TEM data is not FAIR, how to make it play well with others?

Andy Stewart University of Limerick



andy.stewart@ul.ie

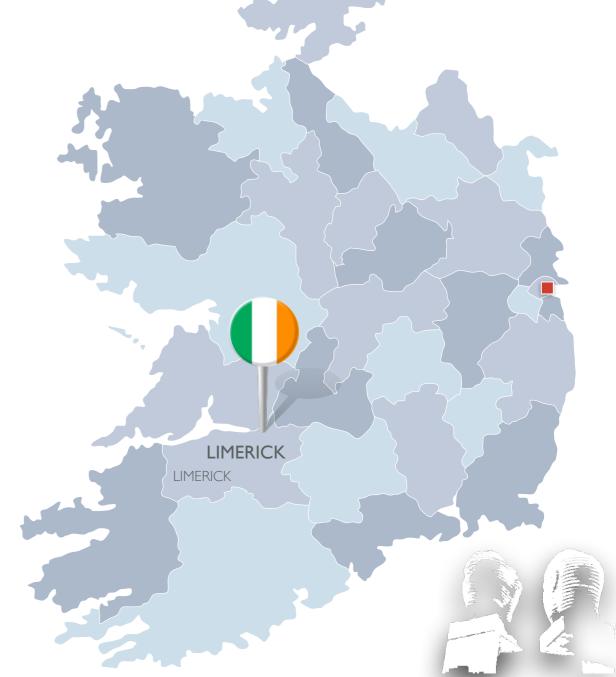


Where is Limerick?





