

```
!pip install -q altair shap causal-learn graphviz pydot plotnine
```

```
0.0/193.0 kB ? eta -:--:--  
193.0/193.0 kB 6.0 MB/s eta  
0:00:00
```

```
import pandas as pd  
import numpy as np  
import matplotlib.pyplot as plt  
import seaborn as sns  
import json
```

```
from sklearn.preprocessing import StandardScaler  
from sklearn.linear_model import LogisticRegression  
from sklearn.ensemble import RandomForestClassifier  
from sklearn.metrics import accuracy_score, precision_score,  
recall_score, f1_score, roc_auc_score  
from sklearn.metrics import classification_report, confusion_matrix,  
roc_auc_score, roc_curve, auc  
from sklearn.model_selection import train_test_split
```

```
import tensorflow as tf  
from tensorflow.keras.models import Sequential  
from tensorflow.keras.callbacks import EarlyStopping  
from tensorflow.keras.layers import Input, Dense, Dropout
```

```
from plotnine.data import mpg  
from plotnine import ggplot, aes, geom_histogram, geom_density,  
theme_bw, labs  
from IPython.display import display
```

```
import networkx as nx
```

```
import altair as alt
```

```
from causallearn.search.ConstraintBased.PC import pc  
from causallearn.utils.cit import fisherz  
from causallearn.graph.Edge import Endpoint
```

```
import shap  
from shap import Explanation, KernelExplainer, summary_plot,  
force_plot
```

```
df = pd.read_csv('/content/gene_expression.csv')
```

```
df.head()
```

```
{"summary": "{\n  \"name\": \"df\", \n  \"rows\": 3000, \n  \"fields\": [\n    {\n      \"column\": \"Gene One\", \n      \"properties\": {\n        \"dtype\": \"number\", \n        \"std\": 1.8283875362049855, \n
```

```

{"min": 1.0, "max": 10.0, "num_unique_values": 89, "samples": [4.7, 2.9, 6.2], "semantic_type": "", "description": "", "column": "Gene Two", "properties": {"dtype": "number", "std": 1.7290807834436468, "min": 1.0, "max": 10.0, "num_unique_values": 88, "samples": [1.4, 3.9, 6.1], "semantic_type": "", "description": "", "column": "Cancer Present", "properties": {"dtype": "number", "std": 0, "min": 0, "max": 1, "num_unique_values": 2, "samples": [0, 1], "semantic_type": "", "description": ""}}
n}, {"type": "dataframe", "variable_name": "df"}

```

```
df.info()
```

```

<class 'pandas.core.frame.DataFrame'>
RangeIndex: 3000 entries, 0 to 2999
Data columns (total 3 columns):
#   Column          Non-Null Count  Dtype
---  -
0   Gene One        3000 non-null   float64
1   Gene Two        3000 non-null   float64
2   Cancer Present  3000 non-null   int64
dtypes: float64(2), int64(1)
memory usage: 70.4 KB

```

```
df.describe()
```

```

{"summary": {"name": "df", "rows": 8, "fields": [{"column": "Gene One", "properties": {"dtype": "number", "std": 1058.8846317002517, "min": 1.0, "max": 3000.0, "num_unique_values": 8, "samples": [5.6001333333333334, 5.6, 3000.0]}, {"column": "Gene Two", "properties": {"dtype": "number", "std": 1058.934579724635, "min": 1.0, "max": 3000.0, "num_unique_values": 8, "samples": [5.410466666666666, 5.4, 3000.0]}, {"column": "Cancer Present", "properties": {"dtype": "number", "std": 1060.4834582292306, "min": 0.0, "max": 3000.0, "num_unique_values": 5, "samples": [1.0, 0.5000833541724554]}]}

```

```

{"semantic_type\\": "\\n",\\n      "\\description\\": "\\n"\\n      }\\n    }\\n  ]\\n}", "type": "dataframe"}

df['Cancer Present'].value_counts(normalize=True)

Cancer Present
1    0.5
0    0.5
Name: proportion, dtype: float64

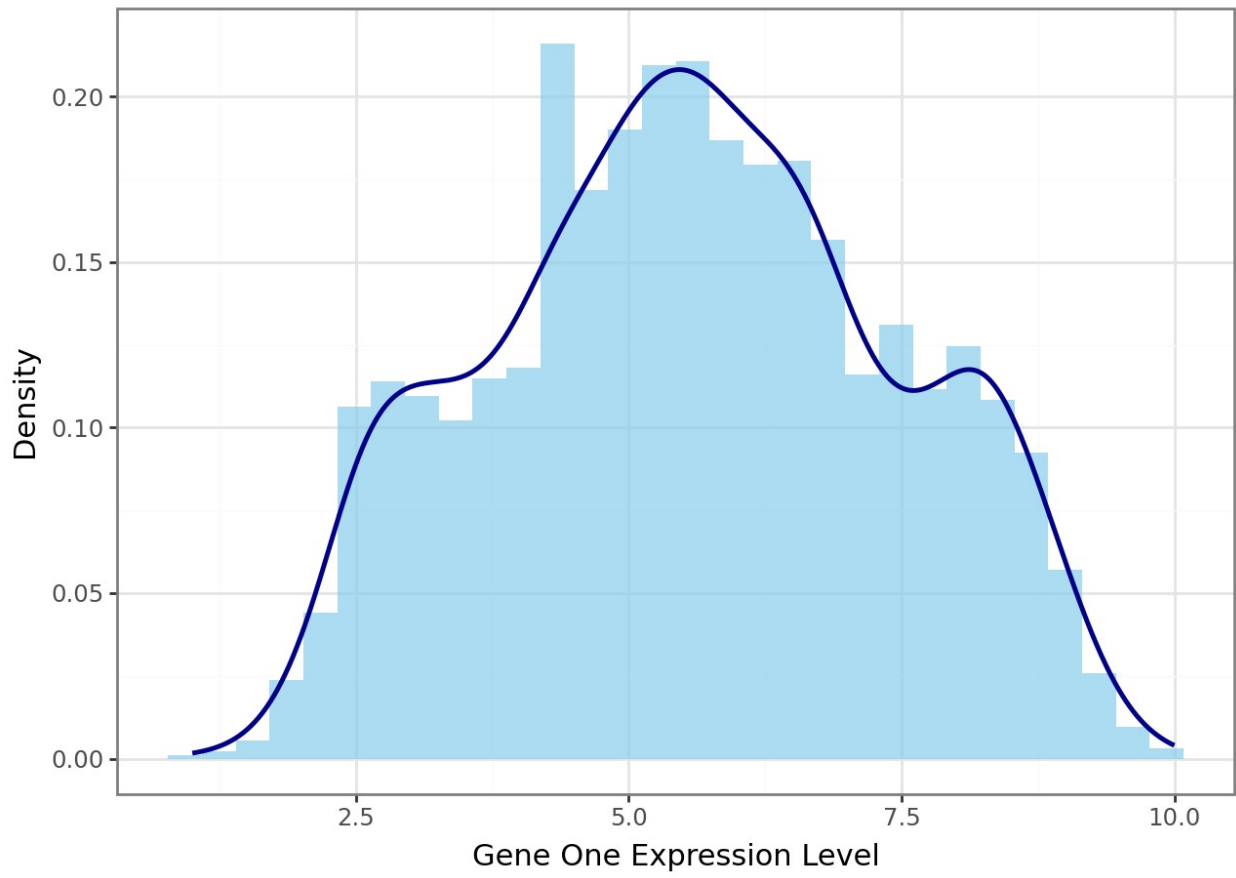
plot_gene_one = (
    ggplot(df, aes(x='Gene One')) +
    geom_histogram(aes(y='..density..'), fill='skyblue', bins=30,
alpha=0.7) +
    geom_density(color='darkblue', size=1.0) +
    labs(title='Distribution of Gene One Expression', x='Gene One
Expression Level', y='Density') +
    theme_bw()
)

plot_gene_two = (
    ggplot(df, aes(x='Gene Two')) +
    geom_histogram(aes(y='..density..'), fill='lightgreen', bins=30,
alpha=0.7) +
    geom_density(color='darkgreen', size=1.0) +
    labs(title='Distribution of Gene Two Expression', x='Gene Two
Expression Level', y='Density') +
    theme_bw()
)

# Display plots
display(plot_gene_one, plot_gene_two)

```

Distribution of Gene One Expression





Gene 1 is expressed consistently like a normal distribution. It may be constitutively expressed.

