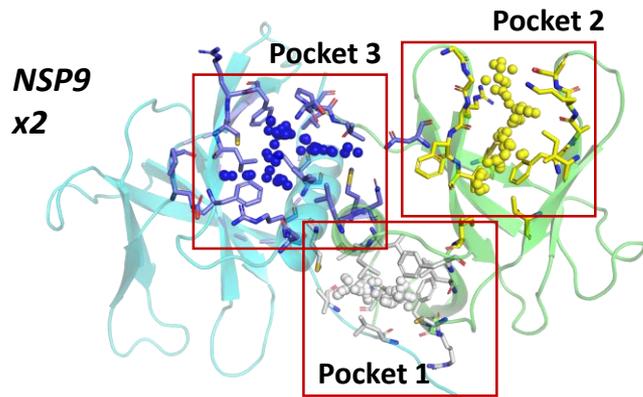


Summary: Some good news

- We understand a lot about Coronaviruses
- A solution will be found. Likely first anti-virals, then vaccines
- We have many good avenues to explore, verify and move forward
- The DOE National labs are working together
- International cooperation is happening
- BNL is part of the fight
 - Drug discovery
 - Computer modeling
 - Virus transport
 - Materials manufacturing

BACKUP

Targets and binding sites



Searching the Literature for the Researchers

Problems
Spurious text
Conjoined text
Out-of-order text

Results of standard PDF-to-text tool

Text	Bbox
bioRxiv preprint first posted online Feb. 5, 2019; doi: https://doi.org/10.1101/537340 . The copyright holder for this preprint (which was not peer-reviewed) is the author/funder, who has granted bioRxiv a license to display the preprint in perpetuity. All rights reserved. No reuse allowed without permission.	5152
Trisomic strains show a commensal phenotype in an oropharyngeal infection model. During oropharyngeal infection in mice, a specific trisomy, Chr6x3, was significantly enriched among strains and recovered from the majority of immunocompromised mice (Forche et al. 2018). The frequency of Chr6x3 increased over the course of infection (Fig. 1A) with the allele combination ABB57 occurring 2-fold more frequently than the AAB combination (Fig. 1B), suggesting that clones with trisomy3 of Chr6 have a general fitness advantage during OPC and that an extra copy of allele B may be more beneficial than an extra copy of allele A in this host niche. To test this hypothesis, we selected several strains that, based on whole genome karyotypes (produced using double digest restriction-site associated DNA sequencing (ddRADseq)), had acquired single trisomies64 as the only change compared to the diploid65 progenitor, strain YB9318. Strains AF127566 and AF1465 both had acquired Chr6x3, the former66 (Chr6ABB) and the latter with allele69 combination AAB (Chr6AAB). Each strain70 was71 oropharynx72. Importantly, these strains had not been73 subjected to any selection regimes (e.g., 74 GAL1) counterselection-induced). A third75 strain, AF1273, that had acquired Chr6x376 (Chr6SAB) and a small LOH on Chr1 (due77 to selection for GAL1 LOH), served as the allele originally of combination recovered the same as from Fig. 1. Chr6 trisomy ABB is overrepresented in isolates recovered from mice with OPC. (A) The frequency of Chr6 trisomy increases over the course of infection. (B) Among Chr6 trisomic strains, genotype ABB is the most frequent allele combination. For each genotype, symbol size is proportional to the frequency of isolation. Results are from the analysis of C. albicans colonies from 3–5 mice per time point as described in (Forche et al. 2018). ABB is the mouse.5	533

- Over **70,000** research papers have been published on aspects of Covid-19
- Having the latest information point to new avenues and avoid wasting time on dead ends.
- However no one can read so many papers - several 1000s a week
- BNL is using Artificial Intelligence to analyse the papers and answer simple questions - what has been written about X, has someone reported symptom Y, has anyone looked at drug compound x?
- We can find relevant text, images and tables in the publications quickly and show them to the researchers.

