# Motion Mapper Framework User Guide

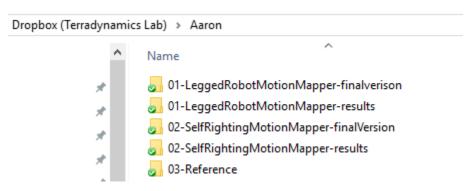
The motion mapper framework can be used to process 1) <u>kinematic data of robot/animal motion</u> and 2) <u>videos of robot/animal motion</u>. The result of the framework would be a 2d embedding space that represents the robot motion. The regions with high point's density in the space probably represent stereotyped behaviors of the robot. The followings are the instruction of the framework processing the two different kinds of inputs.

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# Directory Structure:

### D:\Dropbox (Terradynamics Lab)\Aaron



- 1) ./01-LeggedRobotMotionMapper-finalversion/: Stores the matlab codes of the first section of this manual analysis of kinematic data of the robot motion
- 2) ./01-LeggedRobotMotionMapper-results/: Stores the results of the legged robot motion mapper
- 3) ./02-SelfRightingMotionMapper-finalVersion/: Stores the matlab codes of the second section of this manual analysis of cockroach self-righting videos
- 4) ./02-SelfRightingMotionMapper-results/: Stores the results of the self righting motion mapper
- 5) **./03-Reference/:** The folder in which all study materials and resources mentioned in this manual are stored.

# I. Analysis of kinematic data of the robot motion

This project represents a sample implementation of the motion mapper behavioral analysis by using the kinematic data of the robot crossing the beam. User can easily change the dimension of input data by randomly combining the data of velocity, orientation and position.

### Methods

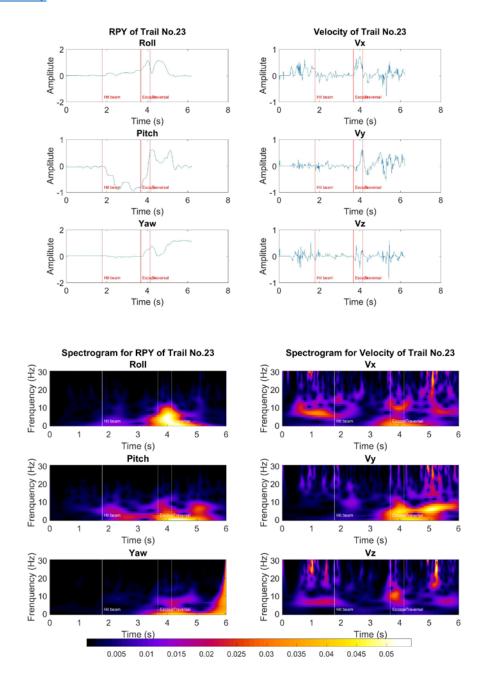
# 1. Spectrogram generation (<u>Interface paper</u> - 3.3)

We compute the amplitudes of the Morlet continuous wavelet transform for each time series. Although similar to a Fourier spectrogram, wavelets possess a multi-resolution time—frequency trade-off, allowing for a more complete description of postural dynamics occurring at several time scales. For the results presented here, we look at 30 frequency channels, dyadically spaced between 1 and 30 Hz. Here is an example of spectrogram for the trial no.23.

### Study materials:

- 1) Study notes: D:\Dropbox (Terradynamics Lab)\Aaron\03-Reference\Wavelet Transform Notes by

  Aaron.pptx
- 2) The wavelet tutorial:
  D:\Dropbox (Terradynamics Lab)\Aaron\03-Reference\WaveletTutorial.pdf or web link
  D:\Dropbox (Terradynamics Lab)\Aaron\03-Reference\Wavelets 2010.pdf or web link
- 3) The Fourier transform: <a href="https://betterexplained.com/articles/an-interactive-guide-to-the-fourier-transform/">https://betterexplained.com/articles/an-interactive-guide-to-the-fourier-transform/</a>



### 2. Spatial embedding (Interface paper - 3.4)

The generated spectrogram comprises 30 frequency channels for each of the six dimensions (R,P,Y,Vx,Vy,Vz), making each point in time represented by a 180 dimensional feature vector. We would like to find a low-dimensional representation that captures the important features of the dataset. t-SNE is used to take data from a high-dimensional space and embed it into a space of much smaller dimensionality.