Gene 2 is expressed in a wider distribution meaning it varies differently in normal and disease states.

```
df_grouped = df.groupby('Cancer Present')[['Gene One', 'Gene Two']].mean().reset_index()

# Melt the DataFrame to long format
df_bar = df_grouped.melt(id_vars='Cancer Present',
                        var_name='Gene',
                        value_name='Expression')

df_bar.rename(columns={'Cancer Present': 'Cancer'}, inplace=True)

# Plot
bar_chart = alt.Chart(df_bar).mark_bar().encode(
    x=alt.X('Gene:N', title='Gene'),
    y=alt.Y('Expression:Q', title='Mean Expression'),
    color=alt.Color('Cancer:N', scale=alt.Scale(range=['#1f77b4', '#d62728'])),
    column=alt.Column('Cancer:N', title='Cancer Status'),
```

```

        tooltip=['Gene:N', 'Expression:Q', 'Cancer:N']
    ).properties(
        title="Mean Gene Expression by Cancer Status",
        width=150,
        height=300
    )

bar_chart
alt.Chart(...)

scatter = alt.Chart(df).mark_circle(size=60).encode(
    x=alt.X('Gene One', title='Gene One Expression'),
    y=alt.Y('Gene Two', title='Gene Two Expression'),
    color=alt.Color('Cancer Present:N',
        scale=alt.Scale(range=['#1f77b4', '#d62728'])),
    legend=alt.Legend(title='Cancer'),
    tooltip=['Gene One', 'Gene Two', 'Cancer Present']
).properties(
    width=500,
    height=400,
    title='Scatter Plot of Gene One vs Gene Two by Cancer Status'
).interactive()

scatter
alt.Chart(...)

scaler = StandardScaler()
X = scaler.fit_transform(df[['Gene One', 'Gene Two']])
y = df['Cancer Present']

X_train, X_test, y_train, y_test = train_test_split(
    X, y, test_size=0.25, stratify=y, random_state=42)

log_reg = LogisticRegression()
log_reg.fit(X_train, y_train)

# Predict on test set
y_pred = log_reg.predict(X_test)
y_prob = log_reg.predict_proba(X_test)[:, 1]

# Evaluate model performance
print("Confusion Matrix:\n", confusion_matrix(y_test, y_pred))
print("\nClassification Report:\n", classification_report(y_test,
y_pred))
print("\nROC AUC Score:", roc_auc_score(y_test, y_prob))

Confusion Matrix:
[[325  50]
 [ 42 333]]

```

```

Classification Report:
              precision    recall  f1-score   support

     0       0.89       0.87       0.88       375
     1       0.87       0.89       0.88       375

 accuracy          0.88
 macro avg         0.88
 weighted avg      0.88

```

ROC AUC Score: 0.9485866666666666

```

rf = RandomForestClassifier(n_estimators=100, random_state=42)
rf.fit(X_train, y_train)

```

Predictions

```

y_pred_rf = rf.predict(X_test)
y_prob_rf = rf.predict_proba(X_test)[:, 1]

```

Evaluation

```

print("Confusion Matrix:\n", confusion_matrix(y_test, y_pred_rf))
print("\nClassification Report:\n", classification_report(y_test,
y_pred_rf))
print("\nROC AUC Score:", roc_auc_score(y_test, y_prob_rf))

```

Confusion Matrix:

```

[[341  34]
 [ 25 350]]

```

```

Classification Report:
              precision    recall  f1-score   support

     0       0.93       0.91       0.92       375
     1       0.91       0.93       0.92       375

 accuracy          0.92
 macro avg         0.92
 weighted avg      0.92

```

ROC AUC Score: 0.9801386666666667

```

feat_df = pd.DataFrame({
    'Feature': ['Gene One', 'Gene Two'],
    'Importance': rf.feature_importances_
})

```

Plotting

```

sns.barplot(data=feat_df, x='Importance', y='Feature',

```

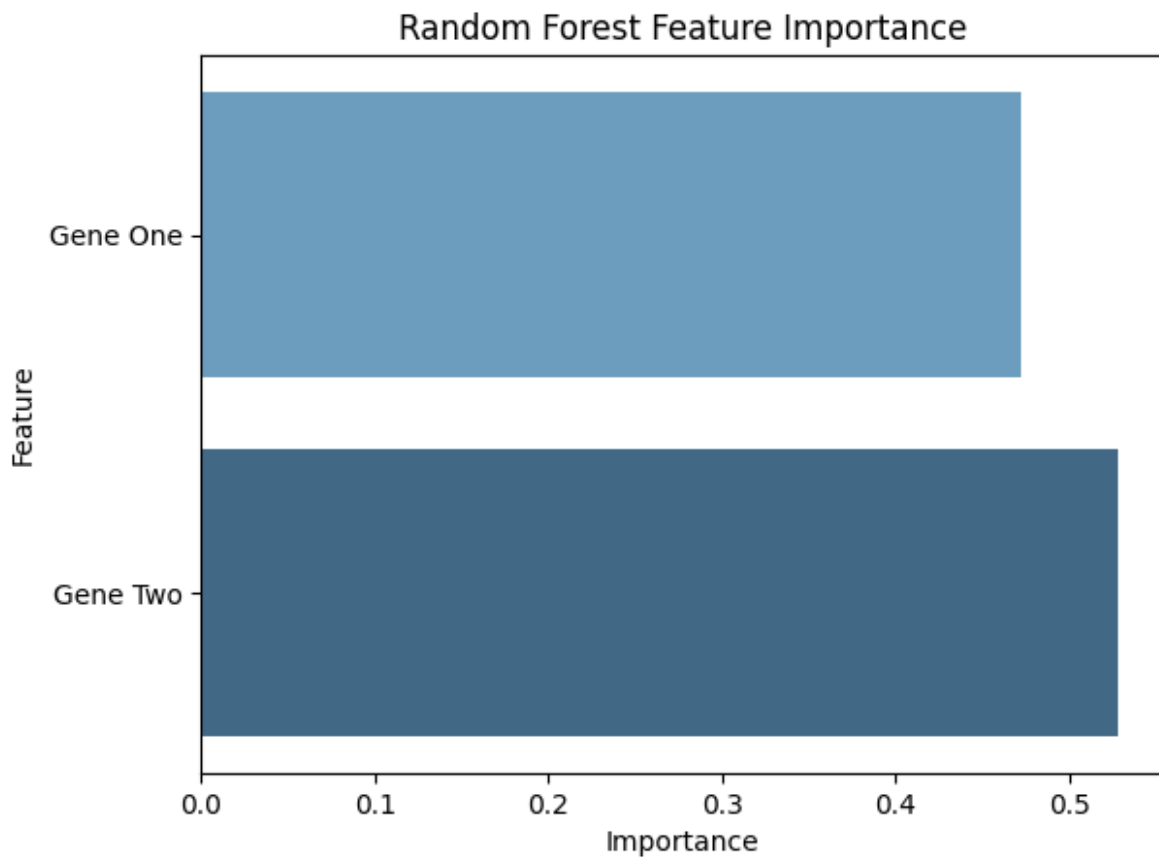
```

palette='Blues_d')
plt.title('Random Forest Feature Importance')
plt.xlabel('Importance')
plt.ylabel('Feature')
plt.show()

```

/tmp/ipython-input-19-1183823344.py:7: FutureWarning:

Passing `palette` without assigning `hue` is deprecated and will be removed in v0.14.0. Assign the `y` variable to `hue` and set `legend=False` for the same effect.



```

tf.random.set_seed(42)

# Define model
model = Sequential([
    Input(shape=(X_train.shape[1],)),
    Dense(16, activation='relu'),
    Dropout(0.3),
    Dense(8, activation='relu'),
    Dropout(0.3),
    Dense(1, activation='sigmoid')
])

