

# Universiteit Leiden Opleiding Informatica

	Developing an integrated environment
	for OPT image reconstruction
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#### Abstract

OPT image reconstruction originally was a very slow process and was completely disjoint from the image acquisition process preceding the reconstruction. This thesis presents an integrated environment to achieve faster reconstruction and bringing the image acquisition and reconstruction closer together. A distributed implementation of the reconstruction algorithm will be presented to improve the reconstruction speed, which significantly decreased reconstruction times. Secondly, a user-friendly prototype GUI application and web service have been developed which allows OPT system operators to easily submit reconstruction jobs of their input images and view a visualization of the reconstructed data within the application. Together these two parts resulted in OPT image reconstruction being significantly faster and more accessible for OPT system operators.

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# Chapter 1

# Introduction

*Optical projection tomography (OPT)* is a form of tomography involving optical microscopy [Shao4]. It uses visible light to image small animals, organs or embryos. In microbiology, *green fluorescent protein (GFP)* is frequently used as a tracer molecule by first inserting the gene for constructing GFP in the gene for a protein. This causes the cell to produce GFP whenever the protein is produced, attached to the protein. Under UV light, these GFP molecules emit green light and this can be used to determine where the protein is located. OPT is one way to image this in 3D.

*Confocal laser scanning microscopy (CLSM)* is an alternative imaging method to construct 3D images using visible light. CLSM has the disadvantage that the light rays cannot reach deep enough in the specimen. With CLSM, the specimen can be sectioned in in several thin layers and the different CLSM results can be stitched together in order to construct a larger 3D image. OPT has the advantage that it allows larger specimens to be imaged without sectioning. OPT basically fills a hole in the resolution spectrum: cells and tissues can be visualized using CLSM, OPT can be used to visualize small organs and embryos and larger organisms can be visualized using for example *magnetic resonance imaging (MRI)*. Figure 1.1 shows some alternative imaging methods for low-resolution objects with the range of resolution they can be applied in.



Figure 1.1: Overview of different small-scale imaging modalities and the range of resolution it can be applied in [KBK<sup>+</sup>09]. Both OPT and CLSM are part of light microscopy.



Figure 1.2: OPT image aquisition: OPT imagine system and OPT images generated by it.



Figure 1.3: Slice-based OPT image reconstruction.

An OPT imaging system, pictured in Figure 1.2, is set up to acquire images of a specimen from hundreds of angles along the vertical axis by rotating the specimen around using a stepping motor. This part of the process is known as image acquisition.

Using the OPT images acquired through such an OPT imaging system, a reconstruction can be executed. This reconstruction is slice-based and visualized in Figure 1.3. These slices can be stacked vertically to obtain a 3D voxel model of the original specimen, shown in Figure 1.4.

The reconstruction process is fairly slow: it takes over 20 minutes on a modern processor using an existing MATLAB implementation. Every slice is reconstructed one after another and takes about a second per slice. As actual input sizes result in 1360 slices being reconstructed and input consists of 400 angles, this is a large bottleneck. It is desired to reconstruct more quickly to, for example, determine whether or not taken OPT



Figure 1.4: 3D visualization of a zebrafish produced by Amira [FEI]. Two channels are visualized: a brightfield channel visualizing the zebrafish itself and a fluorecent channel, highlighting points of interest.

images are of acceptable quality or not as during image acquisition there are several parameters that can affect the quality of the result, such as exposure time. Besides this, another problem is that since the acquisition and reconstruction processes are fairly disjoint, having your images reconstructed requires transmitting the data to someone who is capable of initiating the reconstruction process in the first place.

There are thus two factors that make this process slow: the computationally intensive reconstruction and the requirement for someone else to initiate the process. This project researches a solution to both of these problems in order to increase the speed at which reconstructions can be acquired.

To improve the execution time of the reconstruction, we have chosen to develop a distributed implementation of the reconstruction algorithm, aiming to reconstruct multiple slices in parallel, to run on the Leiden Life Sciences Cluster (LLSC). Chapter 2 will elaborate further on the LLSC and the implementation of this algorithm. Experiments have been performed demonstrating the performance and scalability of the parallel implementation in Chapter 3.

To supplement bringing the acquisition and reconstruction process closer together, we have developed a prototype web service and GUI application, serving as a proof of concept for more accessible reconstruction job submission to the cluster. Chapter 4 will elaborate further on this application. Finally, in Chapter 5 some conclusions will be drawn about the entire project and suggestions for future work will be enumerated.

# Chapter 2

# Methods

This chapter will cover the approach to the problem, what hardware and software was used for the implementation and the optimizations applied to the baseline distributed implementation.

## 2.1 The Leiden Life Sciences Cluster

The *Leiden Life Sciences Cluster (LLSC)* is used to run the software. It is a cluster consisting of three user nodes, a range of compute nodes and a file server. The Maui scheduler is used to schedule the submitted jobs to the cluster on the user node together with the TORQUE resource manager.

The compute nodes are not homogeneous: different processor types are used across nodes and the amount of physical memory differs per node, but is at least 8 GB. The following kinds of nodes were used during the experiments:

- cpu-5150: 10 nodes, two Intel Xeon 5150 processors per node (4 cores total)
- cpu-x5355: 1 node, two Intel Xeon X5355 processors per node (8 cores total)
- cpu-x5450: 2 nodes, two Intel Xeon X5450 processors per node (8 cores total)
- cpu-e5430: 5 nodes, two Intel Xeon E5430 processors per node (8 cores total)

No further discrimination is made based on the physical memory amount each node had. All nodes are interconnected to each other and the file server using gigabit network interfaces.

# 2.2 Baseline sequential implementation

A simple implementation of the entire algorithm to reconstruct slices from OPT input images already existed in MATLAB [Tan16]. The baseline sequential implementation is a fairly straight-forward conversion of this MATLAB code to Python code, utilizing several libraries replacing the MATLAB toolboxes used:

- NumPy
- scikit-image (built-in iradon function)
- pillow (reading multipage TIFF files)
- tifffile (writing multipage TIFF files)

Python 2.7 has been chosen to implement everything on. There exist extremely powerful scientific libraries for Python, such as NumPy. The Python-package *scikit-image* contains several functions to perform the inverse radon transform [sidt]. Python 2.7 also runs on the LLSC without any hassle, which is a large advantage.

The entire implementation can be divided into three parts:

- 1. Reading the input image data into a large NumPy array, then constructing *sinograms* for each of the output slices from the input image data
- 2. Applying the *inverse radon transform* (iradon) on each of the sinograms, generating *reconstructed output slices*.
- 3. Normalizing the output slices from double to uint8 and writing back the image data to an output file.

