```
In [1]: #Processamento e Análise em Big Data
        #Tarefa 05
        #Grupo 03
        #20/04/2021
In [2]: import pandas as pd
        from pyspark.sql import functions as F
        from pyspark.ml.feature import PCA
        from pyspark.ml.feature import StringIndexer, VectorAssembler
        from pyspark.ml import Pipeline
        from pyspark.sql.functions import col
```

Análise Descritiva

```
In [56]: data_ = spark.read.csv('pima_indian_diabetes.csv', header=True)
 In [4]: data_.limit(5).toPandas()
Out[4]:
              n_pregnant plasma_glucose_concentration blood_pressure triceps_skinfold_thickness 2hours
           0
                       6
                                                 148
                                                                 72
                                                                                          35
                                                  85
                                                                                          29
           1
                       1
                                                                 66
           2
                       8
                                                                                           0
                                                 183
                                                                 64
                       1
                                                  89
                                                                 66
                                                                                          23
                       0
                                                                 40
                                                                                          35
                                                 137
```

Checando o Schema do Dataframe

```
In [5]: | data_.printSchema()
        root
         |-- n_pregnant: string (nullable = true)
         |-- plasma_glucose_concentration: string (nullable = true)
         |-- blood pressure: string (nullable = true)
         |-- triceps skinfold thickness: string (nullable = true)
         |-- 2hours_serum_insulin: string (nullable = true)
         |-- body_mass_index: string (nullable = true)
         |-- diabtes_pedigree_function: string (nullable = true)
         |-- age: string (nullable = true)
         |-- class: string (nullable = true)
```

Vamos converter os valores string para int e float

```
In [57]: | num_cols = ['n_pregnant', 'class', 'age']
         for col in num_cols:
             data = data .withColumn(col, data [col].cast('int'))
         num_cols = ['plasma_glucose_concentration', 'blood_pressure','triceps_skinfold
         _thickness',
                      '2hours_serum_insulin','body_mass_index', 'diabtes_pedigree_functi
         on' 1
         for col in num cols:
             data = data .withColumn(col, data [col].cast('float'))
         data_.printSchema()
         root
          |-- n_pregnant: integer (nullable = true)
          |-- plasma glucose concentration: float (nullable = true)
          |-- blood_pressure: float (nullable = true)
          |-- triceps skinfold thickness: float (nullable = true)
          |-- 2hours serum insulin: float (nullable = true)
          |-- body mass index: float (nullable = true)
          |-- diabtes_pedigree_function: float (nullable = true)
          |-- age: integer (nullable = true)
          |-- class: integer (nullable = true)
```

copiando Dataframe

```
In [7]: | schema = data .schema
        X_pd = data_.toPandas()
        data_model = spark.createDataFrame(X_pd,schema=schema)
        del X pd
        data model.printSchema()
        root
         |-- n pregnant: integer (nullable = true)
         |-- plasma glucose concentration: float (nullable = true)
         |-- blood pressure: float (nullable = true)
         |-- triceps_skinfold_thickness: float (nullable = true)
         |-- 2hours serum insulin: float (nullable = true)
         |-- body_mass_index: float (nullable = true)
         |-- diabtes pedigree function: float (nullable = true)
         |-- age: integer (nullable = true)
         |-- class: integer (nullable = true)
```

Colunas presentes no Dataframe

```
In [8]: for column in data_.limit(5).toPandas().columns:
            print(column)
```

n pregnant plasma_glucose_concentration blood pressure triceps_skinfold_thickness 2hours_serum_insulin body mass index diabtes_pedigree_function age class

n pregnant - Número de vezes que engravidou

plasma_glucose_concentration - Concentração de glicose plasmática a 2 horas em um teste oral de tolerância à glicose

blood pressure - Pressão arterial diastólica (mm Hg)

triceps_skinfold_thickness - Espessura da dobra da pele do tríceps (mm)

2hours_serum_insulin - Insulina sérica de 2 horas (mu U / ml)

body mass index - Índice de massa corporal (peso em kg / (altura em m) ^ 2)

diabtes_pedigree_function - valores que medem a tendência ao desenvolvimento de diabetes com base nas relações genéticas do indivíduo

age - Idade

class - Presença ou não de diabetes

Vamos ver as estatísticas das variáveis

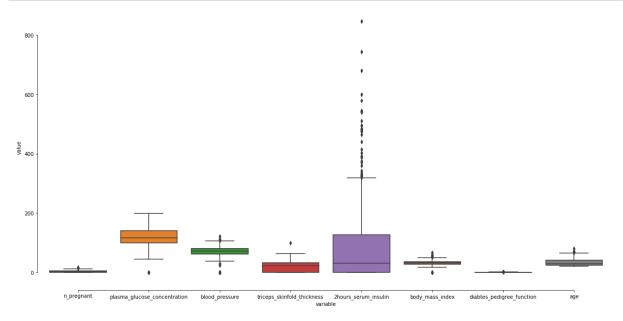
In [9]: data_.summary().toPandas().T

Out[9]:

	0	1	2	3	4	
summary	count	mean	stddev	min	25%	50
n_pregnant	768	3.84505208333333335	3.36957806269887	0	1	
plasma_glucose_concentration	768	120.89453125	31.97261819513622	0.0	99.0	117
blood_pressure	768	69.10546875	19.355807170644777	0.0	62.0	72
triceps_skinfold_thickness	768	20.536458333333333	15.952217567727642	0.0	0.0	23
2hours_serum_insulin	768	79.79947916666667	115.24400235133803	0.0	0.0	29
body_mass_index	768	31.99257813890775	7.884160293010772	0.0	27.3	32
diabtes_pedigree_function	768	0.4718763029280429	0.3313285967924436	0.078	0.243	0.37
age	768	33.240885416666664	11.760231540678689	21	24	1
class	768	0.34895833333333333	0.476951377242799	0	0	
4						•

