

Mantid on parallel computing architectures using MPI

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European Spallation Source

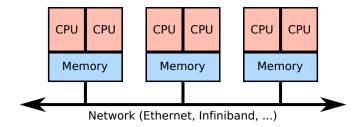
- 1 Introduction: Why and how to run Mantid on a cluster
 - Motivation, goals, and approach
 - Python workflow example with MPI

- 2 Status and results
 - Performance benchmarks
 - Status, plans, and conclusions

Motivation: Why parallelize over many processors?

Computationally challenging problems:

- Exceedingly long computation runtime on a single computer
- More memory required than available on single computer



MPI (Message Passing Interface) is a standard for libraries supporting data exchange between processes running on different computers.

- Speedup compared to current multi-threading Mantid.
- 2 MPI support for Mantid without being visible/explicit for user.
 - Ideally existing Python workflows should work without changes.

Technique areas with many small files

- Process individual files on each compute node.
- Not this talk!

Technique areas with large files (many events and/or spectra)

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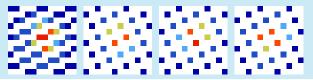


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Example: Python workflow

Consider a typical (simplified) Mantid script:

```
LoadEventNexus (Filename='ABCD_event.nxs',
2
                   OutputWorkspace='ws')
3
   Rebin(InputWorkspace='ws',
4
          OutputWorkspace='ws', Params='100')
5
   SumSpectra (InputWorkspace='ws',
6
               OutputWorkspace='ws')
7
   Rebin (InputWorkspace='ws',
8
          OutputWorkspace='ws', Params='10')
   SaveNexusProcessed (InputWorkspace='ws',
9
10
                        Filename='result.nxs')
```

(replace SumSpectra with Q1D or DiffractionFocussing to obtain SANS or powder diffraction workflow)

I am a workspace

algorithm execution by a process

Run this script with 5 processes using MPI:

```
LoadEventNexus(Filename='ABCD_event.nxs',
OutputWorkspace='ws')
```

```
StorageMode::Distributed
```

```
Rebin(InputWorkspace='ws',
OutputWorkspace='ws', Params='100')
```

```
5 SumSpectra(InputWorkspace='ws',
6 OutputWorkspace='ws')
```

```
Rebin(InputWorkspace='ws',
OutputWorkspace='ws', Params='10')
```

```
9 SaveNexusProcessed(InputWorkspace='ws', 10 Filename='result.nxs')
```

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```

```
Rebin(InputWorkspace='ws',
OutputWorkspace='ws', Params='10')
```

```
9 SaveNexusProcessed(InputWorkspace='ws',
10 Filename='result.nxs')
```

```
StorageMode::Distributed
StorageMode::Distributed
StorageMode::MasterOnly
```

I am a workspace

~ algorithm execution by a process

Run this script with 5 processes using MPI:

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LoadEventNexus(Filename='ABCD_event.nxs',
OutputWorkspace='ws')

Rebin(InputWorkspace='ws',
OutputWorkspace='ws',
Params='100')

SumSpectra(InputWorkspace='ws',
OutputWorkspace='ws')
```

```
Rebin(InputWorkspace='ws',
OutputWorkspace='ws', Params='10')
```

```
SaveNexusProcessed(InputWorkspace='ws',
Filename='result.nxs')
```

```
StorageMode::Distributed
StorageMode::Distributed
StorageMode::MasterOnly
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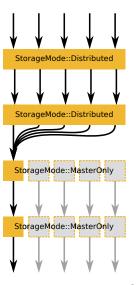
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LoadEventNexus (Filename='ABCD_event.nxs',
OutputWorkspace='ws')

Rebin (InputWorkspace='ws',
```

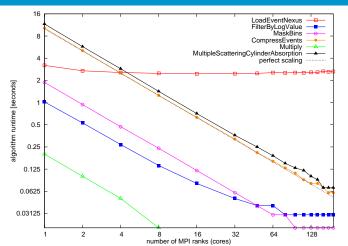
```
OutputWorkspace='ws', Params='100')
```

```
Rebin(InputWorkspace='ws',
OutputWorkspace='ws', Params='10')
```

```
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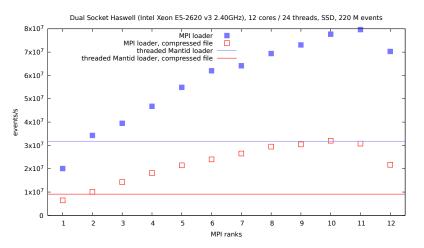


Performance scaling — results for selected algorithms



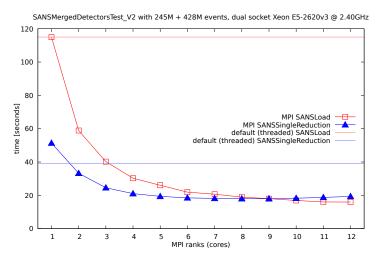
- Algorithms doing I/O generally show limited or no scaling.
- I/O-heavy workflows will exhibit limited scaling.

Optimized MPI version of LoadEventNexus (with Lamar Moore)



- Currently not all flavors of NeXus event files supported.
- Tests on cluster with parallel file system pending.

Performance scaling — SANS2D workflow



 Currently only limited scaling due to expensive algorithm property validation and loading of masking files.

Status and plans (1/2): Python scripts / batch reduction

MatrixWorkspace-based workflows

- Distributed implementation of LoadEventNexus
- Fair number of common algorithms supported
 - Supported does not imply distributed (see SaveNexusProcessed example above)
- Python workflow support without changes?
 - Works for scripts calling only algorithms.
 - Support for lower-level API calls needs more work (in combination with MasterOnly workspaces).
- Basic SNSPowderRedux and SANS2D workflows mostly there.

Workflows including multi-dimensional workspaces

Under investigation (in particular MDEventWorkspace)

Status and plans (2/2): Live reduction and interactive workflows

Live reduction

Reduction of incoming event stream, "live" update of reduced data such as I(Q).

- Should be relatively straightforward.
- Currently not clear whether/when this is required due to lower data rates.
 - Likely to be supported in the future.

Interactive workflows

Most likely no support for:

- Python console
- GUI (Mantidplot)

Distribution and documentation

- No binary release.
 - Usually compilation for specific MPI implementation is necessary, so any binary release would be highly facility- and machine-specific.
 - Not ready for production yet.
- Build instructions and detailed documentation available at http://docs.mantidproject.org/nightly/ development/AlgorithmMPISupport.html.
 - Includes table of supported algorithms.

Conclusion

Running Mantid Python scripts with MPI

- Initial working version nearly there (for MatrixWorkspace).
- Good scaling of most algorithms that do not load or save files.
- (Sometimes greatly) improved workflow performance, depends on file sizes and balance between I/O and computation.
- In some cases MPI reduction is faster than threaded reduction on the same machine.

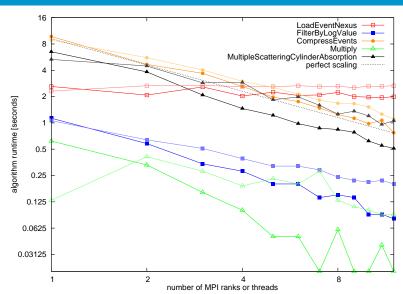
Next steps

- Analyse shortcomings and limitations.
- Acceptance by other developers and power users?
- Not focussed on more optimizations or getting production ready at this stage.

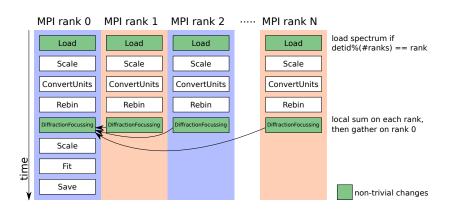
Performance benchmarks Status, plans, and conclusions

Thank you for your attention! Questions?

Backup slide: MPI vs. threading

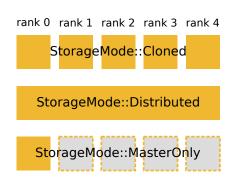


Backup slide: Powder Diffraction example



Backup slide: Workspace storage modes

- Workspace has a new field: StorageMode
- Algorithm behavior can be determined by StorageMode



Backup slide: How...?

Basic idea

- Workspace stored distributed, i.e., across all MPI ranks.¹
- Each rank executes an Algorithm. Typically Algorithm is not aware that it "sees" only parts of the detectors.
- Output of Algorithm is again a distributed Workspace.

The are some Algorithms that do need to know about the other MPI ranks and exchange data with them.

¹Technically, there is an instance of a Workspace on each MPI rank. Logically we interpret these as a single workspace, given by the combination of all spectra from all ranks.