

# *Reduced Order Models (ROMs)*

RAVEN Workshop



[www.inl.gov](http://www.inl.gov)

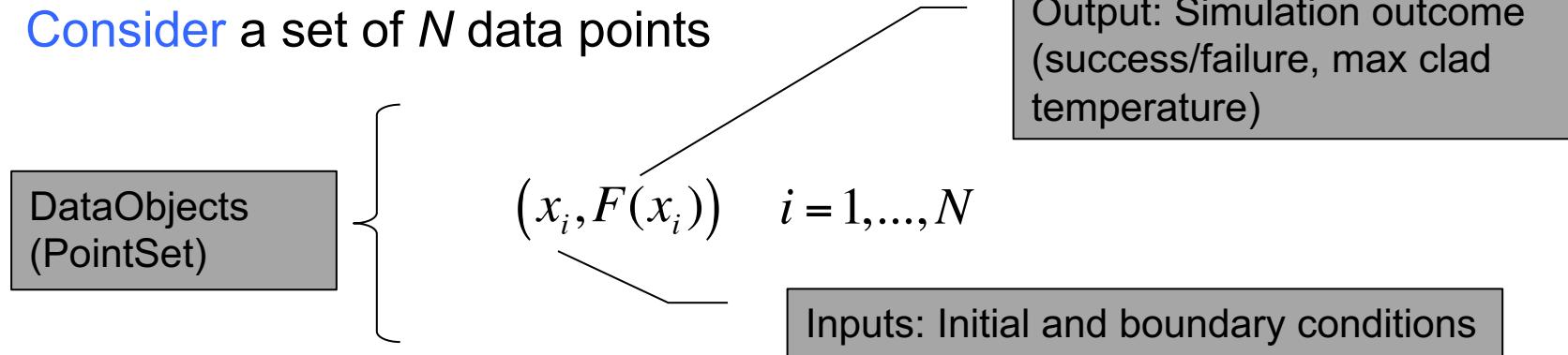


# Outline

- Brief introduction on ROMs
- Application examples of ROMs
- ROMs and RAVEN
  - Available ROMs
  - RAVEN ROM workflow
- RAVEN examples
  - Create ROMs
  - Perform sampling of ROMs

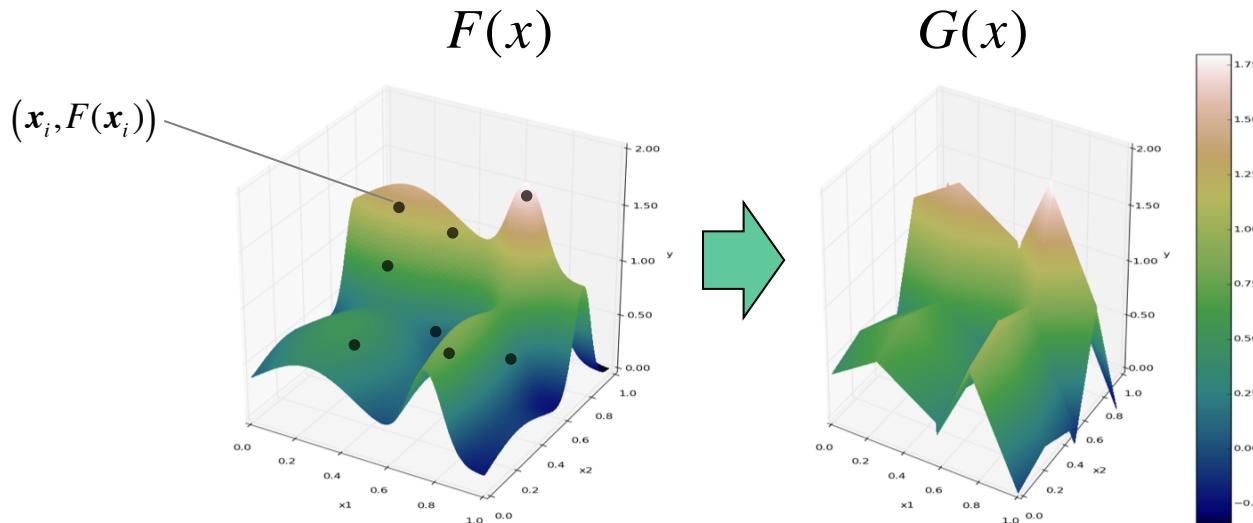
# ROMs: a Quick Introduction

- Consider a set of  $N$  data points



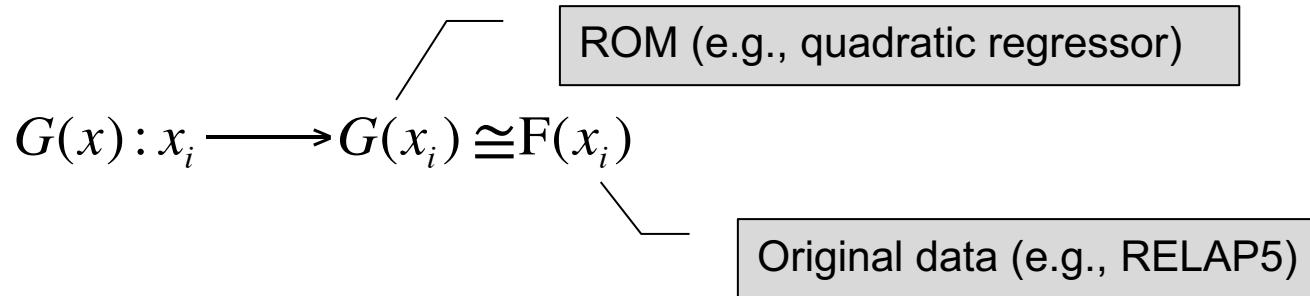
- Build a surrogate model

– Reduced Order Model  $G(x) : x_i \longrightarrow G(x_i) \cong F(x_i)$



# ROMs: a Quick Introduction

- Basically we are trying to **reduce the complexity** of the original model



- Pros:
  - Much **faster computation** of the output variable
- Cons:
  - Presence of **error** in the ROM computed values

# Classes of ROMs

- Model-Based
  - Prediction is performed using a blend of **interpolation and regression** algorithms
  - Examples:
    - Gaussian Process Models (GPMs)
    - Support Vector Machines (SVM)
- Data-Based
  - Prediction is performed by solely considering the input data by using **data searching** algorithms
  - Example:
    - K nearest neighbor (KNN)

