Comparative Analysis of Machine Learning Algorithms in Predicting Smoking and Drinking Behaviors

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1 Trabalho final Sistemas Inteligentes

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```
[1]: # Etapa 1: Carregar e Preparar o Dataset

import pandas as pd

# Caminho do dataset
dataset_path = './smoking_driking_dataset_Ver01.csv' # Ajuste conforme ou
caminho real

# Tentando carregar o dataset
try:
    data = pd.read_csv(dataset_path)
    # Exibindo as primeiras linhas do dataset para verificação
    display_data = data.head()
except Exception as e:
    display_data
display_data
```

```
[1]:
                            weight
                                                sight_left
                                                             sight_right hear_left \
         sex
              age
                   height
                                     waistline
                                75
     0 Male
               35
                       170
                                          90.0
     1 Male
               30
                       180
                                80
                                          89.0
                                                        0.9
                                                                      1.2
                                                                                 1.0
     2 Male
               40
                       165
                                75
                                          91.0
                                                        1.2
                                                                      1.5
                                                                                 1.0
     3 Male
               50
                       175
                                80
                                          91.0
                                                        1.5
                                                                      1.2
                                                                                 1.0
     4 Male
               50
                       165
                                60
                                          80.0
                                                        1.0
                                                                      1.2
                                                                                 1.0
        hear_right
                       SBP
                               LDL chole
                                          triglyceride hemoglobin urine_protein
     0
               1.0
                     120.0
                                    126.0
                                                   92.0
                                                                17.1
     1
               1.0
                     130.0 ...
                                    148.0
                                                  121.0
                                                                15.8
                                                                                 1.0
     2
               1.0
                     120.0 ...
                                     74.0
                                                  104.0
                                                                15.8
                                                                                 1.0
                     145.0 ...
                                    104.0
                                                  106.0
     3
               1.0
                                                                17.6
                                                                                 1.0
               1.0 138.0 ...
                                    117.0
                                                  104.0
                                                                13.8
                                                                                 1.0
```

```
SGOT_AST SGOT_ALT
                                          gamma_GTP
                                                       SMK_stat_type_cd DRK_YN
   serum_creatinine
                                                40.0
                                                                    1.0
0
                          21.0
                                     35.0
                                                                               Y
                1.0
1
                0.9
                          20.0
                                     36.0
                                                27.0
                                                                    3.0
                                                                               N
2
                          47.0
                0.9
                                     32.0
                                                68.0
                                                                    1.0
                                                                               N
3
                1.1
                          29.0
                                     34.0
                                                18.0
                                                                    1.0
                                                                               N
                0.8
                          19.0
                                    12.0
                                                25.0
                                                                    1.0
                                                                               N
```

[5 rows x 24 columns]

```
[2]: from sklearn.impute import SimpleImputer
     from sklearn.preprocessing import StandardScaler, LabelEncoder
     # Imputação dos valores faltantes
     imputer = SimpleImputer(strategy='mean')
     numeric_columns = data.select_dtypes(include=[float, int]).columns
     numeric_columns = numeric_columns.drop(['SMK_stat_type_cd']) # Exclua 'DRK_YN'_u
      ⇔se ela já for categórica
     data[numeric_columns] = imputer.fit_transform(data[numeric_columns])
     # Normalizar os dados numéricos
     scaler = StandardScaler()
     data[numeric_columns] = scaler.fit_transform(data[numeric_columns])
     # Codificação de variáveis categóricas
     label_encoders = {}
     categorical columns = ['sex', 'SMK stat type cd', 'DRK YN'] # Adicione outras__
      →colunas categóricas se necessário
     for col in categorical columns:
         if col in data.columns:
             le = LabelEncoder()
             data[col] = le.fit_transform(data[col])
             label_encoders[col] = le
     # Visualizando os dados após o pré-processamento
     print(data.head())
```

