Reference

class PHETS.PH.Filtration(sig, params[, filename='none'])

- **Parameters:** sig (ndarray or str) Input trajectory as numpy array or path to .txt file.
 - **params** (*dict*) Options for computation of filtration.
 - filename (str) If sig is an ndarray, a filename may be provided for labelling in visualizations.

PHETS.PH.make_movie(filtration, out filename[, color scheme='none', camera angle=(135, 55), alpha=1, dpi=150, save frames=False, framerate=1)

Make filtration movie.

- **Parameters:** filtration (Filtration) -
 - out filename (str) Path/filename for movie. Should end in ".mp4" or other video format.
 - color_scheme (str) 'none', 'highlight new', or 'birth time gradient'
 - camera angle (tuple) for 3D mode. (azimuthal, elevation) in degrees.
 - alpha (float) opacity of simplexes. 0...1 : transparent...opaque
 - **dpi** (*int*) dots per inch (resolution)
 - save frames (bool) save frames to PH/frames/frame*.png for debugging
 - framerate (int) frames per second

PHETS.PH.make_PD(filtration, out filename)

Plot persistence diagram.

- **Parameters:** filtration (Filtration) -
 - out_filename (str) Image path and filename. Should end in ".png" or other image format.

PHETS.PH.make_PRF_plot(filt, out filename[, PRF res=50])

Plot persistent rank function.

- **Parameters:** filtration (Filtration) -
 - out_filename (str) Image path and filename. Should end in ".png" or other image format.
 - PRF res (int) number of divisions per epsilon axis

Todo:

a function for a single frame of the filtration (i.e. fixed epsilon) as an image. (For now, a single frame movie can be used.)

PHETS.DCE.Movies.slide_window(in filename, out filename, window size=.5, step size=.1, tau=10, ds rate=1, max frames=0, save worms=True, save movie=True)

Show embedding of in filename with window start point varied over time.

- **Parameters:** window size (float) seconds
 - **step size** window start point step size (seconds)
 - tau (float) seconds?

PHETS.DCE.Movies.vary_tau(in filename, out filename, tau lims=(1, 15), tau inc=1, embed crop=(1, 2), ds rate=1, save worms=True, save movie=True, m=2)

Show embedding of in filename with tau varied over time.

- **Parameters:** in filename (*str*) Path/filename for text file time series.
 - out filename (str) Path/filename for movie. Should end in ".mp4" or other video format.
 - tau_lims (tuple) tau range (seconds)
 - **tau inc** (*int*) tau stepsize (seconds)
 - embed crop (tuple) Limits for window from input time series (seconds)
 - **ds rate** (*int*) time series downsample rate
 - save_worms (bool) save embeddings to text files in output/DCE /saved worms
 - save movie (bool) If False, no movie will be created. Useful for saving embeddings quickly.
 - **m** (*int*) target embedding dimension

PHETS.DCE.Movies.compare_vary_tau(in filename 1, in filename 2, out filename, tau lims, tau inc=1, embed crop=(1, 2), ds rate=1, m=2, save worms=True, save movie=True) Like vary tau(), but shows embeddings for two time series side by side.

PHETS.DCE.Movies.compare multi(dir1, dir1 base, dir2, dir2 base, out filename, i lims=(1,89), $embed_crop_1 = 'auto', embed_crop_2 = 'auto', auto_crop_length = .3, tau_1 = 'auto ideal', tau_2 = 'auto'$ ideal', tau T=1/np.pi, save worms=True, save movie=True, normalize volume=True, waveform zoom=None, ds rate=1, dpi=200, m=2)

Takes two directories of (eg one with piano notes, another with range of viol notes), and generates a movie over a range note indexes (pitch). Tau and crop may be set explicity or automatically.

- **Parameters:** dir1 (str) Path of first directory to be iterated over
 - dir1 base (str) Base filename for files in dir1
 - dir2 (str) -
 - dir2 base (str) -
 - out filename (str) -
 - i_lims (tuple) (start, stop) index. Default is (1, 89).
 - **embed crop 1** (tuple or str) (start, stop) in seconds or 'auto'
 - embed_crop_2 (tuple or str) -
 - auto_crop_length=.3 (float) seconds
 - tau 1 (str) explicit (seconds) or 'auto detect' or 'auto ideal'
 - tau 2 (str) -
 - tau_T (float) For use with auto tau: tau = period * tau_T
 - save worms (bool) -
 - save movie (bool) -
 - normalize volume (bool) -
 - waveform_zoom -
 - **ds rate** (int) -
 - **dpi** (int) -

Todo:

function for plotting embeddings without varying a parameter or input, as an image. (For now, a single frame movies can be used.)