#### 2.2.1 Constructing sinograms from the input images

Sinograms can be constructed straight from the input images as illustrated in Figure 2.1. One sinogram consists of rows of pixels. To construct sinogram at depth x, all input images (angles) are iterated in order and for each image (angle) the pixels in row x are copied to the sinogram. By stacking these rows of pixels vertically, a sinogram is obtained. A sinogram contains all data needed to reconstruct a slice at a specific depth of the object.

An alternative, useful way of visualizing sinograms is by layering the different OPT images (projection angles) vertically first, as shown in Figure 2.2. Sinograms can then be obtained by taking a cross-section from this stack of OPT images. This clearly illustrates the data dependency a sinogram has: it needs one row of pixels from every OPT image.



Figure 2.1: Constructing sinograms from input OPT images and reconstruction of slices.



Figure 2.2: 3D visualization of the relation between OPT images and sinograms. Each cube represents a pixel.



Figure 2.3: Transposition of the input image.

It should also be noted that the images in Figure 2.1 are transposed compared to the original input images. This is done for consistency in this thesis. In the actual implementation, the image may still be non-transposed, in which rows refer to columns in the input data. A view can be constructed on top of the original data that represents the transposed form as shown in Figure 2.3. Whenever it is relevant that the original data is not this transposed variant, it will be stated.

#### 2.2.2 Applying the inverse radon transform on each of the sinograms

Each sinogram contains data needed to reconstruct a single slice at a specific depth. By applying the inverse radon transform (iradon) on the sinogram, its slice is reconstructed. Figure 2.1 also visualizes this process: it is entirely independent of other sinograms.

The inverse radon transform technique used, known as filtered backprojection, consists of two phases [Tof96]. A Ram-Lak filter is first applied on the sinogram in the frequency domain of the sinogram to avoid blurry results. Then, backprojection is used to actually reconstruct the slice. Due to the internals of these operations, the pixels in the result of the filtering are a floating point type. The pixels after backprojection are of floating point type as well, as backprojection simply sums up all backprojections of rows of pixels in the sinogram at an angle on top of an initially square output image with all values set to o. The length of a side of this square image is usually maximized to  $\lfloor \sin(\frac{1}{2}) \times w_{sinogram} \rfloor$  to ensure all backprojections fully fit inside of the output image even when backprojected diagonally. Figure 2.4 enumerates the different kinds of data the input image is being pulled through and the associated datatypes of the entire process.

#### 2.2.3 Normalizing the reconstructed slices and writing back the result

As described in Section 2.2.2, the result of the inverse radon transform is an image where each pixel is a floating point value. These pixels are normalized to a single byte (uint8) before writing all reconstructed

- 1. OPT images (400 pages (angles),  $1036 \times 1360$  uint16 pixels per page, ~1.1 GB)
- 2. Sinograms (1360 sinograms, 1036  $\times$  400 uint16 pixels per sinogram)  $\downarrow$  (1D FFT)
- 3. Sinograms, frequency domain (1360 sinograms)
   ↓ (Ram-Lak filter)
- 4. Filtered sinograms, frequency domain (1360 sinograms)  $\downarrow$  (1D FFT<sup>-1</sup>)
- 5. Filtered sinograms (1360 sinograms,  $1036 \times 400$  float pixels per sinogram)  $\downarrow$  (backprojection)
- 6. Reconstructed slices (1360 slices,  $732 \times 732$  float pixels per slices)  $\downarrow$  (normalizing)
- 7. Normalized reconstructed slices (1360 slices,  $732 \times 732$  uint8 pixels per slices,  $\sim 0.7$  GB)

Figure 2.4: The different kinds of data the input image is pulled through. Dimensions of typical input data are listed as well and the datatype of each pixel.

slices back. This reduces the file size significantly and makes the result it easier to view in most image viewers. The normalization is performed by first computing the global minimum and maximum pixel values across all slices, *amin*, *amax*. Using these values, all pixel values are rescaled from [*amin*, *amax*] to [0,255] using Equation 2.1 and then converted to uint8 (and floored), making it more suitable for writing it back to an image file.

$$normalizedValue = \frac{value - amin}{amax - amin} \times 255$$
(2.1)

The Python-module *tifffile* is used in order to write the normalized reconstructed slices back to the file server as a multipage TIFF-file.

### 2.3 Baseline parallel implementation

In this section the baseline parallel implementation will be described. This implementation intends to achieve a significant and scalable speedup compared to the sequential implementation by distributing the computationally intensive tasks over multiple worker processes. The baseline parallel implementation distributes reading the input file, applying the inverse radon on a range of sinograms and normalizing the output slices. Once normalized, all data still has to be written back. The write-back to an output file is still done sequentially.

*OpenMPI*, an implementation of *Message Passing Interface (MPI)*, has been chosen to facilitate communication between different processes [Pro]. It is one of the most common systems for *inter-process communication (IPC)* on clusters. There also exists a commonly-used Python-package facilitating bindings to MPI: *MPI4Py* [Dal].

Naturally, this implementation can be divided up into four parts, and is very similar to the separation de-

filtering

scribed in Section 2.2:

- 1. Reading the input image data and a range of sinograms at each worker process (in parallel).
- 2. Applying iradon on the range of sinograms assigned to each worker process (in parallel).
- 3. Normalizing the data to uint8 (in parallel).
- 4. Gathering the data to a single node and writing everything back to an output file.

Before the actual program can run on the cluster, it has to be scheduled for execution. Jobs can be submitted to the scheduler using qsub. A TORQUE submission script (an extension to many scripting languages) is passed to the submission application. It contains both directives for TORQUE in script comments, indicating job parameters such as the job name and the actual script. Once scheduled, the actual contents of the script is run. A simple bash script calling mpiexec suffices in this case. mpiexec is responsible for starting the Python script on multiple nodes in parallel, collecting output, handling proper termination and other related tasks.

#### 2.3.1 Parallel input file reading and constructing sinograms

Constructing sinograms from the input file is relatively straight-forward, but all data required for a single sinogram is scattered across the entire input file since each row of pixels in a sinogram is retrieved from another page in the original TIFF file. Thus, practically the entire input file has to be scanned in order to just generate a small range of sinograms, making this not a trivial parallelization.

There are multiple approaches to rapidly reading the input file and ensuring each worker process has a range of sinograms to reconstruct, of which three have been taken into consideration. This section will describe those three methods and list some expected pros and cons for each of them. The performance of these three methods will be evaluated in Section 3.3.

#### Method 1: All processes read the entire input file

A simple way to construct sinograms for all worker processes is by simply having all processes read all pages in the original TIFF file and only taking rows from each of the pages that it needs to construct its range of sinograms, copying these rows to a memory buffer. This approach does not require any IPC, but requires each process to fetch the entire input file from the file server and iterate through it, which logically does not scale well with more processes. Figure 2.5 visualizes this.



Figure 2.5: An example with two processes: all processes read the entire input file and copy required rows of pixels to their local sinogram array. No IPC is needed.

