## Lab 8

```
In [ ]: ID = 1631938
Name = 'MIN SOE HTUT'
```

# 2 Define the data generator

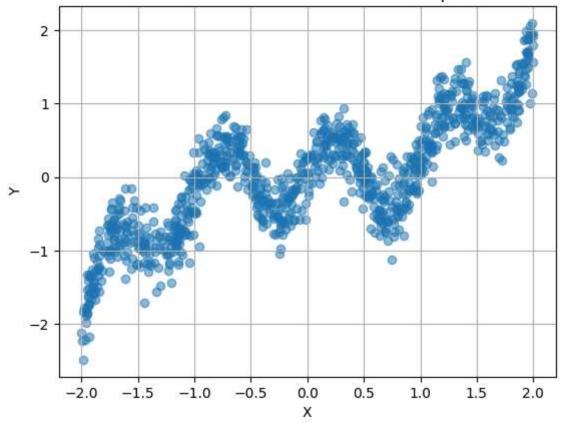
```
In []: import numpy as np

def ground_truth_concept(x):
    y=0.25*x**3+0.5*np.sin(np.pi*2*x)
    return y

def data_generator(n_examples,seed=314159265):
    np.random.seed(seed)
    x=(np.random.rand(n_examples)-0.5)*4
    np.random.seed(seed+1)
    y=ground_truth_concept(x)+np.random.randn(n_examples)*0.25
    x=x.reshape(n_examples,1)
    return x,y
```

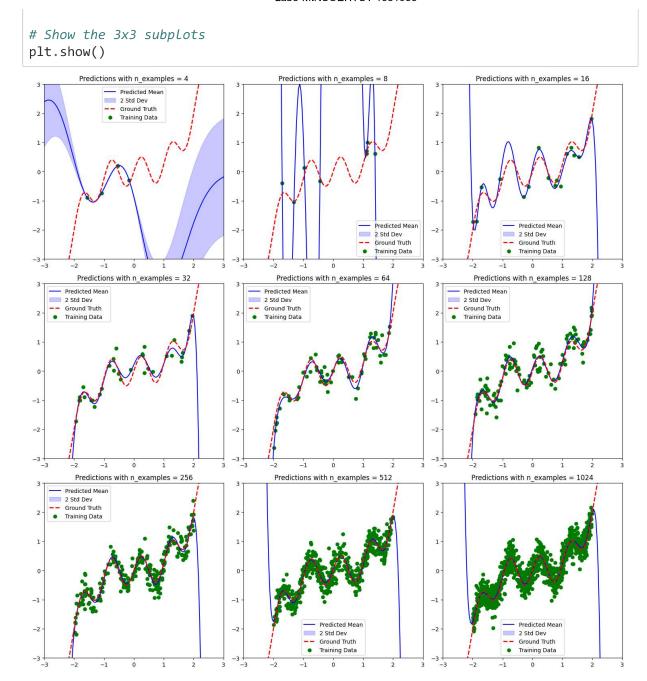
3 Generate 1000 samples using data\_generator and visualize it using a scatter plot

