

## **Feature Engineering**

*It is the process of creating, selecting, or transforming the input variables (features) used in an ML model to improve its performance.*

### **Aspects of Feature Engineering**

- *Creating Features*
- *Selecting features*
- *Transforming features*
- *Handling missing data*
- *Dealing with Outliers*
- *Encoding categorical data*
- *Feature Scaling*
- *Time and date feature engineering*

### **Techniques used in Feature Engineering**

- *Feature selection*
  - *Univariate feature selection*
  - *Recursive feature elimination*
  - *Feature importance from tree-based models*
  - *L1 regularization*
- *Feature Transformation*
  - *Scaling and normalization*
  - *One-hot and label encoding*
  - *Binning and discretization*
  - *Logarithmic, square root, or Box-Cox transformation*
  - *Handling text data (TF-IDF, Word Embeddings)*
- *Feature Creation*
  - *Polynomial features*
  - *Interaction terms*
  - *Aggregation and grouping of data*
- *Handling missing data*
  - *Imputation*
  - *Creating indicator variables for missing data*
- *Outliers*
  - *Identifying outliers through statistical methods*
  - *Winsorizing or clipping outliers*
  - *Transform data to get outliers*
- *Feature Scaling*
  - *Standardization*
  - *Min-max*
  - *Robust*
  - *Scaling with maximum absolute value*
  - *Log transformation*

- **Encoding**
  - One-Hot
  - Label
  - Mean Encoding(Target Encoding)
  - Binary Encoding
  - Frequency Encoding
- **Text data**
  - Tokenization and text preprocessing
  - TF-IDF (Term Frequency-Inverse Document Frequency)
  - Word embeddings (Word2Vec, GloVe, FastText)
  - Text sentiment analysis
  - N-grams and bag of word representations
- **Dimensionality Reduction**
  - PCA
  - ICA
  - LDA
  - t-SNE
  - Non-Negative Matrix Factorization (NMF)
- **AutoML libraries for feature engineering**
  - Featuretools
  - TSMF
  - Featurewiz
  - Pycaret

## **Feature Selection**

*It is the process of choosing a subset of the most relevant features (variables or columns) from the dataset*

### **Filter Methods**

- **Correlation-based feature selection:** Select features that have the strongest correlation with the target
- **Variance Threshold:** Remove features with low variance, as they might not provide much information
- **Information Gain:** Calculates the reduction in entropy from the transformation of a dataset
- **Chi-square Test:** Used for categorical features in a dataset. Calculate between each feature and the target and decide the features
- **Fisher's Score:** Fisher's score is one of the most widely used supervised feature selection methods.
- **Mean Absolute difference (MAD):** computes the absolute difference between the mean value. MAD is like a variance.

- *Dispersion Ratio: Another measure of dispersion applies the arithmetic mean and geometric mean.*

$$AM_i = \overline{X_i} = \frac{1}{n} \sum_{j=1}^n X_{ij} , \quad GM_i = \left( \prod_{j=1}^n X_{ij} \right)^{\frac{1}{n}} ,$$

$RM = AM / GM$ . This is called the dispersion measure and the higher the value, the higher the relevance to a feature

### **Wrapper Methods**

- *Forward Selection: Start with an empty set of features and iteratively add the most informative feature until a stopping criterion is met.*
- *Backward Elimination: Start with all features and iteratively remove the least informative feature until a stopping criterion is met.*
- *Exhaustive Feature Selection: This is a brute-force evaluation of each feature subset. It tries every combination of the variables and returns the best performance*
- *Recursive Feature Elimination (RFE): Repeatedly fit the model and remove the least important feature in each iteration until the desired number of features is reached.*

### **Embedded Methods**

- *L1 Regularization (Lasso): Features with coefficients that become zero are removed from the list*
- *Tree-based Feature Importance: Decision trees and tree-based models can provide feature importances that help identify the most relevant features.*

### **Sklearn Things**

- *SelectKBest: Can select the top k best features.*
- *SelectPercentile: Can select the features by the score above a certain percentage*

### **Univariate vs Multivariate selection**

- ❖ *Univariate considers each feature independently of the others. Statistical tests like ANOVA, chi-square test, and correlation coefficients are used in here.*
- ❖ *Multivariate feature selection methods take into account interactions or dependencies between multiple features simultaneously. These methods typically involve more complex techniques, such as recursive feature elimination, and wrapper methods.*

### **Mutual Information**

*It is a way to measure how much one random variable tells you about another. If the mutual information between 2 variables is high, it means changes in one variable give a lot of information about the changes in the other.*

$$I(X; Y) = \sum_{x \in \mathcal{X}} \sum_{y \in \mathcal{Y}} p(x, y) \log \frac{p(x, y)}{p(x)p(y)}$$

- ★  $I(X; Y)$ : Mutual information between random variables  $X$  and  $Y$
- ★  $\sum \sum$ : Represents a summation of all possible values of  $X$  and  $Y$
- ★  $p(x, y)$ : The joint probability of  $X$  and  $Y$  occurring together.
- ★  $p(x)$ : The probability of  $X$  occurring on its own
- ★  $p(y)$ : probability of  $Y$  occurring on its own
- ★  $\log 2$ : This is the base 2 logarithm

## Feature Transformation

### Scaling

Feature scaling is a method used to normalize the range of independent variables or features of data.

