

Predicting Mobile Game Success

Milestone 1

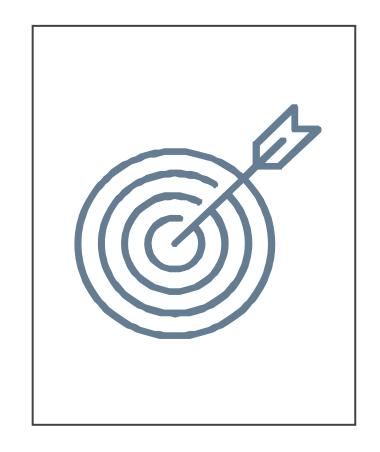
By: Team ID: CS 29



Objective

The objective of the Milestone 1 is to clean the data, applying preprocessing, feature selection, feature scaling, feature engineering, regression.

The objective of the Milestone 2 is to Classify games according to its Rate.



TEAM

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Feature Selection DATA **Feature CLEANING Scaling**

DATA CLEANING

- Dataset Size = 5214
- Number of duplicates = 43
- Number of Nan values = 5799
- URL, ID, Name, Icon URL, Description (All these Columns are Unique)
- Subtitle Column Contain almost Nan Values "3749."
- In-app Purchases Contains 2039 Nan Values.
- Languages Column Contains 11 Nan Values.

	Name	dtypes	Missing	Uniques
0	URL	object	0	5171
1	ID	int64	0	5171
2	Name	object	0	5171
3	Subtitle	object	3749	1399
4	Icon URL	object	0	5171
5	User Rating Count	int64	0	1410
6	Price	float64	0	17
7	In-app Purchases	object	2039	2052
8	Description	object	0	5099
9	Developer	object	0	3084
10	Age Rating	object	0	4
11	Languages	object	11	580
12	Size	int64	0	5081
13	Primary Genre	object	0	19
14	Genres	object	0	507
15	Original Release Date	object	0	2281
16	Current Version Release Date	object	0	1976
17	Average User Rating	float64	0	9

Check duplicates, then remove them.

```
dataset.duplicated().sum() # Duplicates = 43
# Drop duplicate rows
dataset = dataset.drop_duplicates() # Duplicates = 0
```

• Check missing Values, then replace them with Mean or Mode or Zero.

URL	0
ID	0
Name	0
Subtitle	3717
Icon URL	0
User Rating Count	0
Price	0
In-app Purchases	2025
Description	0
Developer	0
Age Rating	0
Languages	11
Size	0
Primary Genre	0
Genres	0
Original Release Date	0
Current Version Release Date	0
Average User Rating	0

each cell in the 'In-app Purchases' column contains prices of available in-app purchases, First, we sum the prices listed in each cell of the "In-app Purchases" column and calculate the mean value of the column. Then, we replace any missing values in the column with this mean value.

dataset['In-app Purchases'].isnull().sum() missing values = 2025

Before	After
In-app Purchases	In-app Purchases
29.99, 19.99,	307.840000
9.99, 29.99, 29.99, 8.99, 4.99,	61.674269
NaN	61.674269
NaN	61.674269
IVAIV	61.674269
NaN	
NaN	53.860000
1.777	60.920000
1.99, 1.99, 1.99, 1.99, 1.99, 1.99, 2.99, 7.99	61.674269
2.99, 4.99, 1.99, 2.99, 5.99, 9.99,	121.900000
11.99, 19.99	88.930000

each cell in the 'Languages' column contains a lot of languages that available in the game First, we separate each language in each cell and calculate the mode value of all languages in the column. Then, we replace any missing values in the column with this mode value which it is English

dataset['Languages'].isnull().sum()

missing values = 11

Before	After
Languages	Languages
EN, FR, DE, JA, KO, ZH, ES, TH, ZH, VI	EN, FR, DE, JA, KO, ZH, ES, TH, ZH, VI
EN	EN
EN, ZH	EN, ZH
NaN	EN
NaN	EN

