# Project ARI3205 Interpretable AI for Deep Learning Models (Part 2)

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# **Importing Necessary Libraries**

```
In [1]: # Check and install required libraries from the libraries.json file
        import json
        # Read the libraries from the text file
        with open('../Libraries/Part2_Lib.json', 'r') as file:
            libraries = json.load(file)
        # ANSI escape codes for colored output
        GREEN = "\033[92m" # Green text
        RED = "\033[91m" # Red text]
        RESET = "\033[0m" # Reset to default color
        # Function to check and install libraries
        def check_and_install_libraries(libraries):
            for lib, import_name in libraries.items():
                try:
                    # Attempt to import the library
                     __import__(import_name)
                    print(f"[{GREEN}√{RESET}] Library '{lib}' is already installed.")
                except ImportError:
                    # If import fails, try to install the library
                    print(f"[{RED}X*{RESET}] Library '{lib}' is not installed. Installing...
                    %pip install {lib}
        # Execute the function to check and install libraries
        check_and_install_libraries(libraries)
        # Import necessary libraries for data analysis and modeling
        import warnings
        import pandas as pd
        import numpy as np
        import matplotlib.pyplot as plt
        from sklearn.model_selection import train_test_split
        from tensorflow.keras.models import Sequential
        from tensorflow.keras.layers import Dense, Input
        from tensorflow.keras.optimizers import Adam
        from sklearn.preprocessing import StandardScaler, OneHotEncoder
        from sklearn.impute import SimpleImputer
        from sklearn.neural_network import MLPClassifier
        import lime
        from lime.lime tabular import LimeTabularExplainer
        import shap
        from anchor import anchor_tabular
```

```
# Suppress specific warnings
        warnings.filterwarnings("ignore", message="X does not have valid feature names")
        warnings.filterwarnings("ignore", category=RuntimeWarning)
        warnings.filterwarnings("ignore", category=UserWarning)
        [√] Library 'tensorflow' is already installed.
        [\checkmark] Library 'matplotlib' is already installed.
        [\checkmark] Library 'pandas' is already installed.
        \left[ 
ightharpoonup 
ight] Library 'numpy' is already installed.
        [✓] Library 'lime' is already installed.
        C:\Users\Sean Muscat\AppData\Local\Packages\PythonSoftwareFoundation.Python.3.11 qb
        z5n2kfra8p0\LocalCache\local-packages\Python311\site-packages\tqdm\auto.py:21: Tqdm
        Warning: IProgress not found. Please update jupyter and ipywidgets. See https://ipy
        widgets.readthedocs.io/en/stable/user_install.html
          from .autonotebook import tqdm as notebook_tqdm
        [√] Library 'shap' is already installed.
        [√] Library 'anchor' is already installed.
In [2]: # Define the filenames
        train_filename = '../Datasets/Titanic/train.csv'
        test_filename = '../Datasets/Titanic/test.csv'
        gender_submission_filename = '../Datasets/Titanic/gender_submission.csv'
        # Load the datasets
        try:
            train_data = pd.read_csv(train_filename)
            test_data = pd.read_csv(test_filename)
             gender_submission_data = pd.read_csv(gender_submission_filename)
            print(f"'{train_filename}' dataset loaded successfully.")
            print(f"'{test filename}' dataset loaded successfully.")
            print(f"'{gender_submission_filename}' dataset loaded successfully.")
        except FileNotFoundError as e:
            print(f"Error: {e.filename} was not found. Please ensure it is in the correct di
            exit()
        except pd.errors.EmptyDataError as e:
            print(f"Error: {e.filename} is empty.")
        except pd.errors.ParserError as e:
            print(f"Error: There was a problem parsing {e.filename}. Please check the file f
            exit()
        # Dataset insights
        print("\nTrain Dataset Overview:")
        print(train data.info())
        print("\nTrain Dataset Statistical Summary:")
        print(train_data.describe())
        print("\nTest Dataset Overview:")
        print(test_data.info())
        print("\nTest Dataset Statistical Summary:")
        print(test_data.describe())
        print("\nGender Submission Dataset Overview:")
        print(gender submission data.info())
```

- '../Datasets/Titanic/train.csv' dataset loaded successfully.
- '../Datasets/Titanic/test.csv' dataset loaded successfully.
- '../Datasets/Titanic/gender\_submission.csv' dataset loaded successfully.

