

From_CT_acquisition_to_CAD_models

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```
[1]: from IPython.display import display
      from IPython.display import Image
```

1 Use of fast realistic simulations on GPU to extract CAD models from microtomographic data in the presence of strong CT artefacts:

1.1 Registration of Tungsten Fibres on XCT Images

1.1.1 by Franck P. Vidal, Iwan T. Mitchell and Jean-Michel Létang

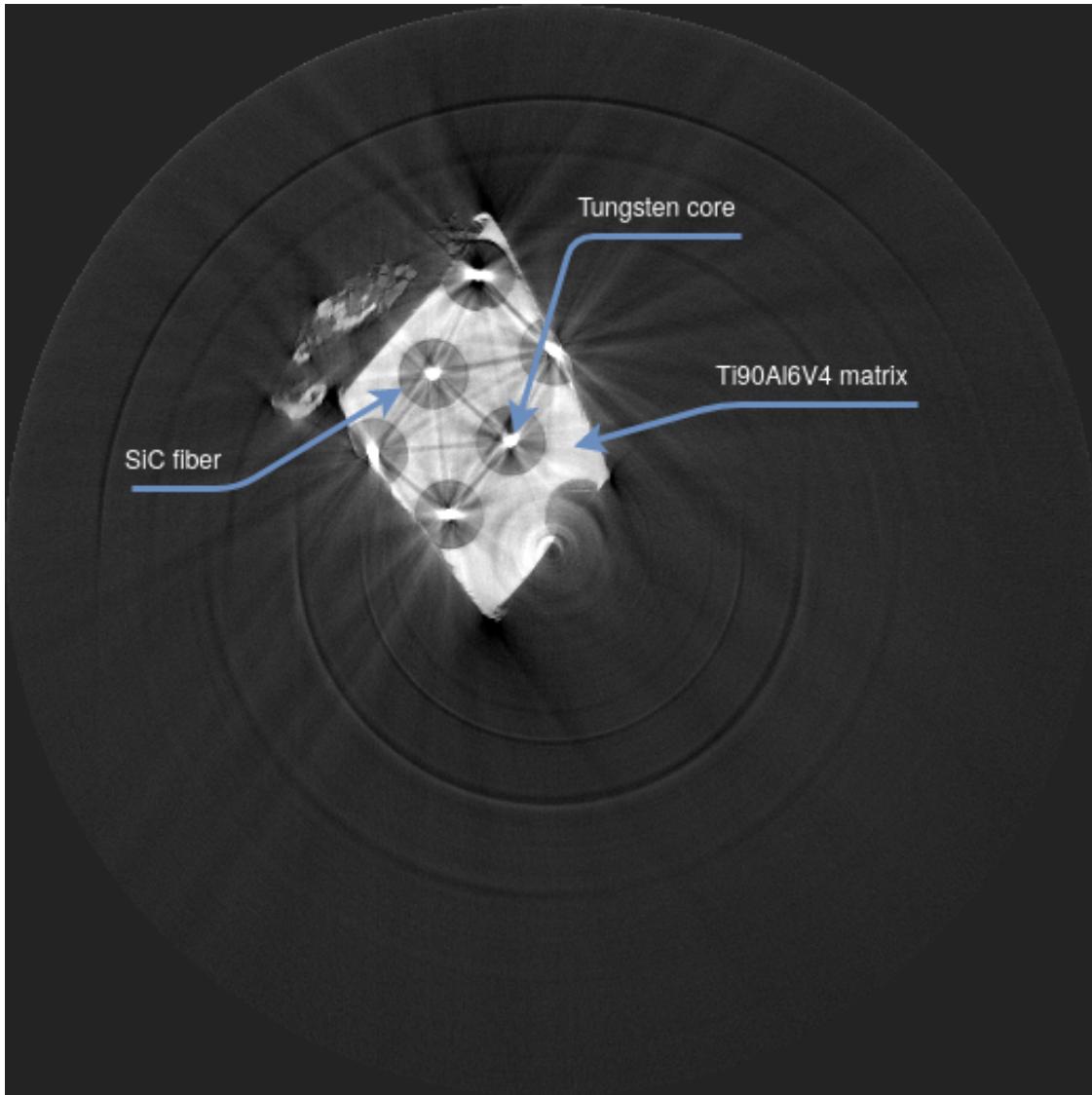
This demo aims to demonstrate how to register polygon meshes onto X-ray microtomography (micro-CT) scans of a tungsten fibre. The code relies on two building blocks:

1. A global optimisation algorithm. We use the [CMA-ES \(Covariance Matrix Adaptation Evolution Strategy\)](#). It is an evolutionary algorithm for difficult non-linear non-convex optimisation problems.
2. A fast X-ray simulation toolkit. We use [gVirtualXRay](#). It is a framework supporting many modern programming languages to generate realistic X-ray images from polygon meshes (triangles or tetrahedrons) on the graphics processor unit (GPU).

Below is an example of CT slice from an experiment we carried out at the [European Synchrotron Radiation Facility \(ESRF\)](#).

```
[2]: Image(filename=".//scanned_object.png")
```

[2]:



In a previous article, on *Investigation of artefact sources in synchrotron microtomography via virtual X-ray imaging* in Nuclear Instruments and Methods in Physics Research Section B: Beam Interactions with Materials and Atoms, we demonstrated that the image above was corrupted by:

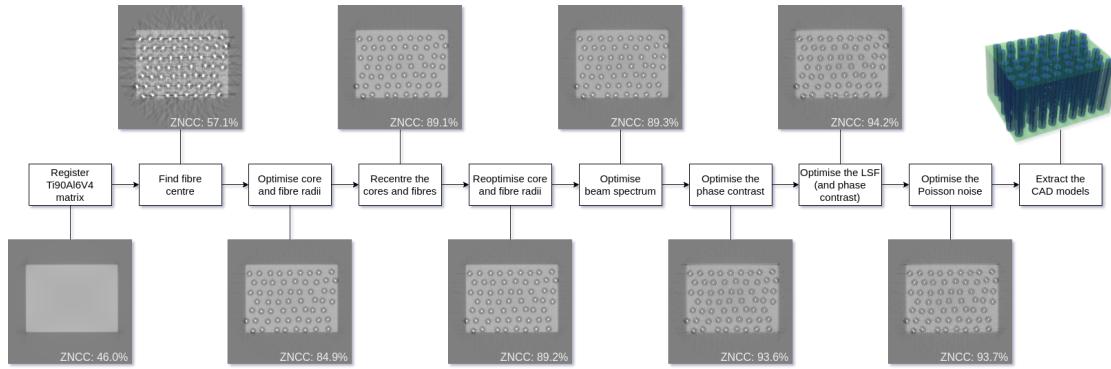
- 1) beam hardening despite the use of a monochromator,
- 2) the response of the camera despite the point spread function (PSF) being almost a Dirac, and
- 3) phase contrast.

That study was published in 2005, when computer were still relatively slow. Since then, massively parallel processors such as graphics processor units (GPUs) have emerged. Using today's hardware, we will demonstrate that we can now finely tuned the virtual experiments by mathematical optimisation to register polygons meshes on XCT data. Our simulations will include beam-hardening due to polychromatism, take into account the response of the detector, and have phase contrast.

1.2 Registration steps

```
[3]: Image(filename=".pipeline.png")
```

[3] :



1. Initialisation

- Import Python packages
- Global variables with values corresponding to known parameters
- Load the image data from the experiment at ESRF
- Reconstruct the corresponding CT data
- Normalise the image data
- Set the X-ray simulation environment
- LSF
- Find circles to identify the centre of fibres

2. Simulate the CT acquisition

3. Registration of the Ti90Al6V4 matrix

4. Optimisation of the cores and fibres radii

5. Recentre each core/fibre

6. Optimisation of the radii after recentring

7. Optimisation of the beam spectrum

8. Optimisation of the phase contrast and the radii

9. Optimisation of the phase contrast and the LSF

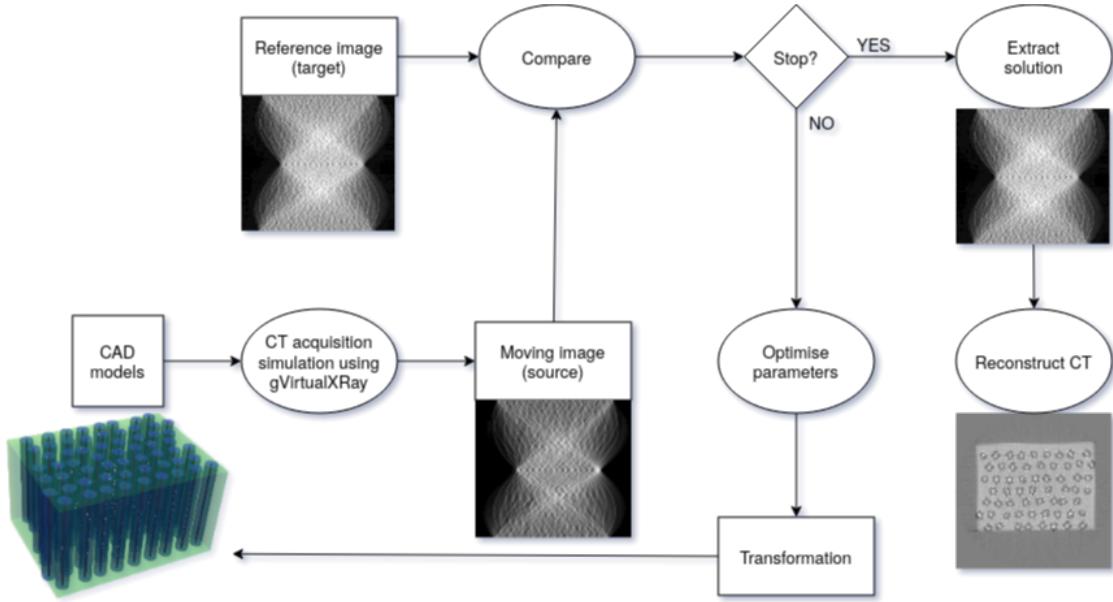
10. Optimisation of the Poisson noise

11. Results in terms of linear attenuation coefficients

In image registration, a *moving object* is geometrically deformed so that its image matches a *target image*. The parameters of the deformation is controlled and iteratively tuned by an optimisation algorithm.

```
[4]: Image(filename=".registration.png")
```

[4] :



In our context, the target is the sinogram provided by the experiment at ESRF. The moving image is created by simulation using the CAD models and [gVirtualXRay](#). The simulation parameters controlling the CAD models are repetitively tuned by a global optimisation algorithm until a stopping criterion is met. The optimisation algorithm will minimise (or maximise) a numerical value, the *objective function*. The comparison between the target and moving images measures how different (or similar) the two images are. It is performed within the objective function.

1.3 Import packages

We need to import a few libraries (called packages in Python). We use:

- `copy`: duplicating images using deepcopies;
- `glob`: retrieving file names in a directory;
- `math`: the `floor` function;
- `os`: creating a new directory;
- `sys`: retrieving the largest possible floating-point value;
- `base64`: to save a PNG screenshot of the 3D visualisation;
- `cma`: non-linear numerical optimization ([CMA-ES](#), [Covariance Matrix Adaptation Evolution Strategy](#));
- ([OpenCV](#)) (`cv2`): Hough transform and bilateral filter (an edge-preserving smoothing filter);
- `imageio`: creating GIF files;
- `IPython.display`: display Pandas'dataframes as HTML tables;
- `matplotlib`: plotting data;
- `mpl_toolkits`: plotting 3D data (final CAD models);
- `numpy`: who doesn't use numpy?
- `pandas`: creating a DataFrame to store μ data;
- `SimpleITK`: image processing and saving volume data;

- `tomopy`: package for CT reconstruction;
- `scipy`: for the convolution of a 2D image by a 1D kernel;
- `skimage`: comparing the reference CT slice and the simulated one;
- `sklearn`: comparing the reference CT slice and the simulated one;
- `stl`: to import STL files (CAD models);
- `k3d`: to visualise the STL files;
- `lsf`: the line spread function to filter the X-ray images; and
- `gvxrPython3`: `gVirtualXRay`'s Python wrapper to simulate X-ray images using the Beer-Lambert law on GPU.

[5]: %matplotlib inline

```

import copy
import glob
import math
import os
import sys
import base64

import cma
import cv2
import imageio
import matplotlib.pyplot as plt
import numpy as np
import pandas as pd
import SimpleITK as sitk
import tomopy

from matplotlib import cm
from mpl_toolkits import mplot3d
from scipy import ndimage
from skimage.metrics import structural_similarity as ssim
from skimage.util import compare_images
from sklearn.metrics import mean_absolute_error, mean_squared_error

from stl import mesh
import k3d

# plt.ioff()
plt.rcParams['figure.figsize'] = [12, 8]
plt.rcParams['figure.dpi'] = 100 # 200 e.g. is really fine, but slower
# plt.rcParams['font.size'] = 15

import gvxrPython3 as gvxr

from lsf import *

```

```
[6]: if not os.path.exists("outputs"):
    os.makedirs("outputs");

if not os.path.exists("plots"):
    os.makedirs("plots");
```

1.4 Global variables

We need some global variables:

- `NoneType`: the type of `None`;
- `pixel_spacing_in_micrometre`: the physical distance between the centre of two successive pixel;
- `pixel_spacing_in_mm`: the physical distance between the centre of two successive pixel;
- `number_of_projections`: the total number of angles in the sinogram;he total number of angles in the sinogram;
- `angular_span_in_degrees`: the angular span covered by the sinogram;
- `angular_step`: the angular step;
- `theta`: the rotation angles in degrees (vertical axis of the sinogram);
- `theta_rad`: the rotation angles in radians (vertical axis of the sinogram);
- `roi_length`: control the size of the ROI when displaying the central fibre;
- `value_range`: control the binning of the Laplacian kernel
- `num_samples`: control the binning of the Laplacian kernel
- `sigma_set`: spread of the Laplacian kernels
- `k_set`: weight of the Laplacian kernels
- `label_set`: label of the structures on which a Laplacian kernel is applied
- `bias`: control the bias of the Poisson noise
- `gain`: control the gain of the Poisson noise: control the bias of the Poisson noise
- `scale`: control the scale of the Poisson noise: control the bias of the Poisson noise
- `use_normalisation`: use or do not use zero-mean, unit-variance normalisation in the objective functions;
- `use_sinogram`: compute the objective functions on the sinogram or flat-field;
- `metrics_type`: type of image comparison used in the objective functions;
- `fibre_radius`: radius of the SiC fibres in um
- `core_radius`: radius of the W fibres in um

```
[7]: NoneType = type(None);
pixel_spacing_in_micrometre = 1.9;
pixel_spacing_in_mm = pixel_spacing_in_micrometre * 1e-3;
number_of_projections = 900;
angular_span_in_degrees = 180.0;
angular_step = angular_span_in_degrees / number_of_projections;
theta = np.linspace(0.,
                    angular_span_in_degrees,
                    number_of_projections,
                    endpoint=False);
theta_rad = theta / 180.0 * math.pi;
```

```

roi_length = 60;

value_range = 6;
num_samples = 15;

sigma_set = None;
k_set = None;
label_set = None;

bias = None;
gain = None;
scale = None;

use_normalisation = True;
use_sinogram = True;

metrics_type = "RMSE";

fibre_radius = 140 / 2; # um
core_radius = 30 / 2; # um

```

1.5 Load the image data

Load and display the reference projections from a raw binary file, i.e. the target of the registration.

```
[8]: # Target of the registration
reference_normalised_projections = np.fromfile("sino.raw", dtype=np.float32);
reference_normalised_projections.shape = [
    number_of_projections,
    int(reference_normalised_projections.shape[0] / number_of_projections)
];
```

We define a function to save raw images in the MHA format:

```
[9]: def saveMHA(fname, image, spacing):
    """
    save the image into a file.

    :param str fname: the filename
    :param 2D_image image: the image to save
    :param [flt, flt, flt] spacing: the space between two successive voxels ↴ along the 3 direction
    """
    volume = sitk.GetImageFromArray(image);
```

```

volume.SetSpacing(spacing);
sitk.WriteImage(volume, fname, useCompression=True);

```

The reference projections in a MHA file

```
[10]: saveMHA('outputs/reference_normalised_projections.mha',_
           ↪reference_normalised_projections, [pixel_spacing_in_mm, angular_step,_
           ↪pixel_spacing_in_mm]);
```

Display the reference projections using Matplotlib

In the literature, a projection is often modelled using the polychromatic version of the Beer-Lambert law:

$$\mathbf{I}(x, y) = \sum_i \mathbf{R}_i \mathbf{N}_i \exp \left(- \sum_j \mu_j(E_i) \mathbf{d}_j(x, y) \right)$$

- $\mathbf{I}(x, y)$ the value of the raw X-ray projection at pixel location (x, y) , and with the sample and with the X-ray beam turned on;
- i the i -th energy channel in the beam spectrum;
- E_i the energy in eV;
- \mathbf{R}_i and \mathbf{N}_i the detector response and the number of photons at that energy respectively;
- j the j -th material being scanned, $\mu_j(E_i)$ its linear attenuation coefficient at energy E_i , and
- $\mathbf{d}_j(x, y)$ path length in cm^{-1} of the ray crossing the j -th material from the X-ray source to pixel (x, y) .

Projections are then corrected to account for variations in beam homogeneity and in the pixel-to-pixel sensitivity of the detector. This is the projection with flat-field correction (**Proj**):

$$\mathbf{Proj} = \frac{\mathbf{I} - \mathbf{D}}{\mathbf{F} - \mathbf{D}}$$

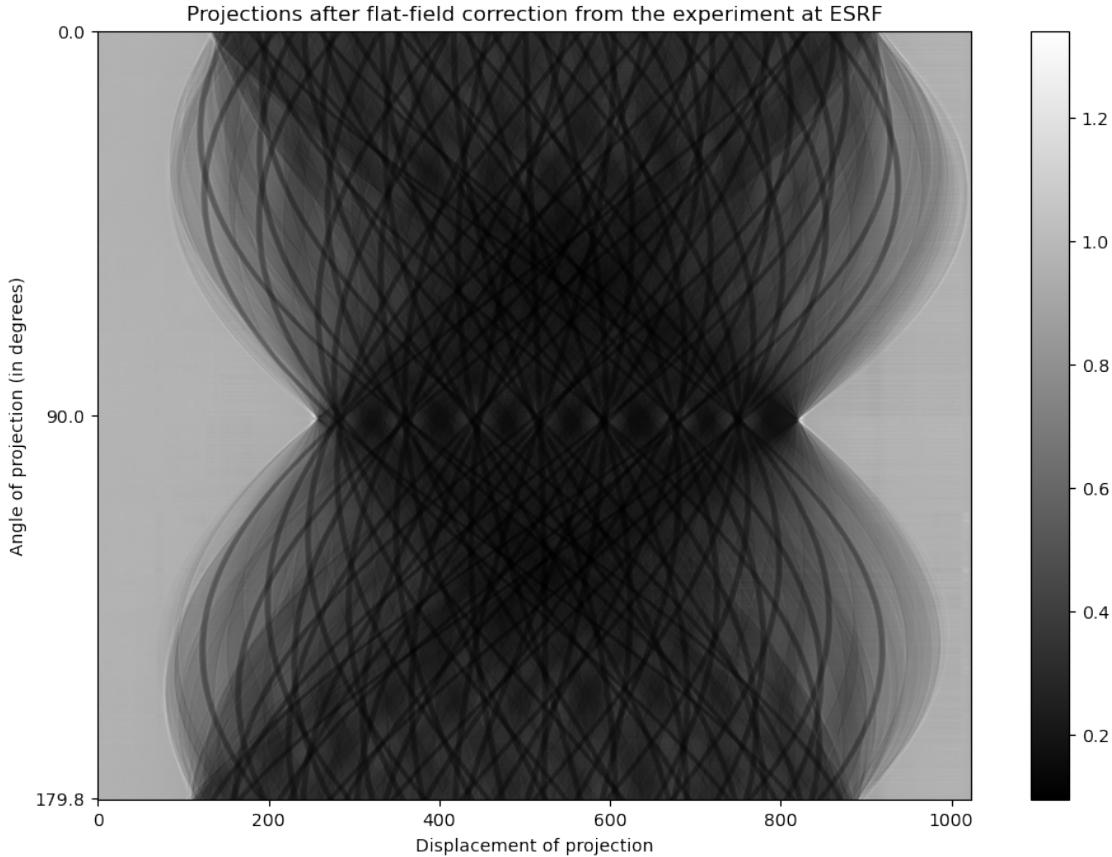
F (full fields) and **D** (dark fields) are projection images without sample and acquired with and without the X-ray beam turned on respectively. Note that with an ideal detector ($\mathbf{R}_i = E_i$), pixels of **D** are null, and pixels of **F** are equal to $\sum_i E_i \mathbf{N}_i$.

```
[11]: labels = [theta[0], theta[reference_normalised_projections.shape[0] // 2],_
            ↪theta[-1]];
tics = [
    0,
    reference_normalised_projections.shape[0] // 2,
    reference_normalised_projections.shape[0]-1
];
fig = plt.figure();
imgplot = plt.imshow(reference_normalised_projections, cmap="gray");
plt.xlabel("Displacement of projection");
plt.ylabel("Angle of projection (in degrees)");
plt.yticks(tics, labels);
plt.title("Projections after flat-field correction from the experiment at_
           ↪ESRF");
```

```

fig.colorbar(imgplot);
plt.savefig('plots/Normalised_projections_from_experiment_ESRF.pdf')
plt.savefig('plots/Normalised_projections_from_experiment_ESRF.png')

```



`reference_normalised_projections` (the figure above) corresponds to the data loaded from the binary file. It corresponds to **Proj**, i.e. the flat-field correction has already been performed.

We can see that when the primary spectrum is not monochromatic the measurement is the sum of several attenuation laws. We could however compute the effective monochromatic attenuation that would give the same measurement:

$$\mathbf{I}(x, y) = \mathbf{I}_0(x, y) \exp \left(- \sum_j \mu_j(E_{\text{eff}}) \mathbf{d}_j(x, y) \right)$$

with $\mathbf{I}_0(x, y) = \sum_i \mathbf{R}_i \mathbf{N}_i$, and where E_{eff} corresponds to the monochromatic energy that would give the same attenuation than the one measured. We are now able to linearise the transmission tomography data, namely **Proj**, and we get the sinogram:

$$\mathbf{Sino} = -\ln(\mathbf{Proj})$$

We define a new function to compute the sinogram from flat-field correction and calls it straight-away.

```
[12]: def computeSinogramFromFlatField(normalised_projections):
    """
    This function apply the minus log normalisation
    on the projections that have been corrected with the flat-field method.

