STUDENTS PERFORMANCE IN EXAMS - ML FINAL PROJECT

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Introducción

El Aprendizaje Automático se ha convertido en una herramienta esencial en distintos ámbitos pues su variedad de técnicas han permitido desarrollar patrones y relaciones en conjuntos de datos grandes. Dada su importancia y relevancia, el objetivo de este trabajo es hacer uso de estas técnicas.

Para desarrollar este notebook se emplea un dataset, obtenido de kaggle, que contiene información sobre el rendimiento de estudiantes en tres tipos de exámenes: Examen de Matemáticas, Examen de Lectura y Examen de Escritura. Uno de los principales objetivos de este proyecto es predecir las calificaciones de los estudiantes en base a las características (features) de background que se le son atribuídas.

Entre las técnicas de implementación que se pueden encontrar en este proyecto están las siguientes:

- **Feature Selection:** con la finalidad de seleccionar un subconjunto de características relevantes y útiles a partir de un conjunto de características grande. El objetivo de feature selection es mejorar la precisión y eficiencia del modelo.
- Regresión Lineal Regularizada y Regresión lineal con Múltiples inputs y múltiples
 outputs: útiles para modelar datos en los que puede existir una relación lineal entre las
 variables de entrada y de salida del dataset.
- Repeated Stratified K-fold cross validation: debido a su utilidad en evaluar la capacidad de generalización de un modelo.
- Pipelines: para simplificar el proceso de construcción de modelos y poder atribuirles distintos parámetros.
- Redes Neuronales: con la finalidad de aprender patrones complejos en los datos e intentar alcanzar un alto nivel de predicción. En el contexto de este proyecto el objetivo de las redes neuronales era optimizar su rendimiento (optimizar las capas ocultas y el número de neuronas de cada capa). Para esto se realizaron las redes neuronales con Pipelines para cada una de las asignaturas usando las bibliotecas de Keras y Scikit-learn.
- Para la optimización se hizo la busqueda de hiperparámetros con GridSearchCV.

```
In [3]: import pandas as pd
   import numpy as np
   import matplotlib.pyplot as plt

# reading the csv file
   dataset = "https://raw.githubusercontent.com/Negatix092/Machine-Learning/main/ML_Cl
   df = pd.read_csv(dataset)

df.head()
```

Out[3]:

	gender	race/ethnicity	parental level of education	lunch	test preparation course	math score	reading score	writing score
) female	group B	bachelor's degree	standard	none	72	72	74
	1 female	group C	some college	standard	completed	69	90	88
2	2 female	group B	master's degree	standard	none	90	95	93
3	3 male	group A	associate's degree	free/reduced	none	47	57	44
4	4 male	group C	some college	standard	none	76	78	75

Dataset

Este dataset tiene un total de datos correspondiente a 1000 alumnos, con 5 características de los mismos. Los outputs que nos interesan son 3 y representan las últimas 3 filas del dataset: calificación de matemáticas, calificación de lectura y calificación de escritura.

Detalle del Dataset

- gender -- ['male', 'female']
- race/ethnicity -- ['group A', 'group B', 'group C', 'group D', 'group E']
- parental level of education -- ['some high school', 'high school', "bachelor's degree", 'some college', "master's degree", "associate's degree"]
- lunch -- ['standard', 'free/reduced']
- test preparation course -- ['none', 'completed']
- math score Student unique math score
- reading score Student unique reading score
- writing score Student unique writin score

```
In [4]:
        #preparacion del dataset
        from sklearn.metrics import classification report
        from sklearn.metrics import confusion matrix
        from sklearn.metrics import accuracy score
        from sklearn.metrics import roc auc score
        from sklearn.preprocessing import StandardScaler, LabelEncoder
        from sklearn.model_selection import train_test_split
        #outputs
        score_output_Math = 'math score'
        score_output_Reading = 'reading score'
        score_output_Writing = 'writing score'
        #en la lista binary feat se almacenan los nombres de las columnas (del dataset) que
        binary_feat = df.nunique()[df.nunique() == 2].keys().tolist()
        #en la lista numeric_feat se almacenan los nombres de las columnas (del dataset) qu
        numeric_feat = [col for col in df.select_dtypes(['float','int']).columns.tolist() i
        #en la lista categorical_feat se almacenan los nombres de las columnas (del dataset
        categorical_feat = [ col for col in df.select_dtypes('object').columns.to_list() if
        #una copa del dataframe (dataset) se almacena en un nuevo dataframe para trabajarlo
        df proc = df.copy()
        #Etiquetas para características binarias
        #aqui se transforman los binary features (solo dos valores diferentes) a valores nul
        le = LabelEncoder()
        for i in binary feat:
          df proc[i] = le.fit transform(df proc[i])
          print(i, '\n', np.unique(df_proc[i].values))
        def get_df_size(df, header='Dataset dimensions'):
          print(header,
                 '\n# Attributes: ', df.shape[1],
                 '\n# Entries: ', df.shape[0],'\n')
        #Dummy variables
        #se crean variables dummy para las categorical features
        #a cada categorical feature se la convierte en multiples binary features, con un va
        # es decir, one-hot encoding se aplica a las categorical features
        get df size(df proc, header='Processed dataset before dummies:')
        df_proc = pd.get_dummies(df_proc, columns=categorical_feat)
        get_df_size(df, header='Original dataset:')
        get_df_size(df_proc, header='Processed dataset:')
        df proc.head()
```

gender

[0 1]

lunch

[0 1]

test preparation course

[0 1]

Processed dataset before dummies:

Attributes: 8
Entries: 1000

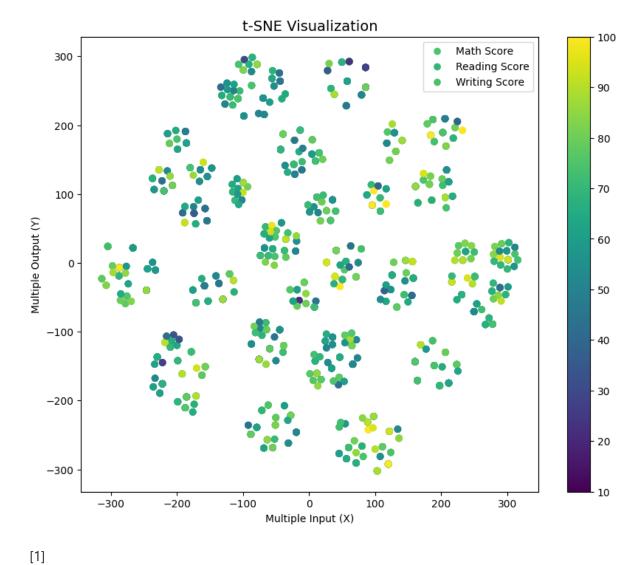
Original dataset: # Attributes: 8 # Entries: 1000

Processed dataset: # Attributes: 17 # Entries: 1000

Out[4]:

•		gender	lunch	test preparation course	math score	reading score	writing score	race/ethnicity_group A	race/ethnicity_group B
_	0	0	1	1	72	72	74	0	1
	1	0	1	0	69	90	88	0	0
	2	0	1	1	90	95	93	0	1
	3	1	0	1	47	57	44	1	0
	4	1	1	1	76	78	75	0	0

```
In [5]:
        import numpy as np
        from sklearn.manifold import TSNE
        import matplotlib.pyplot as plt
        # Seleccionamos las columnas que vamos a utilizar
        \#X = df_{proc.iloc[:, [0,1,2,6,7,8,9,10,11,12,13,14,15,16]].values
        x=df_proc
        X=df_proc.drop(columns =['math score','reading score','writing score'], axis=1)
        #y = df_proc.iloc[:, 3:6].values
        from scipy.stats import zscore
        X_normalized = X.apply(zscore)
        tsne = TSNE(n components=2, perplexity=30, learning rate=200)
        X_tsne = tsne.fit_transform(X_normalized)
        # Get the indices of the output columns
        output_indices = [3, 4, 5]
        output_labels = ['Math Score', 'Reading Score', 'Writing Score']
        # Plot the data points with different colors for each output column
        plt.figure(figsize=(10, 8))
        for i in range(len(output_indices)):
            idx = output_indices[i]
            il = output labels[i]
            plt.scatter(X_tsne[:, 0], X_tsne[:, 1], c=x.iloc[:, idx].values, cmap='viridis'
        plt.colorbar()
        plt.title('t-SNE Visualization', fontsize=14)
        plt.xlabel('Multiple Input (X)')
        plt.ylabel('Multiple Output (Y)')
        plt.legend()
        plt.show()
```



[2]

Punto 1: Multiple Input & Only One Output

```
In [6]: # inputs y outputs
        math_score_Y = df_proc['math score']
        reading_score_Y = df_proc['reading score']
        writing_score_Y = df_proc['writing score']
        all_X=df_proc.drop(columns =['math score', 'reading score', 'writing score'], axis=1)
        #all_Y=df_proc['math score', 'reading score', 'writing score']
        # Dividimos el conjunto de entrenamiento y test
        all_X_values_train = all_X[:-200]
        all_X_values_test = all_X[-200:] #last 200 for test
        #split the targets into training/testing sets
        math_score_Y_train = math_score_Y[:-200]
        math_score_Y_test = math_score_Y[-200:]
        #split the targets into training/testing sets
        writing_score_Y_train = writing_score_Y[:-200]
        writing_score_Y_test = writing_score_Y[-200:]
        #split the targets into training/testing sets
        reading_score_Y_train = reading_score_Y[:-200]
        reading_score_Y_test = reading_score_Y[-200:]
```