### Study materials:

- 1) Comprehensive Guide on t-SNE: <a href="https://www.analyticsvidhya.com/blog/2017/01/t-sne-implementation-r-python/">https://www.analyticsvidhya.com/blog/2017/01/t-sne-implementation-r-python/</a>
- 2) Study notes: D:\Dropbox (Terradynamics Lab)\Aaron\03-Reference\t-SNE Study Notes.pdf

After we got the embedding of our spectral feature vectors into two dimensions, we generated an estimate of the probability density by convolving each point in the embedded map with a Gaussian of relatively small width. The locations of local maxima provide a potential representation for the stereotyped behaviors that the files perform.

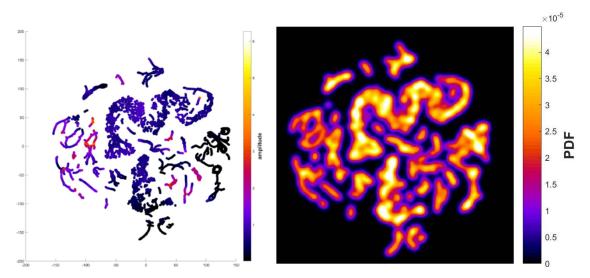


Figure left: spectral points embedded into two dimensions via t-SNE
Figure right: Probability density function (PDF) generated from convolving with a Gaussian

### 3. Behavior states definition (Interface paper - 4.1, 4.2)

The embedded space comprises peaks surrounded by valleys. Finding connected areas in the embedding plane such that climbing up the gradient of probability density always leads to some

local maximum, often referred to as a watershed transform, we delineate 116 regions of the embedded space. Each of these contains a single local maximum of probability density.

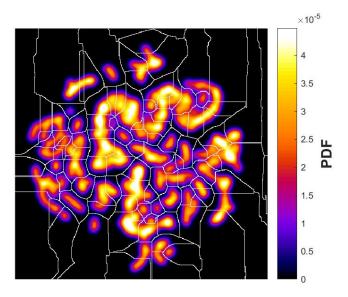


Figure: Boundary lines obtained from performing a watershed transform on the PDF.

Observing the original movies corresponding to pauses in one of the regions, we consistently observe the robot performing a distinct action that corresponds to a recognizable behavior when viewed by eye.

All the mapped videos can be found in: <a href="D:\Dropbox">D:\Dropbox</a> (Terradynamics Lab)\Aaron\Videos\Mapped

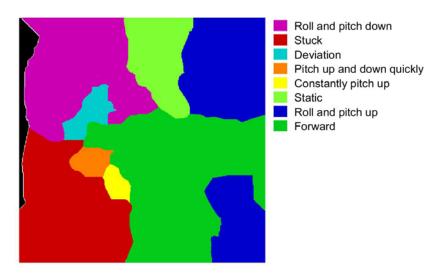
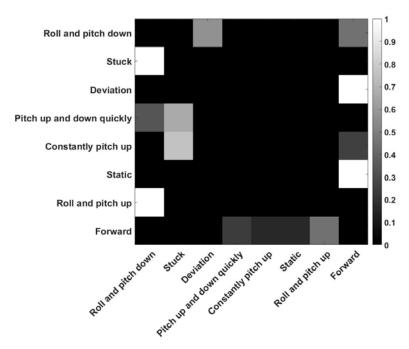


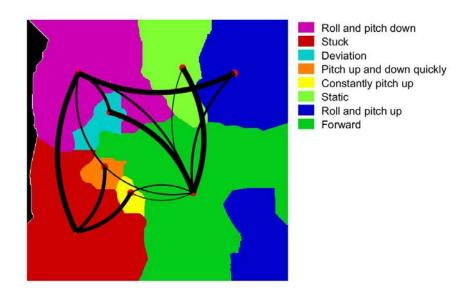
Figure: The organization of behavioral space into regions of similar movement types. Definition of regions is performed through visual assessment of movies

# 4. Transition matrices (PNAS paper - Transition Matrices)

To investigate the temporal pattern of behaviors, we first calculated the behavioral Markov transition matrix over different time scales, which describes the probability that the animal will go from state j to state i.