# Scatter Plot of 1000 Generated Samples



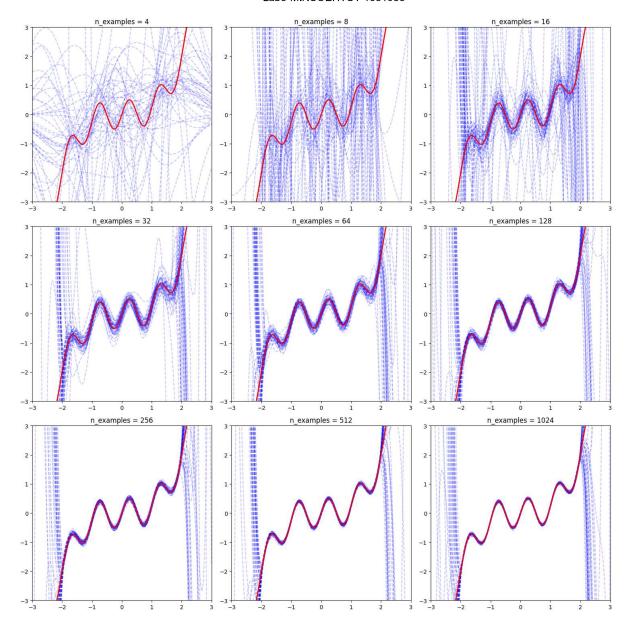
4 Do a 3x3 subplot, in each subplot, you will plot the graphs for one of each results of for n\_examples=4,8,16,32,64,128,256,512,1024

```
In [ ]: from sklearn.gaussian_process import GaussianProcessRegressor
        from sklearn.gaussian process.kernels import RBF
        import numpy as np
        import matplotlib.pyplot as plt
        # Define n examples values as specified
        n_examples_values = [4, 8, 16, 32, 64, 128, 256, 512, 1024]
        # Set up a 3x3 subplot layout
        fig, axes = plt.subplots(3, 3, figsize=(15, 15))
        # Define X test to evaluate the predictions on a fine grid
        X_{\text{test}} = \text{np.linspace}(-3, 3, 151).reshape}(-1, 1)
        y_test = ground_truth_concept(X_test)
        # Loop through each n examples value and generate subplots
        for i, n_examples in enumerate(n_examples_values):
            # Get the current subplot
            ax = axes[i // 3, i % 3]
            # Generate the data
            x_data, y_data = data_generator(n_examples, ID + i)
            # Fit the GaussianProcessRegressor (with the default RBF kernel)
            gp = GaussianProcessRegressor()
            gp.fit(x data, y data)
            # Make predictions on X test and get the standard deviations
            y pred, sigma = gp.predict(X test, return std=True)
            # Plot the predicted mean
            ax.plot(X_test, y_pred, 'b-', label='Predicted Mean')
            # Plot the 2 standard deviation intervals
            ax.fill_between(X_test[:, 0], y_pred - 2 * sigma, y_pred + 2 * sigma, colo
        r='blue', alpha=0.2, label='2 Std Dev')
            # Overlay the ground truth concept
            ax.plot(X test, y test, 'r--', label='Ground Truth', linewidth=2)
            # Scatter plot of the training data
            ax.scatter(x data, y data, color='green', marker='o', label='Training Dat
        a')
            # Set the plot limits for x and y
            ax.set xlim(-3, 3)
            ax.set_ylim(-3, 3)
            # Set the title for the current subplot
            ax.set_title(f'Predictions with n_examples = {n_examples}')
            # Add the legend for all subplots
            ax.legend()
        # Adjust layout to prevent overlapping subplots
        plt.tight layout()
```



5. Do a 3x3 subplot, in each subplot, you will plot the graphs for one of each results for  $n_examples=4,8,16,32,64,128,256,512,1024$ 

```
In [ ]: from sklearn.gaussian_process import GaussianProcessRegressor
        from sklearn.gaussian process.kernels import RBF
        import numpy as np
        import matplotlib.pyplot as plt
        # Define n examples values
        n_examples_values = [4, 8, 16, 32, 64, 128, 256, 512, 1024]
        # Set up a 3x3 subplot layout
        fig, axes = plt.subplots(3, 3, figsize=(15, 15))
        # Define X test for prediction and the ground truth concept
        X_{\text{test}} = \text{np.linspace}(-3, 3, 151).reshape}(-1, 1)
        y_test = ground_truth_concept(X_test)
        # Loop through each n examples value and generate subplots
        for i, n examples in enumerate(n examples values):
            # Get the current subplot
            ax = axes[i // 3, i % 3]
            # Repeat the data generation and GPR fitting 50 times
            for j in range(50):
                # Generate data for each iteration (vary the seed for randomness)
                x data, y data = data generator(n examples, ID + i * 50 + j)
                # Fit the GaussianProcessRegressor (with the default RBF kernel)
                gp = GaussianProcessRegressor()
                gp.fit(x_data, y_data)
                # Make predictions on X test
                y pred = gp.predict(X test)
                # Plot each mean prediction as a dashed line (set alpha to make them s
        emi-transparent)
                ax.plot(X_test, y_pred, 'b--', alpha=0.2)
            # Overlay the ground truth concept in red
            ax.plot(X_test, y_test, 'r-', linewidth=2, label='Ground Truth')
            # Set the plot limits for x and y
            ax.set_xlim(-3, 3)
            ax.set ylim(-3, 3)
            # Set the title for the current subplot
            ax.set title(f'n examples = {n examples}')
        # Adjust layout to prevent overlapping subplots
        plt.tight layout()
        # Show the 3x3 subplots
        plt.show()
```



Part B

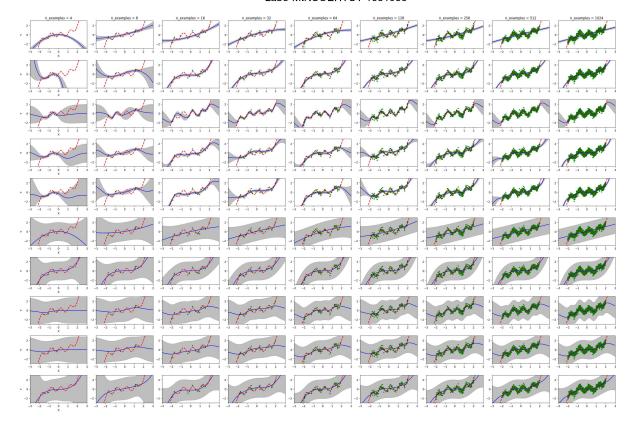
# 1) Define the Kernels

```
In [26]:
         from sklearn.gaussian_process.kernels import DotProduct, WhiteKernel, Matern,
         RBF, Exponentiation
         # Define the kernels
         kernels = [
             Exponentiation(DotProduct(), 2) + WhiteKernel(noise_level=0.01, noise_leve
         l_bounds='fixed'), # Polynomial degree 2 + fixed White noise 0.01
             Exponentiation(DotProduct(), 3) + WhiteKernel(noise level=0.01, noise leve
         l_bounds='fixed'), # Polynomial degree 3 + fixed White noise 0.01
             Matern(length_scale=1.0, length_scale_bounds='fixed') + WhiteKernel(noise_
         level=0.01, noise_level_bounds='fixed'), # Matern with fixed length scale
             RBF(length scale=1.0, length scale bounds='fixed') + WhiteKernel(noise lev
         el=0.01, noise_level_bounds='fixed'), # RBF with fixed Length_scale
             Exponentiation(DotProduct(), 2) + RBF(length_scale=1.0, length_scale_bound
         s='fixed') + WhiteKernel(noise_level=0.01, noise_level_bounds='fixed'), # Pol
         ynomial degree 2 + RBF + fixed length scale
             Exponentiation(DotProduct(), 2) + WhiteKernel(noise_level=1.0, noise_level
         bounds='fixed'), # Polynomial degree 2 + fixed White noise 1.0
             Exponentiation(DotProduct(), 3) + WhiteKernel(noise level=1.0, noise level
         _bounds='fixed'),  # Polynomial degree 3 + fixed White noise 1.0
             Matern(length scale=1.0, length scale bounds='fixed') + WhiteKernel(noise
         level=1.0, noise_level_bounds='fixed'), # Matern with fixed Length_scale
             RBF(length scale=1.0, length scale bounds='fixed') + WhiteKernel(noise lev
         el=1.0, noise_level_bounds='fixed'), # RBF with fixed length_scale
             Exponentiation(DotProduct(), 3) + RBF(length scale=1.0, length scale bound
         s='fixed') + WhiteKernel(noise level=1.0, noise level bounds='fixed') # Polyn
         omial degree 3 + RBF + fixed Length scale
```

2) For each of the kernels in part B.1)Doing a 3x3 subplot, in each subplot, you will plot the graphs for one of each results of for n\_examples=4,8,16,32,64,128,256,512,1024

```
In [27]:
         from sklearn.gaussian_process import GaussianProcessRegressor
          import numpy as np
          import matplotlib.pyplot as plt
         # Define X_test for generating predictions and the ground truth
         X_{\text{test}} = \text{np.linspace}(-3, 3, 151).reshape}(-1, 1)
         y_{test} = 0.25 * X_{test**3} + 0.5 * np.sin(np.pi * 2 * X_{test})
          # Define n_examples values as specified
         n_examples_values = [4, 8, 16, 32, 64, 128, 256, 512, 1024]
          # Kernel names for easy identification in the plots
          kernel names = [
              "Poly (deg 2) + White noise 0.01",
              "Poly (deg 3) + White noise 0.01",
             "Matern + White noise 0.01",
              "RBF + White noise 0.01",
              "Poly (deg 2) + RBF + White noise 0.01",
              "Poly (deg 2) + White noise 1.0",
             "Poly (deg 3) + White noise 1.0",
              "Matern + White noise 1.0",
              "RBF + White noise 1.0",
              "Poly (deg 3) + RBF + White noise 1.0"
          ]
         # Function to generate training data (based on a provided seed)
         def data generator(n examples, seed):
             np.random.seed(seed)
             x = (np.random.rand(n examples) - 0.5) * 4 # Random x-values in range (-
          2, 2)
             np.random.seed(seed + 1)
             y = 0.25 * x**3 + 0.5 * np.sin(np.pi * 2 * x) + np.random.randn(n example)
          s) * 0.25 # Noisy data
             x = x.reshape(n examples, 1)
              return x, y
          # Function to plot for each kernel
          def plot for kernel(kernel, kernel name, axes, row idx):
              # Loop through each n examples value
              for i, n examples in enumerate(n examples values):
                  ax = axes[row idx, i]
                  # Generate the data
                  x data, y data = data generator(n examples, ID + i)
                  # Fit the GaussianProcessRegressor with the current kernel
                  gp = GaussianProcessRegressor(kernel=kernel)
                  gp.fit(x_data, y_data)
                  # Predict the mean and standard deviations
                  y pred, sigma = gp.predict(X test, return std=True)
                  # Plot the mean predictions
                  ax.plot(X_test, y_pred, 'b-', label='Predicted Mean')
                  # Plot the 2 standard deviation intervals
```

```
ax.fill_between(X_test.flatten(), y_pred - 2 * sigma, y_pred + 2 * sig
ma, color='gray', alpha=0.5, label='2 Std Dev')
        # Overlay the ground truth concept
        ax.plot(X_test, y_test, 'r--', label='Ground Truth', linewidth=2)
        # Scatter plot of the training data
        ax.scatter(x_data, y_data, color='green', marker='.', label='Training
Data')
        # Set plot limits
        ax.set_xlim(-3, 3)
        ax.set ylim(-3, 3)
        # Set the title for the current subplot
        if row_idx == 0:
            ax.set title(f'n examples = {n examples}')
    # Add labels and legend only to the first column and row
    if row idx == len(kernels) - 1:
        for ax in axes[:, 0]:
            ax.set(xlabel='X', ylabel='Y')
# Create the large figure for all subplots
fig, axes = plt.subplots(len(kernels), len(n_examples_values), figsize=(30, 2
0))
# Loop through all kernels and generate plots
for idx, kernel in enumerate(kernels):
    plot_for_kernel(kernel, kernel_names[idx], axes, idx)
# Ensure the layout is tight and not overlapping
plt.tight_layout()
plt.show()
```



Part C

#### a) The behavoir of the regressors as the number of examples used to estimate the regressors increase

As seen in the plots, when the number of examples is small n\_examples = 4 or n\_examples = 8 the model exhibits large uncertainty shown by wider 2-standard-deviation intervals. The predictions fluctuate and deviate more significantly from the ground truth indicating that the model has insufficient data to make confident predictions. As the number of examples increases the predictions become more accurate and the uncertainty bands narrow. By n\_examples = 1024, the regressors closely fit the ground truth, and the variance in predictions decreases dramatically. This demonstrates the model's improved confidence with more training data.

#### b) The extrapolation (the estimated curve outside the region of the data) behavior of the regressors

The extrapolation behavior is highly dependent on the kernel. We observe that the RBF and Matern kernels extrapolate more smoothly outside the range of the training data. These kernels revert to the mean and produce reasonable consistent predictions outside the data region. Polynomial kernels with higher degrees exhibit more erratic behavior when extrapolating showing large deviations from the ground truth in the extrapolated regions. This is particularly evident when the data is sparse where polynomial kernels tend to overfit within the data range and fail to generalize beyond it.

## c) The ability for the regressor to "capture" the wriggly behaivoir in the middle of the data

The RBF and Matern kernels perform well in capturing the wriggly behavior in the middle of the data .These kernels are designed to handle smooth and continuous variations, which makes them well-suited to model the oscillations in the data.Polynomial kernels especially lower-degree ones struggle to capture these local variations. They impose global trends leading to a more linear or smooth fit that does not accurately reflect the wriggly nature of the data. Higher degree polynomials can capture some of the wiggles but are prone to overfitting particularly when the data is sparse.

#### d) The amount of estimated (white) noise epsilon

The white noise component significantly affects the model ability to handle deviations in the data. Models with higher noise levels show wider uncertainty intervals indicating that the model is accounting for more stochastic variation. This results in smoother predictions as the model does not attempt to fit every small fluctuation in the data. When the noise level is small or fixed the model fits the data more tightly reducing the apparent noise in the predictions. This is evident in the plots where smaller noise levels lead to tighter confidence intervals and more precise fits.