#### Method 2: One process reads the entire input file

An alternative approach is to assign one process to read the entire input file. This way, the file server no longer has to transmit the entire file to all nodes, which may lead to faster execution times. This process then has to transmit data that is belongs to another process's sinograms to that specific process, though, making it more complex than the previous variant. Figure 2.6 visualizes this method and the IPC required to get rows of pixels at the right process.

Due to the relation between the input images and the sinograms, at most one row of pixels can be transferred from the input image to a sinogram of another process at a time without extra intermediate steps. This results in many small MPI send requests, which could result in a lot of extra overhead. It is also not desired that a receiving process of a row of pixels synchronizes with a sending process. To avoid this, all processes construct a large array representing all sinograms this process should reconstruct. By initializing asynchronous receive requests (Irecv) with location-identifying tags for each row of data in the sinograms before any process sends data, any process can now fill any row in that process's sinogram-array by issuing a Rsend-request to that node with the specific location-identifying tag, which the receiving process processes asynchronously. Such a location-identifying tag consists of two attributes: the index of the sinogram and the row within that sinogram.

#### Method 3: Each process reads a range of pages from the input file

Instead of having each process read all pages or having one process read all data for all processes, a range of pages can be allocated to each process for it to process. A process may read rows from a page of the input



Figure 2.6: An example with two processes: the first process reads the entire input file. IPC is used to transfer rows of pixels to other processes when needed.



Figure 2.7: An example with two processes: each process has a range of sinograms it has to reconstruct and each process reads a range of pages from the input file.

file that belong to a sinogram allocated to another process, though. Just like previous case, these rows of pixels have to be transferred to another process. The same construction can be used where the receiver will asynchronously receive rows of pixels from any other process. Compared to the previous two cases, a single process no longer has to read the entire input file assuming that more than one compute process is started, which can lead to improved scalability. Figure 2.7 illustrates the IPC that has to take place to get rows of pixels at the right process.

Just like the previous option, this results in equally many Irecv and Rsend operations and may negatively affect performance in general. Less time it required to iterate through the entire input, though, which may compensate for this.

#### 2.3.2 Applying the inverse radon transform on each of the sinograms

Once the input file has been transformed into sinograms and each compute process now has a range of sinograms in its memory, each process can reconstruct the assigned sinograms using the inverse radon transform. This works exactly the same as the method described in Section 2.2.2, but since each process has a small subset of all sinograms assigned, it executes a lot faster when multiple processes are used.

#### 2.3.3 Normalizing the reconstructed slices

Normalization to uint8 can be performed in parallel too. First, the local minimum and maximum pixelvalues in the reconstructed slices are computed in parallel. Each process computes these values for their own reconstructed slices. Then, all the local minimum and maximum values are reduced to a global minimum and maximum value. Once these values are computed, the compute processes can normalize all pixel-values to uint8 using the technique described in Section 2.2.3.

Note that the reduce operation is blocking and also serves as a synchronization point for all processes. There is no way any worker process finishes the reduce operation before all processes have started it.

#### 2.3.4 Gathering the result and writing it back to a file

After normalization, the result cannot instantly be written back as the normalized reconstructed slices are spread across the different worker processes. The result has to be collected to a single master worker process before writing it back to a file, often referred to as a gather-operation. The master-node is always chosen to be the last worker process because due to the way sinograms (and thus slices) are allocated to the processes, the last process has most chance of having fewer sinograms allocated compared to all other process and may thus enter the gather-stage of the entire process more quickly as it has to normalize fewer slices, reducing the amount of time other processes have to potentially wait on the master process to be ready.

The master-node preallocates a 3D array in which all reconstructed slices will be placed. MPI has efficient gather-procedures built-in. Gatherv is used to gather data to the master node. After the gather, the multipage TIFF output file can be generated from this 3D array using tifffile as described in Section 2.2.3.

#### 2.3.5 Data flow

The concrete actions of the entire process are enumerated in this section. Additionally, this section presents a data flow diagram illustrating all IPC between different worker processes and file operations with files on



Figure 2.8: Full data flow diagram using the third sinogram construction method described in Section 2.3.1.

the file server. Figure 2.8 shows the data flow diagram. The colors prefixing the actions in the steps listed below match with the line colors illustrating the associated data flow in this figure. The steps enumerated below hold for the third method described in Section 2.3.1, where each process reads part of the input file and sends rows of pixels of sinograms to other processes.

- 1. Through TORQUE, using a job script running mpiexec, all *n* worker processes are started.
- 2. Worker processes read their *pageCount* / *n* pages from the input file from the file server.
- 3. Worker processes distribute the partial sinogram data read from their file and receive all sinogram data this process will compute (rows of pixels).
- 4. Worker processes reconstruct all slices from the sinogram data the process now has in its memory.
- 5. Worker processes compute the maximum and minimum values in their own result images for normalization.
- 6. Worker processes (all)reduce the local maximum and minimum values to global maximum and minimum values.
- 7. Worker processes normalize their own result images to uint8 using the global maximum and minimum values.
- 8.  $\bigcirc$  All normalized result images are gathered at worker process *n* (the master process).
- 9. Worker process *n* creates the output image file and writes all gathered pages to the file server.

The data flow diagram can trivially be modified for the other two methods of constructing the sinograms. For

the method 1 described in Section 2.3.1, where every worker process reads the entire input file, the -colored line (and associated step) can be removed. As each process simply derives all data from the entire input file, no IPC is required for sinogram construction.

For method 2 described in Section 2.3.1, where a single process reads the entire input file but transmits sinogram data to other processes through IPC, the data flow diagram can be modified by ensuring the colored line only goes to one process: the single process that reads the entire input file. IPC is then required to send sinogram data to other processes. The colored line represents this communication and should thus always start at the process reading the entire input file and transmit data to all other worker processes. It will obviously never receive data.

## 2.4 Optimizations

In Section 2.3 the baseline implementation has been described. This implementation can make use of multiple of the LLSC's compute nodes to reconstruct the slices in parallel. In order to speed up execution time more, a number of optimizations have been applied to this implementation, which are described in this section.

### 2.4.1 Backprojection C Python-module

The inverse radon transform consists of two phases: a ramp filter and the actual backprojection. The *scikit-image* implementation was originally used for both of these operations. Whilst the first phase is relatively fast, the second phase is extremely slow due to many large array operations.

To improve the performance, a Python-module written in C has been developed for faster backprojection. The algorithm used within this Python-module is heavily based upon the code used in an existing module for MATLAB [Orco6]. This module iterates the memory of the output image in sequential order, speeding up memory access speeds, avoids unpredictable branches and does not allocate memory in the algorithm's hot loops.