```
height
                            weight waistline sight_left sight_right \
  sex
            age
    1 -0.889514 0.835874 0.936210
                                                 0.031629
                                                             0.035668
0
                                     0.739781
    1 -1.242090 1.913117 1.335755
1
                                     0.655395
                                                -0.133401
                                                             0.366370
    1 -0.536938 0.297252 0.936210
                                     0.824167
                                                 0.361690
                                                             0.862423
    1 0.168215 1.374495 1.335755
3
                                     0.824167
                                                0.856782
                                                             0.366370
    1 0.168215 0.297252 -0.262425 -0.104078
                                                0.031629
                                                             0.366370
                             SBP ... LDL chole
                                               triglyceride hemoglobin \
  hear_left hear_right
0 -0.180329
             -0.177296 -0.167261 ...
                                      0.361643
                                                   -0.392788
                                                               1.810919
1 -0.180329 -0.177296 0.520349 ...
                                      0.975435
                                                   -0.109022
                                                               0.990693
2 -0.180329
             -0.177296 -0.167261 ... -1.089136
                                                   -0.275368
                                                               0.990693
3 -0.180329 -0.177296 1.551763 ... -0.252148
                                                  -0.255798
                                                               2.126391
```

```
4 -0.180329 -0.177296 1.070436 ... 0.110547
                                                        -0.275368
                                                                   -0.271194
       urine_protein serum_creatinine SGOT_AST SGOT_ALT gamma_GTP \
    0
            -0.21526
                              0.290374 -0.212371 0.351404
                                                           0.056791
    1
            -0.21526
                              0.082270 -0.254936 0.389415 -0.201022
    2
            -0.21526
                              0.082270 0.894324 0.237373
                                                             0.612081
    3
            -0.21526
                              0.498477 0.128151 0.313394 -0.379508
    4
            -0.21526
                             -0.125833 -0.297501 -0.522835 -0.240685
       SMK_stat_type_cd DRK_YN
    0
                      0
                              1
    1
                      2
                              0
    2
                      0
                              0
    3
                      0
                              0
    4
    [5 rows x 24 columns]
[3]: from sklearn.model_selection import train_test_split
    # Dividir os dados para prever 'SMK_stat_type_cd'
    X = data.drop(['SMK_stat_type_cd', 'DRK_YN'], axis=1)
    y_smk = data['SMK_stat_type_cd']
    X_train_smk, X_test_smk, y_train_smk, y_test_smk = train_test_split(X, y_smk,_
      →test_size=0.2, random_state=42)
    # Dividir os dados para prever 'DRK_YN'
    y_drk = data['DRK_YN']
    X_train_drk, X_test_drk, y_train_drk, y_test_drk = train_test_split(X, y_drk,_

state=42)

state=42)

state=42)

    import numpy as np
    X_train_smk = np.array(X_train_smk)
    y_train_smk = np.array(y_train_smk)
    X_test_smk = np.array(X_test_smk)
    y_test_smk = np.array(y_test_smk)
    X_train_drk = np.array(X_train_drk)
    y_train_drk = np.array(y_train_drk)
    X_test_drk = np.array(X_test_drk)
    y_test_drk = np.array(y_test_drk)
    # Verificando valores únicos para DRK YN
    print("Valores únicos em DRK_YN:", data['DRK_YN'].unique())
```

```
# Verificando o balanceamento das classes
print("Contagem de classes em DRK_YN:")
print(data['DRK_YN'].value_counts())
# Verificando tipos de dados
print("\nTipos de dados:")
print(data.dtypes)
# Verificando dados faltantes
print("\nDados faltantes por coluna:")
print(data.isnull().sum())
Valores únicos em DRK_YN: [1 0]
Contagem de classes em DRK_YN:
DRK_YN
0
     495858
     495488
Name: count, dtype: int64
Tipos de dados:
                      int32
sex
                    float64
age
                    float64
height
weight
                    float64
waistline
                    float64
sight_left
                    float64
sight_right
                    float64
hear_left
                    float64
hear_right
                    float64
SBP
                    float64
DBP
                    float64
BLDS
                    float64
tot_chole
                    float64
HDL_chole
                    float64
LDL_chole
                    float64
triglyceride
                    float64
hemoglobin
                    float64
urine_protein
                    float64
serum_creatinine
                    float64
SGOT\_AST
                    float64
SGOT_ALT
                    float64
gamma_GTP
                    float64
                      int64
SMK_stat_type_cd
                      int32
DRK_YN
dtype: object
Dados faltantes por coluna:
                    0
sex
```

