```
In [82]: %pip install pandas
%pip install matplotlib
%pip install seaborn
%pip install scikit-learn
%pip install nltk
%pip install wordcloud
%pip install emoji
```

```
Requirement already satisfied: pandas in ./.venv/lib/python3.10/site-packa
ges (2.2.3)
Requirement already satisfied: numpy>=1.22.4 in ./.venv/lib/python3.10/sit
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Requirement already satisfied: kiwisolver>=1.3.1 in ./.venv/lib/python3.1
0/site-packages (from matplotlib!=3.6.1,>=3.4->seaborn) (1.4.8)
Requirement already satisfied: packaging>=20.0 in ./.venv/lib/python3.10/s
ite-packages (from matplotlib!=3.6.1,>=3.4->seaborn) (25.0)
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ckages (from matplotlib!=3.6.1,>=3.4->seaborn) (11.2.1)
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site-packages (from matplotlib!=3.6.1,>=3.4->seaborn) (3.2.3)
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3.10/site-packages (from matplotlib!=3.6.1,>=3.4->seaborn) (2.9.0.post0)
```

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Requirement already satisfied: pytz>=2020.1 in ./.venv/lib/python3.10/sit
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        te-packages (from pandas>=1.2->seaborn) (2025.2)
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        s (from nltk) (4.67.1)
        Note: you may need to restart the kernel to use updated packages.
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        site-packages (from matplotlib->wordcloud) (1.3.2)
        Requirement already satisfied: cycler>=0.10 in ./.venv/lib/python3.10/sit
        e-packages (from matplotlib->wordcloud) (0.12.1)
        Requirement already satisfied: fonttools>=4.22.0 in ./.venv/lib/python3.1
        O/site-packages (from matplotlib->wordcloud) (4.57.0)
        Requirement already satisfied: kiwisolver>=1.3.1 in ./.venv/lib/python3.1
        O/site-packages (from matplotlib->wordcloud) (1.4.8)
        Requirement already satisfied: packaging>=20.0 in ./.venv/lib/python3.10/s
        ite-packages (from matplotlib->wordcloud) (25.0)
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        site-packages (from matplotlib->wordcloud) (3.2.3)
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        3.10/site-packages (from matplotlib->wordcloud) (2.9.0.post0)
        Requirement already satisfied: six>=1.5 in ./.venv/lib/python3.10/site-pac
        kages (from python-dateutil>=2.7->matplotlib->wordcloud) (1.17.0)
        Note: you may need to restart the kernel to use updated packages.
        Requirement already satisfied: emoji in ./.venv/lib/python3.10/site-packag
        es (2.14.1)
        Note: you may need to restart the kernel to use updated packages.
In [83]: import matplotlib.pyplot as plt
```

```
import seaborn as sns
import numpy as np

import pandas as pd
df = pd.read_csv("scitweets_export.tsv", sep="\t")
df.head()
```

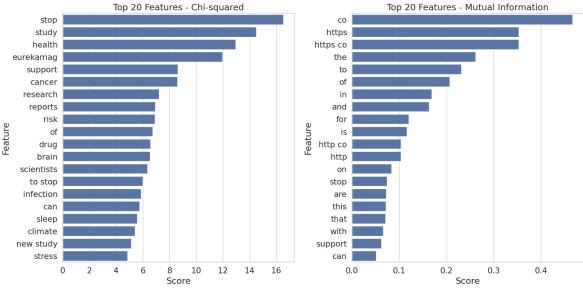
Out[83]:	Unname	d: 0	tweet_id	text	science_related	scientific_claim scie
	0	0	316669998137483264	Knees are a bit sore. i guess that's a sign th	0	0.0
	1	1	319090866545385472	McDonald's breakfast stop then the gym &	0	0.0
	2	2	322030931022065664	Can any Gynecologist with Cancer Experience ex	1	1.0
	3	3	322694830620807168	Couch-lock highs lead to sleeping in the couch	1	1.0
	4	4	328524426658328576	Does daily routine help prevent problems with	1	1.0
In [120	<pre># Remove warning messages import warnings from sklearn.exceptions import ConvergenceWarning warnings.filterwarnings("ignore", category=UserWarning, module="matplotli") warnings.filterwarnings("ignore", category=UserWarning, module="seaborn") warnings.filterwarnings("ignore", category=UserWarning, module="pandas") warnings.filterwarnings("ignore", category=UserWarning, module="wordcloud") warnings.filterwarnings("ignore", category=FutureWarning, module="pandas") warnings.filterwarnings("ignore", category=DeprecationWarning, module="pandas") warnings.filterwarnings("ignore", category=ConvergenceWarning, module="sk")</pre>					
In [85]:	<pre>from sklearn.feature_extraction.text import TfidfVectorizer vectorizer = TfidfVectorizer(stop_words='english', max_features=5000) X_all = vectorizer.fit_transform(df['text'])</pre>					
In [119	<pre># Compare different feature extraction methods from sklearn.feature_extraction.text import CountVectorizer from sklearn.feature_selection import SelectKBest, chi2, mutual_info_clas # Create label for Task 1 y_task1 = df['science_related']</pre>					

```
# 1. Count Vectorizer
count vec = CountVectorizer(max features=5000)
X_count = count_vec.fit_transform(df['text'])
# 2. TF-IDF with more parameters
tfidf vec = TfidfVectorizer(max features=5000,
                           min df=5,
                           max df=0.8,
                           ngram range=(1, 2)
X_tfidf = tfidf_vec.fit_transform(df['text'])
# 3. TF-IDF with preprocessing already done
tfidf processed = TfidfVectorizer(max features=5000)
X_tfidf_processed = tfidf_processed.fit_transform(df['text'])
# Compare feature extraction methods
print(f"Count Vectorizer Features: {X count.shape}")
print(f"TF-IDF Vectorizer Features: {X tfidf.shape}")
print(f"TF-IDF on Preprocessed Text Features: {X tfidf processed.shape}")
# Feature selection using Chi-squared
selector_chi2 = SelectKBest(chi2, k=100)
X_chi2 = selector_chi2.fit_transform(X_tfidf, y_task1)
# Feature selection using Mutual Information
selector mi = SelectKBest(mutual info classif, k=100)
X_mi = selector_mi.fit_transform(X_tfidf, y_task1)
print(f"\nFeatures after Chi-squared selection: {X chi2.shape}")
print(f"Features after Mutual Information selection: {X mi.shape}")
# Get and visualize the most important features
chi2_selected_indices = selector_chi2.get_support(indices=True)
mi_selected_indices = selector_mi.get_support(indices=True)
chi2_feature_names = np.array(tfidf_vec.get_feature_names_out())[chi2_sel
mi feature names = np.array(tfidf vec.get feature names out())[mi selecte
# Plot top 20 features by importance
plt.figure(figsize=(16, 8))
plt.subplot(1, 2, 1)
chi2_scores = selector_chi2.scores_[chi2_selected_indices]
chi2_features_df = pd.DataFrame({'Feature': chi2_feature_names, 'Score':
chi2_features_df = chi2_features_df.sort_values('Score', ascending=False)
sns.barplot(x='Score', y='Feature', data=chi2_features_df)
plt.title('Top 20 Features - Chi-squared')
plt.subplot(1, 2, 2)
mi_scores = selector_mi.scores_[mi_selected_indices]
mi_features_df = pd.DataFrame({'Feature': mi_feature_names, 'Score': mi_s
mi_features_df = mi_features_df.sort_values('Score', ascending=False).hea
sns.barplot(x='Score', y='Feature', data=mi_features_df)
plt.title('Top 20 Features - Mutual Information')
plt.tight layout()
plt.show()
# We'll use the TF-IDF on preprocessed text for subsequent modeling
X selected = X tfidf
```

```
Count Vectorizer Features: (1140, 5000)
TF-IDF Vectorizer Features: (1140, 693)
TF-IDF on Preprocessed Text Features: (1140, 5000)
```

Features after Chi-squared selection: (1140, 100)





```
In [87]:
         # Create labels for all tasks
         df['task1_label'] = df['science_related']
         df_sci = df[df['science_related'] == 1].copy()
         df_sci['task2_label'] = ((df_sci['scientific_claim'] == 1.0) | (df sci['s
         df sci['task3 label'] = df sci[['scientific claim', 'scientific reference
         df_sci['task3_label'] = df_sci['task3_label'].map({
             'scientific claim': 0,
             'scientific reference': 1,
              'scientific context': 2
         })
```

```
In [88]:
         from sklearn.model_selection import train_test_split
         from sklearn.naive bayes import GaussianNB, MultinomialNB, ComplementNB,
         from sklearn.metrics import classification report, confusion matrix, accu
         from sklearn.neighbors import KNeighborsClassifier
         from sklearn.svm import SVC
         from sklearn.linear model import LogisticRegression
         from sklearn.ensemble import RandomForestClassifier
         # Function to evaluate model performance
         def evaluate_model(model, X_train, X_test, y_train, y_test, model_name):
             model.fit(X_train, y_train)
             y_pred = model.predict(X_test)
             accuracy = accuracy_score(y_test, y_pred)
             report = classification_report(y_test, y_pred, output_dict=True)
             cm = confusion_matrix(y_test, y_pred)
             print(f"Model: {model_name}")
             print(f"Accuracy: {accuracy:.4f}")
             print(classification_report(y_test, y_pred))
             return {
                 'model name': model name,
```

```
'accuracy': accuracy,
        'report': report,
        'confusion_matrix': cm,
        'y pred': y pred
    }
# Get a dense version of our features for Gaussian NB
X dense = X selected.toarray()
# Split data for Task 1
X train task1, X test task1, y train task1, y test task1 = train test spl
    X dense, df['task1 label'], test size=0.2, random state=42
# Compare different NB variants for Task 1
nb_models = {
    'Gaussian NB': GaussianNB(),
    'Multinomial NB': MultinomialNB(),
    'Complement NB': ComplementNB(),
    'Bernoulli NB': BernoulliNB(),
           : KNeighborsClassifier(n_neighbors=5),
           : SVC(kernel='linear', C=1),
    'Logistic Regression': LogisticRegression(max_iter=1000),
    'Random Forest': RandomForestClassifier(n estimators=100),
}
task1_results = {}
print("Task 1: Science Related Classification\n" + "="*40)
for name, model in nb models.items():
    task1 results[name] = evaluate model(
        model, X_train_task1, X_test_task1, y_train_task1, y_test_task1,
    print("\n")
```

Task 1: Science Related Classification

Task 1: Science Related Classification					
Model: Gaussian NB Accuracy: 0.6711					
	precision	recall	f1-score	support	
0 1	0.79 0.53	0.66 0.70	0.72 0.60	146 82	
accuracy macro avg	0.66	0.68	0.67 0.66	228 228	
weighted avg	0.70	0.67	0.68	228	
Model: Multin					
•	precision	recall	f1-score	support	
0 1	0.74 0.84	0.96 0.39	0.83 0.53	146 82	
accuracy macro avg	0.79	0.67	0.75 0.68	228 228	
weighted avg	0.77	0.75	0.73	228	
Model: Comple Accuracy: 0.7					
	precision	recall	f1-score	support	
0 1	0.84 0.69	0.82 0.73	0.83 0.71	146 82	
accuracy macro avg	0.77	0.77	0.79 0.77	228 228	
weighted avg	0.79	0.79	0.79	228	
Model: Bernou Accuracy: 0.7					
,	precision	recall	f1-score	support	
0 1	0.80 0.77	0.90 0.59	0.85 0.67	146 82	
accuracy macro avg	0.78	0.74	0.79 0.76	228 228	
weighted avg	0.79	0.79	0.78	228	
Model: KNN Accuracy: 0.7105					
	precision	recall		support	
Ω	0.00	0.72	0.76	1/6	

0.73

0.68

0.76

0.63

146

82

0

1

0.80

0.58

accura	су			0.71	228
macro a	vg G	0.69	0.70	0.70	228
weighted a	vg 6	0.72	0.71	0.71	228

Model: SVM

Accuracy: 0.7895

	precision	recall	f1-score	support
0 1	0.79 0.79	0.92 0.56	0.85 0.66	146 82
accuracy macro avg weighted avg	0.79 0.79	0.74 0.79	0.79 0.75 0.78	228 228 228

Model: Logistic Regression

Accuracy: 0.7675

	precision	recall	f1-score	support
0 1	0.74 0.89	0.97 0.40	0.84 0.55	146 82
accuracy macro avg weighted avg	0.82 0.80	0.69 0.77	0.77 0.70 0.74	228 228 228

Model: Random Forest

Accuracy:	0./	precision	recall	f1-score	support
		precision	recate	11 30010	Suppor c
	0	0.74	0.94	0.83	146
	1	0.80	0.43	0.56	82
accura	асу			0.75	228
macro a	avg	0.77	0.68	0.69	228
weighted a	avg	0.76	0.75	0.73	228

```
In [89]: from sklearn.model_selection import StratifiedKFold
    from sklearn.metrics import accuracy_score, precision_score, recall_score

# Initialize k-fold cross validation
    k_folds = 10
    skf = StratifiedKFold(n_splits=k_folds, shuffle=True, random_state=42)

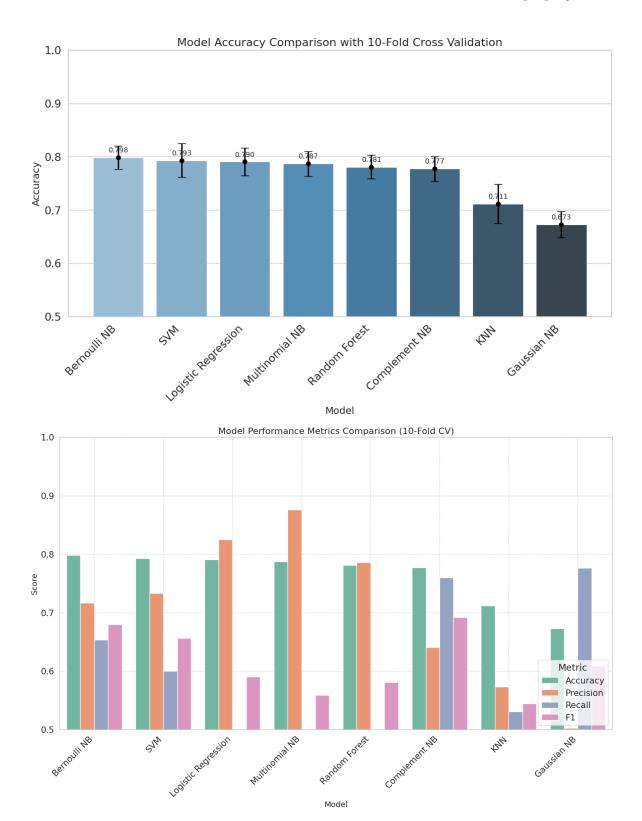
# Dictionary to store results
    cv_results = {}

# Perform k-fold cross validation for each model
    for model_name, model in nb_models.items():
        print(f"Performing {k_folds}-fold cross-validation for {model_name}...
```

```
# Initialize lists to store performance metrics for each fold
   fold_accuracy = []
    fold precision = []
    fold recall = []
    fold f1 = []
    # For each fold
    for fold, (train idx, test idx) in enumerate(skf.split(X dense, y tas
        # Split data
        X train fold, X test fold = X dense[train idx], X dense[test idx]
        y train fold, y test fold = y task1.iloc[train idx], y task1.iloc
        # Train model
        model.fit(X_train_fold, y_train_fold)
        # Make predictions
        y pred_fold = model.predict(X_test_fold)
        # Calculate metrics
        acc = accuracy_score(y_test_fold, y_pred_fold)
        prec = precision_score(y_test_fold, y_pred_fold, zero_division=0)
        rec = recall_score(y_test_fold, y_pred_fold, zero_division=0)
        f1 = f1_score(y_test_fold, y_pred_fold, zero_division=0)
        fold accuracy.append(acc)
        fold_precision.append(prec)
        fold_recall.append(rec)
        fold f1.append(f1)
    # Store average metrics and standard deviations
    cv_results[model_name] = {
        'accuracy': {
            'mean': np.mean(fold_accuracy),
            'std': np.std(fold accuracy)
        },
        'precision': {
            'mean': np.mean(fold_precision),
            'std': np.std(fold_precision)
        },
        'recall': {
            'mean': np.mean(fold_recall),
            'std': np.std(fold_recall)
        },
        'f1': {
            'mean': np.mean(fold_f1),
            'std': np.std(fold f1)
        }
    }
    print(f" Average: Accuracy={cv_results[model_name]['accuracy']['mean
    print(f" Average: F1 Score={cv_results[model_name]['f1']['mean']:.4f
   print()
# Create DataFrame for visualization
results df = pd.DataFrame({
    'Model': [],
    'Metric': [],
    'Mean': [],
    'Std': []
```