    :param 2D_image normalised_projections: The projections after flat-field
    ↪corrections
    :return the sinogram.
    """

    # Create a temporary image to hold the sinogram
    simulated_sinogram = copy.deepcopy(normalised_projections);

    # Make sure no value is negative or null (because of the log function)
    # It should not be the case, however, when the Laplacian is used to simulate
    # phase contrast, negative values can be generated.
    threshold = 0.000001
    simulated_sinogram[simulated_sinogram < threshold] = threshold;

    # Apply the minus log normalisation
    simulated_sinogram = -np.log(simulated_sinogram);

    # Rescale the data taking into account the pixel size
    simulated_sinogram /= pixel_spacing_in_micrometre * gvxr.
    ↪getUnitOfLength("um") / gvxr.getUnitOfLength("cm");

    # Return the new image
    return simulated_sinogram;
```

Compute the sinogram from the flat-field data

```
[13]: reference_sinogram =_
    ↪computeSinogramFromFlatField(reference_normalised_projections);
```

Save the corresponding image

```
[14]: saveMHA('outputs/reference_sinogram.mha', reference_sinogram,_
    ↪[pixel_spacing_in_mm, angular_step, pixel_spacing_in_mm]);
```

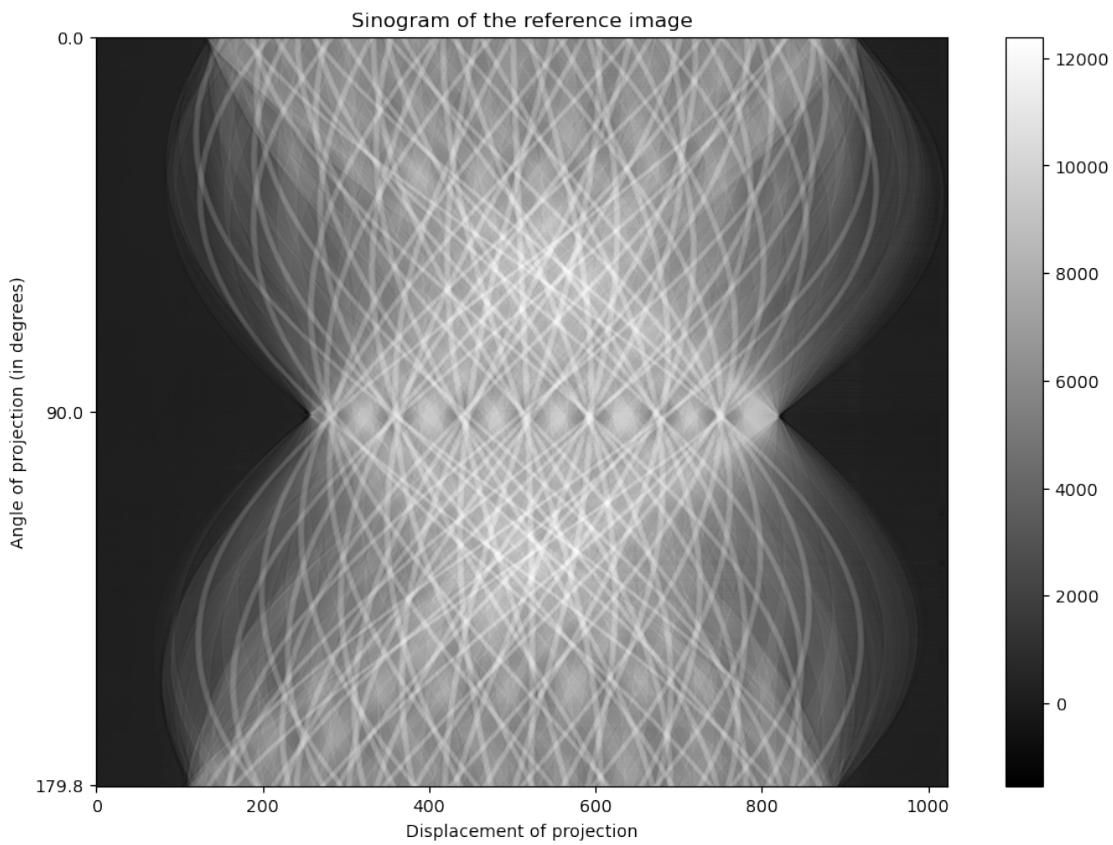
Display the sinogram using Matplotlib

```
[15]: labels=[theta[0], theta[reference_sinogram.shape[0] // 2], theta[-1]];
tics=[0, reference_sinogram.shape[0] // 2, reference_sinogram.shape[0]-1];
fig=plt.figure();
imgplot = plt.imshow(reference_sinogram, cmap="gray");
```

```

plt.xlabel("Displacement of projection");
plt.ylabel("Angle of projection (in degrees)");
plt.yticks(tics, labels);
plt.title("Sinogram of the reference image");
fig.colorbar(imgplot);
plt.savefig('plots/Sinogram_reference_image.pdf');
plt.savefig('plots/Sinogram_reference_image.png');

```



1.6 CT reconstruction

Now we got a sinogram, we can reconstruct the CT slice. As we used a synchrotron, we can assume we have a parallel source. It means we can use a FBP rather than the FDK algorithm. In fact we use the gridrec algorithm, which is much faster:

Dowd BA, Campbell GH, Marr RB, Nagarkar VV, Tipnis SV, Axe L, and Siddons DP. [Developments in synchrotron x-ray computed microtomography at the national synchrotron light source](#). In Proc. SPIE, volume 3772, 224–236. 1999.

```
[16]: reference_sinogram.shape = [
    reference_sinogram.shape[0],
```

```

    1,
    reference_sinogram.shape[1]
];

rot_center = int(reference_sinogram.shape[2]/2);

reference_CT = tomopy.recon(reference_sinogram,
                             theta_rad,
                             center=rot_center,
                             sinogram_order=False,
                             algorithm='gridrec',
                             filter_name='shepp')[0];

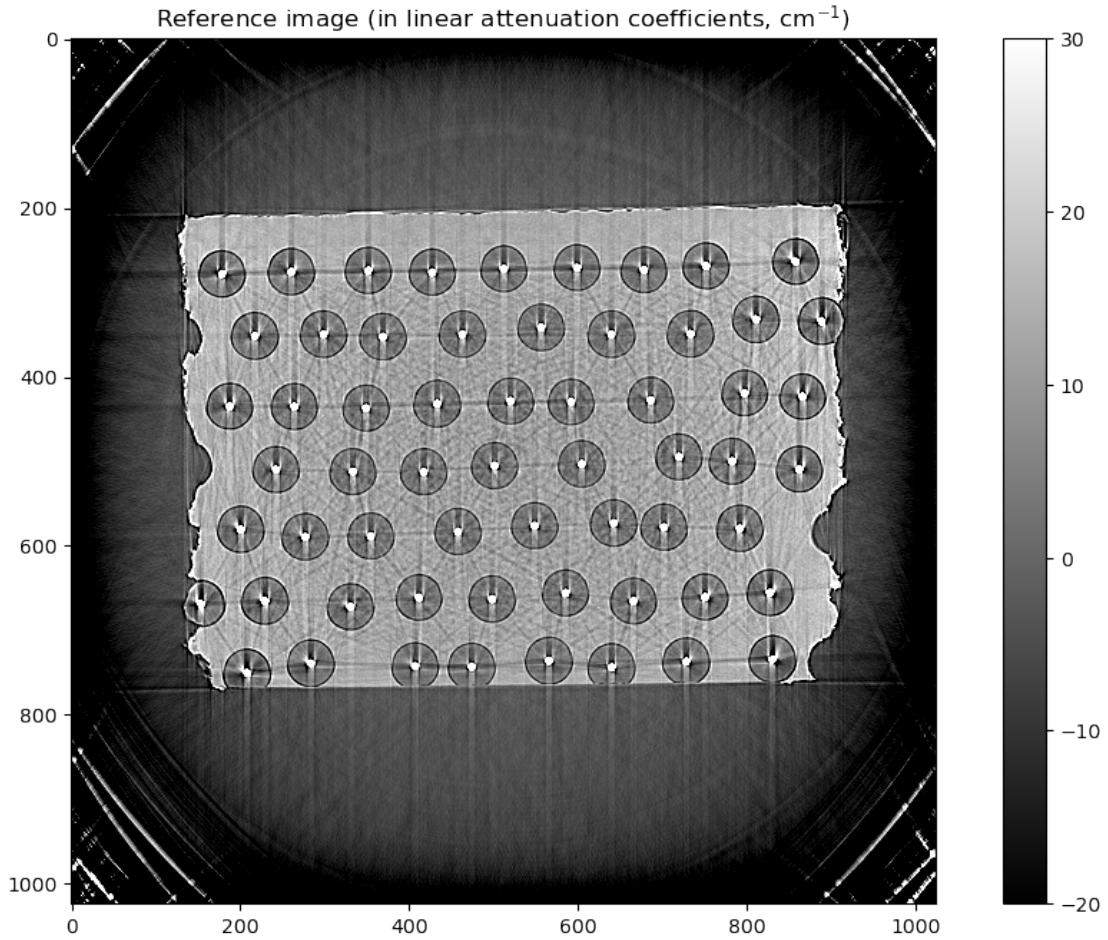
```

Save the reconstruction in a MHA file

```
[17]: saveMHA('outputs/reference_CT.mha', reference_CT, [pixel_spacing_in_mm, ↵
angular_step, pixel_spacing_in_mm]);
```

Plot the CT slice using Matplotlib

```
[18]: fig=plt.figure();
norm = cm.colors.Normalize(vmax=30, vmin=-20)
imgplot = plt.imshow(reference_CT, cmap="gray", norm=norm);
fig.colorbar(imgplot);
plt.title("Reference image (in linear attenuation coefficients, cm$^{-1}$)");
plt.savefig('plots/reference_image_in_mu.pdf');
plt.savefig('plots/reference_image_in_mu.png');
```



1.7 Normalise the image data

Zero-mean, unit-variance normalisation is applied to use the reference images in objective functions and perform the registration. Note that it is called standardisation (or Z-score Normalisation) in machine learning. It is computed as follows:

$$\mathbf{m}_o = \frac{\mathbf{m} - \bar{m}}{\sigma_m}$$

where \mathbf{m}_o is the image after normalisation of Image \mathbf{m} , \bar{m} is the average pixel value of Image \mathbf{m} , and σ_m its standard deviation. After normalisation, the average pixel value is null and the standard deviation of pixel values is equal to one.

We define a function to apply this:

```
[19]: def standardisation(I):
    image = copy.deepcopy(I);
```

```

# Sometimes the CT reconstruction algorithm create NaN on
# the top and right borders, we filter them out using
# a median filter ignoring NaN
nan_index = np.argwhere(np.isnan(image));
if nan_index.shape[0]:
    temp = np.pad(image, 1, "edge");

    for index in nan_index:
        roi = temp[index[0]-1+1:index[0]+1+2, index[1]-1+1:index[1]+1+2];
        image[index[0], index[1]] = np.nanmedian(roi);

return (image - image.mean()) / image.std();

```

Normalise the reference sinogram and CT slice

```
[20]: normalised_reference_sinogram = standardisation(reference_sinogram);
normalised_reference_CT      = standardisation(reference_CT);
```

1.8 Set the X-ray simulation environment

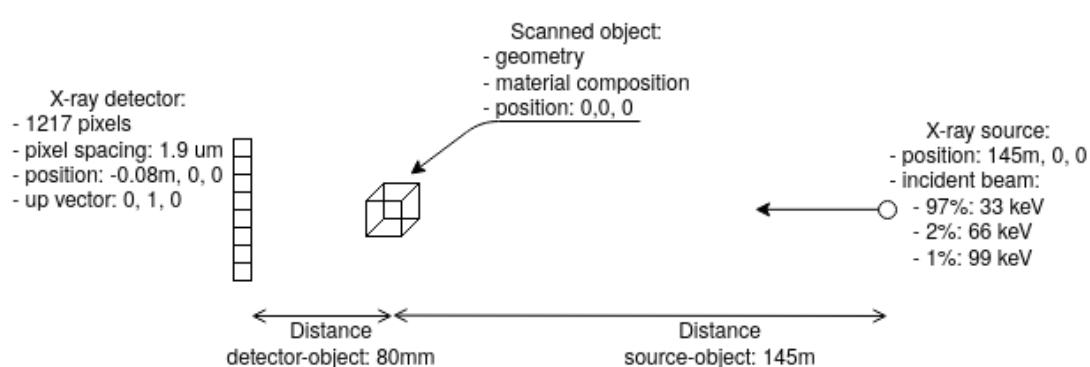
First we create an OpenGL context, here using EGL, i.e. no window.

```
[21]: gvxr.createWindow(0, 1, "EGL");
gvxr.setWindowSize(512, 512);
```

We set the parameters of the X-ray detector (flat pannel), e.g. number of pixels, pixel, spacing, position and orientation:

```
[22]: Image(filename=".\\3d_scene.png")
```

[22]:



```
[23]: detector_width_in_pixels = reference_sinogram.shape[2];
detector_height_in_pixels = 1;
distance_object_detector_in_m = 0.08; # = 80 mm
```

```

gvxr.setDetectorPosition(-distance_object_detector_in_m, 0.0, 0.0, "m");
gvxr.setDetectorUpVector(0, 1, 0);
gvxr.setDetectorNumberOfPixels(detector_width_in_pixels, □
    ↳detector_height_in_pixels);
gvxr.setDetectorPixelSize(pixel_spacing_in_micrometre, □
    ↳pixel_spacing_in_micrometre, "micrometer");

```

And the source parameters (beam shape, source position)

```
[24]: # Set up the beam
distance_source_detector_in_m = 145.0;

gvxr.setSourcePosition(distance_source_detector_in_m -□
    ↳distance_object_detector_in_m, 0.0, 0.0, "m");
gvxr.usePointSource();
# gvxr.useParallelBeam();
```

The beam spectrum. Here we have a polychromatic beam, with 97% of the photons at 33 keV, 2% at 66 keV and 1% at 99 keV.

```
[25]: energy_spectrum = [(33, 0.97, "keV"), (66, 0.02, "keV"), (99, 0.01, "keV")];

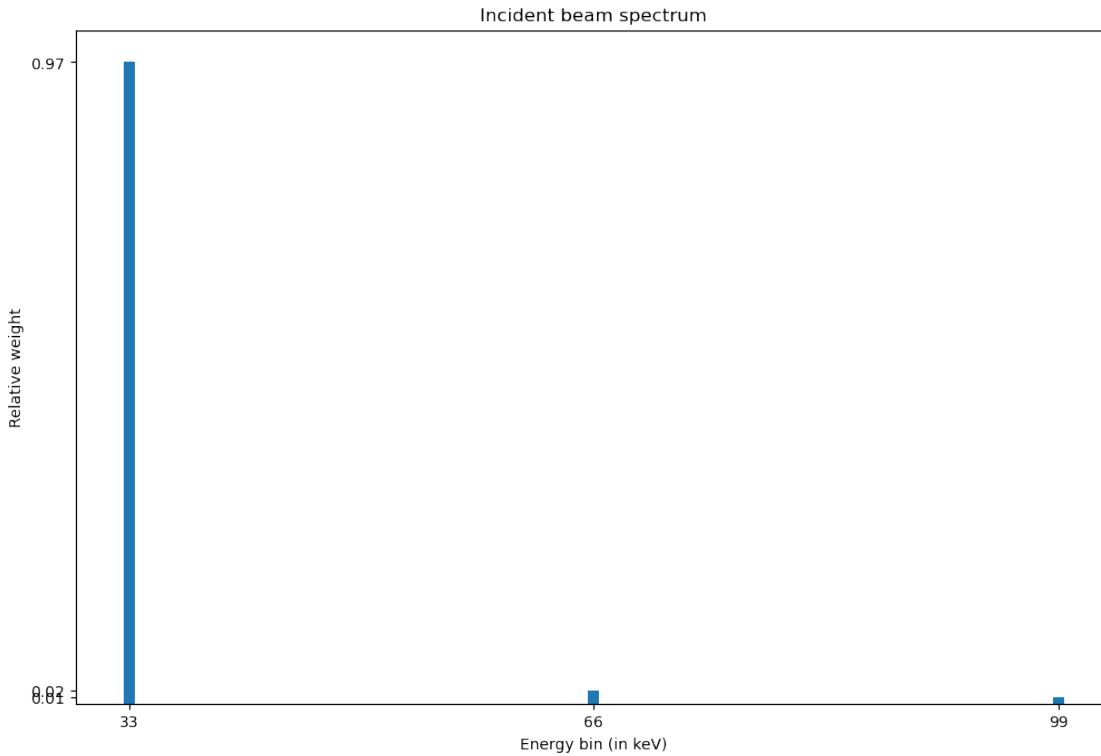
for energy, percentage, unit in energy_spectrum:
    gvxr.addEnergyBinToSpectrum(energy, unit, percentage);
```

Plot the beam spectrum using Matplotlib

```
[26]: energies_in_keV = [];
weights = [];

for energy, percentage, unit in energy_spectrum:
    weights.append(percentage);
    energies_in_keV.append(energy * gvxr.getUnitOfEnergy(unit) / gvxr.
    ↳getUnitOfEnergy("keV"));

fig=plt.figure();
plt.xlabel("Energy bin (in keV)");
plt.ylabel("Relative weight");
plt.xticks(energies_in_keV);
plt.yticks(weights);
plt.title("Incident beam spectrum");
plt.bar(energies_in_keV, weights);
plt.savefig('plots/beam_spectrum.pdf');
plt.savefig('plots/beam_spectrum.png');
```



The material properties (chemical composition and density)

```
[27]: fibre_material = [("Si", 0.5), ("C", 0.5)];
fibre_density = 3.2; # g/cm3

core_radius = 30 / 2; # um
core_material = [("W", 1)];

g_matrix_width = 0;
g_matrix_height = 0;
g_matrix_x = 0;
g_matrix_y = 0;
matrix_material = [("Ti", 0.9), ("Al", 0.06), ("V", 0.04)];
matrix_density = 4.42 # g/cm3
```

1.8.1 The LSF

In a previous study, we experimentally measured the impulse response of the detector as the line spread function (LSF):

F.P. Vidal, J.M. Létang, G. Peix, P. Cloetens, Investigation of artefact sources in synchrotron microtomography via virtual X-ray imaging, *Nuclear Instruments and Methods in Physics Research*

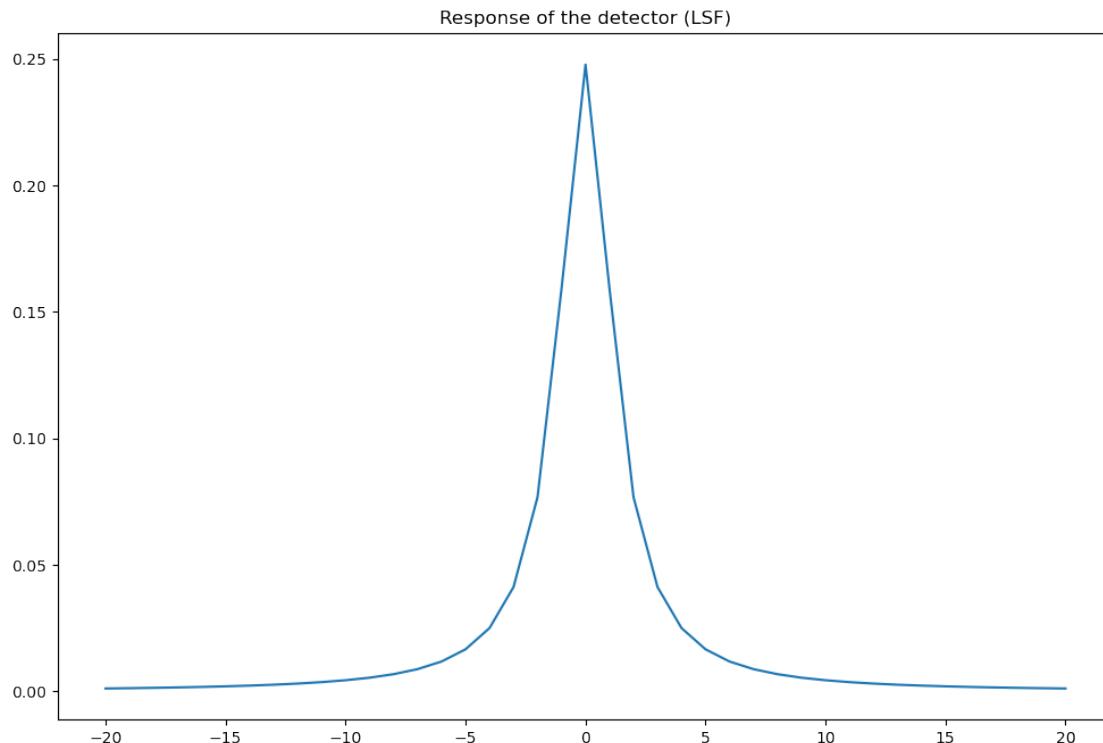
Section B: Beam Interactions with Materials and Atoms, Volume 234, Issue 3, 2005, Pages 333-348, ISSN 0168-583X, DOI [10.1016/j.nimb.2005.02.003](https://doi.org/10.1016/j.nimb.2005.02.003).

We use this model during the initial steps of the registration. The LSF model will be tuned in one of the final steps of the registration.

```
[28]: t = np.arange(-20., 21., 1.);  
lsf_kernel=lsf(t*41)/lsf(0);  
lsf_kernel/=lsf_kernel.sum();
```

Plot the LSF using Matplotlib

```
[29]: fig=plt.figure();  
plt.title("Response of the detector (LSF)");  
plt.plot(t, lsf_kernel);  
plt.savefig('plots/LSF.pdf');  
plt.savefig('plots/LSF.png');
```



1.9 Find circles to identify the centre of fibres

We can use the Hoguh transform to detect where circles are in the image. However, the input image in OpenCV's function must be in UINT. We blur it using a bilateral filter (an edge-preserving smoothing filter).

1.9.1 Convert the image to UINT

We first create a function to convert images in floating point numbers into UINT.

```
[30]: def float2uint8(anImage, min_threshold = None, max_threshold = None):  
  
    uchar_image = copy.deepcopy(anImage);  
  
    if isinstance(min_threshold, NoneType):  
        min_threshold = np.min(uchar_image);  
  
    if isinstance(max_threshold, NoneType):  
        max_threshold = np.max(uchar_image);  
  
    uchar_image[uchar_image < min_threshold] = min_threshold;  
    uchar_image[uchar_image > max_threshold] = max_threshold;  
  
    uchar_image -= min_threshold;  
    uchar_image /= max_threshold - min_threshold;  
    uchar_image *= 255;  
  
    return uchar_image.astype(np.uint8);
```

We blur the CT scan using a bilateral filter. It preserves edges.

```
[31]: uint8_reference_CT = float2uint8(reference_CT, 0, 300);  
blurred_reference_CT = cv2.bilateralFilter(uint8_reference_CT, 9, 75, 75);  
  
saveMHA('outputs/blurred_reference_CT.mha', blurred_reference_CT, ↴  
        [pixel_spacing_in_mm, angular_step, pixel_spacing_in_mm]);
```

1.9.2 Apply the Hough transform

As the fibres and the cores correspond to circles in the CT images, the obvious technique to try is the Hough Circle Transform (HCT). It is a feature extraction technique used in image analysis that can output a list of circles (centres and radii).

```
[32]: circles = cv2.HoughCircles(blurred_reference_CT, cv2.HOUGH_GRADIENT, 2, 80,  
                                param1=150, param2=5, minRadius=5, maxRadius=15);
```

1.9.3 Overlay the detected circles on the top of the image

```
[33]: cimg = cv2.cvtColor(float2uint8(reference_CT, 0, 50), cv2.COLOR_GRAY2BGR);  
circles = np.uint16(np.around(circles));  
  
for i in circles[0,:]:
```

```

# draw the outer circle
cv2.circle(cimg, (i[0], i[1]), i[2], (0, 255, 0), 2);

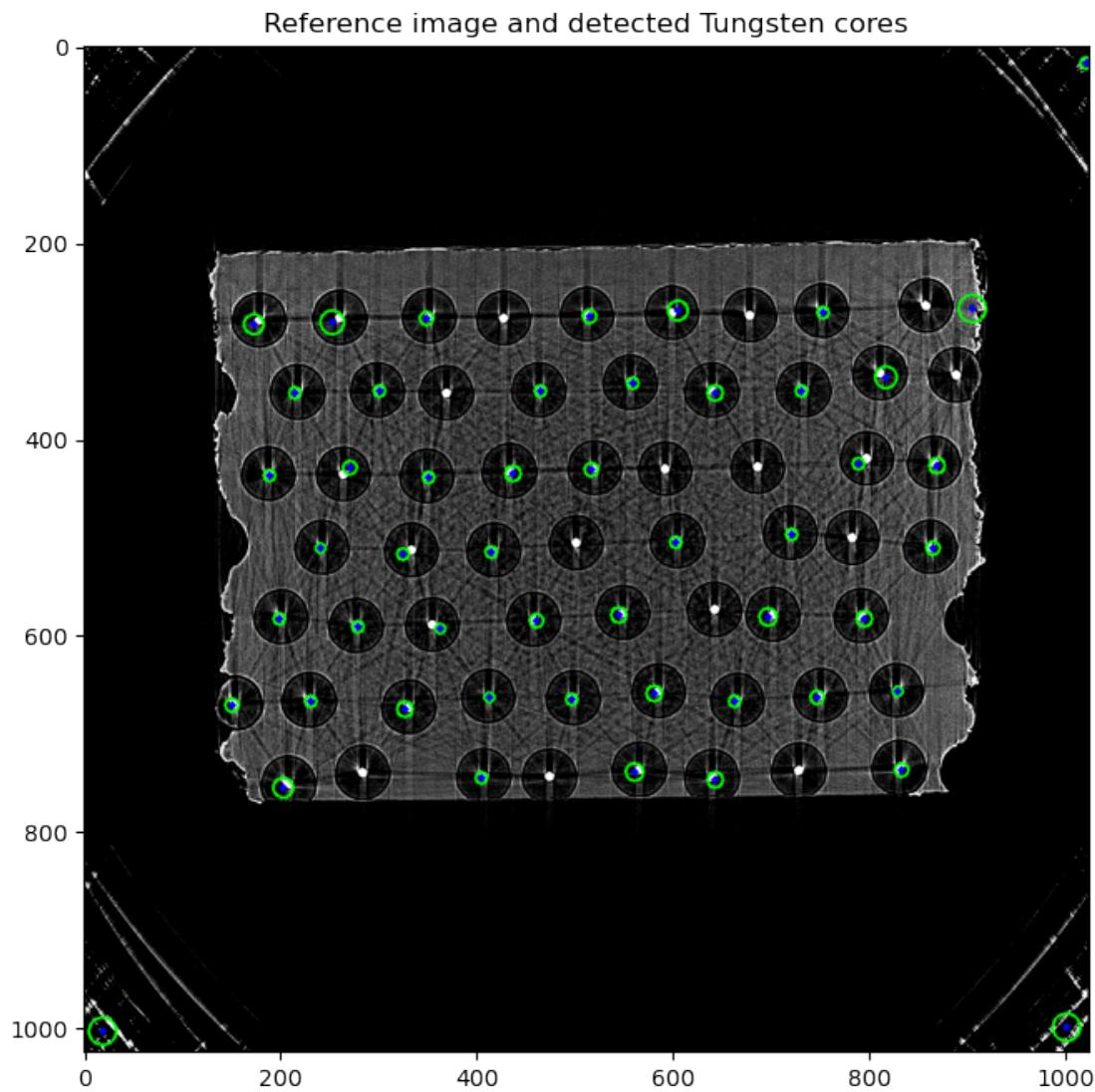
# draw the center of the circle
cv2.circle(cimg, (i[0], i[1]), 2, (0, 0, 255), 3);

```

```

[34]: fig=plt.figure();
imgplot = plt.imshow(cimg);
plt.title("Reference image and detected Tungsten cores");
plt.savefig('plots/fibre_detection_using_Hough_transform.pdf');
plt.savefig('plots/fibre_detection_using_Hough_transform.png');

```



13 fibres were missed and many centres were misplaced. Controlling the meta-parameters of the algorithm can be difficult to employ in a fully-automatic registration framework. We will use another technique to register the fibres, the popular Otsu's method. It creates a histogram and uses a heuristic to determine a threshold value.