Plotting this matrix on the behavioral map itself, we can see the transition distribution on the behavioral map. All transition probabilities less than 0.05 are omitted.

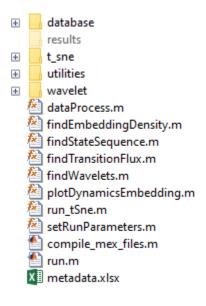


### Study materials:

- 1) MIT open course on Markov Matrics: <a href="https://youtu.be/nnssRe5DewE">https://youtu.be/nnssRe5DewE</a>
- 2) Markov Chains: <a href="https://nlp.stanford.edu/IR-book/html/htmledition/markov-chains-1.html">https://nlp.stanford.edu/IR-book/html/htmledition/markov-chains-1.html</a>

### Matlab Codes

**IMPORTANT:** the path of the final version codes that implement the above-mentioned methods: D:\Dropbox (Terradynamics Lab)\Aaron\01-LeggedRobotMotionMapper-finalverison.



- To run this code, you will first have to compile the associated mex files contained in this repository.
   In order to do this, type compile\_mex\_files in the MATLAB command prompt from the main directory.
- 2) An example run-through of all the portions of the algorithm can be found in **run.m**. Given a mat file storing the kinematic data, this will run through each of the steps in the algorithm.
- 3) All default parameters can be adjusted within **setRunParameters.m**. Additionally, parameters can be set by inputting a struct containing the desired parameter name and value into the function (see code for details).
- 4) The major sub-routines for the method are named as coherently as possible:
  - a) dataProcess.m: Process the data of RPY, XYZ and velocity.

```
function RobData = dataProcess(robotData, shellTagID, position, orientation, velocity)
```

- 1. Input:
  - robotData -> the raw kinematic data got from robot experiment.

- shellTagID -> the corresponding tag for robot.
- position -> boolean variable to decide whether select the position data (true/false).
- orientation -> boolean variable to decide whether select the orientation data (true/false).
- velocity -> Boolean variable to decide whether select the velocity data (true/false)

### 2. Output:

- RobData -> the processed kinematic data. It can be 3d, 6d or 9d depending on user selection.
- b) findWavelets.m: Computes the Morlet wavelet transform for the input kinematic data
  function [amplitudes,f] = findWavelets(timesSeries,numDims,parameters)
  - 1. Input:
    - timeSeries -> the n(time) x d(dimension) kinematic time series.
    - numDims -> the number of dimension of time series.
    - parameters -> struct containing non-default choices for parameters.

### 2. Output:

- amplitudes -> wavelet amplitudes.
- f -> frequencies used in wavelet transforms (Hz).
- c) run\_tSne.m: Runs the t-SNE algorithm

```
function yData = run_tSne(data,parameters)
```

- 1. Input:
  - data -> Nxd array of wavelet amplitudes
  - parameters -> struct containing non-default choices for parameters.

#### 2. Output:

- yData -> Nx2 array of embedding results
- d) findEmbeddingDensity.m: Calculates and plots the density on the embedding space

```
function [PDFAll,PDFStiffness,PDFTrial,PDFWatershed] = findEmbeddingDensity(embeddingValues)
```

- 1. Input:
  - embeddingValues -> 1x 30(no. of trials) cell arrays, each cell contains Nx2 array of embedding results.

### 2. Output:

• PDFAII -> embedding density figure for all the trials.

- PDFStiffness -> embedding density figure for each stiffness (low, stiffness, and high).
- PDFTrial -> embedding density figure for each trails.
- PDFWatershed -> watershed embedding density figure for all the trials.
- e) plotDynamicsEmbedding.m: Plots the trajectories of each trial on the embedding space.

# function plotDynamicsEmbedding(embeddingValues, save\_path)

- 1. Input:
  - embeddingValues -> 1x 30(no. of trials) cell arrays, each cell contains Nx2 array of embedding results.
  - Save\_path -> the path to save the figures.
- f) findTransitionFlux.m: Plots the transition flux for all clusters on the watershed embedding space.