	Before	After
each cell in the 'Age Rating' column contains either 4+, 9+, 12+ or 17+.	Age Rating	Min_Age
We perform the following steps to make to usable:	12+	12
 we remove the positive sign(+) next to each age. 	12+	12
 we change data type of column from 'object' to 'int'. 	4+	4
 we redefined column name from age rating to Min Age for better usages 		
	9+	9
	12+	12

Before

	URL	Name	Subtitle	Icon URL	Description
0	https://apps.apple.com/us/app/heir-of-light/id	HEIR OF LIGHT	Dark Fantasy RPG	https://is3-ssl.mzstatic.com/image/thumb/Purpl	A Dark Fantasy, Collectible RPG\n\nDarkness ha
1	https://apps.apple.com/us/app/endgame- eurasia/	Endgame:Eurasia	NaN	https://is4- ssl.mzstatic.com/image/thumb/Purpl	"This interactive experience is an exploration
2	https://apps.apple.com/us/app/free-solitaire/i	Free Solitaire+	NaN	https://is5-ssl.mzstatic.com/image/thumb/Purpl	Same Solitaire game with classic Solitaire run
3	https://apps.apple.com/us/app/draft-trainer/id	Draft Trainer	NaN	https://is1- ssl.mzstatic.com/image/thumb/Purpl	** Discounted for a limited time **\n\nEver wo
4	https://apps.apple.com/us/app/rogue-knight-inf	Rogue Knight: Infested Lands	Tactical roguelike w/ stealth	https://is2-ssl.mzstatic.com/image/thumb/Purpl	Fight or sneak your way through hordes of mons

We convert the categorical columns, namely, "URL ", "Name ", "Subtitle ", "Icon URL ", and "Description " from strings to numerical values and this by using the Label Encoder function from the Sklearn Package.

After

	URL	Name	Subtitle	Icon URL	Description
0	2336	2379	451	2526	2131
1	1710	1841	1394	3359	1581
2	2010	2123	1394	4863	4082
3	1582	1712	1394	491	1941
4	3839	3882	1154	1604	3053
777	***		***	***	•••
5209	3568	3618	316	1288	378
5210	2610	85	14	348	935
5211	4713	4716	1394	433	1636
5212	2549	2652	1394	4252	584
5213	3401	3468	1394	768	4912

Before

5210

5211

5212

AFEEL, Inc.

Stasis Software LLC

ZEN Studios Ltd.

Supervillain Studios

	Developer	Languages	Primary Genre	Genres
0	GAMEVIL Inc.	EN, FR, DE, JA, KO, ZH, ES, TH, ZH, VI	Games	Games, Role Playing, Strategy
1	Auroch Digital Ltd	EN	Games	Games, Simulation, Strategy, News
2	Chen Zhong Yuan	EN, ZH	Games	Games, Strategy, Entertainment, Card
3	GG Wizards, LLC	EN	Games	Games, Utilities, Card, Strategy
4	Luis Regueira	EN	Games	Games, Role Playing, Strategy
***	***			
5209	Ndemic Creations	EN, FR, DE, IT, JA, KO, NB, PL, PT, RU, ZH, ES	Games	Games, Strategy, Simulation

EN, JA, KO

EN

EN

FN

Games

Utilities

Games

Games

Games, Simulation, Strategy, Entertainment

Utilities, Games, Board, Strategy

Games, Card, Entertainment, Strategy

Games, Strategy, Role Playing

Convert Categorical Data to numeric based on Label (Average User Rating)

" Developer ", " Languages ", " Primary Genre " and "Genres ",

We Give weights to categorial strings based on label column.

```
dataset['Developer'] = dataset.groupby('Developer')['Average User Rating'].transform(lambda x: x.mean())
dataset['Languages'] = dataset.groupby('Languages')['Average User Rating'].transform(lambda x: x.mean())
dataset['Primary Genre'] = dataset.groupby('Primary Genre')['Average User Rating'].transform(lambda x: x.mean())
dataset['Genres'] = dataset.groupby('Genres')['Average User Rating'].transform(lambda x: x.mean())
```