```
[35]: # Convert the numpy array in float32 into uint, then into a SITK image
volume = sitk.GetImageFromArray(blurred_reference_CT);
volume.SetSpacing([pixel_spacing_in_mm, pixel_spacing_in_mm, ↵
    ↵pixel_spacing_in_mm]);

# Apply the Otsu's method
otsu_filter = sitk.OtsuThresholdImageFilter();
otsu_filter.SetInsideValue(0);
otsu_filter.SetOutsideValue(1);
seg = otsu_filter.Execute(volume);

# Print the corresponding threshold
print("Threshold:", otsu_filter.GetThreshold());
```

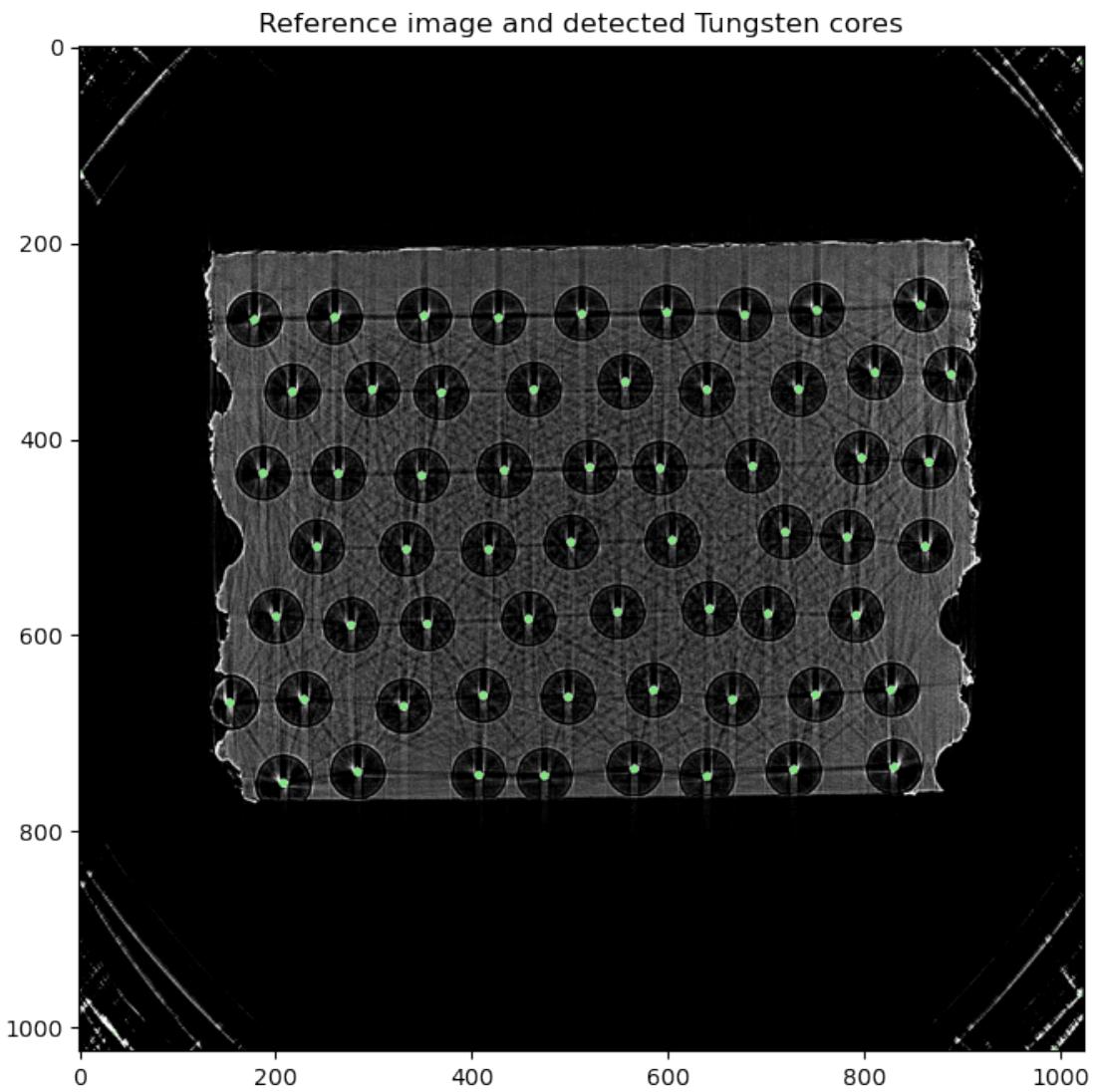
Threshold: 91.0

```
[36]: sitk.WriteImage(seg, "outputs/cores_segmentation.mha", useCompression=True);
```

```
[37]: fig = plt.figure();

volume = sitk.GetImageFromArray(float2uint8(reference_CT, 0, 50));
volume.SetSpacing([pixel_spacing_in_mm, pixel_spacing_in_mm, ↵
    ↵pixel_spacing_in_mm]);

imgplot = plt.imshow(sitk.GetArrayViewFromImage(sitk.LabelOverlay(volume, ↵
    ↵seg)));
plt.title("Reference image and detected Tungsten cores");
plt.savefig('plots/fibre_detection_using_otsu_method.pdf');
plt.savefig('plots/fibre_detection_using_otsu_method.png');
```



1.9.4 Clean up

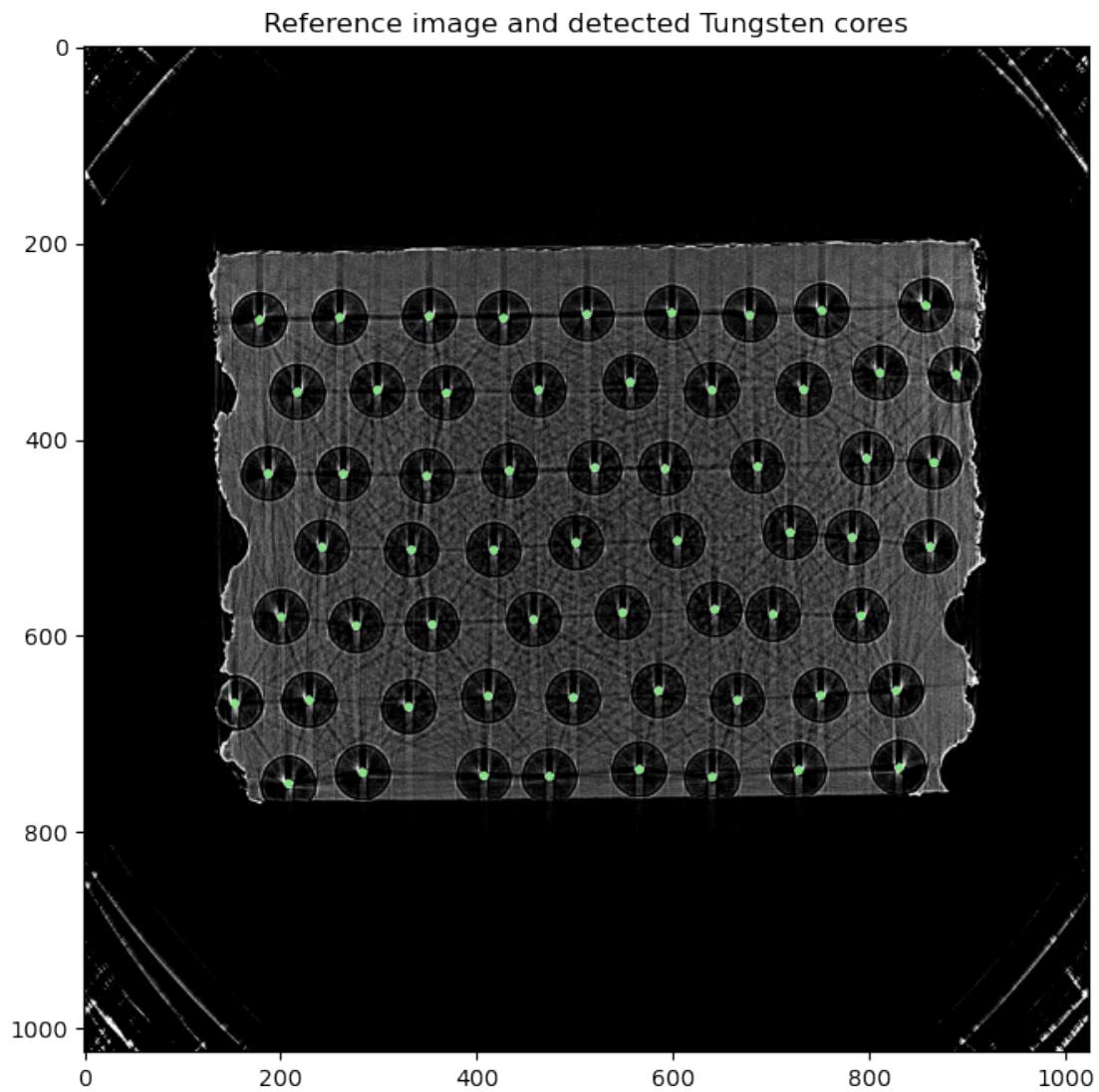
```
[38]: # Clean-up using mathematical morphology
cleaned_thresh_img = sitk.BinaryOpeningByReconstruction(seg, [3, 3, 3])
cleaned_thresh_img = sitk.BinaryClosingByReconstruction(cleaned_thresh_img, [3, 3, 3])

[39]: sitk.WriteImage(cleaned_thresh_img, "outputs/cores_cleaned_segmentation.mha", useCompression=True);
```

```
[40]: fig = plt.figure();

volume = sitk.GetImageFromArray(float2uint8(reference_CT, 0, 50));
volume.SetSpacing([pixel_spacing_in_mm, pixel_spacing_in_mm, ▾
    ↵pixel_spacing_in_mm]);

imgplot = plt.imshow(sitk.GetArrayViewFromImage(sitk.LabelOverlay(volume, ▾
    ↵cleaned_thresh_img)));
plt.title("Reference image and detected Tungsten cores");
plt.savefig('plots/fibre_detection_using_otsu_method_after_cleaning.pdf');
plt.savefig('plots/fibre_detection_using_otsu_method_after_cleaning.png');
```



1.10 Mark each potential tungsten core with unique label

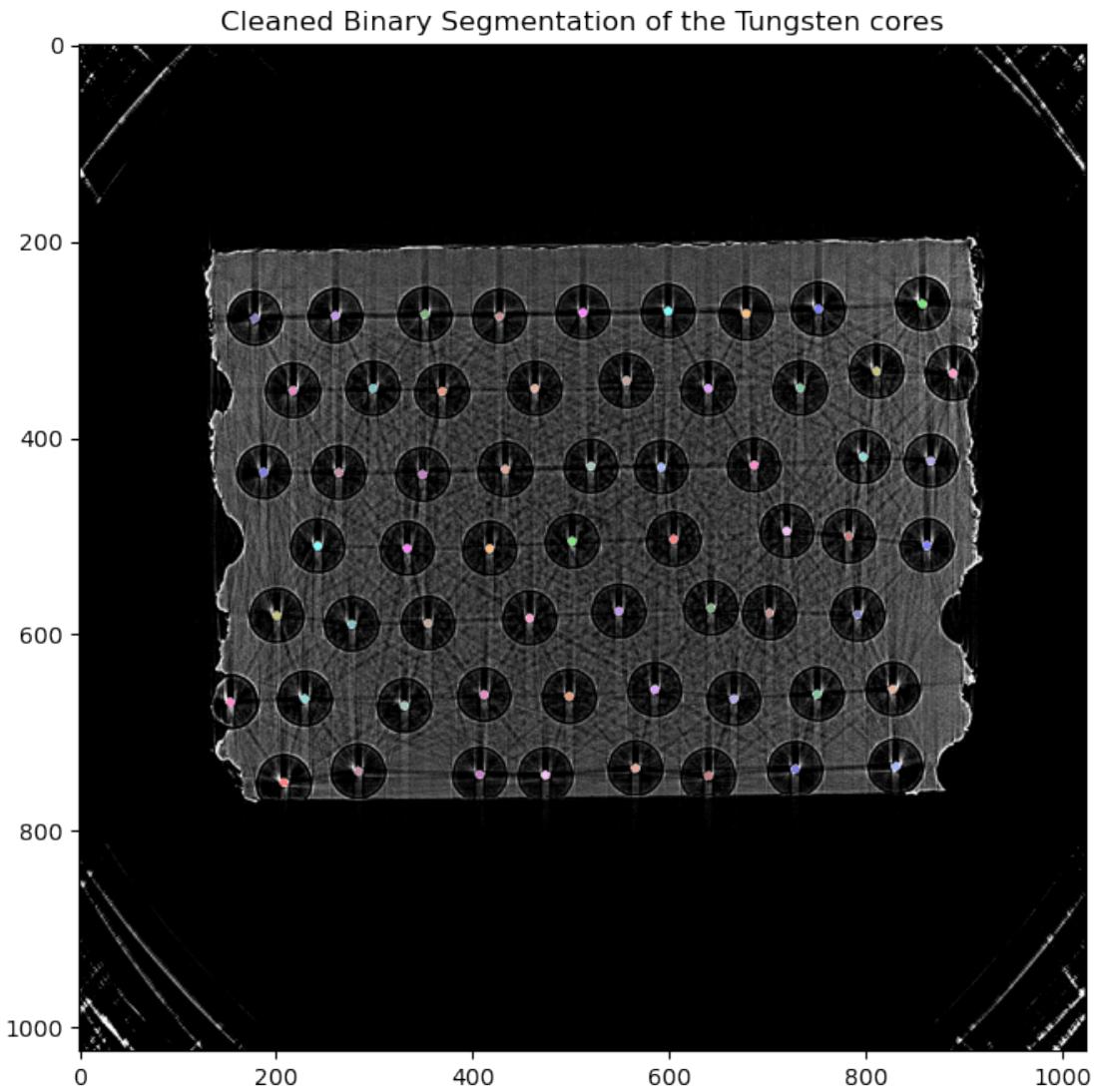
Each distinct tungsten core is assigned a unique label, i.e. a unique pixel intensity

```
[41]: core_labels = sitk.ConnectedComponent(cleaned_thresh_img);
```

```
[42]: fig = plt.figure();

volume = sitk.GetImageFromArray(float2uint8(reference_CT, 0, 50));
volume.SetSpacing([pixel_spacing_in_mm, pixel_spacing_in_mm, ↴
    ↴pixel_spacing_in_mm]);

imgplot = plt.imshow(sitk.GetArrayViewFromImage(sitk.LabelOverlay(volume, ↴
    ↴core_labels)));
plt.title("Cleaned Binary Segmentation of the Tungsten cores");
plt.savefig('plots/fibre_detection_with_label_overlay.pdf');
plt.savefig('plots/fibre_detection_with_label_overlay.png');
```



1.10.1 Object Analysis

Once we have the segmented objects we look at their shapes and the intensity distributions inside the objects. For each labelled tungsten core, we extract the centroid. Note that sizes and positions are given in millimetres.

```
[43]: shape_stats = sitk.LabelShapeStatisticsImageFilter()
shape_stats.ComputeOrientedBoundingBoxOn()
shape_stats.Execute(core_labels)
```

```
[44]: centroid_set = [];
```

```

for i in shape_stats.GetLabels():
    centroid_set.append(cleaned_thresh_img.
    ↪TransformPhysicalPointToIndex(shape_stats.GetCentroid(i)));

```

We now have a list of the centres of all the fibres that can be used as a parameter of the function below to create the cylinders corresponding to the cores and the fibres. For each core, a cylinder is created and translated:

```

gvxr.emptyMesh("core_" + str(i));
gvxr.makeCylinder("core_" + str(i), number_of_sectors, 815.0, core_radius, "micrometre");
gvxr.translateNode("core_" + str(i), y, 0.0, x, "micrometer");

```

For each fibre, another cylinder is created and translated:

```

gvxr.emptyMesh("fibre_" + str(i));
gvxr.makeCylinder("fibre_" + str(i), number_of_sectors, 815.0, fibre_radius, "micrometer");
gvxr.translateNode("fibre_" + str(i), y, 0.0, x, "micrometer");

```

The fibre's cylinder is hollowed to make space for its core:

```
gvxr.subtractMesh("fibre_" + str(i), "core_" + str(i));
```

```
[45]: def setFibres(aCentroidSet):
    """This function loads a cylinders in the GPU memory.
    Some are hollow and represent the fibres, some are not and
    correspond to the cores.

    :param array aCentroidSet: a list of cylinder centres.
    """

    global core_radius;
    global fibre_radius;

    # Create empty geometries
    gvxr.emptyMesh("fibre");
    gvxr.emptyMesh("core");

    # Number of sectors to approximate cylinders with triangle meshes
    # It controls the accuracy of the meshes.
    number_of_sectors = 100;

    # Process all the centres from the input list
    for i, cyl in enumerate(aCentroidSet):

        # Convert the centre position from 2D image coordinates in 3D world coordinates
        x = pixel_spacing_in_micrometre * -(cyl[0] - reference_CT.shape[1] / 2
        ↪+ 0.5);
        y = pixel_spacing_in_micrometre * (cyl[1] - reference_CT.shape[0] / 2 + ↪
        0.5);
```

```

# Create empty geometries (is it needed?)
gvxr.emptyMesh("fibre_" + str(i));
gvxr.emptyMesh("core_" + str(i));

# Create the two corresponding cylinders (fibre and core)
gvxr.makeCylinder("fibre_" + str(i), number_of_sectors, 815.0, □
↳ fibre_radius, "micrometer");
gvxr.makeCylinder("core_" + str(i), number_of_sectors, 815.0, □
↳ core_radius, "micrometer");

# Translate the two cylinders to the position of their centre
gvxr.translateNode("fibre_" + str(i), y, 0.0, x, "micrometer");
gvxr.translateNode("core_" + str(i), y, 0.0, x, "micrometer");

# Apply the local transformation matrix (so that we could save the
↳ corresponding STL files)
gvxr.applyCurrentLocalTransformation("fibre_" + str(i));
gvxr.applyCurrentLocalTransformation("core_" + str(i));

# Subtract the fibre from the matrix
gvxr.subtractMesh("matrix", "fibre_" + str(i));

# Subtract the core from the fibre
gvxr.subtractMesh("fibre_" + str(i), "core_" + str(i));

# Save the corresponding STL files
# gvxr.saveSTLfile("fibre_" + str(i), "Tutorial2/outputs/fibre_" +
↳ str(i) + ".stl");
# gvxr.saveSTLfile("core_" + str(i), "Tutorial2/outputs/core_" +
↳ str(i) + ".stl");

# Add the mesh of the current fibre to the overall fibre mesh
gvxr.addMesh("fibre", "fibre_" + str(i));

# Add the mesh of the current core to the overall core mesh
gvxr.addMesh("core", "core_" + str(i));

# Set the mesh colours (for the interactive visualisation)
gvxr.setColor("fibre", 1.0, 0.0, 0.0, 1.0);
gvxr.setColor("core", 1.0, 0.0, 1.0, 1.0);

# Set the fibre's material properties
# gvxr.setLinearAttenuationCoefficient("fibre", fibre_mu, "cm-1");
gvxr.setCompound("fibre", "SiC");
gvxr.setDensity("fibre", fibre_density, "g/cm3");

```

```

# Set the core's material properties
# gvxr.setLinearAttenuationCoefficient("core", core_mu, "cm-1");
gvxr.setElement("core", "W");

# Add the fibres and cores to the X-ray renderer
gvxr.addPolygonMeshAsInnerSurface("core");
gvxr.addPolygonMeshAsInnerSurface("fibre");

```

1.11 Registration of a cube

We define a function to create the polygon mesh of the Ti90Al6V4 matrix.

```
[46]: def setMatrix(apGeneSet):
    """This function loads a cube in the GPU memory. The cube represents
    the Ti90Al6V4 matrix.

    apGeneSet[0] is a number between -0.5 and 0.5, related to the translation
    vector (X component) of the cube. It can be interpreted as a percentage of
    the detector width.

    apGeneSet[1] is the same as apGeneSet[0], but related to the Y component of
    the translation vector.

    apGeneSet[2] is a number between -0.5 and 0.5, related to the rotation
    angle in degrees

    apGeneSet[3] is a scaling factor between -0.5 and 0.5. It can be
    interpreted as a percentage of the detector width.

    apGeneSet[4] is a scaling factor between -0.5 and 0.5. It can be
    interpreted as a percentage of apGeneSet[3].
    """

    # Remove all the geometries from the whole scenegraph
    gvxr.removePolygonMeshesFromSceneGraph();

    # Make a cube
    gvxr.makeCube("matrix", 1.0, "micrometer");

    # Translation vector
    x = apGeneSet[0] * detector_width_in_pixels * pixel_spacing_in_micrometre;
    y = apGeneSet[1] * detector_width_in_pixels * pixel_spacing_in_micrometre;
    gvxr.translateNode("matrix", x, 0.0, y, "micrometer");

    # Rotation angle
    rotation_angle_in_degrees = (apGeneSet[2] + 0.5) * 180.0;
    gvxr.rotateNode("matrix", rotation_angle_in_degrees, 0, 1, 0);

    # Scaling factors
```

```

w = (apGeneSet[3] + 0.5) * detector_width_in_pixels * pixel_spacing_in_micrometre;
h = (apGeneSet[4] + 0.5) * w;
gvxr.setScaleNode("matrix", w, 815, h);

# Apply the transformation matrix so that we can save the corresponding STLfile
gvxr.applyCurrentLocalTransformation("matrix");

# Set the matrix's material properties
gvxr.setMixture("matrix", "Ti90Al16V4");
gvxr.setDensity("matrix", matrix_density, "g/cm3");

# Add the matrix to the X-ray renderer
gvxr.addPolygonMeshAsInnerSurface("matrix");

```

1.11.1 Simulate the CT acquisition

There are 7 successive steps to simulate the XCT data acquisition:

1. Set the fibre and cores geometries and material properties (Step 45)
2. Set the matrix geometry and material properties (Step 46)
3. Simulate the raw projections for each angle:
 - Without phase contrast (Line 9 of Step 49), or
 - With phase contrast (Lines 12-55 of Step 49)
4. Apply the LSF (Lines 57-60 of Step 49)
5. Apply the flat-field correction (Line 63 of Step 49)
6. Add Poisson noise (Lines 67-76 of Step 49)(Lines 12-55 of Step 49)
7. Apply the minus log normalisation to compute the sinogram (Line 79 of Step 49)

Compute the raw projections and save the data. For this purpose, we define a new function.

```
[47]: def tomographyAcquisition():
    """
    This function simulate a CT acquisition.

    :return the raw projections in keV
    """

    # Create a new array to save every projection in default unit of energy
    raw_projections = [];

    # For each angle, simulate a projection
    for angle_id in range(0, number_of_projections):

        # Reset the transformation matrix and rotate the scnned object
        gvxr.resetSceneTransformation();
```

```

gvxr.rotateScene(-angular_step * angle_id, 0, 1, 0);

# Compute the X-ray image
xray_image = np.array(gvxr.computeXRayImage());

# Add the projection
raw_projections.append(xray_image);

# Convert from the default unit of energy to keV
raw_projections = np.array(raw_projections);
raw_projections_in_keV = raw_projections / gvxr.getUnitOfEnergy("keV");

return raw_projections_in_keV;

```

1.11.2 Flat-field correction

Because the data suffers from a fixed-pattern noise in X-ray imaging in actual experiments, it is necessary to perform the flat-field correction of the raw projections using:

$$\mathbf{Proj} = \frac{\mathbf{I} - \mathbf{D}}{\mathbf{F} - \mathbf{D}}$$

F (full fields) and **D** (dark fields) are projection images without sample and acquired with and without the X-ray beam turned on respectively.

Note that in our example, `raw_projections_in_keV` (**I**), `flat_field_image` (**F**) and `dark_field_image` (**D**) are in keV whereas `corrected_projections` (**Proj**) does not have any unit:

$$0 \leq \mathbf{I} \leq \sum_E N_0(E) \times E \quad 0 \leq \mathbf{Proj} \leq 1$$

We define a new function to compute the flat-field correction.

```
[48]: def flatFieldCorrection(raw_projections_in_keV):
    """
    This function applies the flat-field correction on raw projections.

