Import the libraries

```
import numpy as np
import pandas as pd
from sklearn.linear_model import SGDRegressor
import matplotlib.pyplot as plt
import random
from sklearn.model_selection import learning_curve, cross_val_score
```

Set the seeds

```
In [22]: seed = 2377
    np.random.seed(seed)
    random.seed(seed)
```

Read the data

```
In [23]: df = pd.read_csv("Valhalla23.csv")
    x = df["Celsius"]
    y = df["Valks"]
    df
```

Out[23]:		Celsius	Valks
	0	61.4720	-139.7400
	1	70.5790	-156.6000
	2	-7.3013	73.2690
	3	71.3380	-165.4200
	4	43.2360	-75.8350
	•••		•••
	95	-7.0094	69.6320
	96	36.8820	-71.2400
	97	26.9390	-34.2550
	98	-18.8100	106.4300
	99	13.7120	9.1011

100 rows × 2 columns

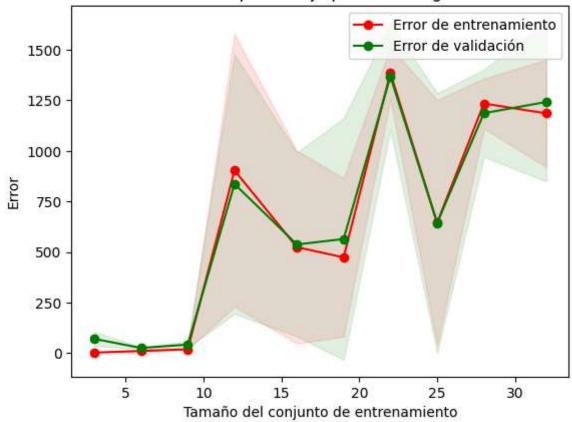
Split the data: training (40%), validation (40%), and test (20%) sets

```
In [24]: train = df.sample(frac=0.8, random_state=seed)

df = df.drop(train.index)
  test = df
```

```
validation = train.sample(frac=0.5, random_state=seed)
         train = train.drop(validation.index)
         len(train), len(validation), len(test)
Out[24]: (40, 40, 20)
In [25]: learning_rate = 1e-4
         model = SGDRegressor(
             max iter=1 000 000,
             learning rate='constant',
             eta0=learning rate,
             random_state=seed
         X train = train.drop(columns=["Valks"])
         y train = train["Valks"]
         X_validation = validation.drop(columns=["Valks"])
         y validation = validation["Valks"]
         X_test = test.drop(columns=["Valks"])
         y test = test["Valks"]
In [26]: # Curva de aprendizaje
         train_sizes, train_scores, val_scores = learning_curve(model, X_train, y_train, cv=
         # Calcular el promedio y la desviación estándar
         train_mean = -np.mean(train_scores, axis=1)
         val_mean = -np.mean(val_scores, axis=1)
         train_std = np.std(train_scores, axis=1)
         val_std = np.std(val_scores, axis=1)
         # Graficar la curva de aprendizaje
         plt.plot(train_sizes, train_mean, 'o-', color="r", label="Error de entrenamiento")
         plt.plot(train_sizes, val_mean, 'o-', color="g", label="Error de validación")
         plt.fill_between(train_sizes, train_mean - train_std, train_mean + train_std, alpha
         plt.fill_between(train_sizes, val_mean - val_std, val_mean + val_std, alpha=0.1, co
         plt.title('Curva de Aprendizaje para SGDRegressor')
         plt.xlabel('Tamaño del conjunto de entrenamiento')
         plt.ylabel('Error')
         plt.legend(loc="best")
         plt.show()
         # Validación cruzada
         cv_scores = cross_val_score(model, X_train, y_train, cv=5, scoring='neg_mean_square
         print(f'Error de validación cruzada (MSE): {-cv_scores.mean():.3f} +/- {cv_scores.s
```

Curva de Aprendizaje para SGDRegressor



Error de validación cruzada (MSE): 1242.342 +/- 392.501

Podemos observar un grado considerable de sesgo.

