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Question 1

```
mypath = "megray.tif";

[me,melut] = imread(mypath);
figure;
subplot(1,1,1);
imshow(me);
```



```
me_info = imfinfo(mypath);
me_info
me_info = struct with fields:
                     Filename: 'C:\Users\LENOVO\Desktop\PSUT\1st sem - Y4\Digital Image processing\images\megray.ti
                 FileModDate: '13-Nov-2023 10:39:17'
                     FileSize: 66250
                      Format: 'tif'
               FormatVersion: []
                       Width: 256
                      Height: 256
                     BitDepth: 8
                   ColorType: 'grayscale'
              FormatSignature: [73 73 42 0]
                   ByteOrder: 'little-endian'
              NewSubFileType: 0
               BitsPerSample: 8
                  Compression: 'PackBits'
   PhotometricInterpretation: 'BlackIsZero'
                StripOffsets: [8 8274 16534 24799 33059 41326 49532 57792]
              SamplesPerPixel: 1
                 RowsPerStrip: 32
              StripByteCounts: [8266 8260 8265 8260 8267 8206 8260 8204]
                 XResolution: 72
```

```
YResolution: 72
ResolutionUnit: 'Inch'
Colormap: []
PlanarConfiguration: 'Chunky'
TileWidth: []
TileLength: []
TileOffsets: []
TileByteCounts: []
Orientation: 1
FillOrder: 1
GrayResponseUnit: 0.0100
MaxSampleValue: 255
MinSampleValue: 0
Thresholding: 1
Offset: 65996
```

Question 2

```
handxray_path = "handxray.tif";

handxray = imread(handxray_path);
figure;
subplot(1,1,1);
imshow(handxray);
```



```
hand_info = imfinfo(handxray_path);
```

hand_info

```
hand_info = struct with fields:
                     Filename: 'C:\Users\LENOVO\Desktop\PSUT\1st sem - Y4\Digital Image processing\images\handxray.
                  FileModDate: '13-Nov-2023 10:40:19'
                     FileSize: 154210
                       Format: 'tif'
                FormatVersion: []
                        Width: 390
                       Height: 476
                     BitDepth: 8
                    ColorType: 'grayscale'
              FormatSignature: [73 73 42 0]
                    ByteOrder: 'little-endian'
               NewSubFileType: 0
                BitsPerSample: 8
                  Compression: 'PackBits'
    PhotometricInterpretation: 'BlackIsZero'
                 StripOffsets: [8 6715 12298 18149 24969 32065 39496 46288 53188 60393 67735 75353 82680 90157 9793
              SamplesPerPixel: 1
                 RowsPerStrip: 21
              StripByteCounts: [6707 5583 5851 6820 7096 7431 6792 6900 7205 7342 7618 7327 7477 7776 7776 6563 641
                  XResolution: 72
                  YResolution: 72
               ResolutionUnit: 'Inch'
                     Colormap: []
          PlanarConfiguration: 'Chunky'
                   TileWidth: []
                   TileLength: []
                  TileOffsets: []
               TileByteCounts: []
                  Orientation: 1
                    FillOrder: 1
             GrayResponseUnit: 0.0100
               MaxSampleValue: 255
               MinSampleValue: 0
                 Thresholding: 1
                       Offset: 153836
```

Show a better contrast image

```
enhancedImg = histeq(handxray);
```

Displaying the difference in the image

```
figure;
subplot(2, 2, 1);
imshow(handxray);
title('Original X-ray Image');

subplot(2, 2, 2);
imshow(enhancedImg);
title('Contrast Enhanced X-ray Image');

subplot(2, 2, 3);
imhist(handxray);
title('Histogram of Original Image');

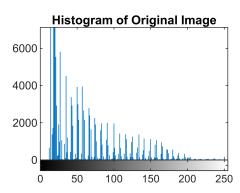
subplot(2, 2, 4);
```

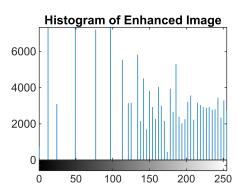
Original X-ray Image



Contrast Enhanced X-ray Image







We can see that the original image is a normal xray image with no enhancment, after applying histeq() function, which shift the pixel values to be more uniform, since the original image has a lot of values consentrated between 0 and 200, after adjusting, the values are shifted from 100 to 255 which resulted in a brighter image with more details and now we can see the edges of the hand and more defined bones.