# Getting Started with ProSimPlus®

# Case 7: Integrating Artificial Intelligence (AI) into ProSimPlus



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Artificial Intelligence (AI) and process simulation can work together to provide significant advances in the field of process engineering.

By exploiting advanced Machine Learning techniques, AI can analyze existing simulation models, extract key information and create faster predictive models.

These AI-powered models, also known as Surrogate Models, can significantly reduce computing times by optimizing complex process engineering operations.

### Introduction

This document outlines the steps involved in creating a Surrogate Model in ProSimPlus.

The steps are as follows:

- 1. Creation of a dataset by ProSimPlus
- 2. Model training
- 3. Model deployment within ProSimPlus

Before delving into this chapter, it is highly recommended to refer to "Getting Started with ProSimPlus, Use Case 1," which introduces the primary features of ProSimPlus

References:

R. Bounaceur, O. Baudouin, "Couplage entre logiciel PSE et modèles fondés sur des algorithmes d'Intelligence Artificielle", tutorial SFGP 2022, Toulouse (2022)

R. Bounaceur *et al.*, "Development of an artificial intelligence model to predict combustion properties, with a focus on auto-ignition delay", J. Eng. Gas Turbines Power., 1-28 (2023)

### Prerequisite

- Firstly, installation of the Python software is required.
- A few libraries for Python can be used, including (but not limited to):

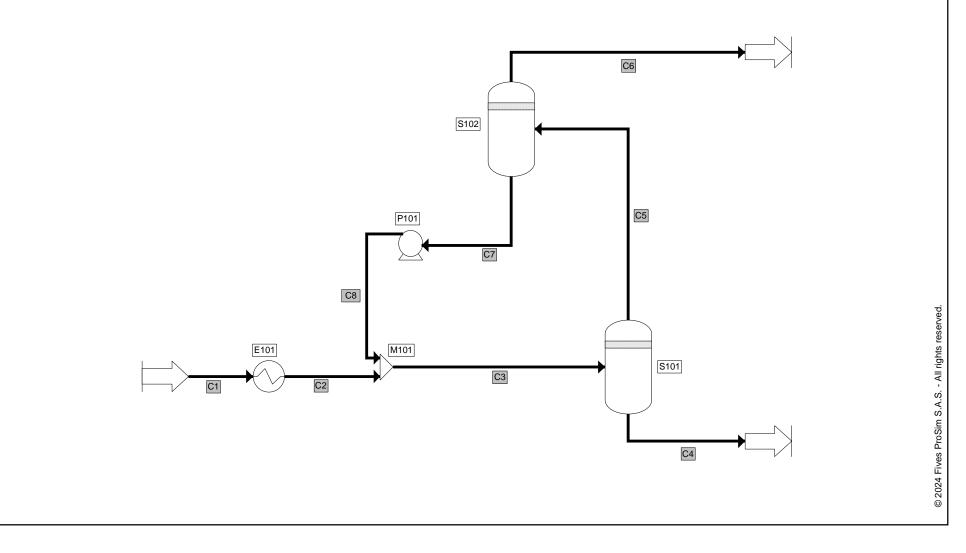
Library	Function
Pywin32	Integrate the surrogate model into ProSimPlus
NumPy	Manipulate matrices or multidimensional arrays in Python
Pandas	Manipulate objects (dataframe)
Matplotlib	Create graphics
Scikit-learn	Open-source Python library for machine learning
Jupyter Create and execute Python code interactively, combine code, visual explanatory text and results	
JobLib	Save trained models and reload them later
	explanatory text and results

To install these libraries, open the command prompt and type:

pip install Library name

### Introduction of the process: Simple Example

For this study, the example is based on a simple process available in the ProSimPlus samples directory under the name "PSPS\_EX\_EN-Simple-Example".



- The first step is to choose the training subject and generate a dataset.
- In the absence of experimental data, we will opt to use ProSimPlus with an external solver to generate the necessary values. An alternative would be to use the sensitivity analysis or the console version, but this would be less practical.