In [7]: all_X_values_train.shape

Out[7]: (800, 14)

```
In [8]: from sklearn.linear model import Lasso, Ridge
        from sklearn.feature selection import SelectFromModel
        from sklearn.pipeline import Pipeline
        from sklearn.model selection import RepeatedStratifiedKFold, GridSearchCV
        from sklearn.preprocessing import PolynomialFeatures, StandardScaler
        from sklearn.metrics import mean_squared_error
        from sklearn.metrics import r2_score
        from sklearn.feature selection import SelectKBest, f regression
        from scipy.stats import ttest rel
        #tenemos que usar pipelines, entonces cada tecnica de ML tendra su pipeline
        #como uno de los requisitos es que al menos una tecnica de ML sea Regresion Lineal
        #hacemos el pipe para Lasso
        # define pipelines
        pipeline_L = Pipeline([
            ('scaler', StandardScaler()),
            ('selector', SelectKBest(score_func=f_regression)),
            ('poly', PolynomialFeatures()),
            #('feature_selection', SelectFromModel(Lasso())),
            ('model', Lasso())
        ])
        #hacemos el pipe para Ridge, con el fin de compararlo con Lasso
        pipeline_R = Pipeline([
            ('scaler', StandardScaler()),
            ('selector', SelectKBest(score_func=f_regression)),
            ('poly', PolynomialFeatures()),
            #('feature_sel', SelectFromModel(Lasso())),
            ('model', Ridge())
        ])
        #estos parametros son necesarios para realizar la optimizacion del modelo(encontrar
        p_L = {'selector_k': range(1, 15),
               'poly__degree':[0,1,2],
               'model__alpha': [0.1, 1, 10, 100],
               'model__max_iter': [1000, 5000],
               'model tol': [1e-3, 1e-5]}
        p_R = {'selector__k': range(1, 15),
               'poly__degree':[0,1,2],
               'model__alpha': [0.1, 1, 10, 100],
               'model__max_iter': [1000, 5000],
               'model tol': [1e-3, 1e-5]}
        #repeated k-fold cross validation
        #Consulte qué quiere decir estratificado y por qué usted debe o no usarlo en su pro
        cv = RepeatedStratifiedKFold(n_splits=5, n_repeats=2, random_state=26)
        #se usa grid search para optimizar los parametros, el conjunto de validacion se uti
        lasso_search = GridSearchCV(pipeline_L, p_L, cv=cv, scoring='neg_mean_squared_error
        ridge_search = GridSearchCV(pipeline_R, p_R, cv=cv, scoring='neg_mean_squared_error
        #se hace el fit de cada regresion
```

```
result_lasso = lasso_search.fit(all_X_values_train, math_score_Y_train)
        result ridge = ridge search.fit(all X values train, math score Y train)
        #se quiere encontrar los mejores hiperparametros
        best_hp_lasso = result_lasso.best_params_
        best_hp_ridge = result_ridge.best_params_
        print("Los mejores hiperparámetros para Lasso son:", best_hp_lasso)
        print("Los mejores hiperparámetros para Ridge son:", best hp ridge)
        /usr/local/lib/python3.10/dist-packages/sklearn/model_selection/_split.py:700: User
        Warning: The least populated class in y has only 1 members, which is less than n_sp
        lits=5.
          warnings.warn(
        /usr/local/lib/python3.10/dist-packages/sklearn/model selection/ split.py:700: User
        Warning: The least populated class in y has only 1 members, which is less than n_sp
        lits=5.
          warnings.warn(
        /usr/local/lib/python3.10/dist-packages/sklearn/model_selection/_split.py:700: User
        Warning: The least populated class in y has only 1 members, which is less than n_sp
        lits=5.
          warnings.warn(
        /usr/local/lib/python3.10/dist-packages/sklearn/model_selection/_split.py:700: User
        Warning: The least populated class in y has only 1 members, which is less than n_sp
        lits=5.
          warnings.warn(
        Los mejores hiperparámetros para Lasso son: {'model__alpha': 0.1, 'model__max_iter
        ': 1000, 'model__tol': 1e-05, 'poly__degree': 1, 'selector__k': 14}
        Los mejores hiperparámetros para Ridge son: {'model__alpha': 10, 'model__max_iter':
        1000, 'model__tol': 0.001, 'poly__degree': 1, 'selector__k': 14}
In [9]: math_score_Y_train.shape
Out[9]: (800,)
        [3]
        [4]
        [5]
```

```
from sklearn.model selection import cross val score
#conociendo ya los modelos con sus mejores parametros, se vuelve a definir sus pipe
pipeline_L_best = Pipeline([
    ('scaler', StandardScaler()),
    ('selector', SelectKBest(score_func=f_regression, k=best_hp_lasso['selector__k'
    ('poly', PolynomialFeatures(degree=best_hp_lasso['poly__degree'])),
    ('model', Lasso(alpha=best_hp_lasso['model__alpha'], max_iter=best_hp_lasso['model', max_iter=best_hp_lasso['model']
    #('model', Lasso(alpha=best hp lasso['model alpha']))
])
#hacemos el pipe para Ridge, con el fin de compararlo con Lasso
pipeline_R_best = Pipeline([
    ('scaler', StandardScaler()),
    ('selector', SelectKBest(score_func=f_regression, k=best_hp_ridge['selector__k'
