

# NISP Toolbox Manual

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## **Abstract**

This document is a brief introduction to the NISP module. We present the installation process of the module in binary from ATOMS or from the sources. We present the configuration functions and the randvar, setrandvar and polychaos classes.

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# Chapter 1

## Introduction

### 1.1 The OPUS project

The goal of this toolbox is to provide a tool to manage uncertainties in simulated models. This toolbox is based on the NISP library, where NISP stands for "Non-Intrusive Spectral Projection". This work has been realized in the context of the OPUS project,

<http://opus-project.fr>

"Open-Source Platform for Uncertainty treatments in Simulation", funded by ANR, the french "Agence Nationale pour la Recherche":

<http://www.agence-nationale-recherche.fr>

The toolbox is released under the Lesser General Public Licence (LGPL), as all components of the OPUS project.

### 1.2 The NISP library

The NISP library is based on a set of 3 C++ classes so that it provides an object-oriented framework for uncertainty analysis. The Scilab toolbox provides a pseudo-object oriented interface to this library, so that the two approaches are consistent. The NISP library is release under the LGPL licence.

The NISP library provides three tools, which are detailed below.

- The "randvar" class allows to manage random variables, specified by their distribution law and their parameters. Once a random variable is created, one can generate random numbers from the associated law.
- The "setrandvar" class allows to manage a collection of random variables. This collection is associated with a sampling method, such as MonteCarlo, Sobol, Quadrature, etc... It is possible to build the sample and to get it back so that the experiments can be performed.
- The "polychaos" class allows to manage a polynomial representation of the simulated model. One such object must be associated with a set of experiments which have been performed. This set may be read from a data file. The object is linked with a collection of random

variables. Then the coefficients of the polynomial can be computed by integration (quadrature). Once done, the mean, the variance and the Sobol indices can be directly computed from the coefficients.

The figure 1.1 presents the NISP methodology. The process requires that the user has a numerical solver, which has the form  $Y = f(X)$ , where  $X$  are input uncertain parameters and  $Y$  are output random variables. The method is based on the following steps.

- We begin by defining normalized random variables  $\xi$ . For example, we may use a random variables in the interval  $[0, 1]$  or a Normal random variable with mean 0 and variance 1. This choice allows to define the basis for the polynomial chaos, denoted by  $\{\Psi_k\}_{k \geq 0}$ . Depending on the type of random variable, the polynomials  $\{\Psi_k\}_{k \geq 0}$  are based on Hermite, Legendre or Laguerre polynomials.
- We can now define a Design Of Experiments (DOE) and, with random variable transformations rules, we get the physical uncertain parameters  $X$ . Several types of DOE are available: Monte-Carlo, Latin Hypercube Sampling, etc... If  $N$  experiments are required, the DOE define the collection of normalized random variables  $\{\xi_i\}_{i=1,N}$ . Transformation rules allows to compute the uncertain parameters  $\{X_i\}_{i=1,N}$ , which are the input of the numerical solver  $f$ .
- We can now perform the simulations, that is compute the collection of outputs  $\{Y_i\}_{i=1,N}$  where  $Y_i = f(X_i)$ .
- The variables  $Y$  are then projected on the polynomial basis and the coefficients  $y_k$  are computed by integration or regression.

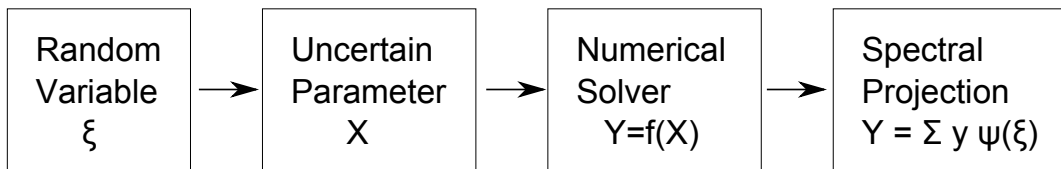


Figure 1.1: The NISP methodology

### 1.3 The NISP module

The NISP toolbox is available under the following operating systems:

- Linux 32 bits,
- Linux 64 bits,
- Windows 32 bits,
- Mac OS X.

The following list presents the features provided by the NISP toolbox.

- Manage various types of random variables:
  - uniform,
  - normal,
  - exponential,
  - log-normal.
- Generate random numbers from a given random variable,
- Transform an outcome from a given random variable into another,
- Manage various Design of Experiments for sets of random variables,
  - Monte-Carlo,
  - Sobol,
  - Latin Hypercube Sampling,
  - various samplings based on Smolyak designs.
- Manage polynomial chaos expansion and get specific outputs, including
  - mean,
  - variance,
  - quantile,
  - correlation,
  - etc...
- Generate the C source code which computes the output of the polynomial chaos expansion.

This User's Manual completes the online help provided with the toolbox, but does not replace it. The goal of this document is to provide both a global overview of the toolbox and to give some details about its implementation. The detailed calling sequence of each function is provided by the online help and will not be reproduced in this document. The inline help is presented in the figure 1.2.

For example, in order to access to the help associated with the `randvar` class, we type the following statements in the Scilab console.

```
help randvar
```

The previous statements opens the Help Browser and displays the helps page presented in figure

Several demonstration scripts are provided with the toolbox and are presented in the figure 1.4. These demonstrations are available under the "?" question mark in the menu of the Scilab console.

Finally, the unit tests provided with the toolbox cover all the features of the toolbox. When we want to know how to use a particular feature and do not find the information, we can search in the unit tests which often provide the answer.

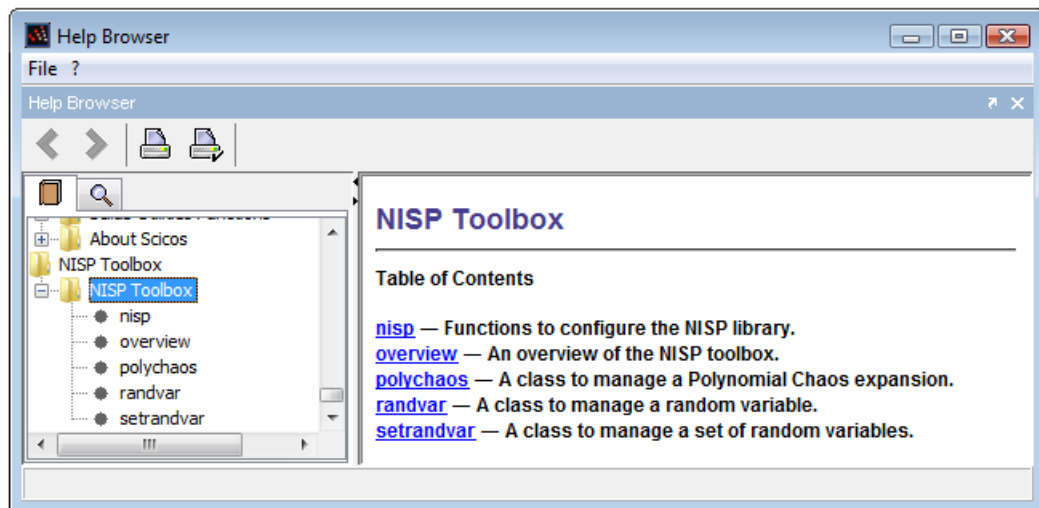


Figure 1.2: The NISP inline help.



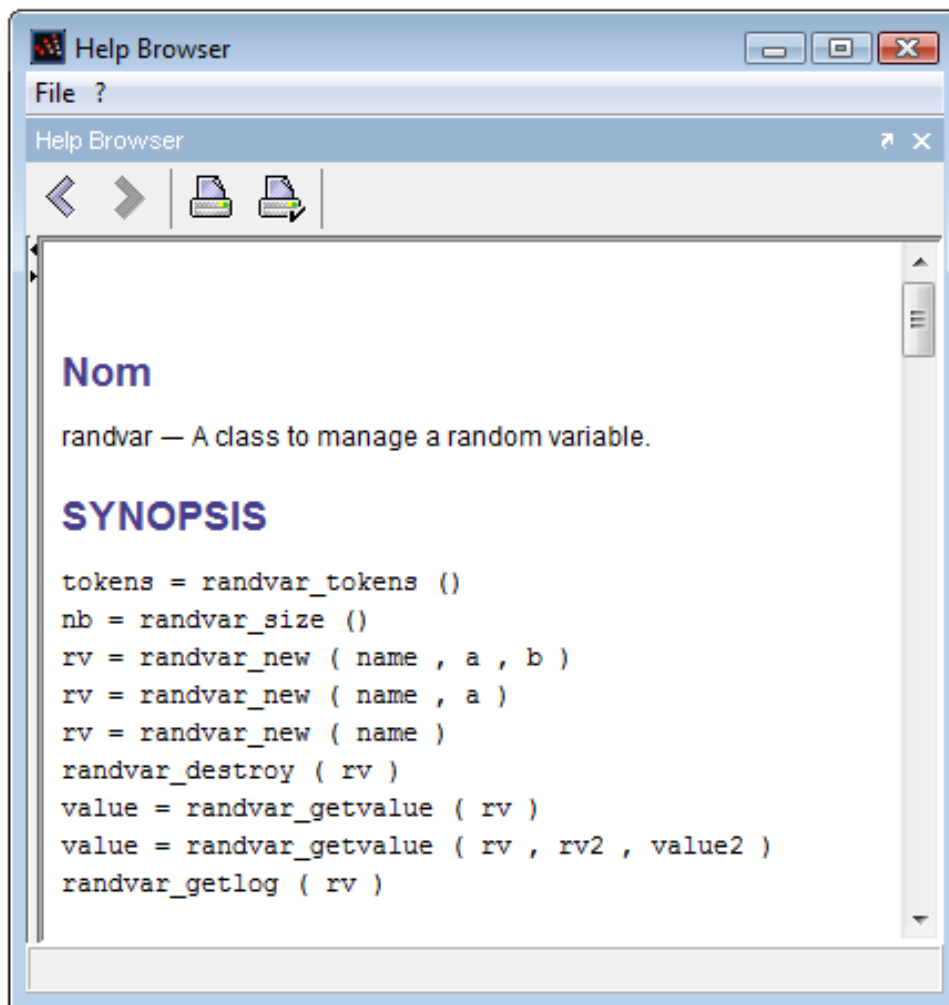


Figure 1.3: The online help of the randvar function.

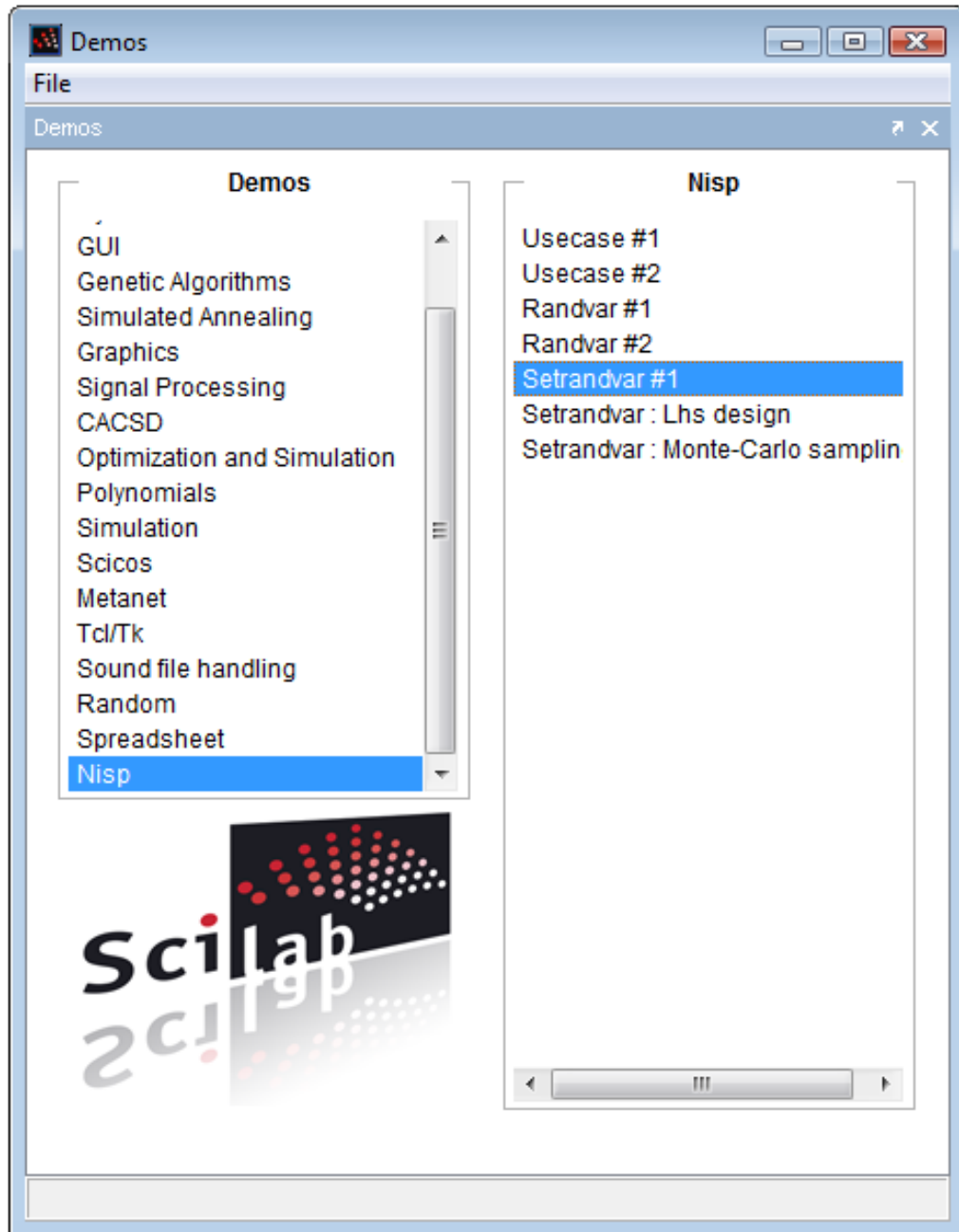


Figure 1.4: Demonstrations provided with the NISP toolbox.

# Chapter 2

## Theory

In this chapter, we (extremely) briefly present the theory which is used in the library. This section is a tutorial introduction to the NISP module.

### 2.1 Sensitivity analysis

In this section, we present the sensitivity analysis and emphasize the difference between global and local analysis.

Consider the model

$$\mathbf{Y} = f(\mathbf{X}), \quad (2.1)$$

where  $\mathbf{X} \in D_X \subset \mathbb{R}^p$  is the input and  $\mathbf{Y} \in D_Y \subset \mathbb{R}^m$  is the output of the model. The mapping  $f$  is presented in figure 2.1.

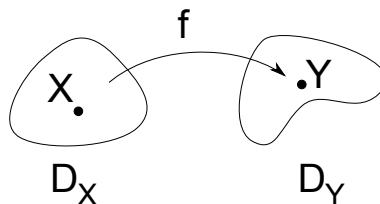


Figure 2.1: Global analysis.

We assume that the input  $\mathbf{X}$  is a random variable, so that the output variable  $\mathbf{Y}$  is also a random variable. We are interested in measuring the sensitivity of the output depending on the uncertainty of the input. More precisely, we are interested in knowing

- the input variables  $X_i$  which generate the most variability in the output  $\mathbf{Y}$ ,
- the input variables  $X_i$  which are not significant,
- a sub-space of the input variables where the variability is maximum,
- if input variables interact.

Consider the mapping presented in figure 2.1. The  $f$  mapping transforms the domain  $D_X$  into the domain  $D_Y$ . If  $f$  is sufficiently smooth, small perturbations of  $X$  generate small perturbations of  $Y$ . The local sensitivity analysis focuses on the behaviour of the mapping in the neighbourhood of a particular point  $X \in D_X$  toward a particular point  $Y \in D_Y$ . The global sensitivity analysis models the propagation of uncertainties transforming the whole set  $D_X$  into the set  $D_Y$ .

In the following, we assume that there is only one output variable so that  $Y \in \mathbb{R}$ .

There are two types of analysis that we are going to perform, that is uncertainty analysis and sensitivity analysis.

