

# **Pipelines**



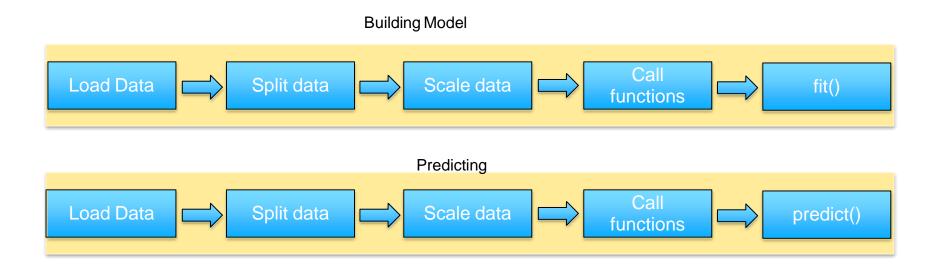
# What is a pipeline-

- 1. Almost always, we need to tie together many different processes that we use to prepare data for machine learning based model
- 2. It is paramount that the stage of transformation of data represented by these processes are standardized
- 3. Pipeline class of sklearn helps simplify the chaining of the transformation steps and the model
- 4. Pipeline, along with the GridsearchCV helps search over the hyperparameter space applicable at each stage



## **Pipelines**

- 1. Sequentially apply a list of transforms and a final estimator.
- 2. Intermediate steps of the pipeline must be 'transforms', that is, they must implement fit and transform methods.
- 3. The final estimator only needs to implement fit
- 4. Helps standardize the model project by enforcing consistency in building testing and production. Ref: https://scikit-learn.org/stable/modules/compose.html





## **Build a pipeline**

- 1. Import the pipeline class
  - a. from sklearn.pipeline import Pipeline
- 2. Instantiate the class into an object by listing out the transformation steps. In the following example, a scaling function is followed by the SVC algorithm
  - a. pipe = Pipeline([(" scaler", MinMaxScaler()), (" Ir", logisticregression())])
- 3. Call the fit() function on the pipeline object
  - a. pipe.fit( X\_train, y\_train)
- 4. Call the score() function on the pipeline object or predict() function
  - a. pipe.score( X\_test, y\_test)

In the step 2b, the pipeline object is created using a dictionary of key:value pairs. The key is specified in strings for e.g. "scaler" followed by the function to be called.

The key is the name given to a step.



## **Build a pipeline (Contd...)**

- 1. The pipeline object requires all the stages included to have a "transform()" function except for the last stage which is an estimator.
- 2. The transform step transforms the input data. The transformed output of a stage is the input to the next stage
- 3. During the call "pipeline.fit()", the pipeline calls the fit and transform functions on each step in sequence. For the last step, only the fit function is called
- While predicting using pipeline, similarly transform function in all the stages followed by a predict function in the last stage is performed
- 5. The pipeline object does not need to have a predict function. It only needs to have a fit function at least



# **Build a pipeline (Contd...)**

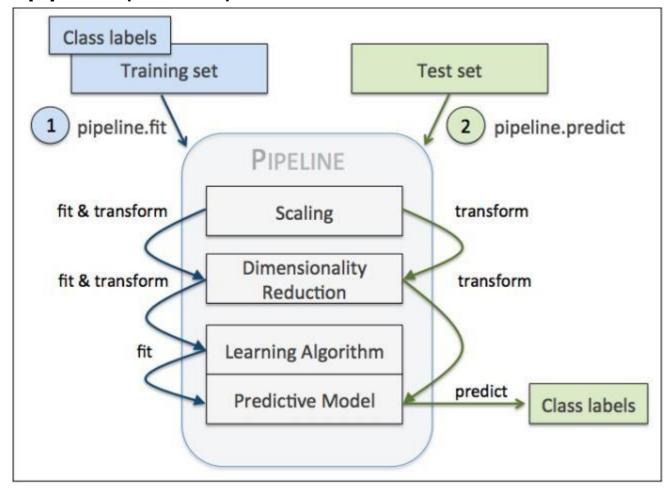


Image Source: Python Deeper Insights Into Machine Learning



### make\_pipeline

- Creating the pipeline could be cumbersome. Specifying a name to each stage may not be necessary
- 2. Alternatively there is a "make\_pipeline()" function that will create the pipeline and automatically name each step. We do not need to specify a name
  - a) from sklearn.pipeline import make\_pipeline
  - b) pipe = make\_pipeline( MinMaxScaler(), (SVC()))
  - c) print(" Pipeline steps:\ n{}". format( pipe.steps))
- 3. Note, we have not specified any name to the stages. The names will be automatically assigned and are usually lowercase of the class names



# **HyperParameter Tuning**



## **Hyper Parameters & Tuning**

- 1. Hyper parameters are like handles available to the modeler to control the behavior of the algorithm used for modeling
- Hyper parameters are supplied as arguments to the model algorithms while initializing them. For e.g. setting the criterion for decision tree building "dt\_model = DecisionTreeClassifier(criterion = 'entropy')"
- 3. To get a list of hyper parameters for a given algorithm, call the function get\_params()...for e.g. to get support vector classifier hyper parameters
  - from sklearn.svm import SVC
  - 2. svc=SVC()
  - svc.get\_params()
- 4. Hyper parameters are not learnt from the data as other model parameters are. For e.g. attribute coefficients in a linear model are learnt from data while cost of error is input as hyper parameter.