```



```

])

model.compile(optimizer='adam', loss='binary_crossentropy',
metrics=['accuracy'])
model.summary()

```

Model: "sequential"

Layer (type) Param #	Output Shape	
dense (Dense) 48	(None, 16)	
dropout (Dropout) 0	(None, 16)	
dense_1 (Dense) 136	(None, 8)	
dropout_1 (Dropout) 0	(None, 8)	
dense_2 (Dense) 9	(None, 1)	

Total params: 193 (772.00 B)

Trainable params: 193 (772.00 B)

Non-trainable params: 0 (0.00 B)

Early stopping

```

early_stop = EarlyStopping(monitor='val_loss', patience=10,
restore_best_weights=True)

```

Train the model

```

history = model.fit(
    X_train, y_train,
    validation_split=0.2,
    epochs=100,
    batch_size=32,

```

```
callbacks=[early_stop],
verbose=1
)

Epoch 1/100
57/57 _____ 5s 8ms/step - accuracy: 0.4681 - loss:
0.7219 - val_accuracy: 0.7156 - val_loss: 0.5741
Epoch 2/100
57/57 _____ 0s 4ms/step - accuracy: 0.6618 - loss:
0.5892 - val_accuracy: 0.8533 - val_loss: 0.4934
Epoch 3/100
57/57 _____ 0s 4ms/step - accuracy: 0.7215 - loss:
0.5367 - val_accuracy: 0.8644 - val_loss: 0.4346
Epoch 4/100
57/57 _____ 0s 5ms/step - accuracy: 0.7719 - loss:
0.4875 - val_accuracy: 0.8644 - val_loss: 0.3850
Epoch 5/100
57/57 _____ 0s 3ms/step - accuracy: 0.8112 - loss:
0.4384 - val_accuracy: 0.8667 - val_loss: 0.3492
Epoch 6/100
57/57 _____ 0s 4ms/step - accuracy: 0.8088 - loss:
0.4400 - val_accuracy: 0.8711 - val_loss: 0.3303
Epoch 7/100
57/57 _____ 0s 4ms/step - accuracy: 0.8293 - loss:
0.3940 - val_accuracy: 0.8711 - val_loss: 0.3186
Epoch 8/100
57/57 _____ 0s 4ms/step - accuracy: 0.8367 - loss:
0.4071 - val_accuracy: 0.8711 - val_loss: 0.3134
Epoch 9/100
57/57 _____ 0s 4ms/step - accuracy: 0.8157 - loss:
0.4122 - val_accuracy: 0.8711 - val_loss: 0.3107
Epoch 10/100
57/57 _____ 0s 3ms/step - accuracy: 0.8394 - loss:
0.3854 - val_accuracy: 0.8711 - val_loss: 0.3068
Epoch 11/100
57/57 _____ 0s 4ms/step - accuracy: 0.8341 - loss:
0.3905 - val_accuracy: 0.8711 - val_loss: 0.3033
Epoch 12/100
57/57 _____ 0s 4ms/step - accuracy: 0.8418 - loss:
0.3760 - val_accuracy: 0.8733 - val_loss: 0.2997
Epoch 13/100
57/57 _____ 0s 5ms/step - accuracy: 0.8298 - loss:
0.3859 - val_accuracy: 0.8733 - val_loss: 0.2979
Epoch 14/100
57/57 _____ 0s 3ms/step - accuracy: 0.8467 - loss:
0.3810 - val_accuracy: 0.8733 - val_loss: 0.2946
Epoch 15/100
57/57 _____ 0s 4ms/step - accuracy: 0.8462 - loss:
0.3578 - val_accuracy: 0.8733 - val_loss: 0.2910
Epoch 16/100
```

57/57 ————— 0s 4ms/step - accuracy: 0.8466 - loss: 0.3519 - val_accuracy: 0.8756 - val_loss: 0.2881
Epoch 17/100
57/57 ————— 0s 5ms/step - accuracy: 0.8413 - loss: 0.3628 - val_accuracy: 0.8778 - val_loss: 0.2848
Epoch 18/100
57/57 ————— 0s 4ms/step - accuracy: 0.8473 - loss: 0.3546 - val_accuracy: 0.8733 - val_loss: 0.2814
Epoch 19/100
57/57 ————— 0s 4ms/step - accuracy: 0.8450 - loss: 0.3589 - val_accuracy: 0.8800 - val_loss: 0.2794
Epoch 20/100
57/57 ————— 0s 5ms/step - accuracy: 0.8467 - loss: 0.3459 - val_accuracy: 0.8756 - val_loss: 0.2769
Epoch 21/100
57/57 ————— 0s 4ms/step - accuracy: 0.8492 - loss: 0.3626 - val_accuracy: 0.8733 - val_loss: 0.2744
Epoch 22/100
57/57 ————— 0s 4ms/step - accuracy: 0.8470 - loss: 0.3408 - val_accuracy: 0.8822 - val_loss: 0.2716
Epoch 23/100
57/57 ————— 0s 4ms/step - accuracy: 0.8600 - loss: 0.3387 - val_accuracy: 0.8800 - val_loss: 0.2691
Epoch 24/100
57/57 ————— 0s 4ms/step - accuracy: 0.8519 - loss: 0.3395 - val_accuracy: 0.8844 - val_loss: 0.2660
Epoch 25/100
57/57 ————— 0s 4ms/step - accuracy: 0.8585 - loss: 0.3302 - val_accuracy: 0.8844 - val_loss: 0.2631
Epoch 26/100
57/57 ————— 0s 4ms/step - accuracy: 0.8622 - loss: 0.3372 - val_accuracy: 0.8844 - val_loss: 0.2611
Epoch 27/100
57/57 ————— 0s 5ms/step - accuracy: 0.8511 - loss: 0.3361 - val_accuracy: 0.8822 - val_loss: 0.2587
Epoch 28/100
57/57 ————— 0s 3ms/step - accuracy: 0.8567 - loss: 0.3321 - val_accuracy: 0.8911 - val_loss: 0.2559
Epoch 29/100
57/57 ————— 0s 7ms/step - accuracy: 0.8627 - loss: 0.3175 - val_accuracy: 0.8933 - val_loss: 0.2533
Epoch 30/100
57/57 ————— 1s 5ms/step - accuracy: 0.8611 - loss: 0.3225 - val_accuracy: 0.8933 - val_loss: 0.2507
Epoch 31/100
57/57 ————— 0s 5ms/step - accuracy: 0.8596 - loss: 0.3291 - val_accuracy: 0.8956 - val_loss: 0.2497
Epoch 32/100
57/57 ————— 1s 7ms/step - accuracy: 0.8519 - loss:

0.3351 - val_accuracy: 0.8933 - val_loss: 0.2479
Epoch 33/100
57/57 _____ 1s 5ms/step - accuracy: 0.8601 - loss:
0.3255 - val_accuracy: 0.8933 - val_loss: 0.2454
Epoch 34/100
57/57 _____ 1s 4ms/step - accuracy: 0.8757 - loss:
0.3050 - val_accuracy: 0.8933 - val_loss: 0.2444
Epoch 35/100
57/57 _____ 0s 4ms/step - accuracy: 0.8540 - loss:
0.3074 - val_accuracy: 0.8956 - val_loss: 0.2430
Epoch 36/100
57/57 _____ 0s 4ms/step - accuracy: 0.8652 - loss:
0.3224 - val_accuracy: 0.8956 - val_loss: 0.2418
Epoch 37/100
57/57 _____ 0s 4ms/step - accuracy: 0.8617 - loss:
0.3240 - val_accuracy: 0.8978 - val_loss: 0.2404
Epoch 38/100
57/57 _____ 0s 3ms/step - accuracy: 0.8580 - loss:
0.3259 - val_accuracy: 0.9022 - val_loss: 0.2392
Epoch 39/100
57/57 _____ 0s 4ms/step - accuracy: 0.8634 - loss:
0.3050 - val_accuracy: 0.9022 - val_loss: 0.2359
Epoch 40/100
57/57 _____ 0s 5ms/step - accuracy: 0.8776 - loss:
0.3013 - val_accuracy: 0.9089 - val_loss: 0.2334
Epoch 41/100
57/57 _____ 0s 4ms/step - accuracy: 0.8770 - loss:
0.3025 - val_accuracy: 0.9133 - val_loss: 0.2312
Epoch 42/100
57/57 _____ 0s 4ms/step - accuracy: 0.8807 - loss:
0.3000 - val_accuracy: 0.9111 - val_loss: 0.2308
Epoch 43/100
57/57 _____ 0s 3ms/step - accuracy: 0.8669 - loss:
0.3002 - val_accuracy: 0.9133 - val_loss: 0.2288
Epoch 44/100
57/57 _____ 0s 4ms/step - accuracy: 0.8577 - loss:
0.3128 - val_accuracy: 0.9133 - val_loss: 0.2271
Epoch 45/100
57/57 _____ 0s 3ms/step - accuracy: 0.8800 - loss:
0.3086 - val_accuracy: 0.9133 - val_loss: 0.2260
Epoch 46/100
57/57 _____ 0s 3ms/step - accuracy: 0.8739 - loss:
0.3033 - val_accuracy: 0.9156 - val_loss: 0.2252
Epoch 47/100
57/57 _____ 0s 5ms/step - accuracy: 0.8754 - loss:
0.2986 - val_accuracy: 0.9156 - val_loss: 0.2248
Epoch 48/100
57/57 _____ 0s 4ms/step - accuracy: 0.8852 - loss:
0.2853 - val_accuracy: 0.9156 - val_loss: 0.2237

Epoch 49/100
57/57 _____ 0s 4ms/step - accuracy: 0.8799 - loss: 0.3010 - val_accuracy: 0.9156 - val_loss: 0.2235
Epoch 50/100
57/57 _____ 0s 4ms/step - accuracy: 0.8719 - loss: 0.2922 - val_accuracy: 0.9178 - val_loss: 0.2210
Epoch 51/100
57/57 _____ 0s 3ms/step - accuracy: 0.8791 - loss: 0.2985 - val_accuracy: 0.9178 - val_loss: 0.2208
Epoch 52/100
57/57 _____ 0s 4ms/step - accuracy: 0.8719 - loss: 0.3000 - val_accuracy: 0.9178 - val_loss: 0.2196
Epoch 53/100
57/57 _____ 0s 3ms/step - accuracy: 0.8839 - loss: 0.2935 - val_accuracy: 0.9200 - val_loss: 0.2191
Epoch 54/100
57/57 _____ 0s 5ms/step - accuracy: 0.8800 - loss: 0.3051 - val_accuracy: 0.9200 - val_loss: 0.2191
Epoch 55/100
57/57 _____ 0s 5ms/step - accuracy: 0.8800 - loss: 0.3083 - val_accuracy: 0.9222 - val_loss: 0.2183
Epoch 56/100
57/57 _____ 0s 4ms/step - accuracy: 0.8735 - loss: 0.2969 - val_accuracy: 0.9222 - val_loss: 0.2183
Epoch 57/100
57/57 _____ 0s 4ms/step - accuracy: 0.8865 - loss: 0.2937 - val_accuracy: 0.9200 - val_loss: 0.2162
Epoch 58/100
57/57 _____ 0s 4ms/step - accuracy: 0.8780 - loss: 0.2990 - val_accuracy: 0.9222 - val_loss: 0.2157
Epoch 59/100
57/57 _____ 0s 5ms/step - accuracy: 0.8871 - loss: 0.2825 - val_accuracy: 0.9222 - val_loss: 0.2135
Epoch 60/100
57/57 _____ 0s 4ms/step - accuracy: 0.8808 - loss: 0.2958 - val_accuracy: 0.9244 - val_loss: 0.2127
Epoch 61/100
57/57 _____ 0s 4ms/step - accuracy: 0.8890 - loss: 0.2913 - val_accuracy: 0.9222 - val_loss: 0.2111
Epoch 62/100
57/57 _____ 0s 3ms/step - accuracy: 0.8874 - loss: 0.2943 - val_accuracy: 0.9244 - val_loss: 0.2115
Epoch 63/100
57/57 _____ 0s 4ms/step - accuracy: 0.8876 - loss: 0.2861 - val_accuracy: 0.9244 - val_loss: 0.2121
Epoch 64/100
57/57 _____ 0s 3ms/step - accuracy: 0.8925 - loss: 0.2900 - val_accuracy: 0.9267 - val_loss: 0.2112
Epoch 65/100

57/57 _____ 0s 5ms/step - accuracy: 0.8749 - loss:
0.2973 - val_accuracy: 0.9267 - val_loss: 0.2110
Epoch 66/100
57/57 _____ 1s 5ms/step - accuracy: 0.8912 - loss:
0.2793 - val_accuracy: 0.9267 - val_loss: 0.2089
Epoch 67/100
57/57 _____ 1s 5ms/step - accuracy: 0.8808 - loss:
0.3021 - val_accuracy: 0.9289 - val_loss: 0.2094
Epoch 68/100
57/57 _____ 1s 7ms/step - accuracy: 0.8855 - loss:
0.3023 - val_accuracy: 0.9289 - val_loss: 0.2089
Epoch 69/100
57/57 _____ 0s 6ms/step - accuracy: 0.8752 - loss:
0.2975 - val_accuracy: 0.9267 - val_loss: 0.2088
Epoch 70/100
57/57 _____ 1s 5ms/step - accuracy: 0.8854 - loss:
0.2871 - val_accuracy: 0.9244 - val_loss: 0.2068
Epoch 71/100
57/57 _____ 0s 4ms/step - accuracy: 0.8892 - loss:
0.2771 - val_accuracy: 0.9244 - val_loss: 0.2071
Epoch 72/100
57/57 _____ 0s 4ms/step - accuracy: 0.8874 - loss:
0.2904 - val_accuracy: 0.9267 - val_loss: 0.2067
Epoch 73/100
57/57 _____ 0s 3ms/step - accuracy: 0.8837 - loss:
0.2855 - val_accuracy: 0.9244 - val_loss: 0.2076
Epoch 74/100
57/57 _____ 0s 5ms/step - accuracy: 0.8892 - loss:
0.2927 - val_accuracy: 0.9244 - val_loss: 0.2088
Epoch 75/100
57/57 _____ 0s 4ms/step - accuracy: 0.8878 - loss:
0.2758 - val_accuracy: 0.9222 - val_loss: 0.2072
Epoch 76/100
57/57 _____ 0s 4ms/step - accuracy: 0.8917 - loss:
0.2924 - val_accuracy: 0.9222 - val_loss: 0.2067
Epoch 77/100
57/57 _____ 0s 3ms/step - accuracy: 0.8868 - loss:
0.2965 - val_accuracy: 0.9222 - val_loss: 0.2073
Epoch 78/100
57/57 _____ 0s 4ms/step - accuracy: 0.8708 - loss:
0.2945 - val_accuracy: 0.9222 - val_loss: 0.2081
Epoch 79/100
57/57 _____ 0s 4ms/step - accuracy: 0.8855 - loss:
0.3055 - val_accuracy: 0.9244 - val_loss: 0.2072
Epoch 80/100
57/57 _____ 0s 4ms/step - accuracy: 0.8746 - loss:
0.2965 - val_accuracy: 0.9244 - val_loss: 0.2086
Epoch 81/100
57/57 _____ 0s 5ms/step - accuracy: 0.8819 - loss:

0.2950 - val_accuracy: 0.9244 - val_loss: 0.2081
Epoch 82/100
57/57 _____ 0s 3ms/step - accuracy: 0.8796 - loss:
0.2988 - val_accuracy: 0.9244 - val_loss: 0.2062
Epoch 83/100
57/57 _____ 0s 3ms/step - accuracy: 0.8942 - loss:
0.2800 - val_accuracy: 0.9222 - val_loss: 0.2041
Epoch 84/100
57/57 _____ 0s 4ms/step - accuracy: 0.8811 - loss:
0.2776 - val_accuracy: 0.9222 - val_loss: 0.2028
Epoch 85/100
57/57 _____ 0s 5ms/step - accuracy: 0.8862 - loss:
0.2976 - val_accuracy: 0.9200 - val_loss: 0.2033
Epoch 86/100
57/57 _____ 1s 4ms/step - accuracy: 0.8890 - loss:
0.2979 - val_accuracy: 0.9222 - val_loss: 0.2033
Epoch 87/100
57/57 _____ 0s 3ms/step - accuracy: 0.8804 - loss:
0.2860 - val_accuracy: 0.9200 - val_loss: 0.2031
Epoch 88/100
57/57 _____ 0s 4ms/step - accuracy: 0.8875 - loss:
0.2918 - val_accuracy: 0.9222 - val_loss: 0.2023
Epoch 89/100
57/57 _____ 0s 4ms/step - accuracy: 0.8892 - loss:
0.2820 - val_accuracy: 0.9244 - val_loss: 0.2019
Epoch 90/100
57/57 _____ 0s 4ms/step - accuracy: 0.9021 - loss:
0.2682 - val_accuracy: 0.9244 - val_loss: 0.1998
Epoch 91/100
57/57 _____ 0s 4ms/step - accuracy: 0.8844 - loss:
0.2870 - val_accuracy: 0.9244 - val_loss: 0.2001
Epoch 92/100
57/57 _____ 0s 4ms/step - accuracy: 0.8900 - loss:
0.2953 - val_accuracy: 0.9244 - val_loss: 0.1996
Epoch 93/100
57/57 _____ 0s 4ms/step - accuracy: 0.8903 - loss:
0.2961 - val_accuracy: 0.9267 - val_loss: 0.1993
Epoch 94/100
57/57 _____ 0s 3ms/step - accuracy: 0.8890 - loss:
0.2656 - val_accuracy: 0.9222 - val_loss: 0.2003
Epoch 95/100
57/57 _____ 0s 3ms/step - accuracy: 0.8870 - loss:
0.2868 - val_accuracy: 0.9222 - val_loss: 0.2008
Epoch 96/100
57/57 _____ 0s 3ms/step - accuracy: 0.8947 - loss:
0.2769 - val_accuracy: 0.9244 - val_loss: 0.2005
Epoch 97/100
57/57 _____ 0s 4ms/step - accuracy: 0.8941 - loss:
0.2807 - val_accuracy: 0.9222 - val_loss: 0.2008

```
Epoch 98/100
57/57 _____ 0s 4ms/step - accuracy: 0.9024 - loss:
0.2739 - val_accuracy: 0.9222 - val_loss: 0.1999
Epoch 99/100
57/57 _____ 0s 3ms/step - accuracy: 0.8788 - loss:
0.2819 - val_accuracy: 0.9244 - val_loss: 0.2020
Epoch 100/100
57/57 _____ 0s 3ms/step - accuracy: 0.8769 - loss:
0.2995 - val_accuracy: 0.9244 - val_loss: 0.2025
```

Predictions

```
y_pred_nn = (model.predict(X_test) > 0.5).astype("int32").flatten()
y_prob_nn = model.predict(X_test).flatten()
```

Metrics

```
print("Confusion Matrix:\n", confusion_matrix(y_test, y_pred_nn))
print("\nClassification Report:\n", classification_report(y_test,
y_pred_nn))
print("\nROC AUC Score:", roc_auc_score(y_test, y_prob_nn))
```

```
24/24 _____ 0s 5ms/step
24/24 _____ 0s 3ms/step
```

Confusion Matrix:

```
[[351  24]
 [ 21 354]]
```

Classification Report:

	precision	recall	f1-score	support
0	0.94	0.94	0.94	375
1	0.94	0.94	0.94	375
accuracy			0.94	750
macro avg	0.94	0.94	0.94	750
weighted avg	0.94	0.94	0.94	750

ROC AUC Score: 0.9806079999999999

```
# Compute false positive rate, true positive rate, and thresholds
fpr, tpr, thresholds = roc_curve(y_test, y_prob_nn)
```

