

PSI

# **Fast GPU-friendly indexing (cell reorientation) algorithm for X-ray crystallography**

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NoBugs Conference 2024 at Grenoble

# Agenda

- 1 Motivation
- 2 Paper on Algorithm, Implementations
- 3 Speed Potential
- 4 Indexing Quality
- 5 Algorithm
- 6 Questions

## New Jungfrau Detector 16M@2kHz

- High data rate - data reduction desirable.
- Fast feedback on measurement data - can a frame be indexed?
- Potential for data reduction - only save indexable frames.
- Potential for data reduction - only save relevant image areas.
- Just speed up indexing.

## Project REDML

- Piero Gasparotto<sup>1</sup>, Luis Barba<sup>2</sup>, HC Stadler<sup>1</sup>, Greta Assman<sup>1</sup>, Henrique Mendonça<sup>3</sup>, Alun Ashton<sup>1</sup>, Markus Janousch<sup>1</sup>, Filip Leonarski<sup>1</sup>, Benjamín Béjar<sup>2</sup>
- PSI<sup>1</sup>, SDSC<sup>2</sup>, CSCS<sup>3</sup>

## Toro (paper/algo/implementation)

- Algorithm
- Pytorch based implementation in Python (Luis Barba)
- Quality and speed measurements
- <https://journals.iucr.org/j/issues/2024/04/00/jo5098/index.html>
- <https://renkulab.io/projects/lfbarba/toro-indexer-for-serial-crystallography>

## Fast feedback indexer (algo/implementation)

- Same algorithm except for details
- CUDA based implementation in C++ (HC Stadler)
- Higher speed for online (frame by frame) processing
- <https://github.com/paulscherrerinstitute/fast-feedback-indexer>

# Speed Potential



On my laptop:

```
$ ./simple-data-bulk-indexer --rep=1000 --quiet --method=ifssr --maxspots=150  
--cells=16 --cands=24 --hsp=$((25*1024)) --triml=.01 --trimh=.3 --delta=.1 --  
minpts=8 --contr=.8 --iter=8 -ths=7 --rblk=3 --ipg=5 --reducalc  
../../data/simple/files/image*_local.txt
```

```
files: 10, operations: 10000  
per operation average timings:  
    clock time: 0.000456699s  
reading time: 4.59597e-07s  
    index time: 0.00226413s  
    refine time: 0.000205426s
```

→ 0.46ms per frame (10 frames indexed 1000 times) for this run

depends on parameters / HW / warmup / disturbances ...

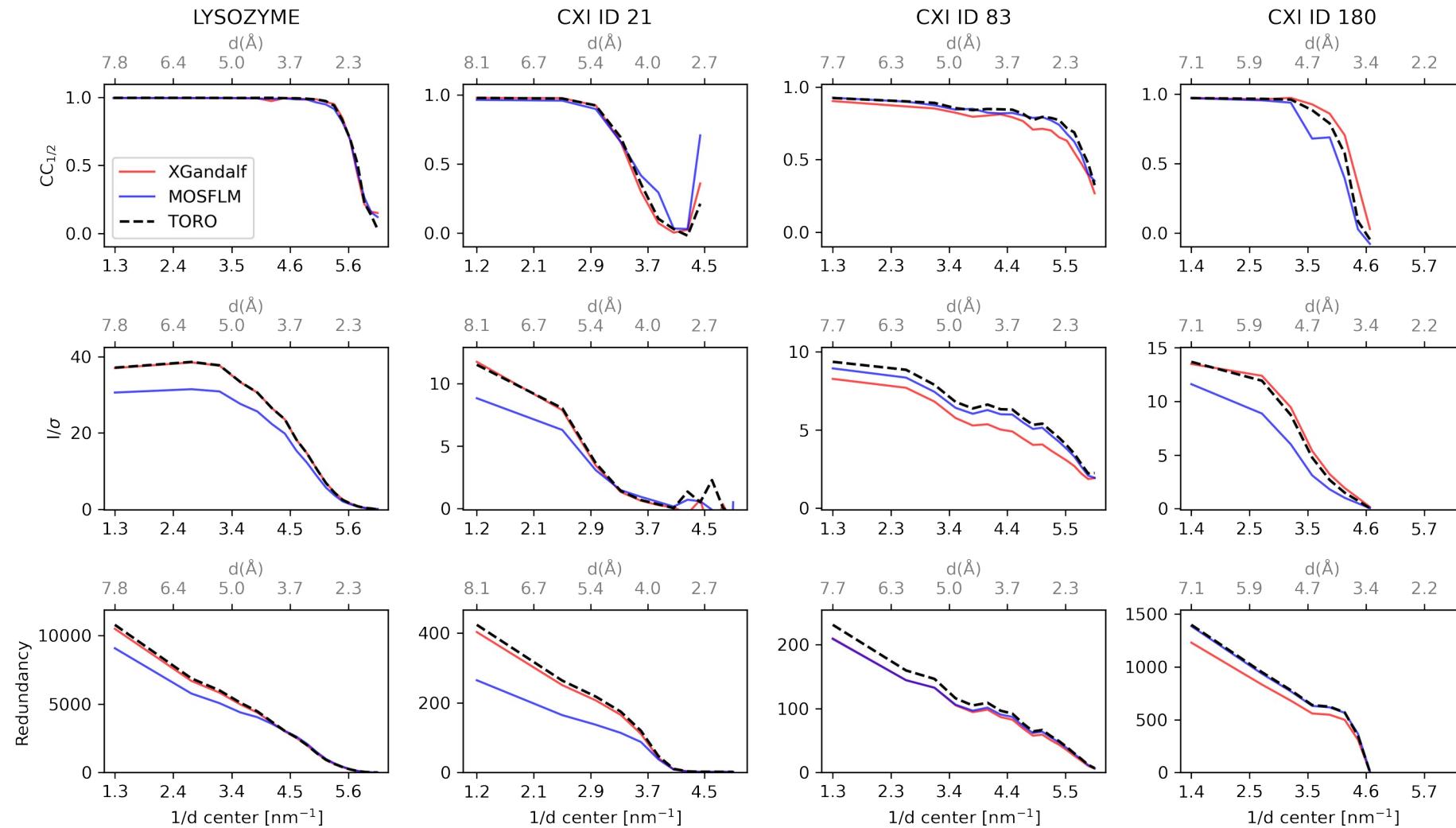
→ try it yourself: examples/cpp-simple-data-bulk-indexer