```
0
    age
                        0
    height
    weight
                        0
    waistline
    sight left
                        0
    sight_right
                        0
    hear_left
    hear_right
    SBP
    DBP
                        0
    BLDS
                        0
                        0
    tot_chole
                        0
    HDL_chole
    LDL_chole
    triglyceride
    {\tt hemoglobin}
    urine_protein
                        0
    serum_creatinine
                        0
    SGOT_AST
    SGOT ALT
                        0
    gamma_GTP
                        0
    SMK_stat_type_cd
    DRK_YN
                        0
    dtype: int64
[4]: from sklearn.neighbors import KNeighborsClassifier
     from sklearn.ensemble import RandomForestClassifier
     from sklearn.neural_network import MLPClassifier
     from sklearn.svm import SVC
     def apply_knn(X_train, y_train, X_test, y_test):
         knn = KNeighborsClassifier(n_neighbors=5)
         knn.fit(X_train, y_train)
         return knn.score(X_test, y_test)
     def apply_random_forest(X_train, y_train, X_test, y_test):
         rf = RandomForestClassifier(n_estimators=100, verbose=True)
         rf.fit(X_train, y_train)
         return rf.score(X_test, y_test)
     def apply_mlp(X_train, y_train, X_test, y_test):
         mlp = MLPClassifier(hidden_layer_sizes=(100,), verbose=True,_
      →early_stopping=True)
         mlp.fit(X_train, y_train)
         return mlp.score(X_test, y_test)
     def apply_svm(X_train, y_train, X_test, y_test):
```

```
svm = SVC()
svm.fit(X_train, y_train)
return svm.score(X_test, y_test)
```

```
[8]: from concurrent.futures import ThreadPoolExecutor, as completed
           import logging
           from sklearn.metrics import confusion_matrix
           import matplotlib.pyplot as plt
           import seaborn as sns
            # Configurando o logging
           logging.basicConfig(level=logging.INFO, format='%(asctime)s - %(levelname)s -

√%(message)s')
           def train and evaluate(model, X_train, y_train, X_test, y_test, model_name,__
              ⇔dataset name):
                     logging.info(f"Iniciando treinamento do modelo {model name} para o dataset,
              model.fit(X_train, y_train)
                     accuracy = model.score(X_test, y_test)
                     y pred = model.predict(X test)
                     cm = confusion_matrix(y_test, y_pred)
                     logging.info(f"Treinamento concluído para o modelo {model_name} no datasetu
              return model_name, dataset_name, accuracy, cm
            # Definindo os modelos
           models = [
                     (KNeighborsClassifier(n_neighbors=5), "KNeighborsClassifier SMK"),
                     (RandomForestClassifier(n estimators=100), "RandomForestClassifier SMK"),
                     (MLPClassifier(hidden_layer_sizes=(64,16,4), verbose=True,_
              ⇔early_stopping=True), "MLPClassifier_SMK"),
                     #(SVC(), "SVC")
           ]
           # Criando tarefas para treinamento
           tasks = []
           for model, model_name in models:
                     tasks append((model, X_train_smk, y_train_smk, X_test_smk, y_test_smk, u_test_smk, y_test_smk, y_test_
              →model_name, "SMK"))
           # Executando treinamento em paralelo
           results = \Pi
           with ThreadPoolExecutor(max_workers=len(models)) as executor:
                     future_to_task = {executor.submit(train_and_evaluate, *task): task for task_
              →in tasks}
```

```
for future in as_completed(future_to_task):
        results.append(future.result())
# Plotando as matrizes de confusão
for model_name, dataset_name, accuracy, cm in results:
    plt.figure(figsize=(6, 5))
    sns.heatmap(cm, annot=True, fmt='d', cmap='Blues')
    plt.title(f"Matriz de Confusão para {model_name} - {dataset_name}\nAcurácia:

    {accuracy:.2f}")