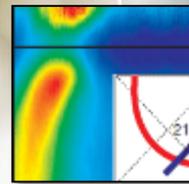
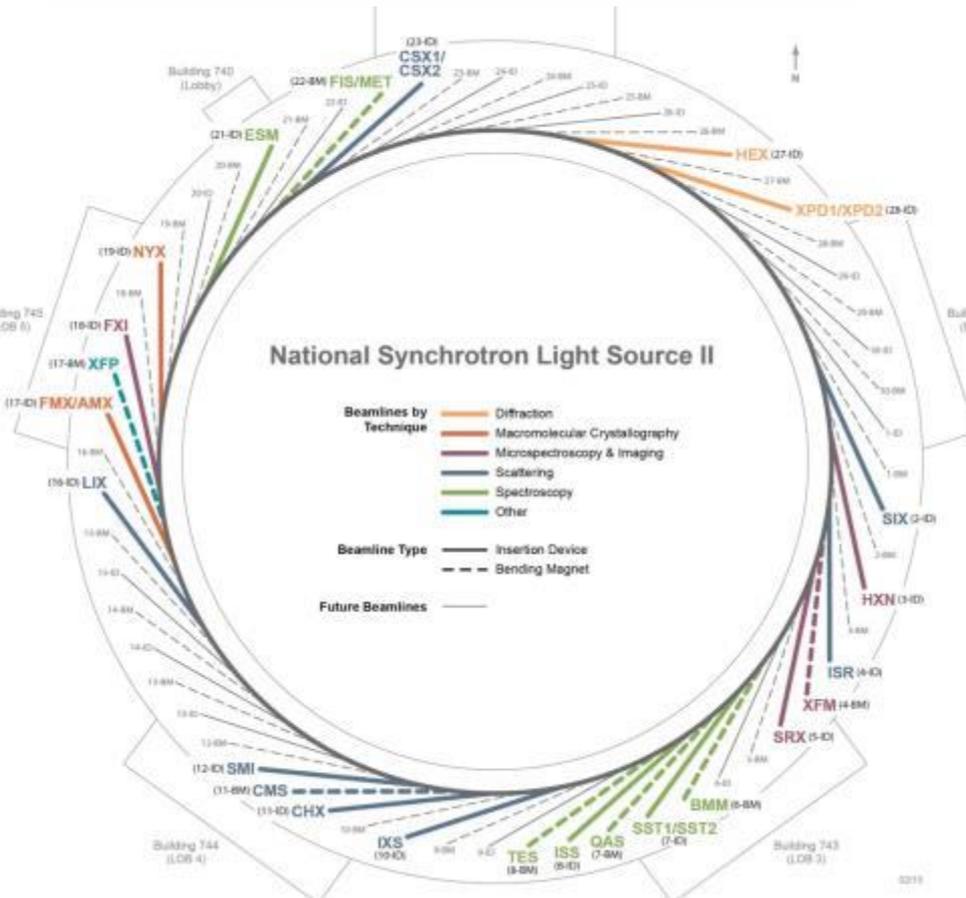
Summary

- NSLS-II provides state-of-the-art tools and the scientist experts to use them
- Having impact in wide ranging scientific areas
- User facility – free to use with intent to publish research
- A number of partnering mechanisms available to develop additional capabilities

NSLS-II Beamlines

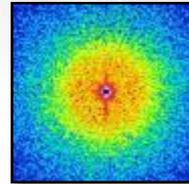
- 28 Operating/Commissioning
- 1 Under Development

