X-RAY PTYCHOGRAPHY RECONSTRUCTION ON DISTRIBUTED GPUS

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OUTLINE

- Introduction
 - X-ray Ptychography
 - Data and Computation Challenges
- ▶ GPU Acceleration
 - Implementation Workflow
 - Results
- Conclusions



INTRODUCTION

MOTIVATION

- X-ray diffraction technique is an essential tool for studying the structure of materials at nanoscale.
- With brighter light sources, faster data acquisition rates, larger data volumes (higher image resolutions), high performance computing (HPC) is critical for enabling in-situ analysis of image data from light sources.
- Efficient interactive graphical user interface (GUI) is also necessary to allow general beam line users to analyze their data in near real time, so that they can adjust their X-ray experiments as needed.



PTYCHOGRAPHY

Ptychography

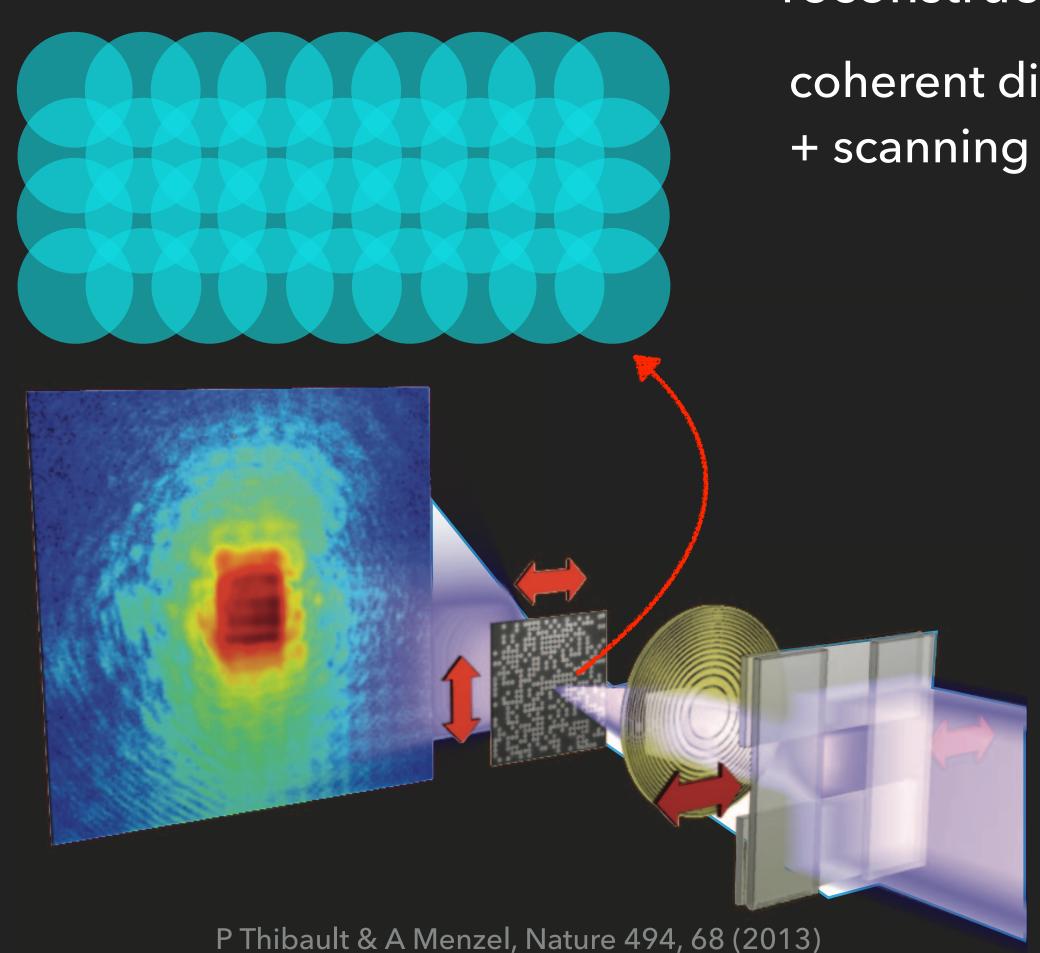
From Wikipedia, the free encyclopedia

Ptychography (/trˈtʃoʊgræfi/ ti-CHOH-graf-ee; πτυχο-γραφία, from πτυχή = fold + γραφή = writing/scripture), also known as scanning diffraction microscopy, is a type of coherent diffraction imaging invented by Walter Hoppe in the $1960s^{[1]}$ that uses a beam of coherent radiation (e.g. x-rays) or particles (e.g. electrons) to illuminate a small sample of material and then the phase information from the beam is recovered by analysing the diffraction pattern produced by the scattered radiation. The resulting microscopic image can show details at the atomic level.

- Ptychography, pronounced as ti-CHOH-graf-ee, from Greek for "fold".
- A coherent diffraction imaging technique invented by Walter Hoppe in 1960s.
- Coherent diffraction images are obtained from overlapping regions of the sample.
- > Sample phase information can be reconstructed from post analysis of the scanning images.