#### 2.4.2 Implicit transposition of the input data

In the original input data, the image is rotated in a way such that the lines of pixels that need to be copied into sinograms are not continuous memory, assuming the default row-major arrays are used, as shown in Figure 2.3. This requires copying each line of pixels, which are in fact columns of pixels in the original image, into a temporary buffer before it can be sent using a MPI message. This can cause a lot of extra overhead. NumPy

supports both row-major and column-major arrays. By reading the input images into a column-major order array, single rows of pixels are continuous memory. Whilst the rows of sinograms are sent from column-major order arrays, they are still received in a row-major order array (the sinogram array), which has the effect of implicitly transposing the input image data. Obviously, reading all data into a column-major array is more expensive than reading it into a row-major array, since the input TIFF images are scanline based (the pixel data is stored row by row). This is a trade-off which can have either a positive or negative effect on the total execution time.

#### 2.4.3 Batching Rsend and Irecv

A potential limitation of performance of the sinogram construction process is the large amount of Rsend and Irecv operations if each process does not fetch all sinogram data on its own. Every single process has to reconstruct a range of sinograms. From any input image (angle), any process needs multiple rows of pixels given that it is assigned multiple sinograms to reconstruct. By ensuring these allocated sinograms are a continuous range, it is possible to batch multiple Rsend and Irecv requests of a rows of pixels to reduce the amount of messages being passed around. This utilizes that adjacent rows of pixels in an OPT image is continuous memory, see also Figure 2.2, assuming that the previous optimization is also applied.

It is important to note that with this optimization it is no longer possible to directly Irecv into the sinogram array of the compute process, as multiple rows of pixels being received at once belong to different sinograms and is thus not continuous memory in the receiving process's sinogram array. It is however still possible to transform this kind of input to an array of sinograms after all data has been received without any extra IPC using array re-interpretation, by layering the sequences of rows of sinogram pixels on top of each other, then taking cross-sections of this stack. This is practically the same as visualized in Figure 2.2, but the depth axis is now only #assignedSinograms deep rather than the total amount of existing sinograms.

#### 2.4.4 Parallel page compression and streamed TIFF write-back

Compression reduces the total amount of the data being transmitted between nodes and the file server. Handling data as soon as it arrives instead of a formal gather can improve execution time more as well, as it avoids blocking for an unnecessary long time on the master process even though this master process can do work with partial data. This section describes this optimization in detail.

#### Compression

In the baseline implementation, after parallel normalization, all data is gathered to a single node, which then writes all data back to a file on the file server at once. Both the gather and write-back are significantly limited in execution speed by I/O times. The TIFF file format supports per page deflate compression. By compressing the normalized page data in parallel on all compute nodes the page data can be compressed quickly. Typically, after compression, the page size is reduced to about 40% of the original output data size. Thanks to compression, the gather and write-back thus get a significant speedup as less data has to be transferred from compute nodes to the master compute node and from the master compute node to the file server.

#### Streamed TIFF write-back

To improve the gather and write-back times even more, some sort of streamed write-back has been implemented. Typically, the gather operation blocks until all data has been gathered to the master node, after which the master node will write all gathered data back at once. This gather operation has been replaced by a mechanism that allows compute nodes to asynchronously "push" TIFF-compatible<sup>1</sup> compressed data to the master node. The master node simply waits for any data to arrive, then immediately writes the compressed data to the output file. Once the master has received all data, it will write back some final metadata and filling in some gaps describing where the compressed page data can be found.

Pushing data to the master node utilizes a similar technique to what is described in the second method in Section 2.3.1. The master process initializes a large 2D array. One dimension is used to index the slice index (which is equal to the output TIFF page index), whilst the other is used to contain the compressed data that should be saved. The master initializes, for each page, an asynchronous receive request using Irecv. To ensure this buffer is ready for receiving before any other process starts sending compressed data to avoid blocking worker processes for a long time, this is initialized before the normalization procedure (Section 2.3.3) starts. Since the normalization procedure contains a synchronization point between all worker processes, it is guaranteed that this buffer is initialized before any process sends data.

Due to compression, the page data size may vary between pages. The buffer the master initializes must be large enough to receive any length of compressed data. It must also be possible to recover the original length of the data once the master receives the data. Other TIFF field values, such as the original image dimensions, are constructed at the worker processes in parallel too and must be transmitted to the master process. This data is all packed using a simple binary format that allows recovery of the original data on the receivers end.

<sup>&</sup>lt;sup>1</sup>Before page data is written to a TIFF file, it is transformed slightly, which is an operation that has been moved to the compute nodes too.

Note that the compressed page data can be written back out of order to the TIFF file now. The TIFF file format is extremely flexible since nearly all data can be written anywhere in the file. Data is often referenced through an offset mechanism, as described in the TIFF file format specification [Inc92]. This mechanism can be exploited to implement such streamed write-back.

In short, as visualized in Figure 2.9, each page in a TIFF file is known as an *image file directory (IFD)*. An IFD is a set of fields, consisting of a TIFF tag, identifying what kind of value it is, and the associated value. Basic TIFF fields are, for example, the compression kind of the data and the image dimensions. If the data is 4 bytes in size or less, it is simply written next to the tag in the IFD. Larger values, such as strings identify the picture name or the actual image data, are written somewhere else in the file, usually right after the page's IFD entry. In these cases, an offset it written next to the tag in the IFD describing that page, the image data can be written in advance of all other data. After writing all the compressed image data in any order, IFDs for all pages are generated and written sequentially at the end of the file, filling in correct data offset values for the images.

In contrast to Figure 2.9, the IFD entries could be written right after or before the associated page data. This has the disadvantage, compared to just writing it all at the end of the file, that the next IFD offset is not always known in advance due to the out of order page data write-back. Thus, an extra pass through the file is then required to patch these offsets.

Finally, it should be noted that compressing the output data is not always desired. In order to read the resulting file, decompression is required. It has been observed that sometimes software processes compressed input data slower than non-compressed data, in which case not compressing the output pages at all may be preferred.



Figure 2.9: Output file structure with parallel compression and streamed TIFF write-back.

# Chapter 3

# Experiments

A set of experiments have been conducted to determine how well the implementation scales with more worker processes and what effect the applied optimizations have on the execution time. All experiments were conducted with warm disk caches by pre-reading the input file twice before running the actual experiment. In order to obtain more consistent results, runs of the experiments were also conducted after one the other, not executing multiple runs in parallel. If multiple runs were to execute in parallel, this could result in multiple jobs doing a lot of I/O at the same time, slowing both runs down significantly. In a practical environment, these things obviously do not matter. All data points in these experiments were obtained by repeating the scenario 10 times, removing the best and worst times and then taking the average of the 8 remaining times. Standard deviations also only use these 8 times.

It should be noted that these experiments were not the only jobs that ran on the cluster. Jobs by other cluster users ran at the same time as these experiments. Whilst the experiment jobs typically had to wait for these jobs to finish, it can still have an impact on the execution times as sometimes these jobs did run concurrently.