Compute AUC

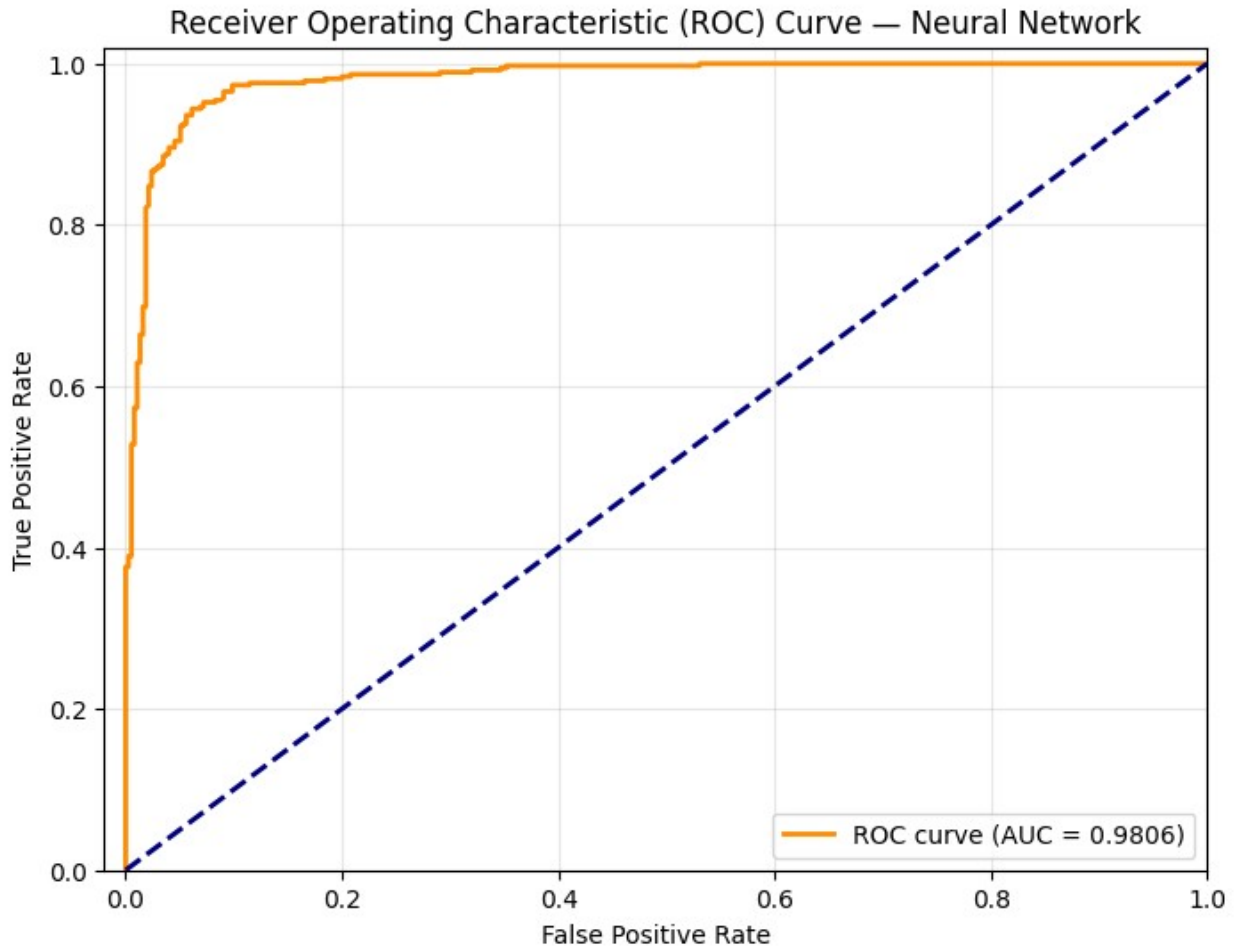
```
roc_auc = auc(fpr, tpr)
```

Plot

```
plt.figure(figsize=(8, 6))
plt.plot(fpr, tpr, color='darkorange', lw=2,
         label=f'ROC curve (AUC = {roc_auc:.4f})')
plt.plot([0, 1], [0, 1], color='navy', lw=2, linestyle='--')
plt.xlim([-0.02, 1.0])
```



```
plt.ylim([0.0, 1.02])
plt.xlabel('False Positive Rate')
plt.ylabel('True Positive Rate')
plt.title('Receiver Operating Characteristic (ROC) Curve — Neural Network')
plt.legend(loc="lower right")
plt.grid(alpha=0.3)
plt.show()
```



```
plt.figure(figsize=(12, 5))

plt.subplot(1, 2, 1)
plt.plot(history.history['accuracy'], label='Train Acc')
plt.plot(history.history['val_accuracy'], label='Val Acc')
plt.title('Model Accuracy')
plt.xlabel('Epochs')
plt.ylabel('Accuracy')
plt.legend()

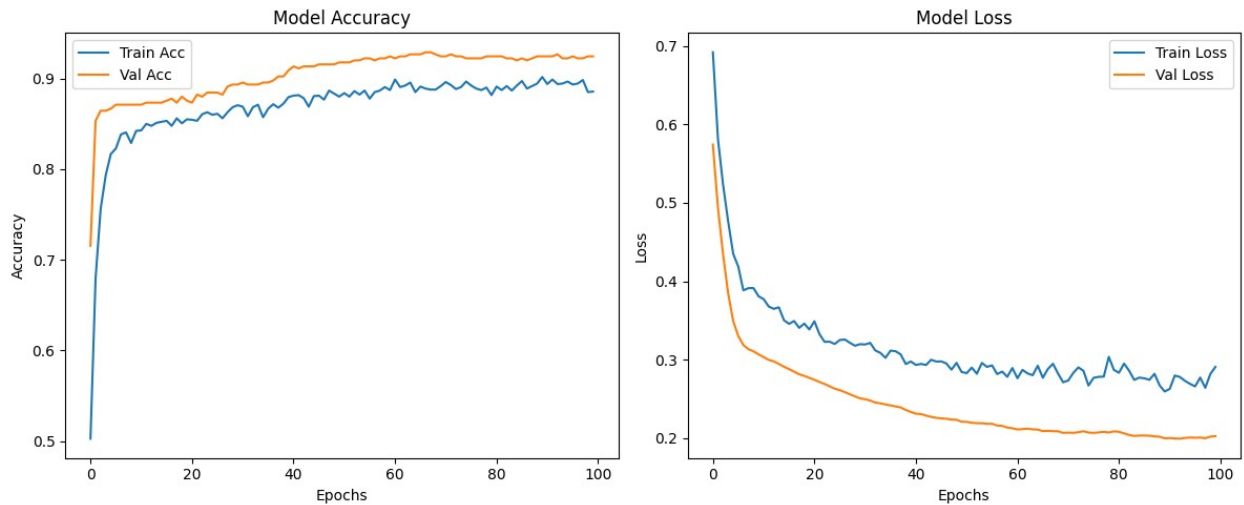
plt.subplot(1, 2, 2)
```

```

plt.plot(history.history['loss'], label='Train Loss')
plt.plot(history.history['val_loss'], label='Val Loss')
plt.title('Model Loss')
plt.xlabel('Epochs')
plt.ylabel('Loss')
plt.legend()

plt.tight_layout()
plt.show()

```

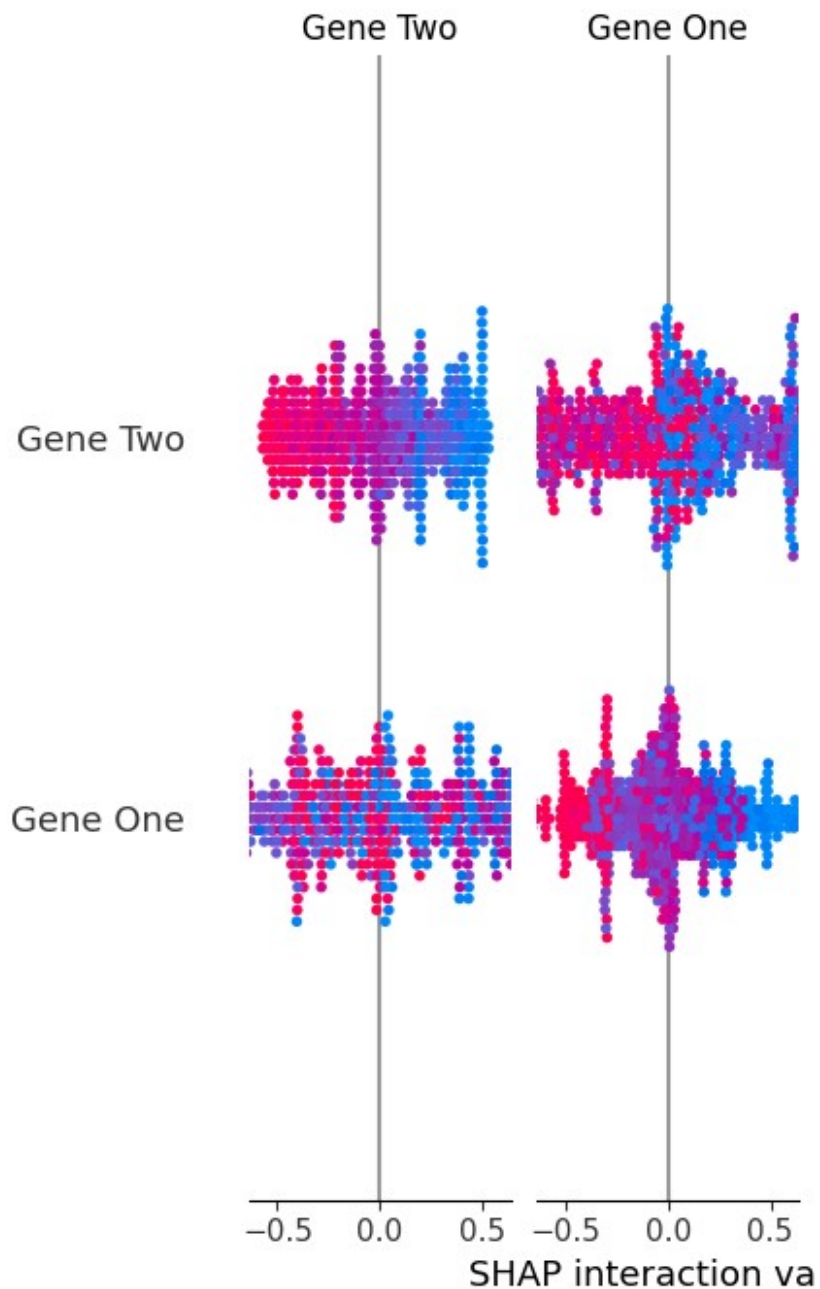


```

explainer_rf = shap.TreeExplainer(rf)
shap_values_rf = explainer_rf.shap_values(X_test)

shap.summary_plot(shap_values_rf, features=X_test,
feature_names=['Gene One', 'Gene Two'])

