

Smdata Quantification Toolbox: Documentation

1. ChangeLog

Sun Aug 16: osporns@indiana.edu, max.lungarella@aist.go.jp

- Released complexity toolbox v0.005 (functions: calcI_det.m, calcC_det.m, smdata_cov.m, smdata_corr.m, smdata_H256.m, smdata_jointH256.m, smdata_MIt.m, smdata_MIcov.m)

Sun Aug 30: max.lungarella@aist.go.jp

- Draft documentation
- Released complexity toolbox v.0.01 (added functions: smdata_corr.m, entropy.m, smdata_Hn.m, smdata_roulstonH256.m, smdata_jointHn.m, smdata_MItn.m, smdata_roulstonMIt.m, ksdensity2d.m, isomap.m, isomapii.m, L2_distance.m)

Tue Sep 16: max.lungarella@aist.go.jp

- Updated documentation
- Released complexity toolbox v.0.02 (added functions: pca_cov.m, pca_svd.m, sortem.m)

Tue Sep 30: osporns@indiana.edu, max.lungarella@aist.go.jp

- Changed title of package and of documentation
- Released complexity toolbox v.0.03 (added functions: discretize.m, normalize.m; removed functions: smdata_H256.m, smdata_jointH256.m, smdata_roulstonH256.m)

Sat Oct 11: max.lungarella@aist.go.jp

- Fixed a bug in matscat_plot.m
- Fixed a bug in test_ksdensity.m

- `matscat_plot.m` is also called a Hinton diagram. Added functions `hinton.m` and `test_hinton.m`. The former was adapted from `hintonw.m` from the neural network toolbox.

Tue Apr 6: `maxl@isi.imi.i.u-tokyo.ac.jp`

- Released complexity toolbox v.0.05 (what happened to v.0.04?). Added functions: `ksdensity1d.m`, `gsi.m`, `test_gsi.m`

Mon May 31: `maxl@isi.imi.i.u-tokyo.ac.jp`

- Released complexity toolbox v.0.06. Many unused functions were purged, some functions were integrated with other ones.

2. Short Introduction

3. Overview of the Functions

Function	
smd_acf.m	Auto-correlation
smd_H.m	Entropy
smd_Himages.m	Entropy
smd_jointH.m	Joint entropy
smd_covcor.m	Covariance and correlation
smd_corrt.m	Correlation
smd_ksdens1d.m	Histogram
smd_ksdens2d.m	Histogram
smd_MIt.m	Mutual information
smd_MIcov.m	Mutual information
smd_calcI_det.m	Integration of a system
smd_calcC_det.m	Complexity of a system
isomap.m	Isometric feature mapping
isomapii.m	Isometric feature mapping
smd_pca_cov.m	Principal component analysis
smd_pca_svd.m	Principal component analysis
smd_gsi.m	Linear separability
smd_matscat_plot.m	Visualization
smd_hinton.m	Visualization
smd_discrete.m	Discretization
smd_norm.m	Normalization
sortem.m	Sort of eigenvalues and vectors
L2_distance.m	Euclidean distance
test_discrete.m	Test function
test_gsi.m	Test function
test_hinton.m	Test function
test_ksdensity.m	Test function
test_matscat.m	Test function
test_pca.m	Test function
test_smdata.m	Test function

Table 1:

4. Entropy

4.1 `smd_H.m`

Synopsis:

```
[H_obs, B_star, H_inf, Sigma_H] = smd_H(Mstates, nst)
```

Input:

`Mstates` = NxM matrix (ensemble of series: columns are realizations of stochastic processes)

`nst` = number of states of input space

Output:

`H_obs` = 1xM (observed entropy)

`B_star` = 1xM (number of states for each column of `Mstates`)

`H_inf` = 1xM vector (corrected estimate of entropy)

`Sigma_H` = 1xM vector (standard deviation from theoretically predicted entropy)

Notes:

(a) `Mstates` must be supplied in discretized format with states ranging from 1 to `nst`. To generate such states refer to `smd_discrete.m`

(b) To obtain a meaningful estimate, a sufficient number of samples of the random variables `Mstates` should be supplied. For example, for 8-bit resolution (`nst` = 256 possible states), $N > 3 \times 256 = 800$ samples, because otherwise there are not enough samples to estimate the histogram.