#### Types of Scaling and Transformation:

- **Min-max scaler:**
  - Scale data into typically  $[0, 1]$
  - For each data point,  $(X - X_{\min}) / (X_{\max} - X_{\min})$
- **Standard Scaler (Z-score scaling):**
  - Also known as standardization
  - Transform data to have a mean of 0 and std of 1
  - For each data point,  $(X - X_{\text{mean}}) / X_{\text{std}}$
- **Robust Scaler:**
  - Robust to outliers
  - For each data,  $(X - X_{\text{median}}) / \text{IQR}$
- **Max Absolute Transformation**
  - Scale the data in such a way that the maximum absolute value becomes 1
  - $X_{\text{scaled}} = X / X_{\text{max\_abs}}$
- **Power Transformation (Box-Cox)**
  - Applies to make the data more Gaussian-like
  - If  $\lambda == 0$ ,  $\log(\text{data})$
  - else,  $(\text{data}^{\lambda} - 1) / \lambda$
- **Quantile Transformation**
  - Maps the data to a uniform or normal distribution using quantiles
- **Log transformation**
  - Useful for data with exponential growth or skewed distributions.
  - $X_{\text{transformed}} = \log(X)$

## Encoding

- *Label Encoding: Assign a unique integer to each category*
- *One-Hot encoding: Creates a binary column for each category*
- *Binary Encoding: Converts categories into binary code*
- *Ordinal Encoding: Assign numerical values to categories based on the predefined order*
- *Frequency Encoding: Replace categories with their frequency in the dataset*
- *Target encoding: Replace categories with the mean of the target variables in that category*

## Binning

- *Equal Width Binning: divides the range of the data into equally sized bins*
- *Equal Frequency Binning (Quantile binning): Divide the data such that each bin contains the same number*
- *K-means clustering binning: Applies the K-means algorithm to group data points into bins*
- *Entropy-based binning: uses entropy to measure the impurity of bins and split the data to minimize entropy within each bin*
- *Custom Binning: Define specific boundaries according to the requirement*

## Feature Creation

- *Polynomial features: Take the powers of the existing features, such as squaring or cubing them*
- *Interaction Features: Combine 2 or more existing features. Eg:- divide 2 features*
- *Cross-product features: Take dot product of 2 features*
- *Frequency Features: Count the frequency of occurrence of certain values in the dataset*

## Dimensionality reduction

- *PCA (Principle component analysis)*

### Steps

- *Standardize the data*
  - *Calculate the covariance matrix. Each element  $(i, j)$  represents the covariance between feature  $i$  and feature  $j$*
  - *Calculate the Eigen vectors and Eigen values of the covariance matrix. These Eigenvalues represent the principal components, and the corresponding eigenvalues represent the variance explained by each component*
  - *Sort the eigenvalues in descending order to identify the principal ones.*
  - *Select the number of principal components.*
  - *Build the projection matrix by selecting the top-k eigenvalues*
  - *Get the dot product of the projected data and the standardized X*
- *LDA (Linear Discriminant Analysis)*

### Steps

- *Calculate the mean vector of each class*
- *Calculate 2 scatter matrices:*
  - *within the class: Sum of the covariance matrix of each class*

- *between class: Sum of outer products of the difference between class mean and the overall mean*
- *Find the eigenvectors and eigenvalues of the matrix  $S_w^{-1} * S_b$ , where  $S_w^{-1}$  is inverse of within-class scatter matrix and  $S_b$  is between class scatter matrix*
- *Sort eigenvalues in descending order*
- *Select the number of components*
- *Build the projection matrix by selecting the top-k eigenvalues*
- *Get the dot product of the projected data and the  $X$*
- **ICA (Independent Component Analysis)**
  - Used to separate independent sources from a mixed signal*
  - Steps**
    - *Subtract the mean from each feature to center the data*
    - *Whiten the data using PCA*
    - *Choose the number of Independent Components*
    - *Initialize random mixing matrices ( $W$  and  $A$ )*
    - *Update the Unmixing Matrix ( $W$ ): Use any optimization such as gradient descent*
      - $\text{Gradient} = (1 - 2 / (1 + \exp(-1 * X \cdot B)^T)) \cdot X$
      - $W -= \text{learning} * \text{Gradient}$
  - *Get the dot product of data and Unmixing Matrix ( $W$ )*
- **t-SNE (t- Distributed Stochastic Neighbor Embedding)**
  - Steps**
    - *Get the pairwise squared distances between the data points. Shape will  $(n \times n)$  where  $n$  is the number of data points*
    - *Calculate pairwise similarities*
    - *Calculate conditional probabilities of distances and perplexity*
    - *Initialize the low-dimensional embedding*
    - *Choose a perplexity value, a higher perplexity value considers more global relationships, with a lower focus on local relations.*
    - *Optimize to minimize the divergence between the conditional probabilities of high dimensional space and low dimensional space*
    - *Return the low-dimensional embedding*
- **NMF (Non-Negative Matrix Factorization)**
  - Steps**
    - *Choose the number of components*
    - *Initialize the matrix  $W$  with shape  $(n, k)$  and  $H$  with shape  $(k, m)$  where  $n$  is the number of data points,  $m$  is the number of features,  $k$  is the desired dimension*
    - *Define the number of iterations and learning rate*
    - *Update the  $W$  and  $H$  iteratively with some optimizers*
    - *Reconstruct data by taking the dot product of  $W$  and  $H$*