Overview of OpenTURNS and its graphical user interface Persalys

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OpenTURNS Overview

Persalys Overview

OpenTURNS: www.openturns.org

OpenTURNS

An Open source initiative for the Treatment of Uncertainties, Risks'N Statistics

- Multivariate probabilistic modeling including dependence
- Numerical tools dedicated to the treatment of uncertainties
- Generic coupling to any type of physical model
- Open source, LGPL licensed, C++/Python library

OpenTURNS: www.openturns.org



AIRBUS







- Linux, Windows, macOS
- First release: 2007
- 5 full time developers
- lacktriangle Users pprox 1000, mainly in France (1 078 000 Total Conda downloads)
- Project size: 800 classes, more than 6000 services

OpenTURNS: content

- Data analysis
 - Distribution fitting Statistical tests
 - Estimate dependency and copulas
 - Estimate stochastic processes

- Probabilistic modeling
 - Dependence modeling Univariate distributions
 - Multivariate distributions

 - Copulas
 - Processes
 - Covariance kernels

- Surrogate models
 - Linear regression Polynomial chaos expansion
 - Gaussian process regression
 - Spectral methods
 - Low rank tensors
 - Fields metamodel

- Reliability, sensitivity
 - Sampling methods
 - Approximation methods

 - Sensitivity analysis Design of experiments

Calibration

- Least squares calibration
- Gaussian calibration
- Bayesian calibration

Numerical methods

- Optimization
- Integration Least squares
- Meshing
- Coupling with external codes



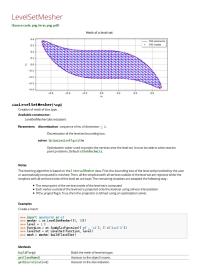








OpenTURNS: documentation



Content:

- Programming interface (API)
- Examples
- ► Theory
- All classes and methods are documented, partly automatically.
- Examples are automatically tested at each update of the code and outputs are checked.

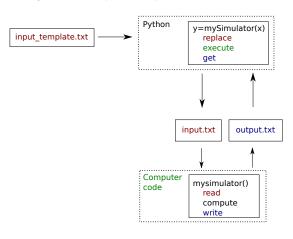
OpenTURNS: practical use

- C++ and Python interface
- ► Parallel computations with shared memory (TBB)
- Optimized linear algebra with LAPACK and BLAS
- Possibility to interface with a computation cluster
- ► Focused towards handling numerical data
- Installation through conda, pip, packages for various Linux distros and source code

```
import numpy as np
import openturns as ot
from openturns.viewer import View
ot.ResourceMap.SetAsString("Contour-DefaultColorMap", "viridis")
ot.ResourceMap.SetAsBool("Contour-DefaultIsFilled", True)
ot.ResourceMap.SetAsUnsignedInteger("Contour-DefaultLevelsNumber", 15)
```

Coupling OpenTURNS with computer codes

OpenTURNS provides a text file exchange based interface in order to perform analyses on complex computer codes



- Replaces the need for input/output text parsers
- Wraps a simulation code under the form of a standard python function
- Allows to interface
 OpenTURNS with a cluster
- otwrapy: interfacing tool to allow easy parallelization

Probabilistic modeling

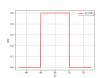
Random variables distributions:

Q: Gumbel(scale=558, mode=1013)>0



d = ot.Gumbel(558.0, 1013.0)
tr = ot.TruncatedDistribution.LOWER
0 = ot.TruncatedDistribution(d, 0.0, tr)

Zv: Uniform(min=49, max=51)



Ks: Normal(mean=30, std=7.5)>0



d = ot.Normal(30.,7.5)
tr = ot.TruncatedDistribution.LOWER
Ks = ot.TruncatedDistribution(d, 0.0, tr)

Zm: Uniform(min=54, max=56)

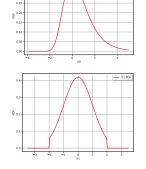


Probabilistic modeling

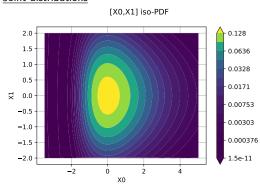
- ▶ We consider a 2-dimensional distribution with the following marginals:
 - Gumbel(min = -1, max = 1)
 - Truncated normal (mean = 0, std = 1, min = -2, max = 2)