# function findTransitionFlux(embeddingValues)

- 1. Input:
  - embeddingValues -> 1x 30(no. of trials) cell arrays, each cell contains Nx2 array of embedding results.
- 5) The function of the folders in the directory:
  - ./Database/ -> The folder saves all the .mat files
  - ./results/ -> The folder saves all the results, including the figures and embedding dynamics. User can change the result destination in the code.
  - ./t-sne/ -> Saves the function files for t-SNE
  - ./utilities/ -> Saves any other utility function files
  - ./wavelet/ -> Saves the function files for Morlet wavelet transform

### How to Run the Codes

Open the run.m at: D:\Dropbox (Terradynamics Lab)\Aaron\01-LeggedRobotMotionMapper-

### finalverison\run.m

```
clear
close all
addpath(genpath('./wavelet/'));
addpath(genpath('./utilities/'));
addpath(genpath('./t_sne/'));
                                                                  1
addpath(genpath('./database/'));
save path = 'D:\MotionMapper\LeggedRobotMotionMapper\results';
if ~exist(save path,'dir')
    mkdir(save path);
end
                                                                  2
% define any desired parameter changes here
parameters.numProcessors = 2;
parameters = setRunParameters(parameters);
numDims = parameters.numDims; % the number of dimenstion of input data
% run the dataProcess first to get the dataset
load('dataBeamRobot.mat');
position = true;
orientation = true;
velocity = true;
RobData = dataProcess(robotData, shellTagID, position, orientation, velocity);
L = length(RobData);
```

- 1. Change the save path
- 2. Define any desired parameters
- 3. Run the dataProcess.m with the selection of position/orientation/velocity

```
%% Caculte wavelet for time series
amp=cell(L,1);
for i = 1 : L
    fprintf(1,'\t Calculating Wavelets for trial #%4i out of %4i\n',i,L);
    [amp{i},~] = findWavelets(RobData{i},numDims,parameters);
end
data = combineCells(amp);
vals = sum(data,2);
data(:) = bsxfun(@rdivide,data,vals); % Normalize
5
```

- 4. Run the findWavelets.m to find the wavelets for the data generated by dataProcess.m
- 5. Normalize the wavelet amplitude.

```
%% Find embedding onto 2D by t-SNE
fprintf(1,'Finding t-SNE Embedding for the data\n');
embedding = run_tSne(data,parameters);
6
```

Run the run\_tSne.m to find the 2d embedding of the wavelet amplitude generated by findWavelets.m

```
%% Make embedding plot
embeddingAll = figure;
scatter(embedding(:,1),embedding(:,2),[],vals,'filled');
colormap(fire)
h = colorbar;
ylabel(h, 'amplitude','FontSize', 15,'fontweight','bold');
```

7. Plot the embedding results.

```
%% Make density plot
[embeddingValues] = groupCells(embedding,RobData);
[PDFAll,PDFStiffness,PDFTrial,PDFWatershed] = ...
findEmbeddingDensity(embeddingValues);
```

8. Run findEmbeddingDensity.m to plot the points' density on the embedding space.

```
%% Make trajectories on the embedding space
save_path = [save_path '\dynamics'];
plotDynamicsEmbedding(embeddingValues,save_path)
9
```

9. Run plotDynamicsEmbedding.m to plot trajectories of each trial on the embedding space.

```
%% Plot transition flux
findTransitionFlux(embeddingValues)
10
```

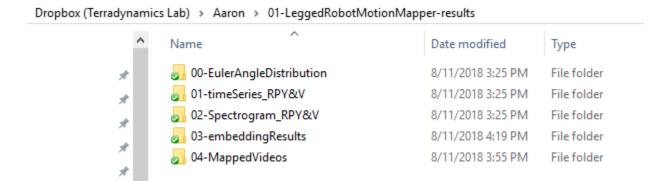
10. Run *findTransitionFlux.m* to plot the transition flux for all clusters on the watershed embedding space.

```
%% Save images
saveas(PDFAll,fullfile(save_path,'PDF_all'),'jpg');
saveas(PDFStiffness,fullfile(save_path,'PDF_stiffness'),'jpg');
saveas(PDFTrial,fullfile(save_path,'PDF_individual'),'jpg');
saveas(PDFWatershed,fullfile(save_path,'PDF_Watershed'),'jpg');
saveas(embeddingAll,fullfile(save_path,'Embedding'),'jpg');
```