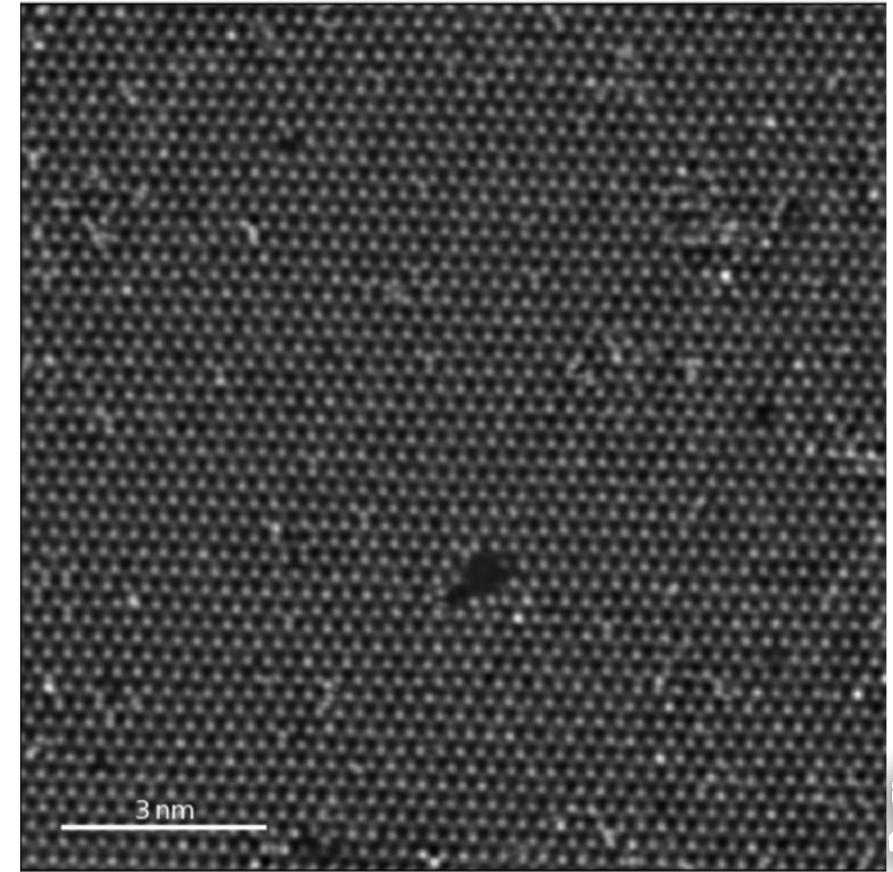




TEM image

Courtesy of Eoghan O'Connell, University of Limerick





Data Processing Metadata

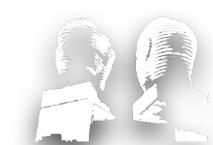




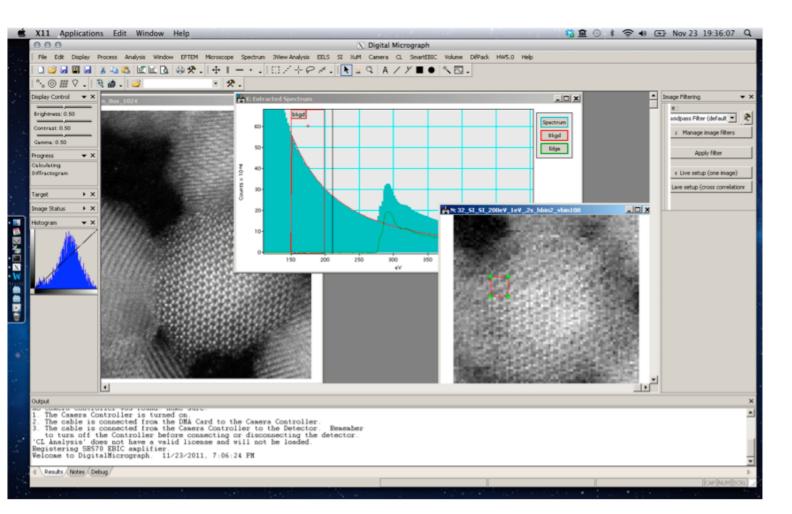
Data Processing Metadata







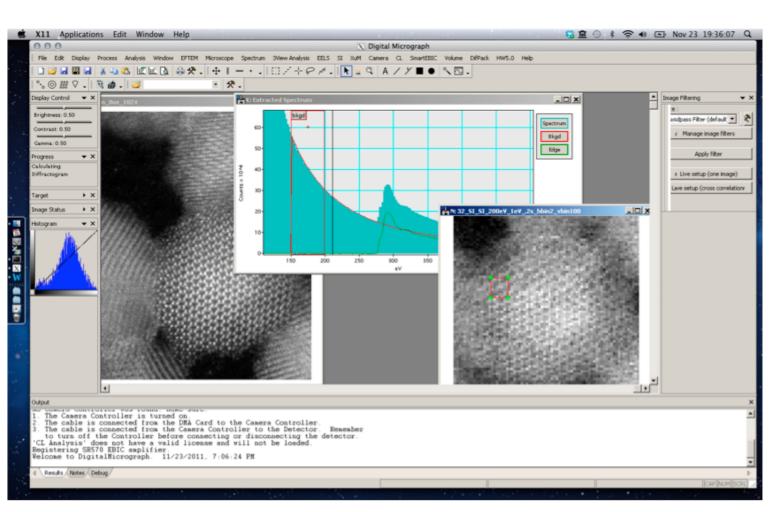
Data formats Currently dm3/dm4







Data formats Currently dm3/dm4





23 versions of the TIFF



TEM Metadata

Acquisition

🛓 Device

- --- Active Size (pixels): (4096,4096)
- ---- Camera Number: 2
- 🚊 CCD
 - Pixel Size (um): 15.0+i15.0
- 🛓 Configuration
 - ---- Location: high_resolution
 - 🛓 Transpose
- --- Name: One View Camera
- --- Source: One View Camera
- --- Source ID: OneView-46565850035
- Source Model: OneView
- Temperature (C): -5.03906

🛓 - Frame

- 🚊 Area
- CCD
- intensity
- 🚋 Reference Images
- 🛓 Parameters
 - Acquisition Write Flags: 4294967295
 - 👜 Base Detector
 - 🛓 Detector
 - 🛓 Environment

🚋 - Environment

- 🚊 High Level
- 🚊 Objects
 - Parameter Set Name: Record
 - ... Parameter Set Tag Path: Imaging:Acquire:Record
- Calibration

∎- DataBar

- --- Acquisition Date: 12/21/2018
- --- Acquisition Time: 11:31:32 AM
- Acquisition Time (OS): 1.31899e+17
- --- Binning: 2
- Collection Number: 1
- Device Name: One View Camera
- --- Exposure Number: 2175
- --- Exposure Time (s): 4.0
- Signal Name: One View Camera
- GMS Version
 - ---- Created: 3.31.2169.1
 - Saved: 3.31.2169.1
- Microscope Info
 - Actual Magnification: 1069.74
 - ---- Cs(mm): 0.0
 - Emission Current (µA): 0.0

Emission Current (µA): 0.0 Formatted Actual Mag: 1kx Formatted Indicated Mag: 580x

- ---- Formatted Voltage: 300kV
- Illumination Mode: TEM
- Illumination Sub-mode: 1
- ... Imaging Mode: DIFFRACTION
- --- Indicated Magnification: 580.0
- Magnification Interpolated: false
- Name: FEI Tecnai Remote
- --- Operation Mode: DIFFRACTION
- Operation Mode Type: 2
- Probe Current (nA): 0.0
- --- Probe Size (nm): 0.0
- STEM Camera Length: 145.0
- Stage Position
 - Stage Alpha: 0.00878402
 - --- Stage X: -90.0588
 - --- Stage Y: -40.6173
 - Stage Z: -35.5413
- Voltage: 300000.0
- 🛓 Session Info
 - 🚊 Items
 - --- Microscope:







Sample

Preparation methods and details Specimen composition Date and time of production Colour Shape Thickness Defects Grain boundaries





Sample

Preparation methods and details Specimen composition Date and time of production Colour Shape Thickness Defects Grain boundaries

Sample Storage

Environment Relative humidity Temperature Light/Dark

Any variable which may cause sample to change over time





Sample

Preparation methods and details Specimen composition Date and time of production Colour Shape Thickness Defects Grain boundaries

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Environment Relative humidity Temperature Light/Dark

Any variable which may cause sample to change over time

Relationship between images and spectroscopy data





Sample

Preparation methods and details Specimen composition Date and time of production Colour Shape Thickness Defects Grain boundaries

Sample Storage

Environment Relative humidity Temperature Light/Dark

Any variable which may cause sample to change over time

Relationship between images and spectroscopy data

Record any value/detail which will be necessary for evaluating with any other dataset or technique