Caculate the mse for each set of data

```
In [27]: model.fit(X_train, y_train)
    mse_train = np.mean((model.predict(X_train) - y_train) ** 2)
    mse_validation = np.mean((model.predict(X_validation) - y_validation) ** 2)
    mse_test = np.mean((model.predict(X_test) - y_test) ** 2)
```

```
In [28]: # Create subplots: 1 row, 3 columns
    fig, axs = plt.subplots(1, 3, figsize=(15, 5))

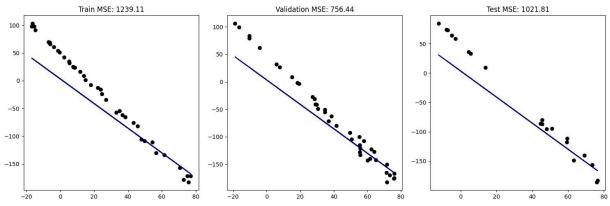
# Train subplot
    axs[0].scatter(X_train, y_train, color="black", label="Train")
    axs[0].plot(X_train, model.predict(X_train), color="navy", label="Model")
    axs[0].set_title(f"Train MSE: {mse_train:.2f}")

# Validation subplot
    axs[1].scatter(X_validation, y_validation, color="black", label="Validation")
    axs[1].plot(X_validation, model.predict(X_validation), color="navy", label="Model")
    axs[1].set_title(f"Validation MSE: {mse_validation:.2f}")
```

```
# Test subplot
axs[2].scatter(X_test, y_test, color="black", label="Test")
axs[2].plot(X_test, model.predict(X_test), color="navy", label="Model")
axs[2].set_title(f"Test MSE: {mse_test:.2f}")

# Adjust layout to prevent overlap
plt.tight_layout()

# Show the plot
plt.show()
```

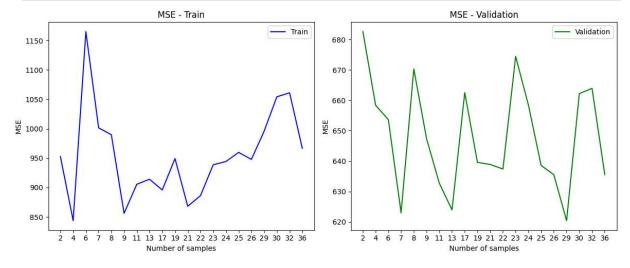


Create a list with numbers between 2 and 40

```
In [29]: numbers = list(range(2, 40))
         # Make sure the number 2 is in the list
         final_list = [2]
         numbers.remove(2)
         final_list += random.sample(numbers, 19)
In [30]: final_list, len(final_list)
Out[30]: ([2, 32, 24, 23, 8, 11, 36, 29, 30, 13, 19, 25, 9, 7, 21, 4, 26, 22, 17, 6],
           20)
In [31]: dict_of_models = {}
         number_models = 100
         for i in final_list:
             models = []
             for model_i in range(number_models):
                 # get a subset of the data from train with the amount of values of i
                 train_subset = train.sample(i, random_state=seed)
                 X_train_subset = train_subset.drop(columns=["Valks"])
                 y_train_subset = train_subset["Valks"]
                 model = SGDRegressor(
                     max_iter=1_000_000,
```

```
learning_rate='constant',
                     eta0=learning_rate,
                  model.fit(X_train, y_train)
                  # Calculate the MSE for the train subset
                  mse_train = np.mean((model.predict(X_train_subset) - y_train_subset) ** 2)
                  # Calculate the MSE for the validation data
                  mse_validation = np.mean((model.predict(X_validation) - y_validation) ** 2)
                  # create a tuple with the model and the mse
                  models.append((model, mse_train, mse_validation))
             dict of models[i] = models
In [32]: # verify the length of the dictionary and the length of the models
         len(dict of models), len(dict of models[2])
Out[32]: (20, 100)
In [33]: # calculate the mean of the mse (train and validation) for each model
         mean_mse = \{\}
         for i in dict of models:
             mean mse[i] = (
                  np.mean([model[1] for model in dict_of_models[i]]),
                  np.mean([model[2] for model in dict_of_models[i]])
             )
         mean_mse # (train mse, validation mse)
Out[33]: {2: (952.6308962791061, 682.6383632930281),
           32: (1061.069290800353, 663.9462800688453),
           24: (944.2521791746015, 658.4850621604311),
           23: (938.6499492302502, 674.4428513447735),
           8: (989.8553832415412, 670.3631150015405),
           11: (905.5652042596297, 632.7420098096321),
           36: (966.7077881822895, 635.5431897315545),
           29: (995.6851455708608, 620.3944372325577),
           30: (1054.2714429732255, 662.209184891686),
           13: (914.1369454402145, 623.9393140097679),
           19: (949.3150822963935, 639.5769253416728),
           25: (959.782982951397, 638.5654907734894),
           9: (856.1029843279877, 647.4112306529497),
           7: (1001.5294963074614, 622.9554009240298),
           21: (868.1717973994014, 638.8877486287344),
           4: (843.7353098521007, 658.36737886336),
           26: (947.8543847967858, 635.5150733617105),
           22: (886.1824792263442, 637.3848570644147),
           17: (895.8928849812838, 662.5266081245579),
           6: (1165.143360530994, 653.6753034252697)}
```

```
In [34]: # Sort the keys of mean mse
         sorted keys = sorted(mean mse.keys())
         # Extract MSE values for train and validation in the order of sorted keys
         mse_train = [mean_mse[i][0] for i in sorted_keys]
         mse_validation = [mean_mse[i][1] for i in sorted_keys]
         # Create subplots: 1 row, 2 columns
         fig, (ax1, ax2) = plt.subplots(1, 2, figsize=(12, 5))
         # Plot mse_train on the first subplot
         ax1.plot(mse_train, label="Train", color='blue')
         ax1.set xlabel("Number of samples")
         ax1.set ylabel("MSE")
         ax1.set title("MSE - Train")
         ax1.set_xticks(range(len(mse_train)))
         ax1.set xticklabels(sorted keys)
         ax1.legend()
         # Plot mse validation on the second subplot
         ax2.plot(mse_validation, label="Validation", color='green')
         ax2.set_xlabel("Number of samples")
         ax2.set ylabel("MSE")
         ax2.set_title("MSE - Validation")
         ax2.set_xticks(range(len(mse_validation)))
         ax2.set_xticklabels(sorted_keys)
         ax2.legend()
         # Adjust layout for better display
         plt.tight layout()
         plt.show()
```



Model Adjustment

- 2 samples: Not bad MSE in the training set but way worse in the validation set because it practially learned nothing.
- 36 samples: Seems to be at an equilibrium point. The model is not overfitting or underfitting. And also, the both training and validation MSE have the same movement.

How the adjustment change as the sample size increases?

- The ideal size for achieving the best results is the train MSE is 4, 9 or 21 samples. But this is not ideal for the validation set because the model might now generalize well.
- The ideal size for achieving the best results is the validation MSE is 29 samples. This is way better because the model can generalize well and predict better unseen data.

What is the best size for the model trainig?

• Around 29 samples. As explained before, this is the best size for the validation set and the model can generalize well.

Bias and variance:

- Model with 2 samples: Low bias and high variance due to the overfitting. Basically a
 good fit for the training set but not for the validation set.
- Model with 40 samples: Seems to be in a good equilibrium point. The model is not overfitting or underfitting. And also, the both training and validation MSE have the same movement.

We can see the improvement in the model in terms of the equilibrium between bias and variance as the sample size increases.

For other models:

- Models with 4 and 9 samples: both have low bias and high variance. The model is overfitting and not generalizing well. There is a high difference between the training and validation MSE.
- Model with 7 samples: low variance and high bias. The model is underfitting and not generalizing well

Train the model with the new sample size

```
In [35]: df = pd.read_csv("Valhalla23.csv")
x = df["Celsius"]
y = df["Valks"]

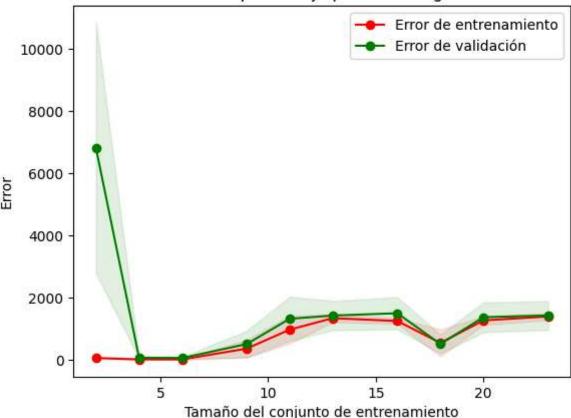
In [36]: train = df.sample(frac=0.8, random_state=seed)

df = df.drop(train.index)
test = df

validation = train.sample(frac=0.5, random_state=seed)
train = train.drop(validation.index)
len(train), len(validation), len(test)
```

```
In [37]: # just use 29 samples for the training
         train = train.sample(29, random_state=seed)
In [38]: learning_rate = 1e-4
         model = SGDRegressor(
             max iter=1 000 000,
             learning_rate='constant',
             eta0=learning_rate,
             random state=seed
         X train = train.drop(columns=["Valks"])
         y train = train["Valks"]
         X_validation = validation.drop(columns=["Valks"])
         y_validation = validation["Valks"]
         X test = test.drop(columns=["Valks"])
         y_test = test["Valks"]
In [39]: model.fit(X train, y train)
         mse_train = np.mean((model.predict(X_train) - y_train) ** 2)
         mse_validation = np.mean((model.predict(X_validation) - y_validation) ** 2)