```
})
for model_name in cv_results:
    for metric in ['accuracy', 'precision', 'recall', 'f1']:
        results df = pd.concat([results df, pd.DataFrame({
            'Model': [model_name],
            'Metric': [metric.capitalize()],
            'Mean': [cv results[model name][metric]['mean']],
            'Std': [cv_results[model_name][metric]['std']]
        })], ignore_index=True)
# Sort models by accuracy
model order = results df[results df['Metric'] == 'Accuracy'].sort values(
# Create plots
plt.figure(figsize=(12, 8))
sns.set style("whitegrid")
# Create bar plot for accuracy
ax = sns.barplot(
    data=results_df[results_df['Metric'] == 'Accuracy'],
    x='Model',
   y='Mean',
    order=model order,
    palette='Blues_d'
)
# Add error bars
for i, model in enumerate(model order):
    row = results df[(results df['Model'] == model) & (results df['Metric
    ax.errorbar(
        i, row['Mean'], yerr=row['Std'],
        fmt='o', color='black', elinewidth=2, capsize=6
    )
# Add value labels on top of bars
for i, bar in enumerate(ax.patches):
    ax.text(
        bar.get_x() + bar.get_width()/2,
        bar.get_height() + 0.01,
        f"{bar.get height():.3f}",
        ha='center',
        fontsize=10
    )
plt.title(f'Model Accuracy Comparison with {k_folds}-Fold Cross Validation
plt.xlabel('Model', fontsize=14)
plt.ylabel('Accuracy', fontsize=14)
plt.xticks(rotation=45, ha='right')
plt.ylim([0.5, 1.0])
plt.tight_layout()
plt.show()
# Create a grouped bar chart for all metrics
plt.figure(figsize=(15, 10))
sns.set_style("whitegrid")
# Create grouped bar plot
ax = sns.barplot(
    data=results_df,
```

```
x='Model',
     y='Mean',
     hue='Metric',
     order=model order,
     palette='Set2'
 plt.title(f'Model Performance Metrics Comparison ({k folds}-Fold CV)', fo
 plt.xlabel('Model', fontsize=14)
 plt.ylabel('Score', fontsize=14)
 plt.xticks(rotation=45, ha='right')
 plt.legend(title='Metric', loc='lower right')
 plt.ylim([0.5, 1.0])
 plt.grid(True, linestyle='--', alpha=0.7)
 plt.tight layout()
 plt.show()
Performing 10-fold cross-validation for Gaussian NB...
 Average: Accuracy=0.6728 \pm 0.0245
 Average: F1 Score=0.6090 \pm 0.0290
Performing 10-fold cross-validation for Multinomial NB...
 Average: Accuracy=0.7868 \pm 0.0232
 Average: F1 Score=0.5587 \pm 0.0653
Performing 10-fold cross-validation for Complement NB...
  Average: Accuracy=0.7772 \pm 0.0229
 Average: F1 Score=0.6921 \pm 0.0204
Performing 10-fold cross-validation for Bernoulli NB...
  Average: Accuracy=0.7982 \pm 0.0215
 Average: F1 Score=0.6795 \pm 0.0355
Performing 10-fold cross-validation for KNN...
  Average: Accuracy=0.7114 \pm 0.0367
 Average: F1 Score=0.5443 \pm 0.0647
Performing 10-fold cross-validation for SVM...
 Average: Accuracy=0.7930 \pm 0.0319
 Average: F1 Score=0.6562 \pm 0.0459
Performing 10-fold cross-validation for Logistic Regression...
  Average: Accuracy=0.7904 \pm 0.0262
 Average: F1 Score=0.5900 \pm 0.0711
Performing 10-fold cross-validation for Random Forest...
 Average: Accuracy=0.7807 \pm 0.0222
 Average: F1 Score=0.5803 \pm 0.0515
/tmp/ipykernel 368067/4283035198.py:93: FutureWarning:
Passing `palette` without assigning `hue` is deprecated and will be remove
d in v0.14.0. Assign the `x` variable to `hue` and set `legend=False` for
the same effect.
 ax = sns.barplot(
```



Parameter optimization

```
In [90]: from sklearn.model_selection import GridSearchCV

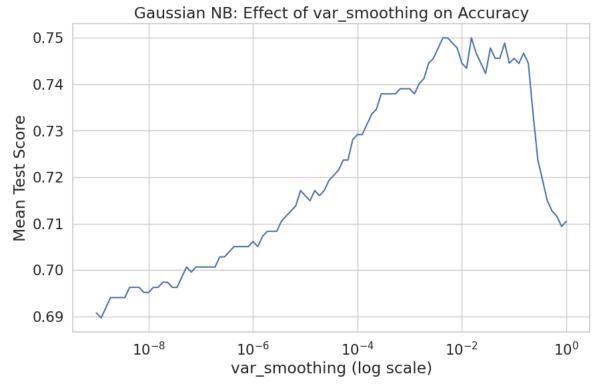
for model_name, model in nb_models.items():
```

```
print(f"Performing Grid Search for {model name}...")
# Define the parameter grid
if model_name == 'Gaussian NB':
    param grid = {
        'var smoothing': np.
        logspace(0, -9, num=100)
elif model_name == 'Multinomial NB':
    param_grid = {
        'alpha': np.logspace(-3, 3, num=100),
        'fit prior': [True, False]
    }
elif model name == 'Complement NB':
    param_grid = {
        'alpha': np.logspace(-3, 3, num=100),
        'fit_prior': [True, False]
elif model name == 'Bernoulli NB':
    param grid = {
        'alpha': np.logspace(-3, 3, num=100),
        'fit_prior': [True, False]
elif model name == 'KNN':
    param grid = {
        'n_neighbors': [3, 5, 7, 9],
        'weights': ['uniform', 'distance'],
        'metric': ['euclidean', 'manhattan']
elif model_name == 'SVM':
    param grid = {
        'C': np.logspace(-3, 3, num=100),
        'kernel': ['linear', 'rbf'],
        'gamma': ['scale', 'auto']
elif model name == 'Logistic Regression':
    param_grid = {
        'C': np.logspace(-3, 3, num=100),
        'penalty': ['l1', 'l2'],
        'solver': ['liblinear', 'saga']
elif model name == 'Random Forest':
    param grid = {
        'n_estimators': [50, 100, 200],
        'max_depth': [None, 10, 20, 30],
        'min_samples_split': [2, 5, 10],
        'min_samples_leaf': [1, 2, 4]
    }
else:
    continue
grid_search = GridSearchCV(
    estimator=model,
    param_grid=param_grid,
    scoring='accuracy',
    cv=StratifiedKFold(n_splits=k_folds, shuffle=True, random_state=4
    n_{jobs=-1}
    verbose=1
)
```