```

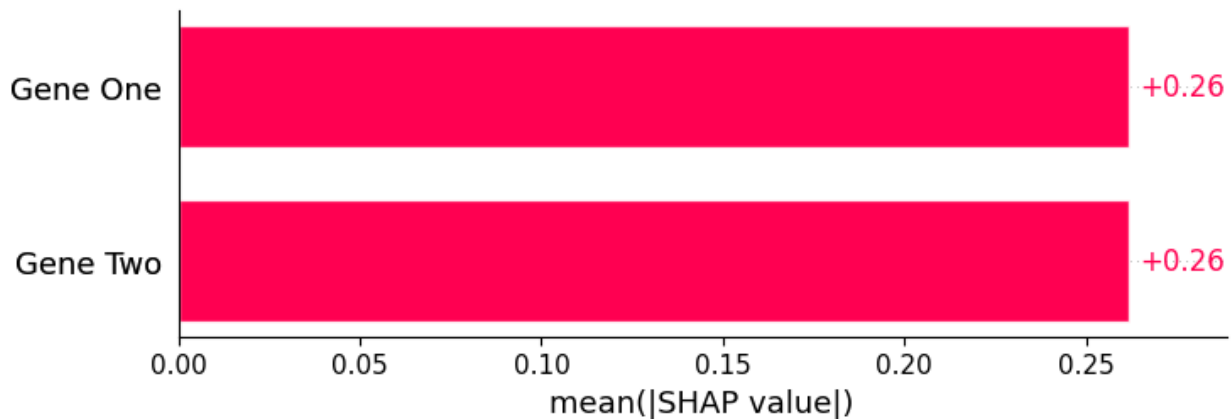


```

explanation = shap.Explanation(
    values=shap_values_rf[1],
    base_values=explainer_rf.expected_value[1],
    data=X_test,
    feature_names=['Gene One', 'Gene Two']
)

# Display the chart
shap.plots.bar(explanation)

```



This bar chart ranks features by their average absolute impact on predictions, providing a clear view of global importance across all predictions.

```
# Subset for KernelExplainer
X_sample = X_test[:100]

# Background dataset for SHAP
background = X_train[np.random.choice(X_train.shape[0], 100,
replace=False)]

# Create KernelExplainer
explainer_nn = shap.KernelExplainer(model.predict, background)
shap_values_nn = explainer_nn.shap_values(X_sample, nsamples=100)

4/4 ————— 0s 23ms/step

{"model_id": "0d0c4aa3530f47a39b2e209d29d77337", "version_major": 2, "version_minor": 0}

1/1 ————— 0s 111ms/step
7/7 ————— 0s 17ms/step
1/1 ————— 0s 144ms/step
7/7 ————— 0s 6ms/step
1/1 ————— 0s 56ms/step
7/7 ————— 0s 6ms/step
1/1 ————— 0s 58ms/step
7/7 ————— 0s 9ms/step
1/1 ————— 0s 59ms/step
7/7 ————— 0s 8ms/step
1/1 ————— 0s 87ms/step
7/7 ————— 0s 7ms/step
1/1 ————— 0s 65ms/step
7/7 ————— 0s 6ms/step
1/1 ————— 0s 57ms/step
7/7 ————— 0s 12ms/step
1/1 ————— 0s 93ms/step
7/7 ————— 0s 6ms/step
```

1/1	_____	0s	56ms/step
7/7	_____	0s	6ms/step
1/1	_____	0s	54ms/step
7/7	_____	0s	7ms/step
1/1	_____	0s	56ms/step
7/7	_____	0s	7ms/step
1/1	_____	0s	54ms/step
7/7	_____	0s	7ms/step
1/1	_____	0s	116ms/step
7/7	_____	0s	6ms/step
1/1	_____	0s	56ms/step
7/7	_____	0s	6ms/step
1/1	_____	0s	81ms/step
7/7	_____	0s	7ms/step
1/1	_____	0s	60ms/step
7/7	_____	0s	8ms/step
1/1	_____	0s	50ms/step
7/7	_____	0s	5ms/step
1/1	_____	0s	39ms/step
7/7	_____	0s	5ms/step
1/1	_____	0s	38ms/step
7/7	_____	0s	4ms/step
1/1	_____	0s	50ms/step
7/7	_____	0s	4ms/step
1/1	_____	0s	37ms/step
7/7	_____	0s	18ms/step
1/1	_____	0s	39ms/step
7/7	_____	0s	4ms/step
1/1	_____	0s	53ms/step
7/7	_____	0s	4ms/step
1/1	_____	0s	42ms/step
7/7	_____	0s	5ms/step
1/1	_____	0s	35ms/step
7/7	_____	0s	5ms/step
1/1	_____	0s	39ms/step
7/7	_____	0s	5ms/step
1/1	_____	0s	35ms/step
7/7	_____	0s	4ms/step
1/1	_____	0s	36ms/step
7/7	_____	0s	4ms/step
1/1	_____	0s	37ms/step
7/7	_____	0s	4ms/step
1/1	_____	0s	36ms/step
7/7	_____	0s	4ms/step
1/1	_____	0s	60ms/step
7/7	_____	0s	4ms/step
1/1	_____	0s	38ms/step
7/7	_____	0s	4ms/step
1/1	_____	0s	39ms/step

7/7	_____	0s 4ms/step
1/1	_____	0s 43ms/step
7/7	_____	0s 5ms/step
1/1	_____	0s 63ms/step
7/7	_____	0s 16ms/step
1/1	_____	0s 60ms/step
7/7	_____	0s 7ms/step
1/1	_____	0s 52ms/step
7/7	_____	0s 7ms/step
1/1	_____	0s 53ms/step
7/7	_____	0s 6ms/step
1/1	_____	0s 54ms/step
7/7	_____	0s 22ms/step
1/1	_____	0s 83ms/step
7/7	_____	0s 14ms/step
1/1	_____	0s 103ms/step
7/7	_____	0s 14ms/step
1/1	_____	0s 57ms/step
7/7	_____	0s 8ms/step
1/1	_____	0s 57ms/step
7/7	_____	0s 7ms/step
1/1	_____	0s 93ms/step
7/7	_____	0s 7ms/step
1/1	_____	0s 91ms/step
7/7	_____	0s 7ms/step
1/1	_____	0s 73ms/step
7/7	_____	0s 7ms/step
1/1	_____	0s 62ms/step
7/7	_____	0s 7ms/step
1/1	_____	0s 70ms/step
7/7	_____	0s 7ms/step
1/1	_____	0s 158ms/step
7/7	_____	0s 13ms/step
1/1	_____	0s 71ms/step
7/7	_____	0s 7ms/step
1/1	_____	0s 55ms/step
7/7	_____	0s 7ms/step
1/1	_____	0s 58ms/step
7/7	_____	0s 7ms/step
1/1	_____	0s 57ms/step
7/7	_____	0s 12ms/step
1/1	_____	0s 57ms/step
7/7	_____	0s 7ms/step
1/1	_____	0s 60ms/step
7/7	_____	0s 7ms/step
1/1	_____	0s 59ms/step
7/7	_____	0s 13ms/step
1/1	_____	0s 56ms/step
7/7	_____	0s 7ms/step