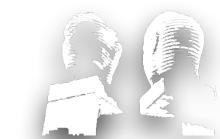


Bernal 👯

Institute

TEM Missing Metadata





TEM Missing Metadata

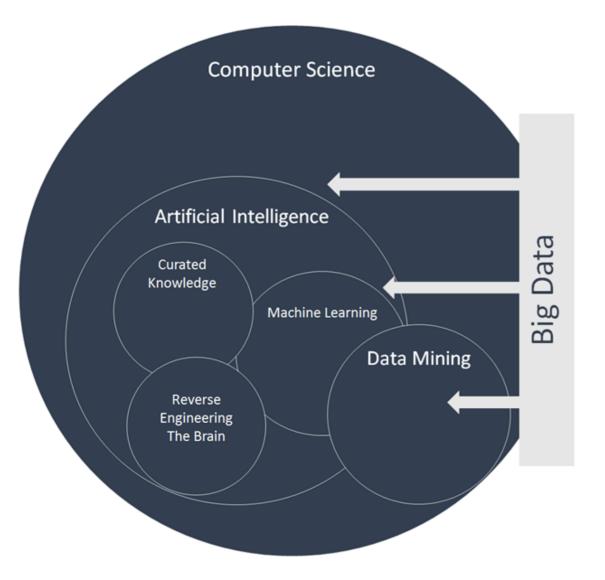
TEM variables

TEM Environment Relative Humidity of room Temperature of room Fields in the room Water temperature Magnet temperature Power Supply variability Beam brightness Detector temperature





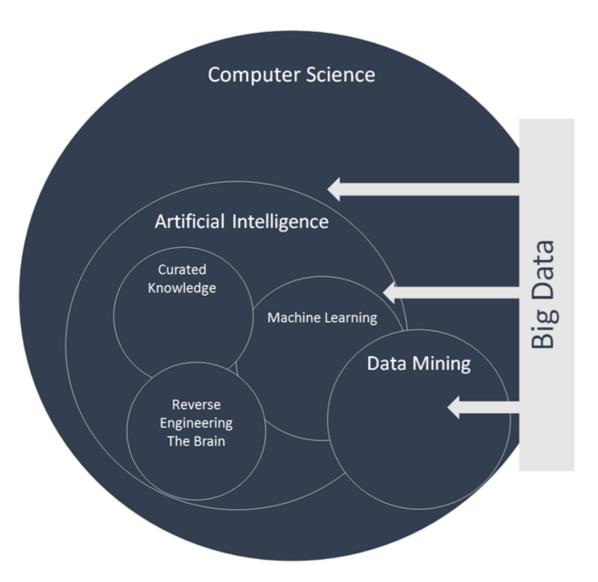
Why should we care?







Why should we care?

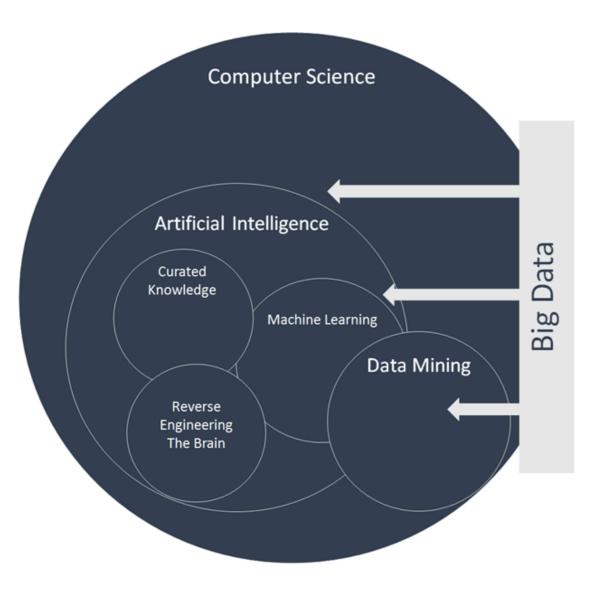


Grow the need for microscopy





Why should we care?



Grow the need for microscopy





It's NOT about you





It's NOT about you













Wellcome Open Research







Engineering and Physical Sciences Research Council



Open Science





It's NOT about you





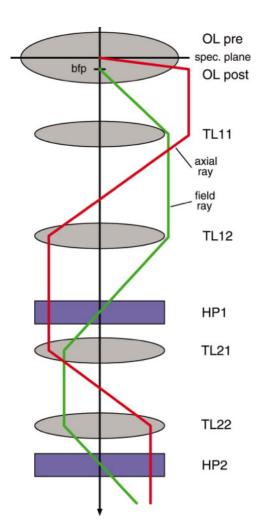








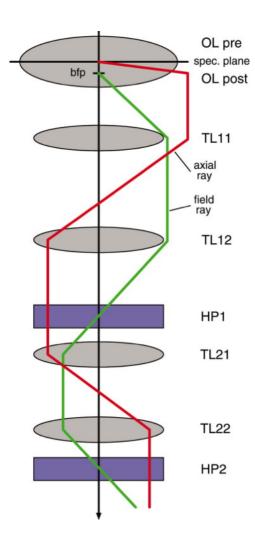
No Corrector aberrations measurements Stored







No Corrector aberrations measurements Stored



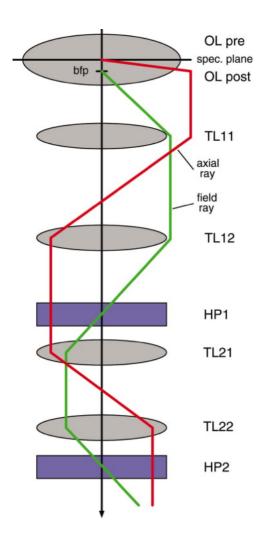


Time between aberration measurement and data acquisition





No Corrector aberrations measurements Stored



Time between aberration measurement and data acquisition



Sample thickness estimate/measurement







We are not the first and will not be the last





We are not the first and will not be the last







CCP4

MTZ format (1970's) Data Harvesting (1990's)

Also Nomad



4		Harvesting Manager	
			Help
Job title			2
	O	en File Browser to select Harvesting Files	
List of harvest files selec	ted		
/nfs/dlpx1/home/proje /nfs/dlpx1/home/proje /nfs/dlpx1/home/proje /nfs/dlpx1/home/proje	cts/cbm2 cts/cbm2 cts/cbm2 cts/cbm2	7/cbm27/Deposit/DepositFiles/cbm27/Peak.refmac 7/cbm27/Deposit/DepositFiles/cbm27/natse_inflection.mlp 7/cbm27/Deposit/DepositFiles/cbm27/natse_inflection.sca 7/cbm27/Deposit/DepositFiles/cbm27/natse_peak.scala 7/cbm27/Deposit/DepositFiles/cbm27/natse_remote.scala 7/cbm27/Deposit/DepositFiles/cbm27/natse_remote.truncat	ala mcate
Un-select all highlighte	d files	Remove selected files from box	
Remove all files from box		View selected file	
Programs			
Run Program to Cross Validate Files	V	alidate Harv. Files 🛁	
Output			
Run	-	Save or Restore 🥌	lose

Data compression

MRCZ

Why GitHub? \sim Enterpri

XFEL

On the file data compression

em-MRCZ / python-mrcz



O Star 10,133

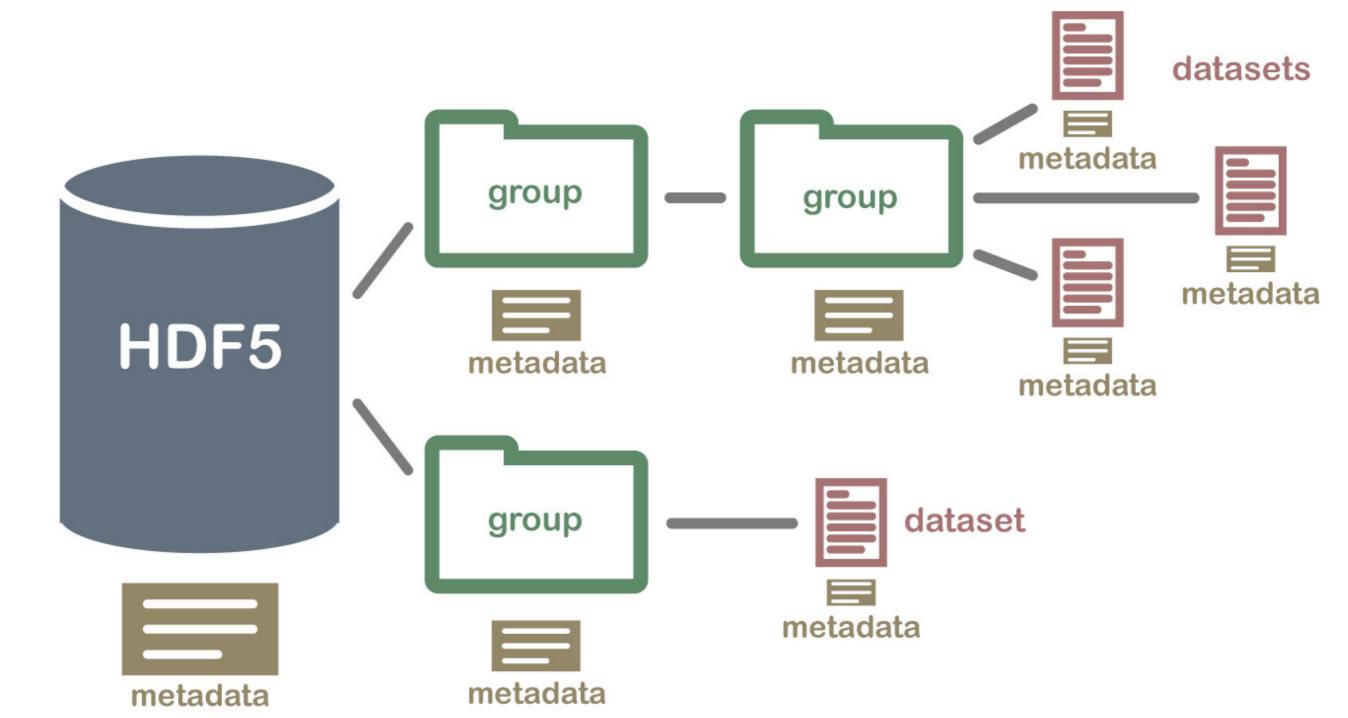
Zstandard is a real-time compression algorithm, providing high compression ratios. It offers a very wide range of compression / speed trade-off, while being backed by a very fast decoder (see benchmarks below). It also offers a special mode for small data, called dictionary compression, and can create dictionaries from any sample set. Zstandard library is provided as open source software using a BSD license.





Recognised need for change Hierarchical Data Format #5

Already >25 variants in the EM community



Data format proliferation







Monty Python, Life of Brian



Data format proliferation







Monty Python, Life of Brian



Adoption

Last thing we need is just another file format

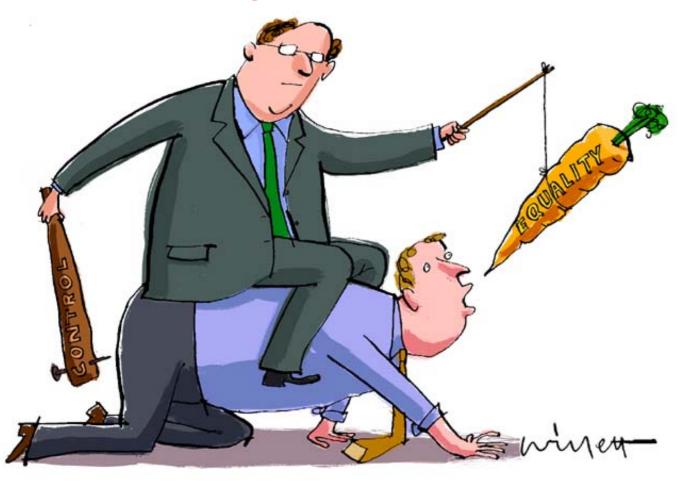






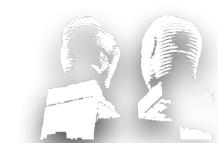
Adoption

Last thing we need is just another file format









Adoption

Last thing we need is just another file format





Public databases Organisations promote use of format





Adoption

Last thing we need is just another file format





 Public databases
 Organisations promote use of format

Journals insist on data submission in a specific format
Funding bodies insist on open data with all meta data





Community Agreement on a single format

Or a series of convertors like Nomad?

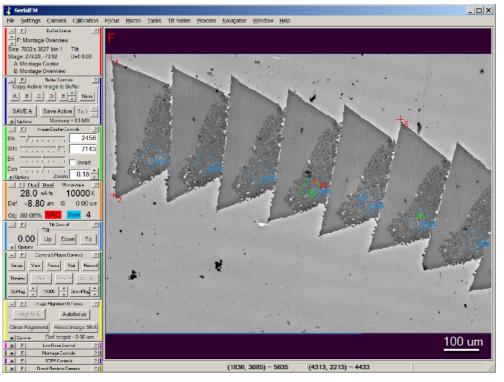
Missing metadata a problem....?





Making Metadata Easier

SerialEM



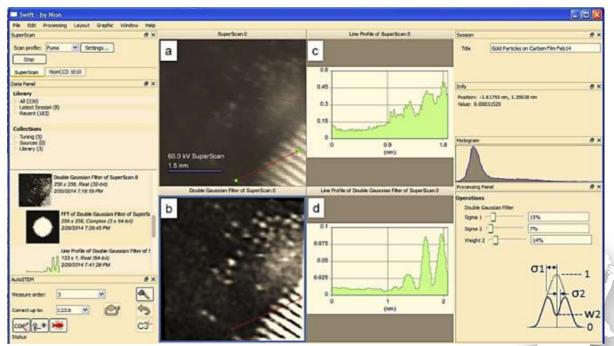




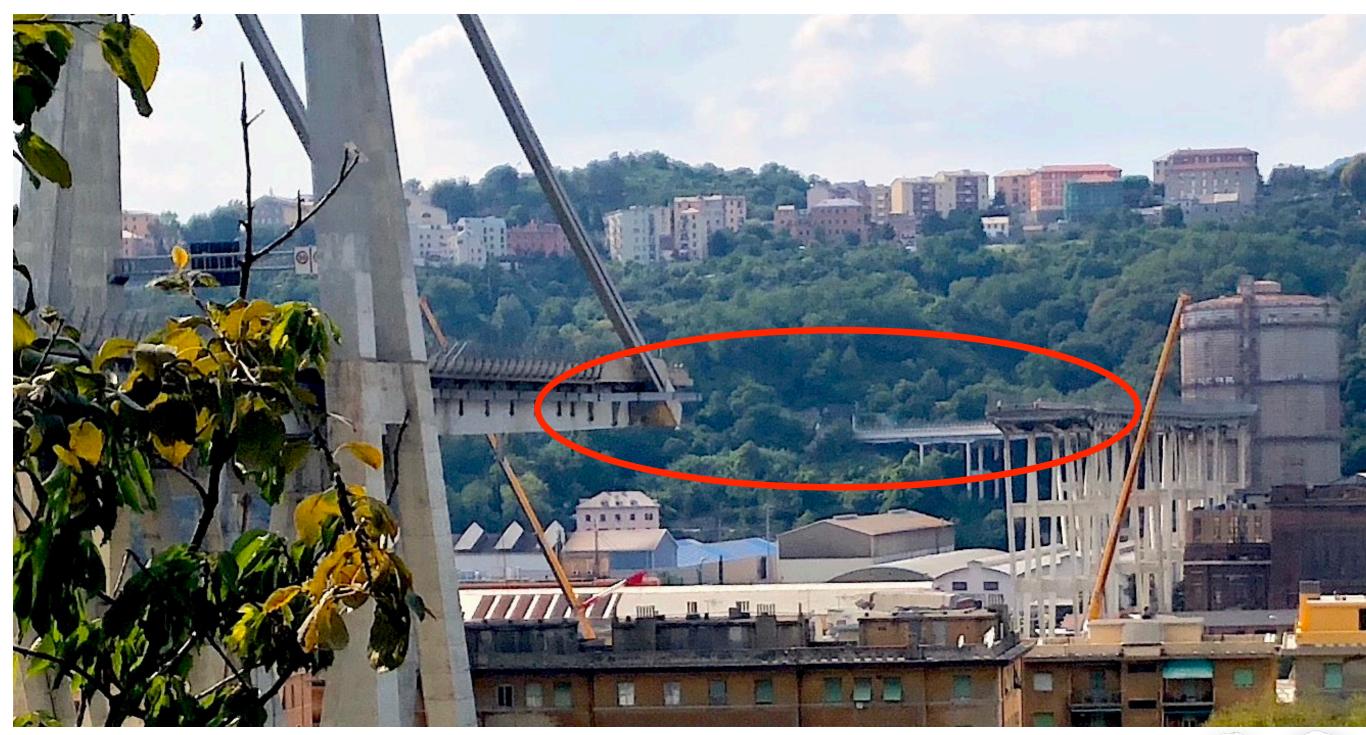
Leginon



Swift Nion



Morandi Bridge Collapse

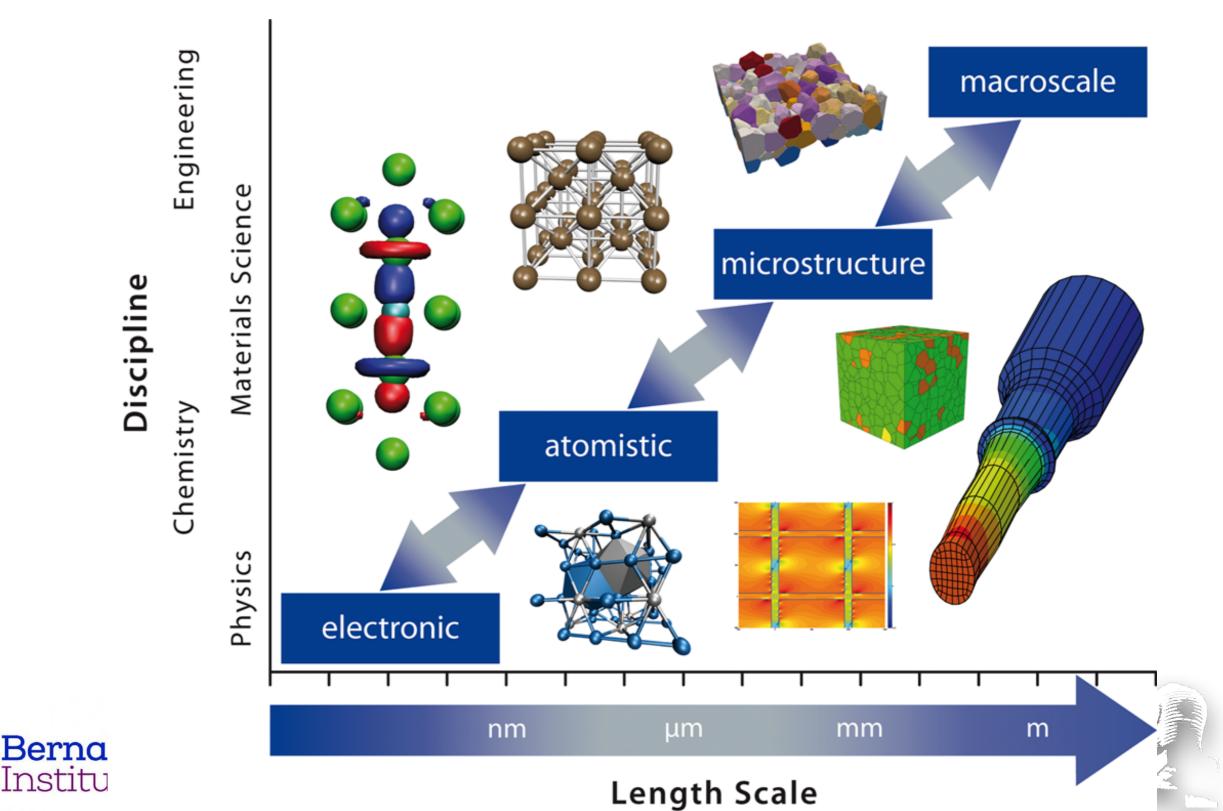








Integration across the length scales

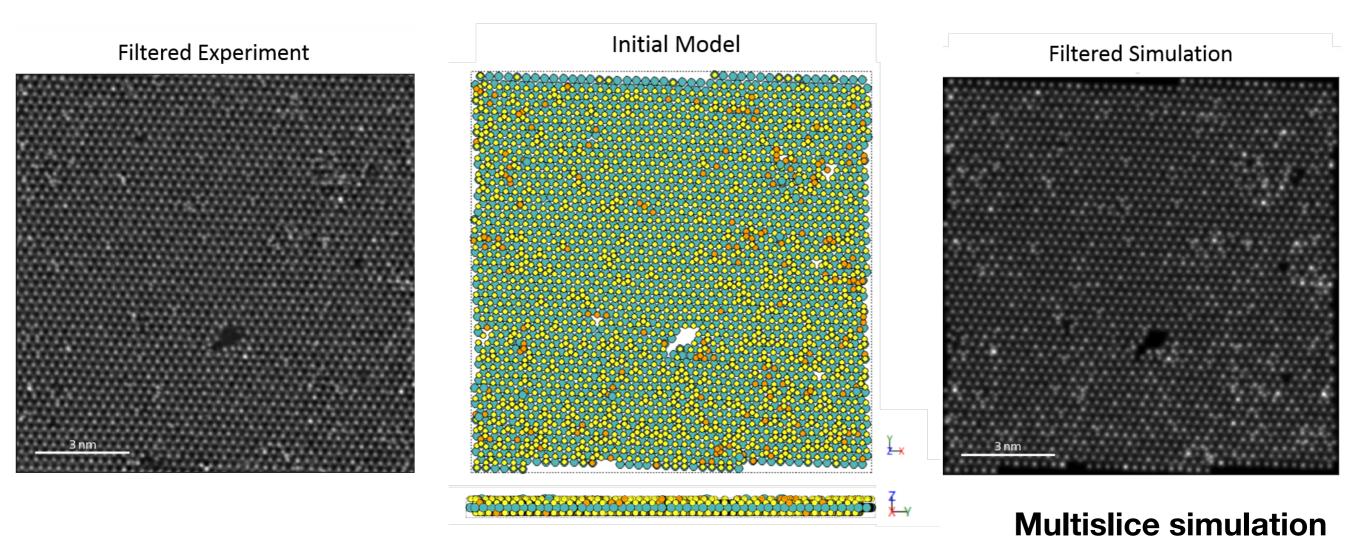


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Mixing fields



For Low Dimensional materials we can easily generate models from TEM data Next link with simulations



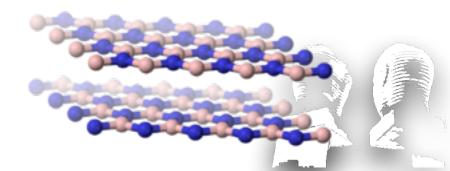


Proposal/Suggestion

Experimentally simulate and test a material across the length scales for a demonstration paper



Suggestion Material? Boron Nitride Magnesium Alloy...



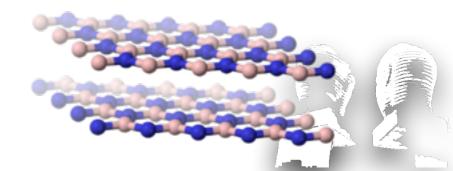
Proposal/Suggestion

Experimentally simulate and test a material across the length scales for a demonstration paper

Additionally use the experience to define and refine the metadata required to make the exchange of data between experiments and simulation as smooth as possible



Suggestion Material? Boron Nitride Magnesium Alloy...



Databases



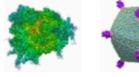
The Electron Microscopy Data Bank (EMDB) at PDBe

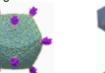
Quick access

opendata

CERN

Click on one of these categories:











Helicase

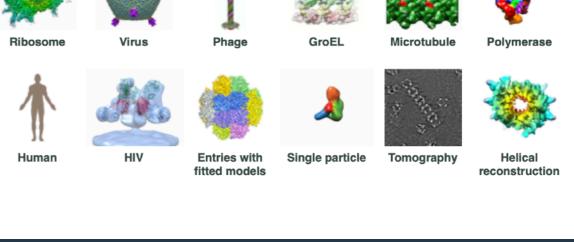
<5Å resolution

About -



153836 Biological Macromolecular Structures Enabling Breakthroughs in **Research and Education**









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EMPIAR

Dataset 🗸	Title 🖨	Authors and references	Size 🛊	Resolution \$	Release date ♦
EMPIAR-10284	Extracellular albumin and endosomal ions prime enterovirus particles for uncoating that can be prevented by fatty acid saturation [multiple data sets in MRC format]	Domanska A, Ruokolainen VP, Pelliccia M, Laajala MA, Marjomäki VS, Butcher SJ [Pubmed: <u>31189702]</u> [DOI: <u>10.1128/JVI.00599-19]</u> EMD-4903, EMD-0565, <u>6rjf</u> , <u>6006</u>	2.4 TB	3.5-3.6	2019-06- 19
EMPIAR-10281	Cryo electron micrographs of human cystic fibrosis transmembrane conductance regulator (CFTR) in complex with GLPG [3335 multi-frame micrographs composed of 50 frames each in TIFF format]	Zhang ZZ, Liu FL, Chen JC [Pubmed: <u>31221859]</u> [DOI: 10.1126/science.aaw7611] EMD-0606, 6o1v	1.4 TB	3.2	2019-06- 21
EMPIAR-10270	Single particle reconstruction of 52 kDa biotin-bound state streptavidin at 3.2 Angstrom resolution [3309 multi-frame micrographs composed of 32 frames each in MRC format]	Fan X, Wang J, Zhang X, Yang Z, Zhang JC, Zhao L, Peng HL, Lei J, Wang HW [Pubmed: <u>31160591]</u> [DOI: <u>10.1038/s41467-019- 10368-w]</u> EMD-0689, 6j6j	5.5 TB	3.2	2019-06- 07



Block Chain database



Caltech Electron Tomography Database

A public repository featuring 11293 electron tomography datasets of intact bacterial and archaeal cells, representing 85 species.

Browse database 🗲

View our terms and conditions

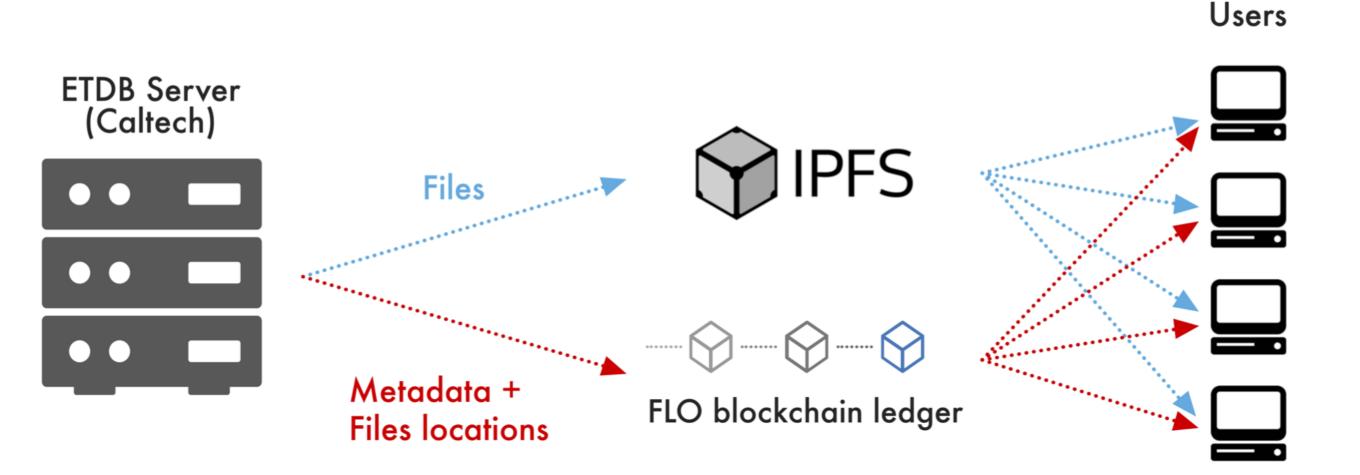
>301 B OT Cryo-Iomography data available, variable quality



ETDB-Caltech: A blockchain-based distributed public database for electron tomography Davi R. Ortega, et al., PLOS ONE <u>https://doi.org/10.1371/journal.pone.0215531</u>



Block Chain database



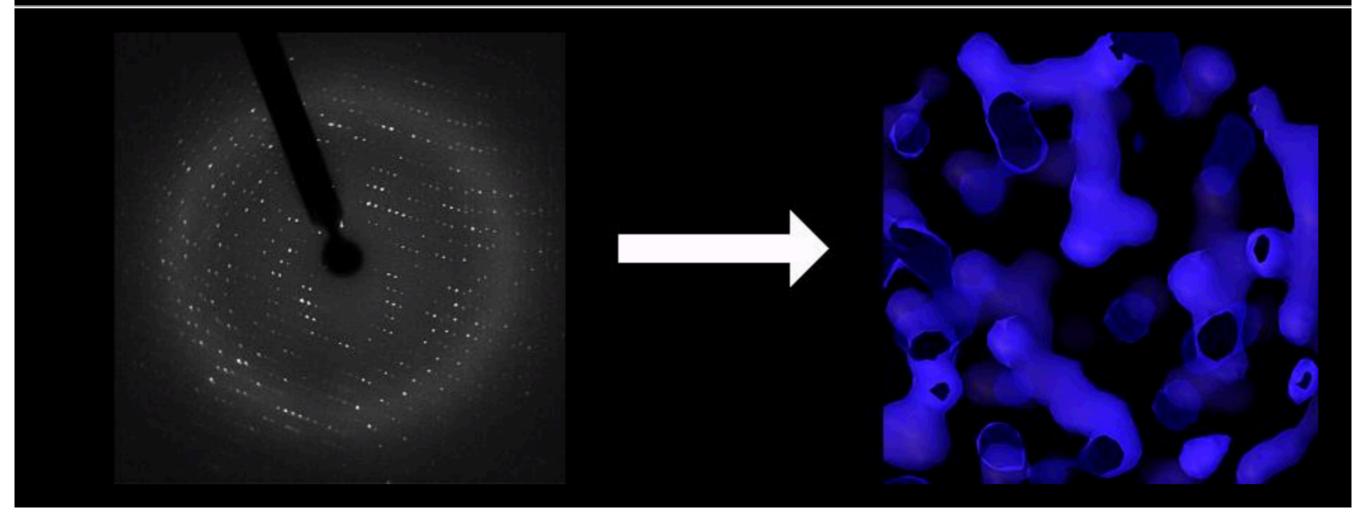
>30TB of Cryo-Tomography data available, variable quality



ETDB-Caltech: A blockchain-based distributed public database for electron tomography Davi R. Ortega, et al., PLOS ONE <u>https://doi.org/10.1371/journal.pone.0215531</u>

ELVES

"Because You Have Better Things To Do" TM





James Holten

Standard Operating Procedures

COMPLIANCE POLICY GUIDE (CPG)

Guidance for Industry, Q7A Good Manufacturing Practice Guidance for Active Pharmaceutical Ingredients



Food and Drug Administration























International Metrology Institutes





European Molecular Biology Laboratory

European Molecular Biology Laboratory



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European Materials Laboratory



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Thank you for listening