The Python source file that will be used to generate the dataset is called "train.py"

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The detailed explanation of the parameters to be modified in the Python file can be found directly within the document: "train.py"

- We need to specify the input values (module parameters) and run simulations to calculate the output values (results).
- We will ensure to avoid creating an input dataset that is entirely random or excessively linear, as it may not necessarily be representative of our model.
- To achieve this, we will use the Sobol sequences method, which will easily provide us with quasi-random values that adequately cover the selected intervals.

- The "sobol.py" python source code is included with this document.
- The invocation of the Sobol sequences method is performed as follows:

tab = sobol.i4\_sobol\_generate(ndim, npoints, npass)

- ndim is the number of dimensions (inputs) (from 1 to 40),
- **npoints** is the total number of points,
- **npass** is the number of initial points in the sequence to skip, in order to avoid encountering the first points that are nearly identical.
- The returned value **tab** is a fixed-size homogeneous NumPy array, with dimensions corresponding to the provided arguments.

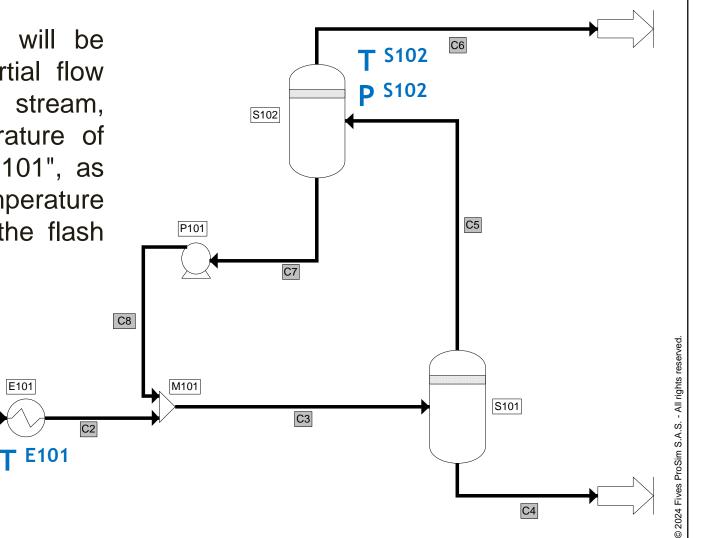


It is possible to use other sampling methods than Sobol sequences, but they are not explained in this document.

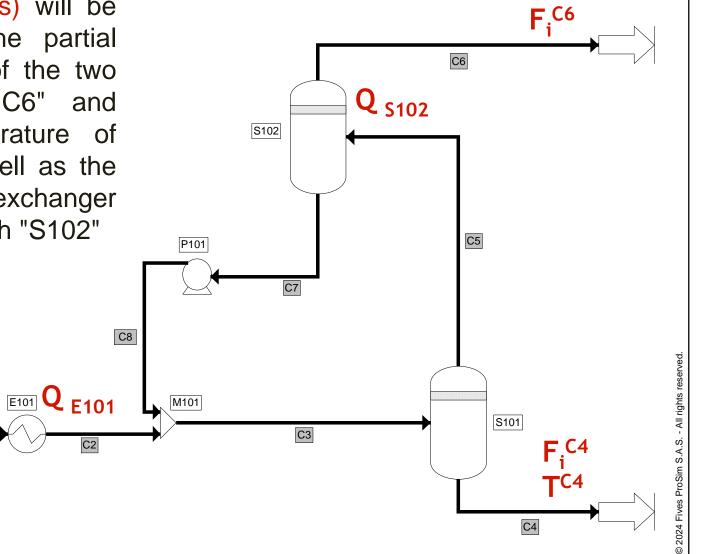
We are going to assume that:

the data (inputs) will be limited to the partial flow rates of the feed stream, the outlet temperature of the exchanger "E101", as well as the temperature and pressure of the flash "S102".