11. Save the above-mentioned figures to the save path.

### **Results**

IMPORTANT: all the figures and the mapped videos are stored at D:\Dropbox (Terradynamics Lab)\Aaron\01-LeggedRobotMotionMapper-results

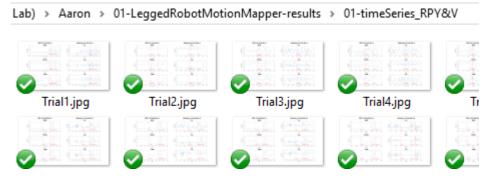


 ./00-EulerAngleDistribution/ contains the euler angle distribution of roll, pitch and yaw for the robot motion

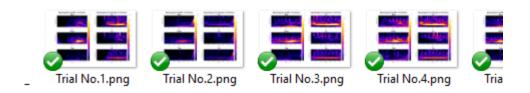
### Lab) > Aaron > 01-LeggedRobotMotionMapper-results > 00-EulerAngleDistribution



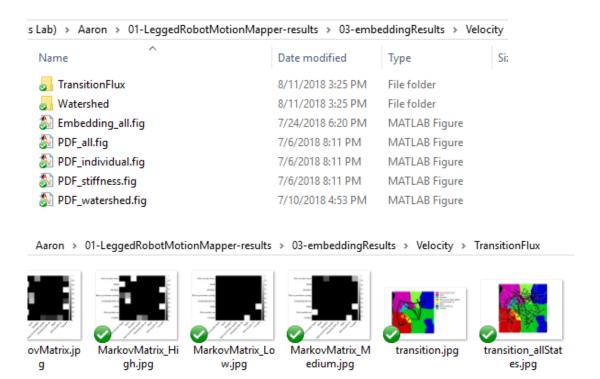
2) ./01-timeSeries\_RPY&V/ contains the figures of input time series



 ./02-Spectrogram\_RPY&V/ contains the figures of sepectrogram generated by wavelet transform 01-LeggedRobotMotionMapper-results > 02-Spectrogram\_RPY&V



4) ./03-embeddingResults/ contains the 2d embedding results for the kinematic data



5) ./04-MappedVideos/ contains the videos mapped with trajectories on embedding space

cs Lab) > Aaron > 01-LeggedRobotMotionMapper-results > 04-MappedVideos

Name	Date modified	Туре
BehaviorSpace_V	8/11/2018 3:53 PM	File folder
■ RPY	8/11/2018 3:53 PM	File folder
■ RPYV	8/11/2018 3:54 PM	File folder
RPYVPosition	8/11/2018 3:55 PM	File folder
V	8/11/2018 3:55 PM	File folder

> Aaron > 01-LeggedRobotMotionMapper-results > 04-MappedVideos > V



Mapped\_30fp s.avi



H2\_Mapped\_30fp s.avi



H3\_Mapped\_30fp s.avi



H4\_Mapped\_30fp s.avi



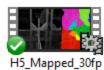
d\_30fp H5\_Mappe s.avi











H

apped\_30fp H2 s.avi

H2\_Mapped\_30f s.avi

H3\_Mapped\_30fp s.avi

H4\_Mapped\_30fp s.avi

ipped\_30fp Hb\_Mapped\_3 s.avi s.avi

# II. Analysis of cockroach self-righting videos

This is a sample implementation of the motion mapper behavioral analysis by using videos of the robot motion. The framework can process single video or multiple videos. The methods of processing videos are pretty similar to that of processing kinematic data. The difference between these two methods is the part of data processing. Here I will introduce the methods that are different from the above-mentioned ones.

#### Methods

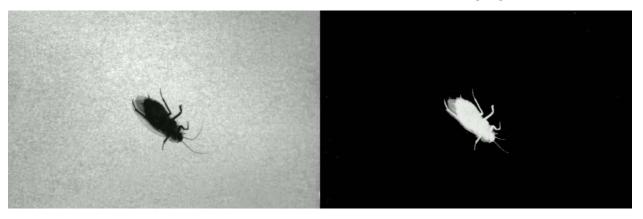
# 1. Video Segmentation (Interface paper - 3.1)

To isolate the fly from the background, we apply Canny Edge Detection method for edge detection, resulting in a binary image containing the edge positions. In order to achieve good segmentation, the contrast between the animal and background should be high. One way we can do is using translucent background and adding a light at the bottom to improve the contrast when doing experiment in the future.