In uncertainty analysis, we assume that  $f_X$  is the probability density function of the variable  $\mathbf{X}$  and we are searching for the probability density function  $f_Y$  of the variable  $Y$  and by its cumulated density function  $F_Y(y) = P(Y \leq y)$ . This problem is difficult in the general case, and this is why we often are looking for an estimate of the expectation of  $Y$ , as defined by

$$E(Y) = \int_{D_X} y f_Y(y) dy, \quad (2.2)$$

and an estimate of its variance

$$Var(Y) = \int_{D_X} (y - E(Y))^2 f_Y(y) dy. \quad (2.3)$$

We might also be interested in the computation of the probability over a threshold.

In sensitivity analysis, we focus on the relative importance of the input variable  $X_i$  on the uncertainty of  $\mathbf{Y}$ . This way, we can order the input variables so that we can reduce the variability of the most important input variables, in order to, finally, reduce the variability of  $\mathbf{Y}$ .

More details on this topic can be found in the papers of Homma and Saltelli [1] or in the work of Sobol [4]. The Thesis by Jacques [2] presents an overview of sensitivity analysis.

## 2.2 Standardized regression coefficients of affine models

In this section, we present the standardized regression coefficients of an affine model.

Assume that the random variables  $X_i$  are independent, with mean  $E(X_i)$  and finite variances  $V(X_i)$ , for  $i = 1, 2, \dots, p$ . Let us consider the random variable  $Y$  as an affine function of the variables  $X_i$ :

$$Y = \beta_0 + \sum_{i=1,2,\dots,p} \beta_i X_i, \quad (2.4)$$

where  $\beta_i$  are real parameters, for  $i = 1, 2, \dots, p$ .

The expectation of the random variable  $Y$  is

$$E(Y) = \beta_0 + \sum_{i=1,2,\dots,p} \beta_i E(X_i). \quad (2.5)$$

Since the variables  $X_i$  are independent, the variance of the sum of variables is the sum of the variances. Hence,

$$V(Y) = V(\beta_0) + \sum_{i=1,2,\dots,p} V(\beta_i X_i), \quad (2.6)$$

which leads to the equality

$$V(Y) = \sum_{i=1,2,\dots,p} \beta_i^2 V(X_i). \quad (2.7)$$

Hence, each term  $\beta_i^2 V(X_i)$  is the part of the total variance  $V(Y)$  which is caused by the variable  $X_i$ .

The standardized regression coefficient is defined as

$$SRC_i = \frac{\beta_i^2 V(X_i)}{V(Y)}, \quad (2.8)$$

for  $i = 1, 2, \dots, p$ .

Hence, the sum of the standardized regression coefficients is one:

$$SRC_1 + SRC_2 + \dots + SRC_p = 1. \quad (2.9)$$

## 2.3 Link with the linear correlation coefficients

In this section, we present the link between the linear correlation coefficients of an affine model, and the standardized regression coefficients.

Assume that the random variables  $X_i$  are independent, with mean  $E(X_i)$  and finite variances  $V(X_i)$ , for  $i = 1, 2, \dots, p$ . Let us consider the random variable  $Y$ , which depends linearly on the variables  $X_i$  by the relationship 2.4.

The linear correlation coefficient between  $Y$  and  $X_i$  is

$$\rho_{Y, X_i} = \frac{Cov(Y, X_i)}{\sqrt{V(Y)}\sqrt{V(X_i)}}, \quad (2.10)$$

for  $i = 1, 2, \dots, p$ . In the particular case of the affine model 2.4, we have

$$Cov(Y, X_i) = Cov(\beta_0, X_i) + \beta_1 Cov(X_1, X_i) + \beta_2 Cov(X_2, X_i) + \dots + \quad (2.11)$$

$$\beta_i Cov(X_i, X_i) + \dots + \beta_p Cov(X_p, X_i). \quad (2.12)$$

$$(2.13)$$

Since the random variables  $X_i$  are independent, we have  $Cov(X_j, X_i) = 0$ , for any  $j \neq i$ . Therefore,

$$Cov(Y, X_i) = \beta_i Cov(X_i, X_i) \quad (2.14)$$

$$= \beta_i V(X_i). \quad (2.15)$$

Hence, the correlation coefficient can be simplified into

$$\rho_{Y, X_i} = \frac{\beta_i V(X_i)}{\sqrt{V(Y)}\sqrt{V(X_i)}} \quad (2.16)$$

$$= \frac{\beta_i \sqrt{V(X_i)}}{\sqrt{V(Y)}}. \quad (2.17)$$

We square the previous equality and get

$$\rho_{Y,X_i}^2 = \frac{\beta_i^2 V(X_i)}{V(Y)}. \quad (2.18)$$

Therefore, the square of the linear correlation coefficient is equal to the first order sensitivity index, i.e.

$$\rho_{Y,X_i}^2 = SRC_i. \quad (2.19)$$

## 2.4 Using scatter plots

In this section, we present an example of an affine model, where the difference between local and global sensitivity is made clearer by the use of scatter plots.

Assume four independent random variables  $X_i$ , for  $i = 1, 2, 3, 4$ . We assume that the variables  $X_i$  are normally distributed, with zero mean and  $i^2$  variance.

Let us consider the affine model

$$Y = X_1 + X_2 + X_3 + X_4. \quad (2.20)$$

Notice that the derivative of  $Y$  with respect to any of its input is equal to one, i.e.

$$\frac{\partial Y}{\partial X_i} = 1, \quad (2.21)$$

for  $i = 1, 2, 3, 4$ . This means that, locally, the inputs all have the same effect on the output.

Let us compute the standardized regression coefficients of this model. By hypothesis, the variance of each variable is

$$V(X_1) = 1, \quad V(X_2) = 4, \quad (2.22)$$

$$V(X_3) = 9, \quad V(X_4) = 16. \quad (2.23)$$

Since the variables are independent, the variance of the output  $Y$  is

$$V(Y) = V(X_1) + V(X_2) + V(X_3) + V(X_4) = 30. \quad (2.24)$$

The standardized regression coefficient is

$$SRC_i = \frac{\beta_i^2 V(X_i)}{V(Y)} = \frac{i^2}{30}, \quad (2.25)$$

for  $i = 1, 2, 3, 4$ . More specifically, we have

$$SRC_1 = \frac{1}{30}, \quad SRC_2 = \frac{4}{30}, \quad (2.26)$$

$$SRC_3 = \frac{9}{30}, \quad SRC_4 = \frac{16}{30}. \quad (2.27)$$

We have the following inequalities:

$$SRC_4 > SRC_3 > SRC_2 > SRC_1. \quad (2.28)$$

This means that the variable which causes the most variance in the output is  $X_4$ , while the variable which causes the least variance in the output is  $X_1$ .

The script below performs the analysis with the NISP module. The sampling is based on a Latin Hypercube Sampling design with 5000 points.

```

function y = Exemple (x)
    y(:) = x(:,1) + x(:,2) + x(:,3) + x(:,4)
endfunction

function r = lincorrcoef ( x , y )
    // Returns the linear correlation coefficient of x and y.
    // The variables are expected to be column matrices with the same size.
    x = x(:)
    y = y(:)
    mx = mean(x)
    my = mean(y)
    sx = sqrt(sum((x-mx).^2))
    sy = sqrt(sum((y-my).^2))
    r = (x-mx)'*(y-my) / sx / sy
endfunction

// Initialisation de la graine aleatoire
nisp_initseed ( 0 );

// Create the random variables.
rvu1 = randvar_new("Normale",0,1);
rvu2 = randvar_new("Normale",0,2);
rvu3 = randvar_new("Normale",0,3);
rvu4 = randvar_new("Normale",0,4);
srvu = setrandvar_new();
setrandvar_addrandvar ( srvu, rvu1);
setrandvar_addrandvar ( srvu, rvu2);
setrandvar_addrandvar ( srvu, rvu3);
setrandvar_addrandvar ( srvu, rvu4);
// Create a sampling by a Latin Hypercube Sampling with size 5000.
nbshots = 5000;
setrandvar_buildsample(srvu, "Lhs",nbshots);
sampling = setrandvar_getsample(srvu);
// Perform the experiments.
y=Exemple(sampling);
// Scatter plots : y depending on X_i
for k=1:4
    scf();
    plot(y,sampling(:,k),'rx');
    xistr="X"+string(k);
    xtitle("Scatter_plot_for_"+xistr,xistr,"Y");
end
// Compute the sample linear correlation coefficients
rho1 = lincorrcoef ( sampling(:,1) , y );
SRC1 = rho1^2;
SRC1expected = 1/30;
mprintf("SRC_1=%.5f (expected=%.5f)\n",SRC1,SRC1expected);
//
rho2 = lincorrcoef ( sampling(:,2) , y );

```

```

SRC2 = rho2^2;
SRC2expected = 4/30;
mprintf("SRC_2=%.5f (expected=%.5f)\n", SRC2, SRC2expected);
//
rho3 = lincorrcoef ( sampling(:,3) , y );
SRC3 = rho3^2;
SRC3expected = 9/30;
mprintf("SRC_3=%.5f (expected=%.5f)\n", SRC3, SRC3expected);
//
rho4 = lincorrcoef ( sampling(:,4) , y );
SRC4 = rho4^2;
SRC4expected = 16/30;
mprintf("SRC_4=%.5f (expected=%.5f)\n", SRC4, SRC4expected);
//
SUM = SRC1 + SRC2 + SRC3 + SRC4;
SUMexpected = 1;
mprintf("SUM=%.5f (expected=%.5f)\n", SUM, SUMexpected);
//
// Clean-up
randvar_destroy(rvu1);
randvar_destroy(rvu2);
randvar_destroy(rvu3);
randvar_destroy(rvu4);
setrandvar_destroy(srvu);

```

The previous script produces the following output.

```

SRC_1=0.03538 (expected=0.03333)
SRC_2=0.12570 (expected=0.13333)
SRC_3=0.31817 (expected=0.30000)
SRC_4=0.54314 (expected=0.53333)
SUM=1.00066 (expected=1.00000)

```

The previous script also produces the plots [2.2](#), [2.3](#), [2.4](#) and [2.5](#).

## 2.5 Sensitivity analysis for nonlinear models

Let us focus on one particular input  $X_i$  of the model  $f$ , with  $i = 1, 2, \dots, p$ . If we set  $X_i$  to a particular value, say  $x_i$  for example, then the variance of the output  $Y$  surely decreases, because the variable  $X_i$  is not random anymore. We can then measure the conditionnal variance given  $X_i$ , denoted by  $V(Y|X_i = x_i)$ .

Since  $X_i$  is a random variable, the conditionnal variance  $V(Y|X_i = x_i)$  is a random variable. We are interested in the average value of this variance, that is, we are interested in  $E(V(Y|X_i))$ . If  $X_i$  has a large weight in the variance  $V(Y)$ , then  $E(V(Y|X_i))$  is small.

The theorem of the total variance states that, if  $V(Y)$  is finite, then

$$V(Y) = V(E(Y|X_i)) + E(V(Y|X_i)). \quad (2.29)$$

If  $X_i$  has a large weight in the variance  $V(Y)$ , then  $V(E(Y|X_i))$  is large.



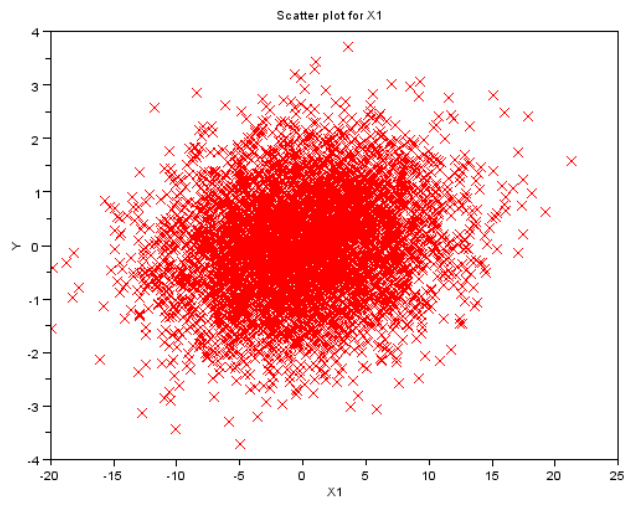


Figure 2.2: Scatter plot for an affine model - Variable  $X_1$ .

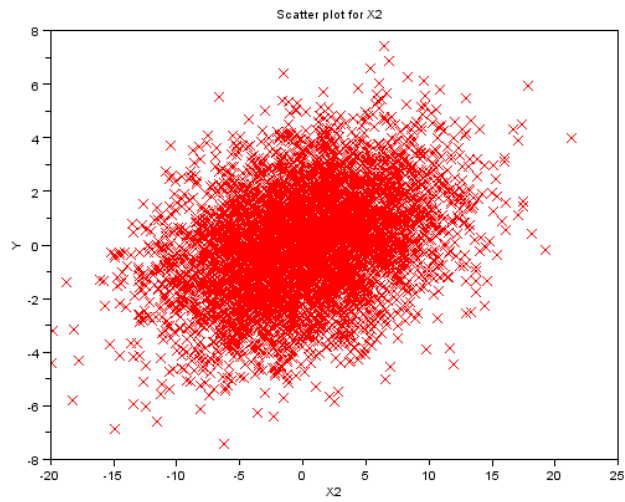


Figure 2.3: Scatter plot for an affine model - Variable  $X_2$ .

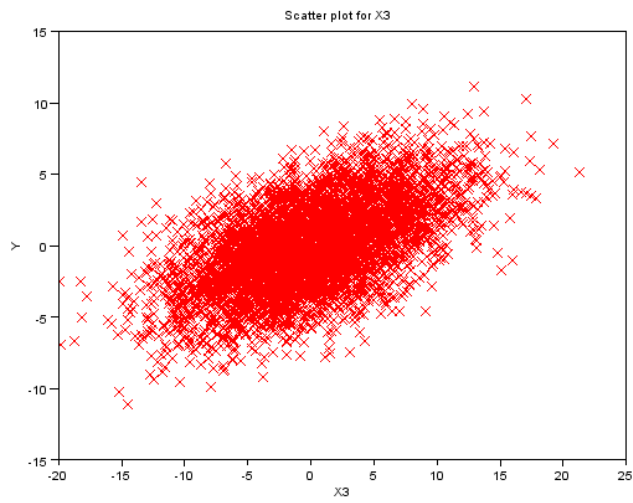


Figure 2.4: Scatter plot for an affine model - Variable  $X_3$ .

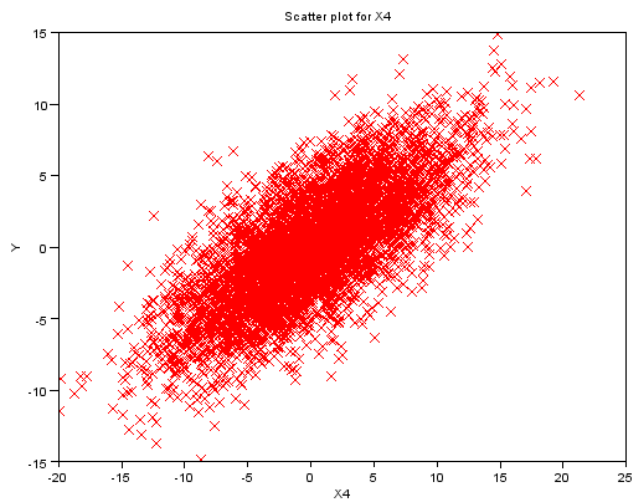


Figure 2.5: Scatter plot for an affine model - Variable  $X_4$ .

The first order sensitivity indice of  $Y$  to the variable  $X_i$  is defined by

$$S_i = \frac{V(E(Y|X_i))}{V(Y)}, \quad (2.30)$$

for  $i = 1, 2, \dots, p$ . The sensitivity indice measures the part of the variance which is caused by the uncertainty in  $X_i$ .

We can compute the sensitivity indice when the function  $f$  is linear. Assume that the output  $Y$  depends linearly on the input  $X_i$ :

$$Y = \beta_0 + \sum_{i=1,2,\dots,p} \beta_i X_i, \quad (2.31)$$

where  $\beta_i \in \mathbb{R}$ , for  $i = 0, 1, 2, \dots, p$ . Then

$$E(Y|X_i) = \beta_0 + \sum_{i=1,2,\dots,i-1,i+1,\dots,p} \beta_i E(X_i) + \beta_i X_i, \quad (2.32)$$

since the expectation of a sum is the sum of expectations. Then,

$$V(E(Y|X_i)) = V(\beta_i X_i) \quad (2.33)$$

$$= \beta_i^2 V(X_i), \quad (2.34)$$

since the variance of a constant term is zero. Therefore, the sensitivity index of  $Y$  to the variable  $X_i$  is

$$S_i = \frac{\beta_i^2 V(X_i)}{V(Y)}, \quad (2.35)$$

for  $i = 1, 2, \dots, p$ .

