# Description of Source Data

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### 1 Structure of source Data

Results are structured in the file:

/vol/biomedic3/bglocker/nnUNet

```
1 bglocker biomedia 236 Sep 24 15:16 exports
-rwxr-xr-x
drwxr-sr-x
             9 bglocker biomedia
                                   9 Nov 25 10:55 nnUNet_preprocessed
drwxr-sr-x
                                  10 Nov 25 10:50 nnUNet_raw
             9 bglocker biomedia
drwxr-sr-x
             9 bglocker biomedia
                                   9 Nov 25 12:20 nnUNet_results
drwxr-sr-x
            11 bglocker biomedia
                                  11 Dec 16 09:10 nnUNet_testing
-rw-r--r-
             1 bglocker biomedia 644 Oct 20 07:20 run_nnunet_0.sh
             1 bglocker biomedia 644 Oct 20 07:20 run_nnunet_1.sh
-rw-r---r--
             1 bglocker biomedia 644 Oct 20 07:20 run_nnunet_2.sh
-rw-r-r-
             1 bglocker biomedia 644 Oct 20 07:21 run_nnunet_3.sh
-rw-r-r--
-rw-r---r--
             1 bglocker biomedia 644 Oct 20 07:21 run_nnunet_4.sh
```

### nnUNet\_raw

nnUNet\_raw has the original (training) images with manual annotations. Each Dataset below is treated as a binary segmentation problem. See Section2

```
drwxr-sr-x
            4 bglocker biomedia
                                  5 Sep 17 13:47 Dataset001_Anorectum
drwxr-sr-x
            3 bglocker biomedia
                                  5 Sep 17 20:24 Dataset002_Bladder
            3 bglocker biomedia
drwxr-sr-x
                                  5 Sep 17 20:27 Dataset003_CTVn
drwxr-sr-x
            3 bglocker biomedia
                                  5 Sep 17 20:28 Dataset004_CTVp
                                  5 Sep 17 20:29 Dataset005_Parametrium
drwxr-sr-x
            3 bglocker biomedia
            1 bglocker biomedia 135 Nov 25 10:50 note
-rw-r-r-
```

### What is a Binary Segmentation Problem?

## 2 Viewing the Data

#### **ItkSnap**

The viewing tool used is ItkSnap, which was developed as an open source tool for viewing medical imaging scans. The view (Figure 1) shows how you would see input data.

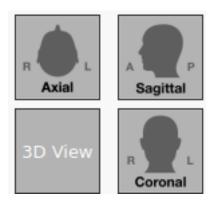


Figure 1: view of all input data

With that we can use this tool to view input data. Here, the R and L stand for right and left respectively, and the A and P stand for Anterior and Posterior. We can provide a few other examples of viewing data displayed below in Figure 2. We are further provided with manual annotation of the substructures. Figure 3 shows an example of the annotation of the Anorectum. You can enable the 3D visual model through Edit > 3D Panel > Toggle 3D view.

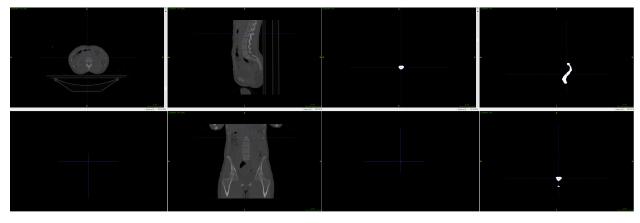


Figure 2: ItkSnap view of the Anorectum Raw Image

Figure 3: ItkSnap view of the Anorectum Raw Image