1/1	_____	0s 59ms/step
7/7	_____	0s 6ms/step
1/1	_____	0s 62ms/step
7/7	_____	0s 7ms/step
1/1	_____	0s 63ms/step
7/7	_____	0s 7ms/step
1/1	_____	0s 62ms/step
7/7	_____	0s 7ms/step
1/1	_____	0s 93ms/step
7/7	_____	0s 4ms/step
1/1	_____	0s 38ms/step
7/7	_____	0s 4ms/step
1/1	_____	0s 38ms/step
7/7	_____	0s 4ms/step
1/1	_____	0s 37ms/step
7/7	_____	0s 4ms/step
1/1	_____	0s 41ms/step
7/7	_____	0s 4ms/step
1/1	_____	0s 37ms/step
7/7	_____	0s 14ms/step
1/1	_____	0s 39ms/step
7/7	_____	0s 5ms/step
1/1	_____	0s 37ms/step
7/7	_____	0s 4ms/step
1/1	_____	0s 59ms/step
7/7	_____	0s 8ms/step
1/1	_____	0s 98ms/step
7/7	_____	0s 6ms/step
1/1	_____	0s 58ms/step
7/7	_____	0s 6ms/step
1/1	_____	0s 57ms/step
7/7	_____	0s 6ms/step
1/1	_____	0s 59ms/step
7/7	_____	0s 6ms/step
1/1	_____	0s 67ms/step
7/7	_____	0s 25ms/step
1/1	_____	0s 57ms/step
7/7	_____	0s 5ms/step
1/1	_____	0s 54ms/step
7/7	_____	0s 9ms/step
1/1	_____	0s 58ms/step
7/7	_____	0s 6ms/step
1/1	_____	0s 57ms/step
7/7	_____	0s 7ms/step
1/1	_____	0s 90ms/step
7/7	_____	0s 8ms/step
1/1	_____	0s 59ms/step
7/7	_____	0s 6ms/step
1/1	_____	0s 142ms/step

```

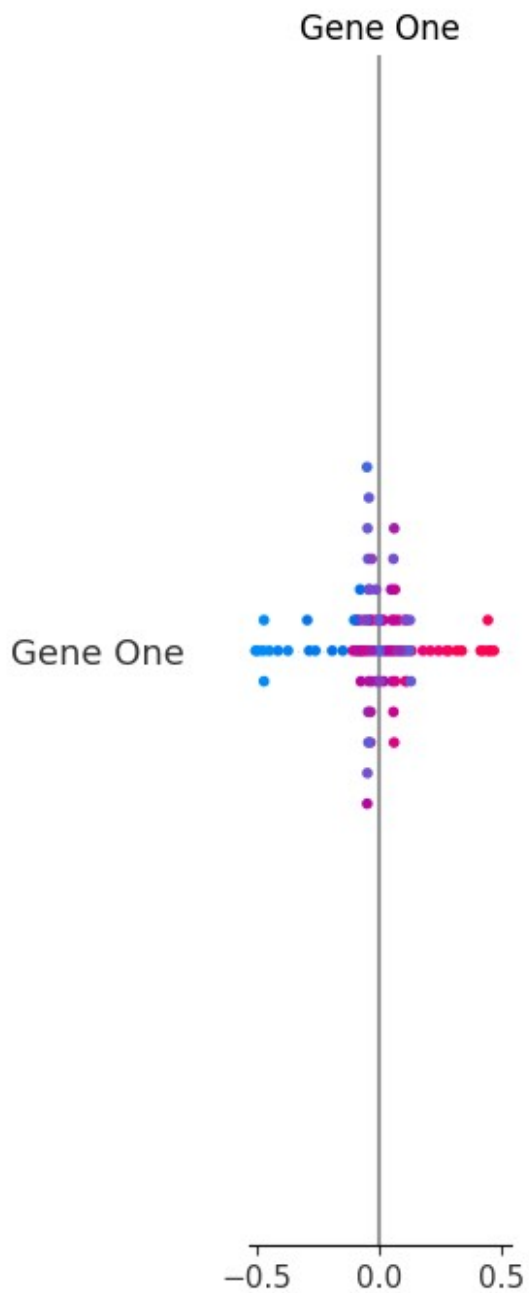
7/7 _____ 0s 6ms/step
1/1 _____ 0s 56ms/step
7/7 _____ 0s 7ms/step
1/1 _____ 0s 68ms/step
7/7 _____ 0s 17ms/step
1/1 _____ 0s 65ms/step
7/7 _____ 0s 8ms/step
1/1 _____ 0s 54ms/step
7/7 _____ 0s 8ms/step
1/1 _____ 0s 101ms/step
7/7 _____ 0s 7ms/step
1/1 _____ 0s 52ms/step
7/7 _____ 0s 7ms/step
1/1 _____ 0s 63ms/step
7/7 _____ 0s 8ms/step
1/1 _____ 0s 112ms/step
7/7 _____ 0s 7ms/step
1/1 _____ 0s 60ms/step
7/7 _____ 0s 7ms/step
1/1 _____ 0s 53ms/step
7/7 _____ 0s 8ms/step
1/1 _____ 0s 64ms/step
7/7 _____ 0s 8ms/step
1/1 _____ 0s 61ms/step
7/7 _____ 0s 14ms/step
1/1 _____ 0s 58ms/step
7/7 _____ 0s 8ms/step
1/1 _____ 0s 76ms/step
7/7 _____ 0s 27ms/step
1/1 _____ 0s 186ms/step
7/7 _____ 0s 20ms/step
1/1 _____ 0s 145ms/step
7/7 _____ 1s 64ms/step
1/1 _____ 0s 263ms/step
7/7 _____ 0s 31ms/step

```

```

# For binary classification, KernelExplainer returns one array:
shap.summary_plot(shap_values_nn, features=X_sample,
feature_names=['Gene One', 'Gene Two'])

```

```
# Compute correlation matrix
gene_corr = df[['Gene One', 'Gene Two']].corr()

# Initialized the graph
G = nx.Graph()

# Add nodes
for gene in gene_corr.columns:
    G.add_node(gene)
```

```

# Add edges for correlations above a threshold
threshold = 0.2
for i in gene_corr.columns:
    for j in gene_corr.columns:
        if i != j and abs(gene_corr.loc[i, j]) > threshold:
            G.add_edge(i, j, weight=gene_corr.loc[i, j])

# Generate positions
pos = nx.spring_layout(G, seed=42)

# Plot nodes
plt.figure(figsize=(6, 4))
nx.draw_networkx_nodes(G, pos, node_color='skyblue', node_size=1000)

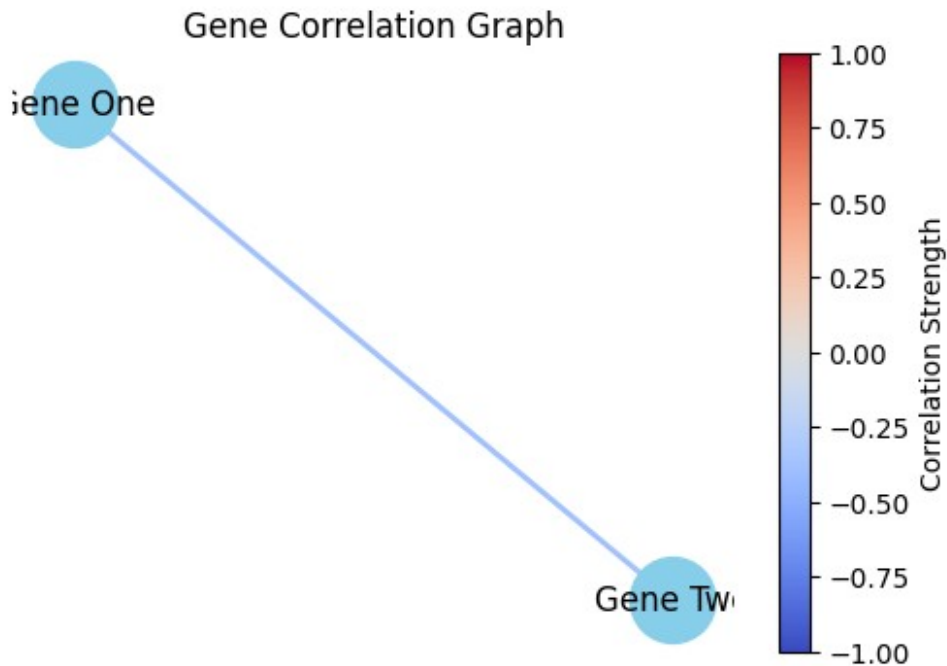
# Add labels to nodes
nx.draw_networkx_labels(G, pos, font_size=12)

# Plot edges with colors
edges = nx.draw_networkx_edges(
    G, pos, width=2,
    edge_color=[G[u][v]['weight'] for u, v in G.edges()],
    edge_cmap=plt.cm.coolwarm,
    edge_vmin=-1, edge_vmax=1
)

# Add colorbar
plt.colorbar(edges, label='Correlation Strength')

plt.title("Gene Correlation Graph")
plt.axis('off')
plt.show()

```



Gene One and Gene Two are negatively correlated, meaning as the expression of one increases, the expression of the other tends to decrease.

The correlation is not extremely strong (not dark blue), but it is still meaningfully negative.

Future Work

To expand causal modeling:

Use Bayesian Networks for probabilistic causal inference.

Introduce counterfactual analysis.

Integrate do-calculus for intervention simulation.

Combine causal graphs with SHAP to build counterfactual explanations.