**Pipeline:** read → index start → index end → refine (→ index start)

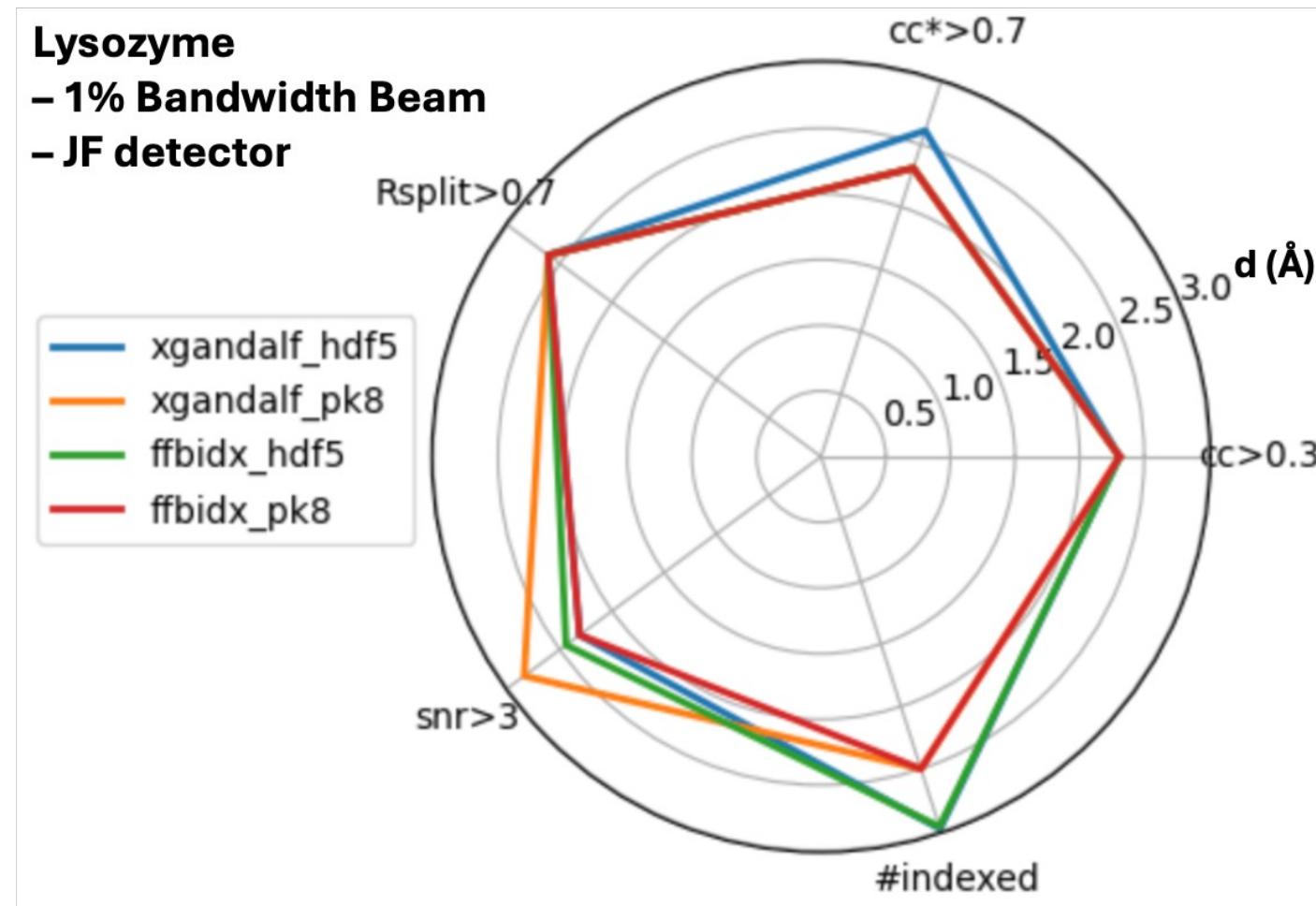
Timings are for **input in memory → output in memory**