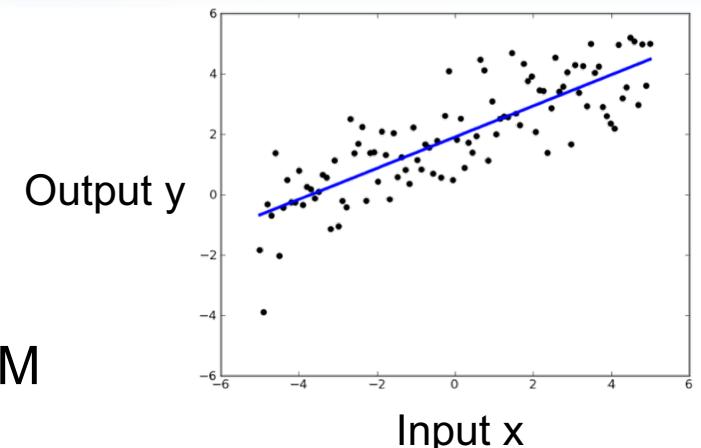
# ROMs: Applications

- Basic steps:

1. Sample original model
2. Train the ROM

$$y = mx + c$$

3. Perform desired analysis with the ROM instead of the original model



- Range of applications:

- Uncertainty quantification / Sensitivity analysis
- Probabilistic Risk Analysis (PRA)
- Accelerator for stochastic analysis (adaptive sampling)
- Prediction models

# *ROMs Available in RAVEN*

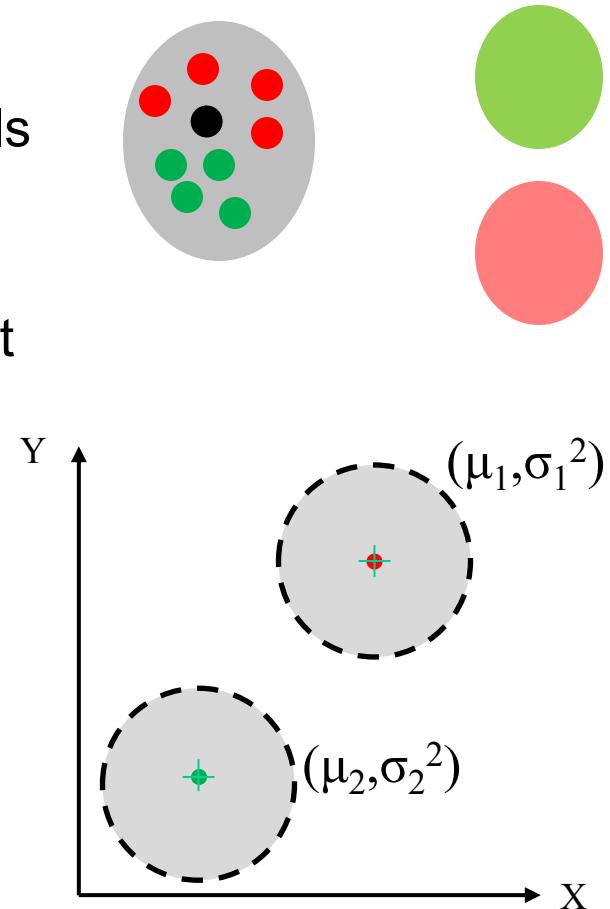
- External dependency: [Scikit-learn](http://scikit-learn.org) (<http://scikit-learn.org>)
  - Open source machine learning library for Python
  - Library for [data mining](#) and [data analysis](#)
  - Built on [NumPy](#) and [SciPy](#)
- Internally C++ developed libraries: [CROW](#)
  - [Generalized Polynomial Chaos](#)
  - [Multi-dimensional interpolators](#)
  - [Synthetic Time-Series Algorithms](#) (e.g. F-ARMA)
  - [Dynamic Mode Decomposition-based algorithms](#)
  - [Polynomial Regression](#) (Poly-Tensor spline, Poly-exponential, etc.)

# *Supervised and Unsupervised Library*

- Available:
  - Classification: identifying to which category an object belongs
  - Regression: predicting a continuous-valued attribute
  - Data clustering: grouping similar objects into sets
  - Dimensionality reduction: reducing the number of random variables
  - Data pre-processing: feature extraction and normalization
- Examples:
  - Linear regression models
  - Support Vector Machines
  - Multi-Class classifiers
  - Naïve Bayes
  - Neighbors classifiers
  - Tree classifiers

# *Supervised and Unsupervised Library*

- **Classification**
  - Starting point: set of data points with labels  
 $[features, class]_i$
  - Objective: identify which class a new point  
 $[features]$  belong
  
- **Clustering**
  - Starting point: set of data points  
 $[features]_i$
  - Objective: group data points based on a specific distance metrics

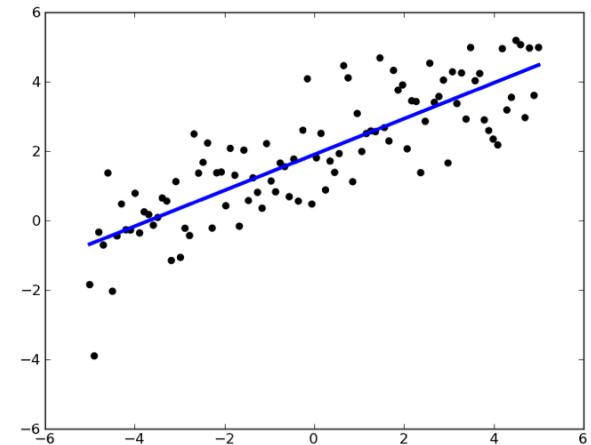


# *Supervised and Unsupervised Library*

Starting point: set of data points

[features]<sub>i</sub>

- **Cardinality reduction**
  - Objective: identify the most relevant features that keep data points unique
  - Outcome: Location of the points on the reduced space (e.g., line)
  
- **Regression**
  - Objective: estimate the relationships among variables via a statistical process
  - Outcome: coefficients of the reduced space (e.g.,  $m$  and  $c$  for linear interpolator  $y = mx + c$ )



Source: scikit-learn.org

# Generalized Polynomial Chaos

- **Objective:** overcome limitations of Monte-Carlo sampling
  - High number of samples
  - Computationally expensive
- **Polynomial representation** of an output variable  $\Phi_k(Y)$ 
  - Simpler to evaluate
  - Easy to get statistical moments
  - Less effort and more accurate than Monte Carlo