Vamos plotar Boxplots das vatiáveis para ter uma visão geral de dimensão e distribuição dos dados

```
In [58]:
                                                                                    import matplotlib.pyplot as plt
                                                                                     import seaborn as sns
                                                                                    plt.rcParams['figure.figsize']=(20,10)
                                                                                     tips = sns.load_dataset("tips")
                                                                                    sns.boxplot(x="variable", y="value", data=pd.melt(data_.toPandas().drop(['clasure of the content of the conte
                                                                                     s'], axis=1)))
                                                                                     sns.despine(offset=10, trim=True)
```



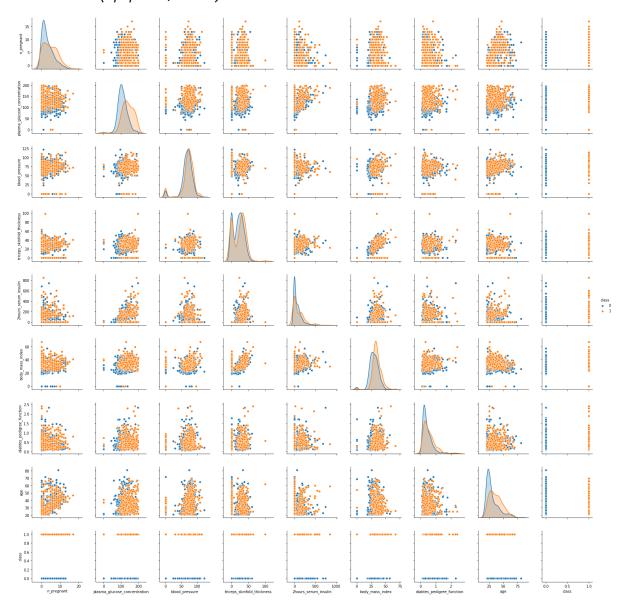
No gráfico acima fica evidente que algumas variáveis apresentam bastantes outliers, o que pode impactar na performance dos modelos de Machine learning

Futuramente iremos tratá-los a fim de verificar o quanto esses outliers interferem no modelo

Vamos plotar uma Scatterplot Matrix para ver se conseguimos enxergar relações entre as variáveis, assim como se existe alguma Decision Boundery aparente aos nossos olhos

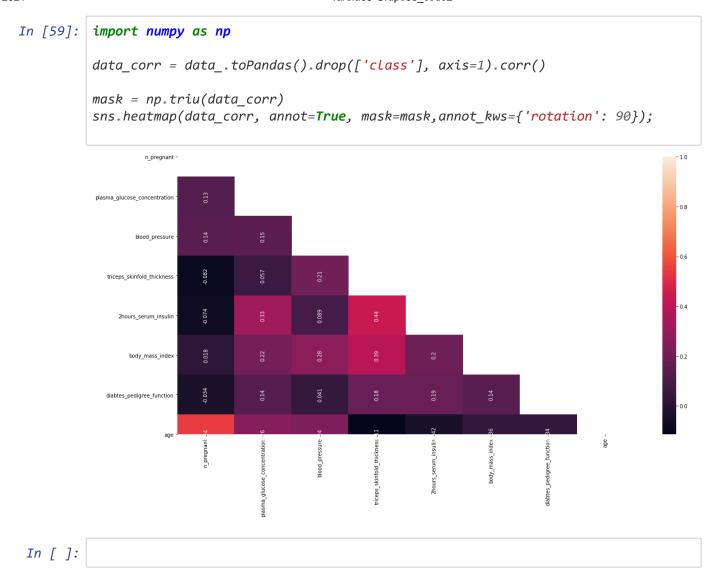
In [11]: sns.pairplot(data_.toPandas(), hue="class");

> /home/matheus/anaconda3/lib/python3.7/site-packages/statsmodels/nonparametri c/kde.py:487: RuntimeWarning: invalid value encountered in true_divide binned = fast_linbin(X, a, b, gridsize) / (delta * nobs) /home/matheus/anaconda3/lib/python3.7/site-packages/statsmodels/nonparametri c/kdetools.py:34: RuntimeWarning: invalid value encountered in double_scalars FAC1 = 2*(np.pi*bw/RANGE)**2



Notas: >>>

Agora que já tivemos um insight visual, vamos mensurar numericalmente o qual forte são as relações entre essas variáveis usando a matrix de correlação



Dentre as variáveis independentes do modelo "n_pregnant" é a única que aparenta ter um comportamento discreto. Abaixo iremos explorar ela um pouco

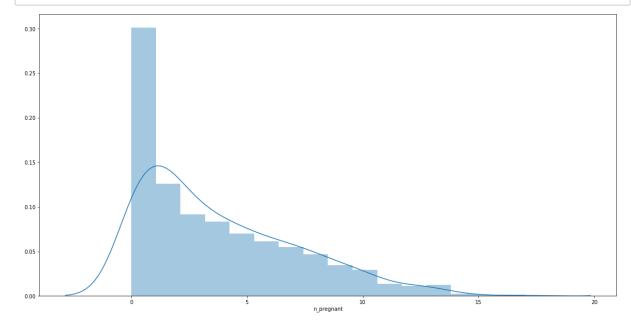
Valores únicos e sua ocorrência:

In [13]: data_.groupby('n_pregnant').count().toPandas().sort_values(by='count', ascendi ng=**False**)

Out[13]:

	n_pregnant	count
1	1	135
16	0	111
15	2	103
4	3	75
9	4	68
5	5	57
3	6	50
11	7	45
10	8	38
7	9	28
12	10	24
13	11	11
2	13	10
0	12	9
14	14	2
6	15	1
8	17	1