Marginals

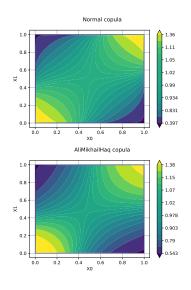
0.30

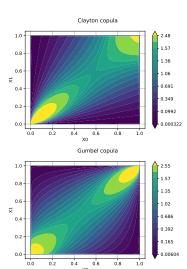


Joint distributions

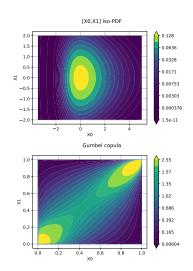


Beyond independent marginals: Copulas

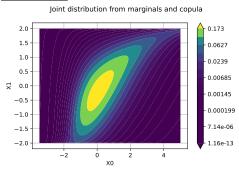




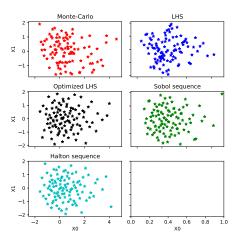
Composing marginal distributions and copulas



We obtain:



Design of experiments

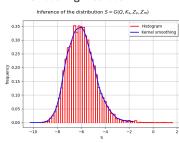


```
dim = 2
X=[ot.Gumbel(),ot.TruncatedNormal(0,1,-2,2)]
dist = ot.JointDistribution(X)
bounds = dist.getRange()
sampleSize = 100
sample1 = dist.getSample(sampleSize)
experiment = ot.LHSExperiment(dist,
  sampleSize, False, False)
sample2 = experiment.generate()
lhs = ot.LHSExperiment(dist, sampleSize)
lhs.setAlwaysShuffle(True) # randomized
space_filling = ot.SpaceFillingC2()
temperatureProfile = ot.GeometricProfile
     (10.0, 0.95, 1000)
algo = ot.SimulatedAnnealingLHS(lhs,
 space_filling, temperatureProfile)
sample3 = algo.generate()
sequence = ot.SobolSequence(dim) # or Halton
experiment = ot.LowDiscrepancyExperiment(
  sequence, dist, sampleSize, False)
```

sample4 = experiment.generate()

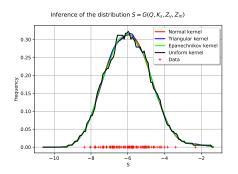
Monte-Carlo sampling

- The input distribution and relative output value are evaluated 10000 times
- The output distribution can be infered as a parametric function or through histogram or kernel smoothing methods



```
Distribution = ot. JointDistribution([Q.Ks.Zv.Zm])
#Python model
def floodFunction(X):
    Q . Ks . Zv . Zm = X
    alpha = (Zm - Zv)/5.0e3
    H = (Q/(300.0*Ks*np.sqrt(alpha)))**0.6
    S = [H + Zv - 58.5]
    return S
fun = ot.PvthonFunction(4.1.floodFunction)
#We define the output as a random vector
inputVector = ot.RandomVector(Distribution)
outputVector = ot.CompositeRandomVector(fun,
     inputVector)
#We sample and infere the output distribution
size = 10000
sampleY = outputVector.getSample(size)
hist = ot. HistogramFactory().build(sampleY)
graph = hist.drawPDF()
loiKS = ot.KernelSmoothing().build(sampleY)
graph2 = loiKS.drawPDF()
```

Distribution and dependence inference



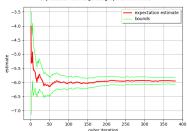
```
size = 100
sampleY = outputVector.getSample(
     size)
graph = ot.KernelSmoothing(ot.
     Normal()).build(sampleY).
     drawPDF()
loiKS = ot.KernelSmoothing(ot.
     Triangular()).build(sampleY)
graph2 = loiKS.drawPDF()
graph.add(graph2)
loiKS = ot.KernelSmoothing(ot.
     Epanechnikov()).build(sampleY)
graph2 = loiKS.drawPDF()
graph.add(graph2)
loiKS = ot.KernelSmoothing(ot.
     Uniform()).build(sampleY)
graph2 = loiKS.drawPDF()
```

- ▶ Parametric (1d Nd) distribution inference
- ► Non-parametric (1*d* − *Nd*) distribution inference
- ▶ Parametric copula inference
- Non-parametric copula inference (Bernstein copula)
- Resampling w.r.t. inferred distributions

Iterative Monte-Carlo: Central tendency anaysis

- The expected value and associated standard deviation are computed iteratively
- Different stopping criteria can be used
- Batch computation can be used

Expectation convergence graph at level 0.95



$$\widehat{\sigma}_{y} = \frac{1}{N} \sum_{1}^{N} G(\mathbf{X}_{i})$$

$$\widehat{\sigma}_{y} = \sqrt{\frac{1}{N} \sum_{1}^{N} (G(\mathbf{X}_{i}) - \hat{m}_{y})}$$

