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### R\_Examples

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Examples of using the toolkit in R

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### R code examples

This page describes a basic set of demonstration scripts for using the toolkit in R. The .r files can be found at demos/r in the distributions (from the V1.1 release onwards).

Please see UseInR for instructions on how to begin using the java toolkit from inside R.

Note that these examples use the rJava R library -- you will need to alter them if you want to use another R-Java interface (though I believe this is the standard one).

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### **Example 1 - Transfer entropy on binary data**

example1TeBinaryData.r - Simple transfer entropy (TE) calculation on binary data using the discrete TE calculator:

```
# Load the rJava library and start the JVM
library("rJava")
.jinit()
```

```
# Change location of jar to match yours:
# IMPORTANT -- If using the default below, make sure you have set the working of
  in R (e.g. with setwd()) to the location of this file (i.e. demos/r) !!
.jaddClassPath("../../infodynamics.jar")
# Generate some random binary data:
sourceArray<-sample(0:1, 100, replace=TRUE)</pre>
destArray<-c(OL, sourceArray[1:99]); # Need OL to keep as integer array
sourceArray2<-sample(0:1, 100, replace=TRUE)</pre>
# Create a TE calculator and run it:
teCalc<-.jnew("infodynamics/measures/discrete/TransferEntropyCalculatorDiscrete"
.jcall(teCalc, "V", "initialise") # V for void return value
.jcall(teCalc,"V","add0bservations",sourceArray, destArray)
result1 <- .jcall(teCalc, "D", "computeAverageLocalOfObservations")</pre>
cat("For copied source, result should be close to 1 bit : ", result1, "\n")
# Now look at the unrelated source:
.jcall(teCalc, "V", "initialise") # V for void return value
.jcall(teCalc, "V", "addObservations", sourceArray2, destArray)
result2 <- .jcall(teCalc, "D", "computeAverageLocalOfObservations")</pre>
cat("For random source, result should be close to 0 bits: ", result2, "\n")
```

### Example 2 - Transfer entropy on multidimensional binary data

example2TeMultidimBinaryData.r - Simple transfer entropy (TE) calculation on multidimensional binary data using the discrete TE calculator.

This example is important for R users, because it shows how to handle multidimensional arrays from R to Java (this is not as simple as single dimensional arrays in example 1 - it requires using extra calls to convert the array).

```
# Load the rJava library and start the JVM
library("rJava")
.jinit()

# Change location of jar to match yours:
# IMPORTANT -- If using the default below, make sure you have set the working c
# in R (e.g. with setwd()) to the location of this file (i.e. demos/r) !!
.jaddClassPath("../../infodynamics.jar")

# Create many columns in a multidimensional array (2 rows by 100 columns),
# where the next time step (row 2) copies the value of the column on the left
# from the previous time step (row 1):
twoDTimeSeriesRtime1 <- sample(0:1, 100, replace=TRUE)</pre>
```

```
twoDTimeSeriesRtime2 <- c(twoDTimeSeriesRtime1[100], twoDTimeSeriesRtime1[1:99])
twoDTimeSeriesR <- rbind(twoDTimeSeriesRtime1, twoDTimeSeriesRtime2)

# Create a TE calculator and run it:
teCalc<-.jnew("infodynamics/measures/discrete/TransferEntropyCalculatorDiscrete"
.jcall(teCalc,"V","initialise") # V for void return value

# Add observations of transfer across one cell to the right per time step:
twoDTimeSeriesJava <- .jarray(twoDTimeSeriesR, "[I", dispatch=TRUE)
.jcall(teCalc,"V","addObservations", twoDTimeSeriesJava, 1L)
result2D <- .jcall(teCalc,"D","computeAverageLocalOfObservations")
cat("The result should be close to 1 bit here, since we are executing copy opera</pre>
```