#### Train Dataset Overview:

<class 'pandas.core.frame.DataFrame'>
RangeIndex: 891 entries, 0 to 890

Data columns (total 12 columns):

#	Column	Non-Null Count	Dtype	
0	PassengerId	891 non-null	int64	
1	Survived	891 non-null	int64	
2	Pclass	891 non-null	int64	
3	Name	891 non-null	object	
4	Sex	891 non-null	object	
5	Age	714 non-null	float64	
6	SibSp	891 non-null	int64	
7	Parch	891 non-null	int64	
8	Ticket	891 non-null	object	
9	Fare	891 non-null	float64	
10	Cabin	204 non-null	object	
11	Embarked	889 non-null	object	
dtypes: $float64(2)$ int64(5) object(5)				

dtypes: float64(2), int64(5), object(5)

memory usage: 83.7+ KB

None

#### Train Dataset Statistical Summary:

	PassengerId	Survived	Pclass	Age	SibSp	\
count	891.000000	891.000000	891.000000	714.000000	891.000000	
mean	446.000000	0.383838	2.308642	29.699118	0.523008	
std	257.353842	0.486592	0.836071	14.526497	1.102743	
min	1.000000	0.000000	1.000000	0.420000	0.000000	
25%	223.500000	0.000000	2.000000	20.125000	0.000000	
50%	446.000000	0.000000	3.000000	28.000000	0.000000	
75%	668.500000	1.000000	3.000000	38.000000	1.000000	
max	891.000000	1.000000	3.000000	80.000000	8.000000	

	Parch	Fare
count	891.000000	891.000000
mean	0.381594	32.204208
std	0.806057	49.693429
min	0.000000	0.000000
25%	0.000000	7.910400
50%	0.000000	14.454200
75%	0.000000	31.000000
max	6.000000	512.329200

#### Test Dataset Overview:

<class 'pandas.core.frame.DataFrame'>
RangeIndex: 418 entries, 0 to 417

Data columns (total 11 columns):

#	Column	Non-Null Count	Dtype
0	PassengerId	418 non-null	int64
1	Pclass	418 non-null	int64
2	Name	418 non-null	object
3	Sex	418 non-null	object
4	Age	332 non-null	float64
5	SibSp	418 non-null	int64
6	Parch	418 non-null	int64

```
7
    Ticket
             418 non-null object
             417 non-null float64
8
    Fare
             91 non-null
                           object
9
    Cabin
10 Embarked 418 non-null object
dtypes: float64(2), int64(4), object(5)
memory usage: 36.1+ KB
None
Test Dataset Statistical Summary:
     PassengerId Pclass
                                          SibSp
                                                    Parch
                                                                Fare
                                Age
count 418.000000 418.000000 332.000000 418.000000 418.000000 417.000000
mean 1100.500000 2.265550 30.272590 0.447368 0.392344 35.627188
     120.810458 0.841838 14.181209 0.896760 0.981429 55.907576
std
     892.000000 1.000000 0.170000 0.000000 0.000000 0.000000
min
25%
     996.250000 1.000000 21.000000 0.000000 0.000000 7.895800
50% 1100.500000 3.000000 27.000000 0.000000 0.000000 14.454200
    1204.750000 3.000000 39.000000 1.000000 0.000000 31.500000
75%
     1309.000000 3.000000 76.000000 8.000000 9.000000 512.329200
max
Gender Submission Dataset Overview:
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 418 entries, 0 to 417
Data columns (total 2 columns):
             Non-Null Count Dtype
   Column
   PassengerId 418 non-null int64
    Survived 418 non-null int64
dtypes: int64(2)
memory usage: 6.7 KB
None
```