Two experiments were conducted which did not actually involve parallel execution: measuring the effect of the backprojection C Python-module optimization and measuring the overhead of MPI with a single process. Three larger experiments followed that which did involve parallelism: measuring which of the three proposed ways to read the input file is the fastest and measuring the effect of the different proposed optimizations on the total execution time. Finally, an experiment is conducted that measures the overall scalability with many nodes.

	Total execution time	Standard deviation
scikit-image backprojection	8764.76 s	12.50 s
Custom backprojection	1017.03 s	0.28 s

Table 3.1: Performance measurement results of the custom backprojection module.

### 3.1 Backprojection C Python-module optimization

The primary single-core optimization applied is the backprojection C Python-module to more quickly reconstruct slices, as the original Python version built into *scikit-image* was deemed too slow. This experiment measures the effect of the module in the sequential implementation.

#### 3.1.1 Experiment set-up

Two variants were constructed, one with the original scikit-image backprojection and one with the custom backprojection. Averages and standard deviations of the total executions time are measured. The times were measured on a cpu-x5450 node.

#### 3.1.2 Results

Table 3.1 shows the results of this experiment. From this table it can be seen that, on a cpu-x5450 node, the custom backprojection module is about 8.6 times faster than the original Python-based backprojection built into scikit-image. This module thus has a lot of impact on the total execution speed of the entire program as a lot of time is spent running the backprojection operation.

### 3.2 MPI overhead

When using the parallel variant with a single process, MPI calls still occur. For example, gathers from and to a single process may come with extra overhead compared to the original sequential version. Within this experiment the difference in execution time between the original, sequential version and the "parallel-ready" version (with MPI calls) executing using one process is measured and compared.

#### 3.2.1 Experiment set-up

Two versions are used: one without MPI calls and one with MPI calls. The total execution time is measured from the point the files are being opened until the point the output file has been written back. In both

	Total execution time	Standard deviation
Non-MPI variant	1017.03 s	0.28 s
MPI variant (parallel ready)	1019.05 s	0.24 s

Table 3.2: Execution time using a single process: the pure sequential variant versus the variant with MPI calls.

versions, the C Python-module for backprojection is used and input data is read into a column-major array, which is thus implicit transposition of the input data. The two other optimizations cannot be applied at all to a non-MPI version and are thus not enabled in the MPI version too. In the MPI version, the implementation for sharing input where each process reads part of the file is used. In the case of a single process, this obviously does not result in IPC, but can still influence execution time. The experiments ran on a cpu-x5450 node.

#### 3.2.2 Results

Table 3.2 shows the results of this experiment. Clearly, the non-MPI variant is measurably faster than the variant with MPI calls. The overhead is, whilst measurable, still fairly minimal. Given that the total execution time nearly halves when using two processes, the MPI overhead is neglectable.

### 3.3 Fast sinogram construction

Three alternatives were discussed for fast input file reading and sinogram construction in Section 2.3.1. This experiment compares the three and it will be determined which of the methods is the fastest. Two parallel optimizations are relevant: implicit transposition of the input data, which is be enabled for all experiments and batching multiple rows of pixels into a single Rsend and Irecv, which is disabled.

#### 3.3.1 Experiment set-up

The three implementation variants that will be measured are:

Variant 1 All processes read the entire input file (Method 1 in Section 2.3.1).

Variant 2 One process reads the entire input file (Method 2 in Section 2.3.1).

Variant 3 Each process reads a range of pages from the input file (Method 3 in Section 2.3.1).

Only cpu-5150-nodes are used in this experiment to eliminate as much noise as possible. This resulted in only 10 nodes being available, each having 4 cores in total. Experiments were conducted on 1 node (1 and 4



Figure 3.1: Performance comparision between the three tested methods to construct a range of sinograms in each process's memory.

		Variant 1		Variant 2		Variant 3	
Processes	Nodes	Total time	Std dev	Total time	Std dev	Total time	Std dev
1	1	7.78 s	0.02 s	8.30 s	0.04 s	8.25 s	0.03 s
4	1	9.64 s	0.06 s	10.89 s	0.03 s	6.38 s	0.03 s
10	10	5.72 s	0.01 s	14.71 s	0.01 s	3.20 s	0.06 s
20	10	6.20 s	0.03 s	13.89 s	0.01 s	2.71 s	0.08 s
30	10	7.63 s	0.03 s	13.87 s	0.01 s	2.92 s	0.19 s
40	10	8.99 s	0.01 s	13.95 s	0.04 s	2.75 s	0.46 s

Table 3.3: Performance comparision between the three tested methods to construct a range of sinograms in each process's memory.

processes) and 10 nodes (10, 20, 30 and 40 processes). This will allow comparing the three variants effectively with both few and many processes and nodes.

#### 3.3.2 Results

Figure 3.1 shows the average time to distribute all sinograms in a plot. Table 3.3 shows the data in a tabular form, including standard deviations. Clearly, the only variant that scales positively with more processes is the third one. The first variant hovers around 8 seconds. When more processes are ran on a node, the time required to distribute the sinograms seems to increase with this variant. Using the second variant will simply lead to increased times when more processes are used, likely due to the increased IPC required to construct

	Variant 1		Variant 2		Variant 3		Variant 4		
Processes	Nodes	Distr time	Std dev	Distr time	Std dev	Distr time	Std dev	Distr time	Std dev
1	1	1156.11 s	0.40 s	1153.92 s	0.29 s	1175.16 s	0.23 s	1173.61 s	0.36 s
4	1	320.84 s	0.33 s	320.62 s	0.20 s	327.76 s	0.47 s	323.73 s	0.23 s
10	10	130.01 s	0.09 s	130.00 s	0.08 s	124.94 s	0.10 s	125.85 s	0.04 s
20	10	73.07 s	0.16 s	72.78 s	0.10 s	66.20 s	0.19 s	68.01 s	0.19 s
30	10	56.63 s	0.18 s	56.56 s	0.29 s	49.50 s	0.25 s	51.49 s	0.24 s
40	10	46.81 s	0.33 s	46.94 s	0.58 s	39.41 s	0.78 s	41.84 s	0.25 s

Table 3.4: Performance comparision of different kinds of optimizations.

all sinograms whilst this IPC all originates from a single process.

# 3.4 Effect of optimizations

Three optimizations that are expected to have an effect on the distributed implementation have been tested. This experiment compares the optimizations and will measure whether or not the optimizations have a positive effect on the total runtime in the first place.

#### 3.4.1 Experiment set-up

Four variants of the implementation were created:

- Variant 1 No optimizations were applied.
- Variant 2 Implicit transposition of the input data was applied.
- Variant 3 Both implicit transposition of the input data and compression of the output and streamed writeback was applied.
- **Variant 4** All optimizations are applied: implicit transposition of the input data, compression of the output and streamed write-back and batching small requests into a single large one.