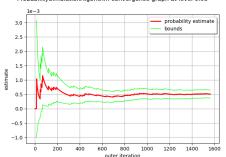
$$\widehat{\sigma}_{my} = \widehat{\sigma}_{y} / \sqrt{N}$$

```
algo = ot.ExpectationSimulationAlgorithm(
    outputVector)
algo.setMaximumOuterSampling(100000)
algo.setBlockSize(1)
algo.setGoefficientOfVariationCriterionType("MAX")
algo.setMaximumCoefficientOfVariation(0.01)
algo.run()
graph = algo.drawExpectationConvergence()
```

Iterative Monte-Carlo: Reliability analysis

- We now consider the probability of flooding: (P(S > 0.))
- ▶ Same as before, but the function $\mathbb{I}_{G(\mathbf{X}_i)>0}$ is considered

ProbabilitySimulationAlgorithm convergence graph at level 0.95



$$\widehat{\rho}_{y} = \frac{1}{N} \sum_{1}^{N} \mathbb{I}_{G(\mathbf{X}_{i}) > 0}$$

$$\widehat{\sigma} = \sqrt{\frac{1}{N} \sum_{1}^{N} (\mathbb{I}_{G(\mathbf{X}_{i}) > 0} - \hat{\rho}_{y})^{2}}$$

$$\widehat{\sigma}_{p_{y}} = \widehat{\sigma} / \sqrt{N}$$

FORM/SORM reliability analysis

- We estimate the probability of flooding through FORM/SORM procedures
- MC estimation requires $\simeq 1500$ function evaluations
- ▶ FORM and SORM only use $\simeq 150$
- Estimated probability:
 - MC: 5.0999999999998 1e-4

FORM: 5.340929030055227 1e-4 SORM: 6.793780433482759 1e-4

#FORM

```
OptAlgo = ot.Cobyla()
startingPoint = Distribution.getMean()
algoFORM = ot.FORM(OptAlgo, eventF,
startingPoint)
algoFORM.run()
```

#SORM

Different types and parameterizations of finite difference gradient computation are available

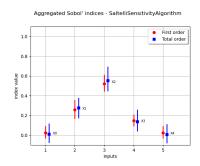
Also:

- Directional sampling
- ► Importance sampling (FORM-IS, NAIS, Adaptive IS-Cross-entropy)
- Subset sampling

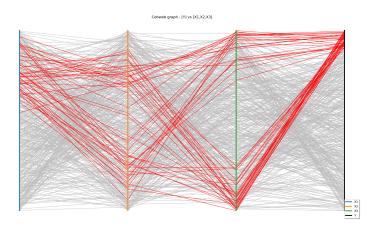
Sensitivity analysis

Various sensitivity analysis methods are available

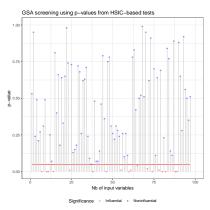
- Graphical analysis
 - Pair plots
 - ► Parallel coordinates plots
 - Cross-cuts
- Quantitative indices
 - ► SRC, SRRC, PRC, PRCC
 - ► Sobol' indices (multiple estimators)
 - FAST indices
 - ANCOVA indices
 - ► HSIC indices
 - Shapley Indices (available as a module)



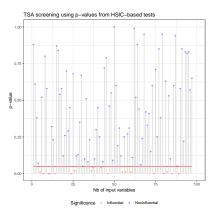
Sensitivity analysis: Parallel coordinates plot



Sensitivity analysis: HSIC indices and associated p-values



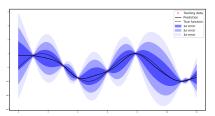
GSA-oriented screening.



TSA-oriented screening.

Surrogate modeling: Gaussian process regression

- Different surrogate modeling methods are available
 - Kriging
 - Polynomial chaos expansion
 - Linear regression & step-wise basis selection
 - Low rank tensors
 - automatic validation tools



Gaussian process regression

- Different types of covariance functions and function basis can be used
- User-defined options are also available
- MLE optimization can be parameterized
- Large number of optimization algorithms available

```
inputSample = Distribution.getSample(100)
outputSample = fun(inputSample)

dimension = 4
basis = ot.ConstantBasisFactory(dimension).
   build()
covarianceModel = ot.SquaredExponential()

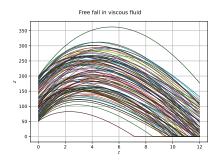
algo = ot.KrigingAlgorithm(inputSample,
   outputSample, covarianceModel, basis)

algo.run()
result = algo.getResult()
KrigingMM = result.getMetaModel()
```