#### Feed-Forward Neural Network

```
In [3]: # Load the Titanic dataset
        train_data = pd.read_csv('.../Datasets/Titanic/train.csv')
        # Preprocessing
        # Separate features and target
        y = train_data['Survived'] # Target
        X = train_data.drop(columns=['Survived', 'PassengerId', 'Name', 'Ticket', 'Cabin'])
        # Handle categorical variables with one-hot encoding
        categorical_features = ['Sex', 'Embarked']
        one hot encoder = OneHotEncoder(sparse output=False, handle unknown='ignore')
        categorical_encoded = one_hot_encoder.fit_transform(X[categorical_features])
        categorical_encoded_df = pd.DataFrame(categorical_encoded, columns=one_hot_encoder.g
        # Drop original categorical columns and append the encoded columns
        X = X.drop(columns=categorical_features)
        X = pd.concat([X.reset index(drop=True), categorical encoded df.reset index(drop=True)
        # Handle missing values with mean imputation
        imputer = SimpleImputer(strategy='mean')
        X_imputed = pd.DataFrame(imputer.fit_transform(X), columns=X.columns)
        # Standardize the features
        scaler = StandardScaler()
        X_scaled = pd.DataFrame(scaler.fit_transform(X_imputed), columns=X.columns)
```

```
# Split the data into training and test sets
        X_train, X_test, y_train, y_test = train_test_split(X_scaled, y, test_size=0.2, rand
        print("Training data shape:", X_train.shape)
        print("Test data shape:", X_test.shape)
        Training data shape: (712, 11)
        Test data shape: (179, 11)
In [4]: # Build the feed-forward neural network
        model = Sequential([
            Input(shape=(X_train.shape[1],)), # Define input shape explicitly
            Dense(64, activation='relu'),
            Dense(32, activation='relu'),
            Dense(1, activation='sigmoid') # Output layer for binary classification
        ])
        # Compile the model
        model.compile(optimizer=Adam(learning_rate=0.001), loss='binary_crossentropy', metri
        # Train the model
        history = model.fit(X_train, y_train, validation_split=0.2, epochs=50, batch_size=32
        # Evaluate the model
        test_loss, test_accuracy = model.evaluate(X_test, y_test, verbose=1)
        print(f"Test Loss: {test_loss:.4f}, Test Accuracy: {test_accuracy:.4f}")
```

```
7346 - val_loss: 0.5474 - val_accuracy: 0.7762
Epoch 2/50
803 - val_loss: 0.4718 - val_accuracy: 0.7972
Epoch 3/50
873 - val_loss: 0.4397 - val_accuracy: 0.7972
Epoch 4/50
961 - val loss: 0.4183 - val accuracy: 0.8182
Epoch 5/50
18/18 [============= ] - 0s 3ms/step - loss: 0.4414 - accuracy: 0.8
049 - val_loss: 0.4095 - val_accuracy: 0.8182
Epoch 6/50
137 - val_loss: 0.4031 - val_accuracy: 0.8252
Epoch 7/50
207 - val_loss: 0.4020 - val_accuracy: 0.8252
Epoch 8/50
190 - val_loss: 0.3981 - val_accuracy: 0.8252
Epoch 9/50
278 - val_loss: 0.3926 - val_accuracy: 0.8112
Epoch 10/50
295 - val_loss: 0.3952 - val_accuracy: 0.8252
Epoch 11/50
278 - val_loss: 0.4027 - val_accuracy: 0.8182
Epoch 12/50
278 - val_loss: 0.3951 - val_accuracy: 0.8322
Epoch 13/50
348 - val_loss: 0.3984 - val_accuracy: 0.8182
Epoch 14/50
383 - val loss: 0.3963 - val accuracy: 0.8252
Epoch 15/50
383 - val_loss: 0.3897 - val_accuracy: 0.8252
Epoch 16/50
348 - val loss: 0.4031 - val accuracy: 0.8252
Epoch 17/50
401 - val_loss: 0.3934 - val_accuracy: 0.8112
Epoch 18/50
313 - val_loss: 0.3901 - val_accuracy: 0.8252
Epoch 19/50
383 - val_loss: 0.3992 - val_accuracy: 0.8392
Epoch 20/50
366 - val_loss: 0.3961 - val_accuracy: 0.8182
```

```
Epoch 21/50
418 - val_loss: 0.3932 - val_accuracy: 0.8112
Epoch 22/50
436 - val_loss: 0.3968 - val_accuracy: 0.8182
Epoch 23/50
401 - val_loss: 0.3990 - val_accuracy: 0.8182
Epoch 24/50
436 - val loss: 0.3930 - val accuracy: 0.8112
Epoch 25/50
18/18 [============= ] - 0s 3ms/step - loss: 0.3776 - accuracy: 0.8
418 - val_loss: 0.3973 - val_accuracy: 0.8462
Epoch 26/50
471 - val_loss: 0.3942 - val_accuracy: 0.8322
Epoch 27/50
436 - val_loss: 0.3979 - val_accuracy: 0.8182
Epoch 28/50
453 - val_loss: 0.3979 - val_accuracy: 0.8322
Epoch 29/50
436 - val_loss: 0.3948 - val_accuracy: 0.8322
Epoch 30/50
453 - val_loss: 0.4008 - val_accuracy: 0.8252
Epoch 31/50
401 - val_loss: 0.4008 - val_accuracy: 0.8252
Epoch 32/50
436 - val_loss: 0.3976 - val_accuracy: 0.8322
Epoch 33/50
366 - val_loss: 0.3964 - val_accuracy: 0.8392
Epoch 34/50
471 - val loss: 0.3966 - val accuracy: 0.8392
489 - val_loss: 0.3994 - val_accuracy: 0.8392
Epoch 36/50
453 - val loss: 0.3971 - val accuracy: 0.8322
Epoch 37/50
418 - val_loss: 0.3993 - val_accuracy: 0.8252
Epoch 38/50
436 - val_loss: 0.4007 - val_accuracy: 0.8392
Epoch 39/50
471 - val_loss: 0.4058 - val_accuracy: 0.8392
Epoch 40/50
471 - val_loss: 0.3996 - val_accuracy: 0.8392
```

```
Epoch 41/50
418 - val_loss: 0.4045 - val_accuracy: 0.8322
Epoch 42/50
471 - val_loss: 0.4043 - val_accuracy: 0.8392
Epoch 43/50
471 - val_loss: 0.4044 - val_accuracy: 0.8392
Epoch 44/50
471 - val loss: 0.4084 - val accuracy: 0.8322
Epoch 45/50
436 - val_loss: 0.4014 - val_accuracy: 0.8392
Epoch 46/50
489 - val_loss: 0.4081 - val_accuracy: 0.8392
Epoch 47/50
453 - val_loss: 0.4007 - val_accuracy: 0.8392
Epoch 48/50
453 - val_loss: 0.4066 - val_accuracy: 0.8322
Epoch 49/50
418 - val_loss: 0.4093 - val_accuracy: 0.8392
Epoch 50/50
489 - val loss: 0.4084 - val accuracy: 0.8392
Test Loss: 0.4422, Test Accuracy: 0.8436
```

## **Surrogate Model - MLPClassifier**

```
In [5]: # Train a surrogate model (MLPClassifier)
surrogate_model = MLPClassifier(hidden_layer_sizes=(32,), activation='logistic', rar
print('Accuracy (MLPClassifier): ' + str(surrogate_model.score(X_train, y_train)))
Accuracy (MLPClassifier): 0.800561797752809
```

## **PART 2.1**

### Set up the LIME explainer

```
In [6]: # Function to visualize LIME explanations as a bar plot
def lime_exp_as_pyplot(exp, label=1, figsize=(8, 5)):
    exp_list = exp.as_list(label=label)
    fig, ax = plt.subplots(figsize=figsize)

# Extract feature names and importance values
    vals = [x[1] for x in exp_list]
    names = [x[0] for x in exp_list]

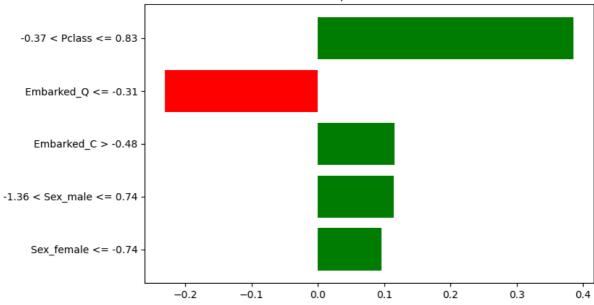
# Reverse for descending order of feature importance
    vals.reverse()
```

```
names.reverse()
   # Color the bars: green for positive, red for negative
   colors = ['green' if x > 0 else 'red' for x in vals]
   # Positions for the bars
   pos = np.arange(len(exp_list)) + .5
   # Plot the bars
   ax.barh(pos, vals, align='center', color=colors)
   plt.yticks(pos, names)
   return fig, ax
# Wrap the Keras model's prediction function for LIME
def predict_proba(X):
    """Custom function for LIME to get model predictions."""
   prob_class_1 = model.predict(X) # Predicted probability for class 1
   prob_class_0 = 1 - prob_class_1 # Predicted probability for class 0
   return np.hstack((prob_class_0, prob_class_1)) # Combine probabilities
# Initialize the LIME Tabular Explainer
explainer = lime.lime_tabular.LimeTabularExplainer(
   X_train.to_numpy(),
   feature_names=X_train.columns.to_list(),
   class_names=['Not Survived', 'Survived'],
   discretize_continuous=True,
   random_state=42
# Example instance index for "Survived" and "Not Survived"
survived_idx = np.where(y_test.to_numpy() == 1)[0][0]
not_survived_idx = np.where(y_test.to_numpy() == 0)[0][0]
# Explanation for "Survived" instance
survived_exp = explainer.explain_instance(
   X_test.iloc[survived_idx].to_numpy(),
   predict_proba,
   num features=5,
   top_labels=1
)
# Dynamically find the label for "Survived" instance
available_label = list(survived_exp.local_exp.keys())[0] # Pick the first available
print(f"Available label for Survived instance: {available_label}")
# Visualize explanation for the "Survived" instance
f, ax = lime exp as pyplot(survived exp, label=available label)
survived_confidence = model.predict(X_test.iloc[survived_idx:survived_idx + 1].to_nd
ax.set_title(f'Survived Case | Model Confidence: {survived_confidence:.2f}')
plt.show()
# Explanation for "Not Survived" instance
not survived exp = explainer.explain instance(
   X_test.iloc[not_survived_idx].to_numpy(),
   predict_proba,
   num_features=5,
   top_labels=1
)
```

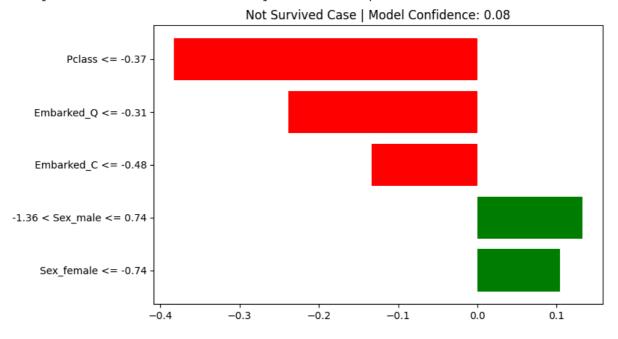
```
# Dynamically find the label for "Not Survived" instance
available_label = list(not_survived_exp.local_exp.keys())[0] # Pick the first avail
print(f"Available label for Not Survived instance: {available_label}")

# Visualize explanation for the "Not Survived" instance
f, ax = lime_exp_as_pyplot(not_survived_exp, label=available_label)
not_survived_confidence = model.predict(X_test.iloc[not_survived_idx:not_survived_id
ax.set_title(f'Not Survived Case | Model Confidence: {not_survived_confidence:.2f}')
plt.show()
```

Survived Case | Model Confidence: 0.13



157/157 [========] - 0s 754us/step Available label for Not Survived instance: 0
1/1 [======] - 0s 22ms/step



## Part 2.1 b

LIME (Local Interpretable Model-agnostic Explanations) is an algorithm designed to provide interpretability for complex, black-box models by approximating their local decision boundaries. It operates by perturbing the input data around a specific instance and observing the model's outputs for these slightly modified samples. The results of these perturbations are used to fit an interpretable surrogate model, typically a linear model, that captures the behaviour of the black-box model within the vicinity of the specific instance.

For our Titanic dataset, LIME highlights the contributions of individual features to the model's decision-making process for specific instances. For example, in the visualisations above:

- In the "Survived" case, features such as "Pclass" (passenger class) and "Sex\_female" have significant positive contributions to the prediction, as indicated by the green bars. On the other hand, features like "Embarked\_Q" negatively influence the outcome, as indicated by the red bars. This suggests that higher socio-economic status and being female are strongly associated with survival, whereas embarking from certain locations may decrease survival probability.
- In the "Not Survived" case, features such as "Pclass" negatively influence the prediction, suggesting that lower socio-economic status correlates with non-survival. Similarly, factors like "Parch" (number of parents/children aboard) might also contribute negatively. Positive influences like "Sex\_female" show a mitigating factor, indicating that the model considers gender but not sufficiently to alter the outcome.

By presenting feature contributions as weights (positive or negative) for each instance, LIME provides a clear interpretative framework. The approximations, while not perfectly reflecting the global decision boundary, give useful insights into how the model uses features locally. This interpretability is crucial for datasets like Titanic, where fairness and historical biases (e.g., gender and class disparity) can be critically analysed.

#### Part 2.2

#### Adding SHAP to Explain Model Predictions

```
In [7]: # Use SHAP's DeepExplainer for neural networks
    explainer = shap.KernelExplainer(model.predict, X_train[:100]) # Use a small sample

# Calculate SHAP values for a set of instances
    shap_values = explainer.shap_values(X_test[:10]) # Explaining the first 10 samples

# Visualize the SHAP values for the first test sample (e.g., index 0)
    shap.initjs()

# Reshape SHAP values if necessary
    shap_values_reshaped = shap_values[0].reshape(1, -1)

# Now plot with reshaped values
    shap.force_plot(
        explainer.expected_value[0],
        shap_values_reshaped[0], # SHAP values for the first sample (class 0)
        X_test.iloc[0], # Actual features for the first sample
```

```
feature names=X.columns # Feature names
      4/4 [======== ] - 0s 997us/step
                | 0/10 [00:00<?, ?it/s]
      1/1 [======] - 0s 21ms/step
      6394/6394 [========== ] - 6s 884us/step
                | 1/10 [00:09<01:28, 9.79s/it]
      1/1 [=======] - 0s 21ms/step
      6394/6394 [========== ] - 5s 848us/step
             2/10 [00:18<01:13, 9.16s/it]
      1/1 [=======] - 0s 21ms/step
      6394/6394 [========== ] - 5s 823us/step
             | 3/10 [00:27<01:02, 8.88s/it]
      1/1 [=======] - 0s 20ms/step
      6394/6394 [========== ] - 5s 814us/step
      40%| 4/10 [00:35<00:52, 8.76s/it]
      6394/6394 [========== ] - 5s 824us/step
               | 5/10 [00:44<00:43, 8.66s/it]
      1/1 [======] - 0s 19ms/step
      6394/6394 [========== ] - 5s 829us/step
      60% | 6/10 [00:52<00:34, 8.63s/it]
      1/1 [======] - 0s 23ms/step
      6394/6394 [========== ] - 6s 859us/step
              7/10 [01:01<00:26, 8.71s/it]
      1/1 [=======] - 0s 23ms/step
      6394/6394 [========== ] - 5s 836us/step
           8/10 [01:10<00:17, 8.77s/it]
      1/1 [=======] - 0s 24ms/step
      6394/6394 [========== ] - 6s 963us/step
          | 9/10 [01:20<00:09, 9.05s/it]
      1/1 [======] - 0s 23ms/step
      6394/6394 [========== ] - 5s 797us/step
      100% | 10/10 [01:28<00:00, 8.87s/it]
                                               Out[7]:
                                                   f(x)
     -0.2019
                  -0.1019
                                -0.001891
                                              0.09811 0.13
                                                            0.1
                      Fare = -0.3415 Parch = 0.7676 Embarked S = -1.615
                                                           Pclas
```

### Part 2.2 b

LIME works by approximating the decision boundary of a model in the vicinity of a particular instance. It perturbs the input data around the instance to generate nearby samples, evaluates the model's predictions on these samples, and fits an interpretable surrogate model, such as a linear regression, to mimic the model's behaviour locally. In the visualisations provided for LIME, we see feature contributions represented as positive or

negative weights, indicating whether each feature supports or opposes a specific outcome. For example, "Pclass" and "Sex\_female" contribute significantly to predicting survival, as denoted by the green bars, while features like "Embarked\_Q" negatively influence the same prediction. LIME's strength lies in its simplicity and ability to provide intuitive explanations for specific instances. However, its reliance on local approximations can sometimes result in less accurate explanations for complex, non-linear models.

SHAP, on the other hand, is based on Shapley values from cooperative game theory, ensuring that feature attributions are both consistent and theoretically sound. SHAP calculates the contribution of each feature to the prediction by considering all possible combinations of feature presence and absence, making it more computationally intensive than LIME. In the SHAP visualisation, feature contributions are displayed on a scale from negative (red, reducing the prediction) to positive (blue, increasing the prediction), with the sum of contributions equalling the model's prediction. For example, "Embarked\_C" strongly increases the prediction score for survival, while "Embarked\_S" has a significant negative impact. SHAP's additive nature and consistency make it particularly suitable for gaining a more holistic and globally consistent understanding of a model's behaviour.

The distinctions between SHAP and LIME are evident when compared. LIME approximates the decision boundary for a particular instance and its surroundings, concentrating only on local explanations. This method might not have the capability to fully capture the intricacies of the global model, although being quicker and frequently intuitive. SHAP, on the other hand, guarantees that feature contributions are uniform throughout the dataset, offering both local and global interpretability. SHAP becomes more resource-intensive but more resilient as a result. In conclusion, SHAP provides a deeper and more trustworthy understanding of feature attributions at the expense of greater computational effort, whereas LIME is useful for rapid and local insights.

#### Part 2.3

# Implementing Anchors to interpret model predictions in specific cases

```
instance_to_explain,
    pred_fn,
    threshold=0.95
)
# Display the results
print('Anchor: %s' % (' AND '.join(exp.names())))
print('Precision: %.2f' % exp.precision())
print('Coverage: %.2f' % exp.coverage())
exp.show_in_notebook()
Anchor: Sex_female <= -0.74 AND Pclass = female
Precision: 0.98
Coverage: 0.40
                                                      Explanation of A.I. prediction
   Example
                                  A.I. pre...
                                                      If ALL of these are true:
                                     Not
         Pclass = female
                                    Survived

✓ Sex_female <= -0.74
</p>
                Age = C
                                                        ✓ Pclass = female
    -0.47 < SibSp <= 0.4
                                                      The A.I. will predict Not
           Parch > -0.47
                                                      Survived 97.7% of the time
    -0.36 < Fare < = -0.03
    Sex_female <= -0.74
    -1.36 < Sex_male <=
                   0.74
```

> Examples where the A.I. agent predicts Not Survived

0.05

> Examples where the A.I. agent DOES NOT predict Not Survived

### Part 2.3 b

 $Embarked_C > -0.48$ 

 $Embarked_Q <= -0.3$ 

 $Embarked_S <= -1.61$ 

Embarked\_nan <= -

Each of SHAP, LIME, and Anchors offers a different way to interpret machine learning predictions. By precisely determining the minimal circumstances that "anchor" a prediction, anchors concentrate on rule-based explanations. Because Anchors generate straightforward, understandable principles, they are therefore very interpretable. However, by disregarding interactions that do not fall under the designated parameters, these rules may oversimplify the decision-making process.

An option is provided by LIME, which approximates the local decision boundary by fitting a linear model and perturbing input data. Individual feature contributions, such "Pclass" or "Sex\_female," are highlighted as either in favour of or against a prediction. Although LIME works well for producing concise and understandable explanations, its unpredictability in perturbations may cause variability and make it difficult to adequately reflect non-linear interactions in the model.

In contrast, SHAP ensures consistency and equity in attribution by employing Shapley values to assign additive contributions to features. Features like "Embarked\_C" greatly support survival predictions, whereas "Embarked\_S" lowers the likelihood of survival, as shown in SHAP visualisations. Although SHAP is more computationally demanding and may be too detailed for consumers, it excels at delivering both local and global explanations.

To sum up, SHAP is best suited for thorough and trustworthy feature attributions, LIME is helpful for quick and easy local explanations, and Anchors are perfect for producing straightforward and actionable rules. The model's complexity and the particular requirements for interpretability will determine which of these approaches is best.