The same nodes and process counts will be used as in the experiment described in the previous section. This will result in a good overview on the effects the optimizations have with just a few processes and with many processes.

#### 3.4.2 Results

Figure 3.2 shows a plot of the average total execution time, Table 3.4 shows it in tabular form including the standard deviation. Comparing the four variants, it is clear that the first and second variant barely differ in



Figure 3.2: Performance comparision of different kinds of optimizations.

execution time. Implicit transposition of the input data thus seems to mainly move the slowness of ensuring rows of pixels are continuous memory to another position in the application.

There is a significant difference between the second and third variant, though. Applying parallel output data compression and streamed TIFF write-back has a significant positive effect on the total execution time when many processes are used. When few processes are used, though, this optimization has a negative effect on the total execution time. It is fairly self-explanatory that in this case the processing time required to compress all output data with few processes exceeds the I/O time gain afterwards.

The fourth variant, where small data requests are batched into a larger one, only seems to yield a significant positive effects with few processes, where batching reduces the total amount of requests by orders of magnitude. With many processes, this optimization leads to an increased execution time, likely due to the overhead the extra array re-interpretation brings in order to extract the actual sinograms from the batches of rows.

### 3.5 General scalability

In this experiment it will be measured how well the execution time scales with amounts of nodes and processes. All (classified) nodes running at the time of this experiment in the cluster are fully utilized.



Figure 3.3: Effect of different processor and node counts on the total execution speed.

#### 3.5.1 Experiment set-up

From the previous experiment (Section 3.4), variant 3 will be used in this experiment. Unlike that experiment, though, we will now not limit ourselves to cpu-5150 nodes. As mixing CPU types will result in less predictable results, allocation of nodes will be ordered based on their CPU type to generate more consistent results. The experiment will be conducted with 1, 2, 8 and 18 nodes. First the two cpu-x5450 nodes will be allocated. When more than two nodes are needed, 5 nodes of type cpu-e5430 are allocated, followed by the single cpu-x5355. All 8-node test cases thus use these three types of nodes. Finally, the 10 nodes of type cpu-5150 are used too in all 18-node measurements. Measured process counts are 1, 2, 4, 8, 18, 32, 64 and 104. The amount of processes assigned to each node is kept equal as much as possible, but since cpu-5150 nodes only have 4 cores, it will be more unbalanced when 104 processes are used.

### 3.5.2 Results

Figure 3.3 shows a plot of these results. Table 3.5 shows the data in tabular form with the standard deviation. It is clear that, while the implementation does not scale perfectly, it arguably scales fairly well with an increased number of nodes and processes. An interesting observation is that there is barely any difference between one and two nodes. Thus, having to transmit data between different nodes did not influence the

Processes	Nodes	Total time	Standard deviation
1	1	1037.41 s	0.41 s
2	1	533.57 s	1.61 s
2	2	536.74 s	0.25 s
4	1	272.63 s	0.36 s
4	2	271.83 s	0.38 s
8	1	147.42 s	0.28 s
8	2	139.69 s	0.30 s
8	8	155.62 s	0.21 s
18	8	72.90 s	0.25 s
18	18	72.79 s	0.22 s
32	8	44.91 s	0.43 s
32	18	44.04 s	0.25 s
64	8	32.65 s	0.51 s
64	18	27.59 s	0.37 s
104	18	22.76 s	0.12 s

Table 3.5: Effect of different processor and node counts on the total execution speed.

total execution time a lot. When jumping to 8 nodes, the nodes with less powerful processors are used too, which could explain the increased execution time. Once 64 processes are used, the gains drop significantly when more processes are allocated.

With 104 processes, the total execution time hovers around 22.76 seconds as shown in Table 3.5, which is 45.6 times faster than when just using a single cpu-x5355 node. About 9 of those seconds are spent reading the input file and distributing the sinograms, compressing, gathering and writing the result back, which is part of the entire process that does not scale as well as the computationally intensive reconstruction part. This likely inhibits continued near-linear speedups.

# Chapter 4

# Application

A prototype GUI application and associated web service have been developed to allow OPT system operators to rapidly submit reconstruction jobs to the cluster to quickly get an indication whether or not the recorded images are of proper quality. This application has been developed to bring the image acquisition at the *Institute of Biology Leiden (IBL)* closer to the image reconstruction. Originally, reconstructing images required the OPT images to be transferred to the Snellius building in Leiden. As these files are fairly large, the data is usually physically transferred using a removable medium, as illustrated in Figure 4.1. At the Snellius building, another person can then initiate the reconstruction process.

By introducing an intermediate application which uploads the data directly to the LLSC and automatically starts the reconstruction process, it is no longer required to physically move data around, as visualized in Figure 4.2. Besides, no second person is needed to initiate the reconstruction process. After the fast reconstruction on the LLSC, it is desired that a simple visualization is shown in the application. The *maximum intensity projection (max-projection)* has been chosen as visualization of the output data. The OPT system



Figure 4.1: Original situation requiring manual file transfers and two people.



Figure 4.2: Desired situation with an intermediate application uploading the data to the LLSC.

operator can use this projection to determine, for example, whether or not the data was properly aligned and if the light intensity needs adjustment.

The application consists of a GUI front-end application and a web service back-end, which this chapter will elaborate upon.

# 4.1 The GUI application

The GUI application serves as a simple and accessible method to submit OPT images to the web service and display the resulting max-projection. It also supports viewing the original OPT images from within the application.

#### 4.1.1 Usability specification and requirements

The goal of the application is to have means to easily submit created OPT images to the cluster web service described in Section 4.2. This service should reconstruct the given OPT images and compute the max-projection. The front-end application should be in the form of a GUI and support dragging and dropping the input TIFF images into the application.

Once an image is loaded, the window should be split into two parts, on the left side a preview of the input OPT images is shown. These TIFF images consist of multiple pages, which can be seen as video frames. From within the application, these frames should be viewable and it should be possible to automatically animate through these frames through clicking a play/pause button. A bar indicates what the currently shown frame is and can be moved around to immediately navigate to a specific frame (similar to a video player).

On the right side, the user has the option to initiate reconstruction of the input OPT images. Since this is a lengthy operation, the application should propagate feedback to the user indicating the state of the reconstruction and assuring the user that the application is still working by showing a progress bar. After this process, a max-projection of the input data should be displayed in this right panel.

The GUI application should run on OS X and should be easily startable through the form of an icon in the OS X dock.

#### 4.1.2 Usability measurements

In order to measure the usability, the application prototypes have been tested with three end-users. Feedback and suggestions were recorded, these were used for further iteration of the prototype. It was ensured the users could accomplish getting the output of the reconstruction procedure on the screen. No objective measurements were performed.