In [60]: sns.distplot(data_.toPandas()["n_pregnant"]);



In []:

Vamos fazer um teste de normalidade

```
In [15]: from scipy import stats
          x = data_.select('n_pregnant').toPandas()['n_pregnant']
          k2, p = stats.normaltest(x)
          alpha = 1e-3
          print("p = {:g}".format(p))
          if p < alpha:</pre>
                print("The null hypothesis can be rejected")
          else:
                print("The null hypothesis cannot be rejected")
         p = 3.91429e - 18
          The null hypothesis can be rejected
```

O teste rejeitou a hipotese nula, consequentemente a coluna "n pregnant" não apresenta uma curva normal

Criando os modelos ML

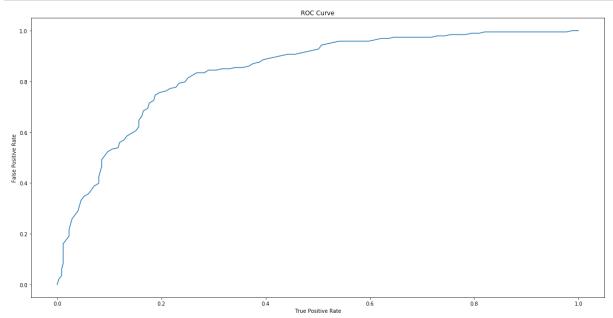
Antes de começar o preprocessamento do Dataframe vamos treinar um modelo sem e verificar as metricas com os dados crus a fim de criar um benchmark

```
In [16]: stages = []
         label stringIdx = StringIndexer(inputCol = 'class', outputCol = 'label')
         stages += [label stringIdx]
         assemblerInputs = ['n_pregnant', 'age', 'plasma_glucose_concentration', 'blood_
         pressure','triceps_skinfold_thickness',
                      '2hours_serum_insulin','body_mass_index', 'diabtes_pedigree_functi
         on']
         assembler = VectorAssembler(inputCols=assemblerInputs, outputCol="features")
         stages += [assembler]
```

```
In [17]: cols = data .columns
         pipeline = Pipeline(stages = stages)
         pipelineModel = pipeline.fit(data )
         data = pipelineModel.transform(data )
         selectedCols = ['label', 'features'] + cols
         data_ = data_.select(selectedCols)
         data .printSchema()
         root
          |-- label: double (nullable = false)
          |-- features: vector (nullable = true)
          |-- n pregnant: integer (nullable = true)
          |-- plasma glucose concentration: float (nullable = true)
          |-- blood pressure: float (nullable = true)
          |-- triceps_skinfold_thickness: float (nullable = true)
          |-- 2hours_serum_insulin: float (nullable = true)
          |-- body_mass_index: float (nullable = true)
          |-- diabtes pedigree function: float (nullable = true)
          |-- age: integer (nullable = true)
          |-- class: integer (nullable = true)
In [18]: # Preparando pros modelos
         train, test = data_.randomSplit([0.7, 0.3], seed = 2019)
         print("Training Dataset Count: " + str(train.count()))
         print("Test Dataset Count: " + str(test.count()))
         Training Dataset Count: 544
         Test Dataset Count: 224
```

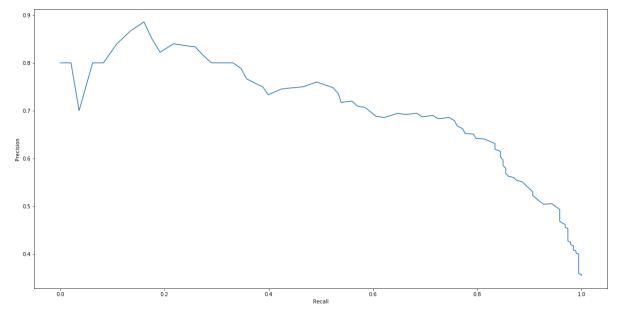
Logistic Regression Model

```
In [62]:
         from pyspark.ml.classification import LogisticRegression
         lr = LogisticRegression(featuresCol = 'features', labelCol = 'label', maxIter=
         10)
         lrModel = lr.fit(train)
         trainingSummary = lrModel.summary
         roc = trainingSummary.roc.toPandas()
         plt.plot(roc['FPR'],roc['TPR'])
         plt.ylabel('False Positive Rate')
         plt.xlabel('True Positive Rate')
         plt.title('ROC Curve')
         plt.show()
         print('Training set areaUnderROC: ' + str(trainingSummary.areaUnderROC))
```



Training set areaUnderROC: 0.8412677324594429

```
In [63]: pr = trainingSummary.pr.toPandas()
         plt.plot(pr['recall'],pr['precision'])
         plt.ylabel('Precision')
         plt.xlabel('Recall')
         plt.show()
```



```
In [21]: from pyspark.ml.evaluation import BinaryClassificationEvaluator
         predictions = lrModel.transform(test)
         evaluator = BinaryClassificationEvaluator()
         print('Test Area Under ROC', evaluator.evaluate(predictions))
```

Test Area Under ROC 0.8171812080536911

Decision Tree Classifier

```
In [22]: | from pyspark.ml.classification import DecisionTreeClassifier
         dt = DecisionTreeClassifier(featuresCol = 'features', labelCol = 'label', maxD
         epth = 3)
         dtModel = dt.fit(train)
         predictions = dtModel.transform(test)
         evaluator = BinaryClassificationEvaluator()
         print("Test Area Under ROC: " + str(evaluator.evaluate(predictions, {evaluator
         .metricName: "areaUnderROC"})))
```

