# Toxin Screening <u>Labeling System</u> <u>Installation and Usage Guide</u>:

# **Installing Anaconda**:

To install, go to <a href="https://www.continuum.io/downloads">https://www.continuum.io/downloads</a>. Follow the download and installation guide for Windows.

You may find this anaconda cheat sheet helpful: <a href="https://conda.io/docs/using/cheatsheet.html">https://conda.io/docs/using/cheatsheet.html</a>

### **Anaconda First Time Use:**

- 1. Navigate to Documents directory
- **2.** Create a new anaconda environment: conda create -n labeltool python=2.7 anaconda
- 3. Activate the environment *activate labeltool*
- 4. Install Django conda install -c anaconda django=1.10.5
- 5. Install OpenCV 3 conda install --channel https://conda.anaconda.org/menpo opencv3

## **Anaconda General Use**:

- 1. Navigate to Documents folder
- 2. Activate the environment activate labeltool
- 2. When finished, deactivate the environment

#### deactivate

## **Labeling System First Time Use**:

- 1, Navigate to the Documents folder
- 2. Start up anaconda (refer to instructions above)
- 3. Download source code:

git clone <a href="https://github.com/aarong4743/">https://github.com/aarong4743/</a>
toxinscreeninglabelingsystem.git

## **Labeling System General Use:**

- 1, Navigate to Documents folder
- 2. Start up anaconda (refer to instructions above)
- 3. Navigate to myproject folder *cd myproject*
- 4. Fire up local server python manage.py runserver
- 5. Go to local host specified in command prompt plus myapp and welcome <a href="http://127.0.0.1:8000/myapp/welcome">http://127.0.0.1:8000/myapp/welcome</a>
- 6. Select "Make a Training Set" or "Make a Test Set"
- 7. Name your set. Be sure that you are not naming it something that already exists
- 8. If you selected training, select all of the files in your training set to upload
  - 1. Click "Upload"
  - 2. Click "Start Labeling"

- 3. Iterate through each image, and enter the number of eyes that you see for the worm in the image. Please enter 0, 1, 2, or -1 if you cannot tell or the number of eyes is different.
- 4. When you are done, it will take you to a page that says "Done Labeling". Click "Home" to go back to the original welcome page.
- 9. If you selected test, select all of the files in your first well to upload
  - 1. Click "Upload"
  - 2. Click "Start Labeling"
  - 3. Look at the list of all of the images in the well, and enter a general consensus on the number of eyes you see for the worm in the well. Please enter 0, 1, 2, or -1 if you cannot tell or the number of eyes is different.
  - 4. Select all of the files in the next well to upload
  - 5. Repeat sub-steps 1-3 of step 9
  - 6. Once you have labeled your last well, click "Finished"
  - 7. Click "Home" to go back to the original welcome page.
- 10. Press "Ctrl + C" to terminate the local server
- 11. Deactivate the conda environment (see instructions above)
- 12. Open up File Explorer
- 13. Navigate to the project foler.
- 14. Go into "myproject"
- 15. Go into "media"

- 16. Go into "training"
- 17. Move all of your labeled training sets to a better location outside of the project folder.
- 18. Go back into media and go into "test"
- 19. Move all of your labeled test sets to a better location outside of the project folder.
- 20. You have now completed all of the instructions for using the labeling system. Congratulations!