$$u(Y) \approx \sum_{k \in \Lambda(L)} u_k \Phi_k(Y)$$

$\Phi_k(Y) = \prod_{n=1}^N \Phi_{k_n}^{(n)}(y_n)$

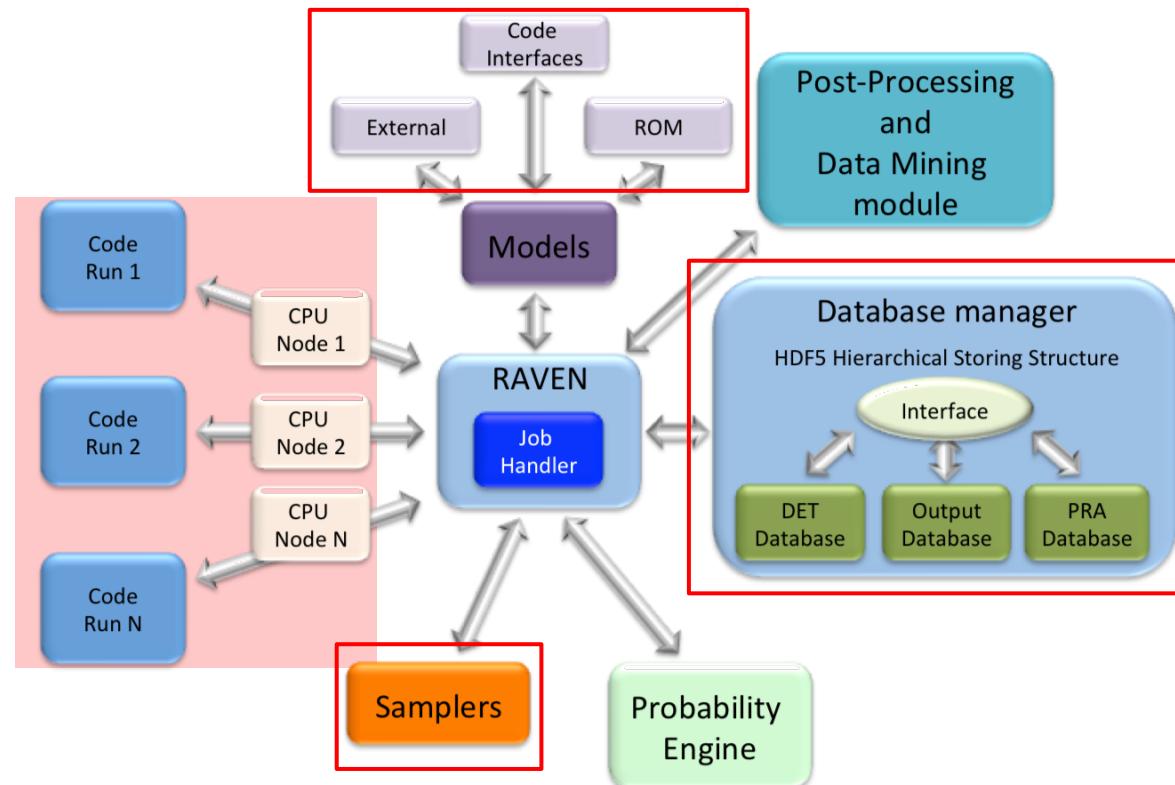
index set of all desired polynomial  
 orders up to order  $L$

## *Multi-Dimensional Interpolators*

- **CROW**: Internally developed C++ library
- Interpolation on any dimension
- Response surface is created as an **interpolation function** given a set of data points defined on:
  - Sparse grid
  - Cartesian grid
- Extension of the known 1-D interpolation schemes

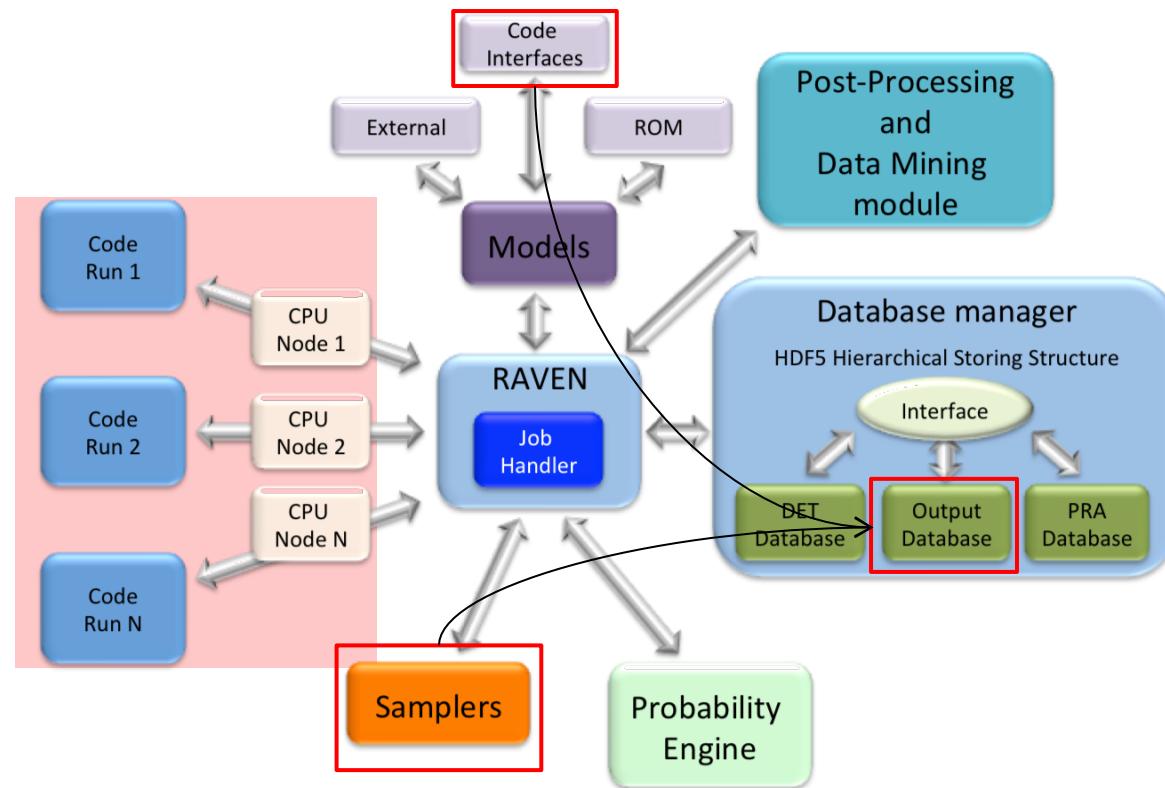
# ROM Modeling Within RAVEN

- All modeling steps that involve ROMs are available in RAVEN
  - Create ROMs from a database
  - Perform statistical analysis using ROMs



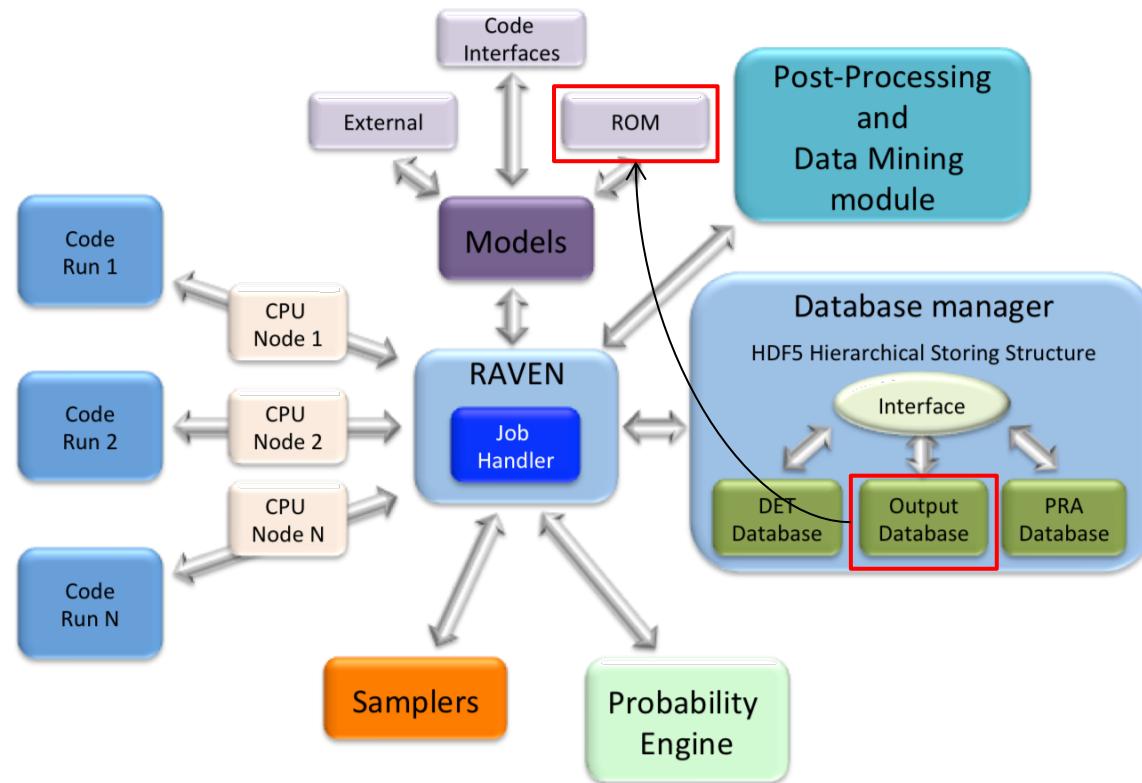
# ROM Modeling Within RAVEN

- Create a Database (PointSet)