    ('poly', PolynomialFeatures(degree=best_hp_ridge['poly__degree'])),
    ('model', Ridge(alpha=best_hp_ridge['model__alpha'], max_iter=best_hp_ridge['mo
    #('model', Ridge(alpha=best_hp_ridge['model__alpha']))
1)
#se desea usar RepeatedStratifiedKfold crossvalidation
rskf = RepeatedStratifiedKFold(n splits=5,n repeats=2, random state=26)
lasso_scores = cross_val_score(pipeline_L_best, all_X_values_train, math_score_Y_tr
ridge scores = cross val score(pipeline R best, all X values train, math score Y tr
print("Lasso: %0.2f (+/- %0.2f)" % (lasso_scores.mean(), lasso_scores.std() * 2))
print("Ridge: %0.2f (+/- %0.2f)" % (ridge_scores.mean(), ridge_scores.std() * 2))
/usr/local/lib/python3.10/dist-packages/sklearn/model_selection/_split.py:700: User
Warning: The least populated class in y has only 1 members, which is less than n_sp
lits=5.
  warnings.warn(
/usr/local/lib/python3.10/dist-packages/sklearn/model_selection/_split.py:700: User
Warning: The least populated class in y has only 1 members, which is less than n_sp
lits=5.
  warnings.warn(
/usr/local/lib/python3.10/dist-packages/sklearn/model_selection/_split.py:700: User
Warning: The least populated class in y has only 1 members, which is less than n_sp
lits=5.
  warnings.warn(
Lasso: 0.22 (+/- 0.12)
Ridge: 0.22 (+/- 0.12)
/usr/local/lib/python3.10/dist-packages/sklearn/model_selection/_split.py:700: User
Warning: The least populated class in y has only 1 members, which is less than n_sp
lits=5.
  warnings.warn(
[5]
[6]
```

```
In [11]: from scipy.stats import wilcoxon
         #diferencia entre cross_val_scores de lasso y ridge
         score diff = lasso scores - ridge scores
         #con la diferencia se hace el test de wilcoxon
         statistic, pvalue = wilcoxon(score_diff)
         print("Test de Wilcoxon:")
         print(f"Statistic: {statistic}")
         print(f"P-value: {pvalue}")
         # interpret
         alpha = 0.05
         if pvalue > alpha:
             print('Same distribution (fail to reject H0)')
         else:
             print('Different distribution (reject H0)')
         Test de Wilcoxon:
         Statistic: 27.0
         P-value: 1.0
         Same distribution (fail to reject H0)
         [6]
In [12]: from scipy.stats import ttest_rel
         #Es posible usar ttest_rel en lugar de wilcoxon
         #se obtiene el t_value y el p_value
         #t_value es la diferencia entre las medias de los scores dividida por la desviacion
         #p_value es la probabilidad de que la diferencia entre las medias sea 0.05
         t_value, p_value = ttest_rel(lasso_scores, ridge_scores)
         print("t-value:", t_value)
         print("p-value:", p_value)
         # interpret
         alpha = 0.05
         if p_value > alpha:
             print('Same distribution (fail to reject H0)')
         else:
             print('Different distribution (reject H0)')
         t-value: -0.5374167752823499
         p-value: 0.6040111423997045
         Same distribution (fail to reject H0)
         [7]
```

```
In [19]:
         #se importan las metricas de sklearn
         from sklearn.metrics import mean squared error, r2 score
         #se reentrena con los mejores hiperparametros encontrados
         pipeline_L_best.fit(all_X_values_train, math_score_Y_train)
         pipeline_R_best.fit(all_X_values_train, math_score_Y_train)
         # Make predictions on the test set for both models
         lasso preds = pipeline L best.predict(all X values test)
         ridge_preds = pipeline_R_best.predict(all_X_values_test)
         # Calculate the MSE for both models
         lasso_mse = mean_squared_error(math_score_Y_test, lasso_preds)
         ridge_mse = mean_squared_error(math_score_Y_test, ridge_preds)
         # Calculate the R2 score for both models
         lasso_r2 = r2_score(math_score_Y_test, lasso_preds)
         ridge_r2 = r2_score(math_score_Y_test, ridge_preds)
         # Calculate the RMSE for both models
         lasso_rmse = mean_squared_error(math_score_Y_test, lasso_preds, squared=False)
         ridge_rmse = mean_squared_error(math_score_Y_test, ridge_preds, squared=False)
         print("Lasso metrics")
         print(f"\nMSE: {lasso_mse}")
         print(f"R2 score: {lasso r2}")
         print(f"RMSE: {lasso rmse}")
         print("\nRidge metrics")
         print(f"\nMSE: {ridge_mse}")
         print(f"R2 score: {ridge_r2}")
         print(f"RMSE: {ridge_rmse}")
         #se grafica el error
         plt.figure(figsize=(10, 10))
         plt.scatter(math_score_Y_test, ridge_preds, c='crimson')
         plt.yscale('log')
         plt.xscale('log')
         p1 = np.amax([np.amax(ridge preds), np.amax(math score Y test)])
         p2 = np.amin([np.amin(ridge_preds), np.amin(math_score_Y_test)])
         plt.plot([p1, p2], [p1, p2], 'b-', label = 'Ideal Line')
         plt.xlabel('True values', fontsize=15)
         plt.ylabel('Predictions', fontsize=15)
         plt.axis('equal')
         plt.legend(fontsize=15)
         plt.show()
```