Hence, if we make the assumption that a model is affine, then the empirical linear correlation coefficient can be used to estimate the sensitivity indices.

## 2.6 The effect of the interactions

In this example, we consider a non-linear, non-additive model made of the product of two independent random variables. The goal of this example is to show that, in some cases, we have to consider the interactions between the variables.

Consider the function

$$Y = X_1 X_2, \quad (2.36)$$

where  $X_1$  and  $X_2$  are two independent normal random variables with mean  $\mu_1$  and  $\mu_2$  and variances  $\sigma_1^2$  and  $\sigma_2^2$ .

Let us compute the expectation of the random variable  $Y$ . The expectation of  $Y$  is

$$E(Y) = \int_{-\infty}^{\infty} \int_{-\infty}^{\infty} X_1 X_2 F(X_1, X_2) dx_1 dx_2, \quad (2.37)$$

where  $F(x_1, x_2)$  is the joint probability distribution function of the variables  $X_1$  and  $X_2$ . Since  $X_1$  and  $X_2$  are independent variables, we have

$$F(x_1, x_2) = F_1(X_1)F_2(X_2), \quad (2.38)$$

where  $F_1$  is the probability distribution function of  $X_1$  and  $F_2$  is the probability distribution function of  $X_2$ . Then, we have

$$E(Y) = \int_{-\infty}^{\infty} \int_{-\infty}^{\infty} X_1 X_2 F_1(X_1) F_2(X_2) dx_1 dx_2 \quad (2.39)$$

$$= \left( \int_{-\infty}^{\infty} X_1 F_1(X_1) dx_1 \right) \left( \int_{-\infty}^{\infty} X_2 F_2(X_2) dx_2 \right). \quad (2.40)$$

$$= E(X_1)E(X_2). \quad (2.41)$$

Therefore,

$$E(Y) = \mu_1 \mu_2. \quad (2.42)$$

The variance of  $Y$  is

$$V(Y) = E(Y^2) - E(Y)^2. \quad (2.43)$$

The expectation  $E(Y^2)$  is

$$E(Y^2) = \int_{-\infty}^{\infty} \int_{-\infty}^{\infty} (X_1 X_2)^2 F(x_1, x_2) dx_1 dx_2 \quad (2.44)$$

$$= \int_{-\infty}^{\infty} \int_{-\infty}^{\infty} (X_1 X_2)^2 F_1(X_1) F_2(X_2) dx_1 dx_2 \quad (2.45)$$

$$= \left( \int_{-\infty}^{\infty} X_1^2 F_1(X_1) dx_1 \right) \left( \int_{-\infty}^{\infty} X_2^2 F_2(X_2) dx_2 \right), \quad (2.46)$$

$$= E(X_1^2)E(X_2^2). \quad (2.47)$$

Now, we have

$$V(X_1) = E(X_1^2) - E(X_1)^2, \quad (2.48)$$

$$V(X_2) = E(X_2^2) - E(X_2)^2, \quad (2.49)$$

which leads to

$$E(X_1^2) = V(X_1) + E(X_1)^2, \quad (2.50)$$

$$E(X_2^2) = V(X_2) + E(X_2)^2. \quad (2.51)$$

Therefore,

$$E(Y^2) = (V(X_1) + E(X_1)^2)(V(X_2) + E(X_2)^2) \quad (2.52)$$

$$= (\sigma_1^2 + \mu_1^2)(\sigma_2^2 + \mu_2^2). \quad (2.53)$$

Finally, we get

$$V(Y) = (\sigma_1^2 + \mu_1^2)(\sigma_2^2 + \mu_2^2) - (\mu_1 \mu_2)^2. \quad (2.54)$$

We can expand the previous equality and get

$$V(Y) = \sigma_1^2\sigma_2^2 + \sigma_1^2\mu_2^2 + \mu_1^2\sigma_2^2 + \mu_1^2\mu_2^2 - (\mu_1\mu_2)^2. \quad (2.55)$$

The last two terms of the previous equality can be simplified, so that we get

$$V(Y) = \sigma_1^2\sigma_2^2 + \sigma_1^2\mu_2^2 + \mu_1^2\sigma_2^2. \quad (2.56)$$

The sensitivity indices can be computed from the definitions

$$S_1 = \frac{V(E(Y|X_1))}{V(Y)}, \quad (2.57)$$

$$S_2 = \frac{V(E(Y|X_2))}{V(Y)}. \quad (2.58)$$

We have  $E(Y|X_1) = E(X_2)X_1 = \mu_2X_1$ . Similarly,  $E(Y|X_2) = \mu_1X_2$ . Hence

$$S_1 = \frac{V(\mu_2X_1)}{V(Y)}, \quad (2.59)$$

$$S_2 = \frac{V(X_2)}{V(Y)}. \quad (2.60)$$

We get

$$S_1 = \frac{\mu_2^2V(X_1)}{V(Y)}, \quad (2.61)$$

$$S_2 = \frac{\mu_1^2V(X_2)}{V(Y)}. \quad (2.62)$$

Finally, the first order sensitivity indices are

$$S_1 = \frac{\mu_2^2\sigma_1^2}{V(Y)}, \quad (2.63)$$

$$S_2 = \frac{\mu_1^2\sigma_2^2}{V(Y)}. \quad (2.64)$$

Since  $\sigma_1^2\sigma_2^2 \geq 0$ , we have

$$\mu_2^2\sigma_1^2 + \mu_1^2\sigma_2^2 \leq V(Y) = \sigma_1^2\sigma_2^2 + \sigma_1^2\mu_2^2 + \mu_1^2\sigma_2^2. \quad (2.65)$$

We divide the previous inequality by  $V(Y)$ , and get

$$\frac{\mu_2^2\sigma_1^2}{V(Y)} + \frac{\mu_1^2\sigma_2^2}{V(Y)} \leq 1. \quad (2.66)$$

Therefore, the sum of the first order sensitivity indices satisfies the inequality

$$S_1 + S_2 \leq 1. \quad (2.67)$$

Hence, in this example, one part of the variance  $V(Y)$  cannot be explained neither by  $X_1$  alone, or by  $X_2$  alone, because it is caused by the interactions between  $X_1$  and  $X_2$ . We define by  $S_{12}$  the sensitivity index associated with the group of variables  $(X_1, X_2)$  as

$$S_{12} = 1 - S_1 - S_2 = \frac{\sigma_1^2 \sigma_2^2}{V(Y)}. \quad (2.68)$$

The following Scilab script performs the sensitivity analysis on the previous example. We consider two normal variables, where the first variable has mean 1.5 and standard deviation 0.5 while the second variable has mean 3.5 and standard deviation 2.5.

```
function y = Exemple (x)
    y(:,1) = x(:,1) .* x(:,2);
endfunction
// First variable
// Normal
mu1 = 1.5;
sigma1 = 0.5;
// Second variable
// Normal
mu2 = 3.5;
sigma2 = 2.5;

// 1. Two stochastic (normalized) variables
vx1 = randvar_new("Normale");
vx2 = randvar_new("Normale");
// 2. A collection of stoch. variables.
srvx = setrandvar_new();
setrandvar_addrandvar ( srvx,vx1);
setrandvar_addrandvar ( srvx,vx2);
// 3. Two uncertain parameters
vu1 = randvar_new("Normale",mu1,sigma1);
vu2 = randvar_new("Normale",mu2,sigma2);
// 4. A collection of uncertain parameters
srvu = setrandvar_new();
setrandvar_addrandvar ( srvu,vu1);
setrandvar_addrandvar ( srvu,vu2);
// 5. Create the Design Of Experiments
degre = 2;
setrandvar_buildsample(srvx,"Quadrature",degre);
setrandvar_buildsample( srvu , srvx );
// 6. Create the polynomial
ny = 1;
pc = polychaos_new ( srvx , ny );
np = setrandvar_getsize(srvx);
mprintf("Number of experiments : %d\n",np);
polychaos_setsizetarget(pc,np);
// 7. Perform the DOE
inputdata = setrandvar_getsample(srvu);
outputdata = Exemple(inputdata);
```

```

polychaos_settarget(pc,outputdata);
// 8. Compute the coefficients of the polynomial expansion
polychaos_setdegree(pc,degree);
polychaos_computexp(pc,srvx,"Integration");
// 9. Get the sensitivity indices
average = polychaos_getmean(pc);
Ey_expectation= mu1*mu2;
var = polychaos_getvariance(pc);
Vy_expectation = mu2^2*sigma1^2 + mu1^2*sigma2^2 + sigma1^2*sigma2^2;
mprintf("Mean_____=%f_(expectation_=%f)\n",average,Ey_expectation);
mprintf("Variance_____=%f_(expectation_=%f)\n",var,Vy_expectation);
mprintf("First_order_sensitivity_index\n");
S1 = polychaos_getindexfirst(pc,1);
S1_expectation = ( mu2^2*sigma1^2 ) / Vy_expectation;
mprintf("_____Variable_X1_=%f_(expectation_=%f)\n",S1,S1_expectation);
re = abs(S1- S1_expectation)/S1_expectation;
mprintf("_____Relative_Error_=%f\n", re);
S2 = polychaos_getindexfirst(pc,2);
S2_expectation = ( mu1^2*sigma2^2 ) / Vy_expectation;
mprintf("_____Variable_X2_=%f_(expectation_=%f)\n",S2,S2_expectation);
re = abs(S2- S2_expectation)/S2_expectation;
mprintf("_____Relative_Error_=%f\n", re);

mprintf("Total_sensitivity_index\n");
ST1 = polychaos_getindextotal(pc,1);
mprintf("_____Variable_X1_=%f\n",ST1);
ST2 = polychaos_getindextotal(pc,2);
mprintf("_____Variable_X2_=%f\n",ST2);
// Clean-up
polychaos_destroy(pc);
randvar_destroy(vu1);
randvar_destroy(vu2);
randvar_destroy(vx1);
randvar_destroy(vx2);
setrandvar_destroy(srvu);
setrandvar_destroy(srvx);

```

The previous script produces the following output.

```

Mean      = 5.250000 (expectation = 5.250000)
Variance  = 18.687500 (expectation = 18.687500)
First order sensitivity index
    Variable X1 = 0.163880 (expectation = 0.163880)
        Relative Error = 0.000000
    Variable X2 = 0.752508 (expectation = 0.752508)
        Relative Error = 0.000000
Total sensitivity index
    Variable X1 = 0.247492
    Variable X2 = 0.836120

```

We see that the polynomial chaos performs an exact computation.

## 2.7 Sobol decomposition

Sobol [4] introduced the sensitivity index based on  $V(E(Y|X_i))$  by decomposing the function  $f$  as a sum of function with an increasing number of parameters.

We consider the function  $f$

$$\mathbf{Y} = f(x_1, x_2, \dots, x_p), \quad (2.69)$$

where  $\mathbf{X} = (x_1, \dots, x_p) \in [0, 1]^p$ . If  $f$  can be integrated in  $[0, 1]^p$ , then there is a unique decomposition

$$\mathbf{Y} = f_0 + \sum_{i=1,2,\dots,p} f_i(x_i) + \sum_{1 \leq i < j \leq p} f_{i,j}(x_i, x_j) + \dots + f_{1,2,\dots,p}(x_1, x_2, \dots, x_p), \quad (2.70)$$

where  $f_0$  is a constant and the function of the decomposition satisfy the equalities

$$\int_0^1 f_{i_1, \dots, i_s}(x_{i_1}, \dots, x_{i_s}) dx_{i_k} = 0, \quad (2.71)$$

for any  $k = 1, 2, \dots, s$  and any indices  $\{i_1, \dots, i_s\} \subset \{1, \dots, p\}$ .

## 2.8 Ishigami function

TODO



# Chapter 3

## Installation

In this section, we present the installation process for the toolbox. We present the steps which are required to have a running version of the toolbox and presents the several checks which can be performed before using the toolbox.

### 3.1 Introduction

There are two possible ways of installing the NISP toolbox in Scilab:

- use the ATOMS system and get a binary version of the toolbox,
- build the toolbox from the sources.

The next two sections present these two ways of using the toolbox.

Before getting into the installation process, let us present some details of the the internal components of the toolbox. The following list is an overview of the content of the directories:

- *tbxnisp/demos* : demonstration scripts
- *tbxnisp/doc* : the documentation
- *tbxnisp/doc/usermanual* : the L<sup>A</sup>T<sub>E</sub>Xsources of this manual
- *tbxnisp/etc* : startup and shutdown scripts for the toolbox
- *tbxnisp/help* : inline help pages
- *tbxnisp/macros* : Scilab macros files \*.sci
- *tbxnisp/sci\_gateway* : the sources of the gateway
- *tbxnisp/src* : the sources of the NISP library
- *tbxnisp/tests* : tests
- *tbxnisp/tests/nonreg\_tests* : tests after some bug has been identified
- *tbxnisp/tests/unit\_tests* : unit tests

The current version is based on the NISP Library v2.1.

## 3.2 Installing the toolbox from ATOMS

The ATOMS component is the Scilab tool which allows to search, download, install and load toolboxes. ATOMS comes with Scilab v5.2. The Scilab-NISP toolbox has been packaged and is provided mainly by the ATOMS component. The toolbox is provided in binary form, depending on the user's operating system. The Scilab-NISP toolbox is available for the following platforms:

- Windows 32 bits,
- Linux 32 bits, 64 bits,
- Mac OS X.

The ATOMS component allows to use a toolbox based on compiled source code, without having a compiler installed in the system.

Installing the Scilab-NISP toolbox from ATOMS requires the following steps:

- `atomsList()`: prints the list of current toolboxes,
- `atomsShow()`: prints informations about a toolbox,
- `atomsInstall()`: installs a toolbox on the system,
- `atomsLoad()`: loads a toolbox.

Once installed and loaded, the toolbox will be available on the system from session to session, so that there is no need to load the toolbox again: it will be available right from the start of the session.

In the following Scilab session, we use the `atomsList()` function to print the list of all ATOMS toolboxes.

```
--> atomsList()
ANN_Toolbox - ANN Toolbox
dde_toolbox - Dynamic Data Exchange client for Scilab
module_lycee - Scilab pour les lycées
  NISP - Non Intrusive Spectral Projection
  plotlib - "Matlab-like" Plotting library for Scilab
  scipad - Scipad 7.20
sndfile_toolbox - Read & write sound files
stixbox - Statistics toolbox for Scilab 5.2
```

In the following Scilab session, we use the `atomsShow()` function to print the details about the NISP toolbox.

```
-->atomsShow("NISP")
Package : NISP
Title : NISP
Summary : Non Intrusive Spectral Projection
Version : 2.1
Depend : Category(ies) : Optimization
Maintainer(s) : Pierre Marechal <pierre.marechal@scilab.org>
                Michael Baudin <michael.baudin@scilab.org>
```

```

Entity : CEA / DIGITEO
WebSite :          License : LGPL
Scilab Version : >= 5.2.0
Status : Not installed
Description : This toolbox allows to approximate a given model,
              which is associated with input random variables.
              This toolbox has been created in the context of the
              OPUS project :
                  http://opus-project.fr/
              within the workpackage 2.1.1:
                  "Construction de mÃl'ta-modÃiles"
              This project has received funding by Agence Nationale
              de la recherche :
                  http://www.agence-nationale-recherche.fr/
              See in the help provided in the help/en_US directory
              of the toolbox for more information about its use.
              Use cases are presented in the demos directory.

```

In the following Scilab session, we use the `atomsInstall()` function to download and install the binary version of the toolbox corresponding to the current operating system.

```

-->atomsInstall ( "NISP" )
ans =
!NISP 2.1 allusers D:\Programs\SC3623~1\contrib\NISP\2.1 I !
The "allusers" option of the atomsInstall function can be used to install the toolbox for all
the users of this computer. We finally load the toolbox with the atomsLoad() function.
-->atomsLoad("NISP")
Start NISP Toolbox
    Load gateways
    Load help
    Load demos
ans =
!NISP 2.1 D:\Programs\SC3623~1\contrib\NISP\2.1 !

```

Now that the toolbox is loaded, it will be automatically loaded at the next Scilab session.