    plt.xlabel('Previsão')
    plt.ylabel('Real')
    plt.show()
2023-12-05 19:13:17,028 - INFO - Iniciando treinamento do modelo
KNeighborsClassifier_SMK para o dataset SMK
2023-12-05 19:13:17,032 - INFO - Iniciando treinamento do modelo
RandomForestClassifier_SMK para o dataset SMK
2023-12-05 19:13:17,037 - INFO - Iniciando treinamento do modelo
MLPClassifier_SMK para o dataset SMK
Iteration 1, loss = 0.66513535
Validation score: 0.693511
Iteration 2, loss = 0.65147124
Validation score: 0.697609
Iteration 3, loss = 0.64913372
Validation score: 0.697571
Iteration 4, loss = 0.64812207
Validation score: 0.699904
Iteration 5, loss = 0.64730937
Validation score: 0.700093
Iteration 6, loss = 0.64678203
Validation score: 0.700119
Iteration 7, loss = 0.64639153
Validation score: 0.699337
Iteration 8, loss = 0.64597520
Validation score: 0.700018
Iteration 9, loss = 0.64577521
Validation score: 0.700661
Iteration 10, loss = 0.64556216
Validation score: 0.700182
Iteration 11, loss = 0.64534725
Validation score: 0.699299
Iteration 12, loss = 0.64520492
Validation score: 0.700509
Iteration 13, loss = 0.64498087
Validation score: 0.700535
Iteration 14, loss = 0.64490875
Validation score: 0.699173
```

Iteration 15, loss = 0.64474973

Validation score: 0.698517

Iteration 16, loss = 0.64476749

Validation score: 0.700081

Iteration 17, loss = 0.64462767

Validation score: 0.700711

Iteration 18, loss = 0.64446855

Validation score: 0.700459

Iteration 19, loss = 0.64442430

Validation score: 0.700068

Iteration 20, loss = 0.64433100

Validation score: 0.700030

Validation score did not improve more than to l=0.000100 for 10 consecutive

epochs. Stopping.

2023-12-05 19:17:45,372 - INFO - Treinamento concluído para o modelo

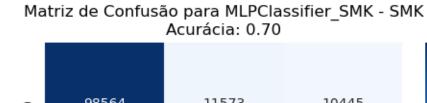
MLPClassifier_SMK no dataset SMK

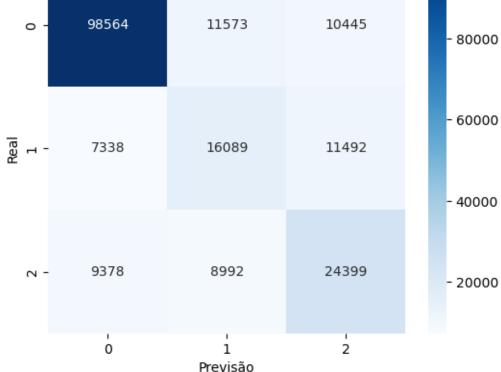
2023-12-05 19:22:57,954 - INFO - Treinamento concluído para o modelo

KNeighborsClassifier_SMK no dataset SMK

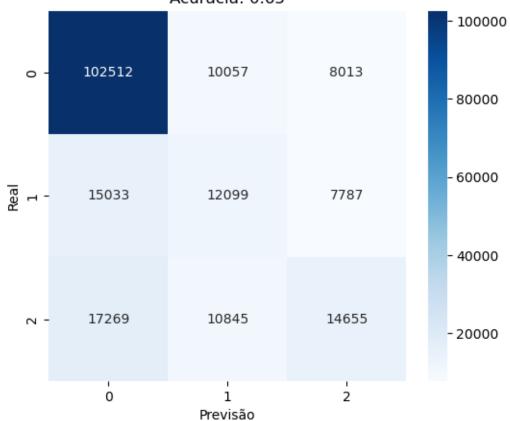
2023-12-05 19:26:07,210 - INFO - Treinamento concluído para o modelo

 ${\tt RandomForestClassifier_SMK}\ \ {\tt no}\ \ {\tt dataset}\ \ {\tt SMK}$

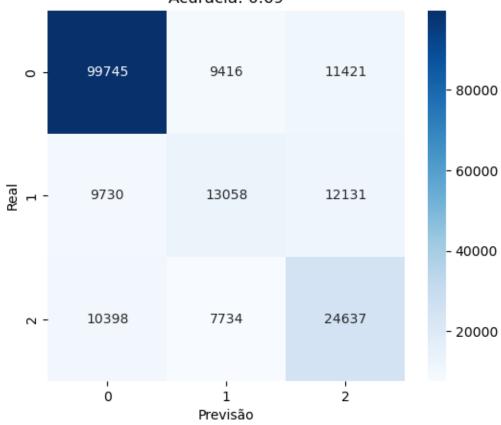




Matriz de Confusão para KNeighborsClassifier_SMK - SMK Acurácia: 0.65







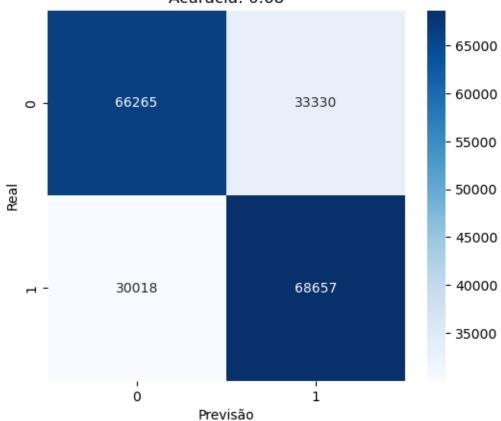
```
[9]: from concurrent.futures import ThreadPoolExecutor, as completed
     from sklearn.metrics import confusion_matrix
     import matplotlib.pyplot as plt
     import seaborn as sns
     def train_and_evaluate(model, X_train, y_train, X_test, y_test, model_name):
        print(f"Iniciando treinamento do modelo {model_name} para o dataset DRK")
        model.fit(X_train, y_train)
        accuracy = model.score(X_test, y_test)
        y_pred = model.predict(X_test)
        cm = confusion_matrix(y_test, y_pred)
        print(f"Treinamento concluído para o modelo {model_name} no dataset DRK")
        return model_name, accuracy, cm
     # Definindo os modelos
     models = [
         (KNeighborsClassifier(n_neighbors=5), "KNeighborsClassifier_DRK"),
         (RandomForestClassifier(n_estimators=100), "RandomForestClassifier_DRK"),
```