Test Area Under ROC: 0.720268456375839

Random Forest Classifier

```
In [23]: from pyspark.ml.classification import RandomForestClassifier
         rf = RandomForestClassifier(featuresCol = 'features', labelCol = 'label')
         rfModel = rf.fit(train)
         predictions = rfModel.transform(test)
         evaluator = BinaryClassificationEvaluator()
         print("Test Area Under ROC: " + str(evaluator.evaluate(predictions, {evaluator
         .metricName: "areaUnderROC"})))
```

Test Area Under ROC: 0.8136912751677854

Gradient-Boosted Tree Classifier

```
In [24]: from pyspark.ml.classification import GBTClassifier
         gbt = GBTClassifier(maxIter=10)
         gbtModel = gbt.fit(train)
         predictions = qbtModel.transform(test)
         evaluator = BinaryClassificationEvaluator()
         print("Test Area Under ROC: " + str(evaluator.evaluate(predictions, {evaluator
         .metricName: "areaUnderROC"})))
```

Test Area Under ROC: 0.7434004474272928

Gradient-Boosted Tree Classifier + Cross Validator + Grid Search

```
from pyspark.ml.tuning import ParamGridBuilder, CrossValidator
In [25]:
         paramGrid = (ParamGridBuilder()
                      .addGrid(gbt.maxDepth, [2, 4, 6])
                       .addGrid(gbt.maxBins, [20, 60])
                       .addGrid(qbt.maxIter, [10, 20])
                       .build())
         cv = CrossValidator(estimator=qbt, estimatorParamMaps=paramGrid, evaluator=eva
         luator, numFolds=5)
         # Run cross validations. This can take about 6 minutes since it is training o
         ver 20 trees!
         cvModel = cv.fit(train)
         predictions = cvModel.transform(test)
         evaluator.evaluate(predictions)
```

Out[25]: 0.8207606263982102

Nosso Benchmark

Logistic Regression Model - Area Under ROC 0.8171812080536911

Decision Tree Classifier - Area Under ROC: 0.720268456375839

Random Forest Classifier - Area Under ROC: 0.8136912751677854

Gradient-Boosted Tree Classifier - Area Under ROC: 0.7434004474272928

Gradient-Boosted Tree Classifier + Cross Validator + Grid Search - Area Under ROC:0.8207606263982102

Criando nosso modelo

Vamos desenvolver algumas funções de processamento, treinamento e teste, para que em seguida possamos rodar todas possibilidades de modelos, considerando varios tipos de processamento difernetes

Funções de processamento:

```
In [26]: from pyspark.ml.classification import RandomForestClassifier
         from pyspark.ml.classification import GBTClassifier
         from pyspark.ml.classification import DecisionTreeClassifier
         from pyspark.ml.classification import MultilayerPerceptronClassifier
         from pyspark.ml.tuning import ParamGridBuilder, CrossValidator
         from pyspark.ml.evaluation import BinaryClassificationEvaluator
         from pyspark.mllib.evaluation import BinaryClassificationMetrics
         from pyspark.ml.classification import LogisticRegression
         from pyspark.ml.feature import MinMaxScaler, StandardScaler
         from pyspark.ml import Pipeline
         from pyspark.sql import functions as f
```

Remover outliers

```
In [44]: # outliers parameters = [False, 'StdDev']
         def remove_outliers(df, outliers_type):
             if outliers type == 'StdDev':
                 data = df.where(df.body_mass_index > 10.0)
                 data = data.where(data.triceps_skinfold_thickness > 0)
                 data = data.where(data['2hours_serum_insulin'] > 0 )
                 data = data.where( data['2hours serum insulin'] < 350)</pre>
                 return data
             eLse:
                 return df
```

Fazer amostragem nas classes - balanceamento

```
In [28]: # sample_parameters = [False, True]
         def balance_dataframe(df, sample_type):
             if sample_type == True:
                 lenght = []
                 sample list = []
                 lenght.append(df.filter(df['class'] == 1).count())
                 lenght.append(df.filter(df['class'] == 0).count())
                 sample size = min(lenght)
                 sample_list.extend(df.filter(df['class'] == 1).rdd.takeSample(False, s
         ample_size, seed=0))
                 sample list.extend(df.filter(df['class'] == 0).rdd.takeSample(False, s
         ample size, seed=0))
                 data = sqlContext.createDataFrame(sample list)
                 return data
             else:
                 return df
```

Remover agrupamento de duplicados remanecentes

```
In [29]: | # remove distinct parameters = [False,['n pregnant','2hours serum insulin', 'a
         ge', 'class'],[ 'age', 'class']]
         def remove dinstinct(df, remove distinct type):
             if remove distinct type == False:
                 return df
             else:
                 data = df.drop_duplicates(subset=remove_distinct_type)
                 return data
```

Normalizar valores - MinMax / Standalizer

```
In [30]: # normalization parameters = [False,'MinMax', 'Standalizer']
         def normalize dataframe(df, normalization type):
             columns to scale = []
             stages = []
             for column in df.columns:
                 if column == "class": continue
                 columns_to_scale.append(column)
             if normalization type == 'MinMax':
                 label stringIdx = StringIndexer(inputCol = 'class', outputCol = 'labe
         L')
                 stages += [label_stringIdx]
                 pipeline = Pipeline(stages = stages)
                 pipelineModel = pipeline.fit(df)
                 df_result = pipelineModel.transform(df)
                 assemblers = VectorAssembler(inputCols=columns to scale, outputCol="fe
         atures_preprocessing")
                 vector = assemblers.transform(df result)
                 standard scaler = MinMaxScaler(inputCol="features preprocessing", outp
         utCol="features")
                 train = standard scaler.fit(vector).transform(vector)
                 return train
             elif normalization_type == 'Standalizer':
                 label_stringIdx = StringIndexer(inputCol = 'class', outputCol = 'labe
         L')
                 stages += [label_stringIdx]
                 pipeline = Pipeline(stages = stages)
                 pipelineModel = pipeline.fit(df)
                 df result = pipelineModel.transform(df)
                 assemblers = VectorAssembler(inputCols=columns_to_scale, outputCol="fe")
         atures preprocessing")
                 vector = assemblers.transform(df result)
                 standard scaler = StandardScaler(inputCol="features preprocessing", ou
         tputCol="features")
                 train = standard scaler.fit(vector).transform(vector)
                 return train
             else:
                 label stringIdx = StringIndexer(inputCol = 'class', outputCol = 'labe
         L')
                 stages += [label_stringIdx]
                 assembler = VectorAssembler(inputCols=columns to scale, outputCol="fea
         tures")
```

```
stages += [assembler]
cols = df.columns
pipeline = Pipeline(stages = stages)
pipelineModel = pipeline.fit(df)
df result = pipelineModel.transform(df)
selectedCols = ['label', 'features'] + cols
df_result = df_result.select(selectedCols)
return df result
```