### 3.3 Installing the toolbox from the sources

In this section, we present the steps which are required in order to install the toolbox from the sources.

In order to install the toolbox from the sources, a compiler is required to be installed on the machine. This toolbox can be used with Scilab v5.1 and Scilab v5.2. We suppose that the archive has been unpacked in the "tbxnisp" directory. The following is a short list of the steps which are required to setup the toolbox.

1. build the toolbox : run the *tbxnisp/builder.sce* script to create the binaries of the library, create the binaries for the gateway, generate the documentation
2. load the toolbox : run the *tbxnisp/load.sce* script to load all commands and setup the documentation

3. setup the startup configuration file of your Scilab system so that the toolbox is known at startup (see below for details),
4. run the unit tests : run the *tbxnisp/runtests.sce* script to perform all unit tests and check that the toolbox is OK
5. run the demos : run the *tbxnisp/rundemos.sce* script to run all demonstration scripts and get a quick interactive overview of its features

The following script presents the messages which are generated when the builder of the toolbox is launched. The builder script performs the following steps:

- compile the NISP C++ library,
- compile the C++ gateway library (the glue between the library and Scilab),
- generate the Java help files from the .xml files,
- generate the loader script.

```
-->exec C:\tbxnisp\builder.sce;
Building sources...
  Generate a loader file
  Generate a Makefile
  Running the Makefile
  Compilation of utils.cpp
  Compilation of blas1_d.cpp
  Compilation of dcdflib.cpp
  Compilation of faure.cpp
  Compilation of halton.cpp
  Compilation of linpack_d.cpp
  Compilation of niederreiter.cpp
  Compilation of reversehalton.cpp
  Compilation of sobol.cpp
  Building shared library (be patient)
  Generate a cleaner file
  Generate a loader file
  Generate a Makefile
  Running the Makefile
  Compilation of nisp_gc.cpp
  Compilation of nisp_gva.cpp
  Compilation of nisp_ind.cpp
  Compilation of nisp_index.cpp
  Compilation of nisp_inv.cpp
  Compilation of nisp_math.cpp
  Compilation of nisp_msg.cpp
  Compilation of nisp_conf.cpp
  Compilation of nisp_ort.cpp
  Compilation of nisp_pc.cpp
  Compilation of nisp_polyrule.cpp
```

```
Compilation of nisp_qua.cpp
Compilation of nisp_random.cpp
Compilation of nisp_smo.cpp
Compilation of nisp_util.cpp
Compilation of nisp_va.cpp
Compilation of nisp_smolyak.cpp
Building shared library (be patient)
Generate a cleaner file
Building gateway...
Generate a gateway file
Generate a loader file
Generate a Makefile: Makelib
Running the makefile
Compilation of nisp_gettoken.cpp
Compilation of nisp_gwsupport.cpp
Compilation of nisp_PolynomialChaos_map.cpp
Compilation of nisp_RandomVariable_map.cpp
Compilation of nisp_SetRandomVariable_map.cpp
Compilation of sci_nisp_startup.cpp
Compilation of sci_nisp_shutdown.cpp
Compilation of sci_nisp_verboselevelset.cpp
Compilation of sci_nisp_verboselevelget.cpp
Compilation of sci_nisp_initseed.cpp
Compilation of sci_randvar_new.cpp
Compilation of sci_randvar_destroy.cpp
Compilation of sci_randvar_size.cpp
Compilation of sci_randvar_tokens.cpp
Compilation of sci_randvar_getlog.cpp
Compilation of sci_randvar_getvalue.cpp
Compilation of sci_setrandvar_new.cpp
Compilation of sci_setrandvar_tokens.cpp
Compilation of sci_setrandvar_size.cpp
Compilation of sci_setrandvar_destroy.cpp
Compilation of sci_setrandvar_freememory.cpp
Compilation of sci_setrandvar_addrandvar.cpp
Compilation of sci_setrandvar_getlog.cpp
Compilation of sci_setrandvar_getdimension.cpp
Compilation of sci_setrandvar_getsize.cpp
Compilation of sci_setrandvar_getsample.cpp
Compilation of sci_setrandvar_setsample.cpp
Compilation of sci_setrandvar_save.cpp
Compilation of sci_setrandvar_buildsample.cpp
Compilation of sci_polychaos_new.cpp
Compilation of sci_polychaos_destroy.cpp
Compilation of sci_polychaos_tokens.cpp
Compilation of sci_polychaos_size.cpp
Compilation of sci_polychaos_setdegree.cpp
Compilation of sci_polychaos_getdegree.cpp
Compilation of sci_polychaos_freememory.cpp
```

```

Compilation of sci_polychaos_getdimoutput.cpp
Compilation of sci_polychaos_setdimoutput.cpp
Compilation of sci_polychaos_getsizetarget.cpp
Compilation of sci_polychaos_setsizetarget.cpp
Compilation of sci_polychaos_freememtarget.cpp
Compilation of sci_polychaos_settarget.cpp
Compilation of sci_polychaos_gettarget.cpp
Compilation of sci_polychaos_getdiminput.cpp
Compilation of sci_polychaos_getdimexp.cpp
Compilation of sci_polychaos_getlog.cpp
Compilation of sci_polychaos_computeexp.cpp
Compilation of sci_polychaos_getmean.cpp
Compilation of sci_polychaos_getvariance.cpp
Compilation of sci_polychaos_getcovariance.cpp
Compilation of sci_polychaos_getcorrelation.cpp
Compilation of sci_polychaos_getindexfirst.cpp
Compilation of sci_polychaos_getindextotal.cpp
Compilation of sci_polychaos_getmultind.cpp
Compilation of sci_polychaos_getgroupind.cpp
Compilation of sci_polychaos_setgroupempty.cpp
Compilation of sci_polychaos_getgroupinter.cpp
Compilation of sci_polychaos_getinvquantile.cpp
Compilation of sci_polychaos_buildsample.cpp
Compilation of sci_polychaos_getoutput.cpp
Compilation of sci_polychaos_getquantile.cpp
Compilation of sci_polychaos_getquantwilks.cpp
Compilation of sci_polychaos_getsample.cpp
Compilation of sci_polychaos_setgroupaddvar.cpp
Compilation of sci_polychaos_computeoutput.cpp
Compilation of sci_polychaos_setinput.cpp
Compilation of sci_polychaos_propagateinput.cpp
Compilation of sci_polychaos_getanova.cpp
Compilation of sci_polychaos_setanova.cpp
Compilation of sci_polychaos_getanovaord.cpp
Compilation of sci_polychaos_getanovaordco.cpp
Compilation of sci_polychaos_realisation.cpp
Compilation of sci_polychaos_save.cpp
Compilation of sci_polychaos_generatecode.cpp
Building shared library (be patient)
Generate a cleaner file
Generating loader_gateway.sce...
Building help...
Building the master document:
    C:\tbxnisp\help\en_US
Building the manual file [javaHelp] in
C:\tbxnisp\help\en_US.
(Please wait building ... this can take a while)
Generating loader.sce...

```

The following script presents the messages which are generated when the loader of the toolbox

is launched. The loader script performs the following steps:

- load the gateway (and the NISP library),
- load the help,
- load the demo.

```
-->exec C:\tbxnisp\loader.sce;  
Start NISP Toolbox  
    Load gateways  
    Load help  
    Load demos
```

It is now necessary to setup your Scilab system so that the toolbox is loaded automatically at startup. The way to do this is to configure the Scilab startup configuration file. The directory where this file is located is stored in the Scilab variable `SCIHOME`. In the following Scilab session, we use Scilab v5.2.0-beta-1 in order to know the value of the `SCIHOME` global variable.

```
-->SCIHOME  
SCIHOME =  
C:\Users\baudin\AppData\Roaming\Scilab\scilab-5.2.0-beta-1
```

On my Linux system, the Scilab 5.1 startup file is located in  
`/home/myname/.Scilab/scilab-5.1/.scilab`.

On my Windows system, the Scilab 5.1 startup file is located in

`C:/Users/myname/AppData/Roaming/Scilab/scilab-5.1/.scilab`.

This file is a regular Scilab script which is automatically loaded at Scilab's startup. If that file does not already exist, create it. Copy the following lines into the `.scilab` file and configure the path to the toolboxes, stored in the `SCILABTBX` variable.

```
exec("C:\tbxnisp\loader.sce");
```

The following script presents the messages which are generated when the unit tests script of the toolbox is launched.

```
-->exec C:\tbxnisp\runtests.sce;  
Tests beginning the 2009/11/18 at 12:47:45  
  TMPDIR = C:\Users\baudin\AppData\Local\Temp\SCI_TMP_6372_  
  001/004 - [tbxnisp] nisp.....passed : ref created  
  002/004 - [tbxnisp] polychaos1.....passed : ref created  
  003/004 - [tbxnisp] randvar1.....passed : ref created  
  004/004 - [tbxnisp] setrandvar1.....passed : ref created  
-----  
Summary  
tests           4 - 100 %  
passed          0 -  0 %  
failed          0 -  0 %  
skipped         0 -  0 %  
length          3.84 sec  
-----
```

Tests ending the 2009/11/18 at 12:47:48\end{verbatim}

# Chapter 4

## Configuration functions

In this section, we present functions which allow to configure the NISP toolbox.

The `nisp_*` functions allows to configure the global behaviour of the toolbox. These functions allows to startup and shutdown the toolbox and initialize the seed of the random number generator. They are presented in the figure 4.1.

<code>nisp_startup ()</code>	Starts up the NISP toolbox.
<code>nisp_shutdown ()</code>	Shuts down the NISP toolbox.
<code>level = nisp_verboselevelget ()</code>	Returns the current verbose level.
<code>nisp_verboselevelset ( level )</code>	Sets the value of the verbose level.
<code>nisp_initseed ( seed )</code>	Sets the seed of the uniform random number generator.
<code>nisp_destroyall</code>	Destroy all current objects.
<code>nisp_getpath</code>	Returns the path to the current module.
<code>nisp_printall</code>	Prints all current objects.

Figure 4.1: Outline of the configuration methods.

The user has no need to explicitly call the `nisp_startup ()` and `nisp_shutdown ()` functions. Indeed, these functions are called automatically by the `etc/NISP.start` and `etc/NISP.quit` scripts, located in the toolbox directory structure.

The `nisp_initseed ( seed )` is especially useful when we want to have reproducible results. It allows to set the seed of the generator at a particular value, so that the sequence of uniform pseudo-random numbers is deterministic. When the toolbox is started up, the seed is automatically set to 0, which allows to get the same results from session to session.



# Chapter 5

## The randvar class

In this section, we present the `randvar` class, which allows to define a random variable, and to generate random numbers from a given distribution function.

### 5.1 The distribution functions

In this section, we present the distribution functions provided by the `randvar` class. We especially present the Log-normal distribution function.

#### 5.1.1 Overview

The table 5.1 gives the list of distribution functions which are available with the `randvar` class [3].

Each distribution functions have zero, one or two parameters. One random variable can be specified by giving explicitly its parameters or by using default parameters. The parameters for all distribution function are presented in the figure 5.2, which also presents the conditions which must be satisfied by the parameters.

Name	$f(x)$	$E(X)$	$V(X)$
"Normale"	$\frac{1}{2\sigma\sqrt{2\pi}} \exp\left(-\frac{1}{2}\frac{(x-\mu)^2}{\sigma^2}\right)$	$\mu$	$\sigma^2$
"Uniforme"	$\begin{cases} \frac{1}{b-a}, & \text{if } x \in [a, b[ \\ 0 & \text{if } x \notin [a, b[ \end{cases}$	$\frac{b+a}{2}$	$\frac{(b-a)^2}{12}$
"Exponentielle"	$\begin{cases} \lambda \exp(-\lambda x), & \text{if } x > 0 \\ 0 & \text{if } x \leq 0 \end{cases}$	$\frac{1}{\lambda}$	$\frac{1}{\lambda^2}$
"LogNormale"	$\begin{cases} \frac{1}{\sigma x \sqrt{2\pi}} \exp\left(-\frac{1}{2}\frac{(\ln(x)-\mu)^2}{\sigma^2}\right), & \text{if } x > 0 \\ 0 & \text{if } x \leq 0 \end{cases}$	$\exp\left(\mu + \frac{1}{2}\sigma^2\right)$	$(\exp(\sigma^2) - 1) \exp(2\mu + \sigma^2)$
"LogUniforme"	$\begin{cases} \frac{1}{x \ln(b)-\ln(a)}, & \text{if } x \in [a, b[ \\ 0 & \text{if } x \notin [a, b[ \end{cases}$	$\frac{b-a}{\ln(b)-\ln(a)}$	$\frac{1}{2} \frac{b^2-a^2}{\ln(b)-\ln(a)} - E(x)$

Figure 5.1: Distributions functions of the `randvar` class. – The expected value is denoted by  $E(X)$  and the variance is denoted by  $V(X)$ .

Name	Parameter #1 : $a$	Parameter #2 : $b$	Conditions
"Normale"	$\mu = 0.$	$\sigma = 1.$	$\sigma > 0$
"Uniforme"	$a = 0.$	$b = 1.$	$a < b$
"Exponentielle"	$\lambda = 1.$	-	-
"LogNormale"	$\mu' = 0.1$	$\sigma = 1.0$	$\mu', \sigma > 0$
"LogUniforme"	$a = 0.1$	$b = 1.0$	$a, b > 0, a < b$

Figure 5.2: Default parameters for distributions functions.

### 5.1.2 Parameters of the Log-normal distribution

For the "LogNormale" law, the distribution function is usually defined by the expected value  $\mu$  and the standard deviation  $\sigma$  of the underlying Normal random variable. But, when we create a LogNormale `randvar`, the parameters to pass to the constructor are the expected value of the LogNormal random variable  $E(X)$  and the standard deviation of the underlying Normale random variable  $\sigma$ . The expected value and the variance of the Log Normal law are given by

$$E(X) = \exp\left(\mu + \frac{1}{2}\sigma^2\right) \quad (5.1)$$

$$V(X) = (\exp(\sigma^2) - 1) \exp(2\mu + \sigma^2). \quad (5.2)$$

It is possible to invert these formulas, in the situation where the given parameters are the expected value and the variance of the Log Normal random variable. We can invert completely the previous equations and get

$$\mu = \ln(E(X)) - \frac{1}{2} \ln\left(1 + \frac{V(X)}{E(X)^2}\right) \quad (5.3)$$

$$\sigma^2 = \ln\left(1 + \frac{V(X)}{E(X)^2}\right). \quad (5.4)$$

In particular, the expected value  $\mu$  of with the Normal random variable satisfies the equation

$$\mu = \ln(E(X)) - \sigma^2. \quad (5.5)$$

### 5.1.3 Uniform random number generation

In this section, we present the generation of uniform random numbers.

The goal of this section is to warn users about a current limitation of the library. Indeed, the random number generator is based on the compiler, so that its quality cannot be guaranteed.

The Uniforme law is associated with the parameters  $a, b \in \mathbb{R}$  with  $a < b$ . It produces real values uniform in the interval  $[a, b]$ .

To compute the uniform random number  $X$  in the interval  $[a, b]$ , a uniform random number in the interval  $[0, 1]$  is generated and then scaled with

$$X = a + (b - a)\overline{X}. \quad (5.6)$$

Let us now analyse how the uniform random number  $\overline{X} \in [0, 1]$  is computed. The uniform random generator is based on the C function `rand`, which returns an integer  $n$  in the interval

$[0, RAND\_MAX[$ . The value of the `RAND_MAX` variable is defined in the file `stdlib.h` and is compiler-dependent. For example, with the Visual Studio C++ 2008 compiler, the value is

$$RAND\_MAX = 2^{15} - 1 = 32767. \quad (5.7)$$

A uniform value  $\bar{X}$  in the range  $[0, 1[$  is computed from

$$\bar{X} = \frac{n}{N}, \quad (5.8)$$

where  $N = RAND\_MAX$  and  $n \in [0, RAND\_MAX[$ .

## 5.2 Methods

In this section, we give an overview of the methods which are available in the `randvar` class.

### 5.2.1 Overview

The figure 5.3 presents the methods available in the `randvar` class. The inline help contains the detailed calling sequence for each function and will not be repeated here.