```
(MLPClassifier(hidden_layer_sizes=(64, 16, 4), max_iter=1000),
"MLPClassifier_DRK"),
#(SVC(), "SVC")
]

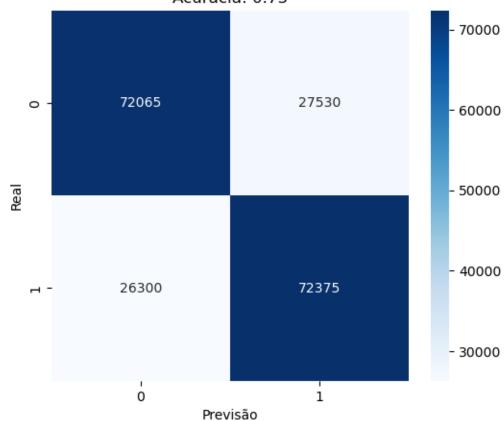
# Executando treinamento em paralelo
results = []
with ThreadPoolExecutor(max_workers=len(models)) as executor:
   future_to_model = {executor.submit(train_and_evaluate, model, X_train_drk, usy_train_drk, X_test_drk, y_test_drk, model_name): model_name for model,
model_name in models}
   for future in as_completed(future_to_model):
        results.append(future.result())
```

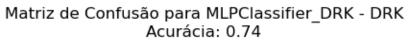
Iniciando treinamento do modelo KNeighborsClassifier_DRK para o dataset DRK Iniciando treinamento do modelo RandomForestClassifier_DRK para o dataset DRK Iniciando treinamento do modelo MLPClassifier_DRK para o dataset DRK Treinamento concluído para o modelo KNeighborsClassifier_DRK no dataset DRK Treinamento concluído para o modelo RandomForestClassifier_DRK no dataset DRK Treinamento concluído para o modelo MLPClassifier_DRK no dataset DRK

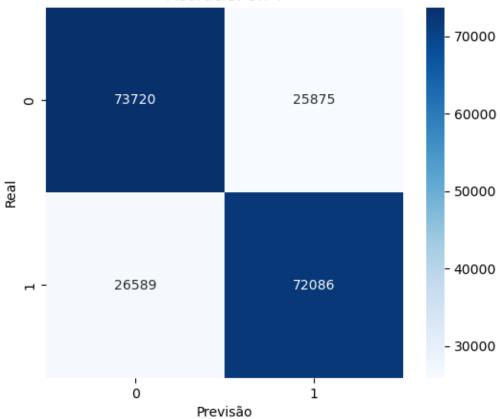
Matriz de Confusão para KNeighborsClassifier_DRK - DRK Acurácia: 0.68



Matriz de Confusão para RandomForestClassifier_DRK - DRK Acurácia: 0.73







[]: