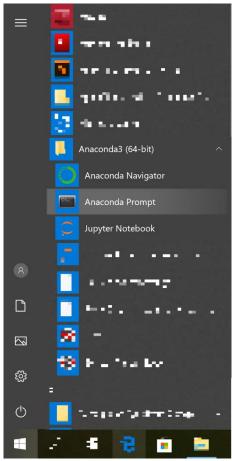
The following are instructions how to operate the code accompanying the paper: <u>Single particle</u> <u>diffusion characterization by deep learning.</u>

We hope that you find this code useful in your work.

### Installation

- Install the Anaconda Python distribution: <a href="https://www.anaconda.com/distribution/">https://www.anaconda.com/distribution/</a>
- Copy the provided anomalous\_diffusion\_DL.yml file to the Anaconda root folder (in Windows systems, it should be C:\Users\UserName)
- Open Anaconda prompt from the start menu



 Import the python environment from the provided yml file by entering the following in the command prompt:

```
conda env create -f anomalous_diffusion_DL.yml
```

Anaconda Prompt - conda env create -f anomalous\_diffusion\_DL.yml

A new environment named anomDiffDL will be created once the process finishes.

- Activate the environment by entering:

```
conda activate anomDiffDL
```

Install Keras into the new environment:

```
conda install -c anaconda keras-gpu
```

Note: this will install the GPU version of Keras, for the CPU version run:

```
conda install -c anaconda keras
```

- Install three more packages required for full functionality by entering the following in the same command prompt:

```
pip install matplotlib
pip install stochastic
pip install seaborn
```

## Training a network

- Open one of the files ending with \_training.py
- In the top of the file, there are several variable definitions which control the simulated trajectories used to train the network.

```
11batchsize = 64
12T = np.arange(19,21,0.1) # this provides another layer of stochasticity to make
13steps = [500] # number of steps to generate in total
14steps_actual = 300 # number of steps the network receives as input out of the number of steps available
```

- batchsize is the number of training samples generated in each instance of the data generator
- T is the total time of the simulation in seconds. The data generator expects to receive an array of T values from which it randomly selects a value in each iteration. This is done to increase the variability in the generated trajectories.
- steps is the number of steps in the simulated trajectory assuming a constant Δt between time points. Similar to T, the data generator expects to receive an array of possible values from which is randomly selects one in each iteration. (When choosing value options for T and steps, please keep the options relatively close to allow the network to train properly)
- steps\_actual is the number of steps the network sees out of the total trajectory length and is the trajectory length the network expects to receive. (This value should always be smaller than the minimal value in steps).
- Choose a file name for the new network model in the bottom of the script (marked in red below)

```
ModelCheckpoint(filepath='new_model.h5',
monitor='val_loss',
save_best_only=True,
mode='min',
save_weights_only=False)]
```

 Once all definitions and parameters have been set, run the script to generate a new network model.

# Analyzing data from a file

- Open and run utils.py to load auxiliary functions to memory
- You can run all analysis functions as is with the network models provided, or replace the loaded model by changing the file name where the following comment is found anywhere in the code:

### change here to load a different network model

- Open and run the appropriate testing file containing the function you want to run
- All file analysis functions receive as input a .mat file, containing one matrix of N dimensional data in the following configuration:

Where x,y represent x,y positions in pixels and N is the trajectory serial number in ascending order, starting from 1 (i.e. all x,y values belonging to trajectory 1 will have N=1 in the third column) See example in the following image:

	1	2	3
1	-27.2493	-59.1920	1
2	-27.6756	-58.8687	1
3	-27.6936	-59.2070	1
4	-28.0118	-59.4451	1
5	-28.0240	-59.2436	1
6	-28.1894	-59.4752	1
7	-28.3665	-59.4824	1
8	-28.2087	-59.0561	1
9	-27.8874	-59.2675	1
10	-27.7105	-59.1068	1
11	-27.4111	-59.0776	1
12	-27.5374	-59.3351	1
13	-27.3369	-59.2576	1
14	-27.8577	-58.8687	1
15	-28.0479	-59.0715	1
16	-28.3585	-59.0283	1
17	-28.2769	-59.1013	1
18	-28.2284	-59.0981	1
19	-28.3001	-58.9508	1
20	-28.2535	-59.5156	1
21	-28.0856	-58.9853	1
22	-27.7788	-58.8111	1
23	-27.7656	-59.0474	1
24	-27.6597	-59.0499	1
25	-27.8396	-58.9466	1
26	-27.8489	-58.7457	1
27	-28.3235	-59.1976	1
28	-28.3782	-59.1863	1

## Other functions

- Trained models can be tested on simulated data with various types of noise, using any of the functions which do not operate on .mat files. More information can be found in the comments above each function.
- Simulations can be generated separately using the functions in utils.py

#### Notes

- The files are separated into \_training and \_testing files (indicated by the end of the file name). Testing files contain functions to analyze .mat files and other functions to test network models on simulated data. Training files contain network structures and the code necessary to train a network model per the specifications in the paper.
- The network models designed to estimate the diffusion coefficient are dependent on physical parameters pixel size and time difference between steps. As such, **new models need to be trained for each new experimental setup** (assuming the experimental parameters are significantly different than one another).
- The networks in general are dependent on the Signal to Noise Ratio (SNR)
  - The signal is defined as the standard deviation of the trajectory derivative, and can be set in simulations by inputting the correct experimental conditions (number of steps and time between steps)
  - The noise is the standard deviation of the trajectory derivative of a fixed cell/bead and can be set in simulations by inputting the standard deviation value to the sigma parameter in each \_training file.

If you find this code useful, please cite our work:

Granik, N., Weiss, L.E., Nehme, E., Levin, M., Chein, M., Perlson, E., Roichman, Y. and Shechtman, Y., 2019. Single particle diffusion characterization by deep learning. Biophysical Journal.

For any questions and comments, please contact:

naorgranik@gmail.com

yoavsh@bm.technion.ac.il