After

	Developer	Languages	Primary Genre	Genres
0	4.250	4.000000	4.038315	4.116667
1	3.250	4.014370	4.038315	4.000000
2	4.000	4.056122	4.038315	4.096774
3	3.500	4.014370	4.038315	3.250000
4	4.500	4.014370	4.038315	4.116667
	***	***	***	***
5209	4.500	4.500000	4.038315	3.947802
5210	4.125	4.195652	4.038315	3.853448
5211	5.000	4.014370	4.016129	5.000000
5212	4.250	4.014370	4.038315	4.182203
5213	3.500	4.014370	4.038315	3.865385

Before	Original Release Date	Current Version Release Date	After	Days Since Release
	2018-03-06	2019-07-31		512
	2013-03-21	2017-06-28		1560
	2013-04-04	2015-04-21		747
	2011-05-26	2019-07-23		2980
	2017-05-19	2019-02-06		628
	2012-05-26	2019-02-08		2449
	2015-01-11	2018-04-16		1191
	2012-08-16	2017-02-21		1650
	2016-06-08	2017-01-30		236
	2012-09-10	2015-03-14		915

We calculate the duration between the "Original Release Date" and the "Current Version Release Date", and this duration is added in a new Column called "Days Since Release", and the "Original Release Date", "Current Version Release Date" columns are removed.

```
dataset[['URL','ID','Subtitle','Icon URL','Name','Description']].nunique()
                                                          URL
                                                                           5171
                                                           ID
                                                                           5171
                                                           Subtitle
                                                                           1400
                                                          Icon URL
                                                                           5171
                                                          Name
                                                                           5171
                                                          Description
                                                                           5099
#We deleted all theses columns because they are Unique (Non-Meaningful)
#Subtitle Column almost all cells are Nan values
#So we will drop this Columns
Columns To Deleted = ['URL', 'ID', 'Subtitle', 'Icon URL', 'Name', 'Description']
dataset = dataset.drop(columns = Columns To Deleted)
```

Selection Feature

Correlation Figure



- 0.8

- 0.6

-04

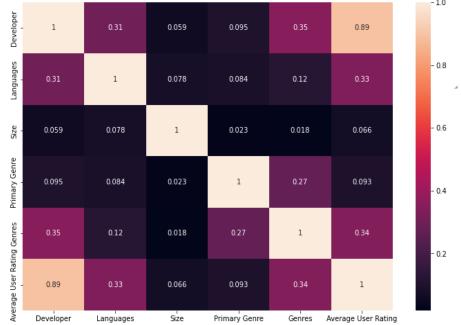
- 0.2

- 0.0

Selection Feature

```
Game_corr = dataset.corr()
# Top 5% Correlation training features with the Value
Common_features = Game_corr.index[abs(Game_corr['Average User Rating']) > 0.05]
# Correlation plot
plt.subplots(figsize=(12, 8))
top_corr = dataset[Common_features].corr()
sns.heatmap(top_corr, annot=True)
plt.show()
Top = Common_features.delete(-1)
data_input = data_input[Top]
```





Feature Scaling using (StandardScaler)

We Scale all Features between range [-1, 1]

```
# Initialize the scaler
scaler = StandardScaler()
# Fit the scaler to the training data
scaler.fit(X train)
# Scale the training, validation, and test data
X train scaled = scaler.transform(X train)
X val scaled = scaler.transform(X val)
X test scaled = scaler.transform(X test)
# Convert the scaled data to a dataframe
X train = pd.DataFrame(data=X train scaled, columns=X train.columns)
X val = pd.DataFrame(data=X val scaled, columns=X val.columns)
X test = pd.DataFrame(data=X test scaled, columns=X test.columns)
```