References:

Roulston (1999)

4.2 `smd_Himages.m`

Synopsis:

```
[Himages] = smd_Himages(Mstates, nst)
```

Input:

`Mstates` = NxM matrix (each row represents a vector-coded image)

`nst` = number of states of input space

Output:

`Himages` = 1xM (each element represents unbiased entropy estimate of one image)

Notes:

(a) `Mstates` must be supplied in discretized format with states ranging from 1 to `nst`. To generate such states refer to `smd_discrete.m`

(b) To obtain a meaningful estimate, a sufficient number of samples of the random variables `Mstates` should be supplied. For example, for 8-bit resolution (`nst = 256` possible states), $N > 3 \times 256 = 800$ samples.

4.3 `smd_jointH.m`

Synopsis:

```
[ jointH ] = smd_jointH( Mstates , units , nst , t )
```

Calculates joint entropy for each pair of units in `Mstates`

Input:

`Mstates` = $N \times M$ matrix (realizations of stochastic process)

`units` = $1 \times L$, $L \leq M$ (columns of `Mdata` used for calculation)

`nst` = number of states of input space

`t` = time offset used to compute joint entropy

Output:

`jointH` = $1 \times M$ vector (time-delayed joint entropy, computed column-wise)

Notes:

`Mstates` must be supplied in discretized format with states ranging from 1 to `nst`.

5. Covariance and Correlation

5.1 `smd_covcor.m`

Synopsis:

```
[COV, COR] = smd_covcor(Mdata, units)
```

Calculates the covariance and the correlation matrix

Input:

`Mdata` = $N \times M$ matrix (realization of stochastic process)

`units` = $1 \times L$, $L \leq M$ (columns of `Mdata` used for calculation)

Output:

`COV` = $L \times L$ matrix (covariance)

`COR` = $L \times L$ matrix (correlation)

5.2 `smd_coort.m`

Synopsis:

```
[CORRt] = smd_coort(Mdata, units, t, T)
```

Calculates the time-delayed correlation matrix

Input:

`Mdata` = $N \times M$ matrix (realization of stochastic process)

`units` = $1 \times L$, $L \leq M$ (columns of `Mdata` used for calculation)

`t` = time offset used to compute correlation

`T` = time window

Output:

`CORRt` = $L \times L$ matrix (time-delayed correlation)

6. Histogram

6.1 `smd_ksdens2d.m`

Synopsis:

```
[D] = smd_ksdens2d(Mdata, gridx1, gridx2, bw)
```

Input:

Mdata = Nx2 matrix

gridx1 = Mx1 vector

gridx2 = Mx1 vector

bw = bandwidth (window parameter)

Output:

D = Kernel density estimation of 2D histogram using Gaussian windows

References:

Moon, Rajagopalan & Lall (1995)

7. Mutual Information

7.1 `smd_MIcov.m`

Synopsis:

```
[MI] = smd_MIcov(Mstates, units)
```

Input:

`Mstates` = NxM matrix (realizations of stochastic process)

`units` = 1xL, $L \leq M$ (columns of `Mstates` used for calculation)

Output:

`MI` = LxL matrix (mutual information between all pairs part of units)

7.2 `smd_MIt.m`

Synopsis:

```
[MI_t_inf, MI_t_obs] = smd_MIt(Mstates, units, nst, t)
```

Input:

`Mstates` = NxM matrix (realizations of stochastic process)

`units` = 1xL, $L \leq M$ (columns of `Mstates` used for calculation)

`nst` = number of states of input space

`t` = time offset used to compute mutual information

Output:

`MI_t_inf` = MxM matrix (corrected time-delay mutual information between all pairs of elements that are part of units)

`MI_t_obs` = MxM matrix (observed mutual information)

References:

Roulston (1999); Steuer, Kurths, Daub, Weise & Selbig (2002)

8. Complexity Measures

8.1 `smd_calcI_det.m`

Synopsis:

```
[ I ] = smd_calcI_det ( COV )
```

Input:

COV = covariance matrix of system X

Output:

I = integration

Notes:

Assumption of Gaussianity for X

References:

Olaf?

8.2 `smd_calcC_det.m`

Synopsis:

```
[ C ] = smd_calcC_det ( COV )
```

Input:

COV = covariance matrix of system X

Output:

C = complexity

Notes:

Uses `smd_calcI_det.m`

References:

Olaf?

9. Dimensionality Reduction

9.1 `smd_pca_cov.m`

Synopsis:

```
[Z,U,lambda] = smd_pca_cov(Mdata,A)
```

Input:

Mdata = NxM matrix (rows: observations, columns: variables)

A = indices of returned principal components

Output:

Z = NxA matrix (z-loading)

U = MxA matrix (principal components)

lambda = Ax1 vector (explained variance)

9.2 `smd_pca_svd.m`

Synopsis:

```
[Z,U,lambda] = smd_pca_svd(Mdata,A)
```

Input:

Mdata = NxM matrix (rows: observations, columns: variables)

A = indices of returned principal components

Output:

U = MxA matrix (principal components)

lambda = Ax1 vector (explained variance)

9.3 `smd_gsi.m`

Synopsis:

```
[GSI] = smd_gsi(Mdata,Catg)
```

Input:

Mdata = NxM matrix (rows: observations, e.g. sensory patterns; columns: variables)

Catg = Nx1 vector (category vector, category to which observations or patterns belong)

Output:

GSI = Geometric Separability Index

References:

Thornton (1997)

9.4 isomap.m**Synopsis:**

```
[Y,R,E] = isomap(D,n_fcn,n_size,options)
```

Input:**Output:****References:**

Tenenbaum, de Silva & Langford (2000)

9.5 isomapii.m**Synopsis:**

```
[Y,R,E] = isomapii(D,n_fcn,n_size,options)
```

Input:**Output:****References:**

Tenenbaum, de Silva & Langford (2000)

10. Visualization

10.1 `smd_matscat_plot.m`

Synopsis:

```
smd_matscat_plot(M,Z,f,marker)
```

Input:

$M = N \times 2$ matrix (sets the position of the patches)

$Z = N \times 1$ vector (sets the size of the patches)

$f =$ number (scale factor)

`marker = marker` ('s':square, 'o':circles)

Example:

```
[X,Y]=meshgrid(1:20,1:20);  
X=reshape(X,400,1);  
Y=reshape(Y,400,1);  
M=0:0.01:3.99;  
M=M'+0.01;  
figure(1); clf;  
smd_matscat_plot([X,Y],M,100,'s');
```

11. Miscellanea

11.1 `smd_discrete.m`

Synopsis:

```
[Mstates] = smd_discrete(Mdata, nst)
```

Discretizes the `Mdata` into `nst` discrete states.

Input:

`Mdata` = NxM matrix (input)

`nst` = scalar (number of states)

Output:

`Mstates` = NxM matrix (ranges from 1 to `nst`)

11.2 `smd_norm.m`

Synopsis:

```
[Mdata_out] = smd_norm(Mdata_in, lo_limit, hi_limit)
```

Normalizes `Mdata_in` between the user-supplied limits `lo_limit` and `hi_limit`.

Input:

`Mdata_in` = NxM matrix

`lo_limit` = scalar (lower limit)

`hi_limit` = scalar (upper limit)

Output:

`Mdata_out` = NxM matrix

11.3 `sortem.m`

Synopsis:

```
[n, v] = sortem(nd, nv)
```

Input:

Output:

11.4 L2_distance.m

Synopsis:

[D] = L2_distance(A,B)

Computes Euclidean distance matrix

Input:

A = NxM matrix

B = NxP matrix

Output:

D = MxP matrix

References

- Moon, Y., Rajagopalan, B. & Lall, U. (1995). Estimation of mutual information using kernel density estimators. *Physical Review E*, **52**, 2318–2321.
- Roulston, M. (1999). Estimating the errors on measured entropy and mutual information. *Physica D*, **125**, 285–294.
- Steuer, R., Kurths, J., Daub, C., Weise, J. & Selbig, J. (2002). The mutual information: detecting and evaluating dependencies between variables. *Bioinformatics*, **18**, 231–240, suppl.2.
- Tenenbaum, J., de Silva, V. & Langford, J. (2000). A global geometric framework for nonlinear dimensionality reduction. *Science*, **290**, 2319–2323.
- Thornton, C. (1997). Separability is a learner's best friend. In *Proc. of 4th Workshop on Neural Computation and Psychology: Connectionists Representations*, 40–47.