Modelos:

```
In [46]: | def apply_Logistic_Regression_Model(df):
             train, test = df.randomSplit([0.8, 0.2], seed = 2019)
             #tocar esssa linha
             model = LogisticRegression(featuresCol = 'features', labelCol = 'label', m
         axIter=10)
             evaluator = BinaryClassificationEvaluator()
              # Create ParamGrid for Cross Validation
             paramGrid = (ParamGridBuilder()
                       .addGrid(model.regParam,[0.02])
                       .addGrid(model.elasticNetParam,[0.2])
                       .build())
                 # Create 5-fold CrossValidator
             cv = CrossValidator(estimator = model,
                                estimatorParamMaps = paramGrid,
                                evaluator = evaluator,
                                numFolds = 5)
             # Run cross validations.
             cvModel = cv.fit(train)
             # Use test set here so we can measure the accuracy of our model on new dat
             predictions = cvModel.transform(test)
             # cvModel uses the best model found from the Cross Validation
             # Evaluate best model
             acc = evaluator.evaluate(predictions)
             auc = BinaryClassificationMetrics(predictions['label', 'prediction'].rdd).a
         reaUnderROC
             print('Accuracy:', acc)
             print('AUC:',auc)
             return acc, auc
```

```
In [47]: | def apply_Decision_Tree_Model(df):
             train, test = df.randomSplit([0.8, 0.2], seed = 2019)
             #tocar esssa linha
             model = DecisionTreeClassifier(featuresCol = 'features', labelCol = 'labe
         L')
             evaluator = BinaryClassificationEvaluator()
              # Create ParamGrid for Cross Validation
             paramGrid = (ParamGridBuilder()
                           .addGrid(model.maxDepth, [2, 20])
                           .addGrid(model.maxBins, [10, 40])
                           .build())
                 # Create 5-fold CrossValidator
             cv = CrossValidator(estimator = model,
                                estimatorParamMaps = paramGrid,
                                evaluator = evaluator,
                                numFolds = 5)
             # Run cross validations.
             cvModel = cv.fit(train)
             # Use test set here so we can measure the accuracy of our model on new dat
             predictions = cvModel.transform(test)
             # cvModel uses the best model found from the Cross Validation
             # Evaluate best model
             acc = evaluator.evaluate(predictions)
             auc = BinaryClassificationMetrics(predictions['label', 'prediction'].rdd).a
         reaUnderROC
             print('Accuracy:', acc)
             print('AUC:',auc)
             return acc, auc
```

```
In [48]: def apply Random Forest Model(df):
             train, test = df.randomSplit([0.8, 0.2], seed = 2019)
             # Create an initial RandomForest model.
             model = RandomForestClassifier(labelCol="label", featuresCol="features")
             # Evaluate model
             evaluator = BinaryClassificationEvaluator()
             # Create ParamGrid for Cross Validation
             paramGrid = (ParamGridBuilder()
                       .addGrid(model.maxDepth, [2 ])
                       .addGrid(model.maxBins, [10,])
                       .addGrid(model.numTrees, [5])
                       .build())
             # Create 5-fold CrossValidator
             cv = CrossValidator(estimator = model,
                                estimatorParamMaps = paramGrid,
                                evaluator = evaluator,
                                numFolds = 5)
             # Run cross validations.
             cvModel = cv.fit(train)
             # Use test set here so we can measure the accuracy of our model on new dat
             predictions = cvModel.transform(test)
             # cvModel uses the best model found from the Cross Validation
             # Evaluate best model
             acc = evaluator.evaluate(predictions)
             auc = BinaryClassificationMetrics(predictions['label', 'prediction'].rdd).a
         reaUnderROC
             print('Accuracy:', acc)
             print('AUC:', auc)
             return acc, auc
```

```
In [49]: def apply Gradient BoostedT Model(df):
             train, test = df.randomSplit([0.8, 0.2], seed = 2019)
             #tocar esssa linha
             model = GBTClassifier(featuresCol = 'features', labelCol = 'label', maxIte
         r=10)
             evaluator = BinaryClassificationEvaluator()
              # Create ParamGrid for Cross Validation
             paramGrid = (ParamGridBuilder()
                       .addGrid(model.maxDepth, [2, ])
                       .addGrid(model.maxBins, [20])
                       .addGrid(model.maxIter, [10])
                       .build())
                 # Create 5-fold CrossValidator
             cv = CrossValidator(estimator = model,
                                estimatorParamMaps = paramGrid,
                                evaluator = evaluator,
                                numFolds = 5)
             # Run cross validations.
             cvModel = cv.fit(train)
             # Use test set here so we can measure the accuracy of our model on new dat
         а
             predictions = cvModel.transform(test)
             # cvModel uses the best model found from the Cross Validation
             # Evaluate best model
             acc = evaluator.evaluate(predictions)
             auc = BinaryClassificationMetrics(predictions['label', 'prediction'].rdd).a
         reaUnderROC
             print('Accuracy:', acc)
             print('AUC:',auc)
             return acc, auc
```

```
In [50]: | def apply MultiLayer Perceptron(df):
             train, test = df.randomSplit([0.8, 0.2], seed = 2019)
             model = MultilayerPerceptronClassifier(featuresCol = 'features', labelCol
         = 'label')
             evaluator = BinaryClassificationEvaluator()
             paramGrid = (ParamGridBuilder()
                  .addGrid(model.maxIter,[10, 20])
                  .addGrid(model.stepSize ,[0.03, 0.01])
                  .addGrid(model.solver ,['l-bfgs', 'gd'])
                  .addGrid(model.layers, [[8, 12, 2], [8, 5, 2], [8, 5, 4, 2]])
                  .build())
                 # Create 5-fold CrossValidator
             cv = CrossValidator(estimator = model, estimatorParamMaps = paramGrid, eva
         luator = evaluator, numFolds = 5)
             # Run cross validations.
             cvModel = cv.fit(train)
             # Use test set here so we can measure the accuracy of our model on new dat
             predictions = cvModel.transform(test)
             # cvModel uses the best model found from the Cross Validation
             # Evaluate best model
             acc = f
             evaluator.evaluate(predictions)
             auc = BinaryClassificationMetrics(predictions['label', 'prediction'].rdd).a
         reaUnderROC
             print('Accuracy:', acc)
             print('AUC:',auc)
             return acc, auc
```