### Study materials:

- 1) Canny edge detection tutorial: <a href="http://www.aishack.in/tutorials/canny-edge-detector/">http://www.aishack.in/tutorials/canny-edge-detector/</a>
- 2) Edge detection in Matlab: https://www.mathworks.com/help/images/edge-detection.html
- a) The raw videos:

b) The videos after being segmented:



### 2. Crop Video / Find Bounding Box

Because the animal goes around in the videos, the center of the fly within the frame vary over time. In order to make sure the animal can locate in the center of each frames, the framework will find the center of the animal body and then crop the frame with a  $450 \times 450$  square window whose center locates at the center of the animal body.

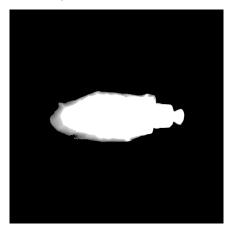
- b) The videos after being segmented:
- b) The videos after being cropped:



# 3. Video Alignment (<u>Interface paper</u> – Appendix A)

To keep the orientation of the animal within the frame same over time, we next register each frame rotationally with respect to a template image. The template image is generated by taking a typical image of the animal and then manually ablating the wings and legs digitally. In Photoshop, use the magnetic lasso tool and select the outline of the animal body, then delete the regions outside the selection region, which finally remains the template image.

### a) Template:



b) The videos after being cropped:

c) The aligned videos:



### 4. Generate Time Series by PCA (Interface paper - 3.2)

The framework will apply principal component analysis (PCA) to the videos and then project individual images onto these principal components, which convert a movie of animal behavior into time series. User can change the number of principal components in the codes.

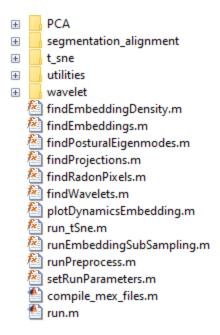
# Study materials:

PCA tutorial: <a href="https://towardsdatascience.com/a-one-stop-shop-for-principal-component-analysis-5582fb7e0a9c">https://towardsdatascience.com/a-one-stop-shop-for-principal-component-analysis-5582fb7e0a9c</a>

After generating the time series, the framework then computes the Morlet wavelet and find the embedding space by t-SNE, which are the same as the methods of processing kinematic data. (See above)

### **Matlab Codes**

**IMPORTANT:** the path of the final version codes that implement the above-mentioned methods: D:\Dropbox (Terradynamics Lab)\Aaron\02-SelfRightingMotionMapper-finalVersion.



- 1) An example run-through of all the portions of the algorithm can be found in **run.m**. Given single video or multiple videos, this will run through each of the steps in the algorithm.
- 2) All default parameters can be adjusted within **setRunParameters.m**. Additionally, parameters can be set by inputting a struct containing the desired parameter name and value into the function (see code for details).
- 3) The major sub-routines for the method are named as coherently as possible:
  - a) runPreprocess.m: Runs the alignment and segmentation routines on a .avi file

```
function runPreprocess(file_path,output_path,crop,register,parameters)
%runPreprocess runs the alignment and segmentation routines on a .avi file
%
% Input variables:
%
% file_path -> avi file to be analyzed
outputPath -> path to which files are saved
crop -> boolean variable to decide crop the frame or not
register -> boolean variable to decide register the frame or not
parameters -> struct containing non-default choices for parameters
%
```

b) findBoundingBox.m: Finds the bounding box of the image and crop it

```
|function imageOut = findBoundingBox(image,boundingWidth)
[%findBoundingBox finds the bouding box of the animal in the frame
     and crop the frame by a boundingWidth x boundingWidth square window
%
%
    Input variables:
%
        image -> image to be cropped
%
        boundingWidth -> the width of the bounding box
%
%
    Output variables:
%
        imageOut -> image after being cropped
%
```

c) findProjections.m: Finds the time series of videos by PCA

```
function projections = findProjections(filePath,vecs,meanValues,pixels,parameters)
%findPosturalEigenmodes finds the projection of a set of images onto
%postural eigenmodes.
%
%
   Input variables:
%
%
        filePath -> a directory containing an aligned .avi file
%
       vecs -> postural eignmodes (L x (M<L) array)
%
        meanValues -> mean value for each of the pixels
%
        pixels -> radon-transform space pixels to use (Lx1 or 1xL array)
%
        parameters -> struct containing non-default choices for parameters
%
%
%
   Output variables:
%
%
        projections -> N x d array of projection values
```