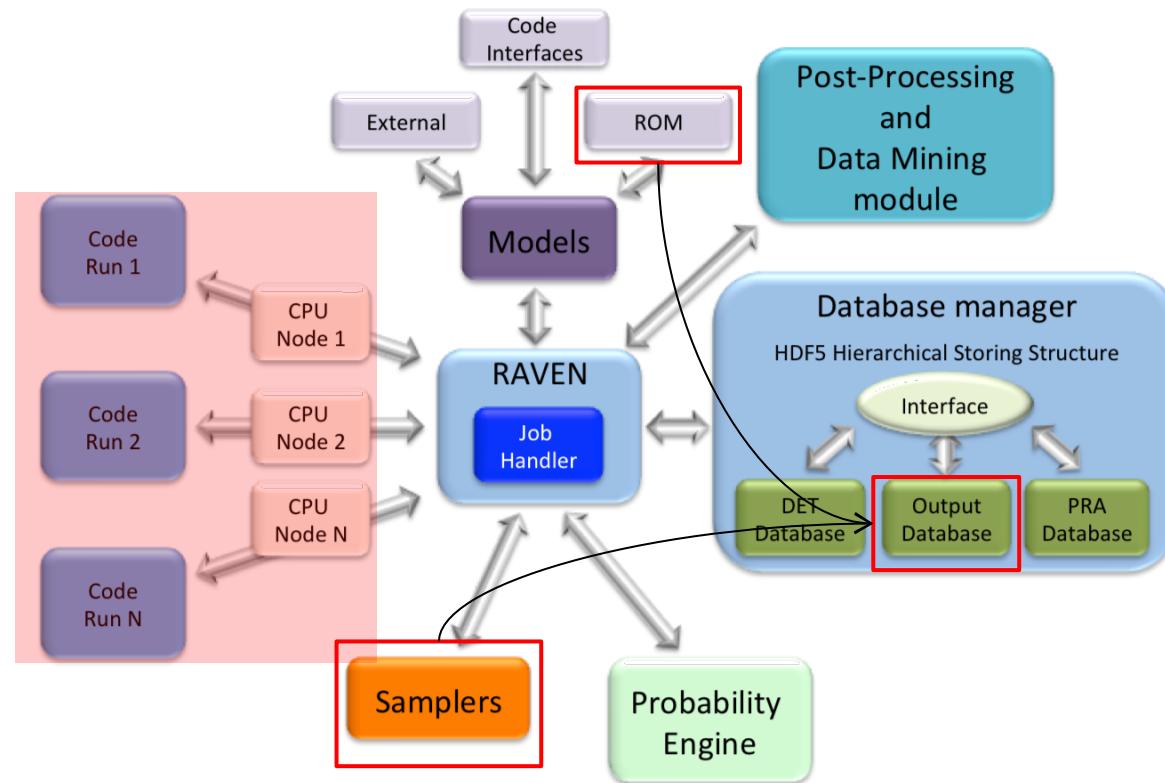
# ROM Modeling Within RAVEN

- Create and train a ROM from a Database



# ROM Modeling Within RAVEN

- Perform statistical analysis using the ROM



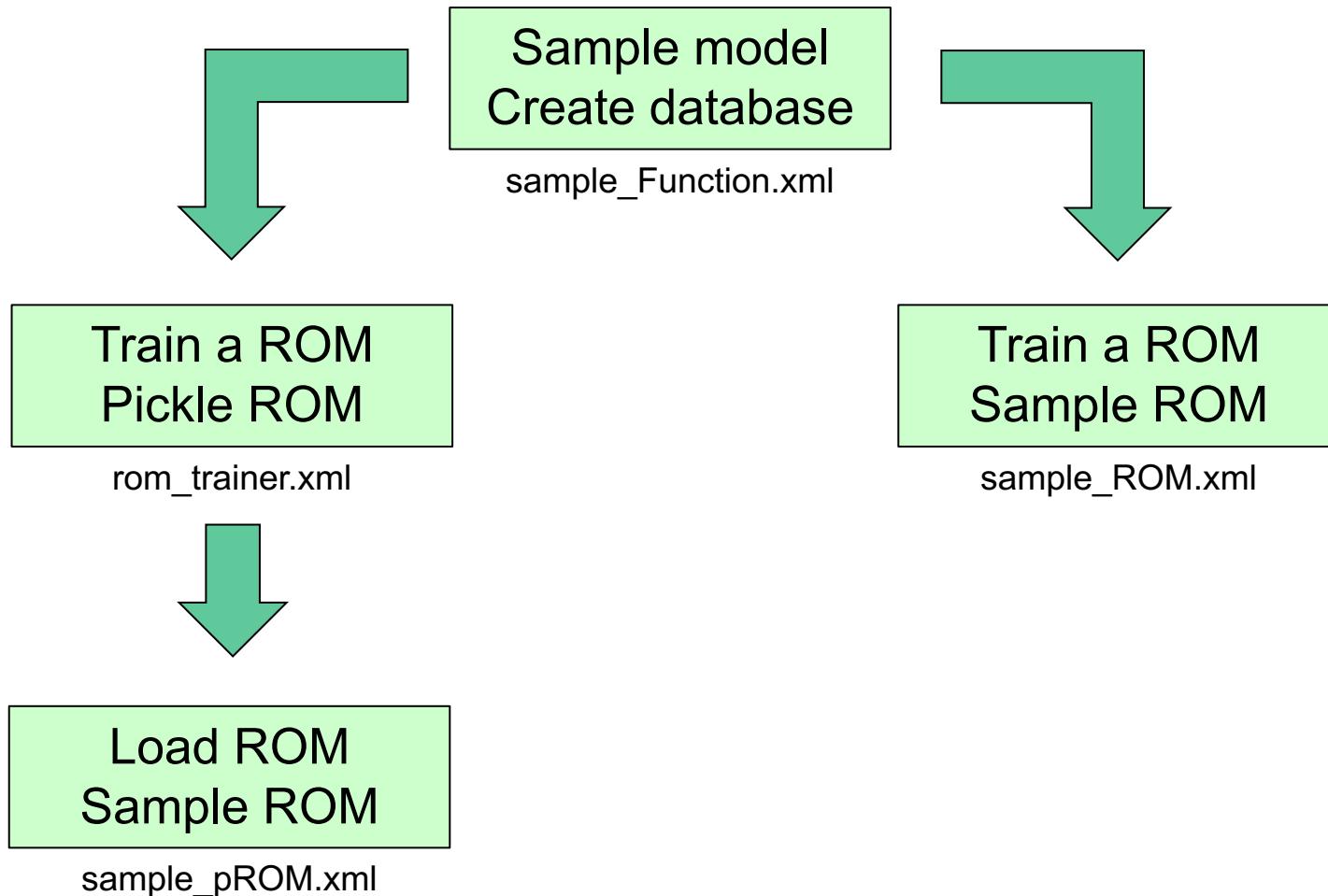
# ROM Pickle

- Compression/serialization scheme
- Pickled object contains all the information necessary to **reconstruct** the object in another python script
- Pickled object can be **saved** as a file
- RAVEN ROMs can be pickled
- **Applications:**
  - Perform statistical analysis on a ROM after they have been generated and/or on a different machine
  - Use pickled ROMs on separate python script (external model for RAVEN)
  - Stochastic analysis for different distributions

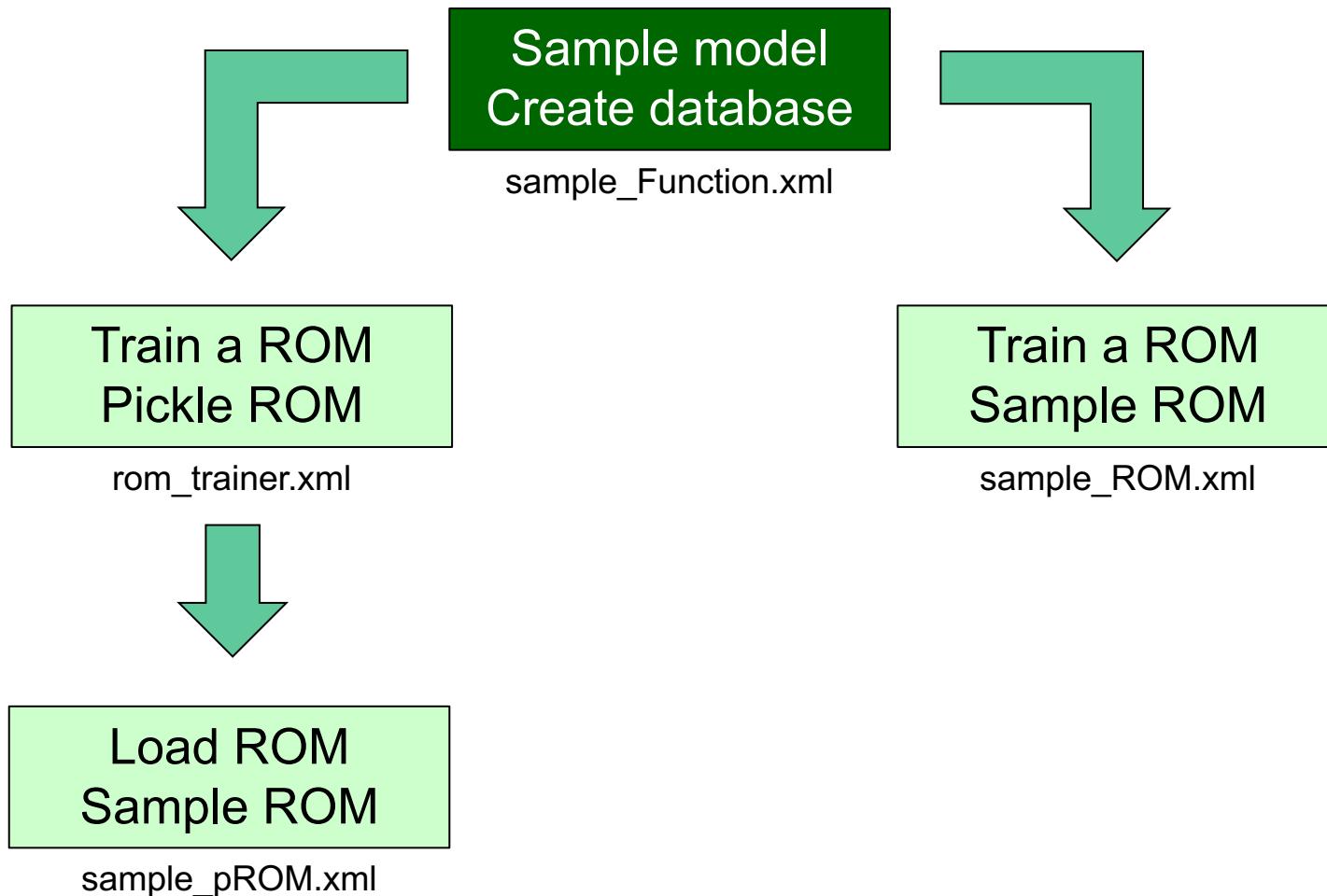
## *RAVEN Examples*

# Workflow

External model employed:  
workshop\_model.py



# Workflow



# Sample a Model and Create a Database

Distributions	Models	Samplers	Databases	DataObjects	Steps
---------------	--------	----------	-----------	-------------	-------