X-RAY PTYCHOGRAPHY



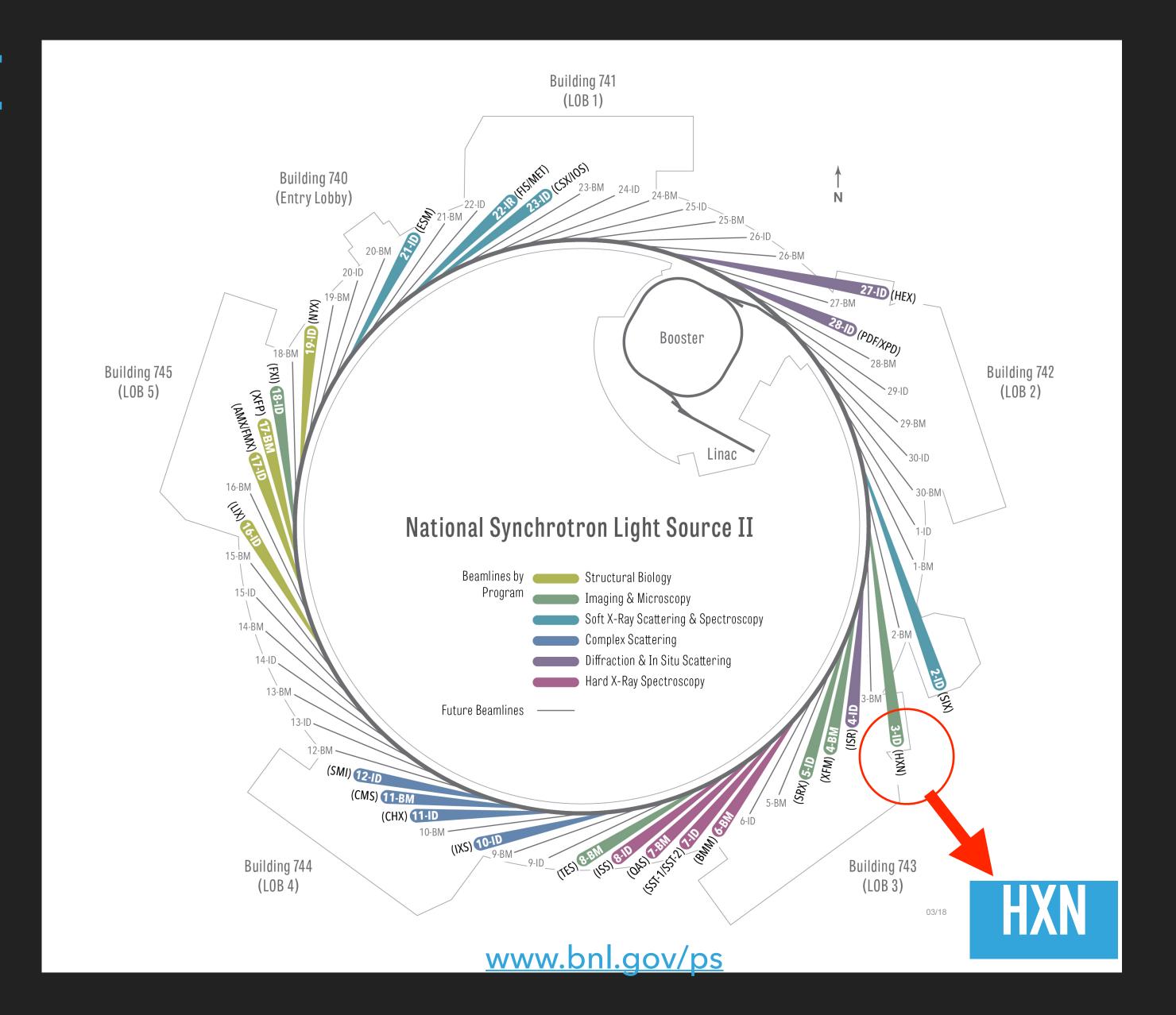
reconstruct high-resolution images from coherent diffraction data coherent diffraction imaging (CDI)

- + scanning transmission X-ray microscopy (STXM)
 - √ no special optics requirement
 - ✓ wavelength-limited resolution
 - ✓ no *a priori* knowledge in illumination
 - √ easy sample preparation
 - diffraction intensity recorded
 - overlapped scanning
 - → sufficient information for object reconstruction



NSLS-II AND THE HXN BEAM LINE

- State-of-the-art, medium-energy (3-billion-electron-volt, or GeV) electron storage ring that produces x-rays up to 10,000 times brighter than the NSLS
- 28 beam lines in operation; 1 under development
- HXN Hard X-ray Nanoprobe: hard x-ray imaging of structure, elements, strain and chemical states with spatial resolution from 10 to 30 nm with an ultimate goal of ~1nm



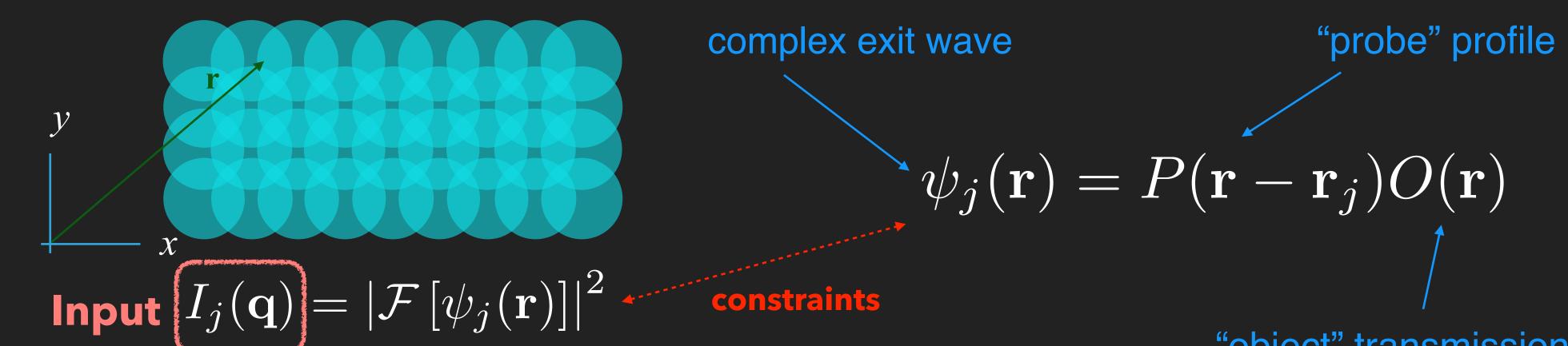


HXN BEAMLINE SETUP

fluorescence detector diffraction detector transmission detector



SINGLE-MODE PTYCHO. RECONSTRUCTION

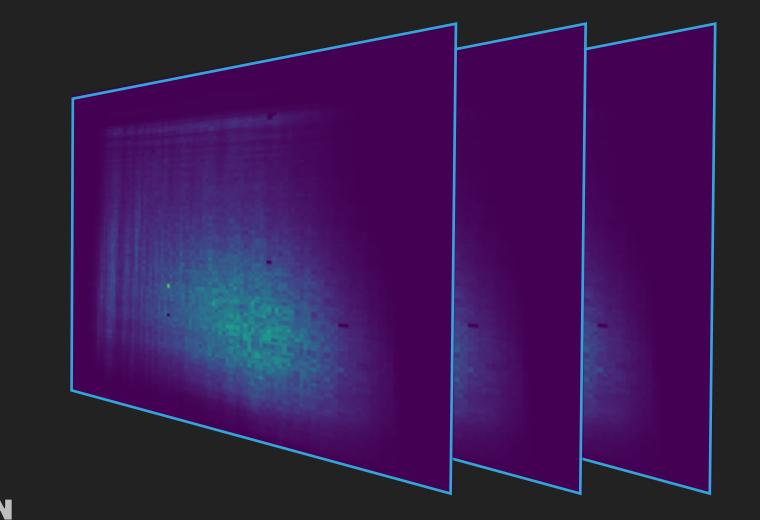




"object" transmission function

difference map algorithm*

$$P(\mathbf{r}) = \frac{\sum_{j} [O(\mathbf{r} + \mathbf{r}_{j})]^{*} \psi_{j}(\mathbf{r} + \mathbf{r}_{j})}{\sum_{j} |O(\mathbf{r} + \mathbf{r}_{j})|^{2}}$$
$$O(\mathbf{r}) = \frac{\sum_{j} [P(\mathbf{r} - \mathbf{r}_{j})]^{*} \psi_{j}(\mathbf{r})}{\sum_{j} |P(\mathbf{r} - \mathbf{r}_{j})|^{2}}$$



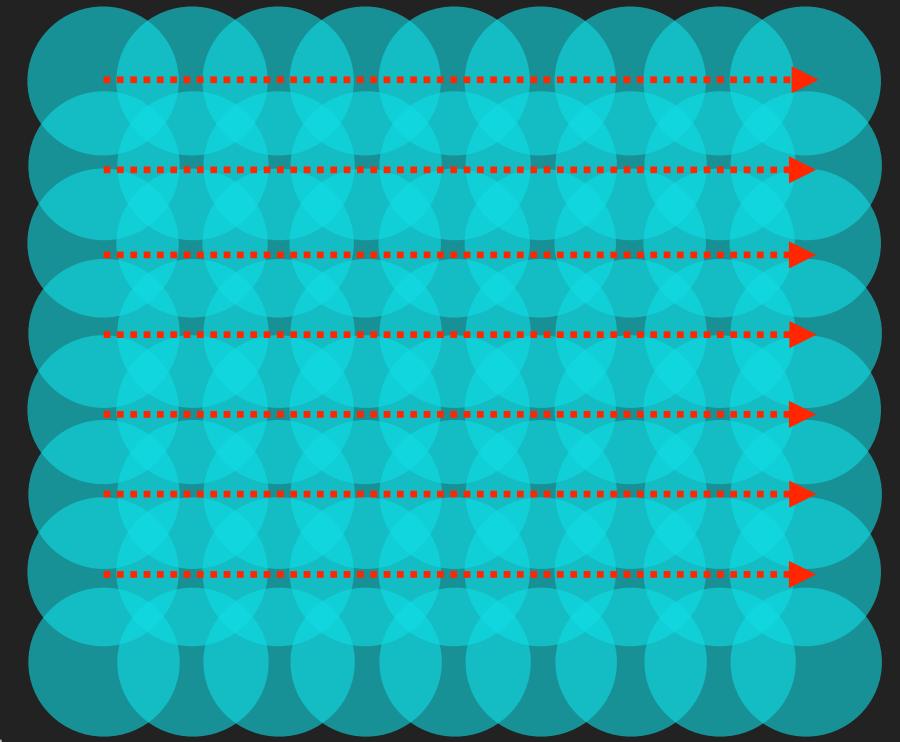
 $j=1,2,\cdots,N$

MULTI-MODE PTYCHO. RECONSTRUCTION

motivation: to reduce motor overhead by performing "fly scan"

blurriness due to motion can be regarded as having incoherent x-ray beam

need to compensate this with multiple probe modes



probe mode index

object mode index

$$\psi_j^{(k,l)}(\mathbf{r}) = P^{(k)}(\mathbf{r} - \mathbf{r}_j)O^{(l)}(\mathbf{r})$$

$$P^{(k)}(\mathbf{r}) = \frac{\sum_{l} \sum_{j} \left[O^{(l)}(\mathbf{r} + \mathbf{r}_{j}) \right]^{*} \psi_{j}^{(k,l)}(\mathbf{r} + \mathbf{r}_{j})}{\sum_{l} \sum_{j} |O^{(l)}(\mathbf{r} + \mathbf{r}_{j})|^{2}}$$

$$O^{(l)}(\mathbf{r}) = \frac{\sum_{k} \sum_{j} \left[P^{(k)}(\mathbf{r} - \mathbf{r}_{j}) \right]^{*} \psi_{j}^{(k,l)}(\mathbf{r})}{\sum_{k} \sum_{j} |P^{(k)}(\mathbf{r} - \mathbf{r}_{j})|^{2}}$$

one more sum to do ⇒ still highly parallelizable!



DATA AND COMPUTATION CHALLENGES

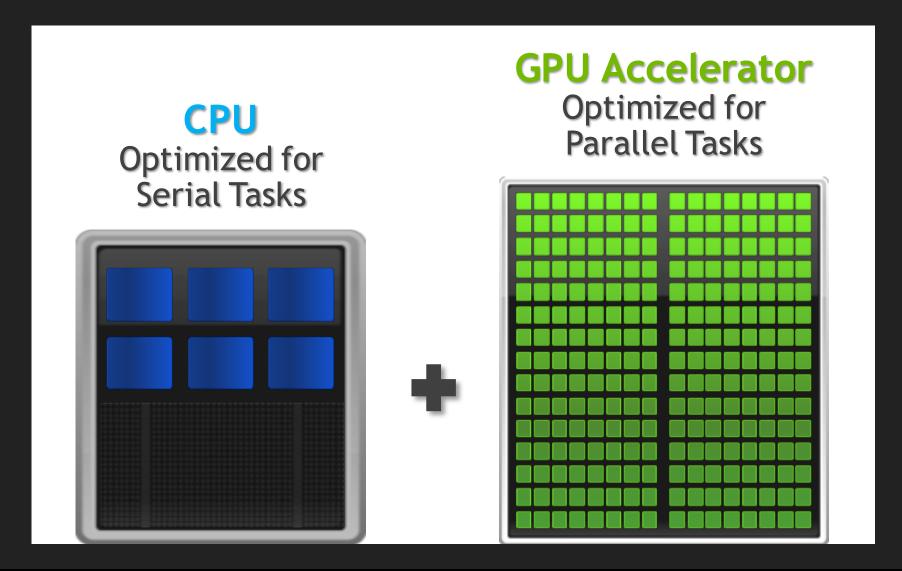
- ▶ For a high-resolution ptychography reconstruction
 - Need O(10,000) scan images
 - ~200x200 pixels in double precision per image
 - ▶ Memory requirement for <u>input</u> images: O(1 GB) to O(10 GB)
- DM algorithm: O(100) iterations
- Memory requirement for the DM algorithm (including temporary buffers):
 - Single-mode: ~4x input
 - Multi-mode: ~10x input
- Serial code: takes hours to complete one ptychography reconstruction.
- ▶ Beamline users have limited access time and possibly limited offline computing resources => Fast *in-situ* ptychography reconstruction software will improve beam line users' productivity.

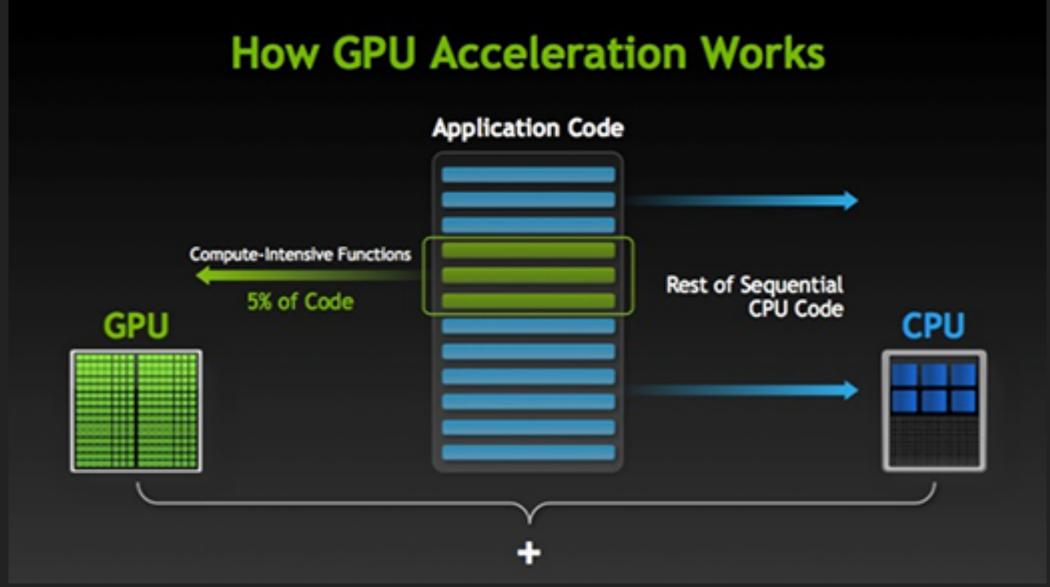


GPU ACCELERATION

GPUS FOR SCIENTIFIC COMPUTING

- Modern GPUs offer massive compute power for parallel tasks
 - Thousands of compute cores vs. O(10) cores on CPUs
- Image processing: straightforward parallelism in pixels.
- ▶ GPU memory still limited: 32 GB in the latest generation of NVIDIA GPUs (Volta V100).
- Need to distribute workload to multiple GPUs







DATA MANAGEMENT FOR GPU COMPUTING

- Host CPUs and the GPU devices have distinct physical memories
- Data transfer between the CPU memory and the device memory could be a bottleneck.
- Want to minimize data transfer while maximizing the computation done on device.
- Difference map algorithm involves mostly local operations. Global sum (Allreduce) needed only at the end of each iteration. Scalable!

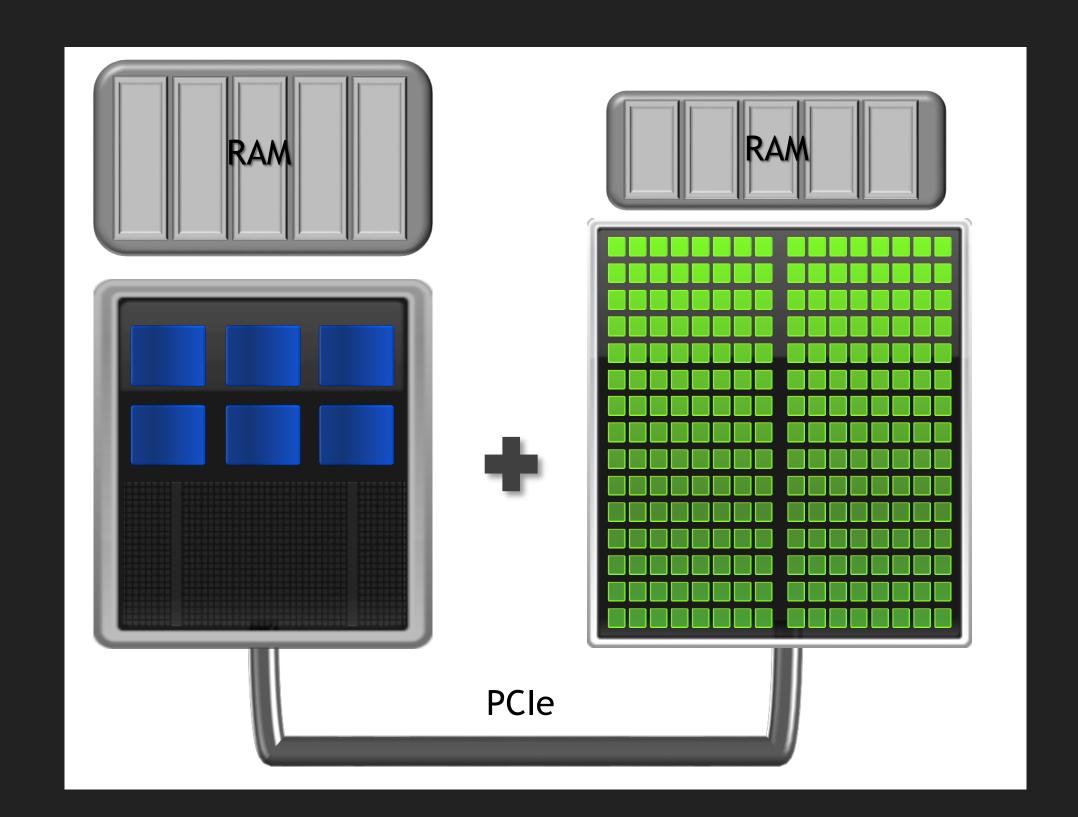
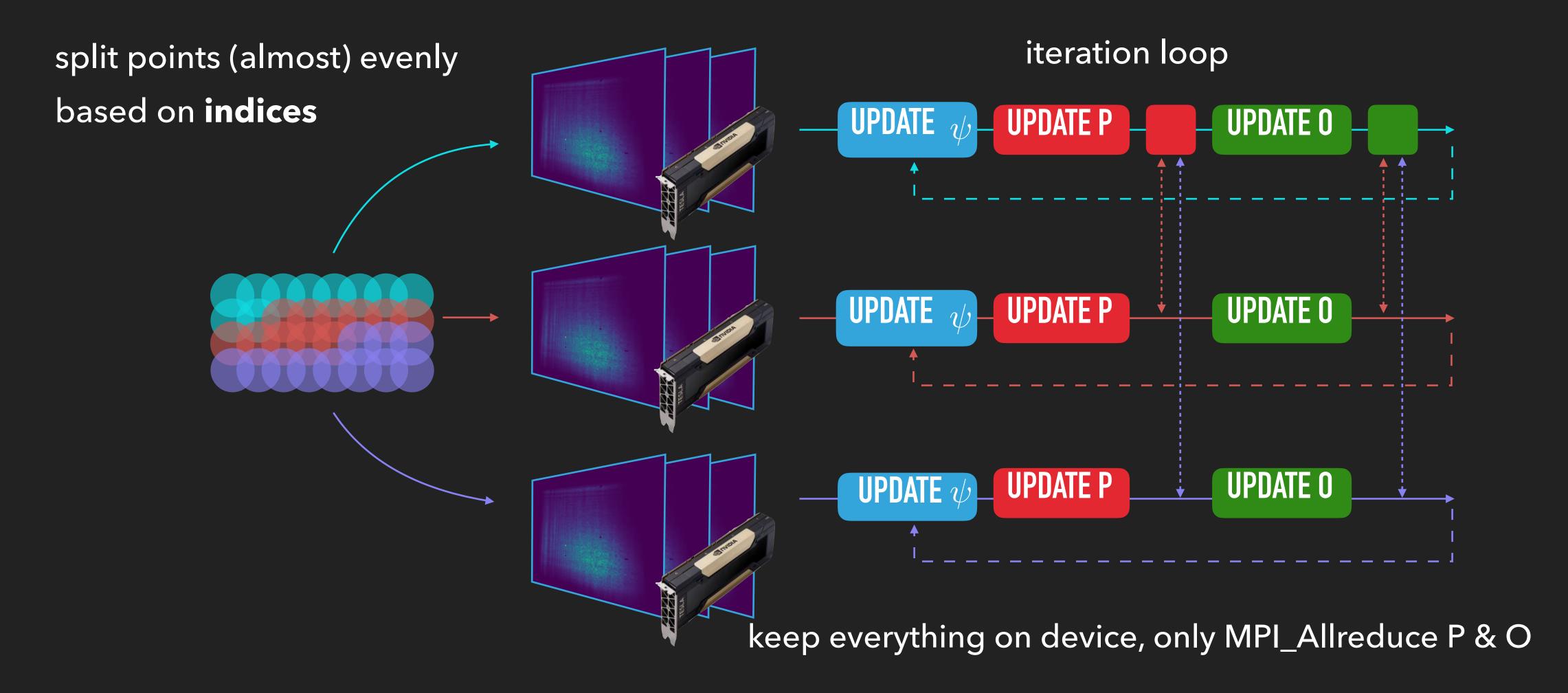


Image from <u>nvidia.com</u>



WORKFLOW FOR PTYCHO MULTI-GPU ACCELERATION



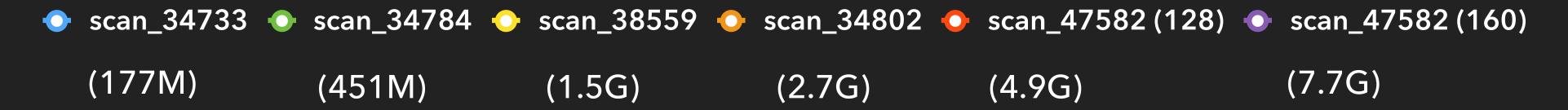


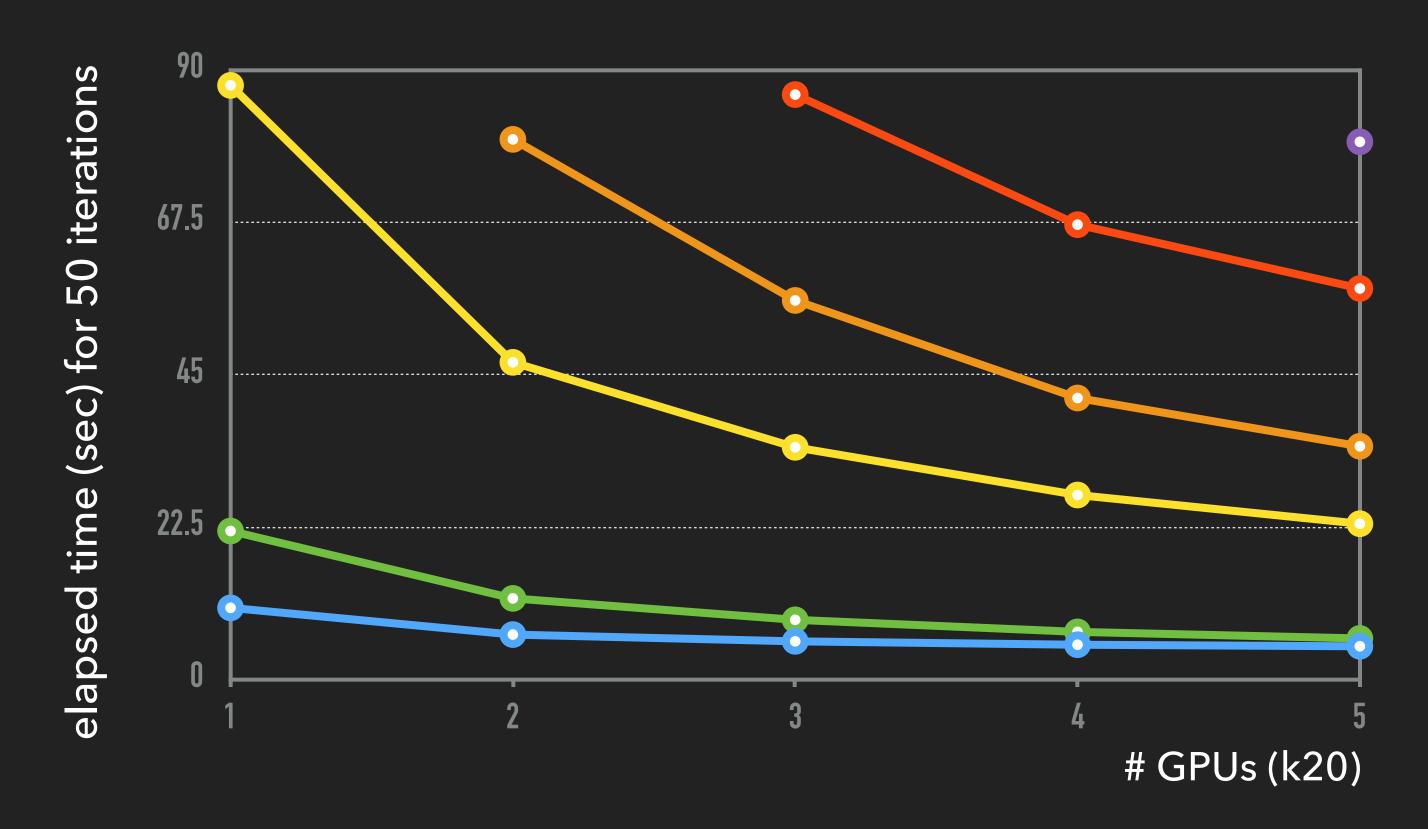
GPU-ACCELERATED PTYCHOGRAPHY SOFTWARE

- Originally written in Python (numpy + scipy + ...)
- ▶ GPU version uses PyCUDA, scikit-cuda, mpi4py
- Computational intensive functions rewritten in CUDA C
- Tremendous speedup: hours => sub-minutes!
- Everything in Python (except for ~ 600 lines of CUDA C) => integration with NSLS-II data acquisition and analysis environment (databroker)...
- Support for distributed GPUs, including GPU clusters and across different servers.
- Graphical user interface (GUI) integrated with multi-GPU backend.