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the results (outputs) will be represented by the partial molar flow rates of the two outlet streams ("C6" and "C4"), the temperature of stream "C4", as well as the heat duties of the exchanger "E101" and the flash "S102"



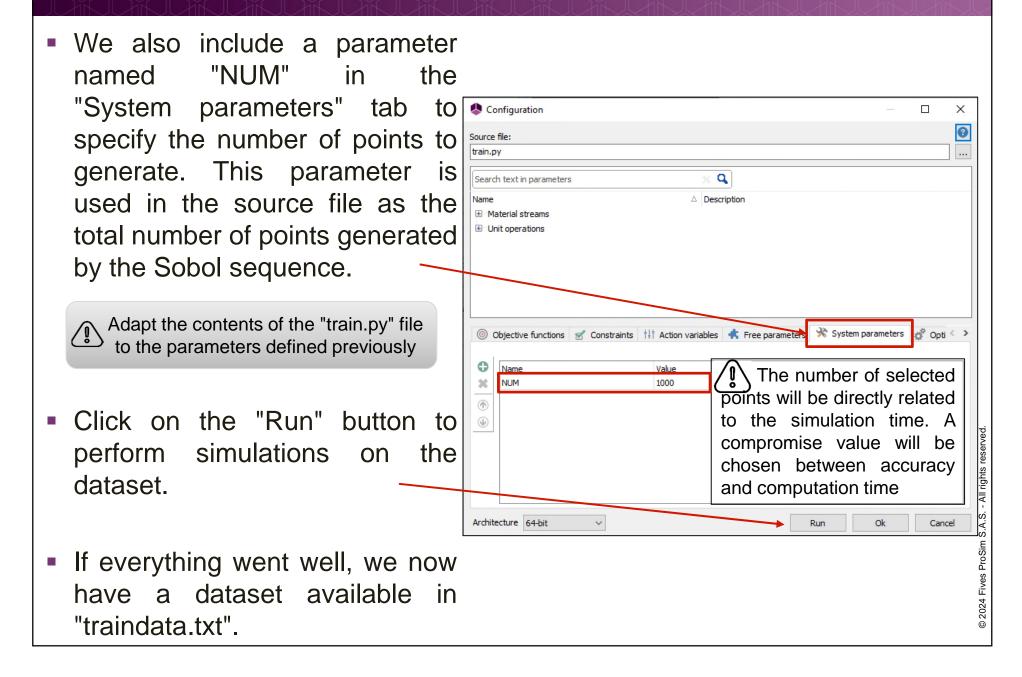
To achieve this, we will configure the external solver by incorporating the results (outputs) into the "objective functions":

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0	Parameter	Indexes	Unit					
	E101.HeatDuty		kcal/h					
-	S102.HeatDuty		kcal/h					
۲	C4.PartialMolarFlowrate.1	1	kmol/h					
٢	C4.PartialMolarFlowrate.2	2	kmol/h					
	C4.PartialMolarFlowrate.3	3	kmol/h					
	C4.PartialMolarFlowrate.4	4	kmol/h					
	C4.PartialMolarFlowrate.5	5	kmol/h					
	C4.PartialMolarFlowrate.6	6	kmol/h					
	C4.PartialMolarFlowrate.7	7	kmol/h					
	C4.PartialMolarFlowrate.8	8	kmol/h					
	C4.PartialMolarFlowrate.9	9	kmol/h					
	C4.Temperature		°C					
	C6.PartialMolarFlowrate.1	1	kmol/h	🗌 🖳 🖓 Results (outputs) = Objective function				
	C6.PartialMolarFlowrate.2	2	kmol/h					
	C6.PartialMolarFlowrate.3	3	kmol/h					
	C6.PartialMolarFlowrate.4	4	kmol/h					
	C6.PartialMolarFlowrate.5	5	kmol/h					
	C6.PartialMolarFlowrate.6	6	kmol/h					
	C6.PartialMolarFlowrate.7	7	kmol/h					
	C6.PartialMolarFlowrate.8	8	kmol/h					
	C6.PartialMolarFlowrate.9	9	kmol/h					

