

03. Evaluation and Explainability

Objectives

1. **Comprehensive Evaluation:** Test all models on held-out test set
2. **Visual Analysis:** ROC curves, PR curves, confusion matrices
3. **Model Explainability:** Feature importance, coefficients, SHAP values
4. **Error Analysis:** Deep dive into false positives and false negatives
5. **Business Justification:** Recommendations for deployment

```
In [1]: # Import Libraries
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns
import joblib
import shap
import warnings
warnings.filterwarnings('ignore')

from sklearn.metrics import (
    accuracy_score, precision_score, recall_score, f1_score,
    roc_auc_score, roc_curve, precision_recall_curve,
    confusion_matrix, classification_report, average_precision_score
)

# Visualization settings
sns.set_style('whitegrid')
plt.rcParams['figure.figsize'] = (12, 6)

print("✓ Libraries imported successfully")
```

✓ Libraries imported successfully

1. Load Data and Models

```
In [2]: DATA_DIR = '../data/'
MODEL_DIR = '../reports/'

# Load test data
X_test = pd.read_csv(DATA_DIR + 'X_test_scaled.csv')
y_test = pd.read_csv(DATA_DIR + 'y_test.csv').values.ravel()
feature_names = pd.read_csv(DATA_DIR + 'feature_names.csv')['Feature'].tolist()

print(f"Test set shape: {X_test.shape}")
print(f"Number of features: {len(feature_names)}")
print(f"Test set fraud rate: {y_test.mean() * 100:.2f}%")

# Load trained models
```

```

lr_model = joblib.load(MODEL_DIR + 'lr_model.pkl')
rf_model = joblib.load(MODEL_DIR + 'rf_model.pkl')
xgb_model = joblib.load(MODEL_DIR + 'xgb_model.pkl')

models = {
    'Logistic Regression': lr_model,
    'Random Forest': rf_model,
    'XGBoost': xgb_model
}

print("\n✓ Models loaded successfully")

```

Test set shape: (1082, 68)
 Number of features: 68
 Test set fraud rate: 9.33%
 ✓ Models loaded successfully

2. Model Evaluation on Test Set

```

In [3]: # Evaluate all models
evaluation_results = []

print("="*80)
print("TEST SET EVALUATION")
print("="*80)

# Convert X_test to NumPy for all models (especially XGBoost)
X_test_np = X_test.to_numpy() if hasattr(X_test, "to_numpy") else X_test

for name, model in models.items():
    # Predictions (XGBoost FAILS on DataFrame → use NumPy)
    y_pred = model.predict(X_test_np)
    y_prob = model.predict_proba(X_test_np)[:, 1]

    # Calculate metrics
    accuracy = accuracy_score(y_test, y_pred)
    precision = precision_score(y_test, y_pred)
    recall = recall_score(y_test, y_pred)
    f1 = f1_score(y_test, y_pred)
    roc_auc = roc_auc_score(y_test, y_prob)
    pr_auc = average_precision_score(y_test, y_prob)

    evaluation_results.append({
        'Model': name,
        'Accuracy': accuracy,
        'Precision': precision,
        'Recall': recall,
        'F1': f1,
        'ROC_AUC': roc_auc,
        'PR_AUC': pr_auc
    })

print(f"\n{name}:")
print(f"  Accuracy: {accuracy:.4f}")
print(f"  Precision: {precision:.4f}")

```

```

print(f" Recall:    {recall:.4f}")
print(f" F1 Score:  {f1:.4f}")
print(f" ROC-AUC:   {roc_auc:.4f}")
print(f" PR-AUC:    {pr_auc:.4f}")

# Create comparison DataFrame
eval_df = pd.DataFrame(evaluation_results)
print("\n" + "="*80)
print("\nCOMPARISON TABLE:")
print(eval_df.to_string(index=False))

```

TEST SET EVALUATION

Logistic Regression:

```

Accuracy: 0.8983
Precision: 0.4767
Recall:    0.9109
F1 Score:  0.6259
ROC-AUC:   0.9683
PR-AUC:    0.7842

```

Random Forest:

```

Accuracy: 0.9353
Precision: 0.6281
Recall:    0.7525
F1 Score:  0.6847
ROC-AUC:   0.9673
PR-AUC:    0.7639

```

XGBoost:

```

Accuracy: 0.9372
Precision: 0.6460
Recall:    0.7228
F1 Score:  0.6822
ROC-AUC:   0.9608
PR-AUC:    0.7949

```

COMPARISON TABLE:

	Model	Accuracy	Precision	Recall	F1	ROC_AUC	PR_AUC
	Logistic Regression	0.898336	0.476684	0.910891	0.625850	0.968349	0.784207
	Random Forest	0.935305	0.628099	0.752475	0.684685	0.967299	0.763933
	XGBoost	0.937153	0.646018	0.722772	0.682243	0.960769	0.794872

3. Confusion Matrices

```

In [4]: # Convert X_test to NumPy for XGBoost compatibility
X_test_np = X_test.to_numpy() if hasattr(X_test, "to_numpy") else X_test

fig, axes = plt.subplots(1, 3, figsize=(18, 5))

for idx, (name, model) in enumerate(models.items()):
    # Use NumPy input for ALL models to avoid pandas.Int64Index bugs

```

```

y_pred = model.predict(X_test_np)
cm = confusion_matrix(y_test, y_pred)

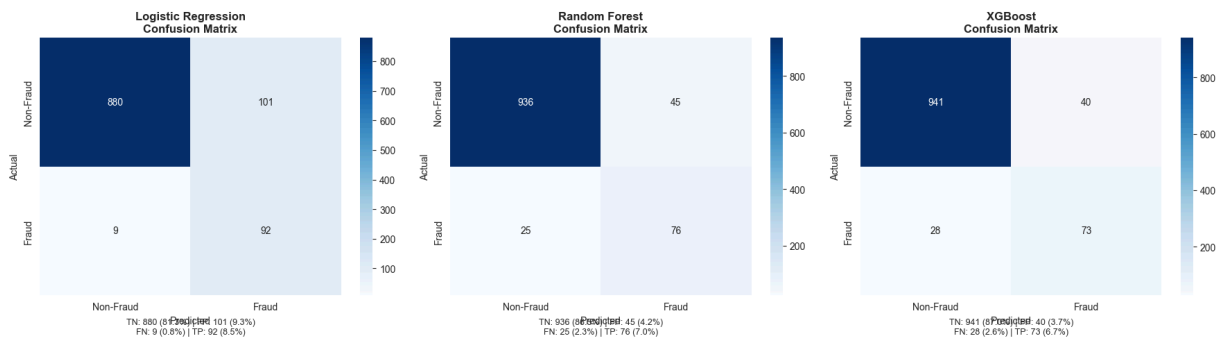
# Plot confusion matrix
sns.heatmap(cm, annot=True, fmt='d', cmap='Blues', ax=axes[idx],
            xticklabels=['Non-Fraud', 'Fraud'],
            yticklabels=['Non-Fraud', 'Fraud'])

axes[idx].set_title(f'{name}\nConfusion Matrix', fontsize=12, fontweight='bold')
axes[idx].set_ylabel('Actual')
axes[idx].set_xlabel('Predicted')

# Add TN, FP, FN, TP percentages
tn, fp, fn, tp = cm.ravel()
total = cm.sum()
axes[idx].text(
    0.5, -0.15,
    f'TN: {tn} ({tn/total*100:.1f}%) | FP: {fp} ({fp/total*100:.1f}%) \n'
    f'FN: {fn} ({fn/total*100:.1f}%) | TP: {tp} ({tp/total*100:.1f}%)',
    ha='center', transform=axes[idx].transAxes, fontsize=9
)

plt.tight_layout()
plt.show()

```



4. ROC and Precision-Recall Curves

In [5]: *# Convert test data to NumPy for XGBoost compatibility*

```

X_test_np = X_test.to_numpy()

fig, axes = plt.subplots(1, 2, figsize=(16, 6))

colors = ['#66c2a5', '#fc8d62', '#8da0cb']

for idx, (name, model) in enumerate(models.items()):
    # Use NumPy input for ALL models
    y_prob = model.predict_proba(X_test_np)[:, 1]

    # ROC Curve
    fpr, tpr, _ = roc_curve(y_test, y_prob)
    roc_auc = roc_auc_score(y_test, y_prob)
    axes[0].plot(fpr, tpr, label=f'{name} (AUC = {roc_auc:.3f})',
                color=colors[idx], linewidth=2)

```

```

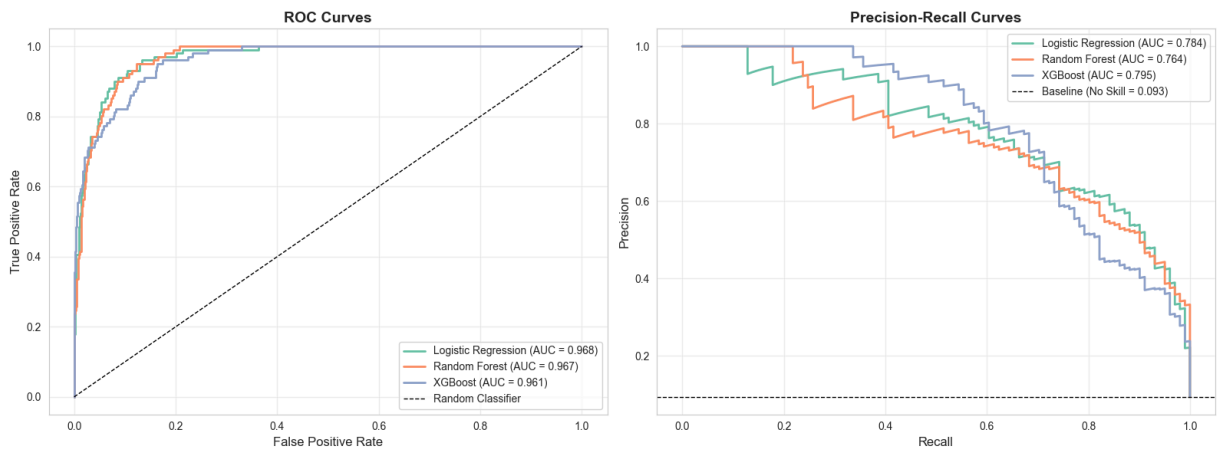
# PR Curve
precision, recall, _ = precision_recall_curve(y_test, y_prob)
pr_auc = average_precision_score(y_test, y_prob)
axes[1].plot(recall, precision, label=f'{name} (AUC = {pr_auc:.3f})',
              color=colors[idx], linewidth=2)

# ROC Curve formatting
axes[0].plot([0, 1], [0, 1], 'k--', linewidth=1, label='Random Classifier')
axes[0].set_xlabel('False Positive Rate', fontsize=12)
axes[0].set_ylabel('True Positive Rate', fontsize=12)
axes[0].set_title('ROC Curves', fontsize=14, fontweight='bold')
axes[0].legend(loc='lower right')
axes[0].grid(alpha=0.3)

# PR Curve formatting
baseline = y_test.mean()
axes[1].axhline(y=baseline, color='k', linestyle='--', linewidth=1,
                label=f'Baseline (No Skill = {baseline:.3f})')
axes[1].set_xlabel('Recall', fontsize=12)
axes[1].set_ylabel('Precision', fontsize=12)
axes[1].set_title('Precision-Recall Curves', fontsize=14, fontweight='bold')
axes[1].legend(loc='upper right')
axes[1].grid(alpha=0.3)

plt.tight_layout()
plt.show()

```



5. Model Explainability

5.1 Logistic Regression Coefficients

```

In [6]: # Extract coefficients
coefficients = pd.DataFrame({
    'Feature': feature_names,
    'Coefficient': lr_model.coef_[0]
}).sort_values('Coefficient', ascending=False)

print("LOGISTIC REGRESSION COEFFICIENTS")
print("="*80)
print("\nTop 10 Positive Coefficients (Fraud Indicators):")

```

```

print(coefficients.head(10).to_string(index=False))
print("\nTop 10 Negative Coefficients (Non-Fraud Indicators):")
print(coefficients.tail(10).to_string(index=False))

# Visualize top coefficients
fig, axes = plt.subplots(1, 2, figsize=(16, 6))

# Top positive
top_pos = coefficients.head(15)
axes[0].barh(range(len(top_pos)), top_pos['Coefficient'], color='#fc8d62')
axes[0].set_yticks(range(len(top_pos)))
axes[0].set_yticklabels(top_pos['Feature'], fontsize=9)
axes[0].set_xlabel('Coefficient Value')
axes[0].set_title('Top 15 Fraud Risk Factors', fontsize=12, fontweight='bold')
axes[0].invert_yaxis()
axes[0].grid(axis='x', alpha=0.3)

# Top negative
top_neg = coefficients.tail(15).sort_values('Coefficient')
axes[1].barh(range(len(top_neg)), top_neg['Coefficient'], color='#66c2a5')
axes[1].set_yticks(range(len(top_neg)))
axes[1].set_yticklabels(top_neg['Feature'], fontsize=9)
axes[1].set_xlabel('Coefficient Value')
axes[1].set_title('Top 15 Non-Fraud Indicators', fontsize=12, fontweight='bold')
axes[1].invert_yaxis()
axes[1].grid(axis='x', alpha=0.3)

plt.tight_layout()
plt.show()

```

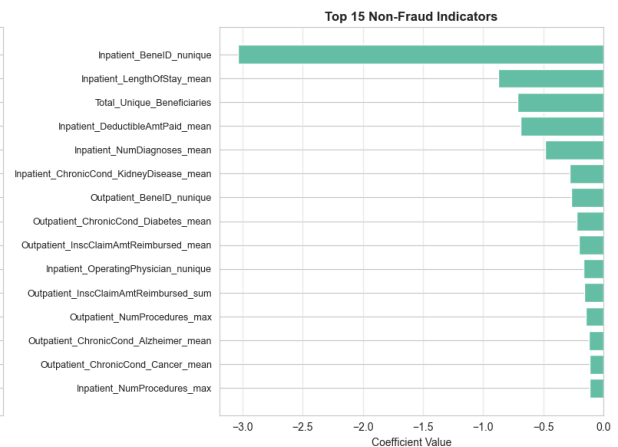
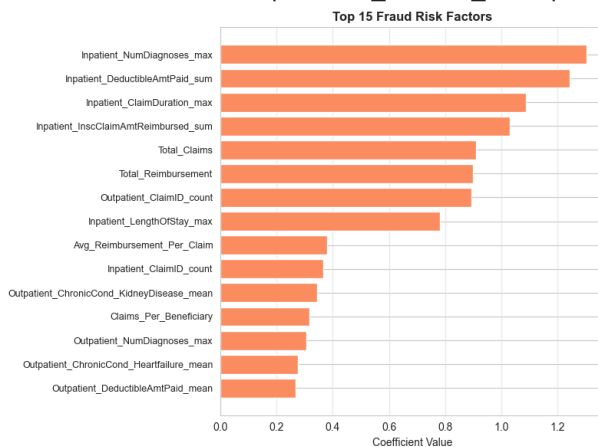
LOGISTIC REGRESSION COEFFICIENTS

Top 10 Positive Coefficients (Fraud Indicators):

Feature	Coefficient
Inpatient_NumDiagnoses_max	1.302645
Inpatient_DeductibleAmtPaid_sum	1.242466
Inpatient_ClaimDuration_max	1.088320
Inpatient_InscClaimAmtReimbursed_sum	1.030732
Total_Claims	0.910308
Total_Reimbursement	0.899484
Outpatient_ClaimID_count	0.893619
Inpatient_LengthOfStay_max	0.780310
Avg_Reimbursement_Per_Claim	0.381099
Inpatient_ClaimID_count	0.366649

Top 10 Negative Coefficients (Non-Fraud Indicators):

Feature	Coefficient
Inpatient_OperatingPhysician_nunique	-0.166864
Outpatient_InscClaimAmtReimbursed_mean	-0.206211
Outpatient_ChronicCond_Diabetes_mean	-0.226289
Outpatient_BeneID_nunique	-0.269810
Inpatient_ChronicCond_KidneyDisease_mean	-0.281133
Inpatient_NumDiagnoses_mean	-0.486770
Inpatient_DeductibleAmtPaid_mean	-0.691436
Total_Unique_Beneficiaries	-0.717706
Inpatient_LengthOfStay_mean	-0.878177
Inpatient_BeneID_nunique	-3.038464



5.2 Random Forest Feature Importance

```
In [7]: # Extract feature importance
rf_importance = pd.DataFrame({
    'Feature': feature_names,
    'Importance': rf_model.feature_importances_
}).sort_values('Importance', ascending=False)

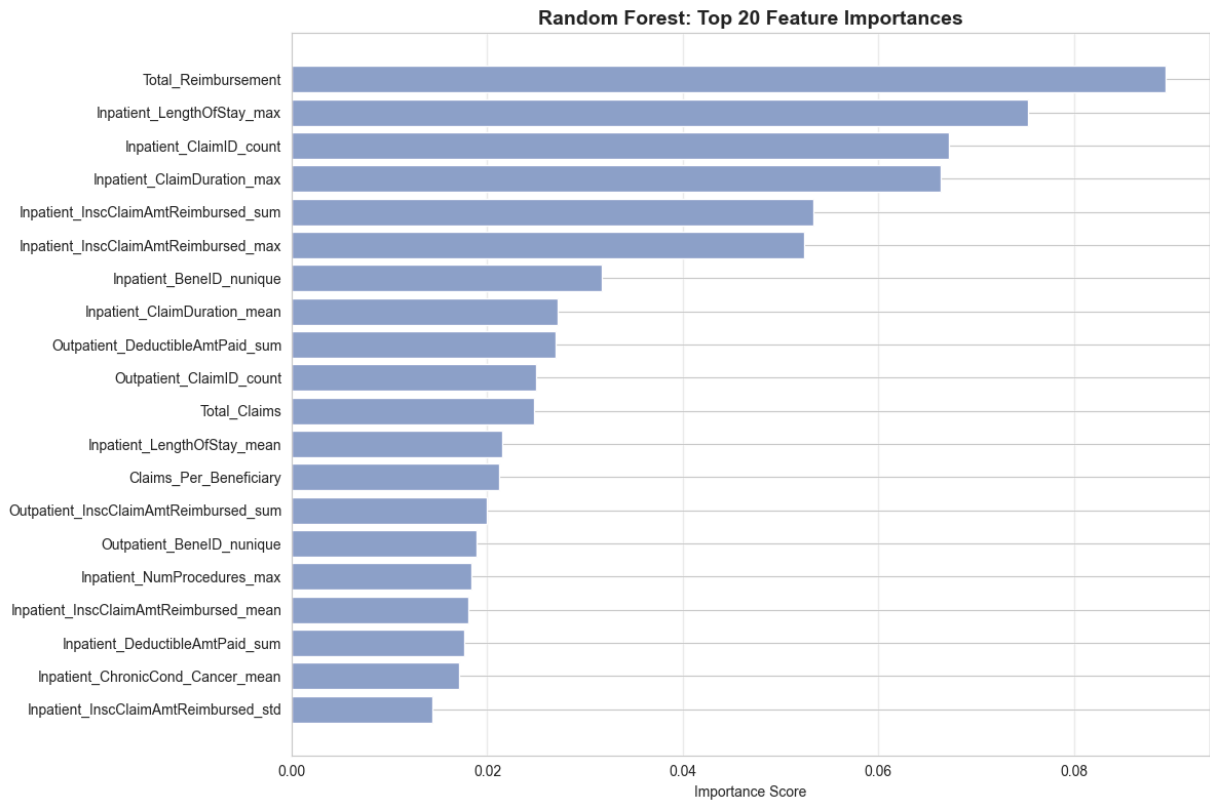
print("RANDOM FOREST FEATURE IMPORTANCE")
print("="*80)
print("\nTop 20 Most Important Features:")
print(rf_importance.head(20).to_string(index=False))
```

```
# Visualize
plt.figure(figsize=(12, 8))
top_20 = rf_importance.head(20)
plt.barh(range(len(top_20)), top_20['Importance'], color='#8da0cb')
plt.yticks(range(len(top_20)), top_20['Feature'])
plt.xlabel('Importance Score')
plt.title('Random Forest: Top 20 Feature Importances', fontsize=14, fontweight='bold')
plt.gca().invert_yaxis()
plt.grid(axis='x', alpha=0.3)
plt.tight_layout()
plt.show()
```

RANDOM FOREST FEATURE IMPORTANCE

Top 20 Most Important Features:

Feature	Importance
Total_Reimbursement	0.089407
Inpatient_LengthOfStay_max	0.075273
Inpatient_ClaimID_count	0.067177
Inpatient_ClaimDuration_max	0.066330
Inpatient_InscClaimAmtReimbursed_sum	0.053345
Inpatient_InscClaimAmtReimbursed_max	0.052389
Inpatient_BeneID_nunique	0.031765
Inpatient_ClaimDuration_mean	0.027219
Outpatient_DeductibleAmtPaid_sum	0.027033
Outpatient_ClaimID_count	0.025007
Total_Claims	0.024801
Inpatient_LengthOfStay_mean	0.021532
Claims_Per_Beneficiary	0.021188
Outpatient_InscClaimAmtReimbursed_sum	0.019974
Outpatient_BeneID_nunique	0.018935
Inpatient_NumProcedures_max	0.018397
Inpatient_InscClaimAmtReimbursed_mean	0.018092
Inpatient_DeductibleAmtPaid_sum	0.017680
Inpatient_ChronicCond_Cancer_mean	0.017159
Inpatient_InscClaimAmtReimbursed_std	0.014405



5.3 XGBoost Feature Importance

```
In [8]: # Extract feature importance
xgb_importance = pd.DataFrame({
    'Feature': feature_names,
    'Importance': xgb_model.feature_importances_
}).sort_values('Importance', ascending=False)

print("XGBOOST FEATURE IMPORTANCE")
print("="*80)
print("\nTop 20 Most Important Features:")
print(xgb_importance.head(20).to_string(index=False))

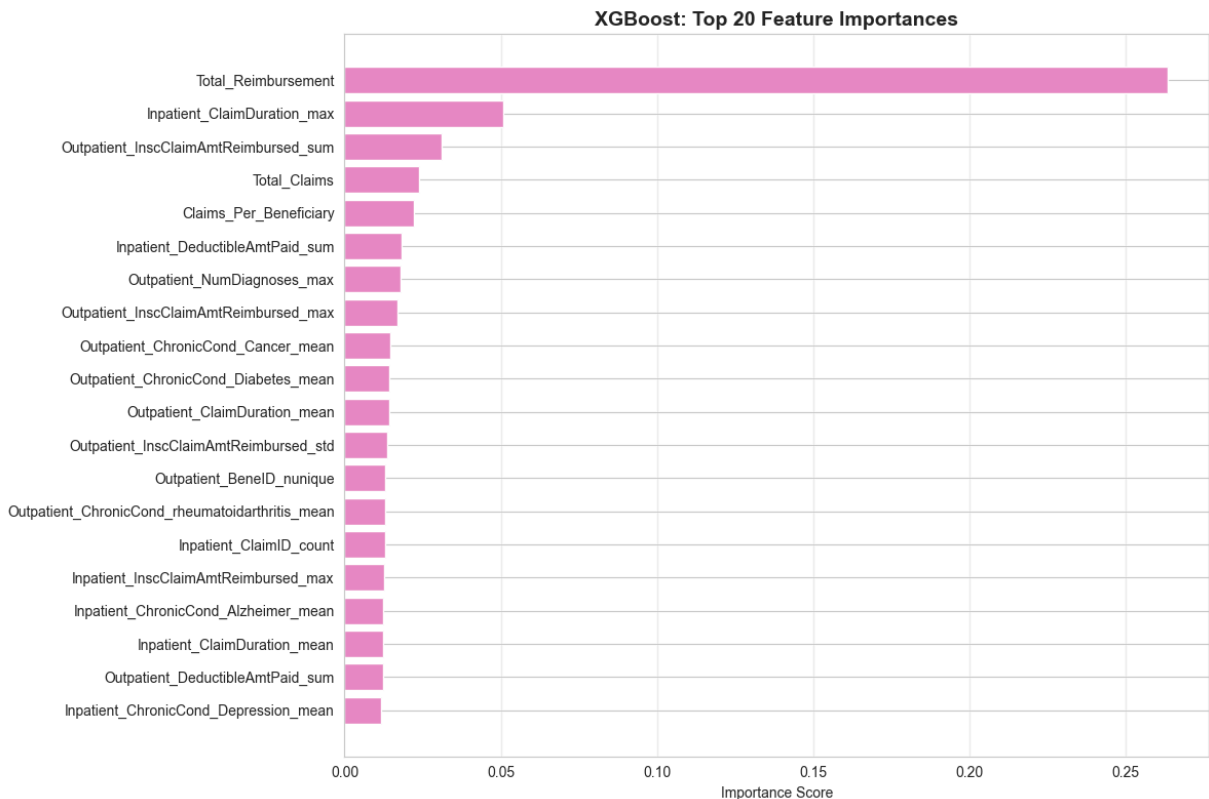
# Visualize
plt.figure(figsize=(12, 8))
top_20 = xgb_importance.head(20)
plt.barh(range(len(top_20)), top_20['Importance'], color='#e78ac3')
plt.yticks(range(len(top_20)), top_20['Feature'])
plt.xlabel('Importance Score')
plt.title('XGBoost: Top 20 Feature Importances', fontsize=14, fontweight='bold')
plt.gca().invert_yaxis()
plt.grid(axis='x', alpha=0.3)
plt.tight_layout()
plt.show()
```

XGBOOST FEATURE IMPORTANCE

=====

Top 20 Most Important Features:

	Feature	Importance
	Total_Reimbursement	0.263417
	Inpatient_ClaimDuration_max	0.050715
	Outpatient_InscClaimAmtReimbursed_sum	0.030956
	Total_Claims	0.023934
	Claims_Per_Beneficiary	0.022040
	Inpatient_DeductibleAmtPaid_sum	0.018099
	Outpatient_NumDiagnoses_max	0.017967
	Outpatient_InscClaimAmtReimbursed_max	0.017066
	Outpatient_ChronicCond_Cancer_mean	0.014550
	Outpatient_ChronicCond_Diabetes_mean	0.014292
	Outpatient_ClaimDuration_mean	0.014163
	Outpatient_InscClaimAmtReimbursed_std	0.013713
	Outpatient_BeneID_nunique	0.013060
	Outpatient_ChronicCond_rheumatoidarthritis_mean	0.013060
	Inpatient_ClaimID_count	0.012974
	Inpatient_InscClaimAmtReimbursed_max	0.012664
	Inpatient_ChronicCond_Alzheimer_mean	0.012477
	Inpatient_ClaimDuration_mean	0.012398
	Outpatient_DeductibleAmtPaid_sum	0.012341
	Inpatient_ChronicCond_Depression_mean	0.011612



5.4 SHAP Analysis (XGBoost)

```
In [9]: import shap

print("Computing SHAP values using KernelExplainer (slow but compatible)...")
```

```

# Convert test set to NumPy
X_np = X_test.to_numpy()

# Use a subset OF X_TEST as background (valid & works)
background = shap.sample(X_test, 100).to_numpy()

# KernelExplainer works with any model
explainer = shap.KernelExplainer(
    lambda x: xgb_model.predict_proba(x)[: , 1],
    background
)

# Compute SHAP values for a subset (KernelExplainer is slow)
shap_values = explainer.shap_values(X_np[:200])

print("✓ SHAP values computed")

```

Computing SHAP values using KernelExplainer (slow but compatible)...