# **Hyper Parameters & Tuning**

- 5. Fine tuning the hyper parameters is done in a sequence of steps
  - 1. Selecting the appropriate model type (regressor or classifier such as sklearn.svm.SVC())
  - 2. Identify the corresponding parameter space
  - 3. Decide the method for searching or sampling parameterspace;
  - 4. Decide the cross-validation scheme to ensure model will generalize
  - 5. Decide a score function to use to evaluate the model
- 6. Two generic approaches to searching hyper parameter space include
  - 1. GridSearchCV which exhaustively considers all parameter combinations
  - 2. RandomizedSearchCV can sample a given number of candidates from a parameter space with a specified distribution.
- 7. While tuning hyper parameters, the data should have been split into three parts
  - Training, validation and testing to prevent data leak
- 8. The testing data should be separately transformed \* using the same functions that were used to transform the rest of the data for model building and hyper parameter tuning

<sup>\*</sup> Any transformation where rows influence each other. For e.g. using zscore. OneHotCode transformation does not come into this category. It can be done before splitting the data

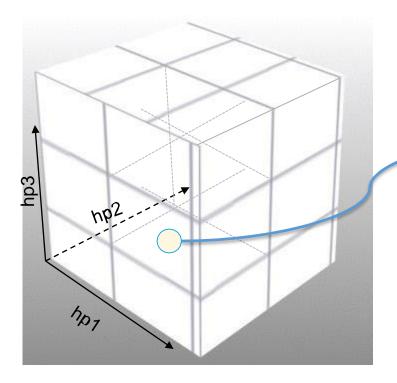
# Hyper Parameters & Tuning (GridsearchCV/ RandomizedSearchCv)

GridsearchCV -

- 1. Is at basic optimal hyperparameter tuning technique.
- 2. It builds a model for each permutation of all of the given hyperparameter values
- 3. Each such model is evaluated and ranked.
- 4. The combination of hyperparameter values that gives the best performing model is chosen
- 5. For every combination, cross validation is used and average score is calculated
- 6. This is an exhaustive sampling of the hyperparameter space and can be quite inefficient



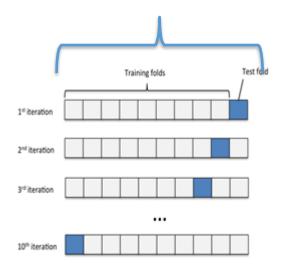
#### **GridSearchCV**



Hyper parameter space

One combination of hyper parameters used K times to train and test. The avg score of the K times is the score associated with this combination

This will repeat for all possible combinations i.e. all the cells in the space.





## Hyper Parameters & Tuning (GridsearchCV/ RandomizedSearchCv)

RandomizedSearchCV -

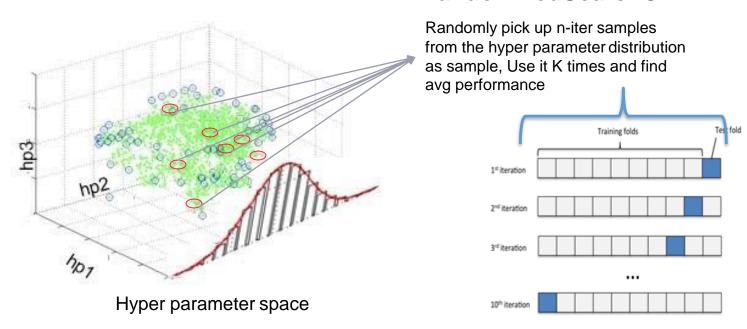
- 1. Random search differs from grid search. Instead of providing a discrete set of values to explore on each hyperparameter (parameter grid), we provide a statistical distribution.
- 2. Values for the different hyper parameters are picked up at random from this combine distribution
- The motivation to use random search in place of grid search is that for many cases, hyperparameters are not equally important.

A Gaussian process analysis of the function from hyper-parameters to validation set performance reveals that for most data sets only a few of the hyper-parameters really matter, but that different hyper-parameters are important on different data sets. This phenomenon makes grid search a poor choice for configuring algorithms for new data sets. - Bergstra, 2012

Picture by Bergstra, 2012



### RandomizedSearchCV



- 4. In contrast to GridSearchCV, not all combinations are evaluated. A fixed number of parameter settings is sampled from the specified distributions.
- 5. The number of parameter settings that are tried is given by n\_iter
- 6. <u>If all parameters are presented as a list, sampling without replacement is performed.</u> <u>If at least one parameter is given as a distribution, sampling with replacement is used</u>. It is highly recommended to use continuous distributions for continuous parameters
- 7. Randomsearch has higher chance of hitting the right combination than gridsearch.