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#### And the data (inputs) into the action variables:

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D	Feed.OutputStreamCompositionSpecValues.4			Real (continuo		7,4	
_	Feed.OutputStreamCompositionSpecValues.5	5		Real (continuo		6,5	
	Feed.OutputStreamCompositionSpecValues.6	-		Real (continuo		3,6	
	Feed.OutputStreamCompositionSpecValues.7			Real (continuo		9,7	
	Feed.OutputStreamCompositionSpecValues.8			Real (continuo		16	
	Feed.OutputStreamCompositionSpecValues.9			Real (continuo		2,5	
	E101.TemperatureSpecValue			Real (continuo		313	
	S102.TemperatureSpecValue			Real (continuo		253	
	S102.PressureSpecValue		atm	Real (continuo	10	70	
							a (inputs) = Action variables



To make things clearer and more understandable, this part will be carried out in Jupyter Notebook

- Open the command prompt and type the following: Jupyter notebook
- Open the notebook titled "Model Training.ipynb"

💭 jupyter		Quit	Log	out
Files Running Clusters				
Select items to perform actions on them.		Upload	New -	C
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🗋 🕘 Model Training.ipynb		il y a 2 minutes	49.1	kB
The explanation regarding the parameters to be modified for the model training process is detailed in this file				

#### 1. Data verification and cleaning

Data cleaning is a crucial step before creating a machine learning model. It involves taking measures to process and correct raw data so that the model can learn effectively and produce high-quality results.

The main steps of the process are as follows:

#### Data Collection

Data Processing (Handling incorrectly labeled data, dealing with "NaN" values...)

#### Normalization or Scaling

If the various data inputs have significantly different scales, it is advisable to normalize or scale them to prevent certain inputs from excessively dominating the model.

#### Data Splitting

Divide the data into <u>training</u> and <u>test</u> sets to evaluate the model's performance in an unbiased way. It is possible to choose the percentage associated to each set.

#### 2. Machine Learning Algorithm

Various learning methods are available, and in this study, we are using a **KernelRidge regression model**, which is a regularized regression technique utilizing kernel methods for predictions. Other learning models could also be employed, such as for instance:

- kNN (k-Nearest Neighborhood)
- DecisionTree
- RandomForest
- MLP (Multi-Layer Perceptron) Artificial Neural Networks

There is no one-size-fits-all solution to determine the best regression algorithm for all scenarios. The most recommended approach is to experiment with several models, evaluate them in terms of performance, and select the one that achieves the best results on the available dataset.

#### 3. Evaluation parameters

Here are some commonly used metrics to evaluate the performance of learning models, especially in the case of regression:

#### Coefficient of Determination (R2 score)

A high R2 score ( $\approx$ 1) indicates that the model effectively explains the data variation, while an R2 score close to 0 or negative suggests that the model does not accurately represent the data.

#### Mean Absolute Error (MAE)

The MAE quantifies the average difference between the model's predicted values and the actual values.

#### Mean Squared Error (MSE)

The MSE measures the average of the squared differences between predicted values and actual values. A lower MSE indicates that the model's predictions are closer to the actual values.

The obtained metrics should be in the same order of magnitude for the training and test sets.



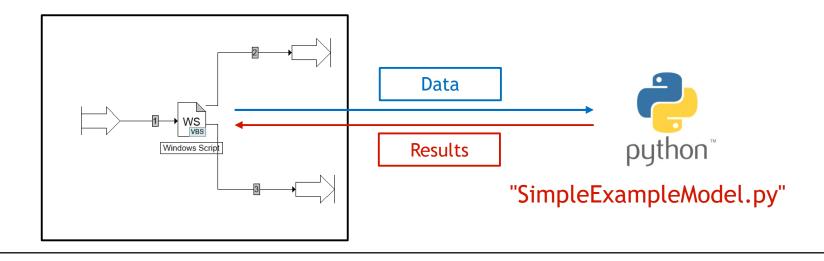
 Additionally, we store the minimum and maximum values of both the input data and the output results. As our model is normalized, these values are needed for the transformation of results back into their original scale.

```
# Saving it with joblib
joblib.dump(minmax, "parameters.joblib")
```

Following the training phase of the model in Jupyter Notebook, a model is created and stored in the "model.joblib" file. The next step involves setting up the necessary framework to integrate it into ProSimPlus.