Lasso metrics

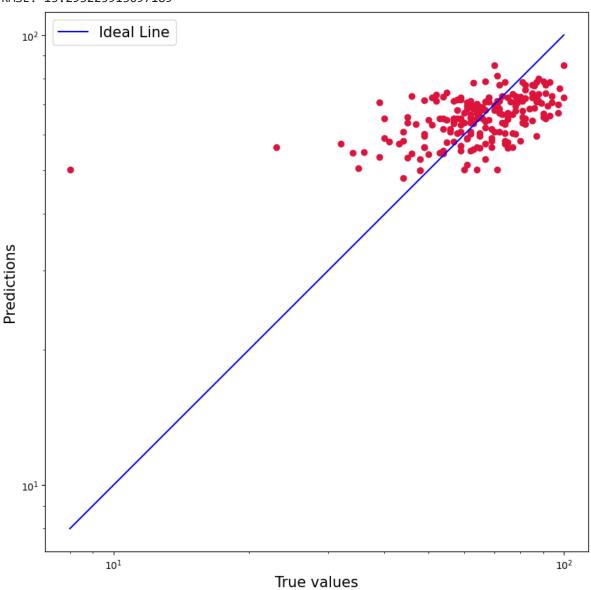
MSE: 176.91955337171063

R2 score: 0.27480097814514415 RMSE: 13.301110982610085

Ridge metrics

MSE: 176.70985517663857

R2 score: 0.27566053788884015 RMSE: 13.293225913097189

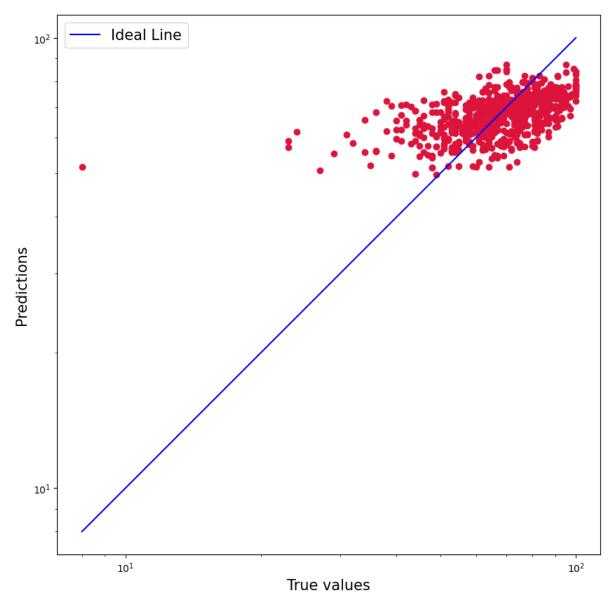


Punto 2: Multiple Inputs & Multiple Outputs

```
In [14]: # inputs y outputs
         math_score_Y = df_proc['math score']
         reading_score_Y = df_proc['reading score']
         writing_score_Y = df_proc['writing score']
         all_X=df_proc.drop(columns =['math score', 'reading score', 'writing score'], axis=1)
         all_Y = df_proc.iloc[:, 3:6].values
         # Dividimos el conjunto de entrenamiento y test
         all_X_values_train = all_X[:-200]
         all_X_values_test = all_X[-200:] #last 200 for test
         #split the targets into training/testing sets
         #math_score_Y_train = math_score_Y[:-200]
         #math_score_Y_test = math_score_Y[-200:]
         all_Y_values_train = all_Y[:-200]
         all_Y_values_test = all_Y[-200:] #last 200 for test
In [15]: all_Y_values_train.shape
Out[15]: (800, 3)
```

```
from sklearn.model selection import RepeatedStratifiedKFold
from sklearn.pipeline import Pipeline
from sklearn.multioutput import MultiOutputRegressor
from sklearn.linear model import Ridge
from sklearn.feature selection import SelectKBest
from sklearn.model_selection import GridSearchCV
from sklearn.preprocessing import PolynomialFeatures, StandardScaler
from sklearn.feature selection import SelectKBest, f regression
# Create a pipeline for the Ridge regression with feature selection
ridge pipeline = Pipeline([
    ('scaler', StandardScaler()),
    ('selector', SelectKBest(score_func=f_regression)),
    ('poly', PolynomialFeatures()),
    ('regressor', Ridge())
])
# Create a dictionary of hyperparameters for the pipeline
ridge_hyperparameters = {
    'estimator selector k': range(1, 15),
    'estimator__poly__degree':[0,1,2],
    'estimator__regressor__alpha': [0.1, 1.0, 10.0, 100]
}
# Create a MultiOutputRegressor wrapper around the Ridge pipeline for multiple outp
multi output ridge pipeline = MultiOutputRegressor(ridge pipeline)
cvm = RepeatedStratifiedKFold(n_splits=5, n_repeats=2, random_state=26)
# Create a grid search object with cross-validation for hyperparameter tuning
ridge_grid_search = GridSearchCV(multi_output_ridge_pipeline, ridge_hyperparameters
result ridge = ridge grid search.fit(all X values train, all Y values train)
#se quiere encontrar los mejores hiperparametros
best_hp_mridge = result_ridge.best_params_
print("Los mejores hiperparámetros para Multi-Ridge son:", best hp mridge)
Los mejores hiperparámetros para Multi-Ridge son: {'estimator_poly_degree': 1, 'e
stimator__regressor__alpha': 10.0, 'estimator__selector__k': 14}
[8]
[9]
```

```
In [18]:
         #conociendo los mejores hiperparametros del modelo, se crea el modelo con esos hipe
         best mridge pipeline = Pipeline([
             ('scaler', StandardScaler()),
             ('selector', SelectKBest(score func=f regression, k=14)),
             ('poly', PolynomialFeatures(degree=1)),
             ('regressor', Ridge(alpha=100))
         ])
         # Create a MultiOutputRegressor wrapper around the Ridge pipeline for multiple outp
         best_multi_output_ridge_pipeline = MultiOutputRegressor(best_mridge_pipeline)
         #se entrena el modelo
         best_multi_output_ridge_pipeline.fit(all_X_values_train, all_Y_values_train)
         #se hacen las predicciones
         all_Y_values_pred = best_multi_output_ridge_pipeline.predict(all_X_values_test)
         #se calcula el error
         from sklearn.metrics import mean_squared_error
         from sklearn.metrics import mean absolute error
         from sklearn.metrics import r2 score
         import math
         print("Multi-Ridge metrics")
         print("\nMean squared error: %.2f" % mean_squared_error(all_Y_values_test, all_Y_va
         print("RMean squared error: %.2f" % math.sqrt(mean squared error(all Y values test,
         print("Mean absolute error: %.2f" % mean_absolute_error(all_Y_values_test, all_Y_va
         print('R2 score: %.2f' % r2_score(all_Y_values_test, all_Y_values_pred))
         #se grafica el error
         plt.figure(figsize=(10, 10))
         plt.scatter(all Y values test, all Y values pred, c='crimson')
         plt.yscale('log')
         plt.xscale('log')
         p1 = np.amax([np.amax(all_Y_values_pred), np.amax(all_Y_values_test)])
         p2 = np.amin([np.amin(all_Y_values_pred), np.amin(all_Y_values_test)])
         plt.plot([p1, p2], [p1, p2], 'b-', label = 'Ideal Line')
         plt.xlabel('True values', fontsize=15)
         plt.ylabel('Predictions', fontsize=15)
         plt.axis('equal')
         plt.legend(fontsize=15)
         plt.show()
         Multi-Ridge metrics
         Mean squared error: 175.31
         RMean squared error: 13.24
         Mean absolute error: 10.74
         R2 score: 0.26
```



```
In [22]: #se calcula el pvalue usando el Wilcoxon test
mstatistic, mp_value = wilcoxon(ridge_preds,all_Y_values_pred[:, 0])#solo considera

print("Test de Wilcoxon:")
print(f"Statistic: {mstatistic}")
print(f"P-value: {mp_value}")

# interpret
alpha = 0.05
if mp_value > alpha:
    print('Same distribution (fail to reject H0)')
else:
    print('Different distribution (reject H0)')

Test de Wilcoxon:
Statistic: 9478.0
P-value: 0.4852093110442015
Same distribution (fail to reject H0)
```

```
In [27]: # Reshape ridge_preds to have the same shape as ridge_preds
    ridge_preds_reshaped = ridge_preds.reshape(-1, 1).repeat(3, axis=1)

# Calculate the t-statistic and p-value for the paired t-test
    mt_statistic, mtp_value = ttest_rel(ridge_preds_reshaped, all_Y_values_pred)

print("t-value:", mt_statistic)
    print("p-value:", mtp_value)

# interpret
    alpha = 0.05
    if np.amax(mtp_value) > alpha:
        print('Same distribution (fail to reject H0)')

else:
        print('Different distribution (reject H0)')

t-value: [-1.04124813 -7.59575454 -4.05461538]
    p-value: [2.99023769e-01 1.16872685e-12 7.20329786e-05]
Same distribution (fail to reject H0)
```

