

1. Navigate to main.py
2. Pass appropriate parameters
 - a. -p: path to dicome files
 - b. -s: settings file
 - c. -u: user name
 - d. -o (optional): output path
3. GUI interface
 - a. Use number keypad to select anatomical label
 - b. Click and drag right mouse button for
 - c. Select location of anatomical label to scroll
 - d. Can also use up and down arrows to scroll
 - i. Label will be reset if clicked again
 - e. Press tab to exit and save

Note: may need to resize cmd window to allow appropriate carriage returns

```
C:\Users\Kevin\Documents\Programming\puml_lablr\puml_lablr>git branch
C:\Users\Kevin\Documents\Programming\puml_lablr\puml_lablr>
C:\Users\Kevin\Documents\Programming\puml_lablr\puml_lablr>
C:\Users\Kevin\Documents\Programming\puml_lablr\puml_lablr>
C:\Users\Kevin\Documents\Programming\puml_lablr\puml_lablr>
C:\Users\Kevin\Documents\Programming\puml_lablr\puml_lablr>
C:\Users\Kevin\Documents\Programming\puml_lablr\puml_lablr>main.py -p data/ex2 -s settings/settings.yaml -u Test
```