						Developer	Languages	Size	Primary Genre	Genres
					1	4.000000	4.000000	19414016	4.038315	4.223776
						4.500000	4.014370	131017728	4.038315	4.017442
						4.666667	4.450000	83177472	4.038315	4.150000
						4.250000	4.014370	59981824	4.038315	3.994444
		After				4.909091	4.785714	15915008	4.038315	3.994444
						***	***	***	***	
Developer	Languages	Size	Primary Genre	Genres	8	4.300000	4.014370	57909248	4.038315	4.203704
-0.049922	-0.157089	-0.455459	0.058217	0.769608	03	5.000000	4.014370	41652224	4.038315	4.320000
0.689308	-0.100639	-0.046105	0.058217	-0.075257	32	4.500000	4.500000	113888256	4.038315	3.926829
0.935718	1.610713	-0.221579	0.058217	0.467521		4.750000	4.014370	6282240	4.038315	3.750000
0.319693	-0.100639	-0.306659	0.058217	-0.169423		4.500000	4.550000	16797696	4.038315	4.380000
1.294132	2.929549	-0.468293	0.058217	-0.169423						
	***		***	***				Bef	ore	
0.393616	-0.100639	-0.314261	0.058217	0.687418						
1.428538	-0.100639	-0.373891	0.058217	1.163609						
0.689308	1.807135	-0.108935	0.058217	-0.446282						
1.058923	-0.100639	-0.503625	0.058217	-1.170334						
0.689308	2.003558	-0.465055	0.058217	1.409287						

1-Linear Regression Model

Linear regression is a statistical method that is used to model the relationship between a dependent variable (also called the response variable or outcome variable) and one or more independent variables (also called predictor variables or explanatory variables). The goal of linear regression is to find the best linear relationship between the variables.

In simple linear regression, there is only one independent variable and one dependent variable, and the relationship between them is modeled as a straight line. The equation for a simple linear regression model is: $y(pred) = theta_0 + theta_1*x + c$

where y is the dependent variable, x is the independent variable, theta₀ is the y-intercept (the value of y when x is zero), theta₁ is the slope (the change in y for a one-unit increase in x), and c is the error term (which represents the part of the variation in y that cannot be explained by the model).

Features which used in ALL Regression Models are (Developer, Languages, Size, Primary Genre, Genres) The following picture show the MSE and R2_Score for Test sets Using linear Regression.

```
Linear Regression:
Mean Square Error on test set: 0.0996
R2 Score on test set: 0.8100
```

2-Random Forest Model

Random Forest is a machine learning algorithm that is used for classification, regression, and other tasks that involve predicting a target variable based on one or more input variables. It is an ensemble learning method that combines multiple decision trees to produce a more accurate and robust model.

Random Forest is based on the idea of bagging, which stands for Bootstrap Aggregating. In bagging, multiple models are trained on different bootstrap samples of the training data, and the predictions from these models are aggregated to produce a final prediction. This helps to reduce the variance of the model and improve its generalization performance.

Random Forest builds on the idea of bagging by adding randomness to the model building process. Instead of using all the features to split each node of the decision tree, it selects a random subset of the features for each split. This helps to decorrelate the trees and reduce overfitting. In addition, Random Forest also uses random sampling of the training data to create each bootstrap sample, further increasing the diversity of the trees.

The following picture show the MSE and R2_Score for Test sets Using Random Forest

Random Forest Regression:
Mean Square Error on test set: 0.1024
R2 Score on test set: 0.8046

3-Polynomial Regression Model

Polynomial regression is a type of regression analysis where the relationship between the independent variable (x) and dependent variable (y) is modeled as an nth degree polynomial. It is used when the linear relationship between the variables does not fit the data well and a more complex relationship is required.

To perform polynomial regression, we first select the degree of the polynomial that best fits the data. We can then use the least squares method to estimate the coefficients of the polynomial equation that best fits the data. The coefficients are estimated by minimizing the sum of the squared errors between the predicted values of y and the actual values of y.

The equation for a polynomial regression model is: $y = b0 + b1x + b2x^2 + b3x^3 + ... + bnx^n + \epsilon$

where y is the dependent variable, x is the independent variable, b0, b1, b2, b3, ..., bn are the coefficients, n is the degree of the polynomial, and ε is the error term.