Ordem que as funções devem rodar

```
data = removeoutliers(data, outliers tag)
data = balance_dataframe(data, sample_tag)
data = remove dinstinct(data, remove distinct tag)
data = normalize dataframe(data, normalization tag)
apply_Logistic_Regression_Model(data)
```

No codigo a baixo iremos fazer um loop entre todos metodos de processamento de modelos de machine learnin para encontrar qual o melhor pipeline de processamento/modelo para nosso conjunto de dados

```
In [51]: from datetime import datetime
         start = datetime.now()
         # models = ["MLP", "GBT", "RDF", "DCT", "LGR"]
         outliers parameters = [False, 'StdDev', 'DropCol']
         sample parameters = [False, True]
         remove_distinct_parameters = [False,['n_pregnant','2hours_serum_insulin', 'ag
         e', 'class'],[ 'age', 'class']]
         normalization parameters = [False, 'MinMax', 'Standalizer']
         models = ["RDF","LGR","GBT"]
         result = []
         for model in models:
             for outliers tag in outliers parameters:
                 for sample_tag in sample_parameters:
                      for remove_distinct_tag in remove_distinct_parameters:
                          for normalization tag in normalization parameters:
                              dic = \{\}
                              data = remove_outliers(data_model, outliers_tag)
                              data = balance_dataframe(data, sample_tag)
                              data = remove dinstinct(data, remove distinct tag)
                              data = normalize dataframe(data, normalization tag)
                              print(model)
                              if model == "MLP":
                                  acc, auc = apply_MultiLayer_Perceptron(data)
                              elif model == "GBT":
                                 acc, auc = apply_Gradient_BoostedT_Model(data)
                              elif model == "RDF":
                                  acc, auc = apply_Random_Forest_Model(data)
                              elif model == "DCT":
                                  acc, auc = apply_Decision_Tree_Model(data)
                              elif model == "LGR":
                                  acc, auc = apply_Logistic_Regression_Model(data)
                              print(" ")
                              dic["Model Name"] = model
                              dic["outliers_tag"] = outliers_tag
                              dic["sample_tag"] = sample_tag
                              dic["remove_distinct_tag"] = remove_distinct_tag
                              dic["normalization_tag"] = normalization_tag
                              dic["Acuracia"] = acc
                              dic["AUC"] = auc
                              result.append(dic)
         df = pd.DataFrame(result)
```

```
df = df.sort_values(by=['AUC'], ascending=False)
stop = datetime.now()
print('Time: ', stop - start)
```

