03 where is my reader tmp

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1 Exercise 03 - Where is my reader?

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Authored by:

[1]: import warnings

#

#

#

1.0.1 2D laboratory micro-CT, fan-beam with no reader

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We don't have readers for every system so you might need to read in your data and create a geometry by hand.

This exercise walks through the steps needed to load in a 2D fan-beam sinogram stored as a single tiff image; create the matching geometry and finally reconstruct it using FDK. The data was acquired on a Nikon system but here we only read the centre slice and do not provide a reader for xtek2dCT files.

Learning objectives are: - Load and investigate a dataset stored as a tiff using TIFFStackReader - Create the CIL geometry using AcquisitionGeometry.Create_Cone2D - Combine the data and geometry to form an AcquisitionData - Find the Centre of rotation to complete the data geometry description - Compute the reconstruction using CIL's FDK

The sample is a plastic tube with a diameter of 25mm filled with uniform Soda-Lime Glass (SiO2-Na2O) beads of diameters 2.5mm (with standard deviation 0.1mm). This example requires the dataset SparseBeads_B12_L1.zip from https://zenodo.org/record/290117:

• https://zenodo.org/record/290117/files/SparseBeads_B12_L1.zip

If running locally please download the data and update the filepath in the filename variable below:

```
[3]: filename = "/mnt/materials/CIL/SparseBeads_B12_L1/CentreSlice/Sinograms"
```

```
[4]: from cil.io import TIFFStackReader
from cil.framework import AcquisitionGeometry, AcquisitionData
from cil.processors import CentreOfRotationCorrector
from cil.recon import FDK
from cil.utilities.display import show2D, show_geometry
import numpy as np
import matplotlib.pyplot as plt
```

2 Read in your data

2.1 Exercise A: Tiff to numpy array

- 1. Load the 2D fan-beam sinogram, using the TIFFStackReader
- 2. Use show2D to visualise your data
- 3. Look at the data type and shape

[]:

If you are stuck uncomment the following line and run the cell to see the solution, to run the lines you'll need to run the cell a second time

```
[5]: # create the TIFF reader by passing the directory containing the files
reader = TIFFStackReader(file_name=filename, dtype=np.float32)

# read in file, and return a numpy array containing the data
data_original = reader.read()

# use show2D to visualise the sinogram
show2D(data_original)
```

```
Exception Traceback (most recent call last)

Cell In[5], line 2

1 # create the TIFF reader by passing the directory containing the files

----> 2 reader = TIFFStackReader(file_name=filename, dtype=np.float32)

4 # read in file, and return a numpy array containing the data

5 data_original = reader.read()
```

```
File ~/miniconda3/envs/cil_test_demos/lib/python3.12/site-packages/cil/io/TIFF.
  →py:269, in TIFFStackReader.__init__(self, file_name, roi, transpose, mode, ⊔
  ⇔dtype)
    266 self.file name = file name
    268 if self.file_name is not None:
             self.set_up(file_name = self.file_name,
 --> 269
     270
                         roi = roi,
     271
                         transpose = transpose,
                         mode = mode, dtype=dtype)
     272
File ~/miniconda3/envs/cil_test_demos/lib/python3.12/site-packages/cil/io/TIFF.
  →py:366, in TIFFStackReader.set_up(self, file_name, roi, transpose, mode, dtyr;)
                 raise Exception("No tiff files were found in the directory \n{} .
     363
  →format(file_name))
     365 else:
 --> 366
             raise Exception("file_name expects a tiff file, a list of tiffs, or

¬a directory containing tiffs.\n{}".format(file name))

     369 for fn in self._tiff_files:
     370
             if '.tif' in fn:
Exception: file name expects a tiff file, a list of tiffs, or a directory
  ⇔containing tiffs.
/mnt/materials/CIL/SparseBeads_B12_L1/CentreSlice/Sinograms
```

```
[]: print("Array stored as:",type(data_original))
    print("Array shape:\t",data_original.shape)
    print("Array contents:\t",data_original.dtype)
```

3 Understand your data

Let us look at a histogram of our data.

```
[]: plt.hist(data_original.ravel(), bins=256)
plt.show()
```

We should notice that the data is X-ray transmission data. We can see that the background peak has the highest value.

The background peak is not at 1. When the data was saved as a tiff it was pre-scaled by 60000 by the scanner and stored as unsigned short which have integer values 0 - 65535. We can see the white level of 60000 in the xtek2dct file so we use this to normalise the data. If you would like to view the xtek2dct open a terminal and type the command cat /mnt/materials/CIL/SparseBeads_B12_L1/CentreSlice/SparseBeads_B12_L1.xtek2dct

```
[]: data_normalised = data_original / 60000
plt.hist(data_normalised.ravel(), bins=256)
plt.show()
```

And now we use Beer-Lambert's law to convert from X-ray transmission data to X-ray absorption data

```
[]: data_absorption = -np.log(data_normalised)
  plt.hist(data_absorption.ravel(), bins=256)
  plt.show()
```

You may notice the background has a small negative attenuation. This is unphysical in an ideal system however it is often caused by systematic errors in the scan. Potentially scatter caused by the object can raise the background value in the background of the scan but is not present in the flat-field image. Or latent image in the scintillator fades between the flat-field collection and the scan. We must be careful not to clip these negative values and bias the reconstruction.

4 Create the geometry

As well as the data itself, we need to create the geometric metadata as an AcquisitionGeometry object.

We start by populating the information we know from the data. For parallel beam data this might be sufficient.

4.1 Exercise B: Use the data

From the data set, extract the number of projections, the number of pixels and the order of the 'horizontal' and 'angle' axes:

```
[]:
```

Uncomment the following line and run the cell to see the solution, to run the lines you'll need to run the cell a second time:

4.2 Exercise C: Create the geometry

Now we create the CIL AcquisitionGeometry object using the method $AcquisitionGeometry.create_Cone2D$

You will have to set the position of the source, rotation axis (object) and detector. Look at the documentation of create_Cone2D as well as the notebook 00_CIL_geometry to help you.

1. Create a fan-beam geometry using the AcquisitionGeometry.create_Cone2D to define how our data was collected. We need to make sure each pixel is mapped to the correct angle of

measurement and voxel in the reconstructed image volume. Note that in CIL, our default system has the rotation axis aligned with the Z axis, the detector rows (Dx) are in the direction of the X-axis, the detector columns (Dy) are in the direction of the Z axis, and the source to object, and source to detector distances are defined along the y axis. E.g. see this illustration of a 3D cone beam geometry, with the axes shown:

You can deviate from the defaults and define the source and the detector along other axes, but you'll need to update the detector_direction_x too. Therefore we recommend using our defaults as much as possible.

- 2. Configure the angles of the data with set_angles
- 3. Configure the number and size of the detector pixels with set_panel
- 4. Configure the order of the data axes using set_labels
- 5. Use the show_geometry method to display the scan set up visually.

Here we give you the information from the scanner metadata, you can also find these values in the xtek2dct. If you would like to view the xtek2dct open a terminal and type the command cat/mnt/materials/CIL/SparseBeads_B12_L1/CentreSlice/SparseBeads_B12_L1.xtek2dct

Have a go at creating the geometry and see how it looks when you reconstruct the data with it. It might take a few attempts to get it right but if you are struggling you can uncomment the snippet and load an example solution.

```
[]: #What we know from the scanner
source_to_detector_distance = 1400.207
source_to_object_distance = 121.932
pixel_size = 0.2
angles = np.linspace(start=0, stop=360, num=number_of_projections,u
endpoint=False)
```

[]:

Uncomment the following line and run the cell to see the solution, to run the lines you'll need to run the cell a second time:

```
# set the angles, remembering to specify the units
geometry.set_angles(angles, angle_unit='degree')

# set the detector shape and size
geometry.set_panel(number_of_pixels, pixel_size)

# set the order of the data
geometry.set_labels(axis_labels)

# display your geometry, does it look like a feasible CT scan set up?
show_geometry(geometry)
```

5 Create the AcquisitionData

Now we have our data data_absorption stored as a numpy array and geometry containing the description of our data, we can use both to create an AcquisitionData that can be used by CIL.

Here we make a copy of the original data with deep_copy=True. For a large data set you may want this to use your existing numpy array with deep_copy=False

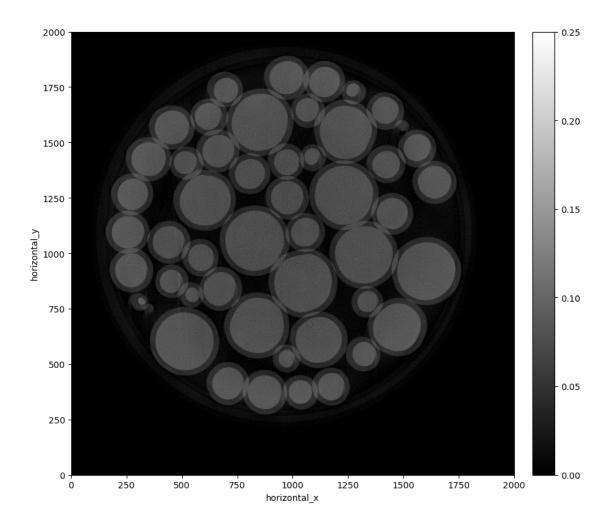
```
[]: acquisition_data = AcquisitionData(data_absorption, deep_copy=True, use geometry=geometry)
```

6 Reconstruct with FDK

Now we will try to reconstruct your AcquisitionData using CIL's recon.FDK. This uses CIL to filter the projections, and then calls the backprojector from TIGRE.

```
[]: reconC = FDK(acquisition_data).run()
[]: show2D(reconC, fix_range=(0,0.25))
```

If your geometry was right, you will now have a reconstruction that looks like this. If something went wrong, then go back to **Exercise C** and have another go at setting up your geometry.



7 Centre of rotation offset

We are nearly there but the double edges show our geometry is still not quite right. These are typical centre of rotation artefacts for a 360degree scan. We can update the geometry by hand until we find the right value.

7.0.1 Exercise D: Fix the centre of rotation offset

Use set_centre_of_rotation on your geometry to update the value and reconstruct your data. Try values between -5 and 5 until you obtain a nice-looking reconstruction.

```
[]: # set the offset between -5 and 5
offset = 5

#apply it to your geometry
acquisition_data.geometry.
⇒set_centre_of_rotation(offset,distance_units='default')

#reconstruct your data with the updated geometry
```

```
reconD = FDK(acquisition_data).run(verbose=0)
show2D(reconD, fix_range=(0,0.25))
```

Often we can use CIL's CentreOfRotationCorrector algorithms to find the centre of rotation offset to subpixel accuracy.

```
[]: processor = CentreOfRotationCorrector.image_sharpness()
    processor.set_input(acquisition_data)
    centred_data = processor.get_output()

recon_centred = FDK(centred_data).run(verbose=0)
    show2D(recon_centred, fix_range=(0,0.25))
```

```
[]: print(centred_data.geometry.get_centre_of_rotation())
```

Some common mistakes when you create your geometry by hand include: - The rotation direction. This might not simply be how the turntable looks to turn (if you can see it!), the detector origin may be interpreted differently and appear to flip your data. - The definition of 0 degree. Look at $show_geometry$ you can see that as default CIL defines 0 degree in the positive Y direction. You might need to add 180 degrees to your angles, or swap the detector and source positions along the y axis. - The pixel size - CIL expects your pixel size in the same units as you defined your system geometry. In this example we used mm. This means our reconstruction has values of mm^{-1}

7.1 Exercise E: Try the common mistakes

Go back to **Exercise C** and try setting the geometry up wrong to see how the reconstruction looks.

- What if the rotation is in the opposite direction? Set the angles to -angles
- What if there's a 180 degree offset in your angles?
- What if you define your angles in the wrong units? Set the units to radian
- What if your pixel size is wrong? How does the cone angle affect the reconstruction?
- CIL's geometry assumes the centre ray of the beam in the positive Y direction. You can configure this along the X axis instead but you will need to update the direction of the detector and redefine your angles accordingly.