Punto 3: Neural Networks

Optimizando el número de capas ocultas y el número de neuronas

```
In []: # inputs y outputs
    math_score_Y = df_proc['math score']
    reading_score_Y = df_proc['reading score']
    writing_score_Y = df_proc['writing score']

all_X=df_proc.drop(columns =['math score','reading score','writing score'], axis=1)

all_Y = df_proc.iloc[:, 3:6].values

# Dividimos el conjunto de entrenamiento y test
all_X_values_train = all_X[:-200]
all_X_values_test = all_X[-200:] #last 200 for test

# split the targets into training/testing sets
# math_score_Y_train = math_score_Y[:-200]
# math_score_Y_test = math_score_Y[:-200:]
all_Y_values_train = all_Y[:-200:] #last 200 for test
```

```
In [28]: from keras.models import Sequential
         from keras.layers import Dense
         from sklearn.pipeline import Pipeline
         from sklearn.model selection import GridSearchCV, train test split, RepeatedStratif
         import numpy as np
         from keras.wrappers.scikit_learn import KerasRegressor
         from sklearn.metrics import get_scorer_names
         from sklearn.metrics import mean squared error, make scorer
         # normalize data to have zero mean and std one (feature scaling)
         from sklearn.preprocessing import StandardScaler
         scaler = StandardScaler()
         scaler.fit(all_X_values_train)# scaler fit is just for training dataset
         all_X_values_train_normalized = scaler.transform(all_X_values_train) #normalize tra
         # Crear un modelo de Keras para utilizar en la búsqueda de Grid
         def create model(neurons=1, hidden layers=1):
             model = Sequential()
             model.add(Dense(neurons, input_dim=all_X_values_train_normalized.shape[1], acti
             # Agregar una cantidad de capas ocultas igual a
             # hidden layers, cada una con 'neurons' neuronas
             for i in range(hidden_layers):
                 model.add(Dense(neurons, activation='relu'))
             model.add(Dense(3, activation='linear'))
             model.compile(loss='mean_squared_error', optimizer='adam')
             return model
         # Crear un pipeline para el modelo de Keras
         model_math = Pipeline([
             ('keras', KerasRegressor(build_fn=create_model, verbose=0))
         ])
         print(get scorer names())
         # Definir una lista de los valores que se probarán para
         # el número de neuronas y el número de capas ocultas
         param_grid = {'keras__neurons': [5, 15, 25, 50, 75], 'keras__hidden_layers': [1, 2,
         # Aplicar el GridSearchCV con Pipeline y el RepeatedStratifiedKFold
         grid_math = GridSearchCV(estimator=model_math, param_grid=param_grid, verbose=4, sc
         grid_result = grid_math.fit(all_X_values_train_normalized, all_Y_values_train)
         # Mostrar los resultados de la búsqueda de Grid para cada materia
         print("Para el modelo de neural networks:")
         means = grid_result.cv_results_['mean_test_score']
         stds = grid_result.cv_results_['std_test_score']
         params = grid_result.cv_results_['params']
         print(f"Mejor: {grid_result.best_score_} usando {grid_result.best_params_}")
```