<http://www.bnl.gov/ps/nsls2/beamlines/map.php>



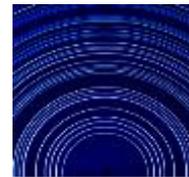
Soft X-Ray Scattering & Spectroscopy

Electronic and magnetic structures and excitations



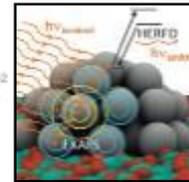
Complex Scattering

Soft materials, structures and dynamics



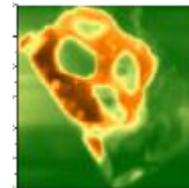
Diffraction & In Situ Scattering

Hard materials, *in operando* Structure



Hard X-Ray Spectroscopy

Chemical reactions *in operando*



Imaging & Microscopy

Chemical, structural and morphological imaging down to 5nm

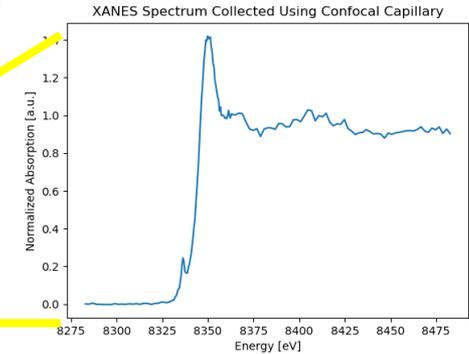
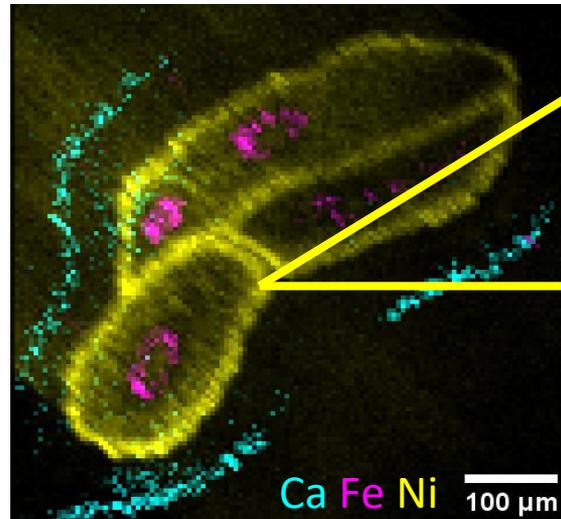
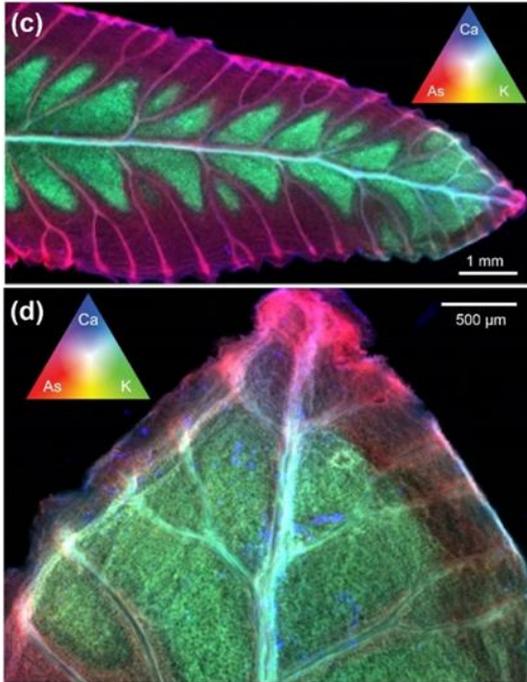


Structural Biology

Protein structures to 1 Å resolution from ~1 micron crystals

Spectroscopic imaging from nm to mm

Arsenic uptake in leaves



Kopittke *et al.*, 2018. *Plant Physiology* (DOI: 10.1104/pp.18.00759).

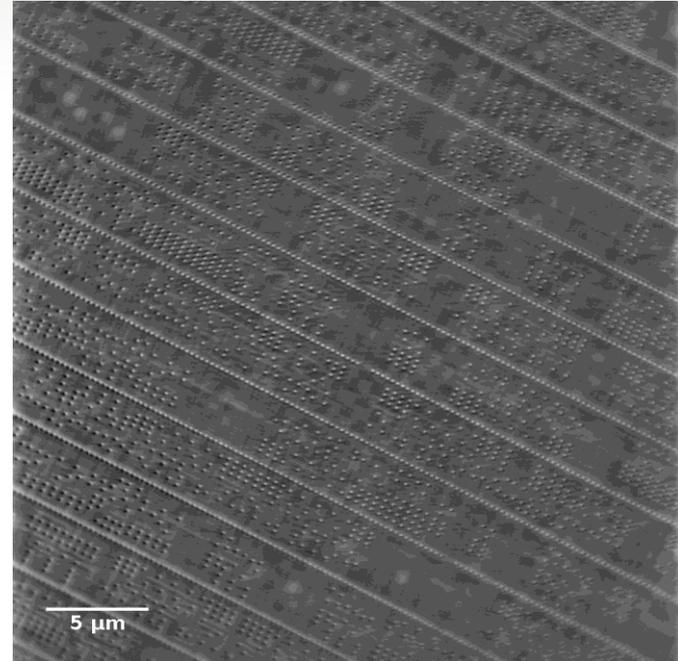
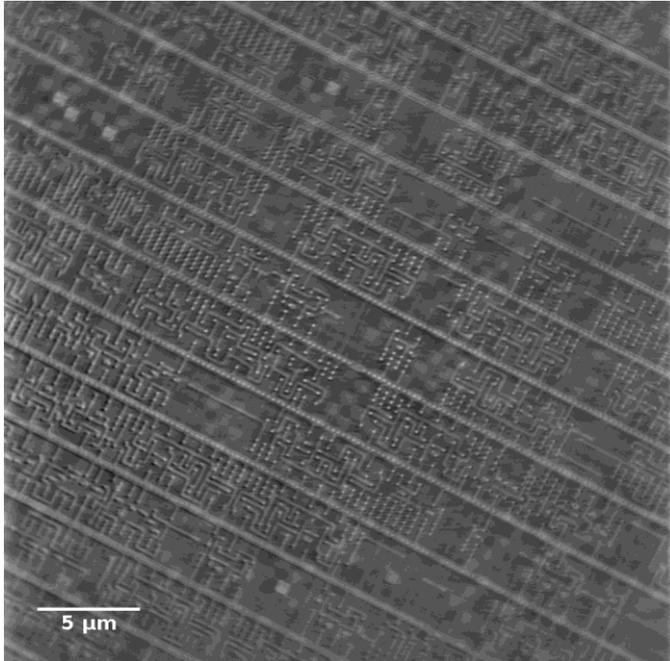
Not just elemental sensitivity, but spectroscopic (chemical) information too with unprecedented resolution and sensitivity