This structure is relatively simple and will consist of:

- A Windows Script unit operation
- A Python program able to make predictions by processing the data and providing the output results



We will now create a Python component following the Microsoft COM standard, which will be used from the VBScript code. The primary advantage of this method is that the Python program can remain in memory throughout the entire simulation.

The source code for this component can be found in the file "SimpleExampleModel.py".

The code is thoroughly described through comments within the file.

Here are some explanations about the codes:

#### \_reg\_progid\_

The ProgID (Programmatic Identifier) is a character string used to uniquely identify a class of COM (Component Object Model) objects that can be created and utilized within a script. This identifier is crucial and plays a pivotal role when instantiating an object in VBScript (CreateObject("My.progid")).

\_reg\_progid\_ = "ProSim.SimpleExampleModel"

#### \_reg\_clsid\_

The GUID(Globally Unique Identifier) is a unique identifier used to reference the class.

```
# https://guidgenerator.com/online-guid-generator.aspx for example. And don't forget the braces!
_reg_clsid_ = "{f8084d10-d072-4b8e-851f-4adeaac4d371}"
```

#### \_public\_methods\_

In this context, we are dealing with an array of strings that define the methods (functions or routines) that will be accessible within our class. For our example, we are introducing only two methods: one for performing calculations (predictions) and another for printing results in the report.

```
_public_methods_ = ["Calculate", "PrintResults"]
```

#### Using the "model.joblib" file

The "Calculate" method will be used for making predictions.

```
def Calculate(self, F1, F2, F3, F4, F5, F6, F7, F8, F9, T_Ech, T_S102, P_S102)
```

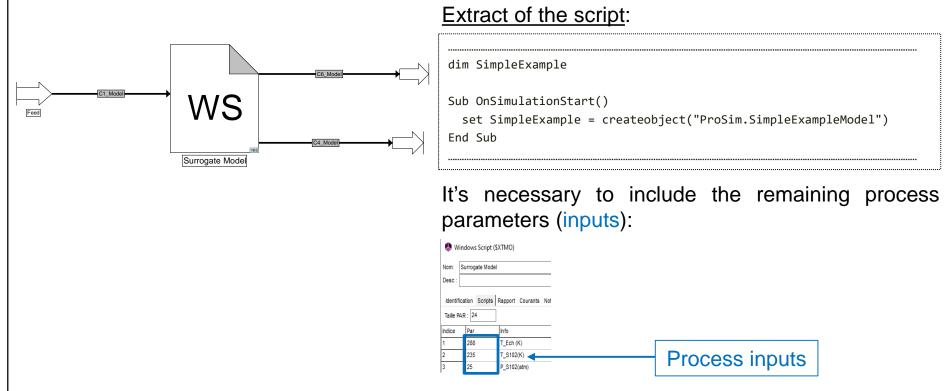
To do this, the process involves retrieving the input data and predicting the results by simply calling the 'predict' method. The use of this 'predict' method becomes possible after loading the model that was previously saved using the joblib file.

```
features = pandas.DataFrame(data, columns=["F1", "F2", "F3", "F4", "F5", "F6", "F7", "F8", "F9", "T_Ech", "T_S102", "P_S102"])
res = model.predict(features)
```