Optimization

- OpenTURNS provides an interface with several optimization libraries
 - Bonmin
 - NLopt
 - dlib
 - pagmo
- Ad-hoc implementation of the COBYLA algorithm
- Constrained and unconstrained optimization
- Gradient-based and derivative-free optimizaiton
- Bounded and unbounded optimization
- Single and multi-objective optimization
- Multi-start wrapper

Field function modeling

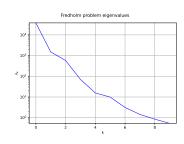


```
def free fall(X):
    g = 9.81
    z0.v0.m.c = X
    tau=m/c
    vinf = -m * g/c
    t = np.array(mesh.getVertices().asPoint())
    z=z0+vinf*t+tau*(v0-vinf)*(1-np.exp(-t/tau))
    z=np.maximum(z,0.0)
    return ot.Field(mesh, z.reshape(-1, 1))
t.min = 0
tmax = 12.
gridsize=100
mesh = ot.IntervalMesher([gridsize-1]).build(
ot.Interval(tmin, tmax))
alti = ot.PythonPointToFieldFunction(4, mesh, 1,
      free_fall)
distZ0 = ot.Uniform(50.0, 200.0)
distV0 = ot.Normal(55.0, 10.0)
distM = ot.Normal(80.0, 8.0)
distC = ot.Uniform(0.0, 30.0)
distX = ot.JointDistribution([distZ0, distV0.
distM, distC])
size = 100
inputSample = distX.getSample(size)
outputField = alti(inputSample)
```

Dimension reduction: Karhunen-Loeve decomposition

- We wish to reduce the dimension of the problem from a infinite dimensional output to a finite dimensional one
- ▶ We can perform a Karhunen-Loeve decomposition with a finite truncature
- ▶ This requires to solve a Fredholm's problem in order to identify the eigenfunctions and associated eigenvalues of the considered process

$$Y(\omega,\underline{t}) = \sum_{k=1}^{\infty} \sqrt{\lambda_k} \xi_k(\omega) \underline{\varphi}_k(\underline{t}) \to \tilde{Y}(\omega,\underline{t}) = \sum_{k=1}^{p} \sqrt{\lambda_k} \xi_k(\omega) \underline{\varphi}_k(\underline{t})$$



```
meanField = outputField.computeMean()
meanFunction = ot.PiLagrangeEvaluation(
meanField)
trend = ot.TrendTransform(meanFunction, mesh
)
invTrend = trend.getInverse()
outputFieldCentered = invTrend(outputField)

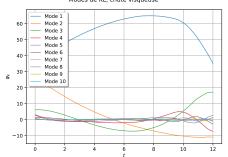
truncThreshold = 1.0e-5
algo = ot.KarhunenLoeveSVDAlgorithm(
outputFieldCentered, truncThreshold)
algo.run()
KLResult = algo.getResult()
eigenValues = KLResult.getEigenvalues()
```

Dimension reduction: Karhunen-Loeve decomposition

$$ilde{Y}(\omega,\underline{t}) = \sum_{k=1}^{p} \sqrt{\lambda_k} \xi_k(\omega) \underline{\varphi}_k(\underline{t})$$

Main modes:

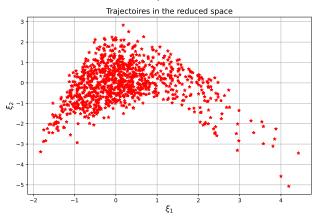
Modes de KL, chute visqueuse



```
scaledModes =
  KLResult.getScaledModesAsProcessSample()
graph = scaledModes.drawMarginal(0)
graph.setTitle('Modes de KL, chute visqueuse
  ')
graph.setXTitle(r'$t$')
graph.setYTitle(r'$t\sqrt{v}\parphi_k$')
leg = ot.Description(['Mode '+str(i +1) for
  i in range(eigenValues.getDimension()) ])
graph.setLegends(leg)
graph.setLegendSoleg)
graph.setLegendPosition('topleft')
view=View(graph)
```

Dimension reduction: Karhunen-Loeve decomposition

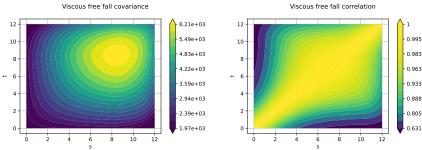
We only consider the first 2 terms of the decomposition:



```
projectionFunction = ot.KarhunenLoeveProjection(KLResult)
sampleKsi = projectionFunction(outputFieldCentered)
sampleKsi = sampleKsi[:,:2]
```