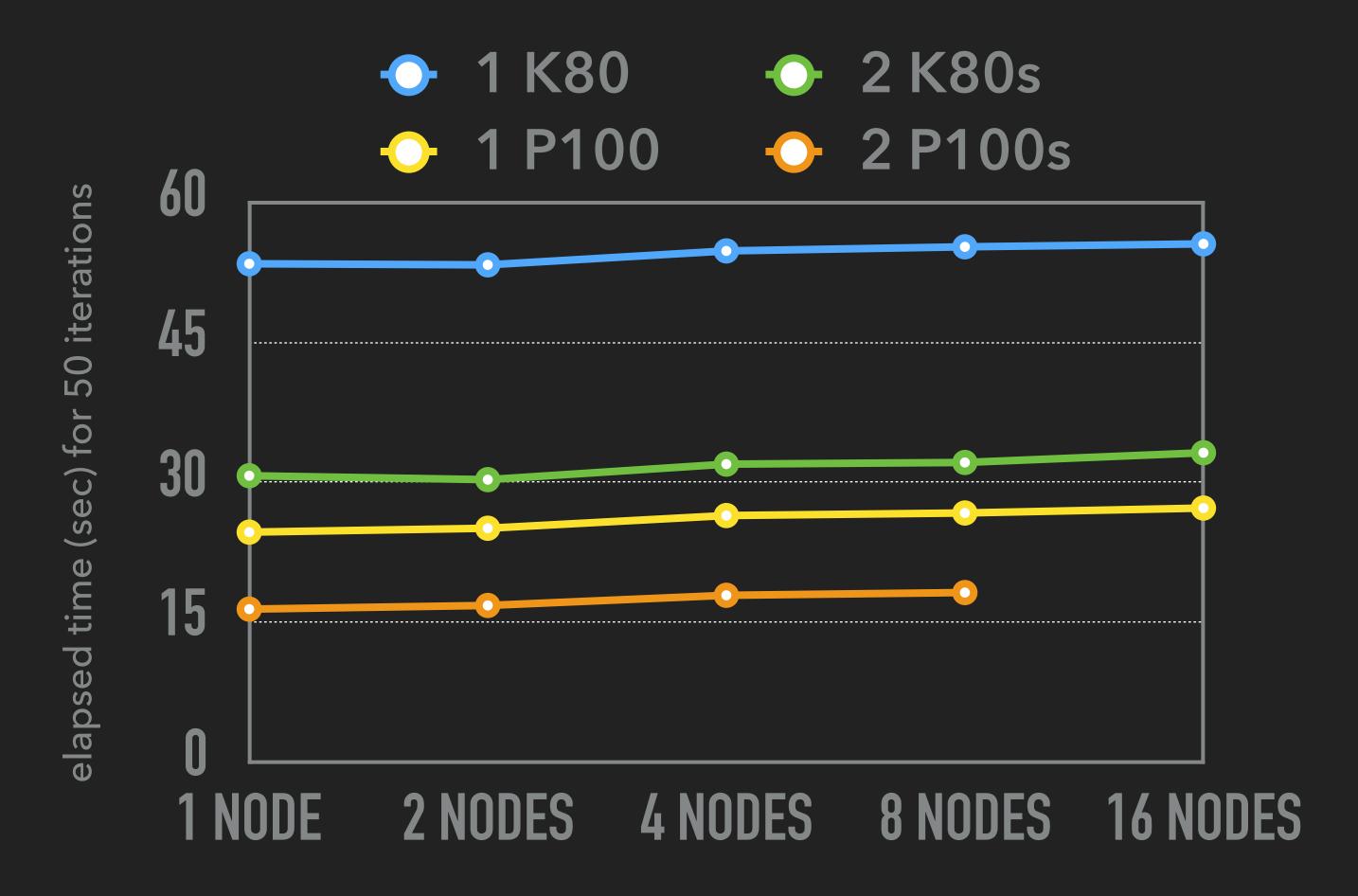
RESULTS: STRONG SCALING







RESULTS: WEAK SCALING

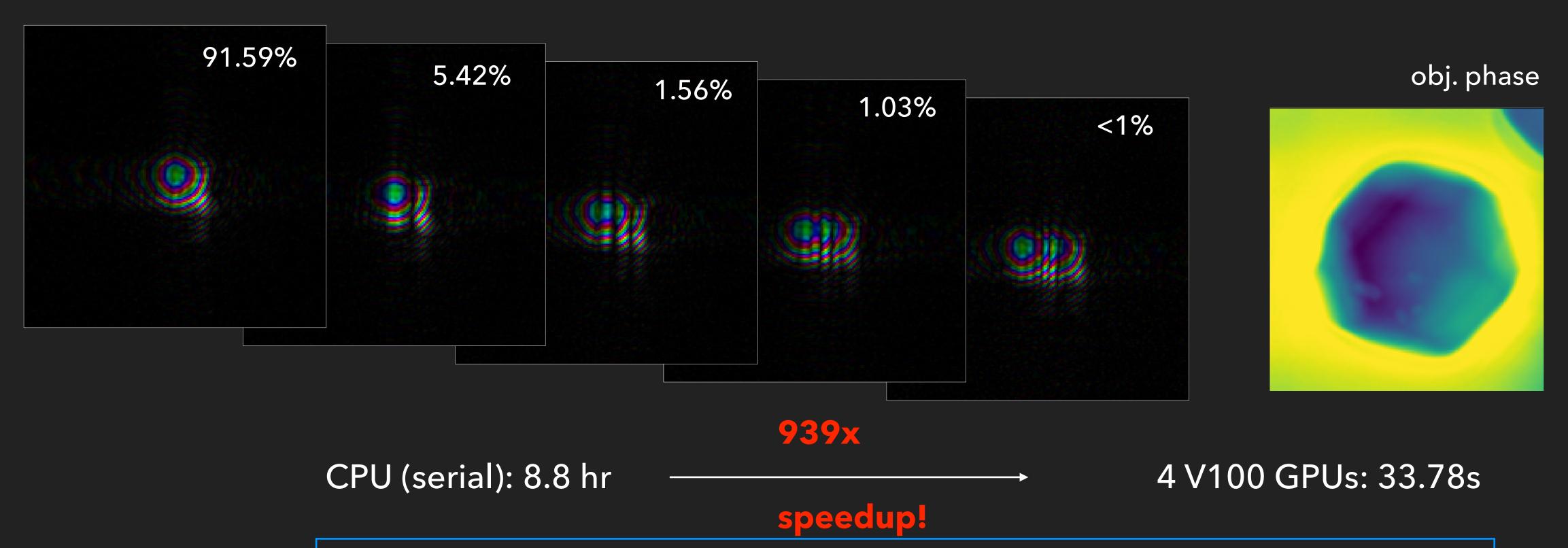




RESULTS: PERFORMANCE IMPROVEMENTS

Test machine: xf03id-srv5@HXN, Intel Xeon CPU E5-2630 v4 @2.20GHz, 256GB RAM, 4 NVIDIA Tesla V100 GPUs. 50 iterations used.

showcase: gold nano-crystal with multi-mode



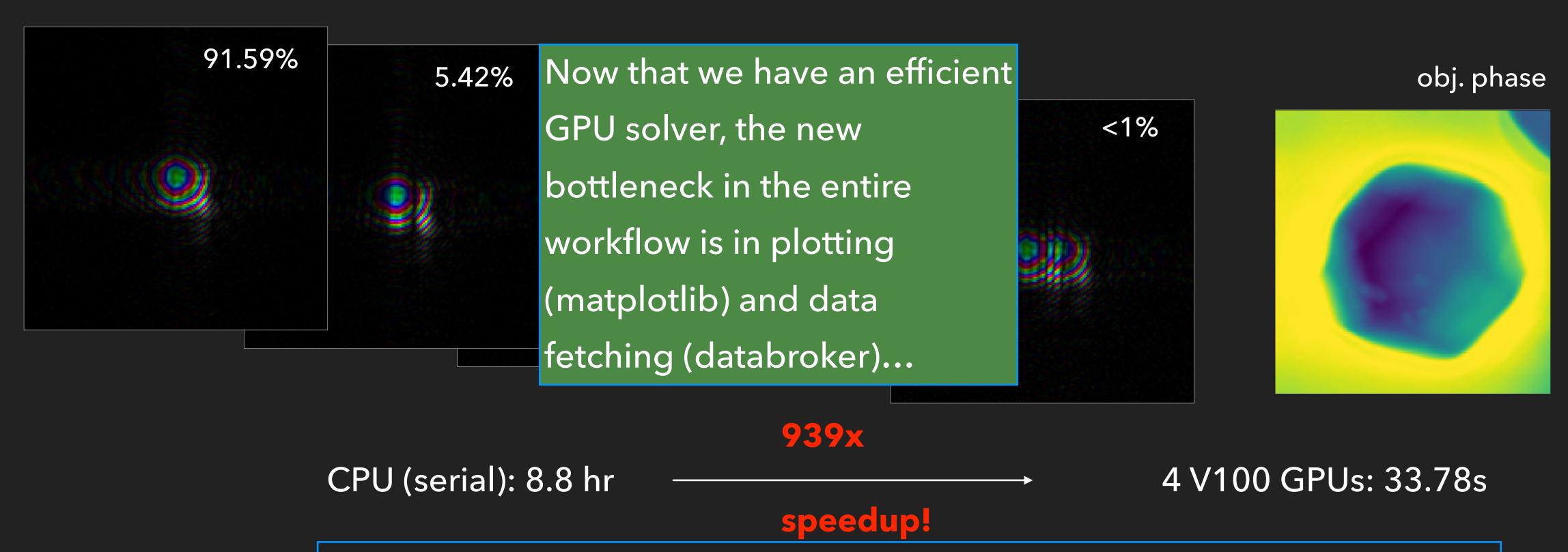


New: MPI-enabled parallel CPU version also reduces the CPU time significantly 10X to 85X with 32 processes depending on data sizes

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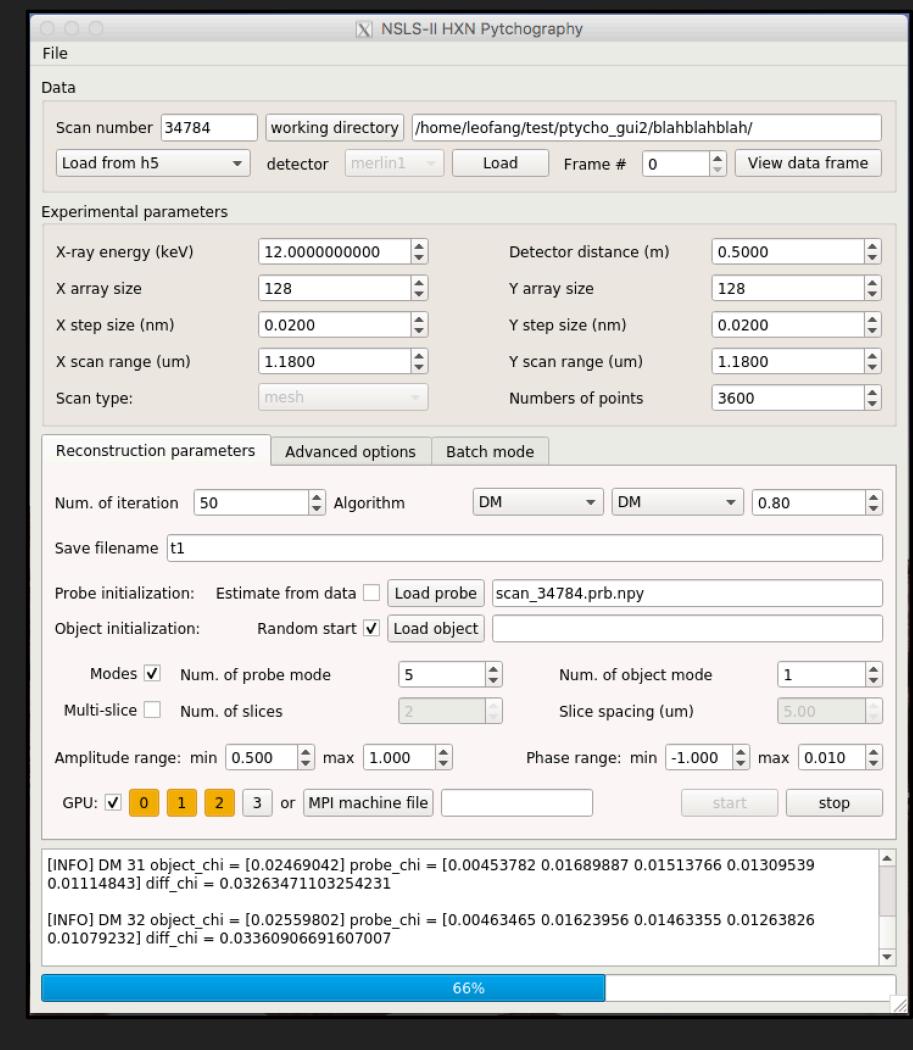
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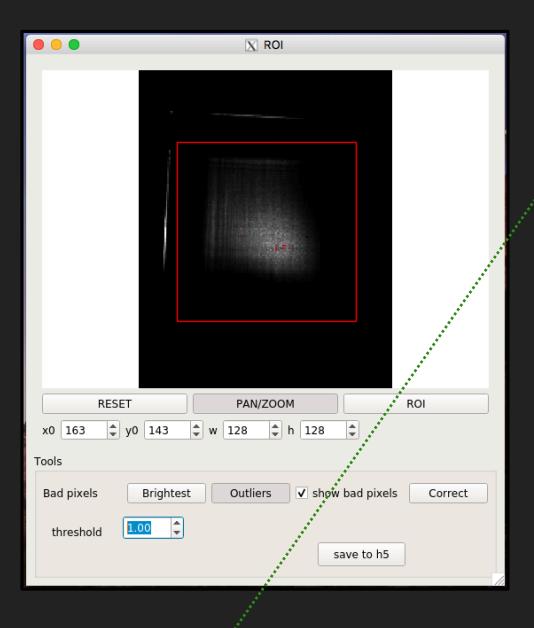


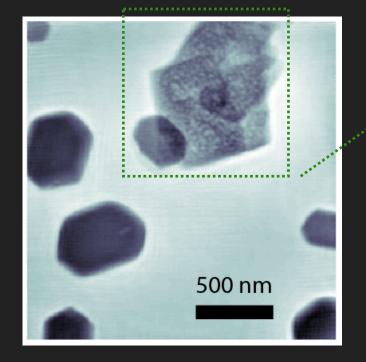


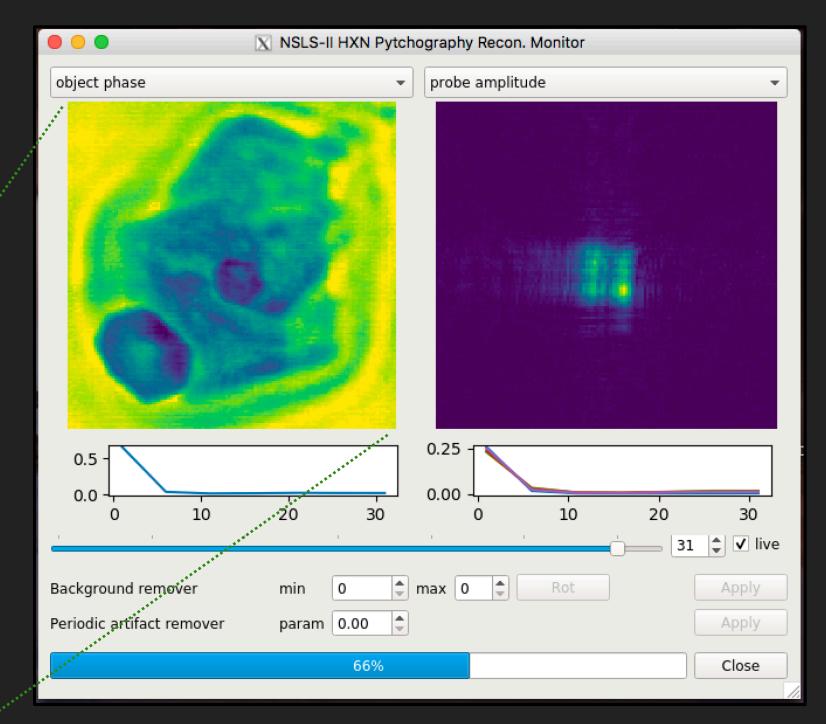
New: MPI-enabled parallel CPU version also reduces the CPU time significantly 10X to 85X with 32 processes depending on data sizes

GRAPHICAL USER INTERFACE









- ★ PyQt5
- ★ Customized event handler
- ★ In-situ processing of raw data
- ★ Efficient realtime monitor
- ★ Clean separation of UI logic, implementation & computation



CONCLUSIONS

CONCLUSIONS

- Fast, in-situ, ptychography reconstruction software is essential for beam line users' productivity.
- We have successfully implemented a GPU-accelerated version of the HXN ptychography reconstruction software with great improvement in performance compared to the serial CPU version.



FUTURE WORK

- Accelerating other solvers (maximum likelihood/error reduction/multi-slice/etc) ongoing
- Further performance fine-tuning (single-precision floating point arithmetics, pinned memory, batch size, CUDA-aware MPI, etc) - ongoing
- Integrating post-processing utilities with GUI
- Optimizing object-update routine
- Modularizing for coupling to SHARP
- Improving quality of amplitude reconstruction



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THANK YOU!