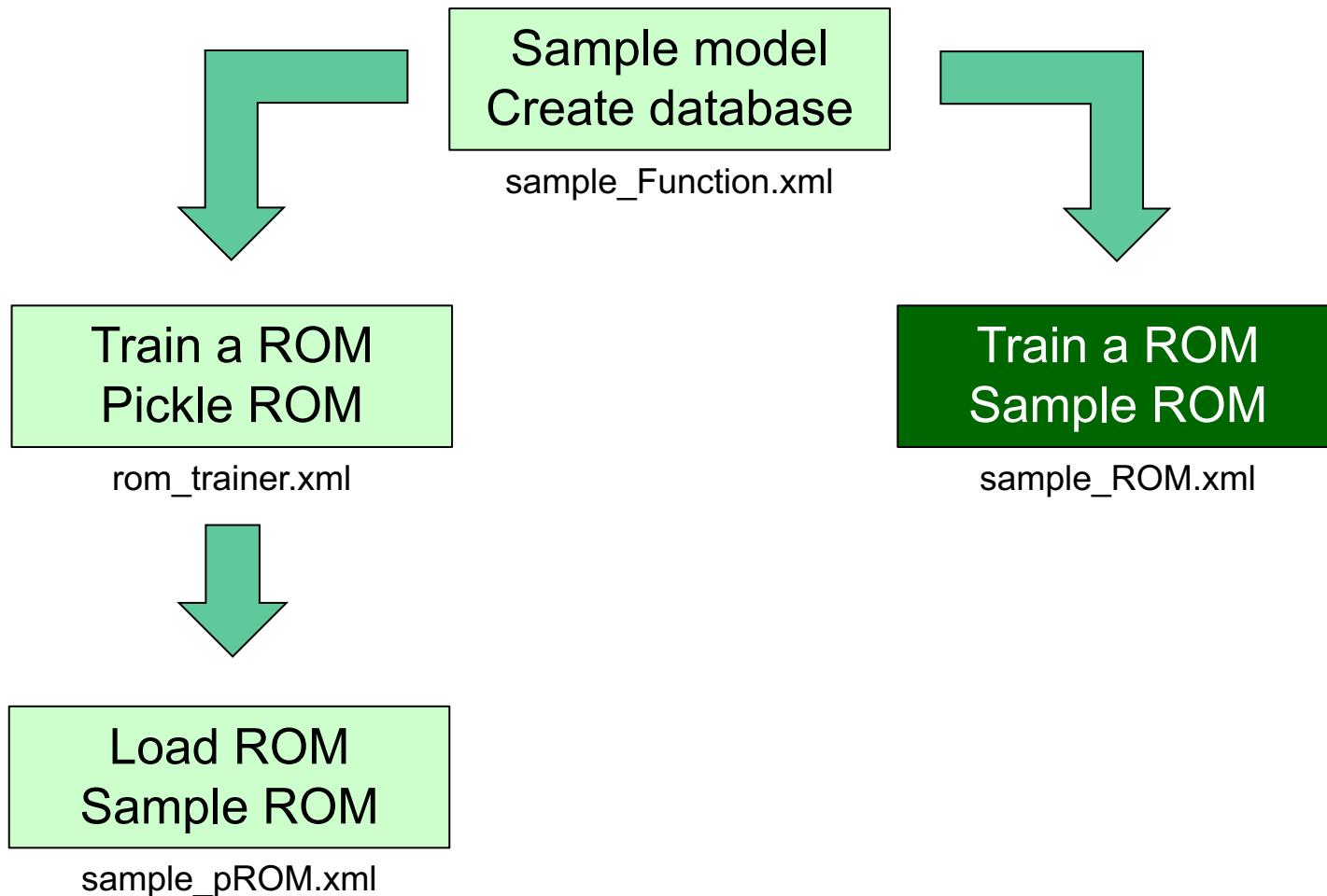
```

<Steps>
  <MultiRun name="FirstMRUn">
    <Input class="DataObjects" type="PointSet"      >inputPlaceHolder</Input>
    <Model  class="Models"      type="ExternalModel ">PythonModule</Model>
    <Sampler class="Samplers"   type="Grid"          >Grid_function</Sampler>
    <Output class="DataObjects" type="PointSet"      >outGRID</Output>
    <Output class="Databases"   type="HDF5"          >out_db</Output>
    <Output class="OutStreams"  type="Print"         >out_dump</Output>
    <Output class="OutStreams"  type="Plot"          >plotResponseFunction</Output>
  </MultiRun>
</Steps>

```



# Workflow



# Train and Sample a ROM

Distributions	Models	Samplers	Databases	DataObjects	Steps
---------------	--------	----------	-----------	-------------	-------

```
<Models>
  <ROM name="ROM4" subType="NDinvDistWeight">
    <Features>x1,x2,x3</Features>
    <Target>y4</Target>
    <p>3</p>
  </ROM>
</Models>
```

Multi-dimensional  
interpolator (CROW)

# *Train and Sample a ROM*

Distributions	Models	Samplers	Databases	DataObjects	Steps
---------------	--------	----------	-----------	-------------	-------

```

<Samplers>
  <Grid name="Grid_ROM">
    <variable name="x1">
      <distribution>normal_trunc</distribution>
      <grid type="value" construction="equal" steps="10">0.0 1.0</grid>
    </variable>
    <variable name="x2">
      <distribution>normal</distribution>
      <grid type="value" construction="equal" steps="10">1.5 2.5</grid>
    </variable>
    <variable name="x3">
      <distribution>uniform</distribution>
      <grid type="value" construction="equal" steps="10">1.0 4.0</grid>
    </variable>
  </Grid>
</Samplers>