#### 4.1.3 Technical components

The implementation is programmed in Python 3. PyQt5 [Lim] is used as the application framework, which are Python bindings for the *Qt framework* [Com]. This framework is event-driven. It consists of a large range of modules cooperating together. This application uses, next to the *Qt Core* module, the *Qt GUI* module for the user interface and the *Qt Network* module for communicating with the web service. The Qt framework was chosen simply because of the author's previous experience with using it. Using Python has the advantage that code to read the input TIFF image for previewing the input data can be re-used. It is also easier to prototype using a scripting language and it is less operating system dependent than C++, which is the language the Qt framework is originally designed for, allowing the application to be initially developed on non-OS X machines.

There is a global tab widget in the application listing the opened files, as shown in the application screenshot in Figure 4.3. The tab widget also contains an initial tab containing a button to browse for files to load and conveying that dragging and dropping files is possible.

When a file is loaded, either by using the browse button in the initial tab or by dragging and dropping a file into the application, the application constructs two components: a controller and a view. The view manages what is shown to the user and can ask the controller to do certain things, such as submitting a reconstruction request to the web service. The controller notifies the view whenever something happens through Qt's signal and slot mechanism. The view gets added to the global tab widget and is immediately displayed after loading a file.



Figure 4.3: Screenshot of the prototype GUI application once reconstruction has finished.



Figure 4.4: Screenshot of the prototype GUI application during upload.

#### The controller

The controller manages data and loading it from both the web server or the local file system. It thus also has an internal model to store some data in, although mostly for caching purposes. The controller has the capability to do the actions listed below.

- Loading a multipage TIFF file representing OPT images from the local file system.
- Requesting to load a specific page from the loaded OPT images asynchronously. The controller uses a
  separate worker thread for this, since seeking and loading OPT images takes a significant amount of
  time and locking up the UI thread is obviously not desired. The controller notifies the view once an
  image has been loaded using a signal. This uses a queue mechanism where the queue has a length of
  one. When the queue overflows, older requests are dropped. The read data from the file is cached in
  the process's memory.
- Starting the reconstruction by uploading the entire OPT image file to the web service. The controller emits signals to the view once a status update is available, like when another chunk of the file has been uploaded, the reconstruction started or when the entire process finishes. The controller also stores an internal state of the reconstruction (not started, uploading, reconstructing, succeeded, failed).
- Canceling an earlier started reconstruction.
- Fetching the max-projection from the web service. The controller notifies the view with the image using a signal once the data arrived.

QNetworkAccessManager is used to communicate with the web service. By hooking to the readyRead signal of a QNetworkReply, partial responses can be read from the HTTP data stream. By reading and parsing lines of data from the network buffer as they arrive in the response of a reconstruction request, the application can read status updates as the web server pushes them (see also Section 4.2.1).

#### The view

The view consists of two sides with a draggable splitter in-between allowing both sides to be resized. The left side contains a view of the input OPT image, displaying one page at a time. A scrollbar below the image can be used to view another image, which signals the controller to load another index. The view connects to a slot of the controller indicating that an image has loaded. When this happens, the view actually updates the displayed image. Left to the scrollbar, a play/pause button is located. By pressing the button, the application will automatically start (or stop) advancing to the next page in the input data every 1/30th of a second. This button uses icons so that the behavior of this button is quickly recognized. These icons were designed by

The right side can show different things and is determined by the state of reconstruction. If the reconstruction has not been started yet, a button is shown to start reconstruction. If it is ongoing, a progress bar and label indicating the status is shown (Figure 4.4). A cancel button is shown too. When the reconstruction finishes, the max-projection will be shown in this panel (Figure 4.3).

with other applications. OS X, however, does not support this by default.

### 4.2 The web service

The web service is programmed in Python and utilizes the *Tornado Web Framework* [Aut]. The module comes with a built-in web service, making it easy to get it up and running. It runs on a user node on the LLSC. There are multiple routes defined which the GUI application calls, described in the next sections. The most fundamental one is the one through which a file can be uploaded to the web service and running the reconstruction request. Whenever this finishes, the max-projection can be requested as a PNG image through another route. There are a few auxiliary routes defined too for downloading the raw reconstruction result, downloading a single slice from the reconstruction result and requesting the cluster its job queue. Within this section, the functionality of the web service will be described.

#### 4.2.1 Submitting a reconstruction request

The route /submitjob can be used to submit reconstruction requests to the web service. It accepts POST requests whose body is a multipart form data (multipart/form-data) request. This body contains a TIFF file, used as input for the reconstruction. Once all data is uploaded, the web service computes the MD5 hash of the entire file. This hash is used to identify the file. The file is saved on the scratch partition on the file server on the LLSC. Once saved, the actual reconstruction process starts.

#### Reconstruction

From the web service, qsub is called to schedule a reconstruction job for the saved file. qsub prints the job id to the standard output stream, which the web service captures. Using this job id, the status of the job can be polled for completion. Using checkjob the status of a given job can be polled. With the -A parameter, the output of checkjob is easier to automatically parse. The State attribute from the output of checkjob is extracted, which indicate that the job may be deferred due to other workload on the cluster, or that it is currently running. Once the job finishes, checkjob will fail to run. Once the web server notices that checkjob



Figure 4.5: Max-projection from a stack of reconstructed slices.

fails, it will verify that the output file exists. If so, the reconstruction is deemed successful, the identifying MD5 hash it written to the output stream too and the request terminates.

#### Max-projection

Once the parallel reconstruction finishes, another Python program is started which computes the maxprojection of all of the reconstructed slices [WM91]. It outputs a single PNG image with the max-projection result. It is computed by loading the entire multipage TIFF image into a large 3D NumPy array, then computing the maximum value along the depth-axis of this array. The result is a 2D image representing the max-projection, which is then saved on the file server and can also be identified using the original file's MD5 hash.

Outlining the entire process: to construct a max-projection from input OPT images, these OPT images are first converted to sinograms (Figure 1.2). Then, slices are reconstructed using these sinograms (Figure 1.3). Finally, using this stack of reconstructed slices, the max-projection can be determined (Figure 4.5).

#### Propagating status to the application

In order to keep the end-user up to date with the current status of the reconstruction (Idle, Running...), the web server occasionally writes the current status as a line in the HTTP response. The request does not terminate until the entire reconstruction either finished or failed.

### 4.2.2 Fetching the max-projection

If the reconstruction finishes successfully, the web service has written the identifying MD5 hash of the input file to the HTTP response body. This MD5 hash can be used with the route /getmproj/([0-9a-f]{32}).png to get the max projection as a PNG image from the web service. This has already been constructed in the previous step, so this route serves a static file.