RDF

Accuracy: 0.8453608247422681 AUC: 0.7697368421052632

RDF

Accuracy: 0.7912234042553192 AUC: 0.6758209822725951

RDF

Accuracy: 0.7912234042553192 AUC: 0.6758209822725951

RDF

Accuracy: 0.7703488372093023 AUC: 0.7881578947368422

RDF

Accuracy: 0.803166666666665 AUC: 0.8331043956043955

RDF

Accuracy: 0.803166666666665 AUC: 0.8331043956043955

RDF

Accuracy: 0.6212121212121212 AUC: 0.5833333333333334

RDF

Accuracy: 0.6262626262626262

AUC: 0.5625

RDF

Accuracy: 0.6262626262626262

AUC: 0.5625

RDF

Accuracy: 0.8359113712374582 AUC: 0.7317124735729387

RDF

Accuracy: 0.8292224080267557 AUC: 0.7609427609427609

RDF

Accuracy: 0.8292224080267557 AUC: 0.7609427609427609

RDF

Accuracy: 0.7361793611793612 AUC: 0.6512422360248447

RDF

Accuracy: 0.7751842751842751

AUC: 0.7111111111111111

RDF

Accuracy: 0.7751842751842751

AUC: 0.7111111111111111

RDF

Accuracy: 0.8 AUC: 0.775

RDF

Accuracy: 0.9861111111111112

AUC: 0.875

RDF

Accuracy: 0.9861111111111112

AUC: 0.875

RDF

Accuracy: 0.8771991555242786 AUC: 0.7384615384615384

RDF

Accuracy: 0.8496428571428571 AUC: 0.8082706766917294

RDF

Accuracy: 0.8496428571428571 AUC: 0.8082706766917294

RDF

Accuracy: 0.8968750000000001

AUC: 0.8875

RDF

Accuracy: 0.950625 AUC: 0.962962962963

RDF

Accuracy: 0.950625 AUC: 0.962962962963

RDF

Accuracy: 0.44696969696969696

AUC: 0.386363636363635

RDF

Accuracy: 0.5857142857142856 AUC: 0.534722222222222

RDF

Accuracy: 0.5857142857142856 AUC: 0.534722222222222

RDF

Accuracy: 0.7391304347826086

AUC: 0.735

RDF

Accuracy: 0.8063241106719368

AUC: 0.7685185185186

RDF

Accuracy: 0.8063241106719368 AUC: 0.7685185185185186

RDF

Accuracy: 0.8527667984189723 AUC: 0.7559288537549407

RDF

Accuracy: 0.8241106719367588 AUC: 0.8003952569169962

RDF

Accuracy: 0.8241106719367588 AUC: 0.8003952569169962

RDF

Accuracy: 0.8 AUC: 0.8125

RDF

Accuracy: 0.666666666666666 AUC: 0.7857142857142857

RDF

Accuracy: 0.666666666666666 AUC: 0.7857142857142857

RDF

Accuracy: 0.8453608247422681 AUC: 0.7697368421052632

RDF

Accuracy: 0.7912234042553192 AUC: 0.6758209822725951

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Accuracy: 0.7912234042553192 AUC: 0.6758209822725951

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Accuracy: 0.803166666666665 AUC: 0.8331043956043955

RDF

Accuracy: 0.6212121212121212 AUC: 0.5833333333333334

RDF

Accuracy: 0.6262626262626262

AUC: 0.5625

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Accuracy: 0.6262626262626262

AUC: 0.5625

RDF

Accuracy: 0.8359113712374582 AUC: 0.7317124735729387

RDF

Accuracy: 0.8292224080267557 AUC: 0.7609427609427609

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RDF

Accuracy: 0.7361793611793612 AUC: 0.6512422360248447

RDF

Accuracy: 0.7751842751842751 AUC: 0.711111111111111

RDF

Accuracy: 0.7751842751842751 AUC: 0.711111111111111

RDF

Accuracy: 0.8 AUC: 0.775

RDF

Accuracy: 0.9861111111111112

AUC: 0.875

RDF

Accuracy: 0.9861111111111112

AUC: 0.875

LGR

Accuracy: 0.7993127147766323 AUC: 0.7573049987925623

LGR

Accuracy: 0.7429078014184399 AUC: 0.6794871794871794

LGR

Accuracy: 0.6914893617021277 AUC: 0.6462201591511936

LGR

Accuracy: 0.721576227390181 AUC: 0.8214285714285714

LGR

Accuracy: 0.76933333333333333 AUC: 0.717391304347826

LGR

Accuracy: 0.74033333333333333 AUC: 0.7083333333333334

LGR

Accuracy: 0.58585858585857 AUC: 0.649999999999999

LGR

Accuracy: 0.6565656565656565

AUC: 0.5625

LGR

Accuracy: 0.6363636363636364 AUC: 0.6499999999999999

LGR

Accuracy: 0.7391304347826085 AUC: 0.6599468320779797

LGR

Accuracy: 0.8189799331103679 AUC: 0.7541806020066888

LGR

Accuracy: 0.774247491638796 AUC: 0.7234800838574424

LGR

Accuracy: 0.6480343980343977 AUC: 0.5785984848484848

LGR

Accuracy: 0.6689189189189 AUC: 0.6328029375764993

LGR

Accuracy: 0.6744471744471747 AUC: 0.5872015915119363

LGR

Accuracy: 0.5714285714285714 AUC: 0.43181818181819

LGR

Accuracy: 0.680555555555556 AUC: 0.5357142857142857

LGR

Accuracy: 0.7361111111111113 AUC: 0.5833333333333334

LGR

Accuracy: 0.7424349049964815 AUC: 0.6825396825396826

LGR

Accuracy: 0.7864285714285715 AUC: 0.7564935064935066

LGR

Accuracy: 0.7228571428571428 AUC: 0.70000000000000001

LGR

Accuracy: 0.7962500000000005 AUC: 0.7639225181598064

LGR

Accuracy: 0.89

AUC: 0.916666666666667

LGR

Accuracy: 0.8700000000000003

AUC: 0.825

LGR

Accuracy: 0.75757575757576 AUC: 0.6083333333333334

LGR

Accuracy: 0.899999999999999 AUC: 0.7285714285714286

LGR

Accuracy: 0.8428571428571427 AUC: 0.7285714285714286

LGR

Accuracy: 0.6996047430830039 AUC: 0.71500000000000001

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Accuracy: 0.7312252964426879 AUC: 0.7601214574898786

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Accuracy: 0.7154150197628459 AUC: 0.6887351778656127

LGR

Accuracy: 0.72727272727273

AUC: 0.66699604743083

LGR

Accuracy: 0.6818181818181819

AUC: 0.67

LGR

Accuracy: 0.6897233201581028

AUC: 0.67

LGR

Accuracy: 0.48

AUC: 0.2222222222222

LGR

Accuracy: 0.625

AUC: 0.75

LGR

Accuracy: 0.45833333333333326

AUC: 0.0

LGR

Accuracy: 0.7993127147766323 AUC: 0.7573049987925623

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Accuracy: 0.7429078014184399 AUC: 0.6794871794871794

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AUC: 0.5625

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Accuracy: 0.5714285714285714 AUC: 0.43181818181819