```
# Fit the grid search to the data
grid search.fit(X train task1, y train task1)
# Get the best parameters and best score
print(f"Best Parameters: {grid search.best params }")
print(f"Best Cross-Validation Score: {grid search.best score :.4f}")
# Evaluate the best model
best model = grid search.best estimator
best_model_accuracy = best_model.score(X_test_task1, y_test_task1)
print(f"Test Accuracy with Best Parameters: {best model accuracy:.4f}
# Visualize parameter impact
results = pd.DataFrame(grid_search.cv_results_)
# Plot effect of parameters
if model name == 'Gaussian NB':
    plt.figure(figsize=(10, 6))
    sns.lineplot(data=results, x='param_var_smoothing', y='mean_test_
    plt.xscale('log')
    plt.title(f'Gaussian NB: Effect of var_smoothing on Accuracy')
    plt.xlabel('var_smoothing (log scale)')
    plt.ylabel('Mean Test Score')
    plt.grid(True)
    plt.show()
elif model_name == 'Multinomial NB':
    plt.figure(figsize=(10, 6))
    sns.lineplot(data=results, x='param_alpha', y='mean_test_score',
    plt.xscale('log')
    plt.title(f'Multinomial NB: Effect of alpha and fit_prior on Accu
    plt.xlabel('alpha (log scale)')
    plt.ylabel('Mean Test Score')
    plt.grid(True)
    plt.show()
elif model_name == 'Complement NB':
    plt.figure(figsize=(10, 6))
    sns.lineplot(data=results, x='param_alpha', y='mean_test_score',
    plt.xscale('log')
    plt.title(f'Complement NB: Effect of alpha and fit_prior on Accur
    plt.xlabel('alpha (log scale)')
    plt.ylabel('Mean Test Score')
    plt.grid(True)
    plt.show()
elif model_name == 'Bernoulli NB':
    plt.figure(figsize=(10, 6))
    sns.lineplot(data=results, x='param_alpha', y='mean_test_score',
    plt.xscale('log')
    plt.title(f'Bernoulli NB: Effect of alpha and fit_prior on Accura
    plt.xlabel('alpha (log scale)')
    plt.ylabel('Mean Test Score')
    plt.grid(True)
    plt.show()
elif model_name == 'KNN':
    plt.figure(figsize=(10, 6))
    sns.lineplot(data=results, x='param_n_neighbors', y='mean_test_sc
    plt.title(f'KNN: Effect of n_neighbors and weights on Accuracy')
    plt.xlabel('n_neighbors')
    plt.ylabel('Mean Test Score')
    plt.grid(True)
```

```
plt.show()
elif model name == 'SVM':
    plt.figure(figsize=(10, 6))
    sns.lineplot(data=results, x='param C', y='mean test score', hue=
    plt.xscale('log')
    plt.title(f'SVM: Effect of C and kernel on Accuracy')
    plt.xlabel('C (log scale)')
    plt.ylabel('Mean Test Score')
    plt.grid(True)
    plt.show()
elif model_name == 'Logistic Regression':
    plt.figure(figsize=(10, 6))
    sns.lineplot(data=results, x='param C', y='mean test score', hue=
    plt.xscale('log')
    plt.title(f'Logistic Regression: Effect of C and penalty on Accur
    plt.xlabel('C (log scale)')
    plt.ylabel('Mean Test Score')
    plt.grid(True)
    plt.show()
elif model_name == 'Random Forest':
    plt.figure(figsize=(10, 6))
    sns.lineplot(data=results, x='param_n_estimators', y='mean_test_s
    plt.title(f'Random Forest: Effect of n estimators and max depth o
    plt.xlabel('n estimators')
    plt.ylabel('Mean Test Score')
    plt.grid(True)
    plt.show()
else:
    print(f"No visualization available for {model name} model.")
```

Performing Grid Search for Gaussian NB... Fitting 10 folds for each of 100 candidates, totalling 1000 fits Best Parameters: {'var_smoothing': np.float64(0.01519911082952933)} Best Cross-Validation Score: 0.7500 Test Accuracy with Best Parameters: 0.7325



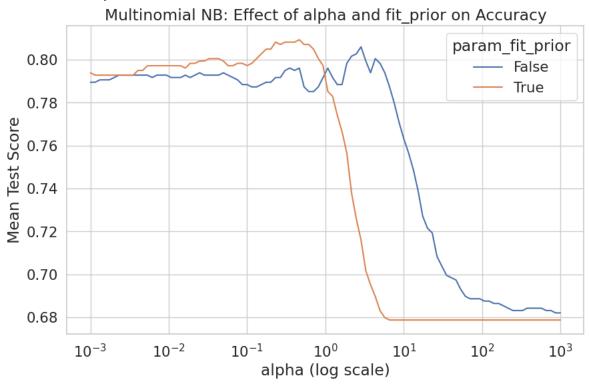
Performing Grid Search for Multinomial NB...

Fitting 10 folds for each of 200 candidates, totalling 2000 fits

Best Parameters: {'alpha': np.float64(0.4641588833612782), 'fit_prior': Tr

Best Cross-Validation Score: 0.8093

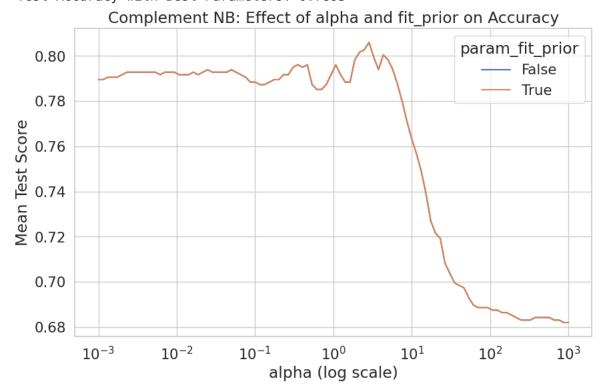
Test Accuracy with Best Parameters: 0.7982



Performing Grid Search for Complement NB... Fitting 10 folds for each of 200 candidates, totalling 2000 fits Best Parameters: {'alpha': np.float64(2.848035868435802), 'fit prior': Tru

Best Cross-Validation Score: 0.8060

Test Accuracy with Best Parameters: 0.7895



Performing Grid Search for Bernoulli NB...

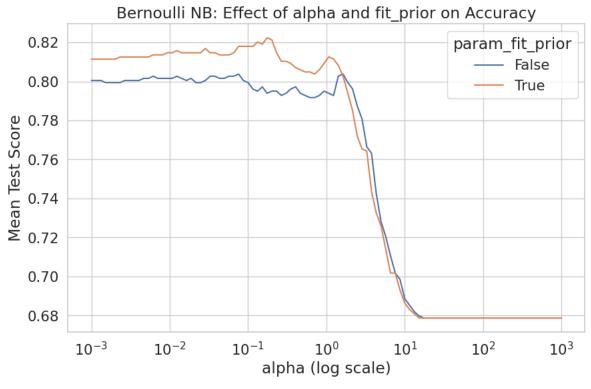
Fitting 10 folds for each of 200 candidates, totalling 2000 fits

Best Parameters: {'alpha': np.float64(0.1747528400007685), 'fit_prior': Tr

e}

Best Cross-Validation Score: 0.8224

Test Accuracy with Best Parameters: 0.8070



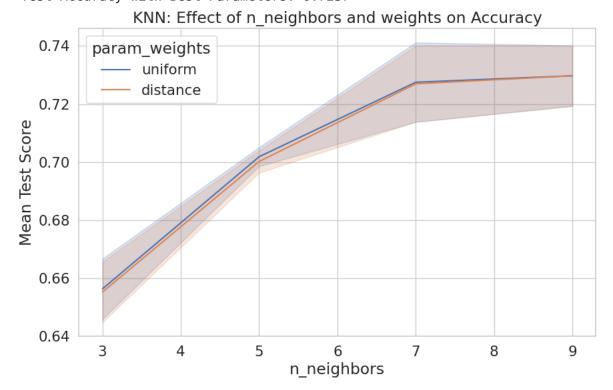
Performing Grid Search for KNN...

Fitting 10 folds for each of 16 candidates, totalling 160 fits

Best Parameters: {'metric': 'euclidean', 'n_neighbors': 7, 'weights': 'uni
form'}

Best Cross-Validation Score: 0.7412

Test Accuracy with Best Parameters: 0.7237



Performing Grid Search for SVM...

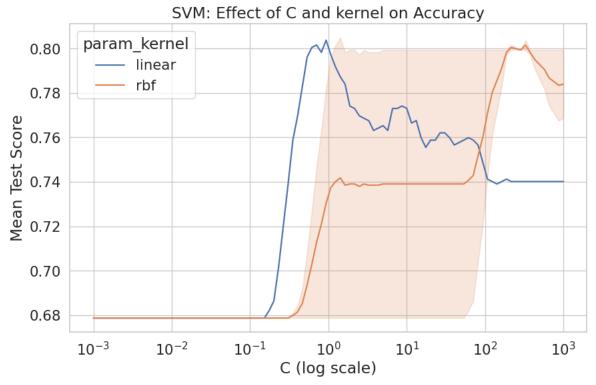
Fitting 10 folds for each of 400 candidates, totalling 4000 fits

Best Parameters: {'C': np.float64(1.4174741629268048), 'gamma': 'scale', '

kernel': 'rbf'}

Best Cross-Validation Score: 0.8049

Test Accuracy with Best Parameters: 0.7719



Performing Grid Search for Logistic Regression... Fitting 10 folds for each of 400 candidates, totalling 4000 fits

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/home/hurel/Documents/repo/projet-ml/.venv/lib/python3.10/site-packages/sk
learn/linear_model/_sag.py:348: ConvergenceWarning: The max_iter was reach
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```

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/home/hurel/Documents/repo/projet-ml/.venv/lib/python3.10/site-packages/sk learn/linear_model/_sag.py:348: ConvergenceWarning: The max_iter was reach ed which means the coef_ did not converge

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/home/hurel/Documents/repo/projet-ml/.venv/lib/python3.10/site-packages/sk learn/linear_model/_sag.py:348: ConvergenceWarning: The max_iter was reach ed which means the coef_ did not converge

warnings.warn(

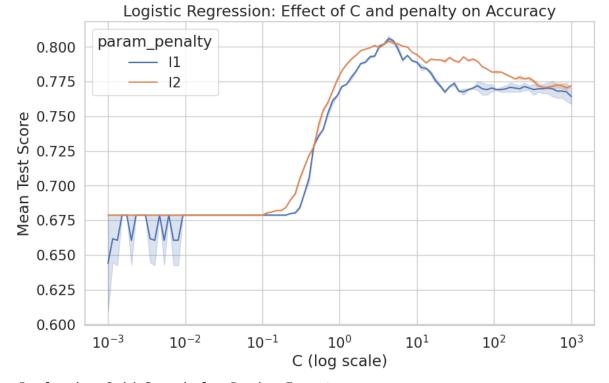
/home/hurel/Documents/repo/projet-ml/.venv/lib/python3.10/site-packages/sk learn/linear_model/_sag.py:348: ConvergenceWarning: The max_iter was reach ed which means the coef_ did not converge

warnings.warn(

Best Parameters: {'C': np.float64(4.328761281083062), 'penalty': 'l1', 'so lver': 'saga'}

Best Cross-Validation Score: 0.8081

Test Accuracy with Best Parameters: 0.7939



Performing Grid Search for Random Forest...

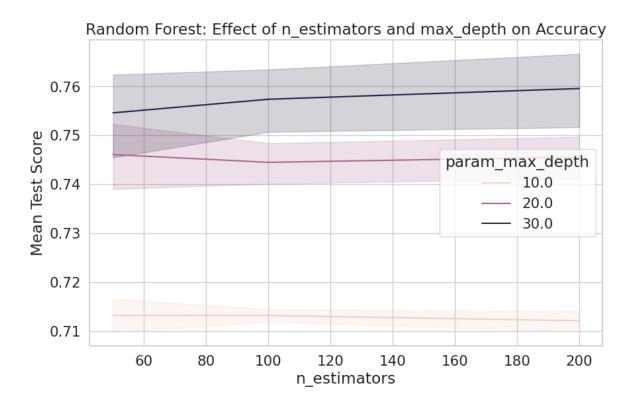
Fitting 10 folds for each of 108 candidates, totalling 1080 fits

Best Parameters: {'max_depth': None, 'min_samples_leaf': 1, 'min_samples_s

plit': 10, 'n_estimators': 50}

Best Cross-Validation Score: 0.7818

Test Accuracy with Best Parameters: 0.7675



Training comparaison with best parameter