         mse test = np.mean((model.predict(X test) - y test) ** 2)
In [40]: # Curva de aprendizaje
         train_sizes, train_scores, val_scores = learning_curve(model, X_train, y_train, cv=
         # Calcular el promedio y la desviación estándar
         train_mean = -np.mean(train_scores, axis=1)
         val_mean = -np.mean(val_scores, axis=1)
         train_std = np.std(train_scores, axis=1)
         val_std = np.std(val_scores, axis=1)
         # Graficar la curva de aprendizaje
         plt.plot(train_sizes, train_mean, 'o-', color="r", label="Error de entrenamiento")
         plt.plot(train_sizes, val_mean, 'o-', color="g", label="Error de validación")
         plt.fill_between(train_sizes, train_mean - train_std, train_mean + train_std, alpha
         plt.fill_between(train_sizes, val_mean - val_std, val_mean + val_std, alpha=0.1, co
         plt.title('Curva de Aprendizaje para SGDRegressor')
         plt.xlabel('Tamaño del conjunto de entrenamiento')
         plt.ylabel('Error')
         plt.legend(loc="best")
         plt.show()
         # Validación cruzada
         cv_scores = cross_val_score(model, X_train, y_train, cv=5, scoring='neg_mean_square
         print(f'Error de validación cruzada (MSE): {-cv_scores.mean():.3f} +/- {cv_scores.s
```

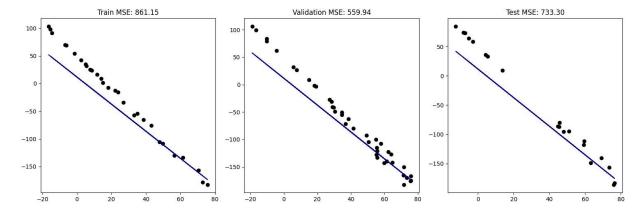
Curva de Aprendizaje para SGDRegressor



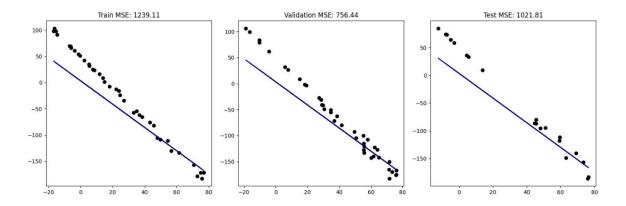
Error de validación cruzada (MSE): 1425.647 +/- 459.991

Results for the new model with 29 samples

```
In [41]: # Create subplots: 1 row, 3 columns
         fig, axs = plt.subplots(1, 3, figsize=(15, 5))
         # Train subplot
         axs[0].scatter(X_train, y_train, color="black", label="Train")
         axs[0].plot(X_train, model.predict(X_train), color="navy", label="Model")
         axs[0].set_title(f"Train MSE: {mse_train:.2f}")
         # Validation subplot
         axs[1].scatter(X_validation, y_validation, color="black", label="Validation")
         axs[1].plot(X_validation, model.predict(X_validation), color="navy", label="Model")
         axs[1].set_title(f"Validation MSE: {mse_validation:.2f}")
         # Test subplot
         axs[2].scatter(X_test, y_test, color="black", label="Test")
         axs[2].plot(X_test, model.predict(X_test), color="navy", label="Model")
         axs[2].set_title(f"Test MSE: {mse_test:.2f}")
         # Adjust layout to prevent overlap
         plt.tight_layout()
         # Show the plot
         plt.show()
```



Results of the base model



Comparison

As you can see, the new model with 29 samples is way better. The new model:

- Has a lower MSE for the training set.
- Has a lower MSE for the validation set.
- Has a lower MSE for the test set.

As we expected, the model with 29 samples improved the results because now the model can generalize way better.