All other routine are the same as those in the analysis of kinematic data.

4) The function of the folders in the directory:

```
./PCA/ -> Saves the function files for PCA
./segmentation_alignment/ -> Saves the function files for registering videos
./t-sne/ -> Saves the function files for t-SNE
./utilities/ -> Saves any other utility function files
./wavelet/ -> Saves the function files for Morlet wavelet transform
```

#### How to Run the Codes

Open the run.m at <u>D:\Dropbox (Terradynamics Lab)\Aaron\02-SelfRightingMotionMapper-finalVersion\run.m</u>

```
close all
%Place path to folder containing example .avi files here
filePath = 'D:\MotionMapper\selfRighting';
                                                    1
%add utilities folder to path
addpath(genpath('./utilities/'));
addpath(genpath('./PCA/'));
addpath(genpath('./segmentation alignment/'));
addpath(genpath('./t sne/'));
addpath(genpath('./wavelet/'));
%find all avi files in 'filePath'
videoFiles = findAllVideosInFolders(filePath, '.avi');
L = length(videoFiles);
numZeros = ceil(log10(L+1e-10));
%define any desired parameter changes here
parameters.samplingFreq = 100;
parameters.trainingSetSize = 5000;
                                                    3
%initialize parameters
parameters = setRunParameters(parameters);
```

- 1. Change the path to the folder containing the .avi videos
- 2. Find all the .avi videos in the path.
- 3. Define any desired parameters.

```
%% Run preprocessing for all videos
% creating preporcessing directory
processedDirectory = [filePath '/preprocess/'];
if ~exist(processedDirectory,'dir')
    mkdir(processedDirectory);
end
fprintf(1, 'Processing videos\n');
processedVideos = cell(L,1);
                                                                5
crop = true;
register = true;
for ii=1:L
    fprintf(1,'\t Processing video #%4i out of %4i\n',ii,L);
    [~,fileName,~] = fileparts(videoFiles{ii});
    processedVideos{ii} = [processedDirectory 'Processed ' fileName '.avi'];
    runPreprocess(videoFiles{ii},processedVideos{ii},crop,register,parameters);
end
```

- 4. Change the path to save the processed videos.
- 5. Decide whether crop and align the videos.
- 6. Change the name of the video to be saved.
- 7. Run runPreprocess.m to process the videos and save the videos to the path.

```
%% Find image subset statistics (a gui will pop-up here)
fprintf(1,'Finding Subset Statistics\n');
numToTest = parameters.pca_batchSize;
[pixels,thetas,means,stDevs,vidObjs] = findRadonPixels(processedDirectory,numToTest,parameters);
```

8. Run findRadonPixels.m to run randon transform to find subset pixels of images

```
%% Find postural eigenmodes
fprintf(1,'Finding Postural Eigenmodes\n');
[vecs,vals,meanValues] = findPosturalEigenmodes(vidObjs,pixels,parameters);
vecs = vecs(:,1:parameters.numProjections);
```

9. Run findPosturalEigenmodes.m to find the principal components / postural eigenmodes.

```
%% Find projections for each data set
projectionsDirectory = [filePath '/projections/'];
if ~exist(projectionsDirectory, 'dir')
    mkdir(projectionsDirectory);
                                                                                 10
end
fprintf(1, 'Finding Projections\n');
for i=1:L
    fprintf(1,'\t Finding Projections for File #%4i out of %4i\n',i,L);
                                                                                  11
    projections = findProjections(processedVideos{i}, vecs, meanValues, pixels, parameters);
    fileNum = [repmat('0',1,numZeros-length(num2str(i))) num2str(i)];
    [~,fileName,~] = fileparts(videoFiles{i});
    save([projectionsDirectory 'projections_' fileName '.mat'], 'projections', 'fileName');
    projectionFiles{i} = projections;
    clear projections
    clear fileNum
    clear fileName
end
```