```
from sklearn.model selection import cross val score
from sklearn.metrics import accuracy_score, precision_score, recall_score
# Create dictionary of models with their best parameters based on grid se
# Define models with their best parameters based on grid search results
best_models = {
     'Gaussian NB': GaussianNB(var smoothing=0.0152), #ample value - updat
    'Multinomial NB': MultinomialNB(alpha=0.4641, fit_prior=True), # Exa
    'Complement NB': ComplementNB(alpha=2.8480, fit_prior=True), # Examp
    'Bernoulli NB': BernoulliNB(alpha=0.17475, fit_prior=True), # Exampl
    'KNN': KNeighborsClassifier(n neighbors=7, weights='uniform', metric=
    'SVM': SVC(C=1.4174742, kernel='rbf', gamma='scale', probability=True
    'Logistic Regression': LogisticRegression(C=np.float64(4.328761281083
    'Random Forest': RandomForestClassifier(n_estimators=200, max_depth=N
}
# Dictionary to store results
best cv results = {}
# Perform k-fold cross validation for each model
for model_name, model in best_models.items():
    print(f"Performing {k_folds}-fold cross-validation for {model_name} w
    # Initialize lists to store performance metrics for each fold
    fold accuracy = []
    fold_precision = []
    fold_recall = []
    fold_f1 = []
    fold_auc = []
```

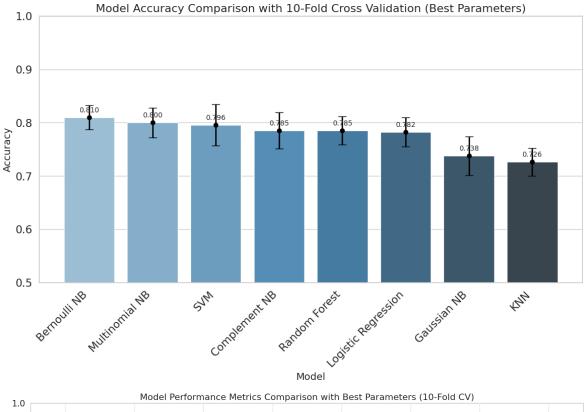
```
# For each fold
for fold, (train idx, test idx) in enumerate(skf.split(X dense, y tas
    # Split data
    X train fold, X test fold = X dense[train idx], X dense[test idx]
    y train fold, y test fold = y task1.iloc[train idx], y task1.iloc
    # Train model
    model.fit(X train fold, y train fold)
    # Make predictions
    y pred fold = model.predict(X test fold)
    # Calculate metrics
    acc = accuracy_score(y_test_fold, y_pred_fold)
    prec = precision_score(y_test_fold, y_pred_fold, zero_division=0)
    rec = recall_score(y_test_fold, y_pred_fold, zero_division=0)
    f1 = f1_score(y_test_fold, y_pred_fold, zero_division=0)
    # For AUC, we need probability estimates
    try:
        if hasattr(model, "predict_proba"):
            y_prob = model.predict_proba(X_test_fold)[:, 1]
            auc_score = roc_auc_score(y_test_fold, y_prob)
            fold auc.append(auc score)
    except:
        pass
    fold_accuracy.append(acc)
    fold precision.append(prec)
    fold recall.append(rec)
    fold_f1.append(f1)
# Store average metrics and standard deviations
best_cv_results[model_name] = {
    'accuracy': {
        'mean': np.mean(fold_accuracy),
        'std': np.std(fold accuracy)
    },
    'precision': {
        'mean': np.mean(fold_precision),
        'std': np.std(fold precision)
    },
    'recall': {
        'mean': np.mean(fold_recall),
        'std': np.std(fold_recall)
    },
    'f1': {
        'mean': np.mean(fold f1),
        'std': np.std(fold_f1)
    }
}
if fold auc:
    best_cv_results[model_name]['auc'] = {
        'mean': np.mean(fold_auc),
        'std': np.std(fold_auc)
    print(f" Average: AUC={best_cv_results[model_name]['auc']['mean'
print(f" Average: Accuracy={best_cv_results[model_name]['accuracy'][
```

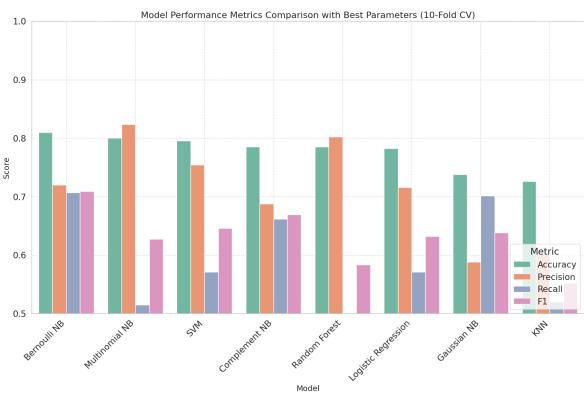
```
print(f" Average: F1 Score={best cv results[model name]['f1']['mean'
    print()
# Create DataFrame for visualization
best results df = pd.DataFrame({
    'Model': [],
    'Metric': [],
    'Mean': [],
    'Std': []
})
for model name in best cv results:
    for metric in ['accuracy', 'precision', 'recall', 'f1']:
        best_results_df = pd.concat([best_results_df, pd.DataFrame({
            'Model': [model name],
            'Metric': [metric.capitalize()],
            'Mean': [best cv results[model name][metric]['mean']],
            'Std': [best cv results[model name][metric]['std']]
        })], ignore index=True)
    if 'auc' in best_cv_results[model_name]:
        best_results_df = pd.concat([best_results_df, pd.DataFrame({
            'Model': [model_name],
            'Metric': ['AUC'],
            'Mean': [best cv results[model name]['auc']['mean']],
            'Std': [best_cv_results[model_name]['auc']['std']]
        })], ignore index=True)
# Sort models by accuracy
best model order = best results df[best results df['Metric'] == 'Accuracy'
# Create plots
plt.figure(figsize=(12, 8))
sns.set_style("whitegrid")
# Create bar plot for accuracy
ax = sns.barplot(
    data=best results df[best results df['Metric'] == 'Accuracy'],
    x='Model',
    y='Mean',
    order=best_model_order,
    palette='Blues d'
)
# Add error bars
for i, model in enumerate(best model order):
    row = best_results_df[(best_results_df['Model'] == model) & (best_res
    ax.errorbar(
        i, row['Mean'], yerr=row['Std'],
        fmt='o', color='black', elinewidth=2, capsize=6
    )
# Add value labels on top of bars
for i, bar in enumerate(ax.patches):
    ax.text(
        bar.get x() + bar.get width()/2,
        bar.get height() + 0.01,
        f"{bar.get_height():.3f}",
        ha='center',
        fontsize=10
    )
```