<b>Constructors</b> <code>rv = randvar_new ( type [, options] )</code>
<b>Methods</b> <code>value = randvar_getvalue ( rv [, options] )</code> <code>randvar_getlog ( rv )</code>
<b>Destructor</b> <code>randvar_destroy ( rv )</code>
<b>Static methods</b> <code>rvlist = randvar_tokens ( )</code> <code>nbrv = randvar_size ( )</code>

Figure 5.3: Outline of the methods of the `randvar` class.

### 5.2.2 The Oriented-Object system

In this section, we present the token system which allows to emulate an oriented-object programming with Scilab. We also present the naming convention we used to create the names of the functions.

The `randvar` class provides the following functions.

- The constructor function `randvar_new` allows to create a new random variable and returns a *token* `rv`.
- The method `randvar_getvalue` takes the token `rv` as its first argument. In fact, all methods takes as their first argument the object on which they apply.

- The destructor `randvar_destroy` allows to delete the current object from the memory of the library.
- The static methods `randvar_tokens` and `randvar_size` allows to query the current object which are in use. More specifically, the `randvar_size` function returns the number of current `randvar` objects and the `randvar_tokens` returns the list of current `randvar` objects.

In the following Scilab sessions, we present these ideas with practical uses of the toolbox.

Assume that we start Scilab and that the toolbox is automatically loaded. At startup, there are no objects, so that the `randvar_size` function returns 0 and the `randvar_tokens` function returns an empty matrix.

```
-->nb = randvar_size()
nb =
    0.
-->tokenmatrix = randvar_tokens()
tokenmatrix =
    []
```

We now create 3 new random variables, based on the Uniform distribution function. We store the tokens in the variables `vu1`, `vu2` and `vu3`. These variables are regular Scilab double precision floating point numbers. Each value is a token which represents a random variable stored in the toolbox memory space.

```
-->vu1 = randvar_new("Uniforme")
vu1 =
    0.
-->vu2 = randvar_new("Uniforme")
vu2 =
    1.
-->vu3 = randvar_new("Uniforme")
vu3 =
    2.
```

There are now 3 objects in current use, as indicated by the following statements. The `tokenmatrix` is a row matrix containing regular double precision floating point numbers.

```
-->nb = randvar_size()
nb =
    3.
-->tokenmatrix = randvar_tokens()
tokenmatrix =
    0.    1.    2.
```

We assume that we have now made our job with the random variables, so that it is time to destroy the random variables. We call the `randvar_destroy` functions, which destroys the variables.

```
-->randvar_destroy(vu1);
-->randvar_destroy(vu2);
-->randvar_destroy(vu3);
```

We can finally check that there are no random variables left in the memory space.

```

-->nb = randvar_size()
nb =
    0.
-->tokenmatrix = randvar_tokens()
tokenmatrix =
    []

```

Scilab is a wonderful tool to experiment algorithms and make simulations. It happens sometimes that we are managing many variables at the same time and it may happen that, at some point, we are lost. The static methods provides tools to be able to recover from such a situation without closing our Scilab session.

In the following session, we create two random variables.

```

-->vu1 = randvar_new("Uniforme")
vu1 =
    3.
-->vu2 = randvar_new("Uniforme")
vu2 =
    4.

```

Assume now that we have lost the token associated with the variable `vu2`. We can easily simulate this situation, by using the `clear`, which destroys a variable from Scilab's memory space.

```

-->clear vu2
-->randvar_getvalue(vu2)
                !--error 4
Undefined variable: vu2

```

It is now impossible to generate values from the variable `vu2`. Moreover, it may be difficult to know exactly what went wrong and what exact variable is lost. At any time, we can use the `randvar_tokens` function in order to get the list of current variables. Deleting these variables allows to clean the memory space properly, without memory loss.

```

-->randvar_tokens()
ans =
    3.    4.
-->randvar_destroy(3)
ans =
    3.
-->randvar_destroy(4)
ans =
    4.
-->randvar_tokens()
ans =
    []

```

## 5.3 Examples

In this section, we present to examples of use of the `randvar` class. The first example presents the simulation of a Normal random variable and the generation of 1000 random variables. The second example presents the transformation of a Uniform outcome into a LogUniform outcome.

### 5.3.1 A sample session

We present a sample Scilab session, where the `randvar` class is used to generate samples from the Normale law.

In the following Scilab session, we create a Normale random variable and compute samples from this law. The `nisp_initseed` function is used to initialize the seed for the uniform random variable generator. Then we use the `randvar_new` function to create a new random variable from the Normale law with mean 1. and standard deviation 0.5. The main loop allows to compute 1000 samples from this law, based on calls to the `randvar_getvalue` function. Once the samples are computed, we use the Scilab function `mean` to check that the mean is close to 1 (which is the expected value of the Normale law, when the number of samples is infinite). Finally, we use the `randvar_destroy` function to destroy our random variable. Once done, we plot the empirical distribution function of this sample, with 50 classes.

```
nisp_initseed ( 0 );
mu = 1.0;
sigma = 0.5;
rv = randvar_new("Normale" , mu , sigma);
nbshots = 1000;
values = zeros(nbshots);
for i=1:nbshots
    values(i) = randvar_getvalue(rv);
end
mymean = mean (values);
mysigma = st_deviation(values);
myvariance = variance (values);
mprintf("Mean is : %f (expected = %f)\n", mymean, mu);
mprintf("Standard deviation is : %f (expected = %f)\n", mysigma, sigma);
mprintf("Variance is : %f (expected = %f)\n", myvariance, sigma^2);
randvar_destroy(rv);
histplot(50, values)
xtitle("Histogram of X", "X", "P(x)")
```

The previous script produces the following output.

```
Mean is : 0.988194 (expected = 1.000000)
Standard deviation is : 0.505186 (expected = 0.500000)
Variance is : 0.255213 (expected = 0.250000)
```

The previous script also produces the figure [5.4](#).

### 5.3.2 Variable transformations

In this section, we present the transformation of uniform random variables into other types of variables. The transformations which are available in the `randvar` class are presented in figure [5.5](#). We begin the analysis by a presentation of the theory required to perform transformations. Then we present some of the many the transformations which are provided by the library.

We now present some additionnal details for the function `randvar_getvalue ( rv , rv2 , value2 )`. This method allows to transform a random variable sample from one law to another. The statement

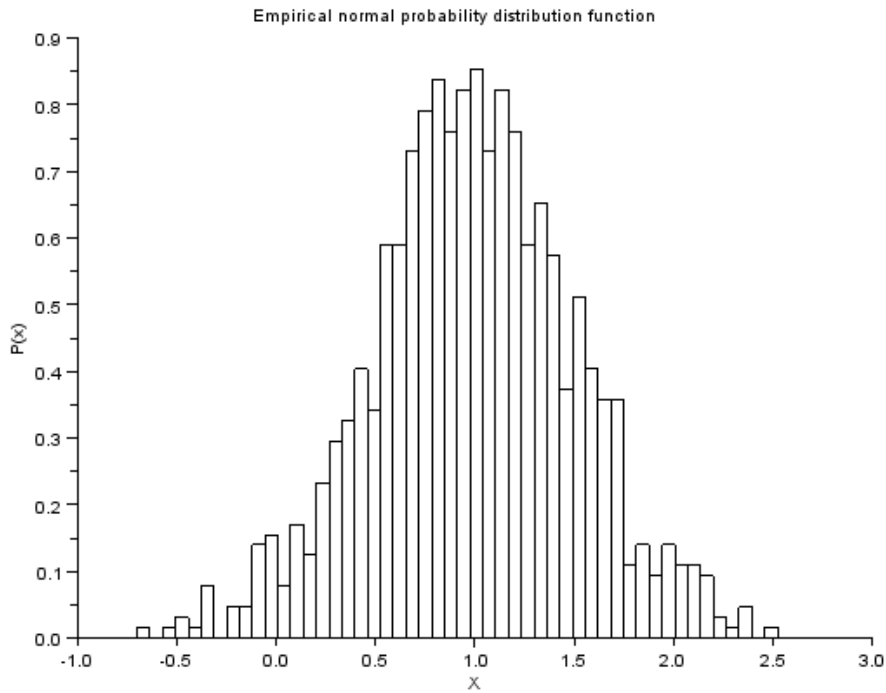


Figure 5.4: The histogram of a Normal random variable with 1000 samples.

Source	Target	Source	Target
Normale	Normale	LogNormale	Normale
	Uniforme		Uniforme
	Exponentielle		Exponentielle
	LogNormale		LogNormale
	LogUniforme		LogUniforme
Uniforme	Uniforme	LogUniforme	Uniforme
	Normale		Normale
	Exponentielle		Exponentielle
	LogNormale		LogNormale
	LogUniforme		LogUniforme
Exponentielle	Exponentielle		

Figure 5.5: Variable transformations available in the `randvar` class.

```
value = randvar_getvalue ( rv , rv2 , value2 )
```

returns a random value from the distribution function of the random variable `rv` by transformation of `value2` from the distribution function of random variable `rv2`.

In the following session, we transform a uniform random variable sample into a LogUniform variable sample. We begin to create a random variable `rv` from a LogUniform law and parameters  $a = 10, b = 20$ . Then we create a second random variable `rv2` from a Uniforme law and parameters  $a = 2, b = 3$ . The main loop is based on the transformation of a sample computed from `rv2` into a sample from `rv`. The `mean` allows to check that the transformed samples have an mean value which corresponds to the random variable `rv`.

```
nisp_initseed ( 0 );
a = 10.0;
b = 20.0;
rv = randvar_new ( "LogUniforme" , a , b );
rv2 = randvar_new ( "Uniforme" , 2 , 3 );
nbshots = 1000;
valuesLou = zeros(nbshots);
for i=1:nbshots
    valuesUni(i) = randvar_getvalue( rv2 );
    valuesLou(i) = randvar_getvalue( rv , rv2 , valuesUni(i) );
end
computed = mean ( valuesLou );
mu = (b-a)/(log(b)-log(a));
expected = mu;
mprintf("Expectation=%.5f (expected=%.5f)\n", computed , expected);
//
scf();
histplot(50, valuesUni);
xtitle("Empirical_histogram-Uniform_variable", "X", "P(X)");
scf();
histplot(50, valuesLou);
xtitle("Empirical_histogram-Log-Uniform_variable", "X", "P(X)");
randvar_destroy(rv);
randvar_destroy(rv2);
```

The previous script produces the following output.

```
Expectation=14.63075 (expected=14.42695)
```

The previous script also produces the figures [5.6](#) and [5.7](#).

The transformation depends on the *mother* random variable `rv1` and on the *daughter* random variable `rv`. Specific transformations are provided for all many combinations of the two distribution functions. These transformations will be analysed in the next sections.



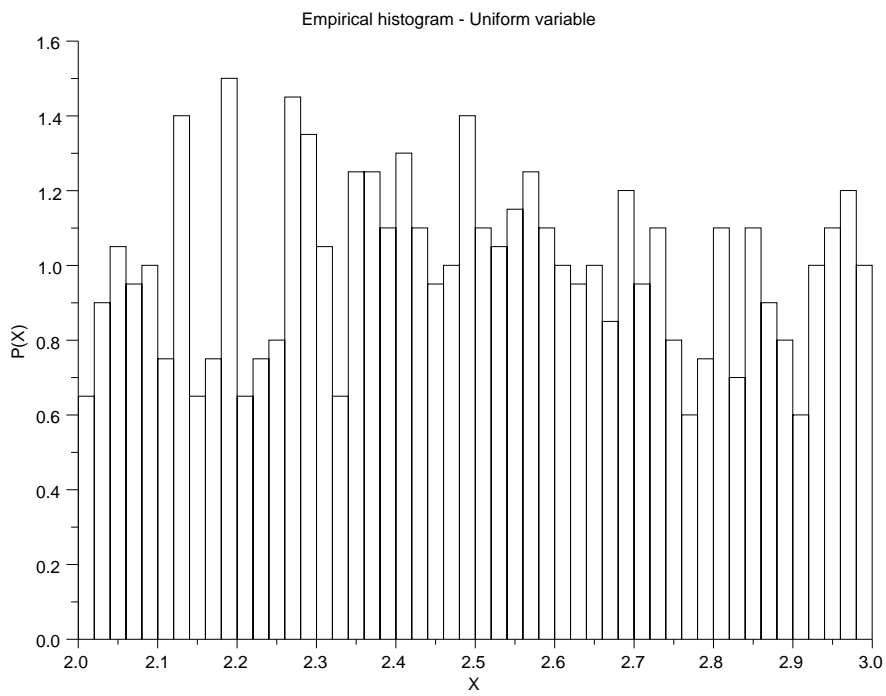


Figure 5.6: The histogram of a Uniform random variable with 1000 samples.

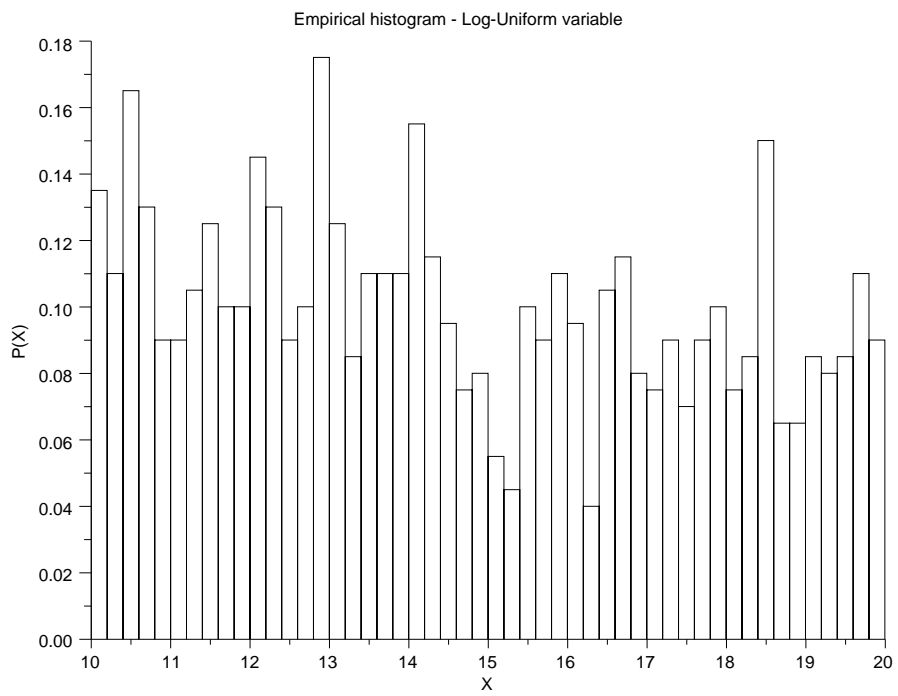


Figure 5.7: The histogram of a Log-Uniform random variable with 1000 samples.

# Chapter 6

## The `setrandvar` class

In this chapter, we present the `setrandvar` class. The first section gives a brief outline of the features of this class and the second section presents several examples.

### 6.1 Introduction

The `setrandvar` class allows to manage a collection of random variables and to build a Design Of Experiments (DOE). Several types of DOE are provided:

- Monte-Carlo,
- Latin Hypercube Sampling,
- Smolyak.

Once a DOE is created, we can retrieve the information experiment by experiment or the whole matrix of experiments. This last feature allows to benefit from the fact that Scilab can natively manage matrices, so that we do not have to perform loops to manage the complete DOE. Hence, good performances can be observed, even if the language still is interpreted.

The figure 6.1 presents the methods available in the `setrandvar` class. A complete description of the input and output arguments of each function is available in the inline help and will not be repeated here.

More informations about the Oriented Object system used in this toolbox can be found in the section 5.2.2.

### 6.2 Examples

In this section, we present examples of use of the `setrandvar` class. In the first example, we present a Scilab session where we create a Latin Hypercube Sampling. In the second part, we present various types of DOE which can be generated with this class.

#### 6.2.1 A Monte-Carlo design with 2 variables

In the following example, we build a Monte-Carlo design of experiments, with 2 input random variables. The first variable is associated with a Normal distribution function and the second

<p><b>Constructors</b></p> <pre> srv = setrandvar_new ( ) srv = setrandvar_new ( n ) srv = setrandvar_new ( file ) </pre>
<p><b>Methods</b></p> <pre> setrandvar_setsample ( srv , name , np ) setrandvar_setsample ( srv , k , i , value ) setrandvar_setsample ( srv , k , value ) setrandvar_setsample ( srv , value ) setrandvar_save ( srv , file ) np = setrandvar_getsize ( srv ) sample = setrandvar_getsample ( srv , k , i ) sample = setrandvar_getsample ( srv , k ) sample = setrandvar_getsample ( srv ) setrandvar_getlog ( srv ) nx = setrandvar_getdimension ( srv ) setrandvar_freememory ( srv ) setrandvar_buildsample ( srv , srv2 ) setrandvar_buildsample ( srv , name , np ) setrandvar_buildsample ( srv , name , np , ne ) setrandvar_addrandvar ( srv , rv ) </pre>
<p><b>Destructor</b></p> <pre> setrandvar_destroy ( srv ) </pre>
<p><b>Static methods</b></p> <pre> tokenmatrix = setrandvar_tokens ( ) nb = setrandvar_size ( ) </pre>

Figure 6.1: Outline of the methods of the `setrandvar` class

variable is associated with a Uniform distribution function. The simulation is based on 1000 experiments.

The function `nisp_initseed` is used to set the value of the seed to zero, so that the results can be reproduced. The `setrandvar_new` function is used to create a new set of random variables. Then we create two new random variables with the `randvar_new` function. These two variables are added to the set with the `setrandvar_addrandvar` function. The `setrandvar_buildsample` allows to build the design of experiments, which can be retrieved as matrix with the `setrandvar_getsample` function. The sampling matrix has `np` rows and 2 columns (one for each input variable).

```
nisp_initseed(0);
rvu1 = randvar_new("Normale",1,3);
rvu2 = randvar_new("Uniforme",2,3);
//
srvu = setrandvar_new();
setrandvar_addrandvar ( srvu, rvu1);
setrandvar_addrandvar ( srvu, rvu2);
//
np = 5000;
setrandvar_buildsample(srvu, "MonteCarlo",np);
sampling = setrandvar_getsample(srvu);
// Check sampling of random variable #1
mean(sampling(:,1)) // Expectation : 1
// Check sampling of random variable #2
mean(sampling(:,2)) // Expectation : 2.5
//
scf();
histplot(50,sampling(:,1));
xtitle("Empirical_histogram_of_X1");
scf();
histplot(50,sampling(:,2));
xtitle("Empirical_histogram_of_X2");
//
// Clean-up
setrandvar_destroy(srvu);
randvar_destroy(rvu1);
randvar_destroy(rvu2);
```

The previous script produces the following output.

```
-->mean(sampling(:,1)) // Expectation : 1
ans =
    1.0064346
-->mean(sampling(:,2)) // Expectation : 2.5
ans =
    2.5030984
```

The previous script also produces the figures [6.2](#) and [6.3](#).

We may now want to add the exact distribution to these histograms and compare. The Normal distribution function is not provided by Scilab, but is provided by the Stixbox module. Indeed, the `dnorm` function of the Stixbox module computes the Normal probability distribution function.

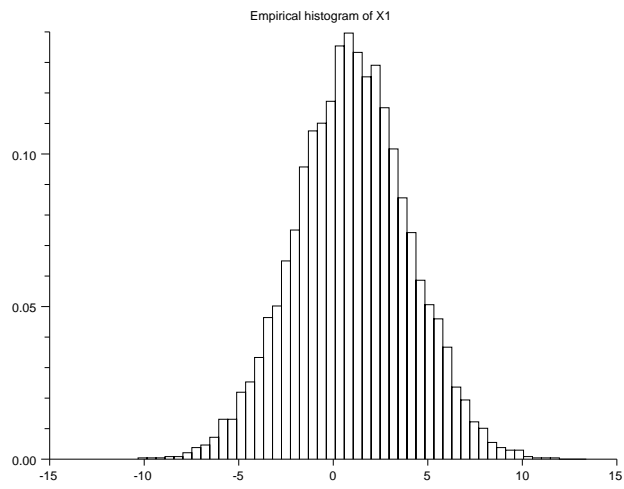


Figure 6.2: Monte-Carlo Sampling - Normal random variable.

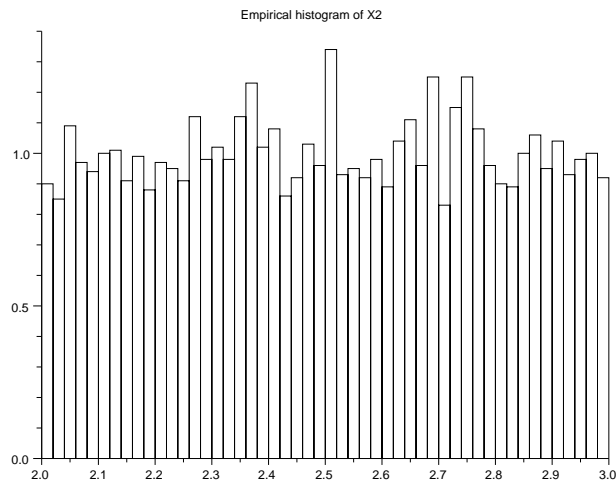


Figure 6.3: Monte-Carlo Sampling - Uniform random variable.

In order to install this module, we can run the `atomsInstall` function, as in the following script.

```
atomsInstall("stibox")
```

The following script compares the empirical and theoretical distributions.

```
scf();  
histplot(50,sampling(:,1));  
xtitle("Empirical histogram of X1");  
x=linspace(-15,15,1000);  
y = dnorm(x,1,3);  
plot(x,y,"r-")  
legend(["Empirical","Exact"]);
```

The previous script produces the figure 6.4.

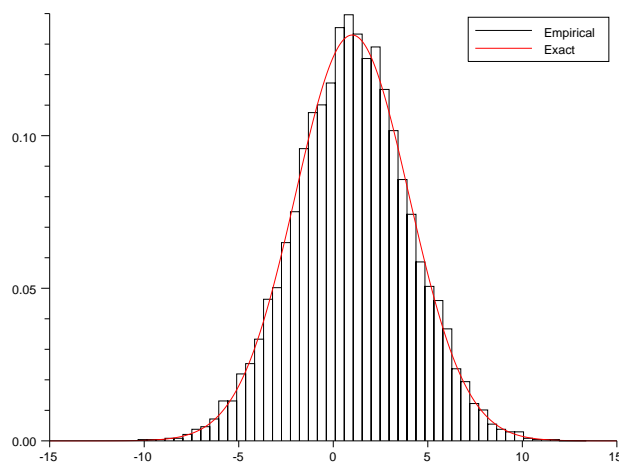


Figure 6.4: Monte-Carlo Sampling - Histogram and exact distribution functions for the first variable.

The following script performs the same comparison for the second variable.

```
scf();  
histplot(50,sampling(:,2));  
xtitle("Empirical histogram of X2");  
x=linspace(2,3,1000);  
y=ones(1000,1);  
plot(x,y,"r-");
```

The previous script produces the figure 6.5.

## 6.2.2 A Monte-Carlo design with 2 variables

In this section, we create a Monte-Carlo design with 2 variables.

We are going to use the exponential distribution function, which is not defined in Scilab. The following `exppdf` function computes the probability distribution function of the exponential distribution function.



Figure 6.5: Monte-Carlo Sampling - Histogram and exact distribution functions for the second variable.

```
function p = exppdf ( x , lambda )
    p = lambda.*exp(-lambda.*x)
endfunction
```

The following script creates a Monte-Carlo sampling where the first variable is Normal and the second variable is Exponential. Then we compare the empirical histogram and the exact distribution function. We use the `dnorm` function defined in the Stibox module.

```
nisp_initseed ( 0 );
rv1 = randvar_new("Normale",1.0,0.5);
rv2 = randvar_new("Exponentielle",5.);
// Definition d'un groupe de variables aleatoires
srv = setrandvar_new ( );
setrandvar_addrandvar ( srv , rv1 );
setrandvar_addrandvar ( srv , rv2 );
np = 1000;
setrandvar_buildsample ( srv , "MonteCarlo" , np );
//
sampling = setrandvar_getsample ( srv );
// Check sampling of random variable #1
mean(sampling(:,1)), variance(sampling(:,1))
// Check sampling of random variable #2
min(sampling(:,2)), max(sampling(:,2))
// Plot
scf();
histplot(40, sampling(:,1))
x = linspace(-1,3,1000)';
p = dnorm(x,1,0.5);
plot(x,p,"r-")
```



```

xtitle("Empirical histogram of X1", "X", "P(X)");
legend(["Empirical", "Exact"]);
scf();
histplot(40, sampling(:,2))
x = linspace(0,2,1000)';
p = exppdf ( x , 5 );
plot(x,p,"r-")
xtitle("Empirical histogram of X2", "X", "P(X)");
legend(["Empirical", "Exact"]);
// Clean-up
setrandvar_destroy(srv);
randvar_destroy(rv1);
randvar_destroy(rv2);

```

The previous script produces the figures 6.6 and 6.7.

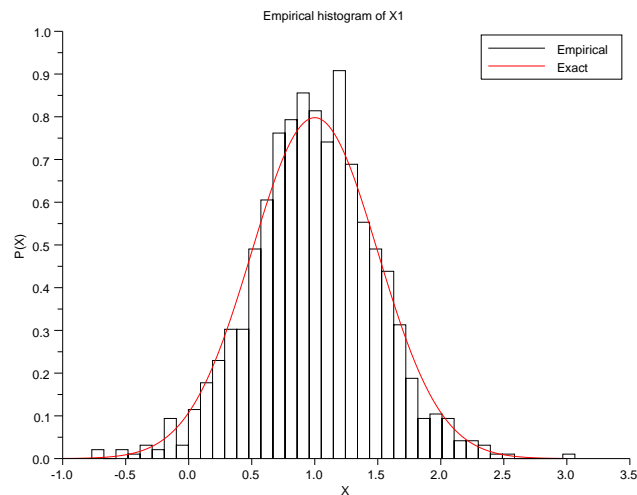


Figure 6.6: Monte-Carlo Sampling - Histogram and exact distribution functions for the first variable.

### 6.2.3 A LHS design

In this section, we present the creation of a Latin Hypercube Sampling. In our example, the DOE is based on two random variables, the first being Normal with mean 1.0 and standard deviation 0.5 and the second being Uniform in the interval [2, 3].

We begin by defining two random variables with the `randvar_new` function.

```

vu1 = randvar_new("Normale", 1.0, 0.5);
vu2 = randvar_new("Uniforme", 2.0, 3.0);

```

Then, we create a collection of random variables with the `setrandvar_new` function which creates here an empty collection of random variables. Then we add the two random variables to the collection.

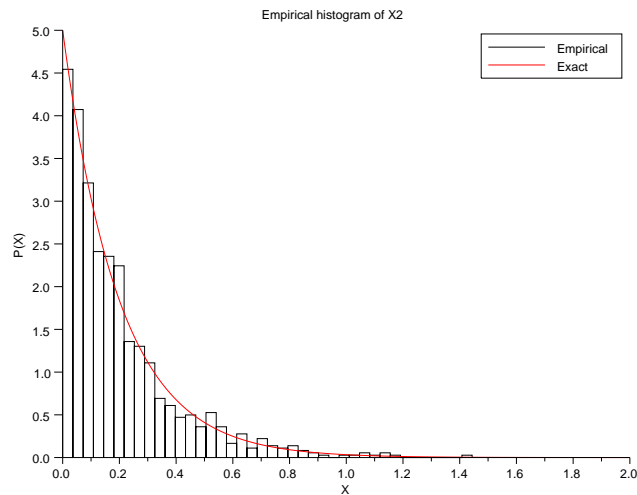


Figure 6.7: Monte-Carlo Sampling - Histogram and exact distribution functions for the second variable.

```

srv = setrandvar_new ( );
setrandvar_addrandvar ( srv , vu1 );
setrandvar_addrandvar ( srv , vu2 );

```

We can now build the DOE so that it is a LHS sampling with 1000 experiments.

```

setrandvar_buildsample ( srv , "Lhs" , 1000 );

```

At this point, the DOE is stored in the memory space of the NISP library, but we do not have a direct access to it. We now call the `setrandvar_getsample` function and store that DOE into the `sampling` matrix.

```

sampling = setrandvar_getsample ( srv );

```

The `sampling` matrix has 1000 rows, corresponding to each experiment, and 2 columns, corresponding to each input random variable.

The following script allows to plot the sampling, which is presented in figure 6.8.

```

my_handle = scf();
clf(my_handle,"reset");
plot(sampling(:,1),sampling(:,2));
my_handle.children.children.children.line_mode = "off";
my_handle.children.children.children.mark_mode = "on";
my_handle.children.children.children.mark_size = 2;
my_handle.children.title.text = "Latin_Hypercube_Sampling";
my_handle.children.x_label.text = "Variable_#1:_Normale,1.0,0.5";
my_handle.children.y_label.text = "Variable_#2:_Uniforme,2.0,3.0";

```

The following script allows to plot the histogram of the two variables, which are presented in figures 6.9 and 6.10.

```

// Plot Var #1

```

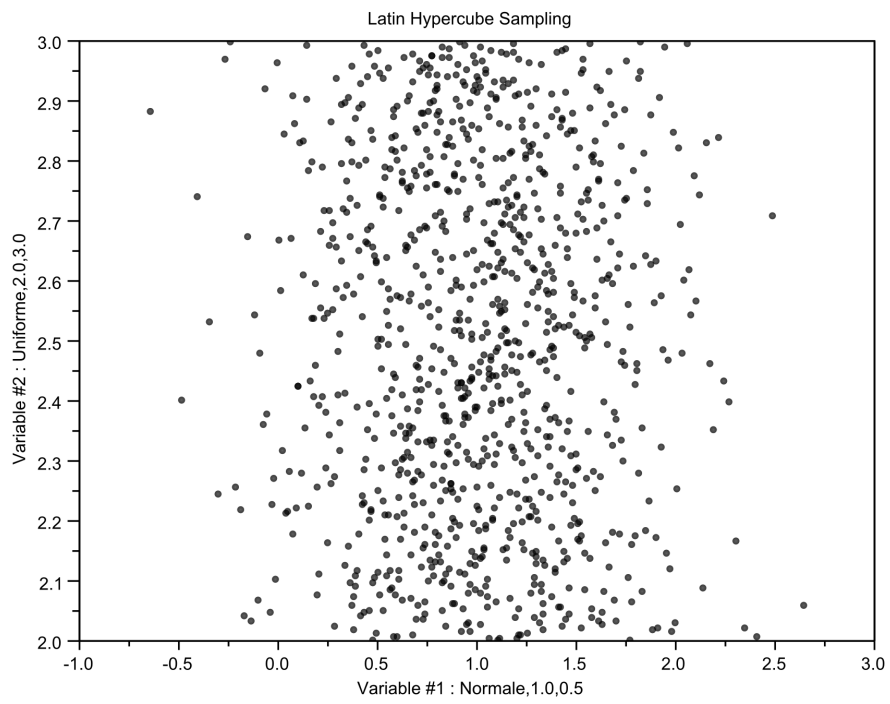


Figure 6.8: Latin Hypercube Sampling - The first variable is Normal, the second variable is Uniform.

```

my_handle = scf();
clf(my_handle,"reset");
histplot ( 50 , sampling(:,1))
my_handle.children.title.text = "Variable_#1:_Normale,1.0,0.5";
// Plot Var #2
my_handle = scf();
clf(my_handle,"reset");
histplot ( 50 , sampling(:,2))
my_handle.children.title.text = "Variable_#2:_Uniforme,2.0,3.0";

```

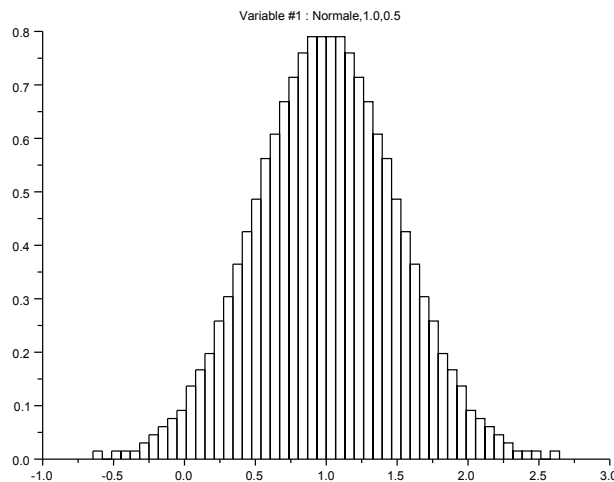


Figure 6.9: Latin Hypercube Sampling - Normal random variable.

We can use the `mean` and `variance` on each random variable and check that the expected result is computed. We insist on the fact that the `mean` and `variance` functions are not provided by the NISP library: these are pre-defined functions which are available in the Scilab library. That means that any Scilab function can be now used to process the data generated by the toolbox.

```

for ivar = 1:2
    m = mean(sampling(:,ivar))
    mprintf("Variable_#%d,_Mean_:_%f\n",ivar,m)
    v = variance(sampling(:,ivar))
    mprintf("Variable_#%d,_Variance_:_%f\n",ivar,v)
end

```

The previous script produces the following output.

```

Variable #1, Mean : 1.000000
Variable #1, Variance : 0.249925
Variable #2, Mean : 2.500000
Variable #2, Variance : 0.083417

```

Our numerical simulation is now finished, but we must destroy the objects so that the memory managed by the toolbox is deleted.

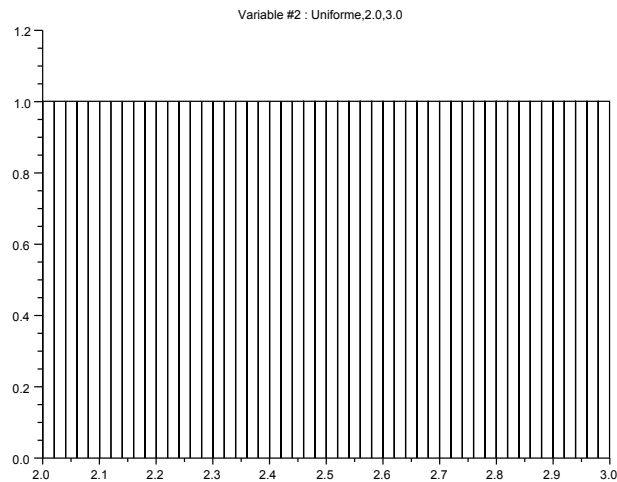


Figure 6.10: Latin Hypercube Sampling - Uniform random variable (the LHS sampling is not a Monte-Carlo sampling).

```

randvar_destroy(vu1)
randvar_destroy(vu2)
setrandvar_destroy(srv)

```

## 6.2.4 Other types of DOEs

The following Scilab session allows to generate a Monte-Carlo sampling with two uniform variables in the interval  $[-1, 1]$ . The figure 6.11 presents this sampling and the figures 6.12 and 6.13 present the histograms of the two uniform random variables.

```

vu1 = randvar_new("Uniforme",-1.0,1.0);
vu2 = randvar_new("Uniforme",-1.0,1.0);
srv = setrandvar_new ( );
setrandvar_addrandvar ( srv , vu1 );
setrandvar_addrandvar ( srv , vu2 );
setrandvar_buildsample ( srv , "MonteCarlo" , 1000 );
sampling = setrandvar_getsample ( srv );
randvar_destroy(vu1);
randvar_destroy(vu2);
setrandvar_destroy(srv);

```

It is easy to change the type of sampling by modifying the second argument of the `setrandvar_buildsample` function. This way, we can create the Petras, Quadrature and Sobol sampling presented in figures 6.14, 6.15 and 6.16.

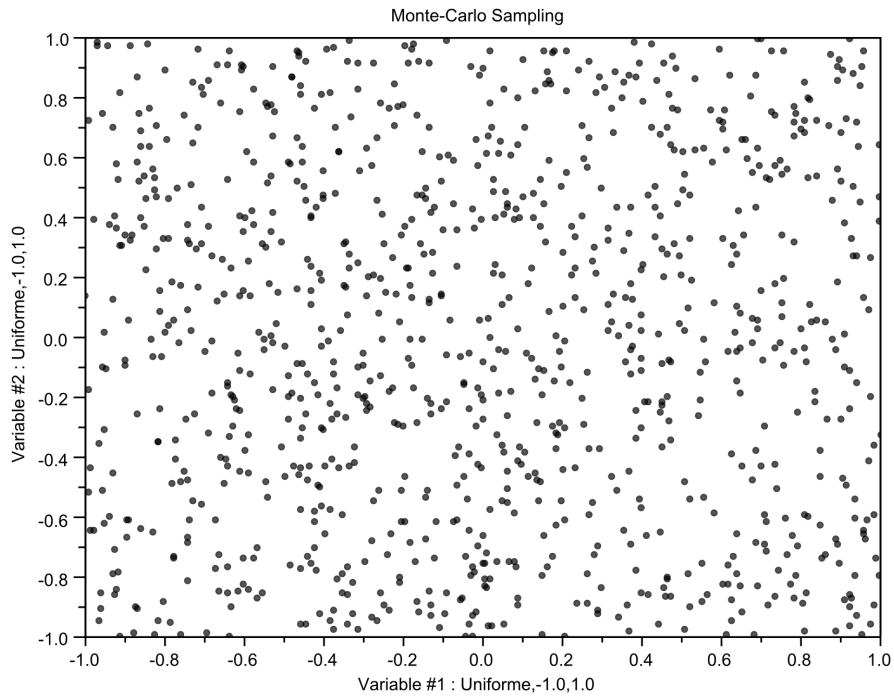


Figure 6.11: Monte-Carlo Sampling - Two uniform variables in the interval  $[-1, 1]$ .

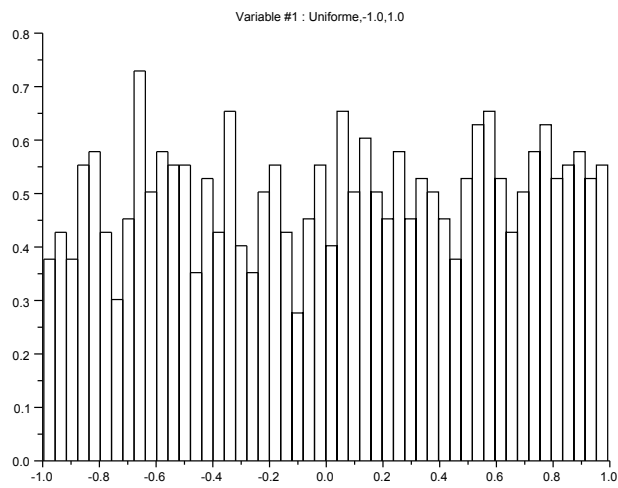


Figure 6.12: Latin Hypercube Sampling - First uniform variable in  $[-1, 1]$ .

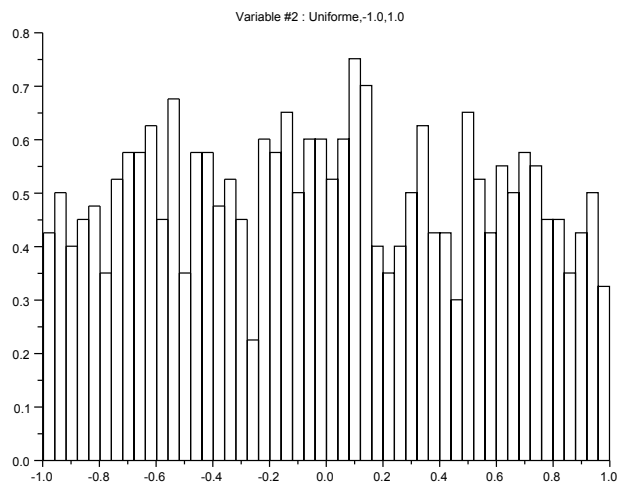


Figure 6.13: Latin Hypercube Sampling - Second uniform variable in  $[-1, 1]$ .

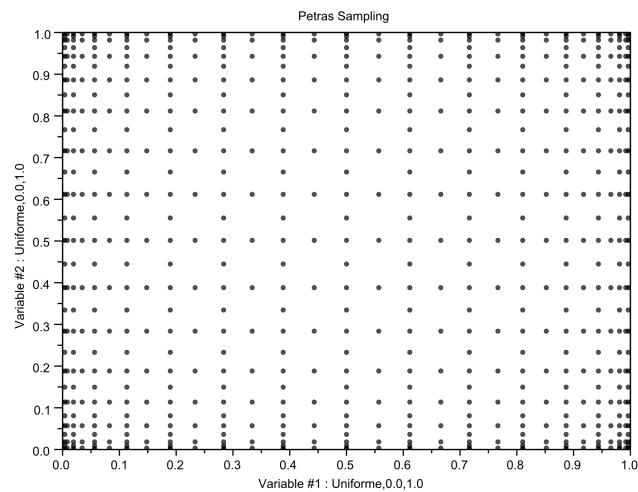


Figure 6.14: Petras sampling - Two uniform variables in the interval  $[-1, 1]$ .

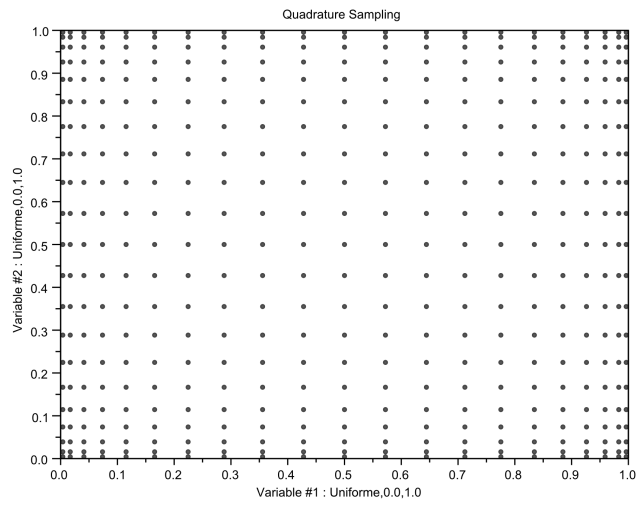


Figure 6.15: Quadrature sampling - Two uniform variables in the interval  $[-1, 1]$ .

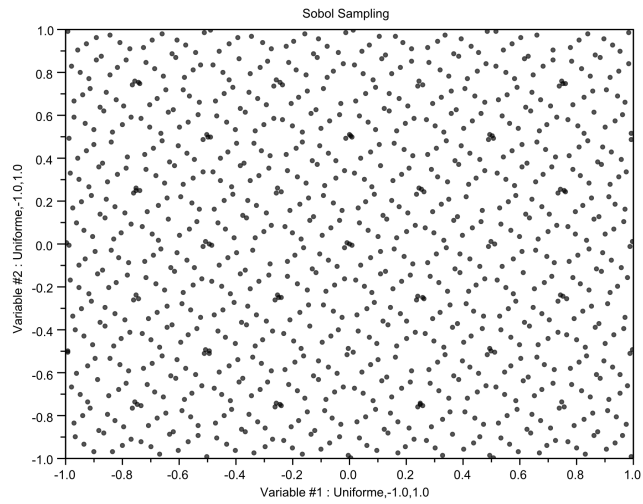


Figure 6.16: Sobol sampling - Two uniform variables in the interval  $[-1, 1]$ .



# Chapter 7

## The polychaos class

### 7.1 Introduction

The `polychaos` class allows to manage a polynomial chaos expansion. The coefficients of the expansion are computed based on given numerical experiments which creates the association between the inputs and the outputs. Once computed, the expansion can be used as a regular function. The mean, standard deviation or quantile can also be directly retrieved.

The tool allows to get the following results:

- mean,
- variance,
- quantile,
- correlation, etc...

Moreover, we can generate the C source code which computes the output of the polynomial chaos expansion. This C source code is stand-alone, that is, it is independent of both the NISP library and Scilab. It can be used as a meta-model.

The figure [7.1](#) presents the most commonly used methods available in the `polychaos` class. More methods are presented in figure [7.2](#). The inline help contains the detailed calling sequence for each function and will not be repeated here. More than 50 methods are available and most of them will not be presented here.

More informations about the Oriented Object system used in this toolbox can be found in the section [5.2.2](#).

### 7.2 Examples

In this section, we present to examples of use of the `polychaos` class.

#### 7.2.1 Product of two random variables

In this section, we present the polynomial expansion of the product of two random variables. We analyse the Scilab script and present the methods which are available to perform the sensi-

<b>Constructors</b>
pc = polychaos_new ( file ) pc = polychaos_new ( srv , ny ) pc = polychaos_new ( pc , nopt , varopt )
<b>Methods</b>
polychaos_setsizetarget ( pc , np ) polychaos_settarget ( pc , output ) polychaos_setinput ( pc , invalue ) polychaos_setdimoutput ( pc , ny ) polychaos_setdegree ( pc , no ) polychaos_getvariance ( pc ) polychaos_getmean ( pc )
<b>Destructor</b>
polychaos_destroy (pc)
<b>Static methods</b>
tokenmatrix = polychaos_tokens () nb = polychaos_size ()

Figure 7.1: Outline of the methods of the polychaos class

tivity analysis. This script is based on the NISP methodology, which has been presented in the Introduction chapter. We will use the figure 1.1 as a framework and will follow the steps in order.

In the following Scilab script, we define the function `Exemple` which takes a vector of size 2 as input and returns a scalar as output.

```
function y = Exemple (x)
    y(:,1) = x(:,1) .* x(:,2)
endfunction
```

We now create a collection of two stochastic (normalized) random variables. Since the random variables are normalized, we use the default parameters of the `randvar_new` function. The normalized collection is stored in the variable `srvx`.

```
vx1 = randvar_new("Normale");
vx2 = randvar_new("Uniforme");
srvx = setrandvar_new();
setrandvar_addrandvar ( srvx , vx1 );
setrandvar_addrandvar ( srvx , vx2 );
```

We create a collection of two uncertain parameters. We explicitly set the parameters of each random variable, that is, the first Normal variable is associated with a mean equal to 1.0 and a standard deviation equal to 0.5, while the second Uniform variable is in the interval [1.0, 2.5]. This collection is stored in the variable `srvu`.

```
vu1 = randvar_new("Normale",1.0,0.5);
vu2 = randvar_new("Uniforme",1.0,2.5);
srvu = setrandvar_new();
setrandvar_addrandvar ( srvu , vu1 );
setrandvar_addrandvar ( srvu , vu2 );
```

### Methods

```
output = polychaos_gettarget ( pc )
np = polychaos_getsizetarget ( pc )
polychaos_getsample ( pc , k , ovar )
polychaos_getquantile ( pc , k )
polychaos_getsample ( pc )
polychaos_getquantile ( pc , alpha )
polychaos_getoutput ( pc )
polychaos_getmultind ( pc )
polychaos_getlog ( pc )
polychaos_getinvquantile ( pc , threshold )
polychaos_getindextotal ( pc )
polychaos_getindexfirst ( pc )
ny = polychaos_getdimoutput ( pc )
nx = polychaos_getdiminput ( pc )
p = polychaos_getdimexp ( pc )
no = polychaos_getdegree ( pc )
polychaos_getcovariance ( pc )
polychaos_getcorrelation ( pc )
polychaos_getanova ( pc )
polychaos_generatecode ( pc , filename , funname )
polychaos_computeoutput ( pc )
polychaos_computeexp ( pc , srv , method )
polychaos_computeexp ( pc , pc2 , invalue , varopt )
polychaos_buildsample ( pc , type , np , order )
```

Figure 7.2: More methods from the polychaos class

The first design of experiment is build on the stochastic set `srvx` and based on a Quadrature type of DOE. Then this DOE is transformed into a DOE for the uncertain collection of parameters `srvu`.

```

degre = 2;
setrandvar_buildsample ( srvx , "Quadrature" , degre );
setrandvar_buildsample ( srvu , srvx );

```

The next steps will be to create the polynomial and actually perform the DOE. But before doing this, we can take a look at the DOE associated with the stochastic and uncertain collection of random variables. We can use the `setrandvar_getsample` twice and get the following output.

```

-->setrandvar_getsample(srvx)
ans =

- 1.7320508    0.1127017
- 1.7320508    0.5
- 1.7320508    0.8872983
  0.          0.1127017
  0.          0.5
  0.          0.8872983
  1.7320508    0.1127017
  1.7320508    0.5
  1.7320508    0.8872983
-->setrandvar_getsample(srvu)
ans =

  0.1339746    1.1690525
  0.1339746    1.75
  0.1339746    2.3309475
  1.          1.1690525
  1.          1.75
  1.          2.3309475
  1.8660254    1.1690525
  1.8660254    1.75
  1.8660254    2.3309475

```

These two matrices are a  $9 \times 2$  matrices, where each line represents an experiment and each column represents an input random variable. The stochastic (normalized) `srvx` DOE has been created first, then the `srvu` has been deduced from `srvx` based on random variable transformations.

We now use the `polychaos_new` function and create a new polynomial `pc`. The number of input variables corresponds to the number of variables in the stochastic collection `srvx`, that is 2, and the number of output variables is given as the input argument `ny`. In this particular case, the number of experiments to perform is equal to `np=9`, as returned by the `setrandvar_getsize` function. This parameter is passed to the polynomial `pc` with the `polychaos_setsizetarget` function.

```

ny = 1;
pc = polychaos_new ( srvx , ny );
np = setrandvar_getsize(srvx);
polychaos_setsizetarget(pc,np);

```

In the next step, we perform the simulations prescribed by the DOE. We perform this loop in the Scilab language and make a loop over the index `k`, which represents the index of the current experiment, while `np` is the total number of experiments to perform. For each loop, we get the input from the uncertain collection `srvu` with the `setrandvar_getsample` function, pass it to the `Exemple` function, get back the output which is then transferred to the polynomial `pc` by the `polychaos_settarget` function.

```
for k=1:np
    inputdata = setrandvar_getsample(srvu,k);
    outputdata = Exemple(inputdata);
    mprintf ( "Experiment_#%d, input_=[%f_%f], output_=%f\n", k, ..
        inputdata(1), inputdata(2) , outputdata )
    polychaos_settarget(pc,k,outputdata);
end
```

The previous script produces the following output.

```
Experiment #1, input =[0.133975 1.169052], output = 0.156623
Experiment #2, input =[0.133975 1.750000], output = 0.234456
Experiment #3, input =[0.133975 2.330948], output = 0.312288
Experiment #4, input =[1.000000 1.169052], output = 1.169052
Experiment #5, input =[1.000000 1.750000], output = 1.750000
Experiment #6, input =[1.000000 2.330948], output = 2.330948
Experiment #7, input =[1.866025 1.169052], output = 2.181482
Experiment #8, input =[1.866025 1.750000], output = 3.265544
Experiment #9, input =[1.866025 2.330948], output = 4.349607
```

There is actually a much faster way of computing the output. Indeed, using vectorisation, we can compute all the outputs in one single call to the `Exemple` function.

```
inputdata = setrandvar_getsample(srvu);
outputdata = Exemple(inputdata);
polychaos_settarget(pc,outputdata);
```

We can compute the polynomial expansion based on numerical integration so that the coefficients of the polynomial are determined. This is done with the `polychaos_computeexp` function, which stands for "compute the expansion".

```
polychaos_setdegree(pc,degree);
polychaos_computeexp(pc,svrx,"Integration");
```

Everything is now ready for the sensitivity analysis. Indeed, the `polychaos_getmean` returns the mean while the `polychaos_getvariance` returns the variance.

```
average = polychaos_getmean(pc);
var = polychaos_getvariance(pc);
mprintf("Mean_=====%f\n",average);
mprintf("Variance_=====%f\n",var);
mprintf("Indice_de_sensibilite_du_1er_ordre\n");
mprintf("====Variable_X1_=%f\n",polychaos_getindexfirst(pc,1));
mprintf("====Variable_X2_=%f\n",polychaos_getindexfirst(pc,2));
mprintf("Indice_de_sensibilite_Totale\n");
mprintf("====Variable_X1_=%f\n",polychaos_getindextotal(pc,1));
mprintf("====Variable_X2_=%f\n",polychaos_getindextotal(pc,2));
```

The previous script produces the following output.

```
Mean      = 1.750000
Variance  = 1.000000
Indice de sensibilite du 1er ordre
  Variable X1 = 0.765625
  Variable X2 = 0.187500
Indice de sensibilite Totale
  Variable X1 = 0.812500
  Variable X2 = 0.234375
```

In order to free the memory required for the computation, it is necessary to delete all the objects created so far.

```
polychaos_destroy(pc);
randvar_destroy(vu1);
randvar_destroy(vu2);
randvar_destroy(vx1);
randvar_destroy(vx2);
setrandvar_destroy(srvu);
setrandvar_destroy(srvx);
```

Prior to destroying the objects, we can inquire a little more about the density of the output of the chaos polynomial. In the following script, we create a Latin Hypercube Sampling made of 10 000 points. Then get the output of the polynomial on these inputs and plot the histogram of the output.

```
polychaos_buildsample(pc,"Lhs",10000,0);
sample_output = polychaos_getsample(pc);
scf();
histplot(50,sample_output);
xtitle("Product_function_Empirical_Histogram","X","P(X)");
```

The previous script produces the figure [7.3](#).

We may explore the following topics.

- Perform the same analysis where the variable  $X_2$  is a normal variable with mean 2 and standard deviation 2.
- Check that the development in polynomial chaos on a Hermite-Hermite basis does not allow to get exact results. See that the convergence can be obtained by increasing the degree.
- Check that the development on a basis Hermite-Legendre allows to get exact results with degree 2.

## 7.2.2 The Ishigami test case

In this section, we present the Ishigami test case.

The function `Exemple` is the model that we consider in this numerical experiment. This function takes a vector of size 3 in input and returns a scalar output.

```
function y = Exemple (x)
  a=7.;
```

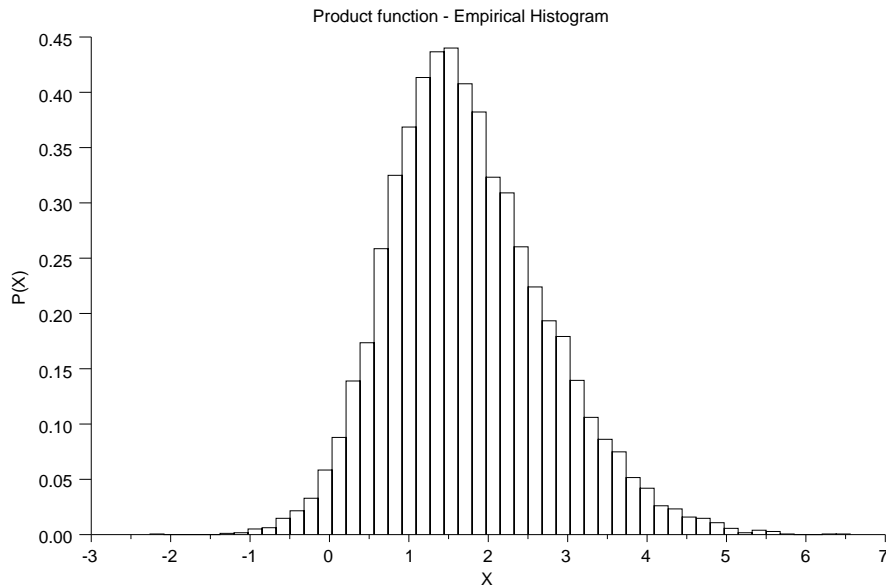


Figure 7.3: Product function - Histogram of the output on a LHS design with 10000 experiments.

```

b=0.1;
s1=sin(x(:,1));
s2=sin(x(:,2));
y(:,1) = s1 + a.*s2.*s2 + b.*x(:,3).*x(:,3).*x(:,3).*x(:,3).*s1;
endfunction

```

We create 3 uncertain parameters which are uniform in the interval  $[-\pi, \pi]$  and put these random variables into the collection `srvu`.

```

rvu1 = randvar_new("Uniform", -%pi, %pi);
rvu2 = randvar_new("Uniform", -%pi, %pi);
rvu3 = randvar_new("Uniform", -%pi, %pi);

srvu = setrandvar_new();
setrandvar_addrandvar ( srvu, rvu1);
setrandvar_addrandvar ( srvu, rvu2);
setrandvar_addrandvar ( srvu, rvu3);

```

The collection of stochastic variables is created with the function `setrandvar_new`. The calling sequence `srvx = setrandvar_new( nx )` allows to create a collection of `nx=3` random variables uniform in the interval  $[0, 1]$ . Then we create a Petras DOE for the stochastic collection `srvx` and transform it into a DOE for the uncertain parameters `srvu`.

```

nx = setrandvar_getdimension ( srvu );
srvx = setrandvar_new( nx );
degre = 9;
setrandvar_buildsample(srvx, "Petras", degre);
setrandvar_buildsample( srvu , srvx );

```

We use the `polychaos_new` function and create the new polynomial `pc` with 3 inputs and 1 output.

```
noutput = 1;
pc = polychaos_new ( srvx , noutput );
```

The next step allows to perform the simulations associated with the DOE prescribed by the collection `srvu`. Here, we must perform `np=751` experiments.

```
np = setrandvar_getsize(srvu);
polychaos_setsizetarget(pc,np);
// This is slow
for k=1:np
    inputdata = setrandvar_getsample(srvu,k);
    outputdata = Exemple(inputdata);
    polychaos_settarget(pc,k,outputdata);
end
```

The previous loop works, but is slow when `np` is large. Instead, the following script is fast because it uses vectorization.

```
inputdata = setrandvar_getsample(srvu);
outputdata = Exemple(inputdata);
polychaos_settarget(pc,outputdata);
```

We can now compute the polynomial expansion by integration.

```
polychaos_setdegree(pc,degree);
polychaos_computexp(pc,srvx,"Integration");
```

Everything is now ready so that we can do the sensitivity analysis, as in the following script.

```
average = polychaos_getmean(pc);
var = polychaos_getvariance(pc);
mprintf("Mean_====_=%f\n",average);
mprintf("Variance_====_=%f\n",var);
mprintf("First_order_sensitivity_index\n");
mprintf("====Variable_X1_=%f\n",polychaos_getindexfirst(pc,1));
mprintf("====Variable_X2_=%f\n",polychaos_getindexfirst(pc,2));
mprintf("====Variable_X3_=%f\n",polychaos_getindexfirst(pc,3));
mprintf("Total_sensitivity_index\n");
mprintf("====Variable_X1_=%f\n",polychaos_getindextotal(pc,1));
mprintf("====Variable_X2_=%f\n",polychaos_getindextotal(pc,2));
mprintf("====Variable_X3_=%f\n",polychaos_getindextotal(pc,3));
```

The previous script produces the following output.

```
Mean          = 3.500000
Variance      = 13.842473
First order sensitivity index
  Variable X1 = 0.313953
  Variable X2 = 0.442325
  Variable X3 = 0.000000
Total sensitivity index
  Variable X1 = 0.557675
```



```
Variable X2 = 0.442326
Variable X3 = 0.243721
```

We now focus on the variance generated by the variables #1 and #3. We set the group to the empty group with the `polychaos_setgroupempty` function and add variables with the `polychaos_setgroupaddvar` function.

```
groupe = [1 3];
polychaos_setgroupempty ( pc );
polychaos_setgroupaddvar ( pc , groupe(1) );
polychaos_setgroupaddvar ( pc , groupe(2) );
mprintf("Fraction of the variance of a group of variables\n");
mprintf("Groupe X1 et X2 = %f\n", polychaos_getgroupind(pc));
```

The previous script produces the following output.

```
Fraction of the variance of a group of variables
Groupe X1 et X2 = 0.557674
```

The function `polychaos_getanova` prints the functional decomposition of the normalized variance.

```
polychaos_getanova(pc);
```

The previous script produces the following output.

```
1 0 0 : 0.313953
0 1 0 : 0.442325
1 1 0 : 1.55229e-009
0 0 1 : 8.08643e-031
1 0 1 : 0.243721
0 1 1 : 7.26213e-031
1 1 1 : 1.6007e-007
```

We can compute the density function associated with the output variable of the function. In order to compute it, we use the `polychaos_buildsample` function and create a Latin Hypercube Sampling with 10000 experiments. The `polychaos_getsample` function allows to query the polynomial and get the outputs. We plot it with the `histplot` Scilab graphic function, which produces the figure 7.4.

```
polychaos_buildsample(pc, "Lhs", 10000, 0);
sample_output = polychaos_getsample(pc);
scf();
histplot(50, sample_output)
xtitle("Ishigami Histogram");
```

We can plot a bar graph of the sensitivity indices, as presented in figure 7.5.

```
for i=1:nx
    indexfirst(i)=polychaos_getindexfirst(pc,i);
    indextotal(i)=polychaos_getindextotal(pc,i);
end
scf();
bar(indextotal, 0.2, 'blue');
bar(indexfirst, 0.15, 'yellow');
```

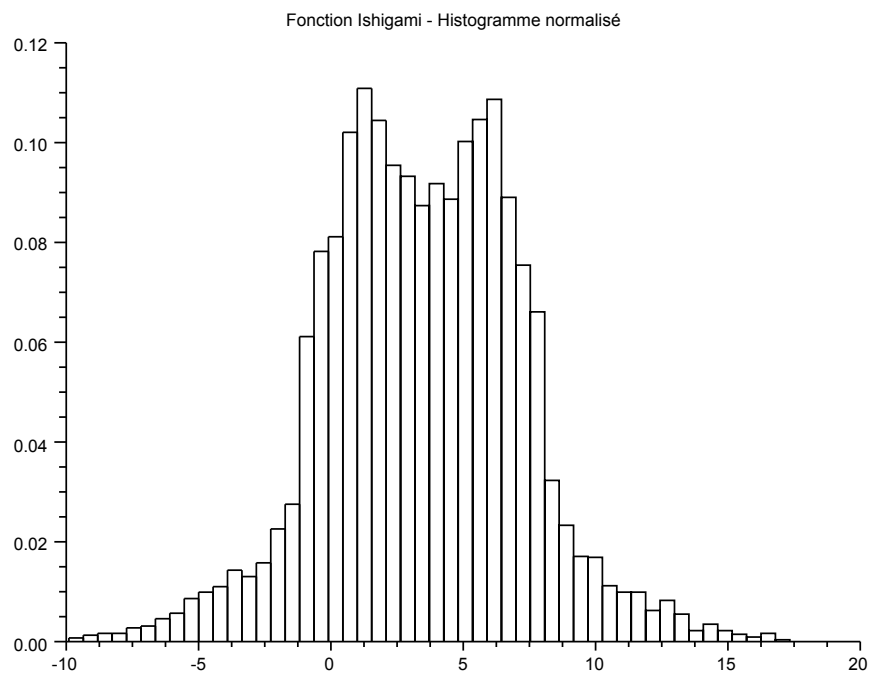


Figure 7.4: Ishigami function - Histogram of the variable on a LHS design with 10000 experiments.

```
legend(["Total" "First_order"], pos=1);  
xtitle("Ishigami - Sensitivity indices");
```

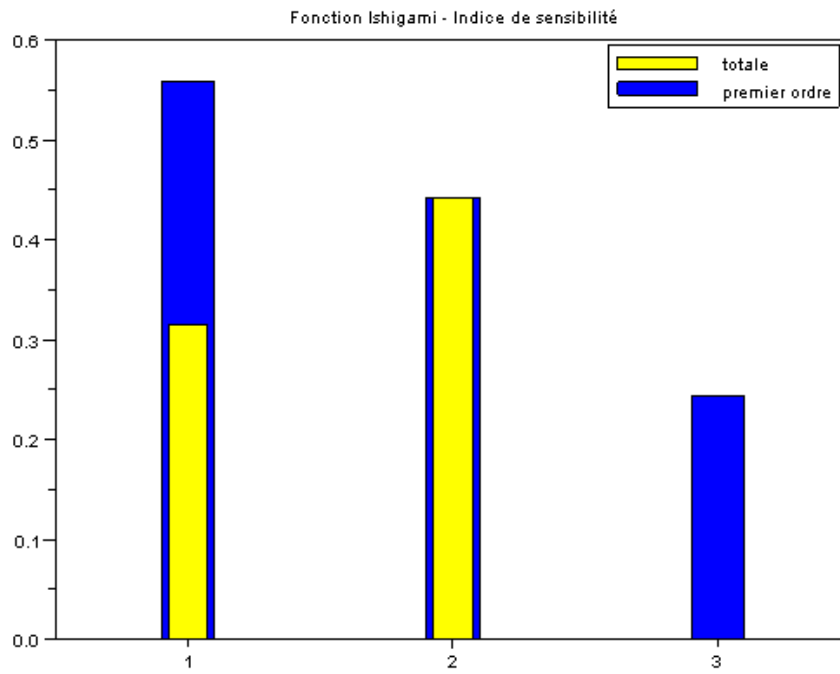


Figure 7.5: Ishigami function - Sensitivity indices.

# Chapter 8

## Thanks

Many thanks to Allan Cornet, who helped us many times in the creation of this toolbox.

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