The following picture show the MSE and R2_score for Test sets Using Polynomial Regression

Polynomial Regression: MSE on test set: 0.0996 R2 Score on test set: 0.8099

1- Logistic Regression

```
log C values = [0.1, 1, 10]
solvers = ['sag', 'liblinear', 'lbfgs']
logreg models = {}
print("-----")
for sol in solvers:
   for C in log C values:
       start = timeit.default timer()
       logreg = LogisticRegression(solver=sol, C=C)
       logreg.fit(X_train, y_train)
       end = timeit.default timer()
       train Time.append(end-start)
       y pred = logreg.predict(X_test)
       logaccuracy = accuracy score(y test, y pred)
       train accurices.append(logaccuracy)
       logmodel name = f'{sol} {C}'
       logreg_models[logmodel_name] = (logreg, logaccuracy)
       print(f"Solver: {sol}, C: {C}")
       print("-----")
       print("Accuracy score:", logaccuracy)
```

```
------Logistic Regression------
Solver: sag, C: 0.1
Accuracy score: 0.8830917874396135
Solver: sag, C: 1
Accuracy score: 0.8917874396135266
Solver: sag, C: 10
Accuracy score: 0.893719806763285
Solver: liblinear, C: 0.1
Accuracy score: 0.8338164251207729
Solver: liblinear, C: 1
Accuracy score: 0.8396135265700483
Solver: liblinear, C: 10
Accuracy score: 0.8917874396135266
Solver: lbfgs, C: 0.1
Accuracy score: 0.8830917874396135
Solver: lbfgs, C: 1
Accuracy score: 0.8917874396135266
Solver: lbfgs, C: 10
Accuracy score: 0.893719806763285
```

3-SVM

```
kernel_values = ['linear', 'rbf', 'sigmoid']
SVM_C_values = [0.1, 1, 10]
svm models = {}
print("----")
for kernel in kernel values:
   for C in SVM C values:
       start = timeit.default timer()
       svm = SVC(kernel=kernel, C=C)
       svm.fit(X train, y train)
       end = timeit.default timer()
       train Time.append(end-start)
       y_pred = svm.predict(X_test)
       symaccuracy = accuracy score(y test, y pred)
       train accurices.append(symaccuracy)
       svmmodel_name = f'{kernel}_{C}'
       svm models[svmmodel name] = (svm, svmaccuracy)
       print(f"Kernel: {kernel}, C: {C}")
       print("----")
       print("Accuracy score:", symaccuracy)
```

```
-----SVM------
Kernel: linear, C: 0.1
Kernel: linear, C: 1
Accuracy score: 0.8966183574879227
Kernel: linear, C: 10
Accuracy score: 0.8966183574879227
Kernel: rbf, C: 0.1
Accuracy score: 0.8946859903381642
Kernel: rbf, C: 1
Accuracy score: 0.8966183574879227
Kernel: rbf, C: 10
Accuracy score: 0.8917874396135266
Kernel: sigmoid, C: 0.1
Accuracy score: 0.49468599033816424
Kernel: sigmoid, C: 1
Accuracy score: 0.4144927536231884
Kernel: sigmoid, C: 10
Accuracy score: 0.088888888888888888
```

3-Naive Bayes

```
var smoothing values = [1e-9, 1e-7, 1e-5]
priors = [None, [0.3, 0.3, 0.4], [0.3, 0.3, 0.4]]
NB_models = \{\}
print("-----")
for prior in priors:
    for var smoothing in var smoothing values:
       nb = GaussianNB(var_smoothing=var_smoothing, priors=prior)
       # Fit the model on the training data
       nb.fit(X train, y train)
       y_pred = nb.predict(X_test)
       # Calculate accuracy score and mean squared error
       NBaccuracy = accuracy score(y test, y pred)
       NBmodel name = f'{prior} {var smoothing}'
       NB_models[NBmodel_name] = (nb, NBaccuracy)
       print(f"Priors: {prior}, Var smoothing: {var smoothing}")
       print("----")
       print("Accuracy score:", NBaccuracy)
```