The result is returned in the form of a two-dimensional array:

return res

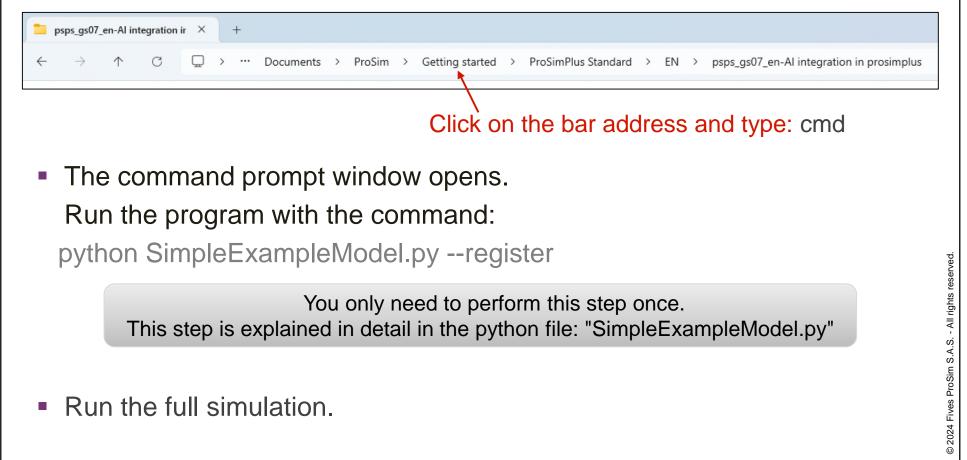
A ProSimPlus test file utilizing this component is also available, named "SimpleExample.pmp3".



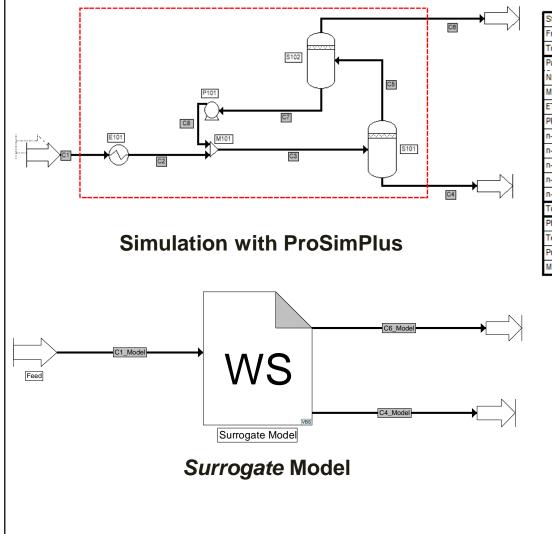
In summary, the script performs the following actions: it creates an instance to obtain the ProgID of the class, retrieves the properties of the current input, incorporates the process parameters (inputs), generates the output streams, and retrieves the results returned by the training model.

The final step consists of simulating the process using the surrogate model we have built. To achieve this:

• Open the file directory:



#### Example of the result:



Streams	C1	C4	C4_Model	C6	C6_Model	
From	Stream dupli	S101	Surrogate M	S102	Surrogate M	
То		E101	Process out	Process out	Process out	Process out
Partial flows (molar)	kmol/h	kmol/h	kmol/h	kmol/h	kmol/h	
NITROGEN		9	1.6583	1.6401	7.3417	7.36
METHANE		41.7	15.493	15.23	26.207	26.471
ETHANE	11.2	8.1088	8.1265	3.0912	3.0727	
PROPANE	6.2	5.5336	5.591	0.66639	0.60812	
n-BUTANE	5.4	5.2748	5.2774	0.12516	0.12263	
n-PENTANE	3	2.9895	2.9863	0.01049	0.013583	
n-HEXANE	8.1	8.0981	8.0962	0.0018615	0.0035282	
n-HEPTANE	13.3	13.3	13.3	0.00033901	0.00067467	
n-OCTANE		2.1	2.1	2.0998	5.2835E-006	1.0028E-00
Total flow (mass)	kg/h	4332.2	3575.5	3573.6	756.65	758.52
Physical state		Liq./Vap.	Liquid	Liquid	Vapor	Liq./Vap.
Temperature	°C	40	14.561	15.598	-38	-38.15
Pressure	atm	75	74.7	75.994	25	25
Molar vapor fraction	Molar vapor fraction			0	1	0.99974





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