# Indexing Quality (Toro)

(see paper)



# Indexing Quality (Fast Feedback Indexer)



Unpublished, measurements and visualization by Duan Jiaxin, PSI

# Algorithm

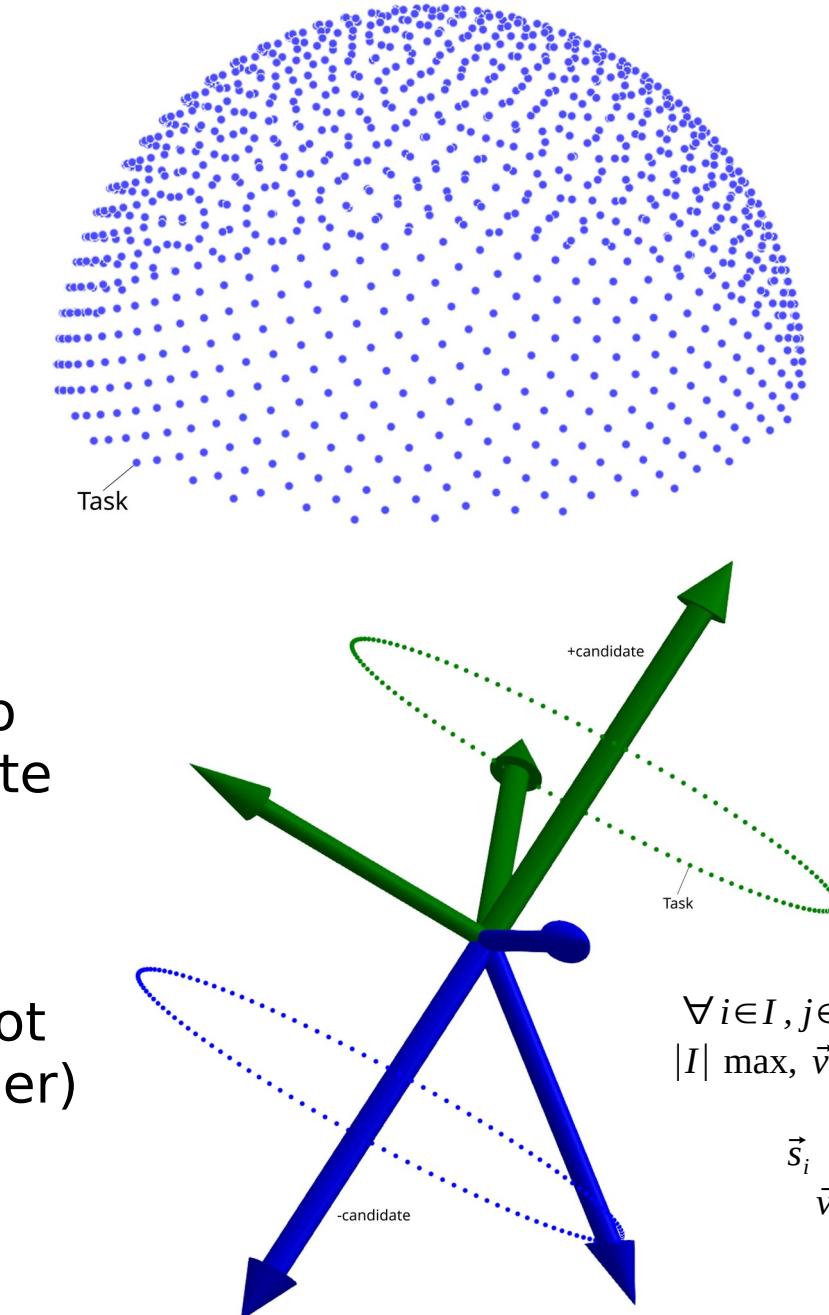
- 1 - Candidate vector sampling**
- 2 - Refinement of best candidates**
- 3 - Candidate cell sampling**
- 4 - Refinement of best candidates**
- 5 - Select solution**

Iterative refinement by least squares fitting to spots that are considered inliers with candidate induced Miller indices.

→ Sampling is GPU friendly (trivially parallel)

Best candidate collection and refinement is not very GPU friendly (synchronization, low number)

→ 1-3 on GPU, 4+5 on CPU



$\forall i \in I, j \in \{1, 2, 3\}: \vec{v}_j \cdot \vec{s}_i$  is ~int  
 $|I|$  max,  $\vec{v}_j$  with given geometry

$\vec{s}_i$  reciprocal spot  
 $\vec{v}_j$  cell vector  
 $I$  inlier set