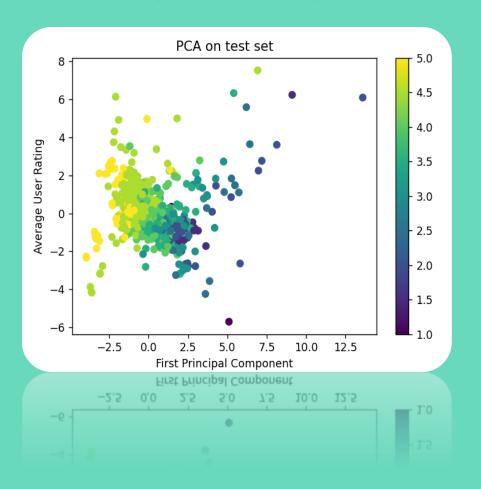
```
-----Baive Bayes-----
Priors: None, Var smoothing: 1e-09
Accuracy score: 0.8714975845410629
Priors: None, Var smoothing: 1e-07
Accuracy score: 0.8714975845410629
Priors: None, Var smoothing: 1e-05
Accuracy score: 0.8714975845410629
Priors: [0.3, 0.3, 0.4], Var smoothing: 1e-09
Accuracy score: 0.8705314009661835
Priors: [0.3, 0.3, 0.4], Var smoothing: 1e-07
Accuracy score: 0.8705314009661835
Priors: [0.3, 0.3, 0.4], Var smoothing: 1e-05
Accuracy score: 0.8705314009661835
Priors: [0.3, 0.3, 0.4], Var smoothing: 1e-09
Accuracy score: 0.8705314009661835
Priors: [0.3, 0.3, 0.4], Var smoothing: 1e-07
Accuracy score: 0.8705314009661835
Priors: [0.3, 0.3, 0.4], Var smoothing: 1e-05
Accuracy score: 0.8705314009661835
```

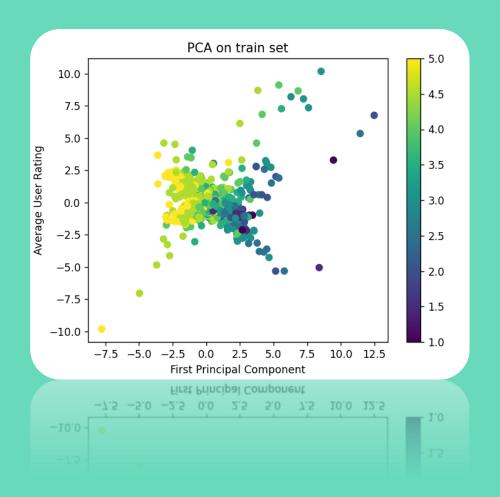
4- Decision Tree

```
max depth values = [2, 5, 10]
min samples split values = [2, 5, 10]
criterion values = ['gini', 'entropy']
dt models = {}
print("-----")
for max depth in max depth values:
    for min_samples_split in min_samples_split_values:
       for criterion in criterion values:
          dt = DecisionTreeClassifier(max depth=max depth, min samples split=min samples split, criterion=criterion)
          dt.fit(X train, y train)
          y pred = dt.predict(X test)
          dt accuracy = accuracy score(y test, y pred)
          dt model name = f'{max depth} {min samples split} {criterion}'
          dt models[dt model name] = (dt, dt accuracy)
          print(f"Max depth: {max_depth}, Min samples split: {min_samples_split}, Criterion: {criterion}")
          print("----")
          print("Accuracy score:", dt_accuracy)
```

```
-----Decision Tree-----
Max depth: 2. Min samples split: 2. Criterion: gini
Accuracy score: 0.8792270531400966
Max depth: 2, Min samples split: 2, Criterion: entropy
Accuracy score: 0.8743961352657005
Max depth: 2. Min samples split: 5. Criterion: gini
Accuracy score: 0.8792270531400966
Max depth: 2, Min samples split: 5, Criterion: entropy
Accuracy score: 0.8743961352657005
Max depth: 2, Min samples split: 10, Criterion: gini
Accuracy score: 0.8792270531400966
Max depth: 2, Min samples split: 10, Criterion: entropy
Accuracy score: 0.8743961352657005
Max depth: 5. Min samples split: 2. Criterion: gini
Accuracy score: 0.8792270531400966
Max depth: 5, Min samples split: 2, Criterion: entropy
Accuracy score: 0.8724637681159421
Max depth: 5, Min samples split: 5, Criterion: gini
Accuracy score: 0.8792270531400966
Max depth: 5, Min samples split: 5, Criterion: entropy
Accuracy score: 0.8724637681159421
Max depth: 5, Min samples split: 10, Criterion: gini
Accuracy score: 0.8792270531400966
Max depth: 5, Min samples split: 10, Criterion: entropy
```