- 10. Change the path to save the time series generated by PCA
- 11. Run findProjections.m to run PCA and generate time series.
- 12. Save the time series to the path.

```
%% Use subsampled t-SNE to find training set
fprintf(1, 'Finding Training Set\n');
                                                                               13
[trainingSetData,trainingSetAmps] = ...
    runEmbeddingSubSampling(projectionsDirectory,parameters);
%% Run t-SNE on time series
fprintf(1, 'Finding t-SNE Embedding for the Training Set\n')|;
[trainingEmbedding,betas,P,errors] = run tSne(trainingSetData,parameters);
%% Find Embeddings for each file
fprintf(1, 'Finding t-SNE Embedding for each file\n');
embeddingValues = cell(L,1);
for i=1:L
    fprintf(1,'\t Finding Embbeddings for File #%4i out of %4i\n',i,L);
                                                                               15
    projections = projectionFiles{i};
    projections = projections(:,1:parameters.pcaModes);
    [embeddingValues{i},~] = ...
        findEmbeddings(projections, trainingSetData, trainingEmbedding, parameters);
    clear projections
end
```

13. Run runEmbeddingSubSampling.m to find training set of time series.

- 14. Run run tSne.m to find the training embeddings on the training set.
- 15. Run findEmbeddings.m to find the 2d embeddings space for all videos

```
%% Plot trajectories on embedding space
save_path = [filePath '\dynamics'];
plotDynam|icsEmbedding(embeddingValues, save_path);
16
```

16. Run plotDynamicsEmbedding.m to plot trajectories of each file on the embedding space.

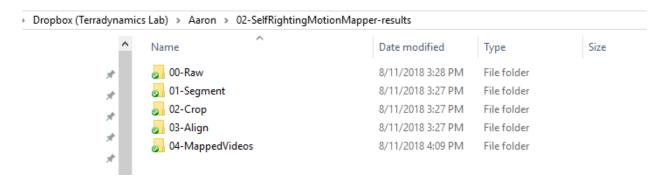
```
%% Make density plots
[PDFAll,PDFIndividual,PDFWatershed] = findEmbeddingDensity(embeddingValues);

saveas(PDFAll,fullfile(save_path,'PDF_all'),'jpg');
saveas(PDFIndividual,fullfile(save_path,'PDF_individual'),'jpg');
saveas(PDFWatershed,fullfile(save_path,'PDF_Watershed'),'jpg');
18
```

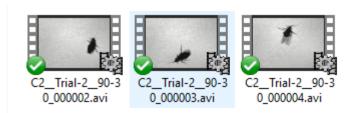
- 17. Run findEmbeddingDensity plot the points' density on the embedding space.
- 18. Save the above-mentioned figures.

### Results

IMPORTANT: all the figures and the mapped videos are stored at D:\Dropbox (Terradynamics Lab)\Aaron\02-SelfRightingMotionMapper-results

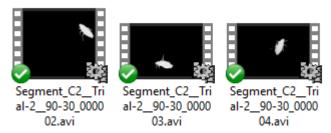


1) ./00-Raw/ contains the raw videos of the cockroach self-righting

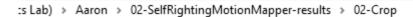


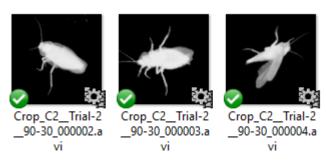
2) ./01-Segment/ contains the videos after being segmented





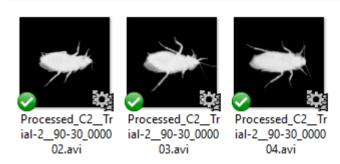
3) ./02-Crop/ contains the video after being cropped by a square window





4) ./03-Align/ contains the alignment videos

s Lab) > Aaron > 02-SelfRightingMotionMapper-results > 03-Align



5) ./04-MappedVideos/ contains the videos mapped with trajectories on the embedding space