<ipython-input-28-d3b80db17790>:30: DeprecationWarning: KerasRegressor is deprecate
d, use Sci-Keras (https://github.com/adriangb/scikeras) instead. See https://www.ad
riangb.com/scikeras/stable/migration.html for help migrating.
 ('keras', KerasRegressor(build_fn=create_model, verbose=0))

```
['accuracy', 'adjusted_mutual_info_score', 'adjusted_rand_score', 'average_precisio
n', 'balanced_accuracy', 'completeness_score', 'explained_variance', 'f1', 'f1_macr
o', 'f1_micro', 'f1_samples', 'f1_weighted', 'fowlkes_mallows_score', 'homogeneity_
score', 'jaccard', 'jaccard_macro', 'jaccard_micro', 'jaccard_samples', 'jaccard_we
ighted', 'matthews_corrcoef', 'max_error', 'mutual_info_score', 'neg_brier_score',
'neg_log_loss', 'neg_mean_absolute_error', 'neg_mean_absolute_percentage_error', 'n
eg_mean_gamma_deviance', 'neg_mean_poisson_deviance', 'neg_mean_squared_error', 'ne
g_mean_squared_log_error', 'neg_median_absolute_error', 'neg_negative_likelihood_ra
tio', 'neg_root_mean_squared_error', 'normalized_mutual_info_score', 'positive like
lihood_ratio', 'precision', 'precision_macro', 'precision_micro', 'precision_sample
s', 'precision_weighted', 'r2', 'rand_score', 'recall', 'recall_macro', 'recall_mic
ro', 'recall_samples', 'recall_weighted', 'roc_auc', 'roc_auc_ovo', 'roc_auc_ovo_we
ighted', 'roc_auc_ovr', 'roc_auc_ovr_weighted', 'top_k_accuracy', 'v_measure_score
Fitting 5 folds for each of 15 candidates, totalling 75 fits
[CV 1/5] END keras_hidden_layers=1, keras_neurons=5;, score=-65.623 total time=
[CV 2/5] END keras_hidden_layers=1, keras_neurons=5;, score=-68.438 total time=
[CV 3/5] END keras_hidden_layers=1, keras_neurons=5;, score=-67.399 total time=
[CV 4/5] END keras_hidden_layers=1, keras_neurons=5;, score=-68.406 total time=
1.2s
[CV 5/5] END keras__hidden_layers=1, keras__neurons=5;, score=-68.375 total time=
0.9s
[CV 1/5] END keras_hidden_layers=1, keras_neurons=15;, score=-65.054 total time=
0.9s
[CV 2/5] END keras_hidden_layers=1, keras_neurons=15;, score=-68.586 total time=
[CV 3/5] END keras__hidden_layers=1, keras__neurons=15;, score=-67.001 total time=
0.9s
[CV 4/5] END keras_hidden_layers=1, keras_neurons=15;, score=-68.322 total time=
0.9s
[CV 5/5] END keras_hidden_layers=1, keras_neurons=15;, score=-68.363 total time=
0.9s
[CV 1/5] END keras__hidden_layers=1, keras__neurons=25;, score=-64.915 total time=
0.9s
[CV 2/5] END keras_hidden_layers=1, keras_neurons=25;, score=-68.243 total time=
[CV 3/5] END keras_hidden_layers=1, keras_neurons=25;, score=-66.773 total time=
0.9s
[CV 4/5] END keras__hidden_layers=1, keras__neurons=25;, score=-67.947 total time=
[CV 5/5] END keras_hidden_layers=1, keras_neurons=25;, score=-67.869 total time=
0.9s
[CV 1/5] END keras hidden_layers=1, keras neurons=50;, score=-64.718 total time=
[CV 2/5] END keras__hidden_layers=1, keras__neurons=50;, score=-67.806 total time=
1.2s
[CV 3/5] END keras_hidden_layers=1, keras_neurons=50;, score=-66.741 total time=
0.9s
[CV 4/5] END keras__hidden_layers=1, keras__neurons=50;, score=-67.854 total time=
0.9s
[CV 5/5] END keras__hidden_layers=1, keras__neurons=50;, score=-67.099 total time=
[CV 1/5] END keras_hidden_layers=1, keras_neurons=75;, score=-64.317 total time=
```

```
0.9s
[CV 2/5] END keras_hidden_layers=1, keras_neurons=75;, score=-67.243 total time=
0.9s
[CV 3/5] END keras_hidden_layers=1, keras_neurons=75;, score=-65.907 total time=
0.9s
[CV 4/5] END keras_hidden_layers=1, keras_neurons=75;, score=-66.944 total time=
0.8s
[CV 5/5] END keras_hidden_layers=1, keras_neurons=75;, score=-66.948 total time=
0.9s
[CV 1/5] END keras_hidden_layers=2, keras_neurons=5;, score=-65.539 total time=
1.0s
[CV 2/5] END keras_hidden_layers=2, keras_neurons=5;, score=-68.731 total time=
1.5s
[CV 3/5] END keras_hidden_layers=2, keras_neurons=5;, score=-66.947 total time=
[CV 4/5] END keras_hidden_layers=2, keras_neurons=5;, score=-67.896 total time=
1.5s
[CV 5/5] END keras_hidden_layers=2, keras_neurons=5;, score=-68.295 total time=
1.1s
[CV 1/5] END keras_hidden_layers=2, keras_neurons=15;, score=-65.137 total time=
1.1s
[CV 2/5] END keras_hidden_layers=2, keras_neurons=15;, score=-68.373 total time=
1.0s
[CV 3/5] END keras_hidden_layers=2, keras_neurons=15;, score=-67.094 total time=
1.0s
[CV 4/5] END keras_hidden_layers=2, keras_neurons=15;, score=-67.948 total time=
1.0s
[CV 5/5] END keras_hidden_layers=2, keras_neurons=15;, score=-68.251 total time=
1.0s
[CV 1/5] END keras__hidden_layers=2, keras__neurons=25;, score=-64.797 total time=
1.1s
[CV 2/5] END keras_hidden_layers=2, keras_neurons=25;, score=-68.213 total time=
1.0s
[CV 3/5] END keras_hidden_layers=2, keras_neurons=25;, score=-67.067 total time=
1.1s
[CV 4/5] END keras__hidden_layers=2, keras__neurons=25;, score=-68.144 total time=
1.3s
[CV 5/5] END keras_hidden_layers=2, keras_neurons=25;, score=-67.952 total time=
2.2s
[CV 1/5] END keras_hidden_layers=2, keras_neurons=50;, score=-64.246 total time=
1.2s
[CV 2/5] END keras__hidden_layers=2, keras__neurons=50;, score=-67.576 total time=
1.0s
[CV 3/5] END keras hidden_layers=2, keras neurons=50;, score=-66.104 total time=
1.0s
[CV 4/5] END keras_hidden_layers=2, keras_neurons=50;, score=-66.702 total time=
1.0s
[CV 5/5] END keras_hidden_layers=2, keras_neurons=50;, score=-67.249 total time=
1.1s
[CV 1/5] END keras hidden layers=2, keras neurons=75;, score=-63.445 total time=
1.0s
[CV 2/5] END keras__hidden_layers=2, keras__neurons=75;, score=-66.700 total time=
1.1s
[CV 3/5] END keras_hidden_layers=2, keras_neurons=75;, score=-64.734 total time=
[CV 4/5] END keras_hidden_layers=2, keras_neurons=75;, score=-65.790 total time=
```

```
1.0s
[CV 5/5] END keras_hidden_layers=2, keras_neurons=75;, score=-64.579 total time=
[CV 1/5] END keras_hidden_layers=3, keras_neurons=5;, score=-65.354 total time=
1.7s
[CV 2/5] END keras_hidden_layers=3, keras_neurons=5;, score=-68.561 total time=
1.3s
[CV 3/5] END keras_hidden_layers=3, keras_neurons=5;, score=-67.293 total time=
[CV 4/5] END keras_hidden_layers=3, keras_neurons=5;, score=-68.288 total time=
1.8s
[CV 5/5] END keras_hidden_layers=3, keras_neurons=5;, score=-68.313 total time=
1.3s
[CV 1/5] END keras_hidden_layers=3, keras_neurons=15;, score=-65.007 total time=
[CV 2/5] END keras_hidden_layers=3, keras_neurons=15;, score=-68.560 total time=
1.2s
[CV 3/5] END keras_hidden_layers=3, keras_neurons=15;, score=-66.744 total time=
[CV 4/5] END keras_hidden_layers=3, keras_neurons=15;, score=-68.116 total time=
[CV 5/5] END keras hidden_layers=3, keras neurons=15;, score=-68.141 total time=
1.7s
[CV 1/5] END keras_hidden_layers=3, keras_neurons=25;, score=-65.199 total time=
1.2s
[CV 2/5] END keras_hidden_layers=3, keras_neurons=25;, score=-68.389 total time=
1.2s
[CV 3/5] END keras_hidden_layers=3, keras_neurons=25;, score=-67.223 total time=
1.2s
[CV 4/5] END keras_hidden_layers=3, keras_neurons=25;, score=-67.923 total time=
1.2s
[CV 5/5] END keras_hidden_layers=3, keras_neurons=25;, score=-68.122 total time=
1.2s
[CV 1/5] END keras_hidden_layers=3, keras_neurons=50;, score=-63.488 total time=
1.8s
[CV 2/5] END keras_hidden_layers=3, keras_neurons=50;, score=-67.268 total time=
1.3s
[CV 3/5] END keras_hidden_layers=3, keras_neurons=50;, score=-65.200 total time=
1.4s
[CV 4/5] END keras_hidden_layers=3, keras_neurons=50;, score=-66.717 total time=
1.8s
[CV 5/5] END keras__hidden_layers=3, keras__neurons=50;, score=-67.242 total time=
[CV 1/5] END keras_hidden_layers=3, keras_neurons=75;, score=-63.143 total time=
1.2s
[CV 2/5] END keras hidden_layers=3, keras neurons=75;, score=-64.026 total time=
1.2s
[CV 3/5] END keras_hidden_layers=3, keras_neurons=75;, score=-64.456 total time=
1.2s
[CV 4/5] END keras_hidden_layers=3, keras_neurons=75;, score=-64.089 total time=
1.2s
[CV 5/5] END keras__hidden_layers=3, keras__neurons=75;, score=-64.923 total time=
1.2s
Para el modelo de neural networks:
Mejor: -64.12741714941959 usando {'keras_hidden_layers': 3, 'keras_neurons': 75}
```