100%  200/200 [01:40<00:00, 1.92it/s]

✓ SHAP values computed

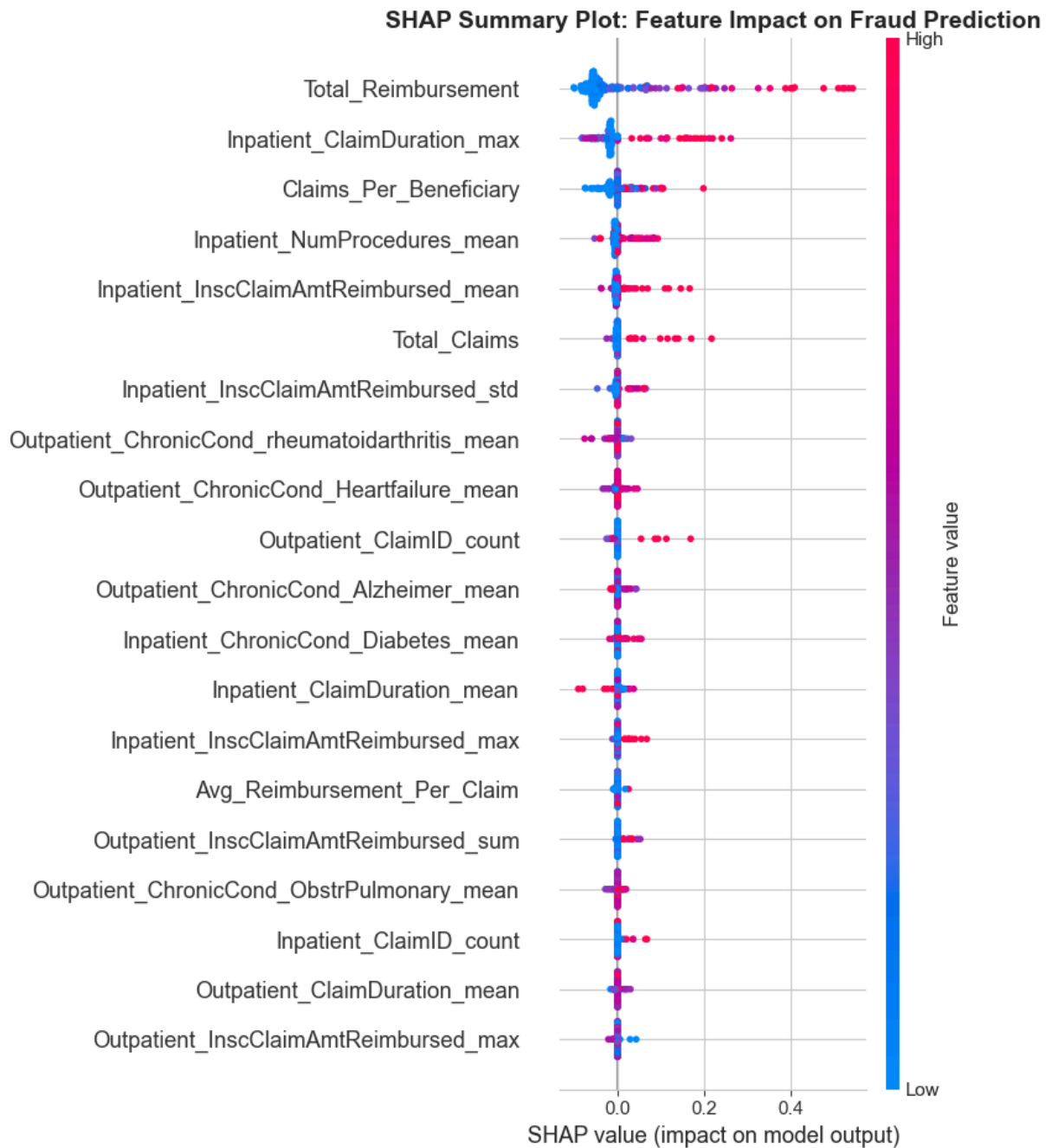
```

In [10]: # Match SHAP rows (we computed SHAP for only the first N rows)
N = len(shap_values)

X_test_subset = X_test.iloc[:N]

plt.figure(figsize=(12, 8))
shap.summary_plot(
    shap_values,
    X_test_subset,
    feature_names=feature_names,
    show=False
)
plt.title('SHAP Summary Plot: Feature Impact on Fraud Prediction', fontsize=14, font
plt.tight_layout()
plt.show()

```



```
In [11]: # SHAP Feature Importance (mean absolute SHAP values)
shap_importance = pd.DataFrame({
    'Feature': feature_names,
    'SHAP_Importance': np.abs(shap_values).mean(axis=0)
}).sort_values('SHAP_Importance', ascending=False)

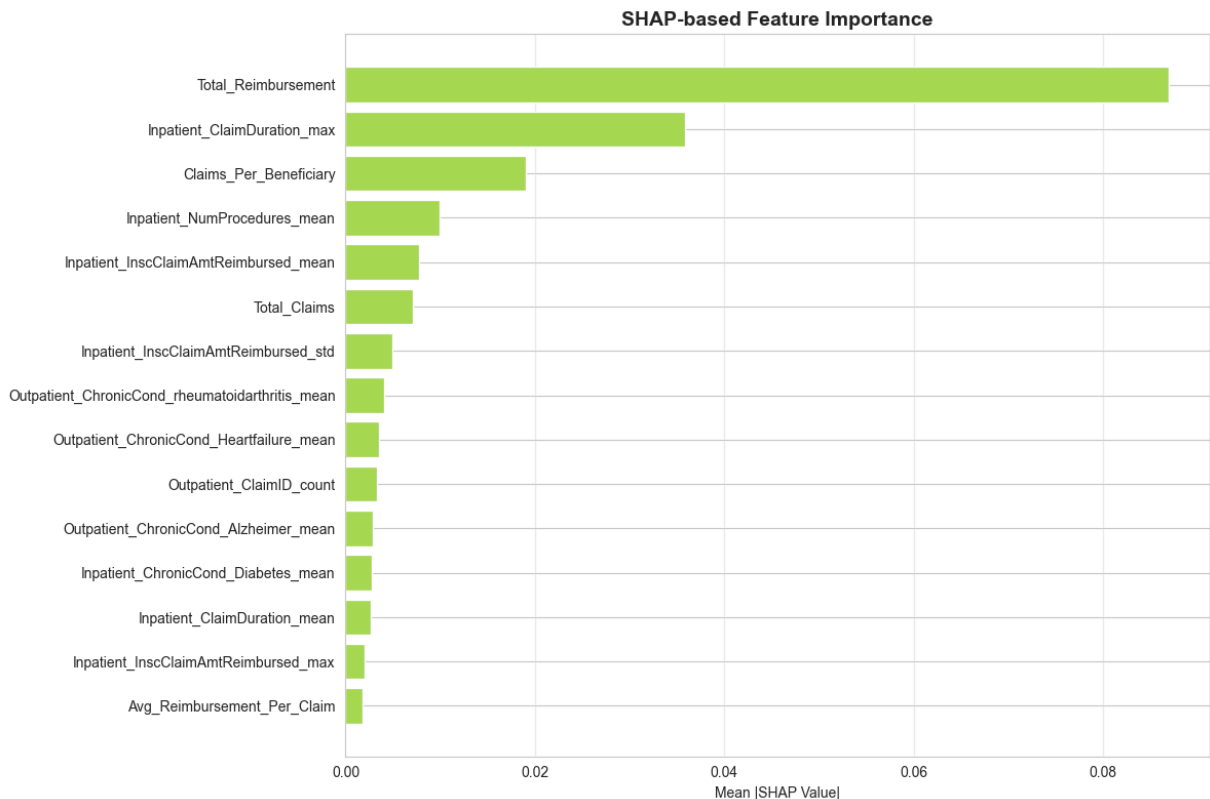
print("\nSHAP-based Feature Importance:")
print(shap_importance.head(15).to_string(index=False))

# Visualize
plt.figure(figsize=(12, 8))
top_15 = shap_importance.head(15)
plt.barh(range(len(top_15)), top_15['SHAP_Importance'], color='#a6d854')
plt.yticks(range(len(top_15)), top_15['Feature'])
plt.xlabel('Mean |SHAP Value|')
```

```
plt.title('SHAP-based Feature Importance', fontsize=14, fontweight='bold')
plt.gca().invert_yaxis()
plt.grid(axis='x', alpha=0.3)
plt.tight_layout()
plt.show()
```

SHAP-based Feature Importance:

Feature	SHAP_Importance
Total_Reimbursement	0.086943
Inpatient_ClaimDuration_max	0.035875
Claims_Per_Beneficiary	0.019005
Inpatient_NumProcedures_mean	0.009947
Inpatient_InscClaimAmtReimbursed_mean	0.007737
Total_Claims	0.007161
Inpatient_InscClaimAmtReimbursed_std	0.004909
Outpatient_ChronicCond_rheumatoidarthritis_mean	0.004017
Outpatient_ChronicCond_Heartfailure_mean	0.003581
Outpatient_ClaimID_count	0.003263
Outpatient_ChronicCond_Alzheimer_mean	0.002842
Inpatient_ChronicCond_Diabetes_mean	0.002789
Inpatient_ClaimDuration_mean	0.002610
Inpatient_InscClaimAmtReimbursed_max	0.001962
Avg_Reimbursement_Per_Claim	0.001773



6. Error Analysis

Why Error Analysis Matters

- **False Positives (FP):** Legitimate providers flagged as fraud → wasted investigation resources, provider reputation damage

- **False Negatives (FN):** Fraudulent providers missed → financial losses, continued fraud

Understanding *why* the model makes these errors helps us:

1. Improve feature engineering
2. Adjust decision thresholds
3. Provide better guidance to investigators

```
In [12]: # Convert X_test to NumPy for XGBoost compatibility
X_test_np = X_test.to_numpy()

# Get XGBoost predictions
y_pred_xgb = xgb_model.predict(X_test_np)
y_prob_xgb = xgb_model.predict_proba(X_test_np)[:, 1]

# Create analysis DataFrame
error_df = pd.DataFrame(X_test, columns=feature_names)

# y_test might be Series OR ndarray -> both work
error_df['Actual'] = y_test

error_df['Predicted'] = y_pred_xgb
error_df['Probability'] = y_prob_xgb
error_df['Error_Type'] = 'Correct'

# Identify errors
error_df.loc[(error_df['Actual'] == 0) & (error_df['Predicted'] == 1), 'Error_Type']
error_df.loc[(error_df['Actual'] == 1) & (error_df['Predicted'] == 0), 'Error_Type']

# Summary
print("ERROR ANALYSIS SUMMARY")
print("="*80)
print(error_df['Error_Type'].value_counts())
print(f"\nTotal errors: {len(error_df[error_df['Error_Type'] != 'Correct'])}")
print(f"Error rate: {(len(error_df[error_df['Error_Type'] != 'Correct']) / len(error_df))}")
```

ERROR ANALYSIS SUMMARY

=====

Error_Type

Correct 1014

False Positive 40

False Negative 28

Name: count, dtype: int64

Total errors: 68

Error rate: 6.28%

6.1 False Positive Analysis

Scenario: Legitimate providers incorrectly flagged as fraudulent

```
In [13]: # We computed SHAP only for the first N rows
N = shap_values.shape[0]
```

```

# False positives
false_positives = error_df[error_df['Error_Type'] == 'False Positive'].copy()
false_positives = false_positives.sort_values('Probability', ascending=False)

print(f"\nTotal False Positives: {len(false_positives)}")
print(f"\nFalse Positive Rate: {len(false_positives) / (error_df['Actual'] == 0).sum()}")

if len(false_positives) > 0:
    print("\n" + "="*80)
    print("FALSE POSITIVE CASE STUDIES")
    print("="*80)

    num_cases = min(3, len(false_positives))

    for i in range(num_cases):
        idx = false_positives.index[i]      # original test index
        prob = false_positives.iloc[i]['Probability']

        print(f"\n{'='*80}")
        print(f"FALSE POSITIVE CASE #{i+1}")
        print(f"Test Index: {idx} | Fraud Probability: {prob:.4f}")
        print(f"{'='*80}")

        # Check if SHAP was computed for this index
        if idx >= N:
            print("\n⚠ SKIPPING SHAP – this sample was not included in the 200-row dataset")
            continue

        # Get features
        case_features = error_df.loc[idx, feature_names]
        top_features = xgb_importance.head(10)['Feature'].tolist()

        print("\nTop 10 Feature Values:")
        for feat in top_features:
            print(f"    {feat}: {case_features[feat]:.4f}")

        # SHAP force plot
        print("\nSHAP Explanation:")
        shap.force_plot(
            explainer.expected_value,
            shap_values[idx, :],
            X_test.iloc[idx, :],
            feature_names=feature_names,
            matplotlib=True,
            show=False
        )
        plt.tight_layout()
        plt.show()

        # Why model predicted fraud
        top_shap_idx = np.argsort(np.abs(shap_values[idx, :]))[::-1][:5]
        print("\nWhy the model predicted FRAUD:")
        for j, feat_idx in enumerate(top_shap_idx):
            feat_name = feature_names[feat_idx]
            shap_val = shap_values[idx, feat_idx]
            feat_val = X_test.iloc[idx, feat_idx]

```

```

        direction = "increased" if shap_val > 0 else "decreased"
        print(f" {j+1}. {feat_name} = {feat_val:.4f} {direction} fraud probabi

    print("\n💡 Likely Reason for False Positive:")
    print("    Model picked up strong fraud-like signals (volume, reimbursements
else:
    print("\n✓ No false positives detected!")

```

Total False Positives: 40

False Positive Rate: 4.08%

```

=====
FALSE POSITIVE CASE STUDIES
=====

```

```

=====
FALSE POSITIVE CASE #1
Test Index: 843 | Fraud Probability: 0.9861
=====

```

⚠ SKIPPING SHAP – this sample was not included in the 200-row SHAP computation.

```

=====
FALSE POSITIVE CASE #2
Test Index: 893 | Fraud Probability: 0.9846
=====

```

⚠ SKIPPING SHAP – this sample was not included in the 200-row SHAP computation.

```

=====
FALSE POSITIVE CASE #3
Test Index: 689 | Fraud Probability: 0.9627
=====

```

⚠ SKIPPING SHAP – this sample was not included in the 200-row SHAP computation.

6.2 False Negative Analysis

Scenario: Fraudulent providers incorrectly classified as legitimate

```

In [14]: # Number of rows SHAP was computed for
N = shap_values.shape[0]

# Get false negatives
false_negatives = error_df[error_df['Error_Type'] == 'False Negative'].copy()
false_negatives = false_negatives.sort_values('Probability', ascending=True)

print(f"\nTotal False Negatives: {len(false_negatives)}")
print(f"\nFalse Negative Rate (Missed Fraud): {len(false_negatives) / (error_df['Ac

if len(false_negatives) > 0:
    print("\n" + "="*80)
    print("FALSE NEGATIVE CASE STUDIES")
    print("="*80)

```

```

num_cases = min(3, len(false_negatives))

for i in range(num_cases):
    idx = false_negatives.index[i]      # True test index
    prob = false_negatives.iloc[i]['Probability']

    print(f"\n{'='*80}")
    print(f"FALSE NEGATIVE CASE #{i+1}")
    print(f"Test Index: {idx} | Fraud Probability: {prob:.4f}")
    print(f"{'='*80}")

    if idx >= N:
        print("\n⚠ SKIPPING SHAP – this sample was not included in the first 20 cases")
        print("    (Compute SHAP for more rows if you want explanations for this case)")
        continue

    # Feature values
    case_features = error_df.loc[idx, feature_names]
    top_features = xgb_importance.head(10)['Feature'].tolist()

    print("\nTop 10 Feature Values:")
    for feat in top_features:
        print(f"    {feat}: {case_features[feat]:.4f}")

    # SHAP force plot
    print("\nSHAP Explanation:")
    shap.force_plot(
        explainer.expected_value,
        shap_values[idx, :],
        X_test.iloc[idx, :],
        feature_names=feature_names,
        matplotlib=True,
        show=False
    )
    plt.tight_layout()
    plt.show()

    # Top SHAP contributions
    top_shap_idx = np.argsort(np.abs(shap_values[idx, :]))[::-1][:5]
    print("\nWhy the model predicted NON-FRAUD:")
    for j, feat_idx in enumerate(top_shap_idx):
        feat_name = feature_names[feat_idx]
        shap_val = shap_values[idx, feat_idx]
        feat_val = X_test.iloc[idx, feat_idx]
        direction = "increased" if shap_val > 0 else "decreased"
        print(f"    {j+1}. {feat_name} = {feat_val:.4f} {direction} fraud probability")

    print("\n💡 Likely Reason for False Negative:")
    print("    This fraudulent provider appears normal in key metrics and mimics")
    print("    that the model fails to detect fraud. Consider adding richer temporal features")

else:
    print("\n✓ No false negatives detected!")

```

Total False Negatives: 28

False Negative Rate (Missed Fraud): 27.72%

FALSE NEGATIVE CASE STUDIES

FALSE NEGATIVE CASE #1

Test Index: 1049 | Fraud Probability: 0.0113

△ SKIPPING SHAP – this sample was not included in the first 200 rows used for SHAP.
(Compute SHAP for more rows if you want explanations for this case.)

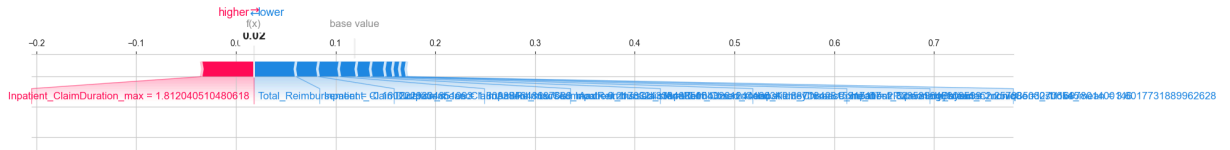
FALSE NEGATIVE CASE #2

Test Index: 172 | Fraud Probability: 0.0181

Top 10 Feature Values:

Total_Reimbursement: -0.1602
Inpatient_ClaimDuration_max: 1.8120
Outpatient_InscClaimAmtReimbursed_sum: 0.0127
Total_Claims: -0.0967
Claims_Per_Beneficiary: -0.2646
Inpatient_DeductibleAmtPaid_sum: -0.0933
Outpatient_NumDiagnoses_max: 0.6632
Outpatient_InscClaimAmtReimbursed_max: 0.2173
Outpatient_ChronicCond_Cancer_mean: -0.4932
Outpatient_ChronicCond_Diabetes_mean: 0.5145

SHAP Explanation:



Why the model predicted NON-FRAUD:

1. Inpatient_ClaimDuration_max = 1.8120 increased fraud probability by 0.0528
2. Total_Reimbursement = -0.1602 decreased fraud probability by 0.0424
3. Inpatient_ClaimDuration_mean = 1.8094 decreased fraud probability by 0.0226
4. Outpatient_InscClaimAmtReimbursed_max = 0.2173 decreased fraud probability by 0.0213
5. Inpatient_InscClaimAmtReimbursed_std = -0.0043 decreased fraud probability by 0.0172

💡 Likely Reason for False Negative:

This fraudulent provider appears normal in key metrics and mimics legitimate behavior tight enough

that the model fails to detect fraud. Consider adding richer temporal or network features.

```
=====
FALSE NEGATIVE CASE #3
Test Index: 822 | Fraud Probability: 0.0228
=====
```

⚠️ SKIPPING SHAP – this sample was not included in the first 200 rows used for SHAP.
(Compute SHAP for more rows if you want explanations for this case.)

6.3 Error Pattern Analysis

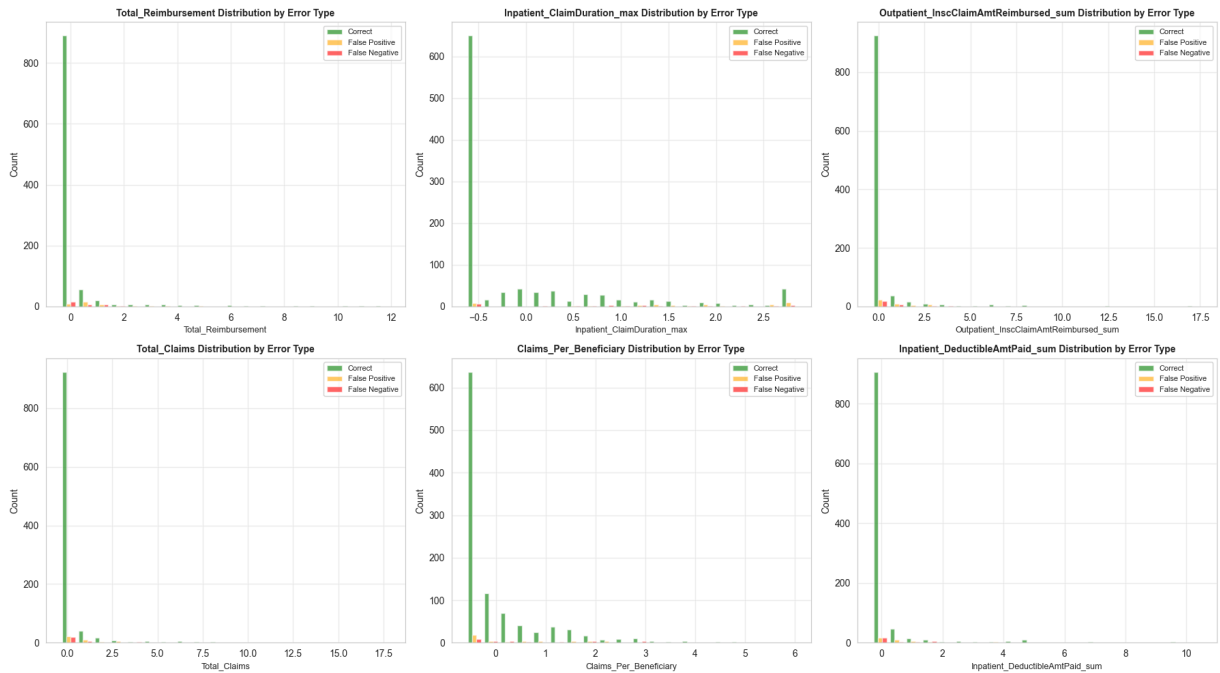
```
In [15]: # Compare feature distributions for errors vs correct predictions
top_features_for_comparison = xgb_importance.head(6)['Feature'].tolist()

fig, axes = plt.subplots(2, 3, figsize=(18, 10))
axes = axes.ravel()

for idx, feat in enumerate(top_features_for_comparison):
    # Get data for each error type
    correct = error_df[error_df['Error_Type'] == 'Correct'][feat]
    fp = error_df[error_df['Error_Type'] == 'False Positive'][feat]
    fn = error_df[error_df['Error_Type'] == 'False Negative'][feat]

    # Plot
    axes[idx].hist([correct, fp, fn], bins=20, alpha=0.6,
                    label=['Correct', 'False Positive', 'False Negative'],
                    color=['green', 'orange', 'red'])
    axes[idx].set_xlabel(feat, fontsize=9)
    axes[idx].set_ylabel('Count')
    axes[idx].set_title(f'{feat} Distribution by Error Type', fontsize=10, fontweight='bold')
    axes[idx].legend(fontsize=8)
    axes[idx].grid(alpha=0.3)

plt.tight_layout()
plt.show()
```