PHETS.PRFCompare.plot dists vs ref(dir, base filename, fname format, out filename, filt params, i_ref=15, i_arr=np.arange(10, 20, 1), weight_func=lambda i, j: 1, metric='L2', dist_scale='none', PRF res=50, load saved PRFs=False, see samples=5)

Takes range of time-series files and a reference file. Generates PRF for each, and finds distances to reference PRF, plots distance vs index.

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Parameters: • dir (str) – input directory
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- base filename (str) input base filename
- fname format (str) input filename format: 'base i or 'i base'
- out filename (str) output filename
- filt params -
- i ref (int) -
- i_arr (arr) -
- weight func (lambda) Default is lambda i, j: 1
- metric (str) 'L1' (abs) or 'L2' (euclidean). Default is 'L2'.
- **dist scale** (*str*) 'none', 'a', or 'a + b'. Default is 'none'.
- **PRF** res (int) number of divisions used for PRF. Default is 50.
- load saved PRFs (bool) reuse previously computed PRF set. Default is
- see_samples (int) interval to generate PRF plots, PDs, and filtration movies when generating PRF set. 0 is none, 1 is all samples, 2 is every other sample,

PHETS.PRFCompare.plot dists vs mean(filename 1, filename 2, out filename, filt params, load saved PRFs=False, time units='seconds', crop 1='auto', crop 2='auto', auto crop length=.3, window size=.05, num windows=10, mean samp num=5, tau 1=.001, tau 2=.001, tau T=np.pi, note_index=None, normalize_volume=True, PRF_res=50, dist_scale='none', metric='L2', weight func=lambda i, j: 1, see samples=5)

Takes two time-series or 2D trajectory files. For each input, slices each into a number of windows. If inputs are time-series, embeds each window. Generates PRF for each window. selects subset of window PRFs, computes their mean, plots distance to mean PRF vs time.

- Parameters: filename 1 (str) -
 - **filename 2** (str) -
 - out filename (str) -
 - filt_params (dict) -
 - load saved PRFs (bool) -
 - time units (str) -
 - crop 1 (str or tuple) -
 - crop 2 (str or tuple) -
 - auto crop length (float) -
 - window size (float) (time units)
 - num_windows (int) per file
 - mean samp num (int) per file
 - tau 1 (str or float) -
 - tau_2 (str or float) -
 - tau T (float) -
 - note index (int) -
 - normalize volume (bool) -
 - PRF res (int) number of divisions used for PRF
 - dist scale (str) 'none', 'a', or 'a + b'
 - **metric** (*str*) 'L1' (abs) or 'L2' (euclidean)
 - weight func (lambda) -
 - see samples (int) -

PHETS.PRFCompare.plot clusters(*args, **kwargs)

See plot_dists_vs_mean for call signature.

PHETS.PRFCompare.plot_variances(filename, out_filename, filt_params, vary_param_1, vary_param_2, load_saved_PRFs=False, time_units='seconds', crop=(100, 1100), auto_crop_length=.3, window_size=1000, num_windows=5, tau=.001, tau_T=np.pi, note_index=None, normalize_volume=True, PRF_res=50, dist_scale='none', metric='L2', weight_func=lambda_i, j: 1, see_samples=5)

Parameters: • filename (str) -

- out_filename (str) -
- filt_params (dict) -
- **vary_param_1** (*str*) -
- **vary_param_2** (*str*) -
- load saved PRFs (bool) -
- time units='seconds' (str) -
- crop=(100, 1100) (tuple) -
- auto_crop_length=.3 (float) -
- window_size=1000 -
- num windows (int) -
- tau (float) -
- tau_T (float) -
- note_index (int) -
- normalize_volume (bool) -
- PRF res (int) -
- dist scale (str) 'none', 'a', or 'a + b'
- metric (str) 'L1' (abs) or 'L2' (euclidean)
- weight_func (lambda) -
- see_samples (bool) -