```
plt.title(f'Model Accuracy Comparison with {k folds}-Fold Cross Validatio
plt.xlabel('Model', fontsize=14)
plt.ylabel('Accuracy', fontsize=14)
plt.xticks(rotation=45, ha='right')
plt.ylim([0.5, 1.0])
plt.tight layout()
plt.show()
# Create a grouped bar chart for all metrics
plt.figure(figsize=(15, 10))
sns.set style("whitegrid")
# Create grouped bar plot
ax = sns.barplot(
   data=best_results_df[best_results_df['Metric'] != 'AUC'],
   x='Model',
   y='Mean',
   hue='Metric',
   order=best_model_order,
   palette='Set2'
plt.title(f'Model Performance Metrics Comparison with Best Parameters ({k
plt.xlabel('Model', fontsize=14)
plt.ylabel('Score', fontsize=14)
plt.xticks(rotation=45, ha='right')
plt.legend(title='Metric', loc='lower right')
plt.ylim([0.5, 1.0])
plt.grid(True, linestyle='--', alpha=0.7)
plt.tight layout()
plt.show()
# Create a comparison table between original results and optimized result
comparison df = pd.DataFrame({
    'Model': [],
    'Metric': [],
    'Original Mean': [],
    'Original Std': [],
    'Optimized Mean': [],
    'Optimized Std': [],
    'Improvement': []
})
for model_name in best_cv_results:
    if model_name in cv_results:
        for metric in ['accuracy', 'precision', 'recall', 'f1']:
            orig mean = cv results[model name][metric]['mean']
            orig_std = cv_results[model_name][metric]['std']
            opt_mean = best_cv_results[model_name][metric]['mean']
            opt_std = best_cv_results[model_name][metric]['std']
            improvement = ((opt_mean - orig_mean) / orig_mean) * 100
            comparison df = pd.concat([comparison df, pd.DataFrame({
                'Model': [model name],
                'Metric': [metric.capitalize()],
                'Original Mean': [orig_mean],
                'Original Std': [orig_std],
                'Optimized Mean': [opt_mean],
                'Optimized Std': [opt_std],
```

```
'Improvement': [improvement]
            })], ignore index=True)
# Plot the improvement in accuracy
plt.figure(figsize=(15, 10))
sns.set style("whitegrid")
# Filter for accuracy only
acc comparison = comparison df[comparison df['Metric'] == 'Accuracy']
acc comparison = acc comparison.sort values('Improvement', ascending=Fals
# Create bar chart of improvements
ax = sns.barplot(
    data=acc_comparison,
    x='Model',
   y='Improvement',
    palette='RdYlGn',
    dodge=False
plt.axhline(y=0, color='black', linestyle='-', alpha=0.3)
# Add value labels on bars
for i, bar in enumerate(ax.patches):
    if bar.get_height() >= 0:
        ax.text(
            bar.get_x() + bar.get_width()/2,
            bar.get_height() + 0.5,
            f"{bar.get height():.2f}%",
            ha='center',
            fontsize=10
    else:
        ax.text(
            bar.get x() + bar.get width()/2,
            bar.get_height() - 1.0,
            f"{bar.get height():.2f}%",
            ha='center',
            fontsize=10
        )
plt.title('Percentage Improvement in Accuracy After Parameter Optimization
plt.xlabel('Model', fontsize=14)
plt.ylabel('Improvement (%)', fontsize=14)
plt.xticks(rotation=45, ha='right')
plt.grid(True, linestyle='--', alpha=0.7)
plt.tight_layout()
plt.show()
```

```
Performing 10-fold cross-validation for Gaussian NB with best parameter
  Average: AUC=0.7703 \pm 0.0396
  Average: Accuracy=0.7377 \pm 0.0363
  Average: F1 Score=0.6381 \pm 0.0403
Performing 10-fold cross-validation for Multinomial NB with best parameter
  Average: AUC=0.8660 \pm 0.0281
  Average: Accuracy=0.8000 \pm 0.0277
 Average: F1 Score=0.6273 \pm 0.0495
Performing 10-fold cross-validation for Complement NB with best parameter
S...
  Average: AUC=0.8549 \pm 0.0274
  Average: Accuracy=0.7851 \pm 0.0343
  Average: F1 Score=0.6695 \pm 0.0449
Performing 10-fold cross-validation for Bernoulli NB with best parameter
  Average: AUC=0.8681 \pm 0.0237
  Average: Accuracy=0.8096 \pm 0.0229
  Average: F1 Score=0.7092 \pm 0.0316
Performing 10-fold cross-validation for KNN with best parameters...
  Average: AUC=0.7673 \pm 0.0471
  Average: Accuracy=0.7263 \pm 0.0260
  Average: F1 Score=0.5524 \pm 0.0539
Performing 10-fold cross-validation for SVM with best parameters...
  Average: AUC=0.8655 \pm 0.0213
  Average: Accuracy=0.7956 \pm 0.0390
  Average: F1 Score=0.6462 \pm 0.0696
Performing 10-fold cross-validation for Logistic Regression with best para
meters...
  Average: AUC=0.8541 \pm 0.0249
  Average: Accuracy=0.7825 \pm 0.0277
  Average: F1 Score=0.6320 \pm 0.0482
Performing 10-fold cross-validation for Random Forest with best parameter
S...
  Average: AUC=0.8466 \pm 0.0334
  Average: Accuracy=0.7851 \pm 0.0264
  Average: F1 Score=0.5831 \pm 0.0627
/tmp/ipykernel 368067/2330540877.py:127: FutureWarning:
Passing `palette` without assigning `hue` is deprecated and will be remove
d in v0.14.0. Assign the `x` variable to `hue` and set `legend=False` for
the same effect.
  ax = sns.barplot(
```

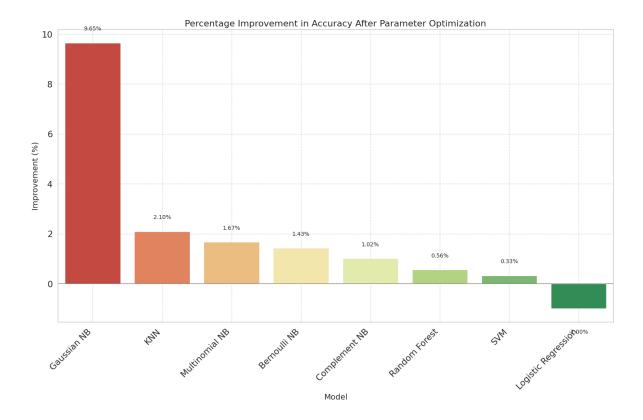




/tmp/ipykernel_368067/2330540877.py:224: FutureWarning:

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

ax = sns.barplot(

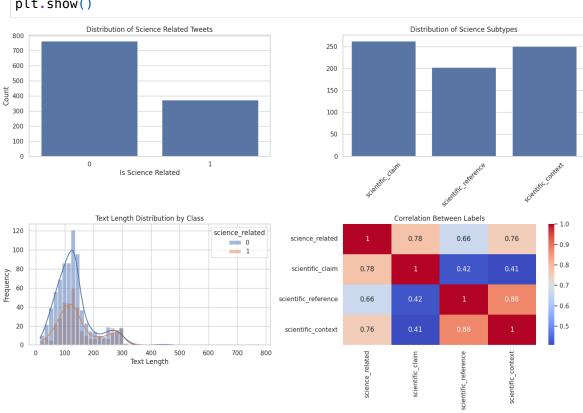


Preprocessing

```
In [92]: import numpy as np
         # Set style
         sns.set(style="whitegrid")
         plt.figure(figsize=(15, 10))
         # Plot distribution of science_related tweets
         plt.subplot(2, 2, 1)
         sns.countplot(x='science_related', data=df)
         plt.title('Distribution of Science Related Tweets')
         plt.xlabel('Is Science Related')
         plt.ylabel('Count')
         # Plot distribution of science subtypes for science-related tweets
         sci df = df[df['science related'] == 1]
         plt.subplot(2, 2, 2)
         subtypes = ['scientific_claim', 'scientific_reference', 'scientific_conte
         sns.barplot(x=subtypes, y=[sci_df[col].sum() for col in subtypes])
         plt.title('Distribution of Science Subtypes')
         plt.xticks(rotation=45)
         plt.tight layout()
         # Plot text length distribution
         plt.subplot(2, 2, 3)
         df['text length'] = df['text'].apply(len)
         sns.histplot(data=df, x='text_length', hue='science_related', bins=50, kd
         plt.title('Text Length Distribution by Class')
         plt.xlabel('Text Length')
         plt.ylabel('Frequency')
         # Plot correlation between features
         plt.subplot(2, 2, 4)
```

```
corr_cols = ['science_related', 'scientific_claim', 'scientific_reference'
sns.heatmap(df[corr_cols].corr(), annot=True, cmap='coolwarm')
plt.title('Correlation Between Labels')

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



```
In [93]: import re
         import nltk
         from nltk.corpus import stopwords
         from nltk.stem import PorterStemmer, WordNetLemmatizer
         import emoji
         # Download required NLTK resources
         nltk.download('stopwords')
         nltk.download('wordnet')
         nltk.download('punkt')
         nltk.download('punkt_tab')
         def preprocess_text(text):
             # Convert to lowercase
             text = text.lower()
             # Demojize text
             text = emoji.demojize(text)
             # Remove URLs
             text = re.sub(r'(http\S+|www\S+)', '', text)
             # Remove mentions and hashtags execept for the word uurekamag
             text = re.sub(r'@\w+', '', text)
             text = re.sub(r'#eurekamag', 'eurekamag', text)
             text = re.sub(r'#\w+', '', text)
```