The Scatter For (Test set, Train set)

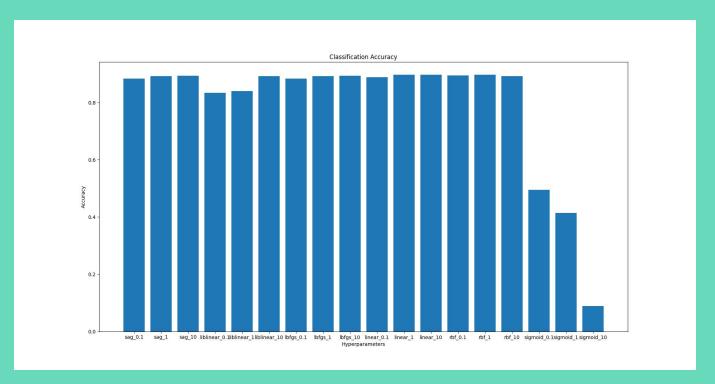




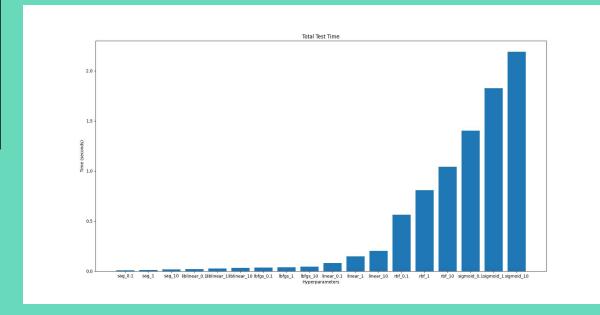
- Split The data into (0.20) For Test sets (0.20) For Validate sets and (0.60) For Training set.
- Apply (PCA) Technique on both (Test set, Train set) on features (Developer, Languages, Size, Primary genre, Genres)

Which is used for analyzing datasets containing a high number of dimensions (Features) per observation, Increasing the interpretability of data while preserving the maximum amount of information, and enabling The visualization of multidimensional data.

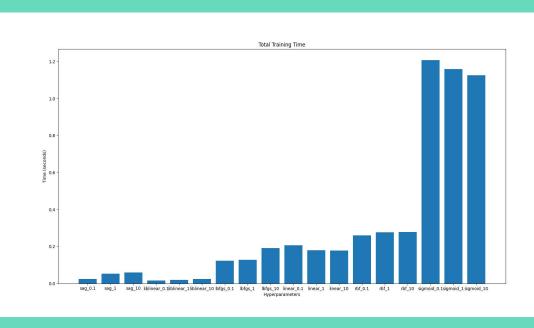
Classification Accuracy



Total Test Time



Total Training Time



- Delete all duplicate rows
- Check missing Values, then replace them with mean or mode or zero
- remove the positive sign (+) next to each age in (Age column) and change its name to (Min Age)
- convert the categorical columns namely (URL, Name, Sub title, Icon URL, Description) from Strings to numerical using the label encoder based on Label (Average User Rating)
- calculate the duration between the "The original Release Date" and the current "Current Version Release Date", and this duration is added in a new Column called" Days Since Release" and the "Original Release Date "and remove each Columns (Original Release Date, Current Version Release Date)
- Delete ALL Unique columns (URL, ID, Sub title, Icon URL, Name, Description)
- Apply Feature Scaling in All features between Range [-1,1]
- Apply Three Regression Models (Linear Regression, Random Forest Regression, Polynomial Regression)
- Visualizing The Scatter for (Test set, Train set) Using PCA technique

Our intuition was that Polynomial linear regression would be better, but it was disproved by the error of the output.

Multivariable linear regression has proven to be better than Polynomial linear regression according to the features that we have selected.

GAME OVER