LGR

Accuracy: 0.680555555555556 AUC: 0.5357142857142857

LGR

Accuracy: 0.7361111111111113 AUC: 0.58333333333333334

GBT

Accuracy: 0.8324169530355098 AUC: 0.7327127659574468

GBT

Accuracy: 0.8262411347517731 AUC: 0.6997549019607843

GBT

Accuracy: 0.8257978723404256 AUC: 0.6997549019607843

GBT

Accuracy: 0.8338178294573643 AUC: 0.7282608695652174

GBT

Accuracy: 0.84433333333333333 AUC: 0.8219533275713051

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Accuracy: 0.84433333333333333 AUC: 0.8219533275713051

GBT

Accuracy: 0.6565656565656566

AUC: 0.696969696969697

GBT

Accuracy: 0.7828282828282829

AUC: 0.75

GBT

Accuracy: 0.7828282828282829

AUC: 0.75

GBT

Accuracy: 0.8520066889632106 AUC: 0.7670004171881519

GBT

Accuracy: 0.8808528428093647

AUC: 0.7874426366291197

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AUC: 0.7874426366291197

GBT

Accuracy: 0.7733415233415232

AUC: 0.73611111111111112

GBT

Accuracy: 0.7733415233415232

AUC: 0.7361111111111112

GBT

Accuracy: 0.7571428571428571

AUC: 0.7571428571428572

GBT

Accuracy: 0.9305555555555555

AUC: 0.9090909090909091

GBT

Accuracy: 0.9305555555555555

AUC: 0.9090909090909091

GBT

Accuracy: 0.8662913441238564

AUC: 0.8712121212121212

GBT

Accuracy: 0.8492857142857142

AUC: 0.75

GBT

Accuracy: 0.8485714285714286

AUC: 0.75

GBT

Accuracy: 0.8706250000000001 AUC: 0.8101903695408734

GBT

Accuracy: 0.9012500000000004 AUC: 0.8607843137254902

GBT

Accuracy: 0.9012500000000004 AUC: 0.8607843137254902

GBT

Accuracy: 0.72727272727273 AUC: 0.6439393939393939

GBT

Accuracy: 0.8785714285714286 AUC: 0.791666666666667

GBT

Accuracy: 0.8785714285714286 AUC: 0.791666666666667

GBT

Accuracy: 0.7885375494071146 AUC: 0.7351190476190477

GBT

Accuracy: 0.7658102766798419 AUC: 0.6887351778656127

GBT

Accuracy: 0.7658102766798419 AUC: 0.6887351778656127

GBT

Accuracy: 0.8448616600790514 AUC: 0.7559523809523809

GBT

Accuracy: 0.8043478260869567 AUC: 0.7114624505928854

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Accuracy: 0.8043478260869567 AUC: 0.7114624505928854

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Accuracy: 0.679999999999999

AUC: 0.8125

GBT

Accuracy: 0.625 AUC: 0.625

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AUC: 0.625

GBT

Accuracy: 0.8324169530355098 AUC: 0.7327127659574468

GBT

Accuracy: 0.8262411347517731 AUC: 0.6997549019607843

GBT

Accuracy: 0.8257978723404256 AUC: 0.6997549019607843

GBT

Accuracy: 0.8338178294573643 AUC: 0.7282608695652174

GBT

Accuracy: 0.84433333333333333 AUC: 0.8219533275713051

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Accuracy: 0.84433333333333333 AUC: 0.8219533275713051

GBT

Accuracy: 0.6565656565656566 AUC: 0.696969696969697

GBT

Accuracy: 0.7828282828282829

AUC: 0.75

GBT

Accuracy: 0.7828282828282829

AUC: 0.75

GBT

Accuracy: 0.8520066889632106 AUC: 0.7670004171881519

GBT

Accuracy: 0.8808528428093647 AUC: 0.7874426366291197

GBT

Accuracy: 0.8808528428093647 AUC: 0.7874426366291197

GBT

Accuracy: 0.7699631449631451 AUC: 0.7014742014742015

GBT

Accuracy: 0.7733415233415232 AUC: 0.7361111111111112

GBT

Accuracy: 0.7733415233415232 AUC: 0.7361111111111112

GBT

Accuracy: 0.7571428571428571 AUC: 0.7571428571428572

GBT

Accuracy: 0.9305555555555555 AUC: 0.9090909090909091

GBT

Accuracy: 0.930555555555555 AUC: 0.9090909090909091

Time: 2:51:16.107462

	Model Name	outliers_tag	sample_tag	remove_distinct_tag	normalization_tag	Acuracia	AUC
22	RDF	StdDev	False	[n_pregnant, 2hours_serum_insulin, age, class]	MinMax	0.950625	0.962963
23	RDF	StdDev	False	[n_pregnant, 2hours_serum_insulin, age, class]	Standalizer	0.950625	0.962963
76	LGR	StdDev	False	[n_pregnant, 2hours_serum_insulin, age, class]	MinMax	0.890000	0.916667
124	GBT	False	True	[age, class]	MinMax	0.930556	0.909091
125	GBT	False	True	[age, class]	Standalizer	0.930556	0.909091
161	GBT	DropCol	True	[age, class]	Standalizer	0.930556	0.909091
160	GBT	DropCol	True	[age, class]	MinMax	0.930556	0.909091
21	RDF	StdDev	False	[n_pregnant, 2hours_serum_insulin, age, class]	False	0.896875	0.887500
52	RDF	DropCol	True	[age, class]	MinMax	0.986111	0.875000
53	RDF	DropCol	True	[age, class]	Standalizer	0.986111	0.875000
16	RDF	False	True	[age, class]	MinMax	0.986111	0.875000
17	RDF	False	True	[age, class]	Standalizer	0.986111	0.875000
126	GBT	StdDev	False	False	False	0.866291	0.871212
131	GBT	StdDev	False	[n_pregnant, 2hours_serum_insulin, age, class]	Standalizer	0.901250	0.860784
130	GBT	StdDev	False	[n_pregnant, 2hours_serum_insulin, age, class]	MinMax	0.901250	0.860784
4							•