Field function analysis

We center the trajectories with respect to the mean field:



```
cov = KLResult.getCovarianceModel()

# As a covariance function
isStationary = False
asCorrelation = False
graph = cov.draw(0, 0, tmin, tmax, 128, isStationary, asCorrelation)

# As a correlation function
asCorrelation = True
graph = cov.draw(0, 0, tmin, tmax, 128, isStationary, asCorrelation)
```

Contents

OpenTURNS Overview

Persalys Overview

Project overview

- ▶ Partnership between EDF and Phimeca since 2015
 - ▶ Developed in C++ using Qt
 - ► Aiming at maximizing the use of OpenTURNS through a dedicated GUI for engineers/researchers without a strong coding experience
 - ► As easy to use as possible while providing the user with help and guidelines
 - ▶ Benefit from the advanced visualization capability of Paraview

- ► Features:
 - Uncertainty quantification:
 - Probabilistic model definition
 - Distribution fitting
 - Probability estimate
 - Metamodeling
 - Screening
 - Optimization
 - Design of experiments
 - As generic as possible
 - ► Allows for a wide variety of models
 - Can be coupled to external code
 - ► GUI language in both English and French
- LGPL license
- Two releases per year, follows OpenTURNS development
- Available for free on demand at https://persalys.fr

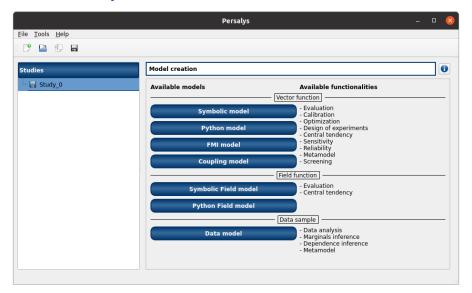
Persalys installation

- Github: sources
- You can request executables at https://persalys.fr/obtenir.php?la=en
- Depending on your OS
 - ▶ Linux → .AppImage (600 Mo)
 - ightharpoonup Windows ightarrow .exe will create a shortcut on your Desktop (program is 1.45 Go)
- Also distributed by Debian-based GNU/Linux distributions (e.g. Debian, Ubuntu...)

Open Persalys and click on "New study"



Create a study



Definition/Evaluation

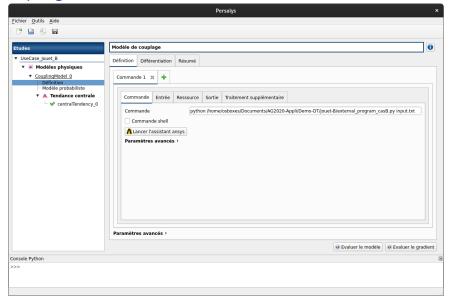
Models are viewed as a "black box" with inputs, outputs and a transfer function (TF). Persalys supports two model categories:

- ▶ vector to vector $(X_i \rightarrow Y_i$, emphasized here)
- ▶ vector to 1D field $(X_i \rightarrow Y_i(t))$

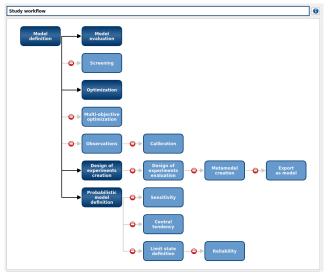
Vector to vector models can be of the following type:

- ▶ Symbolic, TF is a mathematical formula
- Python, TF is a Python function
- FMI, TF is provided by an FMU model
- Coupling, TF is an executable command which reads/writes input/output files

Coupling model definition



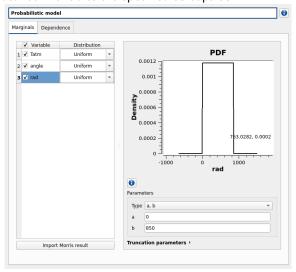
Study workflow



Blocks become available as study content grows and prerequisites are fulfilled.

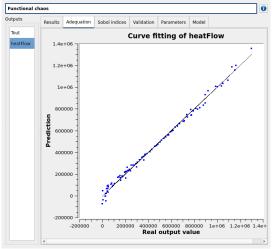
Probabilistic models

Each input variable can be associated to a distribution. Dependencies between variables are specified as copulas.



Metamodel (surrogate model) creation

Using an evaluated design of experiments, the user can build a surrogate model (linear regression, functional chaos or Gaussian process regression). Validation tests are run to check the approximation being made.



OpenTURNS resources

- Website and documentation: www.openturns.org
- GitHub: https://github.com/openturns/openturns
- Forum: https://openturns.discourse.group

Persalys resources

- ▶ Website and documentation: https://persalys.fr/?la=en
- ► Forum: https://persalys.discourse.group