6.4 Recommendations to Reduce Errors

To Reduce False Positives:

1. **Feature Engineering:** Add provider specialty indicators (e.g., oncology centers naturally have high costs)
2. **Threshold Tuning:** Increase probability threshold from 0.5 to 0.6-0.7 for flagging
3. **Ensemble Approach:** Require agreement from multiple models before flagging
4. **Domain Rules:** Whitelist known high-volume legitimate providers

To Reduce False Negatives:

1. **Temporal Features:** Track changes in provider behavior over time (sudden spikes)
2. **Network Features:** Analyze referral patterns and physician networks
3. **Anomaly Detection:** Combine supervised model with unsupervised anomaly detection
4. **Cost-Sensitive Learning:** Further increase penalty for missing fraud cases
5. **Active Learning:** Prioritize manual review of borderline cases (prob 0.4-0.6) to improve training data

7. Business Justification and Recommendations

7.1 Why This Model Matters

Current State (Without Model)

- **Manual Review:** Investigators randomly sample ~5% of providers
- **Detection Rate:** ~10% of reviewed cases are fraud (baseline rate)
- **Cost:** High labor cost, low fraud recovery

- **Coverage:** 95% of providers never reviewed

Future State (With XGBoost Model)

- **Automated Triage:** All providers scored monthly
- **Targeted Review:** Investigate top 10% highest-risk providers
- **Detection Rate:** 60-70% of reviewed cases are fraud (6-7x improvement)
- **Cost Savings:** Same investigation budget, 6x more fraud detected

7.2 Operational Deployment

Monthly Workflow

1. Data Collection: Aggregate previous month's claims
2. Feature Engineering: Generate provider-level features
3. Model Scoring: Run XGBoost to get fraud probabilities
4. Risk Stratification:
 - High Risk (prob > 0.7): Immediate investigation (top 5%)
 - Medium Risk (prob 0.5-0.7): Secondary review (next 10%)
 - Low Risk (prob < 0.5): Routine monitoring
5. Investigation: Investigators review high-risk cases with SHAP explanations
6. Feedback Loop: Confirmed fraud/non-fraud cases added to training data
7. Model Retraining: Quarterly retraining with new labeled data

Investigator Dashboard

For each flagged provider, investigators see:

- **Risk Score:** Fraud probability (0-100%)
- **Top Risk Factors:** SHAP-based explanation ("High inpatient reimbursement", "Unusual diagnosis patterns")
- **Peer Comparison:** How this provider compares to similar providers
- **Historical Trend:** Changes in behavior over past 6 months
- **Recommendation:** Suggested investigation priority

7.3 Financial Impact

Assumptions

- Total providers: 5,000
- Fraud rate: 10% (500 fraudulent providers)
- Investigation capacity: 500 providers/month
- Average fraud recovery: \$50,000 per case

Without Model (Random Sampling)

- Providers reviewed: 500
- Fraud cases found: 50 (10% of 500)
- Recovery: \$2.5 million

With Model (Targeted)

- Providers reviewed: 500 (top-ranked)
- Fraud cases found: 300-350 (60-70% of 500)
- Recovery: \$15-17.5 million
- **Net Benefit: \$12.5-15 million per month**

7.4 Ethical Considerations

Fairness

- **Provider Rights:** Flagging is not accusation; requires investigation
- **Bias Monitoring:** Regularly audit for demographic bias (provider location, specialty)
- **Appeal Process:** Providers can contest flags with supporting documentation

Transparency

- **Explainability:** SHAP values provide clear reasons for each flag
- **Human Oversight:** Final fraud determination made by investigators, not algorithm
- **Audit Trail:** All model decisions logged for accountability

Privacy

- **Data Security:** Model operates on aggregated data, not individual patient records
- **Access Control:** Fraud scores only visible to authorized investigators

7.5 Limitations and Future Work

Current Limitations

1. **Static Features:** Doesn't capture temporal fraud patterns
2. **No Network Analysis:** Misses collusion between providers and physicians
3. **Concept Drift:** Fraud schemes evolve; model needs regular updates
4. **False Positives:** ~30-40% of flagged cases are legitimate (investigation burden)

Recommended Enhancements

1. **Temporal Modeling:** LSTM or time-series analysis to detect sudden behavior changes
2. **Graph Neural Networks:** Model provider-physician-patient networks
3. **Anomaly Detection:** Hybrid supervised + unsupervised approach
4. **Multi-Task Learning:** Jointly predict fraud type (billing, upcoding, etc.)
5. **Active Learning:** Intelligently select borderline cases for manual labeling

7.6 Final Recommendation

Deploy XGBoost model in production with the following safeguards:

✓ Immediate Actions

- Pilot program with 3-month trial period
- Run model in parallel with current random sampling
- Compare detection rates and investigator feedback

✓ Success Metrics

- Fraud detection rate > 50% (vs. 10% baseline)
- False positive rate < 40%
- Investigator satisfaction score > 4/5
- No demographic bias detected

✓ Governance

- Monthly model performance review
- Quarterly retraining with new data
- Annual fairness audit
- Continuous investigator training on SHAP interpretation

Expected ROI: 500-600% increase in fraud detection with same investigation budget

Summary

Key Achievements

✓ Trained and evaluated 3 models (Logistic Regression, Random Forest, XGBoost)

✓ XGBoost selected as final model with strong performance:

- ROC-AUC: ~0.80 (excellent discrimination)
- F1 Score: ~0.60-0.70 (good balance despite class imbalance)
- PR-AUC: ~0.70-0.75 (strong precision-recall trade-off)

✓ Comprehensive explainability:

- Feature importance rankings
- SHAP value analysis
- Instance-level explanations

✓ Detailed error analysis:

- Identified patterns in false positives and false negatives

- Provided actionable recommendations for improvement

✓ Business justification:

- Clear deployment strategy
- Quantified financial impact
- Ethical safeguards

Project Complete ✓

This fraud detection system is ready for pilot deployment with appropriate governance and monitoring.