```

Finer grid

# *Train and Sample a ROM*

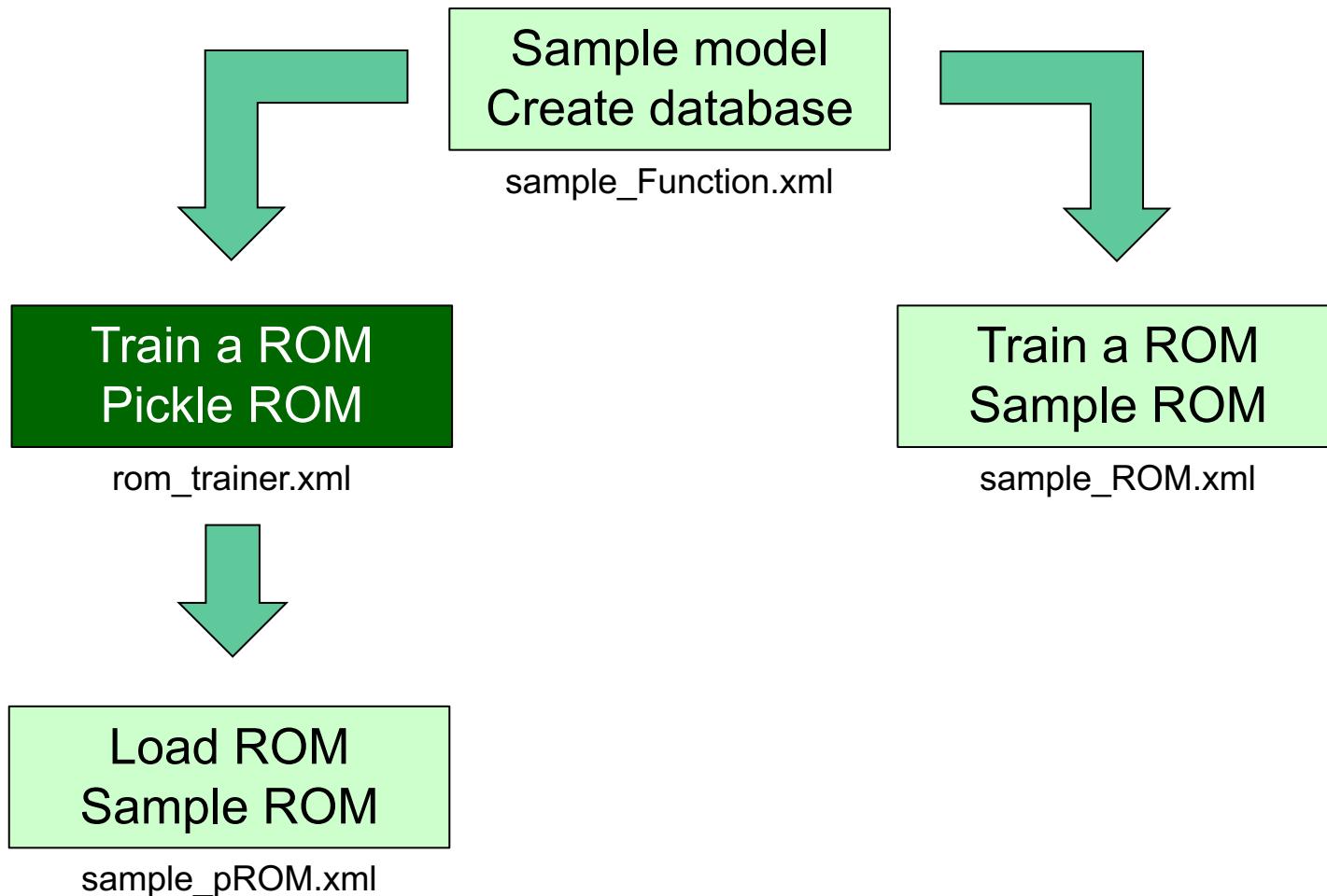
Distributions	Models	Samplers	Databases	DataObjects	Steps
---------------	--------	----------	-----------	-------------	-------

```

<Steps>
  <IOStep name="extract_data4">
    <Input class="Databases" type="HDF5"      >out_db</Input>
    <Output class="DataObjects" type="PointSet" >outGRID_y4</Output>
  </IOStep>
  <RomTrainer name="rom_trainer4">
    <Input class="DataObjects" type="PointSet" >outGRID_y4</Input>
    <Output class="Models"      type="ROM"       >ROM4</Output>
  </RomTrainer>
  <MultiRun name="RunRom4">
    <Input class="DataObjects" type="PointSet" >Data1</Input>
    <Model class="Models"      type="ROM"       >ROM4</Model>
    <Sampler class="Samplers"   type="Grid"      >Grid_ROM</Sampler>
    <Output class="DataObjects" type="PointSet" >outROM_y4</Output>
  </MultiRun>
  ...
</Steps>

```

# Workflow



# Train and Pickle a ROM

Models	Databases	DataObjects	Steps
--------	-----------	-------------	-------

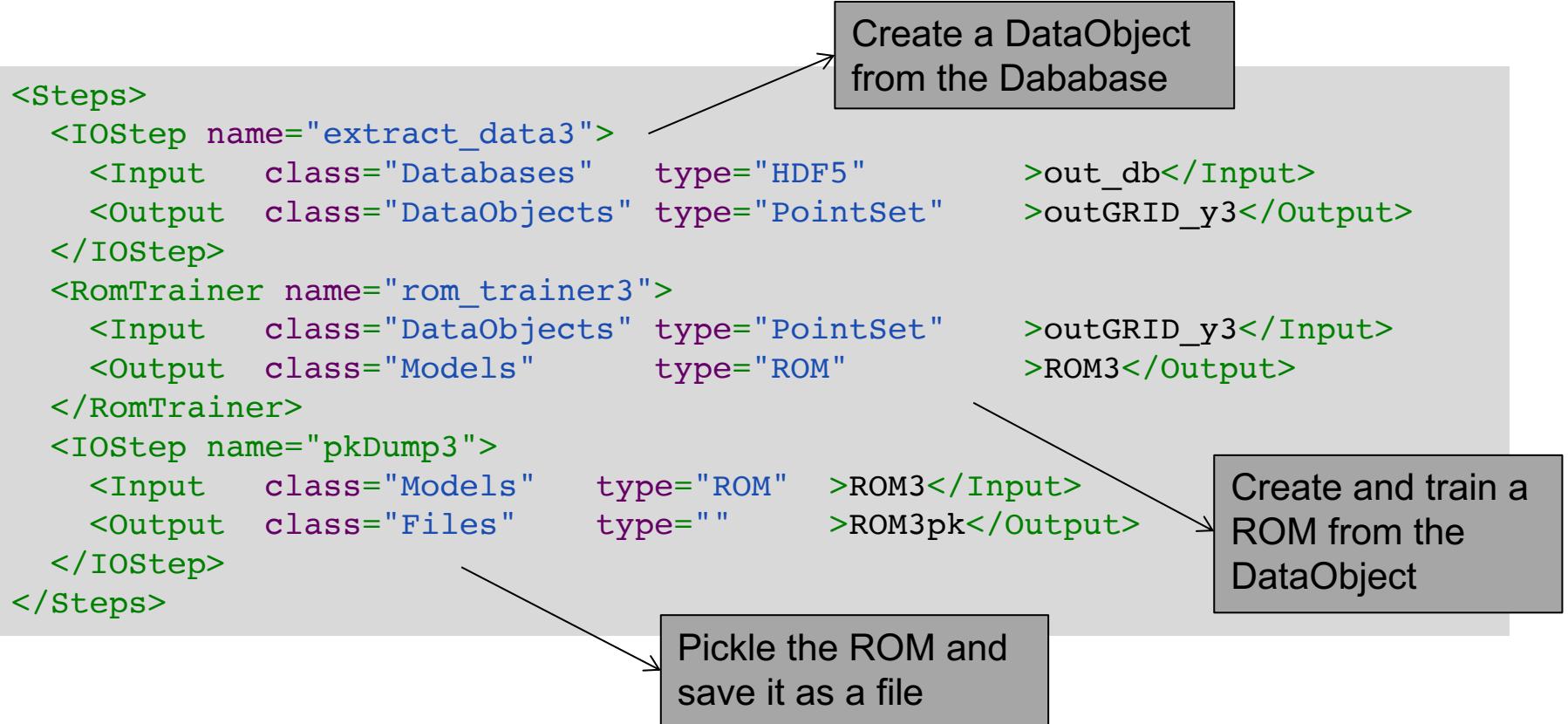
```
<Models>
    <ROM  name="ROM3" subType="SciKitLearn">
        <Features>x1,x2,x3</Features>
        <Target>y3</Target>
        <SKLtype>linear_model|LinearRegression</SKLtype>
        <fit_intercept>True</fit_intercept>
        <normalize>False</normalize>
    </ROM>
</Models>
```

Inputs / Output

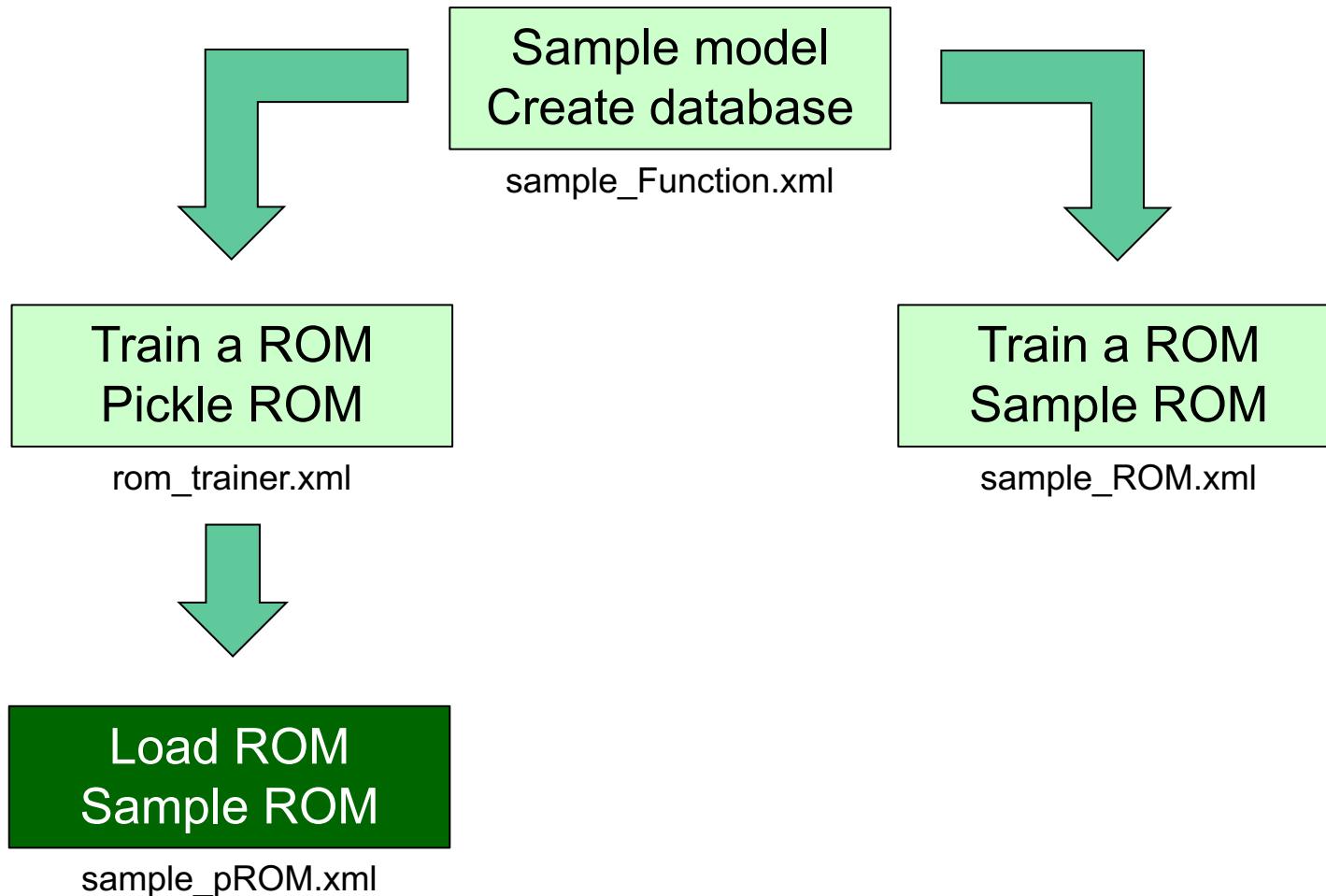
ROM type and parameters

# Train and Pickle a ROM

Models	Databases	DataObjects	Steps
--------	-----------	-------------	-------



# Workflow



# Load and Sample a Pickled ROM

Distributions	Models	Samplers	DataObjects	Steps
---------------	--------	----------	-------------	-------

```

<Models>
    <ROM  name="ROM3" subType="SciKitLearn">
        <Features>x1,x2,x3</Features>
        <Target>y3</Target>
        <SKLtype>linear_model|LinearRegression</SKLtype>
        <fit_intercept>True</fit_intercept>
        <normalize>False</normalize>
    </ROM>
</Models>

```

Inputs / Output

ROM type and parameters

# Load and Sample a Pickled ROM

Distributions

Models

Samplers

DataObjects

Steps

```

<Steps>
  ...
  <IOStep name="pk3Load">
    <Input class="Files" type="">ROM3pk</Input>
    <Output class="Models" type="ROM">pROM3</Output>
  </IOStep>
  <MultiRun name="RunPROM3">
    <Input class="DataObjects" type="PointSet" >Data1</Input>
    <Model class="Models" type="ROM" >pROM3</Model>
    <Sampler class="Samplers" type="Grid" >Grid_ROM</Sampler>
    <Output class="DataObjects" type="PointSet" >outPROM_y3</Output>
    <Output class="Databases" type="HDF5" >out_ROM3_db</Output>
  </MultiRun>
  ...
</Steps>
```

Load the pickled ROM