[10]

Conclusiones

Regresión Lineal Regularizada (Multiple Inputs - One Output)

Tras hacer uso de Pipelines y GridSearchCV se concluye que los mejores hiperparámetros para el modelo de regresión son:

- Lasso -> α = 0.1, iteraciones = 1000, tolerancia: 1e-05, grado del polinomio = 1, features selected = 14.
- Ridge -> α = 10, iteraciones = 1000, tolerancia: 1e-03, grado del polinomio = 1, features selected = 14.

Tras hacer uso del test de Wilcoxon y del TTest, se concluye que, dado α = 0.05 y una hipótesis nula de que no existe evidencia de una diferencia significativa entre modelos:

- Wilcoxon -> para un p-value de 1.0 (> α), no se rechaza la hipótesis nula.
- Ttest -> para un p-value de 0.6040 (> α), no se rechaza la hipótesis nula.

Regresión Lineal (Multiple Inputs - Multiple Outputs)

Tras hacer uso de Pipelines y GridSearchCV se concluye que los mejores hiperparámetros para el modelo de multi-regresión con múltiples outputs son:

• Ridge -> α = 10, iteraciones = 1000, tolerancia: 1e-03, grado del polinomio = 1, features selected = 14.

Tras hacer uso del test de Wilcoxon y del TTest, se concluye que, dado α = 0.05 y una hipótesis nula de que no existe evidencia de una diferencia significativa entre modelos:

- Wilcoxon -> para un p-value de 0.4852 (> α), no se rechaza la hipótesis nula.
- Ttest -> para un p-value (maximo) de 0.2990 (> α), no se rechaza la hipótesis nula.

Redes Neuronales

Para este proyecto el objetivo de las redes neuronales era optimizar su rendimiento (optimizar las capas ocultas y el número de neuronas de cada capa. Para esto se realizaron las redes neuronales con Pipelines para cada una de las asignaturas usando las bibliotecas de Keras y Scikit-learn. Para la optimización se hizo la busqueda de hiperparámetros con GridSearchCV. Validamos el modelo utilizando el Kfold-cross validation.

Cuando optimizamos las capas ocultas y el número de neuronas. Probamos con 1, 2 o 3 capas ocultas. Cada una con 5, 15, 25, 50 y 75 neuronas. El resultado varió en el número de capas de la red de cada asignatura (matemáticas con 1, reading con 2 y writing con 3) mientras que el número de neuronas se mantuvo óptimo en 75. Como se trataba de un

problema de regresión, la capa de salida solo fue una que tiene como función de activación la función sigmoide. El optimizador que usamos fue Adam, que tras prueba y error, fue el que mejor rendimiento tuvo. Como loss function tuvimos el mse que mide el error promedio entre las predicciones que hace el modelo con las etiquetas.

Finalmente, obtuvimos los resultados de la busqueda de los mejores hiperparámetros (# de capas y # de neuronas) y realizamos la prueba de hipótesis para hallar el valor p.

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