    :param 2D_image raw_projections_in_keV: the raw X-ray projections in keV
    :return the projections (raw_projections_in_keV) after flat-field correction
    """

    # Create a mock dark field image
    dark_field_image = np.zeros(raw_projections_in_keV.shape);

    # Create a mock flat field image
    flat_field_image = np.ones(raw_projections_in_keV.shape);
```

```

# Retrieve the total energy
total_energy = 0.0;
energy_bins = gvxr.getEnergyBins("keV");
photon_count_per_bin = gvxr.getPhotonCountEnergyBins();

for energy, count in zip(energy_bins, photon_count_per_bin):
    total_energy += energy * count;
flat_field_image *= total_energy;

# Apply the actual flat-field correction on the raw projections
corrected_projections = (raw_projections_in_keV - dark_field_image) / (flat_field_image - dark_field_image);

return corrected_projections;

```

The function below is used to simulate a sinogram acquisition. Phase contrast in the projections can be taken into account or not. Also, Poisson noise can be added.

```
[49]: def simulateSinogram(sigma_set=None, k_set=None, name_set=None):

    global lsf_kernel;

    # Do not simulate the phase contrast using a Laplacian
    if isinstance(sigma_set, NoneType) or isinstance(k_set, NoneType) or
       isinstance(name_set, NoneType):

        # Get the raw projections in keV
        raw_projections_in_keV = tomographyAcquisition();

    # Simulate the phase contrast using a Laplacian
    else:

        # Create the convolution filter
        pixel_range = np.linspace(-value_range, value_range, num=int(num_samples), endpoint=True)
        laplacian_kernels = {};

        # Store the L-buffers
        L_buffer_set = {};

        # Look at all the children of the root node
        for label in ["core", "fibre", "matrix"]:
            # Get its L-buffer
            L_buffer_set[label] = getLBuffer(label);

        # Create blank images

```

```

raw_projections_in_keV = np.zeros(L_buffer_set["fibre"].shape);
phase_contrast_image = np.zeros(L_buffer_set["fibre"].shape);

for label, k, sigma in zip(name_set, k_set, sigma_set):
    laplacian_kernels[label] = k * laplacian(pixel_range, sigma);

    for z in range(phase_contrast_image.shape[0]):
        for y in range(phase_contrast_image.shape[1]):
            phase_contrast_image[z][y] += ndimage.
    convolve((L_buffer_set[label])[z][y], laplacian_kernels[label], mode='wrap');

    for energy, photon_count in zip(gvxr.getEnergyBins("keV"), gvxr.
    getPhotonCountEnergyBins()):

        # Create a blank image
        attenuation = np.zeros(L_buffer_set["fibre"].shape);

        # Look at all the children of the root node
        # for label in ["core", "fibre", "matrix"]:
        for label in ["core", "fibre", "matrix"]:
            # Get mu for this object for this energy
            mu = gvxr.getLinearAttenuationCoefficient(label, energy, "keV");

            # Compute sum mu * x
            attenuation += L_buffer_set[label] * mu;

        # Store the projection for this energy channel
        raw_projections_in_keV += energy * photon_count * np.
    exp(-attenuation);

        # Apply the phase contrast
        raw_projections_in_keV -= phase_contrast_image;

        # Apply the LSF line by line
        for z in range(raw_projections_in_keV.shape[0]):
            for y in range(raw_projections_in_keV.shape[1]):
                raw_projections_in_keV[z][y] = ndimage.
    convolve(raw_projections_in_keV[z][y], lsf_kernel, mode='wrap');

        # Flat-field correction
        normalised_projections = flatFieldCorrection(raw_projections_in_keV);
        normalised_projections[normalised_projections < 0] = 0;

        # Add noise
        if not isinstance(bias, NoneType) and not isinstance(gain, NoneType) and
    not isinstance(scale, NoneType):

```

```

    map = (normalised_projections + (bias + 1)) * gain;
    temp = np.random.poisson(map).astype(float);
    temp /= gain;
    temp -= bias + 1;

    # Noise map
    noise_map = (normalised_projections - temp) * scale;
    normalised_projections += noise_map;

    # Linearise
    simulated_sinogram = computeSinogramFromFlatField(normalised_projections);

    return simulated_sinogram, normalised_projections, raw_projections_in_keV;

```

The function below is used quantify the differences between two images. It is used in the objective function.

```
[50]: def metrics(ref, test):

    normalised_ref = ref.flatten();
    normalised_test = test.flatten();

    if use_normalisation or metrics_type == "ZNCC":
        normalised_ref = standardisation(normalised_ref);
        normalised_test = standardisation(normalised_test);

    # Mean absolute error
    if metrics_type == "MAE":
        return mean_absolute_error(normalised_ref, normalised_test);
    # RMSE
    elif metrics_type == "RMSE":
        return math.sqrt(mean_squared_error(normalised_ref, normalised_test));
    # Mean relative error
    elif metrics_type == "MRE" or metrics_type == "MAPE":

        # Prevent division by zero
        denominator = np.abs(np.subtract(normalised_ref, normalised_test)) + ↵
        ↵1e-6;
        divisor = np.abs(normalised_ref) + 1e-6;

        return np.mean(np.divide(denominator, divisor));
    elif metrics_type == "SSIM" or metrics_type == "DSSIM":
        normalised_ref.shape = [900, 1024];
        normalised_test.shape = [900, 1024];
        return (1.0 - ssim(normalised_ref, normalised_test,
                           data_range=normalised_ref.max() - normalised_ref.min())) / 2.
        ↵0;
```

```

    elif metrics_type == "ZNCC":
        return (1.0 - np.mean(np.multiply(normalised_ref, normalised_test))) / ↵2.0;
    else:
        raise "Unknown metrics";

```

The function below is the objective function used to register the matrix.

```
[51]: def fitnessFunctionCube(x):
    global best_fitness;
    global best_fitness_id;
    global prefix;

    global reference_sinogram;
    global centroid_set;
    global use_fibres;

    global core_radius;
    global fibre_radius;

    # Load the matrix geometrical properties from x
    setMatrix(x);

    # Simulate a sinogram
    simulated_sinogram, normalised_projections, raw_projections_in_keV = ↵simulateSinogram(sigma_set, k_set, label_set);

    # Compute the objective value
    if use_sinogram:
        objective = metrics(reference_sinogram, simulated_sinogram);
    else:
        objective = metrics(reference_normalised_projections, ↵normalised_projections);

    # The block below is not necessary for the registration.
    # It is used to save the data to create animations.
    if best_fitness > objective:
        best_fitness = objective;

        gvxr.saveSTLfile("matrix", "outputs/matrix_" + str(best_fitness_id) + ".stl");

    # Reconstruct the CT slice
    simulated_CT = tomopy.recon(simulated_sinogram,
                                theta_rad,
                                center=rot_center,
                                sinogram_order=False,
```

```

        algorithm='gridrec',
        filter_name='shepp',
        ncore=40)[0];

    # Save the simulated sinogram
    simulated_sinogram.shape = (simulated_sinogram.size //_
    ↪simulated_sinogram.shape[2], simulated_sinogram.shape[2]);
    saveMHA("outputs/" + prefix + "simulated_sinogram_" +_
    ↪str(best_fitness_id) + ".mha",
            simulated_sinogram,
            [pixel_spacing_in_mm, angular_step, pixel_spacing_in_mm]);

    # Save the simulated CT slice
    saveMHA("outputs/" + prefix + "simulated_CT_" + str(best_fitness_id) +_
    ↪".mha",
            simulated_CT,
            [pixel_spacing_in_mm, pixel_spacing_in_mm,_
    ↪pixel_spacing_in_mm]);

    np.savetxt("outputs/" + prefix + str(best_fitness_id) + ".dat", x,_
    ↪header='x,y,rotation_angle,w,h');

    best_fitness_id += 1;

    return objective

```

[52]: # The registration has already been performed. Load the results.

```

if os.path.isfile("outputs/cube.dat"):
    matrix_geometry_parameters = np.loadtxt("outputs/cube.dat");
# Perform the registration using CMA-ES
else:
    best_fitness = sys.float_info.max;
    best_fitness_id = 0;
    prefix = "cube_";

    opts = cma.CMAOptions()
    opts.set('tolfun', 1e-2);
    opts['tolx'] = 1e-2;
    opts['bounds'] = [5*[-0.5], 5*[0.5]];

    es = cma.CMAEvolutionStrategy([0.0, 0.0, 0.0, 0.256835938, 0.232903226], 0.-
    ↪5, opts);
    es.optimize(fitnessFunctionCube);

    matrix_geometry_parameters = copy.deepcopy(es.result.xbest);

```

```

np.savetxt("outputs/cube.dat", matrix_geometry_parameters, u
→header='x,y,rotation_angle,w,h');

# Release memory
del es;

```

(4_w,8)-aCMA-ES ($\mu_w=2.6, w_1=52\%$) in dimension 5 (seed=723641, Wed Aug 11
17:07:25 2021)

Iterat	#Fevals	function	value	axis	ratio	sigma	min&max	std	t[m:s]
1	8	1.094954737592163e+00	1.0e+00	4.27e-01	4e-01	4e-01	0:04.7		
2	16	7.847555640186077e-01	1.2e+00	3.57e-01	3e-01	4e-01	0:08.1		
3	24	1.115136747209895e+00	1.4e+00	3.64e-01	3e-01	4e-01	0:11.3		
4	32	1.231106093423999e+00	1.4e+00	4.03e-01	3e-01	5e-01	0:14.6		
6	48	8.833336352985762e-01	1.8e+00	3.71e-01	3e-01	4e-01	0:21.0		
8	64	1.056062914736723e+00	1.9e+00	2.99e-01	2e-01	3e-01	0:27.2		
10	80	9.854065226044122e-01	2.3e+00	2.59e-01	1e-01	3e-01	0:33.3		
13	104	6.638930606599894e-01	2.2e+00	1.87e-01	1e-01	2e-01	0:42.7		
16	128	5.974250371930541e-01	2.5e+00	1.49e-01	7e-02	1e-01	0:52.6		
19	152	6.176599538384471e-01	2.2e+00	1.11e-01	5e-02	9e-02	1:02.1		
22	176	5.260765226564615e-01	2.5e+00	1.66e-01	7e-02	2e-01	1:12.3		
26	208	3.408282972258348e-01	3.2e+00	1.01e-01	3e-02	9e-02	1:25.4		
30	240	3.253546297790291e-01	3.1e+00	6.05e-02	2e-02	5e-02	1:37.9		
35	280	2.736482631917837e-01	3.4e+00	3.37e-02	9e-03	2e-02	1:53.8		
40	320	2.484147692010300e-01	2.8e+00	1.76e-02	4e-03	9e-03	2:10.5		

1.11.3 Apply the result of the registration

```

[53]: # Save the result
setMatrix(matrix_geometry_parameters);
gvxr.saveSTLfile("matrix", "outputs/matrix.stl");

# Translation vector
x = matrix_geometry_parameters[0] * detector_width_in_pixels * u
→pixel_spacing_in_micrometre;
y = matrix_geometry_parameters[1] * detector_width_in_pixels * u
→pixel_spacing_in_micrometre;

# Rotation angle
rotation_angle_in_degrees = (matrix_geometry_parameters[2] + 0.5) * 180.0;

# Scaling factors
w = (matrix_geometry_parameters[3] + 0.5) * detector_width_in_pixels * u
→pixel_spacing_in_micrometre;
h = (matrix_geometry_parameters[4] + 0.5) * w;

print("Matrix");

```

```

print("\tposition:", x, y, "um");
print("\trotation:", rotation_angle_in_degrees, "deg");
print("\tsize:", w, h, "um");

```

Matrix

```

position: -61.21655372351931 -18.65939030141249 um
rotation: 90.5658762058477 deg
size: 1429.3872912018892 1074.742903753673 um

```

1.11.4 Simulate the correspond CT acquisition

```
[54]: # Simulate a sinogram
simulated_sinogram, normalised_projections, raw_projections_in_keV =
    →simulateSinogram();

# Reconstruct the CT slice
simulated_CT = tomopy.recon(simulated_sinogram,
                             theta_rad,
                             center=rot_center,
                             sinogram_order=False,
                             algorithm='gridrec',
                             filter_name='shepp',
                             ncore=40)[0];
normalised_simulated_CT = standardisation(simulated_CT);

# Compute the ZNCC
print("ZNCC matrix registration:",
      "{:.2f}".format(100.0 * np.mean(np.multiply(normalised_reference_CT,
                                                 →normalised_simulated_CT))));
```

ZNCC matrix registration: 73.26

1.11.5 Display the result of the registration as an animation

```
[55]: def createAnimation(aPrefix, anOutputFile):
    # Find all the images from the output directory
    files = sorted(
        glob.glob(aPrefix + "[0-9]*.mha"))

    # Store the images
    registration_image_set = [];

    # Create the GIF file
    with imageio.get_writer(anOutputFile, mode='I') as writer:
```

```

# Store the PNG filenames
png_filename_set = [];

# Process all the images
for i in range(len(files)):
    # Create the filenames
    mha_fname = aPrefix + str(i) + ".mha";
    png_filename_set.append(aPrefix + str(i) + ".png");

    # Open the MHA file
    float_image = sitk.ReadImage(mha_fname);

    # Convert in a Numpy array
    narray = sitk.GetArrayFromImage(float_image);

    offset = 60;
    roi_ref = reference_CT[505 - offset:505 + offset + 1,501 - offset:
    ↪501 + offset + 1];
    roi_sim = narray[505 - offset:505 + offset + 1,501 - offset:501 + ↪
    ↪offset + 1];

    narray = standardisation(narray);
    registration_image_set.append(narray);

    # Create the figure
    fig, axs = plt.subplots(3, 3)

    # Display the reference, registration and error map
    fig.suptitle('Registration: Result ' + str(i+1) + "/" + ↪
    ↪str(len(files)))
    plt.tight_layout();
    norm = cm.colors.Normalize(vmax=1.25, vmin=-0.5)

    comp_equalized = compare_images(normalised_reference_CT, narray, ↪
    ↪method='checkerboard');

    roi_normalised_ref = normalised_reference_CT[505 - offset:505 + ↪
    ↪offset + 1,501 - offset:501 + offset + 1];
    roi_normalised_sim = narray[505 - offset:505 + offset + 1,501 - ↪
    ↪offset:501 + offset + 1];
    roi_normalised_compare = comp_equalized[505 - offset:505 + offset + ↪
    ↪1,501 - offset:501 + offset + 1];

    # Reference
    axs[0, 0].set_title("Reference image");

```

```

    axs[0, 0].imshow(normalised_reference_CT, cmap="gray", norm=norm);
    axs[1, 0].imshow(roi_normalised_ref, cmap="gray", norm=norm);
    axs[2, 0].axis('off');

    # Registration
    axs[0, 1].set_title("Simulated CT slice after automatic\u
→registration");
    axs[0, 1].imshow(narray, cmap='gray', norm=norm);
    axs[1, 1].imshow(roi_normalised_sim, cmap="gray", norm=norm);

    y_ref = np.diag(roi_ref)
    y_test = np.diag(roi_sim)
    len_y = len(roi_ref)

    zero_offset = round(57.32 + (63.64 - 57.32) / 2.0)
    x = np.array(range(len_y)).astype(float)
    x -= zero_offset + 0.5
    x = x * math.sqrt(2.0 * pixel_spacing_in_micrometre * \u
→pixel_spacing_in_micrometre)

    axs[2, 1].set_title("Diagonal profiles");
    axs[2, 1].plot(x, y_ref, label="Reference");
    axs[2, 1].plot(x, y_test, label="Simulated");
    axs[2, 1].set_xlabel("Distance along profile (in $\mathbf{\mu}m$)");
    axs[2, 1].set_ylabel("Linear attenuation coefficients\n(in\u
→cm$^{-1}$)")

    # axs[2, 1].plot(np.diag(roi_ref) - np.diag(roi_sim), \u
→label="Error");
    axs[2, 1].legend();

    # Error map
    ZNCC = 100.0 * np.mean(np.multiply(normalised_reference_CT,\u
→narray));
    axs[0, 2].set_title("Checkboard comparison between\nthe reference\u
→and simulated images\nZNCC: " + "{:.2f}".format(ZNCC));
    axs[0, 2].imshow(comp_equalized, cmap='gray', norm=norm);
    axs[1, 2].imshow(roi_normalised_compare, cmap='gray', norm=norm);
    axs[2, 2].axis('off');

    plt.tight_layout();

    # Save the figure as a PNG file
    plt.savefig(png_filename_set[i])

    # Close the figure
    plt.close()

```

```
# Open the PNG file with imageio and add it to the GIF file
image = imageio.imread(png_filename_set[i])
writer.append_data(image)

# Delete the PNG file
os.remove(png_filename_set[i]);

for i in range(15):
    writer.append_data(image)

return registration_image_set, image;
```

```
[56]: if not os.path.exists("plots/cube_registration.gif"):
    cube_registration_image_set, image = createAnimation("outputs/
→cube_simulated_CT_",
    'plots/cube_registration.gif');
```

View the GIF animation (no preview on GitHub .ipynb files)

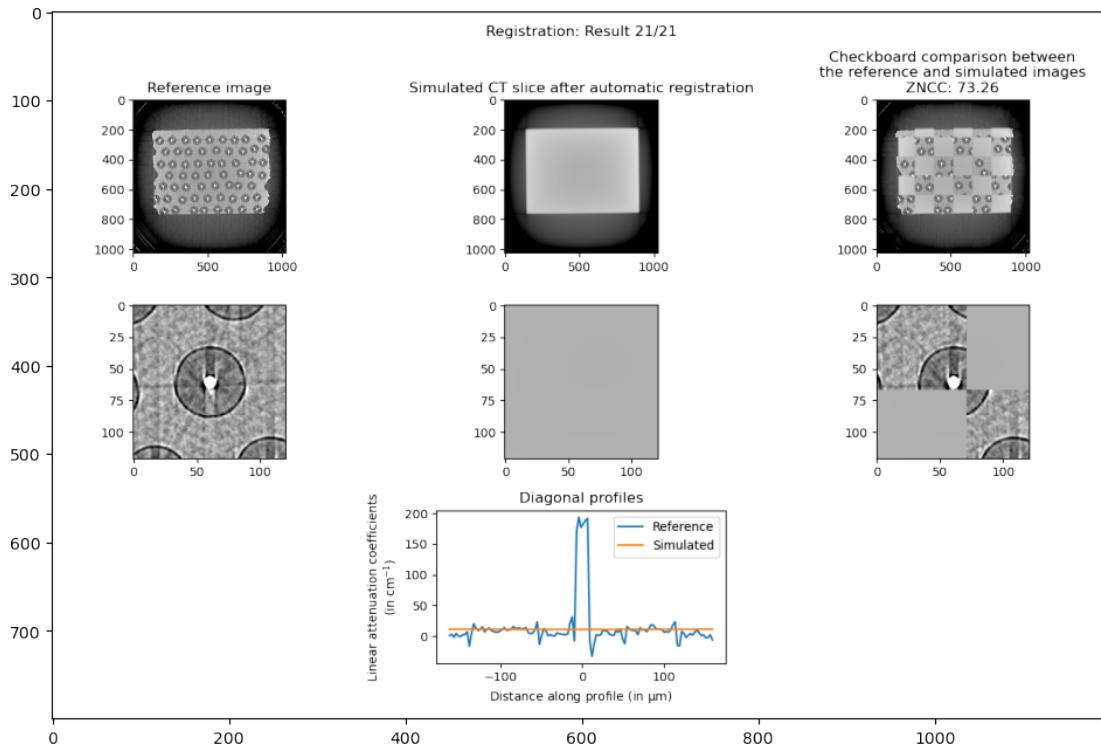
```
[57]: Image(url="plots/cube_registration.gif")
```

```
[57]: <IPython.core.display.Image object>
```

View the last frame

```
[58]: plt.imshow(image)
```

```
[58]: <matplotlib.image.AxesImage at 0x7f3dee224160>
```



1.11.6 Adding the fibres

The radius of a tungsten core is $30 / 2 \text{ um}$. The pixel spacing is 1.9 um . The radius in number of pixels is $15/1.9 \approx 7.89$. The area of a core is $(15/1.9)^2\pi \approx 196$ pixels.

```
[59]: setMatrix(matrix_geometry_parameters);
setFibres(centroid_set);
```

```
[60]: # Simulate a sinogram
simulated_sinogram, normalised_projections, raw_projections_in_keV =
    simulateSinogram(sigma_set, k_set, label_set);
```

```
[61]: # Reconstruct the CT slice
simulated_CT = tomopy.recon(simulated_sinogram,
                             theta_rad,
                             center=rot_center,
                             sinogram_order=False,
                             algorithm='gridrec',
                             filter_name='shepp',
                             ncore=40)[0];
normalised_simulated_CT = standardisation(simulated_CT);
```

```
# Compute the ZNCC
print("ZNCC matrix registration with fibres:",
      "{:.2f}".format(100.0 * np.mean(np.multiply(normalised_reference_CT, □
→normalised_simulated_CT))));
```

ZNCC matrix registration with fibres: 69.25

```
[62]: simulated_sinogram.shape = (simulated_sinogram.size //□
→simulated_sinogram.shape[2], simulated_sinogram.shape[2]);
normalised_projections.shape = (normalised_projections.size //□
→normalised_projections.shape[2], normalised_projections.shape[2]);
raw_projections_in_keV.shape = (raw_projections_in_keV.size //□
→raw_projections_in_keV.shape[2], raw_projections_in_keV.shape[2]);

saveMHA("outputs/simulated_sinogram_with_fibres.mha",
        simulated_sinogram,
        [pixel_spacing_in_mm, angular_step, pixel_spacing_in_mm]);

saveMHA("outputs/normalised_projections_with_fibres.mha",
        normalised_projections,
        [pixel_spacing_in_mm, angular_step, pixel_spacing_in_mm]);

saveMHA("outputs/raw_projections_in_keV_with_fibres.mha",
        raw_projections_in_keV,
        [pixel_spacing_in_mm, angular_step, pixel_spacing_in_mm]);
```

```
[63]: saveMHA("outputs/simulated_CT_with_fibres.mha",
            simulated_CT,
            [pixel_spacing_in_mm, pixel_spacing_in_mm, pixel_spacing_in_mm]);

saveMHA("outputs/normalised_simulated_CT_with_fibres.mha",
        normalised_simulated_CT,
        [pixel_spacing_in_mm, pixel_spacing_in_mm, pixel_spacing_in_mm]);
```

```
[64]: norm = cm.colors.Normalize(vmax=1.25, vmin=-0.5)

fig, (ax1, ax2, ax3) = plt.subplots(1, 3)
plt.tight_layout()
fig.suptitle('CT slice with fibres after the registration')

ax1.set_title("Reference image");
imgplot1 = ax1.imshow(normalised_reference_CT, cmap="gray",
                      norm=norm);

ax2.set_title("Simulated CT slice after automatic registration");
imgplot2 = ax2.imshow(normalised_simulated_CT,
                      cmap='gray',
```

```

        norm=norm);

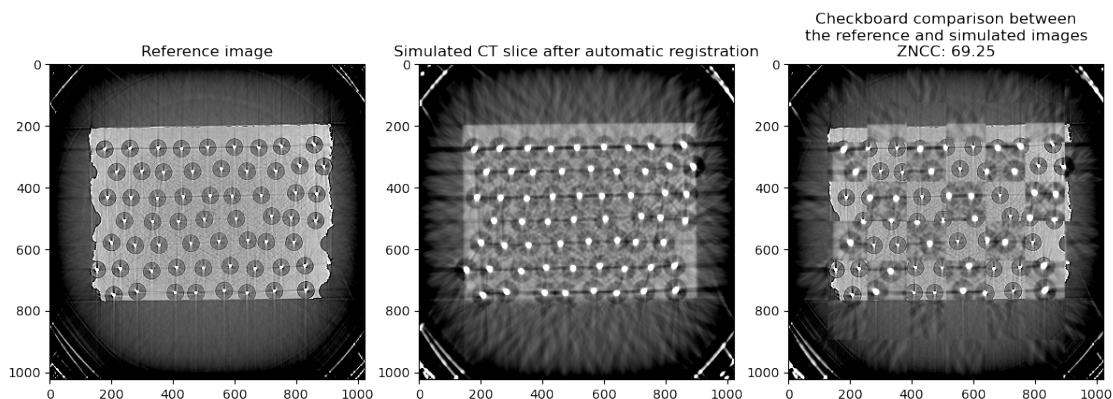
comp_equalized = compare_images(normalised_reference_CT,
                                 normalised_simulated_CT, method='checkerboard');
ax3.set_title("Checkboard comparison between\n" +
              "the reference and simulated images\nZNCC: " +
              "{:.2f}".format(100.0 * np.mean(np.
                                 multiply(normalised_reference_CT, normalised_simulated_CT))));

imgplot3 = ax3.imshow(comp_equalized,
                      cmap='gray',
                      norm=norm);

plt.savefig('plots/simulated_CT_slice_with_fibres_after_cube_registration.pdf');
plt.savefig('plots/simulated_CT_slice_with_fibres_after_cube_registration.png');

```

CT slice with fibres after the registration



1.12 Optimisation of the cores and fibres radii

The function below is the objective function used to optimise the radii of the cores and fibres.

```
[65]: def fitnessFunctionFibres(x):
    global best_fitness;
    global best_fitness_id;
    global fibre_radius;
    global core_radius;
    global prefix;

    # Get the radii
    fibre_radius = x[0];
```

```

core_radius = fibre_radius * x[1];

# Load the matrix
setMatrix(matrix_geometry_parameters);

# Load the cores and fibres
setFibres(centroid_set);

# Simulate a sinogram
simulated_sinogram, normalised_projections, raw_projections_in_keV =
    simulateSinogram(sigma_set, k_set, label_set);

# Compute the objective value
if use_sinogram:
    objective = metrics(reference_sinogram, simulated_sinogram);
else:
    objective = metrics(reference_normalised_projections,
    normalised_projections);

# The block below is not necessary for the registration.
# It is used to save the data to create animations.
if best_fitness > objective:
    best_fitness = objective;

    gvxr.saveSTLfile("core", "outputs/" + prefix + str(best_fitness_id) +
        "_cores.stl");
    gvxr.saveSTLfile("fibre", "outputs/" + prefix + str(best_fitness_id) +
        "_fibres.stl");

# Reconstruct the CT slice
simulated_CT = tomopy.recon(simulated_sinogram,
                            theta_rad,
                            center=rot_center,
                            sinogram_order=False,
                            algorithm='gridrec',
                            filter_name='shepp',
                            ncore=40)[0];

# Save the simulated sinogram
simulated_sinogram.shape = (simulated_sinogram.size //_
    simulated_sinogram.shape[2], simulated_sinogram.shape[2]);
    saveMHA("outputs/" + prefix + "simulated_sinogram_" +_
    str(best_fitness_id) + ".mha",
            simulated_sinogram,
            [pixel_spacing_in_mm, angular_step, pixel_spacing_in_mm]);

```

```

# Save the simulated CT slice
saveMHA("outputs/" + prefix + "simulated_CT_" + str(best_fitness_id) + ".mha",
        simulated_CT,
        [pixel_spacing_in_mm, pixel_spacing_in_mm, pixel_spacing_in_mm]);

np.savetxt("outputs/" + prefix + str(best_fitness_id) + ".dat", x,
           header='x,y,rotation_angle,w,h');

best_fitness_id += 1;

return objective

```

[66]: # The registration has already been performed. Load the results.

```

if os.path.isfile("outputs/fibre1_radii.dat"):
    temp = np.loadtxt("outputs/fibre1_radii.dat");
    core_radius = temp[0];
    fibre_radius = temp[1];
# Perform the registration using CMA-ES
else:
    ratio = core_radius / fibre_radius;

x0 = [fibre_radius, ratio];
bounds = [[5, 0.01], [1.5 * fibre_radius, 0.95]];

best_fitness = sys.float_info.max;
best_fitness_id = 0;
prefix = "fibre1_";

opts = cma.CMAOptions()
opts.set('tolfun', 1e-3);
opts['tolx'] = 1e-3;
opts['bounds'] = bounds;

es = cma.CMAEvolutionStrategy(x0, 0.9, opts);
es.optimize(fitnessFunctionFibres);
fibre_radius = es.result.xbest[0];
core_radius = fibre_radius * es.result.xbest[1];

np.savetxt("outputs/fibre1_radii.dat", [core_radius, fibre_radius],
           header='core_radius_in_um,fibre_radius_in_um');

# Release memory
del es;

```

```
(3_w,6)-aCMA-ES (mu_w=2.0,w_1=63%) in dimension 2 (seed=826019, Wed Aug 11
17:09:53 2021)
Iterat #Fevals function value axis ratio sigma min&max std t[m:s]
 1      6 2.133708698445436e-01 1.0e+00 7.09e-01 6e-01 6e-01 0:08.1
 2     12 3.884900300610271e-01 1.3e+00 7.12e-01 6e-01 7e-01 0:14.7
 3     18 3.505666956237065e-01 1.5e+00 8.88e-01 8e-01 1e+00 0:21.3
 4     24 1.908870801154281e-01 1.9e+00 9.15e-01 7e-01 1e+00 0:28.3
 5     30 4.094047797452126e-01 2.2e+00 8.09e-01 6e-01 1e+00 0:34.9
 6     36 2.178027339316914e-01 2.4e+00 6.54e-01 4e-01 8e-01 0:41.5
 7     42 2.848424211802671e-01 2.9e+00 5.46e-01 3e-01 7e-01 0:48.2
 9     54 2.081527321038656e-01 3.1e+00 5.26e-01 2e-01 7e-01 1:01.7
11    66 1.993601513846019e-01 3.6e+00 3.82e-01 1e-01 4e-01 1:15.2
13    78 2.509825026536547e-01 3.3e+00 2.79e-01 7e-02 2e-01 1:28.3
15    90 2.086870907711814e-01 3.4e+00 1.96e-01 4e-02 1e-01 1:41.4
17   102 2.042247123904917e-01 4.2e+00 2.60e-01 4e-02 2e-01 1:54.6
19   114 1.990669251567721e-01 8.1e+00 2.23e-01 2e-02 2e-01 2:07.8
21   126 1.950387888441152e-01 1.1e+01 3.52e-01 3e-02 3e-01 2:21.2
24   144 1.929368860330429e-01 1.5e+01 3.05e-01 2e-02 3e-01 2:41.5
27   162 1.794416550710599e-01 2.9e+01 8.90e-01 4e-02 1e+00 3:02.8
30   180 1.944530796732157e-01 4.1e+01 4.74e-01 1e-02 5e-01 3:23.1
33   198 1.771589387587338e-01 4.8e+01 7.40e-01 2e-02 1e+00 3:44.3
36   216 1.773296409555877e-01 9.0e+01 1.03e+00 2e-02 2e+00 4:05.2
39   234 1.724468003965145e-01 1.5e+02 4.94e-01 6e-03 6e-01 4:25.0
42   252 1.644278423948802e-01 1.4e+02 4.86e-01 4e-03 5e-01 4:45.1
45   270 1.483282044257309e-01 2.1e+02 1.35e+00 1e-02 2e+00 5:08.3
49   294 1.290201572100871e-01 3.0e+02 2.21e+00 2e-02 4e+00 5:35.4
53   318 1.253218839744022e-01 3.9e+02 1.80e+00 1e-02 3e+00 6:02.8
57   342 1.249622693692700e-01 5.4e+02 5.70e-01 2e-03 5e-01 6:31.4
61   366 1.249538083261566e-01 4.9e+02 2.06e-01 5e-04 1e-01 6:58.8
65   390 1.249531854848102e-01 4.4e+02 1.04e-01 2e-04 4e-02 7:25.6
69   414 1.249527863407902e-01 4.3e+02 9.64e-02 1e-04 3e-02 7:53.6
70   420 1.249526838072387e-01 3.5e+02 9.70e-02 1e-04 3e-02 8:00.4
```

```
[67]: if not os.path.exists("plots/fibre1_registration.gif"):
    registration_image_set, image = createAnimation("outputs/
→fibre1_simulated_CT",
                                                'plots/fibre1_registration.gif');
```

View the GIF animation (no preview on GitHub .ipynb files)

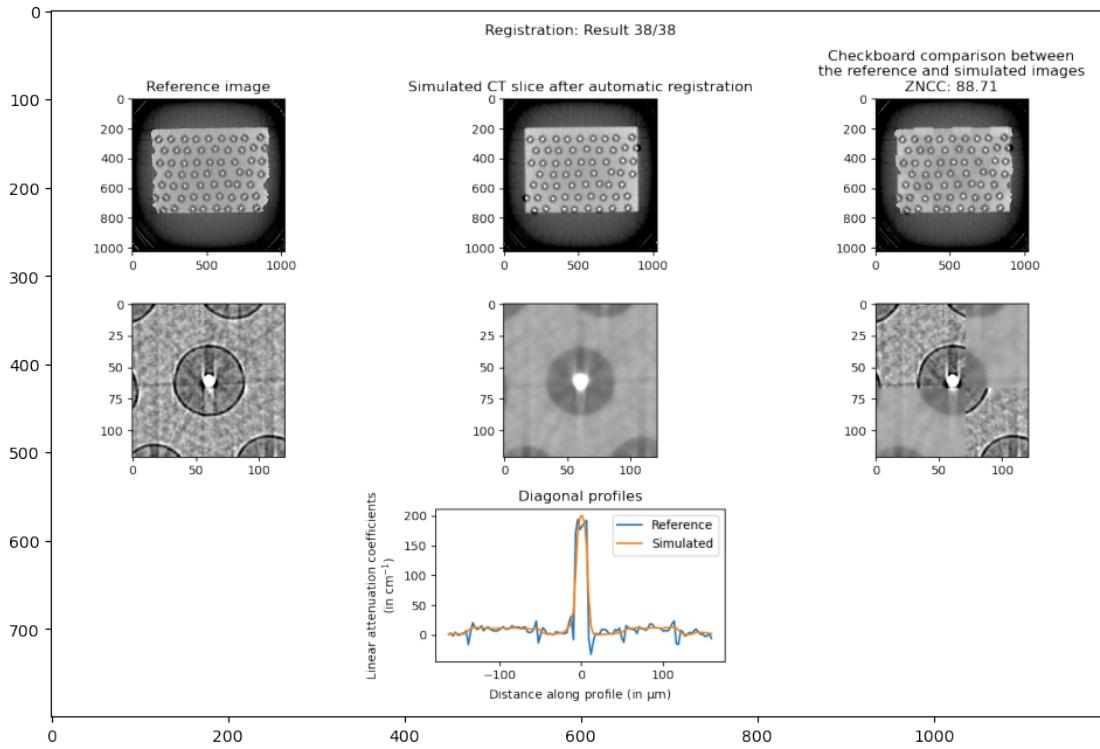
```
[68]: Image(url="plots/fibre1_registration.gif")
```

```
[68]: <IPython.core.display.Image object>
```

View the last frame

```
[69]: plt.imshow(image)
```

[69]: <matplotlib.image.AxesImage at 0x7f3ddac52f10>



1.12.1 Apply the result of the registration

```
[70]: # Load the matrix
setMatrix(matrix_geometry_parameters);

# Load the cores and fibres
setFibres(centroid_set);

gvxr.saveSTLfile("fibre", "outputs/fibre1_fibre.stl");
gvxr.saveSTLfile("core", "outputs/fibre1_core.stl");

print("Core diameter:", round(core_radius * 2), "um");
print("Fibre diameter:", round(fibre_radius * 2), "um");

# Simulate the corresponding CT aquisition
simulated_sinogram, normalised_projections, raw_projections_in_keV = simulateSinogram(sigma_set, k_set, label_set);

# Reconstruct the CT slice
simulated_CT = tomopy.recon(simulated_sinogram,
```

```

        theta_rad,
        center=rot_center,
        sinogram_order=False,
        algorithm='gridrec',
        filter_name='shepp',
        ncore=40)[0];
normalised_simulated_CT = standardisation(simulated_CT);

# Compute the ZNCC
print("ZNCC radii registration 1:",
      "{:.2f}".format(100.0 * np.mean(np.multiply(normalised_reference_CT, ▾
      normalised_simulated_CT))));
```

Core diameter: 16 um
Fibre diameter: 103 um
ZNCC radii registration 1: 88.71

The 3D view of the registration looks like:

[71]: `Image(url='./3d-view.png')`

[71]: <IPython.core.display.Image object>

1.13 Recentre each core/fibre

Each fibre is extracted from both the reference CT slice and simulated CT slice. The displacement between the corresponding fibres is computed to maximise the ZNCC between the two. The centre of the fibre is then adjusted accordingly.

[72]: `def refineCentrePositions(centroid_set, reconstruction_CT_fibres):`

```

# Exhaustive local search to refine the centre of each cylinder
roi_length = 40;
new_centroid_set = [];
for i, cyl in enumerate(centroid_set):

    centre = [
        cyl[0],
        cyl[1]
    ];

    # extract ROI from reference image
    reference_image = copy.deepcopy(reference_CT[centre[1] - roi_length:
      ↵centre[1] + roi_length, centre[0] - roi_length:centre[0] + roi_length]);

    # Normalise ROI
```

```

reference_image = standardisation(reference_image);

best_ZNCC = -1;
best_x_offset = 0;
best_y_offset = 0;

for y in range(-10, 11):
    for x in range(-10, 11):

        centre = [
            cyl[0] + x,
            cyl[1] + y
        ];

        # extract ROI from test image
        test_image = copy.deepcopy(reconstruction_CT_fibres[centre[1] - roi_length:centre[1] + roi_length, centre[0] - roi_length:centre[0] + roi_length]);

        # Normalise ROI
        test_image = standardisation(test_image);

        # Compare the ROIs
        zncc = np.mean(np.multiply(reference_image.flatten(), test_image.flatten()));

        if best_ZNCC < zncc:
            best_ZNCC = zncc;
            best_x_offset = x;
            best_y_offset = y;

        # Correct the position of the centre of the fibre
        new_centroid_set.append([cyl[0] - best_x_offset, cyl[1] - best_y_offset]);

    return new_centroid_set;

```

[73]: centroid_set = refineCentrePositions(centroid_set, normalised_simulated_CT);

1.13.1 Applying the result of recentring

[74]: # Load the matrix
setMatrix(matrix_geometry_parameters);
Load the cores and fibres

```

setFibres(centroid_set);

gvxr.saveSTLfile("fibre", "outputs/fibre2_fibre.stl");
gvxr.saveSTLfile("core", "outputs/fibre2_core.stl");

# Simulate the corresponding CT aquisition
simulated_sinogram, normalised_projections, raw_projections_in_keV = ↵
    simulateSinogram(sigma_set, k_set, label_set);

# Reconstruct the CT slice
simulated_CT = tomopy.recon(simulated_sinogram,
                             theta_rad,
                             center=rot_center,
                             sinogram_order=False,
                             algorithm='gridrec',
                             filter_name='shepp',
                             ncore=40)[0];
normalised_simulated_CT = standardisation(simulated_CT);

# Compute the ZNCC
print("ZNCC recentring registration:",
      "{:.2f}".format(100.0 * np.mean(np.multiply(normalised_reference_CT, ↵
      normalised_simulated_CT))));
```

ZNCC recentring registration: 89.36

1.14 Optimisation the radii after recentring

After recentring the centres, another run of optimisation is executed to refine the radii of the fibres and cores.

[75]: # The registration has already been performed. Load the results.

```

if os.path.isfile("outputs/fibre3_radii.dat"):
    temp = np.loadtxt("outputs/fibre3_radii.dat");
    core_radius = temp[0];
    fibre_radius = temp[1];
# Perform the registration using CMA-ES
else:
    ratio = core_radius / fibre_radius;

    x0 = [fibre_radius, ratio];
    bounds = [[5, 0.01], [1.5 * fibre_radius, 0.95]];

    best_fitness = sys.float_info.max;
    best_fitness_id = 0;
    prefix = "fibre3_";
```

```

opts = cma.CMAOptions()
opts.set('tolfun', 1e-2);
opts['tolx'] = 1e-2;
opts['bounds'] = bounds;

es = cma.CMAEvolutionStrategy(x0, 0.9, opts);
es.optimize(fitnessFunctionFibres);
fibre_radius = es.result.xbest[0];
core_radius = fibre_radius * es.result.xbest[1];

np.savetxt("outputs/fibre3_radii.dat", [core_radius, fibre_radius],  

header='core_radius_in_um,fibre_radius_in_um');

# Release memory
del es;

```

(3_w,6)-aCMA-ES ($\mu_w=2.0, w_1=63\%$) in dimension 2 (seed=857651, Wed Aug 11 17:18:23 2021)

Iterat	#Fevals	function	value	axis	ratio	sigma	min&max	std	t[m:s]
1	6	1.397162791772818e-01	1.0e+00	7.75e-01	6e-01	8e-01	0:08.2		
2	12	2.829841649034579e-01	1.2e+00	9.47e-01	6e-01	1e+00	0:14.9		
3	18	1.312412721152448e-01	2.0e+00	8.07e-01	5e-01	1e+00	0:21.9		
4	24	1.213296684608990e-01	2.1e+00	7.44e-01	5e-01	9e-01	0:29.1		
5	30	1.460846033036106e-01	2.0e+00	6.30e-01	3e-01	7e-01	0:35.8		
6	36	1.220726931919518e-01	2.1e+00	5.92e-01	3e-01	7e-01	0:42.4		
7	42	1.335337579037218e-01	2.7e+00	7.39e-01	3e-01	1e+00	0:49.0		
9	54	1.743395980441586e-01	4.5e+00	4.84e-01	1e-01	7e-01	1:02.2		
11	66	1.216276720300113e-01	4.8e+00	3.91e-01	1e-01	5e-01	1:15.3		
13	78	1.235513576922263e-01	6.9e+00	4.47e-01	7e-02	8e-01	1:28.5		
15	90	1.227070346213213e-01	1.7e+01	7.33e-01	9e-02	2e+00	1:41.7		
17	102	1.217109170133060e-01	2.4e+01	5.28e-01	5e-02	1e+00	1:54.9		
19	114	1.226576045135483e-01	3.6e+01	3.49e-01	2e-02	9e-01	2:08.1		
21	126	1.218677043346403e-01	5.7e+01	2.65e-01	1e-02	7e-01	2:21.6		
24	144	1.222155996383534e-01	9.6e+01	1.94e-01	7e-03	6e-01	2:41.8		
27	162	1.216900698872530e-01	1.3e+02	1.62e-01	4e-03	5e-01	3:01.9		
30	180	1.212588209154846e-01	2.0e+02	2.99e-01	6e-03	1e+00	3:22.3		
33	198	1.212529095953656e-01	2.2e+02	1.81e-01	2e-03	4e-01	3:42.5		
36	216	1.212469697315100e-01	2.6e+02	1.07e-01	8e-04	2e-01	4:02.9		
38	228	1.212455245326126e-01	2.8e+02	8.24e-02	7e-04	1e-01	4:17.1		

```
[76]: if not os.path.exists("plots/fibre3_registration.gif"):
    registration_image_set, image = createAnimation("outputs/
→fibre3_simulated_CT_",
    'plots/fibre3_registration.gif');
```

View the GIF animation (no preview on GitHub .ipynb files)

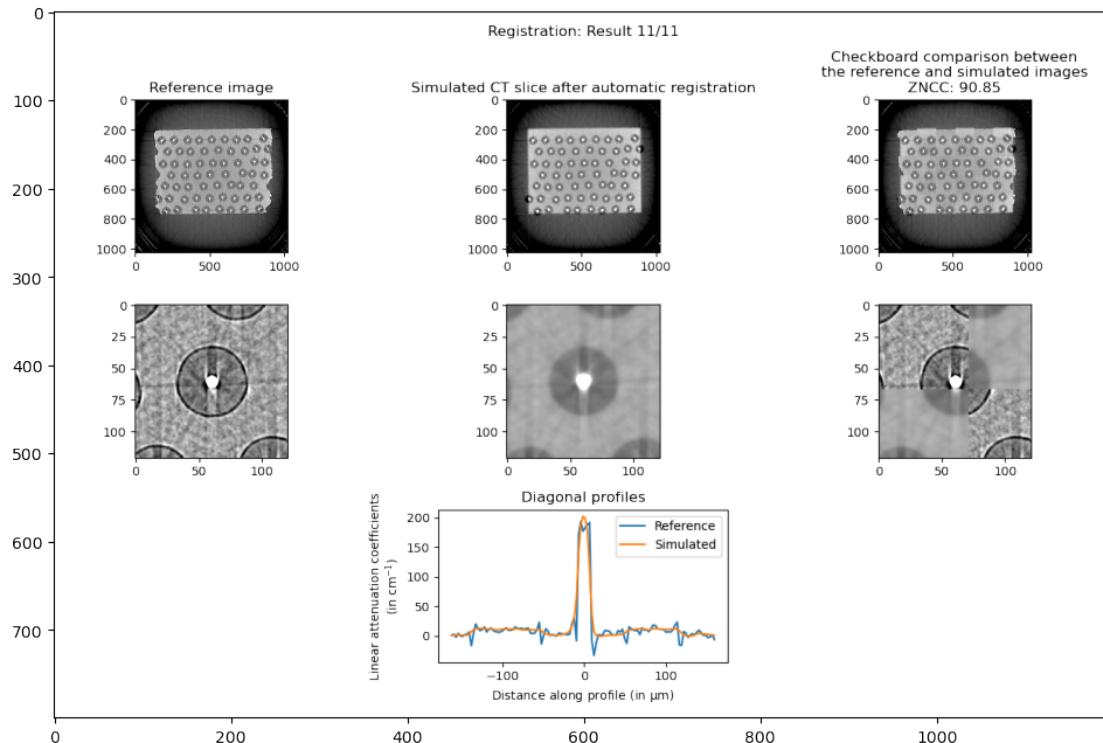
```
[77]: Image(url="plots/fibre3_registration.gif")
```

```
[77]: <IPython.core.display.Image object>
```

View the last frame

```
[78]: plt.imshow(image)
```

```
[78]: <matplotlib.image.AxesImage at 0x7f3dd8a7fac0>
```



1.14.1 Apply the result of the registration

```
[79]: # Load the matrix
setMatrix(matrix_geometry_parameters);

# Load the cores and fibres
setFibres(centroid_set);

gvxr.saveSTLfile("fibre", "outputs/fibre3_fibre.stl");
gvxr.saveSTLfile("core", "outputs/fibre3_core.stl");

print("Core diameter:", round(core_radius * 2), "um");
```

```
print("Fibre diameter:", round(fibre_radius * 2), "um");
```

Core diameter: 16 um
Fibre diameter: 103 um

```
[80]: # Simulate the corresponding CT aquisition
simulated_sinogram, normalised_projections, raw_projections_in_keV =
    ↳simulateSinogram(sigma_set, k_set, label_set);

# Reconstruct the CT slice
simulated_CT = tomopy.recon(simulated_sinogram,
                             theta_rad,
                             center=rot_center,
                             sinogram_order=False,
                             algorithm='gridrec',
                             filter_name='shepp',
                             ncore=40)[0];
normalised_simulated_CT = standardisation(simulated_CT);

# Compute the ZNCC
print("ZNCC radii registration 2:",
      "{:.2f}".format(100.0 * np.mean(np.multiply(normalised_reference_CT,
                                                 ↳normalised_simulated_CT))));
```

ZNCC radii registration 2: 90.85

1.15 Optimisation of the beam spectrum

```
[81]: def fitnessHarmonics(x):

    global energy_spectrum;

    global use_normalisation;

    global best_fitness;
    global best_fitness_id;
    global prefix;

    energy_33_keV = x[0];
    first_order_harmonics = x[1];
    second_order_harmonics = x[2];

    # Normalise the beam spectrum
    total = energy_33_keV + first_order_harmonics + second_order_harmonics;
    energy_33_keV /= total;
    first_order_harmonics /= total;
```

```

second_order_harmonics /= total;

# The beam spectrum. Here we have a polychromatic beam.
gvxr.resetBeamSpectrum();
energy_spectrum = [(33, energy_33_keV, "keV"), (66, first_order_harmonics, "keV"),
                    (99, second_order_harmonics, "keV")];

for energy, percentage, unit in energy_spectrum:
    gvxr.addEnergyBinToSpectrum(energy, unit, percentage);

# Simulate a sinogram
simulated_sinogram, normalised_projections, raw_projections_in_keV =
    simulateSinogram(sigma_set, k_set, label_set);

# Compute the objective value (no normalisation here)
old_normalisation = use_normalisation;
use_normalisation = False;
if use_sinogram:
    objective = metrics(reference_sinogram, simulated_sinogram);
else:
    objective = metrics(reference_normalised_projections,
normalised_projections);
use_normalisation = old_normalisation;

# The block below is not necessary for the registration.
# It is used to save the data to create animations.
if best_fitness > objective:
    best_fitness = objective;

# Reconstruct the CT slice
simulated_CT = tomopy.recon(simulated_sinogram,
                             theta_rad,
                             center=rot_center,
                             sinogram_order=False,
                             algorithm='gridrec',
                             filter_name='shepp',
                             ncore=40)[0];

# Save the simulated sinogram
simulated_sinogram.shape = (simulated_sinogram.size // 2,
                            simulated_sinogram.shape[2], simulated_sinogram.shape[2]);
saveMHA("outputs/" + prefix + "simulated_sinogram_" +
str(best_fitness_id) + ".mha",
        simulated_sinogram,
        [pixel_spacing_in_mm, angular_step, pixel_spacing_in_mm]);

# Save the simulated CT slice

```

```

        saveMHA("outputs/" + prefix + "simulated_CT_" + str(best_fitness_id) + ".mha",
                 simulated_CT,
                 [pixel_spacing_in_mm, pixel_spacing_in_mm, pixel_spacing_in_mm]);

        np.savetxt("outputs/" + prefix + str(best_fitness_id) + ".dat", np.array(x) / total, header='33keV,66keV,99keV');

        best_fitness_id += 1;

    return objective

```

```

[82]: # The registration has already been performed. Load the results.
if os.path.isfile("outputs/spectrum1.dat"):
    temp = np.loadtxt("outputs/spectrum1.dat");

    # The beam spectrum. Here we have a polychromatic beam.
    energy_spectrum = [(33, temp[0], "keV"), (66, temp[1], "keV"), (99, temp[2], "keV")];

# Perform the registration using CMA-ES
else:
    ratio = core_radius / fibre_radius;

    x0 = [0.97, 0.2, 0.1];
    bounds = [[0.0, 0.0, 0.0], [1.0, 1.0, 1.0]];

    best_fitness = sys.float_info.max;
    best_fitness_id = 0;
    prefix = "spectrum1_";

    opts = cma.CMAOptions()
    opts.set('tolfun', 1e-2);
    opts['tolx'] = 1e-2;
    opts['bounds'] = bounds;

    es = cma.CMAEvolutionStrategy(x0, 0.25, opts);
    es.optimize(fitnessHarmonics);

    total = es.result.xbest[0] + es.result.xbest[1] + es.result.xbest[2];
    energy_spectrum = [(33, es.result.xbest[0] / total, "keV"), (66, es.result.xbest[1] / total, "keV"), (99, es.result.xbest[2] / total, "keV")];

    np.savetxt("outputs/spectrum1.dat", [es.result.xbest[0] / total, es.result.xbest[1] / total, es.result.xbest[2] / total], header='weight of main energy, weight of first order harmonics, weight of second order harmonics');

```

```
# Release memory
del es;
```

```
(3_w,7)-aCMA-ES (mu_w=2.3,w_1=58%) in dimension 3 (seed=804237, Wed Aug 11
17:22:50 2021)
Iterat #Fevals    function value  axis ratio   sigma min&max std t[m:s]
 1      7 2.657445985885815e+03 1.0e+00 2.36e-01 2e-01 2e-01 0:09.2
 2     14 1.547085602295766e+03 1.3e+00 1.90e-01 2e-01 2e-01 0:17.4
 3     21 7.753806764553844e+02 1.5e+00 1.70e-01 1e-01 2e-01 0:25.2
 4     28 4.871960904937286e+02 1.5e+00 1.83e-01 1e-01 2e-01 0:33.4
 5     35 5.161828270010666e+02 1.8e+00 1.92e-01 1e-01 2e-01 0:40.9
 6     42 4.298763536256206e+02 1.8e+00 1.66e-01 1e-01 2e-01 0:48.8
 7     49 4.464960307268555e+02 1.7e+00 1.48e-01 7e-02 1e-01 0:56.3
 8     56 4.208880062347967e+02 2.0e+00 1.39e-01 6e-02 1e-01 1:04.1
10    70 5.466637199902613e+02 2.6e+00 1.18e-01 5e-02 1e-01 1:19.2
12   84 4.289769830325816e+02 3.1e+00 9.90e-02 4e-02 1e-01 1:33.8
14   98 4.768015653782354e+02 4.3e+00 8.05e-02 3e-02 8e-02 1:48.7
16  112 4.246221317070134e+02 5.5e+00 6.40e-02 2e-02 6e-02 2:03.7
18  126 4.222919760504752e+02 5.3e+00 4.92e-02 2e-02 4e-02 2:18.8
20  140 4.227664325625593e+02 5.9e+00 3.46e-02 9e-03 3e-02 2:33.8
22  154 4.218591855416741e+02 6.7e+00 2.68e-02 5e-03 2e-02 2:48.6
25  175 4.215517713816474e+02 1.5e+01 2.85e-02 6e-03 3e-02 3:10.6
28  196 4.206550736473947e+02 2.3e+01 2.49e-02 5e-03 3e-02 3:34.1
31  217 4.202160352267793e+02 2.6e+01 2.95e-02 5e-03 3e-02 3:57.9
34  238 4.196228577448755e+02 3.9e+01 3.26e-02 6e-03 4e-02 4:21.5
37  259 4.194933304054617e+02 3.4e+01 3.49e-02 5e-03 3e-02 4:44.7
40  280 4.191565783864131e+02 3.1e+01 3.86e-02 4e-03 4e-02 5:08.2
43  301 4.190216885499082e+02 4.3e+01 6.02e-02 5e-03 7e-02 5:30.9
46  322 4.190261311562708e+02 6.9e+01 4.76e-02 3e-03 7e-02 5:53.9
50  350 4.190403358228058e+02 7.6e+01 3.30e-02 2e-03 4e-02 6:23.9
54  378 4.189568048864857e+02 1.1e+02 2.37e-02 1e-03 3e-02 6:53.9
58  406 4.186925548857706e+02 1.2e+02 3.82e-02 2e-03 3e-02 7:25.1
62  434 4.185991670234870e+02 9.6e+01 5.96e-02 3e-03 4e-02 7:56.6
66  462 4.185436297565211e+02 6.9e+01 5.68e-02 2e-03 3e-02 8:26.4
70  490 4.184839430773974e+02 8.8e+01 5.20e-02 1e-03 3e-02 8:57.4
74  518 4.184573915529784e+02 9.1e+01 6.47e-02 1e-03 2e-02 9:28.6
78  546 4.184611760365249e+02 1.1e+02 5.07e-02 8e-04 2e-02 10:08.4
81  567 4.184465031823069e+02 1.3e+02 3.90e-02 6e-04 1e-02 10:41.2
84  588 4.184446224948985e+02 1.5e+02 3.04e-02 4e-04 9e-03 11:24.6
```

```
[83]: if not os.path.exists("plots/spectrum1_registration.gif"):
    registration_image_set, image = createAnimation("outputs/
→spectrum1_simulated_CT",
                                                'plots/spectrum1_registration.gif');
```

View the GIF animation (no preview on GitHub .ipynb files)

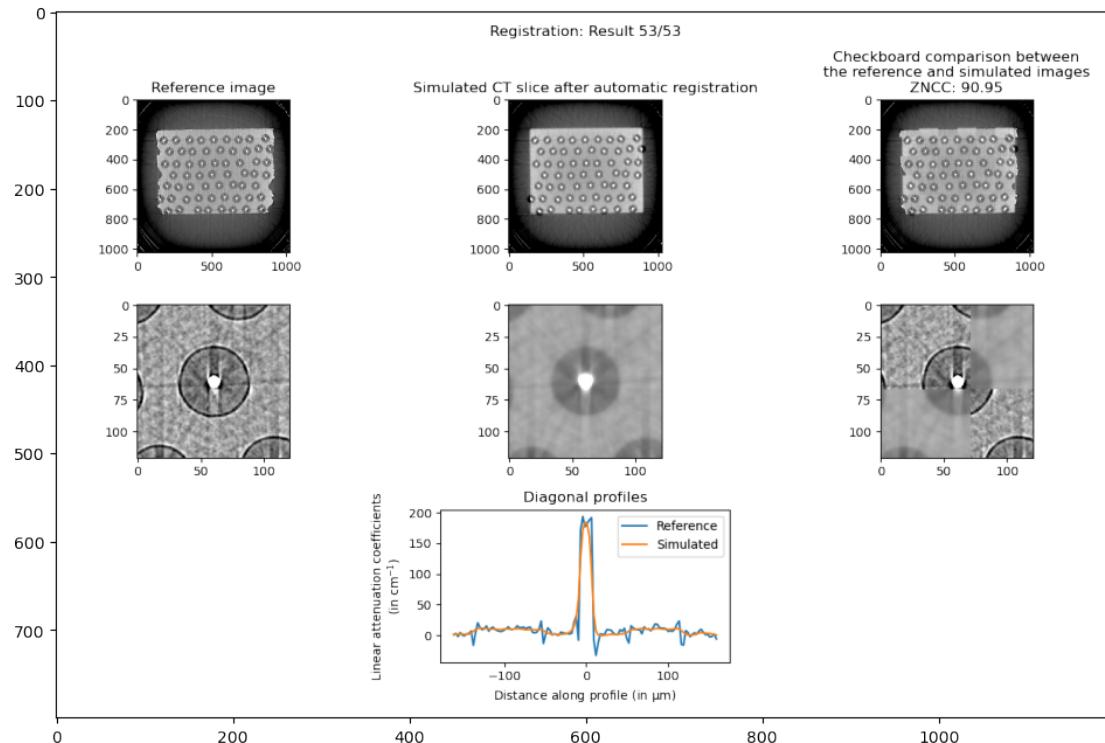
```
[84]: Image(url="plots/spectrum1_registration.gif")
```

```
[84]: <IPython.core.display.Image object>
```

View the last frame

```
[85]: plt.imshow(image)
```

```
[85]: <matplotlib.image.AxesImage at 0x7f3dd3d3f430>
```



```
[86]: # Apply the result of the registration
gvxr.resetBeamSpectrum();
for energy, percentage, unit in energy_spectrum:
    gvxr.addEnergyBinToSpectrum(energy, unit, percentage);
```

```
[87]: for channel in energy_spectrum:
    print(channel);
```

```
(33, 0.962750170494123, 'keV')
(66, 0.0038501802571996205, 'keV')
(99, 0.0333996492486773, 'keV')
```

```
[88]: # Simulate the corresponding CT aquisition
simulated_sinogram, normalised_projections, raw_projections_in_keV =_
    simulateSinogram(sigma_set, k_set, label_set);

# Reconstruct the CT slice
simulated_CT = tomopy.recon(simulated_sinogram,
                             theta_rad,
                             center=rot_center,
                             sinogram_order=False,
                             algorithm='gridrec',
                             filter_name='shepp',
                             ncore=40)[0];
normalised_simulated_CT = standardisation(simulated_CT);

# Compute the ZNCC
print("ZNCC spectrum registration 1:",
      "{:.2f}".format(100.0 * np.mean(np.multiply(normalised_reference_CT,_
    normalised_simulated_CT))));
```

ZNCC spectrum registration 1: 90.95

1.16 Optimisation of the phase contrast and the radii

```
[89]: def laplacian(x, sigma):
    """
    This function create a Laplacian kernel with

    
$$g''(x) = \frac{x^2}{\sigma^4} - \frac{1}{\sigma^2}$$

    
$$\exp(-\frac{x^2}{2\sigma^2})$$


    :param array x:
    :param float sigma:
    :return the convolution kernel
    """

    return (np.power(x, 2.) / math.pow(sigma, 4) - 1. / math.pow(sigma, 2)) *_
        np.exp(-np.power(x, 2.) / (2. * math.pow(sigma, 2)));
```

```
[90]: def getLBuffer(object):
    """
    This function compute the L-buffer of the object over all the angles

    :param str object: the name of the object
    :return the L-buffer over all the angles
    """


```

```

# An empty L-buffer
L_buffer = [] ;

# Get the line of L-buffer for each angle
for angle_id in range(0, number_of_projections):
    gvxr.resetSceneTransformation();
    gvxr.rotateScene(-angular_step * angle_id, 0, 1, 0);

    # Compute the X-ray image
    line_of_L_buffer = np.array(gvxr.computeLBuffer(object));

    # Add the projection
    L_buffer.append(line_of_L_buffer);

# Return as a numpy array
return np.array(L_buffer);

```

```

[91]: def fitnessFunctionLaplacian(x):
    global best_fitness;
    global best_fitness_id;
    global prefix;

    global fibre_radius;
    global core_radius;

    sigma_core = x[0];
    k_core = x[1];
    sigma_fibre = x[2];
    k_fibre = x[3];
    sigma_matrix = x[4];
    k_matrix = x[5];
    core_radius = x[6];
    fibre_radius = x[7];

    # Load the matrix
    setMatrix(matrix_geometry_parameters);

    # Load the cores and fibres
    setFibres(centroid_set);

    # Simulate a sinogram
    simulated_sinogram, normalised_projections, raw_projections_in_keV =simulateSinogram(
        [sigma_core, sigma_fibre, sigma_matrix],
        [k_core, k_fibre, k_matrix],
        ["core", "fibre", "matrix"])

```

```

);

# Compute the objective value
if use_sinogram:
    objective = metrics(reference_sinogram, simulated_sinogram);
else:
    objective = metrics(reference_normalised_projections, □
↪normalised_projections);

# The block below is not necessary for the registration.
# It is used to save the data to create animations.
if best_fitness > objective:
    best_fitness = objective;

# Reconstruct the CT slice
simulated_CT = tomopy.recon(simulated_sinogram,
                             theta_rad,
                             center=rot_center,
                             sinogram_order=False,
                             algorithm='gridrec',
                             filter_name='shepp',
                             ncore=40)[0];

# Save the simulated sinogram
simulated_sinogram.shape = (simulated_sinogram.size //□
↪simulated_sinogram.shape[2], simulated_sinogram.shape[2]);
saveMHA("outputs/" + prefix + "simulated_sinogram_" +□
↪str(best_fitness_id) + ".mha",
        simulated_sinogram,
        [pixel_spacing_in_mm, angular_step, pixel_spacing_in_mm]);

# Save the simulated CT slice
saveMHA("outputs/" + prefix + "simulated_CT_" + str(best_fitness_id) +□
↪".mha",
        simulated_CT,
        [pixel_spacing_in_mm, pixel_spacing_in_mm, □
↪pixel_spacing_in_mm]);

np.savetxt("outputs/" + prefix + str(best_fitness_id) + ".dat", □
↪[sigma_core, k_core, sigma_fibre, k_fibre, sigma_matrix, k_matrix, □
↪core_radius, fibre_radius], header='sigma_core, k_core, sigma_fibre, □
↪k_fibre, sigma_matrix, k_matrix, core_radius_in_um, fibre_radius_in_um');

best_fitness_id += 1;

return objective

```

```
[92]: # The registration has already been performed. Load the results.
if os.path.isfile("outputs/laplacian1.dat"):
    temp = np.loadtxt("outputs/laplacian1.dat");
    sigma_core = temp[0];
    k_core = temp[1];
    sigma_fibre = temp[2];
    k_fibre = temp[3];
    sigma_matrix = temp[4];
    k_matrix = temp[5];
    core_radius = temp[6];
    fibre_radius = temp[7];

# Perform the registration using CMA-ES
else:

    sigma_core = 5.;
    sigma_fibre = 0.75;
    sigma_matrix = 0.6;

    k_core = 1000;
    k_fibre = 1000;
    k_matrix = 1000.0;

    x0 = [
        sigma_core, k_core,
        sigma_fibre, k_fibre,
        sigma_matrix, k_matrix,
        core_radius, fibre_radius
    ];

    bounds = [
        [
            0.005, 0.0,
            0.005, 0.0,
            0.005, 0.0,
            0.95 * core_radius, 0.95 * fibre_radius
        ],
        [
            10.0, 2000,
            2.5, 2000,
            2.5, 2000,
            1.15 * core_radius, 1.15 * fibre_radius
        ]
    ];

    best_fitness = sys.float_info.max;
    best_fitness_id = 0;
```

```

prefix = "laplacian1_";

opts = cma.CMAOptions();
opts.set('tolfun', 1e-4);
opts['tolx'] = 1e-4;
opts['bounds'] = bounds;
opts['CMA_stds'] = [0.25, 20.25, 0.25, 20.25, 0.25, 20.25, core_radius * 0.
˓→1, fibre_radius * 0.1];

es = cma.CMAEvolutionStrategy(x0, 0.25, opts);
es.optimize(fitnessFunctionLaplacian);

sigma_core = es.result.xbest[0];
k_core = es.result.xbest[1];
sigma_fibre = es.result.xbest[2];
k_fibre = es.result.xbest[3];
sigma_matrix = es.result.xbest[4];
k_matrix = es.result.xbest[5];
core_radius = es.result.xbest[6];
fibre_radius = es.result.xbest[7];

np.savetxt("outputs/laplacian1.dat", [sigma_core, k_core, sigma_fibre, ˓→
˓→k_fibre, sigma_matrix, k_matrix, core_radius, fibre_radius], ˓→
˓→header='sigma_core, k_core, sigma_fibre, k_fibre, sigma_matrix, k_matrix, ˓→
˓→core_radius_in_um, fibre_radius_in_um');

# Release memory
del es;

```

(5_w,10)-aCMA-ES ($\mu_w=3.2, w_1=45\%$) in dimension 8 (seed=818251, Wed Aug 11
17:34:52 2021)

Iterat	#Fevals	function	value	axis	ratio	sigma	min&max	std	t[m:s]
1	10	1.128627785320182e-01	1.0e+00	2.44e-01	6e-02	5e+00	0:22.4		
2	20	1.127577655634415e-01	1.2e+00	2.46e-01	6e-02	5e+00	0:45.2		
3	30	1.117705289072551e-01	1.4e+00	2.37e-01	6e-02	5e+00	1:08.2		
4	40	1.110767877201964e-01	1.5e+00	2.14e-01	5e-02	4e+00	1:30.5		
5	50	1.110032970112093e-01	1.5e+00	2.05e-01	5e-02	4e+00	1:53.1		
6	60	1.109513954554905e-01	1.5e+00	2.01e-01	4e-02	4e+00	2:16.4		
7	70	1.109221935247666e-01	1.6e+00	2.05e-01	5e-02	4e+00	2:39.2		
8	80	1.109699000391871e-01	1.7e+00	2.01e-01	4e-02	4e+00	3:01.2		
9	90	1.110527898230884e-01	1.7e+00	2.06e-01	4e-02	4e+00	3:23.8		
10	100	1.110083670338569e-01	1.7e+00	2.23e-01	5e-02	5e+00	3:46.1		
11	110	1.112950586663904e-01	2.0e+00	2.15e-01	4e-02	4e+00	4:08.4		
12	120	1.110456851204059e-01	2.1e+00	1.95e-01	4e-02	4e+00	4:30.9		
13	130	1.111047675055508e-01	2.1e+00	1.80e-01	3e-02	4e+00	4:53.2		
14	140	1.109220228188846e-01	2.2e+00	1.82e-01	3e-02	4e+00	5:15.9		
15	150	1.109134667244638e-01	2.4e+00	1.79e-01	3e-02	4e+00	5:39.1		

16	160	1.109861101562811e-01	2.4e+00	1.66e-01	3e-02	4e+00	5:55.6
18	180	1.109374358548848e-01	2.7e+00	1.66e-01	3e-02	4e+00	6:22.8
19	190	1.108764585808633e-01	2.8e+00	1.59e-01	3e-02	4e+00	6:44.1
20	200	1.108645684680414e-01	2.9e+00	1.51e-01	3e-02	4e+00	7:06.9
21	210	1.108830743479258e-01	2.9e+00	1.41e-01	2e-02	3e+00	7:29.7
22	220	1.108866260290788e-01	3.0e+00	1.31e-01	2e-02	3e+00	7:52.1
23	230	1.108401633765322e-01	3.2e+00	1.25e-01	2e-02	3e+00	8:14.8
24	240	1.108120536496777e-01	3.1e+00	1.33e-01	2e-02	4e+00	8:37.6
25	250	1.108091057869685e-01	3.3e+00	1.45e-01	2e-02	4e+00	9:00.7
27	270	1.107163861967545e-01	3.8e+00	1.95e-01	3e-02	6e+00	9:46.7
29	290	1.105640459628051e-01	4.5e+00	2.42e-01	4e-02	8e+00	10:33.1
31	310	1.104100919779799e-01	5.3e+00	3.76e-01	6e-02	1e+01	11:19.4
33	330	1.100825720219020e-01	5.7e+00	4.87e-01	7e-02	2e+01	12:05.0
35	350	1.103064651907788e-01	5.8e+00	4.80e-01	7e-02	2e+01	12:50.4
37	370	1.096782158836618e-01	5.8e+00	5.01e-01	7e-02	2e+01	13:36.6
39	390	1.095472784975143e-01	6.6e+00	5.27e-01	7e-02	2e+01	14:15.3
41	410	1.092778362588422e-01	7.9e+00	6.13e-01	8e-02	3e+01	15:00.6
43	430	1.090681828306402e-01	9.0e+00	5.89e-01	7e-02	2e+01	15:45.6
45	450	1.085896369094835e-01	9.3e+00	4.97e-01	6e-02	2e+01	16:31.3
47	470	1.085101837709156e-01	1.0e+01	4.58e-01	5e-02	2e+01	17:17.7
49	490	1.081389059321324e-01	1.1e+01	4.56e-01	5e-02	2e+01	18:04.5
51	510	1.081009646291037e-01	1.4e+01	4.97e-01	5e-02	2e+01	18:49.7
53	530	1.076342392279071e-01	1.6e+01	5.08e-01	5e-02	2e+01	19:35.2
55	550	1.074369785376788e-01	1.7e+01	5.39e-01	6e-02	3e+01	20:20.8
57	570	1.074035634960981e-01	1.8e+01	4.57e-01	5e-02	2e+01	21:05.6
60	600	1.071615686113444e-01	2.0e+01	3.44e-01	3e-02	2e+01	21:52.5
64	640	1.068091998172524e-01	2.4e+01	4.78e-01	4e-02	2e+01	22:55.8
66	660	1.068588866198564e-01	2.8e+01	4.45e-01	4e-02	2e+01	23:41.4
68	680	1.068113665878803e-01	3.3e+01	4.64e-01	4e-02	3e+01	24:26.3
70	700	1.067603640321114e-01	3.6e+01	4.42e-01	4e-02	3e+01	25:11.7
72	720	1.067353155739850e-01	3.7e+01	3.65e-01	3e-02	2e+01	25:57.2
75	750	1.067639020182058e-01	3.8e+01	3.20e-01	2e-02	2e+01	27:05.5
78	780	1.067080926284579e-01	3.8e+01	3.04e-01	2e-02	1e+01	28:13.5
81	810	1.066761555739905e-01	4.3e+01	3.08e-01	2e-02	2e+01	29:21.5
84	840	1.066441208414410e-01	4.3e+01	2.46e-01	2e-02	1e+01	30:20.1
87	870	1.066453136884896e-01	3.9e+01	2.01e-01	1e-02	8e+00	31:27.1
90	900	1.066160462269583e-01	3.9e+01	2.09e-01	1e-02	9e+00	32:35.0
93	930	1.065904077563996e-01	3.8e+01	2.79e-01	1e-02	1e+01	33:43.9
96	960	1.065896762864872e-01	4.1e+01	2.89e-01	1e-02	1e+01	34:50.7
99	990	1.065642906580179e-01	5.2e+01	2.45e-01	1e-02	1e+01	35:57.4
100	1000	1.065609401073685e-01	5.2e+01	2.41e-01	1e-02	1e+01	36:20.3
105	1050	1.065210142541943e-01	5.6e+01	2.08e-01	9e-03	8e+00	37:23.1
109	1090	1.065055389221411e-01	7.8e+01	2.42e-01	1e-02	1e+01	38:22.4
114	1140	1.064813097275918e-01	1.0e+02	2.34e-01	1e-02	1e+01	39:29.0
119	1190	1.064518154461284e-01	1.2e+02	2.66e-01	1e-02	1e+01	40:45.5
123	1230	1.064450749109974e-01	1.5e+02	3.26e-01	1e-02	2e+01	41:53.2
129	1290	1.064335830424190e-01	1.7e+02	3.20e-01	1e-02	2e+01	43:05.9
135	1350	1.064159526850854e-01	1.7e+02	2.89e-01	9e-03	1e+01	44:18.2

```

141 1410 1.063899467350692e-01 1.6e+02 3.31e-01 9e-03 2e+01 45:31.1
147 1470 1.063845416892303e-01 1.6e+02 4.12e-01 1e-02 2e+01 46:43.8
149 1490 1.063941809658485e-01 1.7e+02 3.57e-01 8e-03 2e+01 47:07.8

```

```
[93]: if not os.path.exists("plots/laplacian1_registration.gif"):
    registration_image_set, image = createAnimation("outputs/
    ↴laplacian1_simulated_CT_",
    'plots/laplacian1_registration.gif');
```

View the GIF animation (no preview on GitHub .ipynb files)

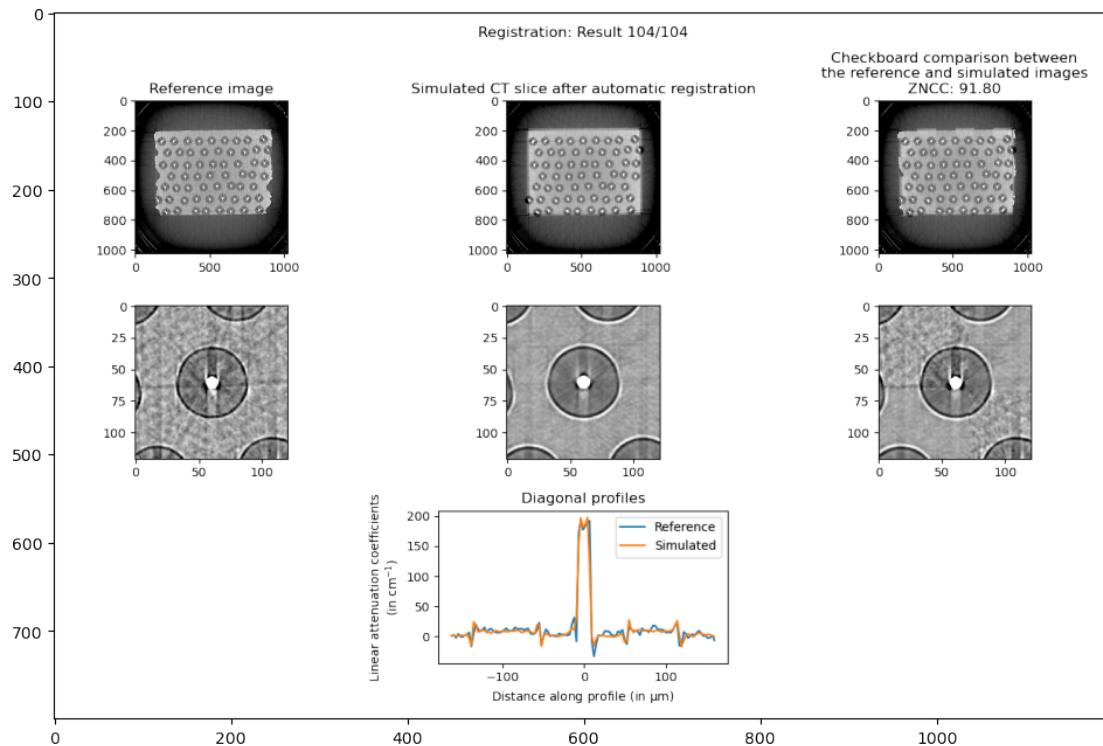
```
[94]: Image(url="./plots/laplacian1_registration.gif")
```

```
[94]: <IPython.core.display.Image object>
```

View the last frame

```
[95]: plt.imshow(image)
```

```
[95]: <matplotlib.image.AxesImage at 0x7f3dc1f65e80>
```



1.16.1 Apply the result of the registration

```
[96]: # Load the matrix
setMatrix(matrix_geometry_parameters);

# Load the cores and fibres
setFibres(centroid_set);

gvxr.saveSTLfile("fibre", "outputs/laplacian1_fibre.stl");
gvxr.saveSTLfile("core", "outputs/laplacian1_core.stl");

print("Core diameter:", round(core_radius * 2), "um");
print("Fibre diameter:", round(fibre_radius * 2), "um");
```

Core diameter: 16 um
Fibre diameter: 107 um

```
[97]: # Simulate the corresponding CT aquisition
sigma_set = [sigma_core, sigma_fibre, sigma_matrix];
k_set = [k_core, k_fibre, k_matrix];
label_set = ["core", "fibre", "matrix"];

simulated_sinogram, normalised_projections, raw_projections_in_keV =
    simulateSinogram(sigma_set, k_set, label_set);

# Reconstruct the CT slice
simulated_CT = tomopy.recon(simulated_sinogram,
                             theta_rad,
                             center=rot_center,
                             sinogram_order=False,
                             algorithm='gridrec',
                             filter_name='shepp',
                             ncore=40)[0];
normalised_simulated_CT = standardisation(simulated_CT);

# Compute the ZNCC
print("ZNCC phase contrast registration 1:",
      "{:.2f}".format(100.0 * np.mean(np.multiply(normalised_reference_CT,
                                                 normalised_simulated_CT))));
```

ZNCC phase contrast registration 1: 93.18

1.17 Optimisation of the phase contrast and the LSF

```
[98]: old_lsf = copy.deepcopy(lsf_kernel);
```

```
[99]: def fitnessFunctionLaplacianLSF(x):
    global best_fitness;
    global best_fitness_id;
    global prefix;

    global lsf_kernel;

    # sigma_core = x[0];
    k_core = x[0];
    # sigma_fibre = x[2];
    k_fibre = x[1];
    # sigma_matrix = x[4];
    k_matrix = x[2];

    b2 = x[3];
    c2 = x[4];
    e2 = x[5];
    f2 = x[6];

    # The response of the detector as the line-spread function (LSF)
    t = np.arange(-20., 21., 1.);
    lsf_kernel=lsf(t*41, b2, c2, e2, f2);
    lsf_kernel/=lsf_kernel.sum();

    # Simulate a sinogram
    simulated_sinogram, normalised_projections, raw_projections_in_keV =simulateSinogram(
        [sigma_core, sigma_fibre, sigma_matrix],
        [k_core, k_fibre, k_matrix],
        ["core", "fibre", "matrix"]
    );

    # Compute the objective value
    if use_sinogram:
        objective = metrics(reference_sinogram, simulated_sinogram);
    else:
        objective = metrics(reference_normalised_projections,normalised_projections);

    # The block below is not necessary for the registration.
    # It is used to save the data to create animations.
    if best_fitness > objective:
        best_fitness = objective;

    # Reconstruct the CT slice
    simulated_CT = tomopy.recon(simulated_sinogram,
                                theta_rad,
```

```

        center=rot_center,
        sinogram_order=False,
        algorithm='gridrec',
        filter_name='shepp',
        ncore=40)[0];

    # Save the simulated sinogram
    simulated_sinogram.shape = (simulated_sinogram.size //_
→simulated_sinogram.shape[2], simulated_sinogram.shape[2]);
    saveMHA("outputs/" + prefix + "simulated_sinogram_" +_
→str(best_fitness_id) + ".mha",
            simulated_sinogram,
            [pixel_spacing_in_mm, angular_step, pixel_spacing_in_mm]);

    # Save the simulated CT slice
    saveMHA("outputs/" + prefix + "simulated_CT_" + str(best_fitness_id) +_
→".mha",
            simulated_CT,
            [pixel_spacing_in_mm, pixel_spacing_in_mm,_
→pixel_spacing_in_mm]);

    np.savetxt("outputs/" + prefix + "laplacian_" + str(best_fitness_id) +_
→".dat", [k_core, k_fibre, k_matrix], header='k_core, k_fibre, k_matrix');
    np.savetxt("outputs/" + prefix + "LSF_" + str(best_fitness_id) + "._
→dat", [b2, c2, e2, f2], header='b2, c2, e2, f2');

    best_fitness_id += 1;

    return objective;

```

```

[100]: # The registration has already been performed. Load the results.
if os.path.isfile("outputs/laplacian2.dat") and os.path.isfile("outputs/lsf2.
→dat"):
    temp = np.loadtxt("outputs/laplacian2.dat");
    k_core = temp[0];
    k_fibre = temp[1];
    k_matrix = temp[2];

    temp = np.loadtxt("outputs/lsf2.dat");
    b2 = temp[0];
    c2 = temp[1];
    e2 = temp[2];
    f2 = temp[3];

# Perform the registration using CMA-ES

```

```

else:

    b2 = 54.9359;
    c2 = -3.58452;
    e2 = 6.32561e+09;
    f2 = 1.0;

    x0 = [
        k_core,
        k_fibre,
        k_matrix,
        b2, c2, e2, f2
    ];

    bounds = [
        [
            k_core-500,
            k_fibre-500,
            k_matrix-500,
            b2 - b2 / 4.,
            c2 + c2 / 4.,
            e2 - e2 / 4.,
            f2 - f2/ 4.
        ],
        [
            k_core+500,
            k_fibre+500,
            k_matrix+500,
            b2 + b2 / 4.,
            c2 - c2 / 4.,
            e2 + e2 / 4.,
            f2 + f2/ 4.
        ]
    ];

```

best_fitness = sys.float_info.max;
 best_fitness_id = 0;
 prefix = "laplacian2_"

```

opts = cma.CMAOptions()
opts.set('tolfun', 1e-4);
opts['tolx'] = 1e-4;
opts['bounds'] = bounds;
#opts['seed'] = 987654321;
# opts['maxiter'] = 5;
opts['CMA_stds'] = [1250 * 0.2, 1250 * 0.2, 1250 * 0.2,
    b2 * 0.2, -c2 * 0.2, e2 * 0.2, f2 * 0.2];

```

```

es = cma.CMAEvolutionStrategy(x0, 0.25, opts);
es.optimize(fitnessFunctionLaplacianLSF);

k_core = es.result.xbest[0];
k_fibre = es.result.xbest[1];
k_matrix = es.result.xbest[2];

b2 = es.result.xbest[3];
c2 = es.result.xbest[4];
e2 = es.result.xbest[5];
f2 = es.result.xbest[6];

np.savetxt("outputs/laplacian2.dat", [k_core, k_fibre, k_matrix], □
←header='k_core, k_fibre, k_matrix');
np.savetxt("outputs/lst2.dat", [b2, c2, e2, f2], header='b2, c2, e2, f2');

# Release memory
del es;

```

(4_w,9)-aCMA-ES ($\mu_w=2.8, w_1=49\%$) in dimension 7 (seed=739029, Wed Aug 11
18:23:07 2021)

Iterat	#Fevals	function	value	axis	ratio	sigma	min&max	std	t[m:s]
1	9	1.065718826341313e-01	1.0e+00	2.19e-01	4e-02	3e+08	0:11.1		
2	18	1.065347508428139e-01	1.2e+00	2.07e-01	4e-02	2e+08	0:21.9		
3	27	1.062627385578101e-01	1.3e+00	1.99e-01	4e-02	2e+08	0:32.7		
4	36	1.061839301954330e-01	1.4e+00	1.98e-01	4e-02	2e+08	0:43.5		
5	45	1.060179154615556e-01	1.5e+00	2.00e-01	4e-02	2e+08	0:54.0		
6	54	1.061024402607409e-01	1.6e+00	1.84e-01	4e-02	2e+08	1:04.1		
7	63	1.061278422545926e-01	1.7e+00	1.71e-01	4e-02	2e+08	1:14.4		
8	72	1.059865839821377e-01	1.7e+00	1.59e-01	4e-02	2e+08	1:25.2		
9	81	1.059098690799496e-01	1.8e+00	1.61e-01	4e-02	2e+08	1:35.8		
10	90	1.058751838664796e-01	1.8e+00	1.85e-01	5e-02	2e+08	1:46.6		
11	99	1.057957056064688e-01	2.2e+00	2.12e-01	6e-02	2e+08	1:57.7		
13	117	1.058362531632262e-01	2.9e+00	2.38e-01	7e-02	3e+08	2:18.2		
15	135	1.057260557244933e-01	3.6e+00	2.60e-01	8e-02	3e+08	2:40.1		
17	153	1.057921617970607e-01	4.0e+00	2.42e-01	8e-02	3e+08	3:00.9		
19	171	1.056599034169730e-01	4.4e+00	1.84e-01	6e-02	2e+08	3:21.6		
21	189	1.055709083247843e-01	4.5e+00	1.72e-01	5e-02	2e+08	3:42.5		
23	207	1.055598500263744e-01	4.9e+00	1.56e-01	5e-02	2e+08	4:03.5		
25	225	1.055371673516115e-01	5.8e+00	1.56e-01	5e-02	2e+08	4:24.5		
27	243	1.055416933602332e-01	6.2e+00	1.39e-01	4e-02	2e+08	4:45.1		
29	261	1.055446827082976e-01	6.6e+00	1.16e-01	4e-02	1e+08	5:05.3		
31	279	1.055397395553711e-01	6.3e+00	1.09e-01	3e-02	1e+08	5:26.3		
33	297	1.055405193942963e-01	7.2e+00	9.72e-02	3e-02	1e+08	5:47.5		
36	324	1.055316386524157e-01	8.5e+00	7.73e-02	2e-02	9e+07	6:18.8		
39	351	1.055311912693103e-01	9.2e+00	7.94e-02	3e-02	9e+07	6:49.5		

```

42    378 1.055298021293029e-01 9.2e+00 7.28e-02 2e-02 9e+07 7:21.0
45    405 1.055310577700169e-01 9.9e+00 7.76e-02 2e-02 1e+08 7:52.4
48    432 1.055297526468582e-01 1.1e+01 7.29e-02 2e-02 1e+08 8:23.5
51    459 1.055302051926598e-01 1.2e+01 5.51e-02 1e-02 9e+07 8:54.2
52    468 1.055295590885513e-01 1.3e+01 6.45e-02 2e-02 1e+08 9:04.8

```

```
[101]: if not os.path.exists("plots/laplacian2_registration.gif"):
    registration_image_set, image = createAnimation("outputs/
→laplacian2_simulated_CT_",
    'plots/laplacian2_registration.gif');
```

View the GIF animation (no preview on GitHub .ipynb files)

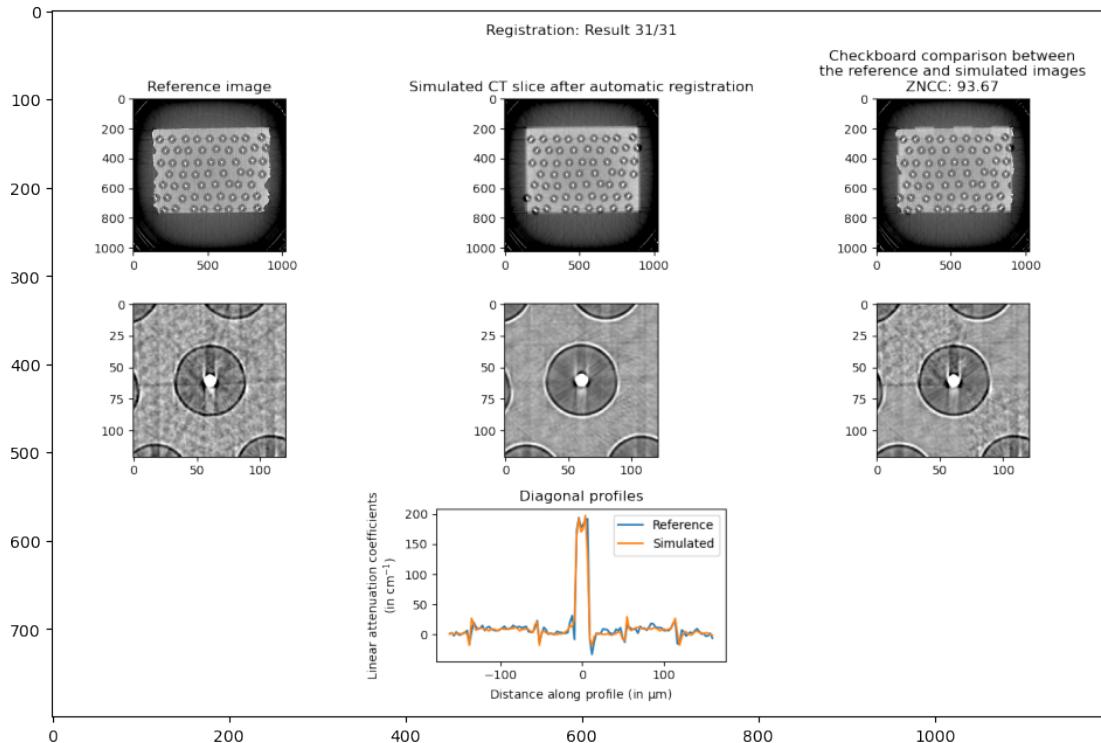
```
[102]: Image(url="plots/laplacian2_registration.gif")
```

```
[102]: <IPython.core.display.Image object>
```

View the last frame

```
[103]: plt.imshow(image)
```

```
[103]: <matplotlib.image.AxesImage at 0x7f3dd97085b0>
```



1.17.1 Apply the result of the registration

```
[104]: # The response of the detector as the line-spread function (LSF)
t = np.arange(-20., 21., 1.);
lsf_kernel=lsf(t*41, b2, c2, e2, f2);
lsf_kernel/=lsf_kernel.sum();
np.savetxt("outputs/LSF_optimised.txt", lsf_kernel);

[105]: # Simulate the corresponding CT aquisition
sigma_set = [sigma_core, sigma_fibre, sigma_matrix];
k_set = [k_core, k_fibre, k_matrix];
label_set = ["core", "fibre", "matrix"];

simulated_sinogram, normalised_projections, raw_projections_in_keV =
    simulateSinogram(sigma_set, k_set, label_set);

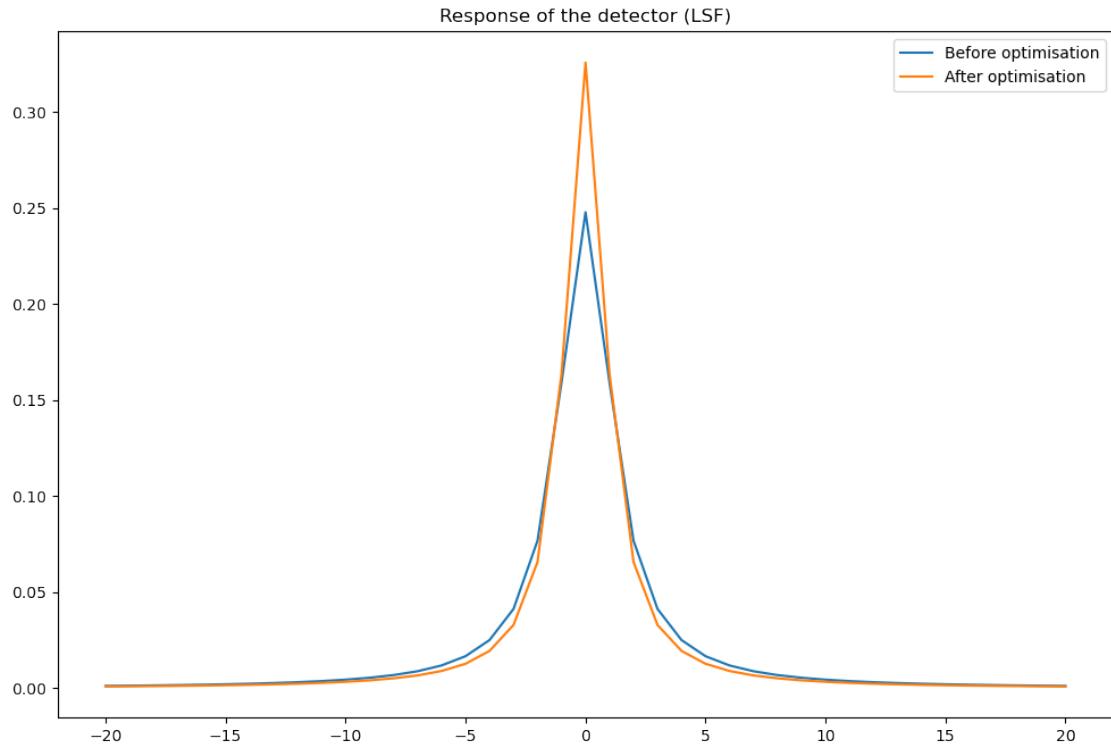
# Reconstruct the CT slice
simulated_CT = tomopy.recon(simulated_sinogram,
                             theta_rad,
                             center=rot_center,
                             sinogram_order=False,
                             algorithm='gridrec',
                             filter_name='shepp',
                             ncore=40)[0];
normalised_simulated_CT = standardisation(simulated_CT);

offset1 = 86;
offset2 = reference_CT.shape[0] - offset1;
profile_test_whole_image_without_Poisson_noise = copy.deepcopy(np.
    diag(simulated_CT[offset1:offset2, offset1:offset2]));

# Compute the ZNCC
print("ZNCC phase contrast and LSF registration:",
      "{:.2f}".format(100.0 * np.mean(np.multiply(normalised_reference_CT,
                                                normalised_simulated_CT))));
```

ZNCC phase contrast and LSF registration: 93.67

```
[106]: fig=plt.figure();
plt.title("Response of the detector (LSF)");
plt.plot(t, old_lsf, label="Before optimisation");
plt.plot(t, lsf_kernel, label="After optimisation");
plt.legend();
plt.savefig('plots/LSF_optimised.pdf');
plt.savefig('plots/LSF_optimised.png');
```



1.17.2 Extract the fibre in the centre of the CT slices

```
[107]: def findFibreInCentreOfCtSlice():
    global centroid_set;
    global reference_CT;
    global cylinder_position_in_centre_of_slice;

    # Find the cylinder in the centre of the image
    cylinder_position_in_centre_of_slice = None;
    best_distance = sys.float_info.max;

    for centre in centroid_set:
        distance = math.pow(centre[0] - reference_CT.shape[1] / 2, 2) + math.
        ↪pow(centre[1] - reference_CT.shape[0] / 2, 2);

        if best_distance > distance:
            best_distance = distance;
            cylinder_position_in_centre_of_slice = copy.deepcopy(centre);

    return cylinder_position_in_centre_of_slice;
```

```
[108]: findFibreInCentreOfCtSlice();

reference_fibre_in_centre = np.array(copy.
→deepcopy(reference_CT[cylinder_position_in_centre_of_slice[1] - roi_length:
→cylinder_position_in_centre_of_slice[1] + roi_length, ↴
→cylinder_position_in_centre_of_slice[0] - roi_length:
→cylinder_position_in_centre_of_slice[0] + roi_length]));
test_fibre_in_centre      = np.array(copy.
→deepcopy(simulated_CT[cylinder_position_in_centre_of_slice[1] - roi_length:
→cylinder_position_in_centre_of_slice[1] + roi_length, ↴
→cylinder_position_in_centre_of_slice[0] - roi_length:
→cylinder_position_in_centre_of_slice[0] + roi_length]));

profile_reference = copy.deepcopy(np.diag(reference_fibre_in_centre));
profile_test_without_Poisson_noise = copy.deepcopy(np.
→diag(test_fibre_in_centre));

reference_fibre_in_centre = standardisation(reference_fibre_in_centre);
test_fibre_in_centre = standardisation(test_fibre_in_centre);
```

```
[109]: norm = cm.colors.Normalize(vmax=1.25, vmin=-0.5)

fig, (ax1, ax2, ax3) = plt.subplots(1, 3)
plt.tight_layout()
fig.suptitle('Fibre in the centre of the CT slices')

ax1.set_title("Reference image");
imgplot1 = ax1.imshow(reference_fibre_in_centre, cmap="gray",
                      norm=norm);

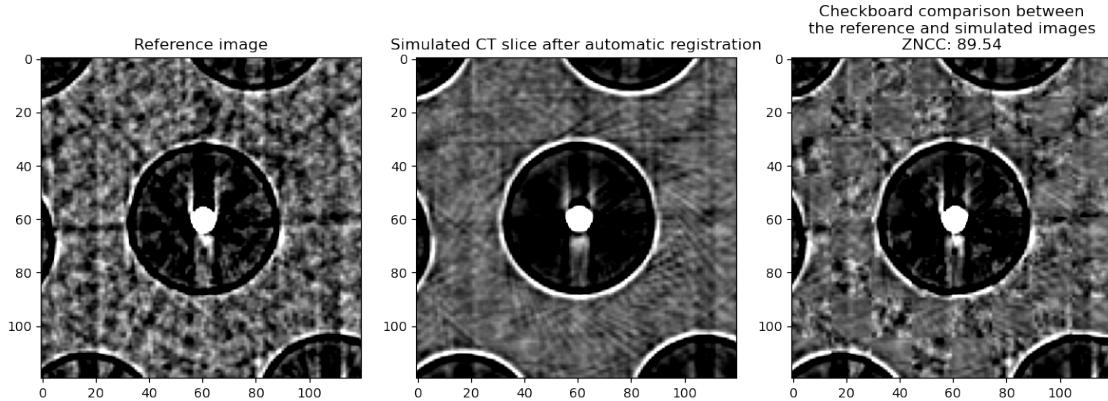
ax2.set_title("Simulated CT slice after automatic registration");
imgplot2 = ax2.imshow(test_fibre_in_centre,
                      cmap='gray',
                      norm=norm);

comp_equalized = compare_images(reference_fibre_in_centre, ↴
→test_fibre_in_centre, method='checkerboard');
ax3.set_title("Checkboard comparison between\n" +
              "the reference and simulated images\nZNCC: " +
              "{:.2f}".format(100.0 * np.mean(np.
→multiply(reference_fibre_in_centre, test_fibre_in_centre))));

imgplot3 = ax3.imshow(comp_equalized,
                      cmap='gray',
                      norm=norm);

plt.savefig('plots/Fibre_in_centre_CT_slices_before_noise.pdf');
plt.savefig('plots/Fibre_in_centre_CT_slices_before_noise.png');
```

Fibre in the centre of the CT slices



1.18 Optimisation of the Poisson noise

```
[110]: def fitnessFunctionNoise(x):
    global best_fitness;
    global best_fitness_id;
    global prefix;

    bias = x[0];
    gain = x[1];
    scale = x[2];

    # Poisson noise
    map = (normalised_projections_ROI + (bias + 1)) * gain;
    temp = np.random.poisson(map).astype(float);
    temp /= gain;
    temp -= bias + 1;

    # Noise map
    noise_map = normalised_projections_ROI - temp;
    noise_map *= scale;
    noisy_image = normalised_projections_ROI + noise_map;

    # Compute the standard deviation of the pixel values in the ROI extracted
    ↪from the simulated image with noise
    noisy_image_noise_ROI_stddev = 0;
    for y in range(noisy_image.shape[0]):
        noisy_image_noise_ROI_stddev += noisy_image[y].std();
```

```

noisy_image_noise_ROI_stddev /= noisy_image.shape[0];

# Difference of std dev between the reference and the simulated image
diff = reference_noise_ROI_stddev - noisy_image_noise_ROI_stddev;
objective = diff * diff;

# The block below is not necessary for the registration.
# It is used to save the data to create animations.
if best_fitness > objective:
    best_fitness = objective;

# Save the simulated CT slice
saveMHA("outputs/" + prefix + "noisy_image_" + str(best_fitness_id) + ".  

→mha",
        noisy_image,
        [pixel_spacing_in_mm, pixel_spacing_in_mm,  

→pixel_spacing_in_mm]);

np.savetxt("outputs/" + prefix + str(best_fitness_id) + ".dat", [bias,  

→gain, scale], header='bias, gain, scale');

best_fitness_id += 1;

return objective

```

```

[111]: # The registration has already been performed. Load the results.
if os.path.isfile("outputs/poisson-noise.dat"):
    temp = np.loadtxt("outputs/poisson-noise.dat");
    bias = temp[0];
    gain = temp[1];
    scale = temp[2];

# Perform the registration using CMA-ES
else:

# Extract a ROI from the reference where no object is
reference_noise_ROI = copy.deepcopy(reference_normalised_projections[450:  

→550,0:125]);

saveMHA("outputs/reference_noise_ROI.mha",
        reference_noise_ROI,
        [pixel_spacing_in_mm, angular_step, pixel_spacing_in_mm]);

# Compute the standard deviation of the pixel values in the ROI extracted  

→from the reference
reference_noise_ROI_stddev = 0;
for y in range(reference_noise_ROI.shape[0]):

```

```

    reference_noise_ROI_stddev += reference_noise_ROI[y].std();
reference_noise_ROI_stddev /= reference_noise_ROI.shape[0];

# Copy the simulated projection in a temporary variable
temp = copy.deepcopy(normalised_projections);
temp.shape = reference_normalised_projections.shape

# Extract the corresponding ROI
normalised_projections_ROI = temp[450:550,0:125];

saveMHA("outputs/normalised_projections_ROI.mha",
        normalised_projections_ROI,
        [pixel_spacing_in_mm, angular_step, pixel_spacing_in_mm]);

# Initialise the values
bias = 0.0;
gain = 255.0;
scale = 1;

x0 = [bias, gain, scale];

bounds = [
    [-1.0, 0.0, 0.0],
    [ 5.0, 255.0, 255.0]
];

opts = cma.CMAOptions()
opts.set('tolfun', 1e-8);
opts['tolx'] = 1e-8;
opts['bounds'] = bounds;
opts['CMA_stds'] = [1, 10, 10];

best_fitness = sys.float_info.max;
best_fitness_id = 0;
prefix = "poisson-noise_";

es = cma.CMAEvolutionStrategy(x0, 0.25, opts);
es.optimize(fitnessFunctionNoise);

bias = es.result.xbest[0];
gain = es.result.xbest[1];
scale = es.result.xbest[2];

np.savetxt("outputs/poisson-noise.dat", [bias, gain, scale], header='bias,'
           'gain, scale');

# Release memory

```

```
del es;
```

```
(3_w,7)-aCMA-ES (mu_w=2.3,w_1=58%) in dimension 3 (seed=857594, Wed Aug 11  
18:32:35 2021)  
Iterat #Fevals function value axis ratio sigma min&max std t[m:s]  
1 7 2.982920191919707e-06 1.0e+00 2.24e-01 2e-01 2e+00 0:00.0  
2 14 6.353849645182939e-04 1.3e+00 2.07e-01 2e-01 2e+00 0:00.0  
3 21 1.233589661382649e-03 1.3e+00 2.32e-01 2e-01 2e+00 0:00.1  
71 497 1.379917233179754e-12 1.2e+03 3.72e-02 9e-04 3e-01 0:01.2
```

```
[112]: print("Noise parameters: ", bias, gain, scale)
```

```
Noise parameters: -0.9755832910752753 253.37403431706466 0.07949086560131374
```

1.18.1 Apply the result of the optimisation

```
[113]: # Simulate the corresponding CT aquisition
```

```
simulated_sinogram, normalised_projections, raw_projections_in_keV =  
    simulateSinogram(sigma_set, k_set, label_set);
```

```
[114]: # Reconstruct the CT slice
```

```
simulated_CT = tomopy.recon(simulated_sinogram,  
                             theta_rad,  
                             center=rot_center,  
                             sinogram_order=False,  
                             algorithm='gridrec',  
                             filter_name='shepp',  
                             ncore=40)[0];  
normalised_simulated_CT = standardisation(simulated_CT);  
profile_test_whole_image_with_Poisson_noise = copy.deepcopy(np.  
    diag(simulated_CT[offset1:offset2, offset1:offset2]));  
  
# Compute the ZNCC  
print("ZNCC noise registration:",  
      "{:.2f}".format(100.0 * np.mean(np.multiply(normalised_reference_CT,  
                                         normalised_simulated_CT))));
```

```
ZNCC noise registration: 93.02
```

```
[115]: norm = cm.colors.Normalize(vmax=1.25, vmin=-0.5)
```

```
fig, (ax1, ax2, ax3) = plt.subplots(1, 3)  
plt.tight_layout()  
fig.suptitle('CT slice with fibres after the registration')  
ax1.set_title("Reference image");
```

```

imgplot1 = ax1.imshow(normalised_reference_CT, cmap="gray",
                      norm=norm);

ax2.set_title("Simulated CT slice after automatic registration");
imgplot2 = ax2.imshow(normalised_simulated_CT,
                      cmap='gray',
                      norm=norm);

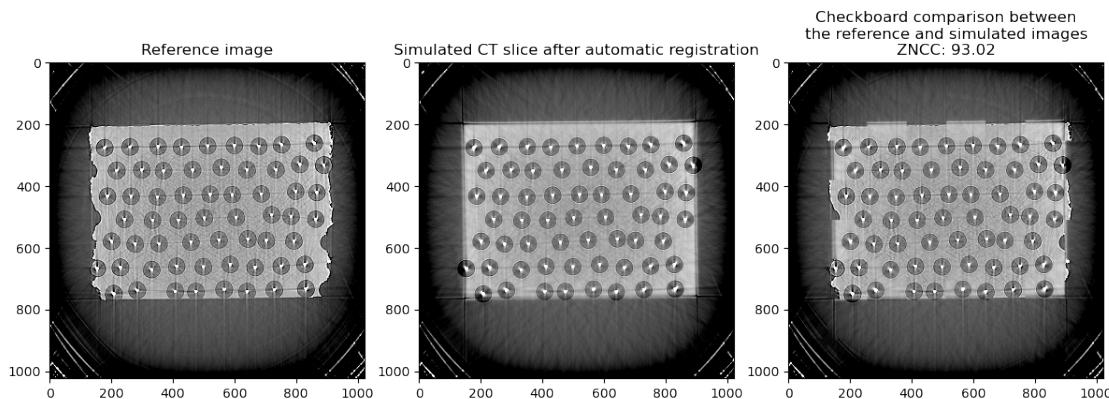
comp_equalized = compare_images(normalised_reference_CT, □
→normalised_simulated_CT, method='checkerboard');
ax3.set_title("Checkboard comparison between\n" +
              "the reference and simulated images\nZNCC: " +
              "{:.2f}".format(100.0 * np.mean(np.
→multiply(normalised_reference_CT, normalised_simulated_CT))));

imgplot3 = ax3.imshow(comp_equalized,
                      cmap='gray',
                      norm=norm);

plt.savefig('plots/Fibre_in_centre_CT_slices_after_noise.pdf');
plt.savefig('plots/Fibre_in_centre_CT_slices_after_noise.png');

```

CT slice with fibres after the registration



```

[116]: test_fibre_in_centre      = np.array(copy.
→deepcopy(simulated_CT[cylinder_position_in_centre_of_slice[1] - roi_length:
→cylinder_position_in_centre_of_slice[1] + roi_length, □
→cylinder_position_in_centre_of_slice[0] - roi_length:
→cylinder_position_in_centre_of_slice[0] + roi_length]));

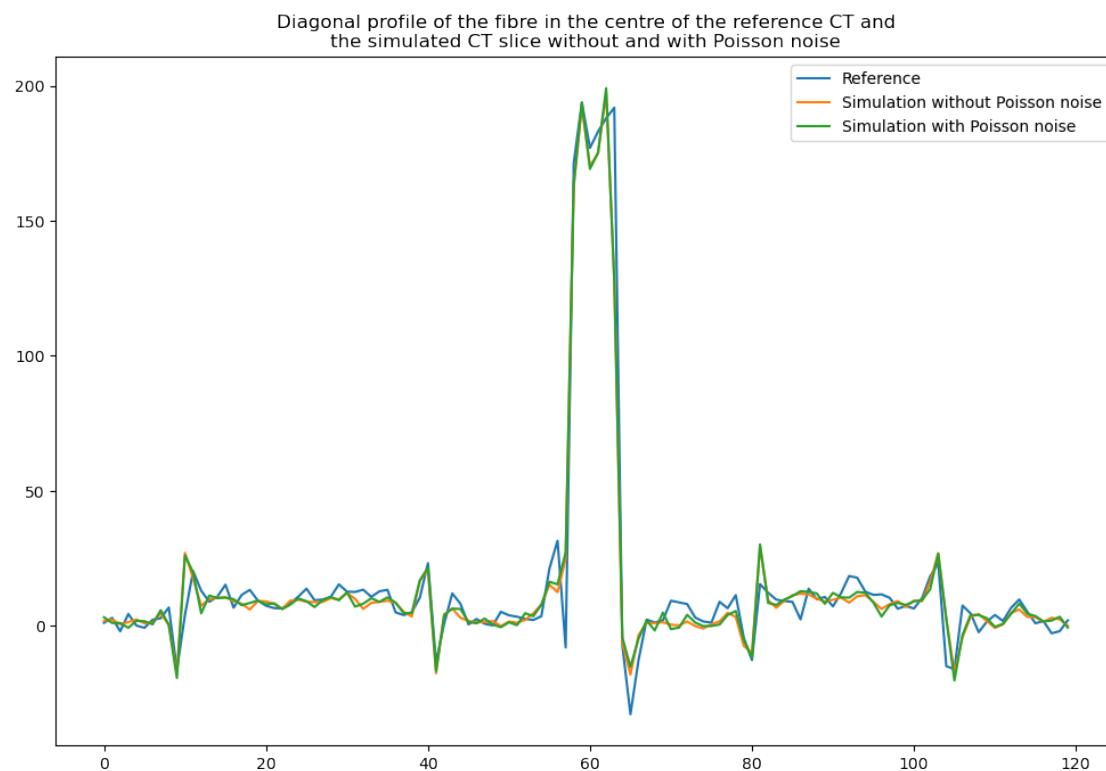
profile_test_with_Poisson_noise = copy.deepcopy(np.diag(test_fibre_in_centre));

```

```
plt.figure()

plt.title("Diagonal profile of the fibre in the centre of the reference CT\u
↪and\nthe simulated CT slice without and with Poisson noise")
plt.plot(profile_reference, label="Reference");
plt.plot(profile_test_without_Poisson_noise, label="Simulation without Poisson\u
↪noise");
plt.plot(profile_test_with_Poisson_noise, label="Simulation with Poisson\u
↪noise");
plt.legend();

plt.savefig('plots/profiles.pdf');
plt.savefig('plots/profiles.png');
```



1.19 Results in terms of linear attenuation coefficients

Reduce the ROI size to focus on a single fibre and its surrounding matrix

```
[117]: roi length = 40
```

Extract the ROIs

```
[118]: reference_fibre_in_centre = np.array(copy.  
    ↪deepcopy(reference_CT[cylinder_position_in_centre_of_slice[1] - roi_length:  
    ↪cylinder_position_in_centre_of_slice[1] + roi_length, □  
    ↪cylinder_position_in_centre_of_slice[0] - roi_length:  
    ↪cylinder_position_in_centre_of_slice[0] + roi_length]));  
test_fibre_in_centre = np.array(copy.  
    ↪deepcopy(simulated_CT[cylinder_position_in_centre_of_slice[1] - roi_length:  
    ↪cylinder_position_in_centre_of_slice[1] + roi_length, □  
    ↪cylinder_position_in_centre_of_slice[0] - roi_length:  
    ↪cylinder_position_in_centre_of_slice[0] + roi_length]));
```

Save the ROIs

```
[119]: saveMHA("outputs/reference_fibre_in_centre.mha", reference_fibre_in_centre, □  
    ↪[pixel_spacing_in_mm, pixel_spacing_in_mm]);  
saveMHA("outputs/test_fibre_in_centre.mha", test_fibre_in_centre, □  
    ↪[pixel_spacing_in_mm, pixel_spacing_in_mm]);
```

A function to create a circular binary mask

```
[120]: def create_circular_mask(h, w, center=None, radius=None):  
  
    if center is None: # use the middle of the image  
        center = (int(w/2), int(h/2))  
    if radius is None: # use the smallest distance between the center and image  
    ↪walls  
        radius = min(center[0], center[1], w-center[0], h-center[1])  
  
    Y, X = np.ogrid[:h, :w]  
    dist_from_center = np.sqrt((X - center[0])**2 + (Y-center[1])**2)  
  
    mask = dist_from_center <= radius  
    return np.array(mask, dtype=bool);
```

A function to create the binary masks for the core, fibre and matrix

```
[121]: def createMasks(mask_shape):  
    fibre_radius_in_px = fibre_radius / pixel_spacing_in_micrometre  
    core_radius_in_px = core_radius / pixel_spacing_in_micrometre  
  
    core_mask = create_circular_mask(mask_shape[1], mask_shape[0], None, □  
    ↪core_radius_in_px);  
  
    fibre_mask = create_circular_mask(mask_shape[1], mask_shape[0], None, □  
    ↪fibre_radius_in_px);  
    matrix_mask = np.logical_not(fibre_mask);  
  
    #fibre_mask = np.subtract(fibre_mask, core_mask);
```

```

fibre_mask = np.bitwise_xor(fibre_mask, core_mask);

#TypeError: numpy boolean subtract, the ` - ` operator, is not supported, use
→the bitwise_xor, the ` ^ ` operator, or the logical_xor function instead.

return core_mask, fibre_mask, matrix_mask

```

Create binary masks for the core, fibre and matrix

```
[122]: mask_shape = reference_fibre_in_centre.shape;
core_mask, fibre_mask, matrix_mask = createMasks(mask_shape);

core_mask = ndimage.binary_erosion(core_mask).astype(core_mask.dtype);

for i in range(4):
    fibre_mask = ndimage.binary_erosion(fibre_mask).astype(fibre_mask.dtype);
    matrix_mask = ndimage.binary_erosion(matrix_mask, border_value=1).
    ↪astype(matrix_mask.dtype);

core_mask.shape = [core_mask.shape[0], core_mask.shape[1]]
fibre_mask.shape = [fibre_mask.shape[0], fibre_mask.shape[1]]
matrix_mask.shape = [matrix_mask.shape[0], matrix_mask.shape[1]]
```

Save the binary masks

```
[123]: saveMHA("outputs/core_mask.mha", core_mask.astype(np.uint8), ↪
    [pixel_spacing_in_mm, pixel_spacing_in_mm, pixel_spacing_in_mm]);
saveMHA("outputs/fibre_mask.mha", fibre_mask.astype(np.uint8), ↪
    [pixel_spacing_in_mm, pixel_spacing_in_mm, pixel_spacing_in_mm]);
saveMHA("outputs/matrix_mask.mha", matrix_mask.astype(np.uint8), ↪
    [pixel_spacing_in_mm, pixel_spacing_in_mm, pixel_spacing_in_mm]);
```

Display the masks

```
[124]: norm = cm.colors.Normalize(vmax=1, vmin=0)

fig, (ax1, ax2, ax3) = plt.subplots(1, 3)
plt.tight_layout()
fig.suptitle('Binary mask for every structure')

ax1.set_title("W core");
imgplot1 = ax1.imshow(core_mask, cmap="gray",
                      norm=norm);

ax2.set_title("SiC fibre");
imgplot2 = ax2.imshow(fibre_mask,
                      cmap='gray',
                      norm=norm);
```

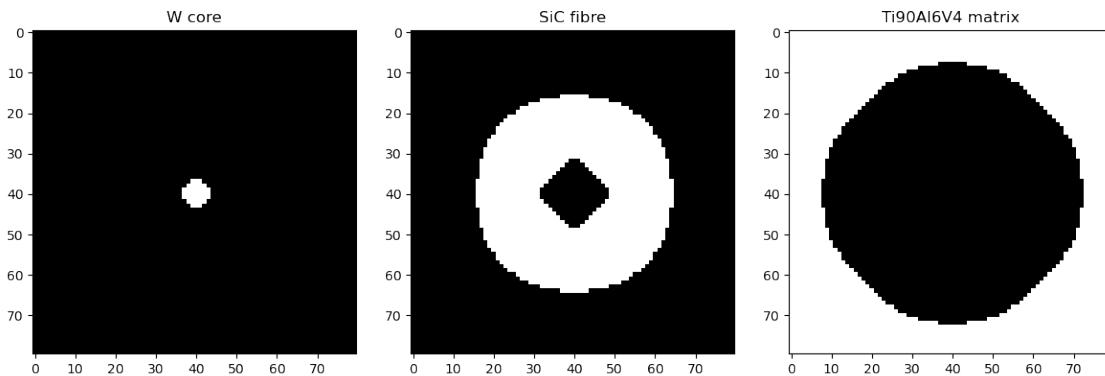
```

ax3.set_title("Ti90Al6V4 matrix");
imgplot3 = ax3.imshow(matrix_mask,
                      cmap='gray',
                      norm=norm);

plt.savefig('plots/masks.pdf');
plt.savefig('plots/masks.png');

```

Binary mask for every structure



A function to collect all the μ statistics from the masks.

```

[125]: def getMuStatistics(reference_fibre_in_centre, test_fibre_in_centre, core_mask,
                        fibre_mask, matrix_mask):

    data = [];
    index = np.nonzero(core_mask);

    data.append(["Theoretical",
                 "Core",
                 "W",
                 341.61,
                 341.61,
                 341.61,
                 0.0]);

    data.append(["Experimental",
                 "Core",
                 "W",
                 np.min(reference_fibre_in_centre[index]),

```

```

        np.max(reference_fibre_in_centre[index]),
        np.mean(reference_fibre_in_centre[index]),
        np.std(reference_fibre_in_centre[index]))];

data.append(["Simulated",
            "Core",
            "W",
            np.min(test_fibre_in_centre[index]),
            np.max(test_fibre_in_centre[index]),
            np.mean(test_fibre_in_centre[index]),
            np.std(test_fibre_in_centre[index]))];

index = np.nonzero(fibre_mask);

data.append(["Theoretical",
            "Fibre",
            "SiC",
            2.736,
            2.736,
            2.736,
            0.0]);

data.append(["Experimental",
            "Fibre",
            "SiC",
            np.min(reference_fibre_in_centre[index]),
            np.max(reference_fibre_in_centre[index]),
            np.mean(reference_fibre_in_centre[index]),
            np.std(reference_fibre_in_centre[index]))];

data.append(["Simulated",
            "Fibre",
            "SiC",
            np.min(test_fibre_in_centre[index]),
            np.max(test_fibre_in_centre[index]),
            np.mean(test_fibre_in_centre[index]),
            np.std(test_fibre_in_centre[index]))];

index = np.nonzero(matrix_mask);
data.append(["Theoretical",
            "Matrix",
            "Ti90Al6V4",
            13.1274,
            13.1274,
            13.1274,
            0.0]);

```

```

data.append(["Experimental",
             "Matrix",
             "Ti90Al6V4",
             np.min(reference_fibre_in_centre[index]),
             np.max(reference_fibre_in_centre[index]),
             np.mean(reference_fibre_in_centre[index]),
             np.std(reference_fibre_in_centre[index]));

data.append(["Simulated",
             "Matrix",
             "Ti90Al6V4",
             np.min(test_fibre_in_centre[index]),
             np.max(test_fibre_in_centre[index]),
             np.mean(test_fibre_in_centre[index]),
             np.std(test_fibre_in_centre[index]]);

return pd.DataFrame(data,
                     index=None,
                     columns=['CT', 'Structure', "Composition", 'min', 'max', 'mean', ↴
                     'stddev'])

```

Get the dataframe with all the values and display it as a table

```
[126]: df = getMuStatistics(reference_fibre_in_centre, test_fibre_in_centre, ↴
                           core_mask, fibre_mask, matrix_mask);

display(df)
```

	CT	Structure	Composition	min	max	mean	\
0	Theoretical	Core	W	341.610000	341.610000	341.610000	
1	Experimental	Core	W	120.325165	212.735718	188.644455	
2	Simulated	Core	W	123.703079	207.287216	185.433456	
3	Theoretical	Fibre	SiC	2.736000	2.736000	2.736000	
4	Experimental	Fibre	SiC	-32.709011	31.958403	3.217891	
5	Simulated	Fibre	SiC	-17.909393	21.187803	2.500488	
6	Theoretical	Matrix	Ti90Al6V4	13.127400	13.127400	13.127400	
7	Experimental	Matrix	Ti90Al6V4	-3.869796	25.127810	10.856583	
8	Simulated	Matrix	Ti90Al6V4	-0.209934	18.534582	9.812553	
			stddev				
0	0.000000						
1	16.044945						
2	16.951956						
3	0.000000						
4	6.269166						
5	4.004930						
6	0.000000						
7	4.368093						

```
8    2.854526
```

1.20 Save the CAD models and plot them in 3D

```
[127]: gvxr.saveSTLfile("fibre", "outputs/final_SiC_fibres.stl");
gvxr.saveSTLfile("core", "outputs/final_W_cores.stl");
gvxr.saveSTLfile("matrix", "outputs/final_Ti90Al6V4_matrix.stl");
```

```
[128]: np.savetxt("outputs/centroids.dat", centroid_set);
```

Load the STL files

```
[129]: fibre_mesh = mesh.Mesh.from_file('outputs/final_SiC_fibres.stl')
core_mesh = mesh.Mesh.from_file('outputs/final_W_cores.stl')
matrix_mesh = mesh.Mesh.from_file('outputs/final_Ti90Al6V4_matrix.stl')
```

Create the k3d geometries

```
[130]: matrix_geometry = k3d.mesh(matrix_mesh.vectors.flatten(),
                                range(int(matrix_mesh.vectors.flatten().shape[0] / 3)),
                                color=0x7fc97f,
                                wireframe=False,
                                flat_shading=False,
                                name="Ti90Al6V4 Matrix",
                                opacity = 0.4)

fibres_geometry = k3d.mesh(fibre_mesh.vectors.flatten(),
                           range(int(fibre_mesh.vectors.flatten().shape[0] / 3)),
                           color=0xbeaed4,
                           wireframe=False,
                           flat_shading=False,
                           name="SiC Fibres",
                           opacity = 0.9)

cores_geometry = k3d.mesh(core_mesh.vectors.flatten(),
                          range(int(core_mesh.vectors.flatten().shape[0] / 3)),
                          color=0xfdc086,
                          wireframe=False,
                          flat_shading=False,
                          name="W Cores",
                          opacity = 0.4)
```

Create a new plot

```
[131]: plot = k3d.plot()
plot.background_color = 0xffffffff
```

```
plot += matrix_geometry  
plot += fibres_geometry  
plot += cores_geometry  
  
plot.display()
```

Output()

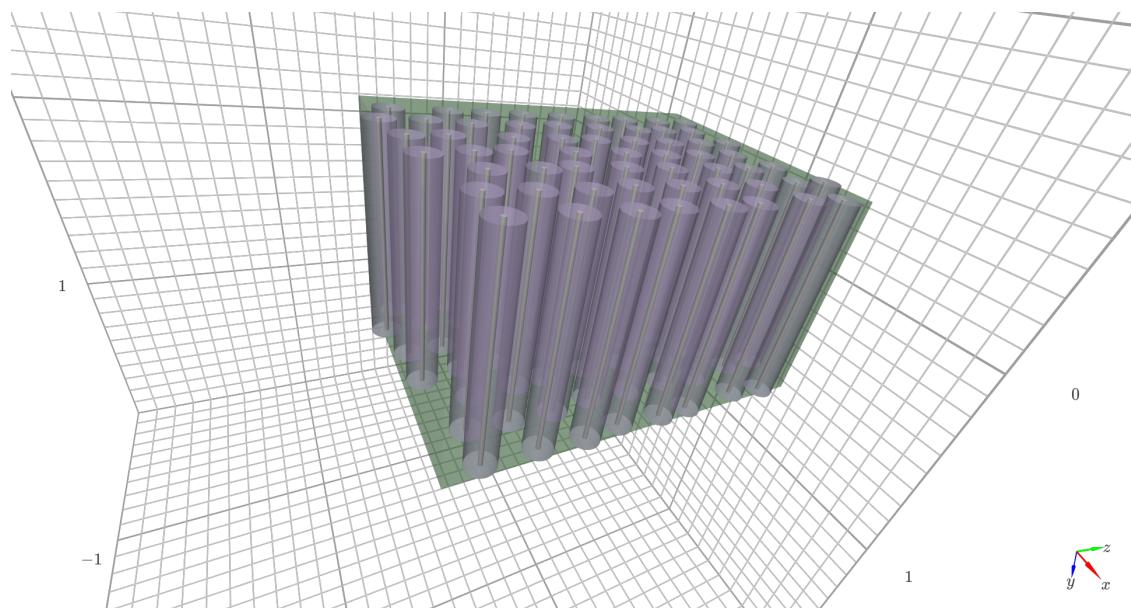
Get a screenshot

```
[132]: plot.fetch_screenshot()  
  
data = base64.b64decode(plot.screenshot)  
with open('outputs/CAD_models_3d.png','wb') as fp:  
    fp.write(data)
```

Save the figure

```
[133]: Image(filename='outputs/CAD_models_3d.png')
```

```
[133]:
```



```
[ ]:
```