# Chapter 5

# **Conclusions and discussion**

### 5.1 Conclusions

In this thesis it has been researched how well the reconstruction of a sizable set of large OPT images can be distributed on the LLSC. Experiments have shown that distribution definitely yields significant and scalable speedups. The gains of distributing the reconstruction in general do reduce when a lot of processes are allocated, though, as then relatively a lot of time is spent doing I/O-related tasks rather than performing the computationally expensive inverse radon transform.

Some optimizations to the implementation were developed, ranging from improving the execution speed of the computationally intensive inverse radon transform, which has a lot positive effect, to optimizations attempting to improve I/O and IPC times. These optimizations typically have less effect in general, but mostly compression and streamed TIFF write-back reduce the total execution time significantly.

A usable prototype GUI application has been developed to interface with a developed web service for submitting reconstruction jobs to the cluster and viewing max-projections of the input OPT images. It allows OPT system operators to, for example, quickly get an indication whether or not the input OPT images are of acceptable quality.

### 5.2 Future work

From this research, several suggestions for future research can be derived. The execution speed and parallelization of the slice reconstruction potentially still has a lot of room for improvement and further research. On the other hand, the prototype web service and corresponding GUI application also have some ways to go.

#### 5.2.1 Slice reconstruction

To improve the reconstruction speed of the slices, another range of different optimizations can be experimented with. Solutions for parallel reconstruction of a single slice exist, for example. Utilizing this technique to reconstruct a single slice on an entire node very fast can be compared with the performance of the algorithm in this thesis, where a single process uses one thread to reconstruct a smaller amount of slices. Different implementations of the inverse radon transform exist in general too, such as a reconstruction based on the Fourier Slice Theorem instead of a reconstruction based on filtered backprojection as used in this thesis [Tof96]. These implementations may improve sequential execution speed too.

An entirely different approach can be attempted too, such as using one or multiple GPUs to reconstruct slices in parallel. Many implementations for filtered backprojections have already been developed and compared for CUDA [SKKH07, PZM11]. These implementations significantly differ from a implementation for CPUs, as used in this project. These GPU implementations can potentially be adapted to perform better with a large amount of sinograms that have to be reconstructed. It could be interesting to research whether or not using GPUs for slice reconstruction reduces the total execution time and if so, by which factor. I/O times could minimize potential gains significantly, as observed in the experiment in Section 3.5.

As shown in Section 3.4, parallel compression and streamed TIFF write-back was an optimization that had most positive influence on the total execution speed of the reconstruction. The performance gains can likely be improved even more. Currently, the compressed data is first sent to a specific master node that also has to compress its own data itself. An extra worker process could be introduced living at the file server, to which the compressed data is immediately sent. This will eliminate the extra data transfer from all worker nodes to the master node and likely improves performance in general as this process does not have to compress data itself like other processes living on compute nodes.

Instead of handling the streamed write-back of the TIFF file manually though Irecv and Rsend calls, direct parallel file writes to the file server can be used too. Instead of allocating large buffers for the output data in the master process's memory, a large file can be allocated in which enough space is available for each process to write their compressed data to, at a pre-determined offset in this file. This obviously results in a very large file with large parts of the file being unused. Assuming either a Ext4 or a XFS filesystem is used and that the file server runs Linux kernel 3.15 or newer, the fallocate Linux-function can be used with the FALLOC\_FL\_COLLAPSE\_RANGE flag to deallocate these unused preallocated file blocks, which then reduces the file size.

The compression level is also something to consider. Increasing the compression level will reduce the data size and thus reduce the amount of I/O time, but increases the amount of processing power needed to actually perform the compression. The experiment in Section 3.4 has shown that with only a few processes,

the used compression level increases the total execution time. This balance is something that has yet to be researched and tuned if execution speed is to be maximized.

Finally, because the cluster is not homogeneous, splitting the workload such that each node receives about an equal amount of slices to reconstruct may not be optimal for the total execution time. Some processors are significantly faster than others in the cluster. Allocating slices to processes while taking core speed into account can improve performance even more, especially when many slices are to be reconstructed.

In Section 2.3.1 three alternatives are discussed for constructing a range of sinograms in each of the processes memory. A limitation of Python modules that read the TIFF files is that they try to serve a general purpose and full pages at a time, not a select range of scanlines, for performance purposes. A Python-module written in C can be developed that does something similarly to the first considered variant, in which each process reads the entire input file and only extracts the rows of pixels it needs. Instead of reading all rows of pixels, it can skip past scanlines in the input data that worker process does not need at all. This way, less time is spent reading the file, which may result in less total execution time required for constructing the sinograms. Each process still seeks through the entire file, but less file reads are issued in total compared to the tested variant, which could make this optimization effective.

#### 5.2.2 Application

Further work on the GUI application and web service can be done too. Uploading data from the application to the cluster is fairly slow as it needs to be sent over the internet. Compression on the data stream can improve the uploading times significantly.

The max-projection is currently computed after the entire reconstruction is written back to an output file on the file server. It first needs to read this entire file, then compute the max-projection sequentially and then write back the output max-projection. The distributed slice reconstruction can be extended to also compute the max-projection in parallel. A local max-projection of the locally reconstructed slices can be computed at each worker node instead of collecting the reconstructed slices into a TIFF file. These locally reconstructed slices can then rapidly be reduced to a single global max-projection, which can be saved and sent to the GUI application. This modification can improve construction of the max-projection significantly.

Currently, the application only shows the max-projection. Alternative views such as the min-projection can be introduced as well.

The web framework used heavily utilizes coroutines to serve multiple requests at the same time. The controllers written for the web service do not properly yield during long operations, such as computing the MD5 hash or writing back the input file to the file system and thus lock the entire web server up during these operations. Converting these procedures to proper asynchronous operations will enable the server to sustain multiple requests at once better.

The web service currently uses an in-memory buffer to store partial uploads in. When the upload finishes, it is all written back at once to the file server which can take about 10 seconds. Writing partially uploaded data back to a file on the file server as it arrives can improve total execution time of such a reconstruction request. Besides, with many ongoing uploads, the web server can run out of memory using the current implementation.

Reconstruction jobs are currently scheduled by the web server in a very crude way. It tries to schedule the reconstruction job on a fixed amount of nodes. Under load or with few active nodes, this may result in the job not scheduling in a timely manner. Some metrics can be used to schedule these jobs more cleverly, such as the amount of nodes currently available. Prioritizing these kind of reconstructions over other batch jobs can be implemented too. Allocating too few nodes to these reconstruction jobs when more nodes may become available in the near future may result in the reconstruction finishing significantly later than it could finish. Research can be performed to schedule these jobs more efficiently.

Currently, the application processes a single scan (one TIFF file with all angles) at a time. Multiple kinds of scans can be made of the same specimen. These scans belong together and can be visualized as such too, which a future iteration of the application could integrate. Background masks are often generated too. An additional preprocessing step could be applied to subtract the background to reduce noise in the output.

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