# **Example 3 - Transfer entropy on continuous data using kernel estimators**

example3TeContinuousDataKernel.r - Simple transfer entropy (TE) calculation on continuous-valued data using the (box) kernel-estimator TE calculator.

```
# Load the rJava library and start the JVM
library("rJava")
.jinit()
# Change location of jar to match yours:
# IMPORTANT -- If using the default below, make sure you have set the working of
  in R (e.g. with setwd()) to the location of this file (i.e. demos/r) !!
.jaddClassPath("../../infodynamics.jar")
# Generate some random normalised data.
numObservations<-1000
covariance <- 0.4
sourceArray<-rnorm(numObservations)</pre>
destArray = c(0, covariance*sourceArray[1:numObservations-1] + (1-covariance)*rr
sourceArray2<-rnorm(numObservations) # Uncorrelated source</pre>
# Create a TE calculator and run it:
teCalc<-.jnew("infodynamics/measures/continuous/kernel/TransferEntropyCalculator
.jcall(teCalc,"V","setProperty", "NORMALISE", "true") # Normalise the individual
.jcall(teCalc, "V", "initialise", 1L, 0.5) # Use history length 1 (Schreiber k=1),
.jcall(teCalc,"V","setObservations", sourceArray, destArray)
# For copied source, should give something close to expected value for correlate
result <- .jcall(teCalc, "D", "computeAverageLocalOfObservations")</pre>
cat("TE result ", result, "bits; expected to be close to ", log(1/(1-covariance
.jcall(teCalc, "V", "initialise") # Initialise leaving the parameters the same
.jcall(teCalc,"V","setObservations", sourceArray2, destArray)
# For random source, it should give something close to 0 bits
result2 <- .jcall(teCalc, "D", "computeAverageLocalOfObservations")</pre>
cat("TE result ", result2, "bits; expected to be close to 0 bits for uncorrelat
```

# Example 4 - Transfer entropy on continuous data using Kraskov estimators

example4TeContinuousDataKraskov.r - Simple transfer entropy (TE) calculation on continuous-valued data using the Kraskov-estimator TE calculator.

```
# Load the rJava library and start the JVM
library("rJava")
.jinit()
# Change location of jar to match yours:
# IMPORTANT -- If using the default below, make sure you have set the working of
   in R (e.g. with setwd()) to the location of this file (i.e. demos/r) !!
.jaddClassPath("../../infodynamics.jar")
# Generate some random normalised data.
numObservations<-1000
covariance <- 0.4
sourceArray<-rnorm(numObservations)</pre>
destArray = c(0, covariance*sourceArray[1:numObservations-1] + (1-covariance)*rr
sourceArray2<-rnorm(numObservations) # Uncorrelated source</pre>
# Create a TE calculator:
teCalc<-.jnew("infodynamics/measures/continuous/kraskov/TransferEntropyCalculatc
.jcall(teCalc,"V","setProperty", "k", "4") # Use Kraskov parameter K=4 for 4 nea
# Perform calculation with correlated source:
.jcall(teCalc,"V","initialise", 1L) # Use history length 1 (Schreiber k=1)
.jcall(teCalc,"V","setObservations", sourceArray, destArray)
result <- .jcall(teCalc, "D", "computeAverageLocalOfObservations")</pre>
# Note that the calculation is a random variable (because the generated
# data is a set of random variables) - the result will be of the order
# of what we expect, but not exactly equal to it; in fact, there will
# be a large variance around it.
cat("TE result ", result, "nats; expected to be close to ", log(1/(1-covariance
# Perform calculation with uncorrelated source:
.jcall(teCalc,"V","initialise") # Initialise leaving the parameters the same
.jcall(teCalc,"V","setObservations", sourceArray2, destArray)
```

### Example 5 - Multivariate transfer entropy on binary data

example5TeBinaryMultivarTransfer.r - Multivariate transfer entropy (TE) calculation on binary data using the discrete TE calculator.

```
# Load the rJava library and start the JVM
library("rJava")
.jinit()
# Change location of jar to match yours:
# IMPORTANT -- If using the default below, make sure you have set the working of
  in R (e.g. with setwd()) to the location of this file (i.e. demos/r) !!
.jaddClassPath("../../infodynamics.jar")
# Generate some random binary data.
numObservations <- 100
sourceArray<-matrix(sample(0:1, numObservations*2, replace=TRUE), numObservations,</pre>
sourceArray2<-matrix(sample(0:1, numObservations*2, replace=TRUE), numObservations</pre>
# Destination variable takes a copy of the first bit of the source in bit 1,
# and an XOR of the two bits of the source in bit 2:
destArray <- cbind( c(0L, sourceArray[1:numObservations-1,1]), # column 1</pre>
                    c(OL, 1L*xor(sourceArray[1:numObservations-1,1],
                                  sourceArray[1:numObservations-1,2]))) # column
# Convert the 2D arrays to Java format:
sourceArrayJava <- .jarray(sourceArray, "[I", dispatch=TRUE)</pre>
sourceArray2Java <- .jarray(sourceArray2, "[I", dispatch=TRUE)</pre>
destArrayJava <- .jarray(destArray, "[I", dispatch=TRUE)</pre>
# Create a TE calculator and run it:
teCalc<-.jnew("infodynamics/measures/discrete/TransferEntropyCalculatorDiscrete"
.jcall(teCalc, "V", "initialise") # V for void return value
# We need to construct the joint values for the dest and source before we pass t
# and need to use the matrix conversion routine when calling from Matlab/Octave
mUtils<-.jnew("infodynamics/utils/MatrixUtils")</pre>
.jcall(teCalc, "V", "addObservations",
                .jcall(mUtils,"[I","computeCombinedValues", sourceArrayJava, 2L)
                 .jcall(mUtils,"[I","computeCombinedValues", destArrayJava, 2L))
result<-.jcall(teCalc, "D", "computeAverageLocalOfObservations")</pre>
cat("For source which the 2 bits are determined from, result should be close to
```

### **Example 6 - Dynamic dispatch with Mutual info calculator**

example6DynamicCallingMutualInfo.r - This example shows how to write R code to take advantage of the common interfaces defined for various information-theoretic calculators. Here, we use the common form of the

infodynamics.measures.continuous.MutualInfoCalculatorMultiVariate interface (which is never named here) to write common code into which we can plug one of three concrete implementations (kernel estimator, Kraskov estimator or linear-Gaussian estimator) by dynamically supplying the class name of the concrete implementation.

```
# Load the rJava library and start the JVM
library("rJava")
.jinit()
# Change location of jar to match yours:
# IMPORTANT -- If using the default below, make sure you have set the working of
    in R (e.g. with setwd()) to the location of this file (i.e. demos/r) !!
.jaddClassPath("../../infodynamics.jar")
# 1. Properties for the calculation (these are dynamically changeable, you could
     load them in from another properties file):
# The name of the data file (relative to this directory)
datafile <- "../data/4ColsPairedNoisyDependence-1.txt"</pre>
# List of column numbers for variables 1 and 2:
# (you can select any columns you wish to be contained in each variable)
variable1Columns \leftarrow c(1,2) # array indices start from 1 in R
variable2Columns \leftarrow c(3,4)
# The name of the concrete implementation of the interface
# infodynamics.measures.continuous.MutualInfoCalculatorMultiVariate
# which we wish to use for the calculation.
# Note that one could use any of the following calculators (try them all!):
  implementingClass <- "infodynamics/measures/continuous/kraskov/MutualInfoCalc</pre>
  implementingClass <- "infodynamics/measures/continuous/kernel/MutualInfoCalcu</pre>
   implementingClass <- "infodynamics/measures/continuous/gaussian/MutualInfoCal</pre>
implementingClass <- "infodynamics/measures/continuous/kraskov/MutualInfoCalcula"</pre>
```

```
#-----
# 2. Load in the data
data <- read.csv(datafile, header=FALSE, sep="")</pre>
# Pull out the columns from the data set which correspond to each of variable 1
variable1 <- data[, variable1Columns]</pre>
variable2 <- data[, variable2Columns]</pre>
# Extra step to extract the raw values from these data.frame objects:
variable1 <- apply(variable1, 2, function(x) as.numeric(x))</pre>
variable2 <- apply(variable2, 2, function(x) as.numeric(x))</pre>
#-----
# 3. Dynamically instantiate an object of the given class:
# (in fact, all java object creation in octave/matlab is dynamic - it has to be,
# since the languages are interpreted. This makes our life slightly easier at t
# point than it is in demos/java/example6LateBindingMutualInfo where we have to
miCalc<-.jnew(implementingClass)</pre>
#-----
# 4. Start using the MI calculator, paying attention to only
# call common methods defined in the interface type
# infodynamics.measures.continuous.MutualInfoCalculatorMultiVariate
# not methods only defined in a given implementation class.
# a. Initialise the calculator to use the required number of
   dimensions for each variable:
.jcall(miCalc,"V","initialise", length(variable1Columns), length(variable2Column
# b. Supply the observations to compute the PDFs from:
.jcall(miCalc, "V", "setObservations",
        .jarray(variable1, "[D", dispatch=TRUE),
        .jarray(variable2, "[D", dispatch=TRUE))
# c. Make the MI calculation:
miValue <- .jcall(miCalc, "D", "computeAverageLocalOfObservations")</pre>
cat("MI calculator", implementingClass, "\n computed the joint MI as ",
        miValue, "\n")
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