Microelectronics

NSLS-II and Microelectronics

- Legacy electronics – mapping chip architecture of existing chip sets
- State-of-the-art electronics – understanding the factors limiting the performance of current technologies
- ‘Beyond-Moore’s law’ - carrying out the basic science behind the next generation information processing and storage, including technologies such as neuromorphics, spintronics, magnonics, ...

Imaging of microelectronics



Two different layers in the integrated circuit

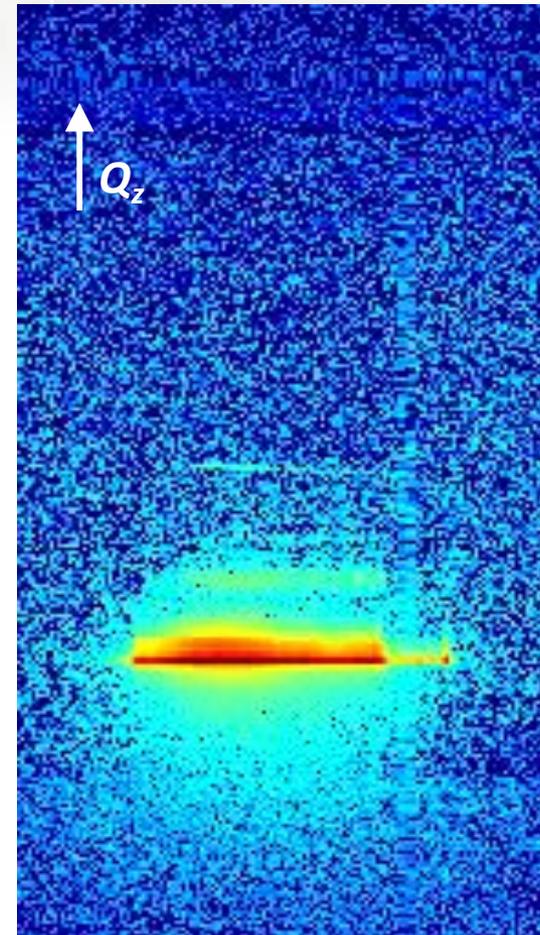
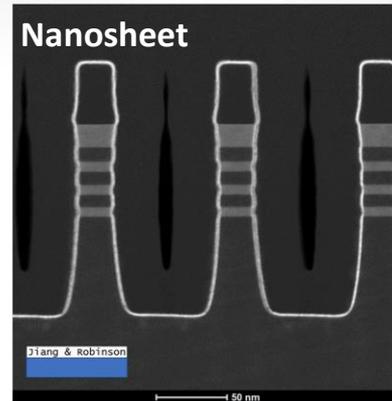
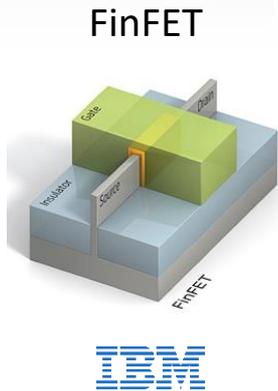
Smallest features are 90 nm

Complete 3D tomography data set taken in 15 seconds* (~300 projections)

~ 500-1000X faster compared with lab-based TXM

*limited by camera frame speed, not photon flux. With a faster camera, total time will be < 10 s

Scanning imaging: Nano Diffraction from Nanosheet



*C. Lavoie, C. Murray, J. Jordan-Sweet (IBM);
H. Yan, X. Huang, Y. Chu (NSLS-II)*

- Strain determines performance of the next-generation microprocessors *e.g.* in IBM's new nanosheet technology
- Studied 7 nm thick and 100 nm wide nanosheet
- Can do such measurements while current is flowing - operando

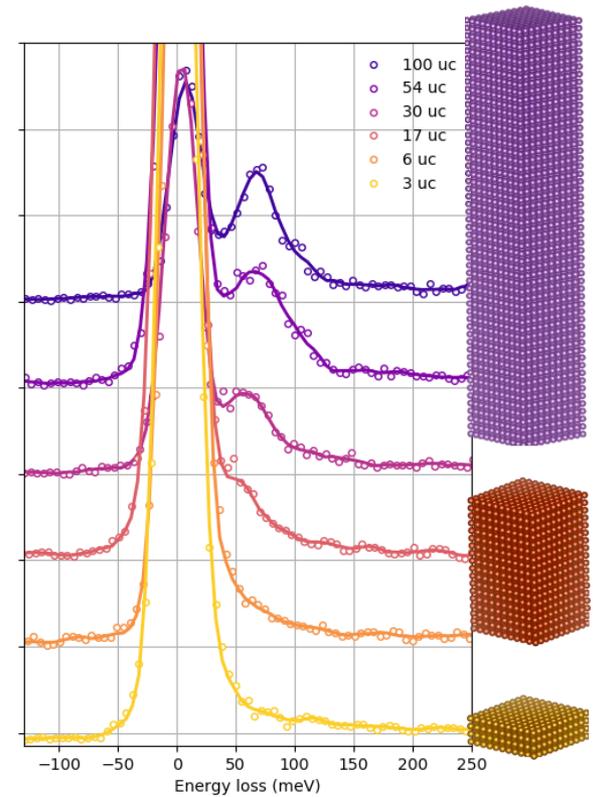
Nano-diffraction at HXN (5 sec/frame)

IBM Research. Samples fabricated at SUNY-Poly

Beyond Moore's law: Using spin-waves for information processing

Spin waves are potentially much more energy efficient

First results: Spin waves in ultra-thin films of Iron

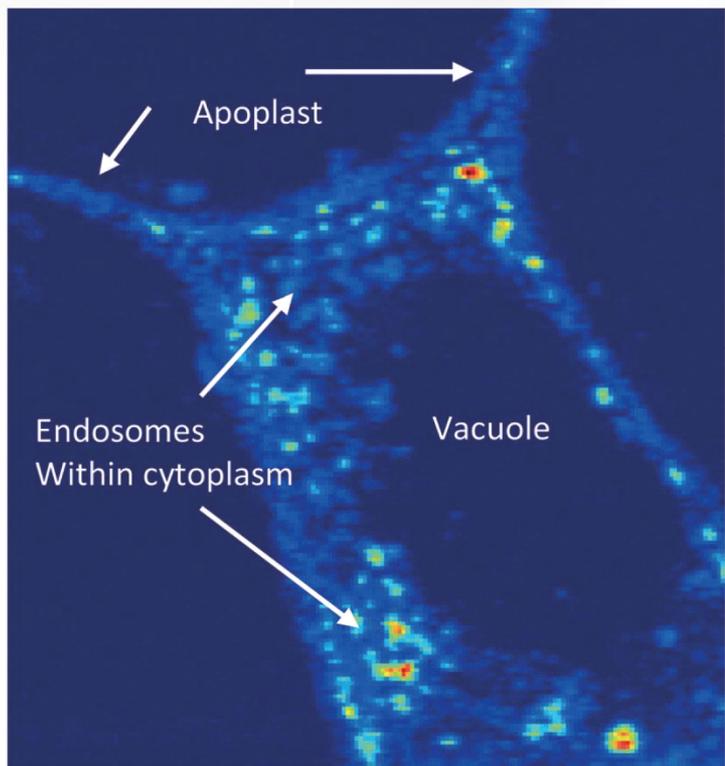


SIX is only beamline in the world with the resolution to carry out these measurements

Valentina Bisogni

Environmental Science

A Nanomaterial's Journey through a Tomato Plant



Scientific Achievement

Revealed how a manufactured nanomaterial (MNM) based on Ce travels through a tomato plant on a subcellular level.

Significance and Impact

This study will enhance our ability to predict how properties of MNM such as CeO_2 – used in rechargeable batteries – influence the uptake, transformation, and transfer of nanomaterials in terrestrial food webs.

J. Li, R. V. Tappero, A. S. Acerbo, H. Yan, Y. Chu, G. V. Lowry, J. M. Unrine. *Environ. Sci.: Nano* 6, 273 (2019).