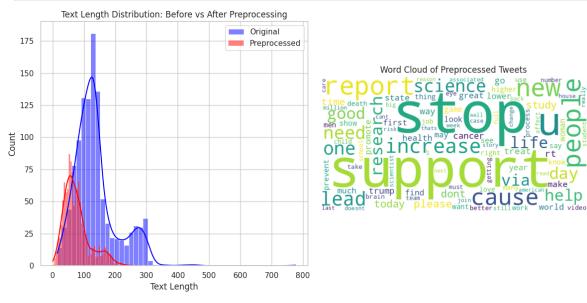
```
# Remove special characters and numbers
              text = re.sub(r'[^a-zA-Z\s]', '', text)
              # Tokenize
              tokens = nltk.word tokenize(text)
              # Remove stopwords
              stop words = set(stopwords.words('english'))
              tokens = [word for word in tokens if word not in stop words]
              # Lemmatize
              lemmatizer = WordNetLemmatizer()
              tokens = [lemmatizer.lemmatize(word) for word in tokens]
              # Rejoin
              return ' '.join(tokens)
          # Apply preprocessing to the dataset
          df['processed_text'] = df['text'].apply(preprocess_text)
          # Compare before and after
          comparison_df = pd.DataFrame({
              'Original': df['text'].head(5),
              'Preprocessed': df['processed_text'].head(5)
          })
          comparison_df
         [nltk_data] Downloading package stopwords to /home/hurel/nltk_data...
         [nltk data] Package stopwords is already up-to-date!
         [nltk data] Downloading package wordnet to /home/hurel/nltk data...
         [nltk data] Package wordnet is already up-to-date!
         [nltk_data] Downloading package punkt to /home/hurel/nltk_data...
         [nltk data]
                       Package punkt is already up-to-date!
         [nltk_data] Downloading package punkt_tab to /home/hurel/nltk_data...
         [nltk data] Package punkt tab is already up-to-date!
Out[93]:
                                        Original
                                                                         Preprocessed
               Knees are a bit sore. i guess that's a sign
                                                       knee bit sore guess thats sign recent
          0
                                                                             treadmil...
              McDonald's breakfast stop then the gym
                                                            mcdonalds breakfast stop gym
          1
                                                                     basketballflexedb...
                    Can any Gynecologist with Cancer
                                                     gynecologist cancer experience explain
          2
                                  Experience ex...
                                                                              danger ...
               Couch-lock highs lead to sleeping in the
                                                   couchlock high lead sleeping couch got ta
          3
                                                   daily routine help prevent problem bipolar
             Does daily routine help prevent problems
                                          with ...
                                                                                  dis...
In [94]: # Visualize the impact of preprocessing
          plt.figure(figsize=(12, 6))
          # Text length before and after preprocessing
          df['original_length'] = df['text'].apply(len)
          df['processed length'] = df['processed text'].apply(len)
          plt.subplot(1, 2, 1)
```

```
sns.histplot(data=df, x='original_length', color='blue', bins=50, kde=Tru
sns.histplot(data=df, x='processed_length', color='red', bins=50, kde=Tru
plt.title('Text Length Distribution: Before vs After Preprocessing')
plt.xlabel('Text Length')
plt.legend()

# Word cloud for processed text
from wordcloud import WordCloud

plt.subplot(1, 2, 2)
wordcloud = WordCloud(width=800, height=400, background_color='white', ma
plt.imshow(wordcloud, interpolation='bilinear')
plt.axis('off')
plt.title('Word Cloud of Preprocessed Tweets')

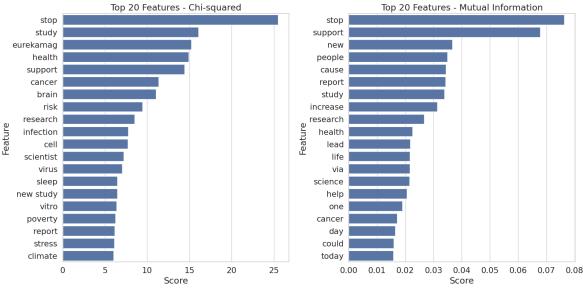
plt.tight_layout()
plt.show()
```



```
In [95]: from sklearn.feature_extraction.text import TfidfVectorizer

vectorizer = TfidfVectorizer(stop_words='english', max_features=5000)
X_all = vectorizer.fit_transform(df['text'])
```

```
tfidf processed = TfidfVectorizer(max features=5000)
 X tfidf processed = tfidf processed.fit transform(df['processed text'])
 # Compare feature extraction methods
 print(f"Count Vectorizer Features: {X count.shape}")
 print(f"TF-IDF Vectorizer Features: {X tfidf.shape}")
 print(f"TF-IDF on Preprocessed Text Features: {X tfidf processed.shape}")
 # Feature selection using Chi-squared
 selector chi2 = SelectKBest(chi2, k=100)
 X chi2 = selector chi2.fit transform(X tfidf, y task1)
 # Feature selection using Mutual Information
 selector_mi = SelectKBest(mutual_info_classif, k=100)
 X mi = selector mi.fit transform(X tfidf, y task1)
 print(f"\nFeatures after Chi-squared selection: {X chi2.shape}")
 print(f"Features after Mutual Information selection: {X mi.shape}")
 # Get and visualize the most important features
 chi2_selected_indices = selector_chi2.get_support(indices=True)
 mi_selected_indices = selector_mi.get_support(indices=True)
 chi2 feature names = np.array(tfidf vec.get feature names out())[chi2 sel
 mi feature names = np.array(tfidf vec.get feature names out())[mi selecte
 # Plot top 20 features by importance
 plt.figure(figsize=(16, 8))
 plt.subplot(1, 2, 1)
 chi2_scores = selector_chi2.scores_[chi2_selected_indices]
 chi2_features_df = pd.DataFrame({'Feature': chi2_feature_names, 'Score':
 chi2_features_df = chi2_features_df.sort_values('Score', ascending=False)
 sns.barplot(x='Score', y='Feature', data=chi2_features_df)
 plt.title('Top 20 Features - Chi-squared')
 plt.subplot(1, 2, 2)
 mi_scores = selector_mi.scores_[mi_selected_indices]
 mi_features_df = pd.DataFrame({'Feature': mi_feature_names, 'Score': mi_s
 mi_features_df = mi_features_df.sort_values('Score', ascending=False).hea
 sns.barplot(x='Score', y='Feature', data=mi features df)
 plt.title('Top 20 Features - Mutual Information')
 plt.tight layout()
 plt.show()
 # We'll use the TF-IDF on preprocessed text for subsequent modeling
 X selected = X tfidf
Count Vectorizer Features: (1140, 5000)
TF-IDF Vectorizer Features: (1140, 465)
TF-IDF on Preprocessed Text Features: (1140, 5000)
Features after Chi-squared selection: (1140, 100)
Features after Mutual Information selection: (1140, 100)
```



```
In [97]: | from sklearn.model_selection import train_test_split
         from sklearn.naive bayes import GaussianNB, MultinomialNB, ComplementNB,
         from sklearn.metrics import classification_report, confusion_matrix, accu
         # Function to evaluate model performance
         def evaluate_model(model, X_train, X_test, y_train, y_test, model_name):
             model.fit(X train, y train)
             y_pred = model.predict(X_test)
             accuracy = accuracy_score(y_test, y_pred)
              report = classification_report(y_test, y_pred, output_dict=True)
             cm = confusion_matrix(y_test, y_pred)
             print(f"Model: {model name}")
             print(f"Accuracy: {accuracy:.4f}")
             print(classification_report(y_test, y_pred))
              return {
                  'model_name': model_name,
                  'accuracy': accuracy,
                  'report': report,
                  'confusion matrix': cm,
                  'y pred': y pred
             }
         # Get a dense version of our features for Gaussian NB
         X_dense = X_selected.toarray()
         # Split data for Task 1
         X_train_task1, X_test_task1, y_train_task1, y_test_task1 = train_test_spl
             X_dense, df['task1_label'], test_size=0.2, random_state=42
         # Define models with their best parameters based on grid search results
         nb models = {
              'Gaussian NB': GaussianNB(var smoothing=0.0152), #ample value - updat
              'Multinomial NB': MultinomialNB(alpha=0.4641, fit prior=True), # Exa
              'Complement NB': ComplementNB(alpha=2.8480, fit prior=True), # Examp
              '<mark>Bernoulli NB</mark>': BernoulliNB(alpha=0.17475, fit_prior=True),                  # Exampl
              'KNN': KNeighborsClassifier(n_neighbors=7, weights='uniform', metric=
              'SVM': SVC(C=1.4174742, kernel='rbf', gamma='scale', probability=True
              'Logistic Regression': LogisticRegression(C=np.float64(4.328761281083
```

```
'Random Forest': RandomForestClassifier(n_estimators=200, max_depth=N)

task1_results = {}
print("Task 1: Science Related Classification\n" + "="*40)
for name, model in nb_models.items():
    task1_results[name] = evaluate_model(
        model, X_train_task1, X_test_task1, y_train_task1, y_test_task1,
    )
    print("\n")
```

Task 1: Science Related Classification

Task 1: Science Related Classification						
Model: Gaussian NB Accuracy: 0.6798						
	precision	recall	f1-score	support		
0 1	0.80 0.54	0.66 0.71	0.73 0.61	146 82		
accuracy			0.68	228		
macro avg weighted avg	0.67 0.71	0.69 0.68	0.67 0.69	228 228		
Model: Multir Accuracy: 0.7						
	precision	recall	f1-score	support		
0 1	0.75 0.71	0.89 0.48	0.82 0.57	146 82		
accuracy			0.74	228		
macro avg	0.73	0.68	0.69	228		
weighted avg	0.74	0.74	0.73	228		
Model: Comple						
·	precision	recall	f1-score	support		
0	0.82 0.64	0.78 0.70	0.80 0.67	146 82		
accuracy macro avg	0.73	0.74	0.75 0.73	228 228		
weighted avg	0.76	0.75	0.75	228		
Model: Bernou	ılli NR					
Accuracy: 0.7	7500	11	6.1			
	precision	recall	f1-score	support		
0 1	0.78 0.69	0.86 0.56	0.81 0.62	146 82		
accuracy			0.75	228		
macro avg weighted avg	0.73 0.74	0.71 0.75	0.72 0.74	228 228		
weighted dig	017.	0.75	0171	220		
Model: KNN						
Accuracy: 0.4	1737 precision	recall	f1-score	support		

0.20

0.96

0.33

0.57

146

82

0

1

0.91

0.40

accuracy			0.47	228
macro avg	0.65	0.58	0.45	228
weighted avg	0.73	0.47	0.41	228

Model: SVM

Accuracy: 0.7763

support	f1-score	recall	precision	
146 82	0.84 0.64	0.90 0.56	0.78 0.75	0 1
228 228 228	0.78 0.74 0.77	0.73 0.78	0.77 0.77	accuracy macro avg weighted avg

Model: Logistic Regression

Accuracy: 0.7675

	precision	recall	f1-score	support
6	0.79	0.88	0.83	146
1	0.72	0.57	0.64	82
accuracy	,		0.77	228
macro avo	0.75	0.72	0.73	228
weighted avg	0.76	0.77	0.76	228

Model: Random Forest Accuracy: 0.7588

precision recall f1-score support 0.83 0.79 0.81 146 1 0.65 0.71 0.68 82 0.76 228 accuracy macro avg 0.74 0.75 weighted avg 0.76 0.76 0.74 228 0.76 0.76 228

```
X chi2 processed = selector chi2.fit transform(X tfidf processed, y task1
# Get a dense version of our features for models that require it
X processed dense = X tfidf processed.toarray()
# Initialize k-fold cross validation
k folds = 10
skf = StratifiedKFold(n splits=k folds, shuffle=True, random state=42)
# Define models with their best parameters based on grid search results
best models = {
    'Gaussian NB': GaussianNB(var smoothing=0.0152),
    'Multinomial NB': MultinomialNB(alpha=0.4641, fit prior=True),
    'Complement NB': ComplementNB(alpha=2.8480, fit_prior=True),
    'Bernoulli NB': BernoulliNB(alpha=0.17475, fit prior=True),
    'KNN': KNeighborsClassifier(n_neighbors=7, weights='uniform', metric=
    'SVM': SVC(C=1.4174742, kernel='rbf', gamma='scale', probability=True
    'Logistic Regression': LogisticRegression(C=np.float64(4.328761281083
    'Random Forest': RandomForestClassifier(n estimators=200, max depth=N
# Dictionary to store results
best cv results = {}
# Perform k-fold cross validation for each model
for model name, model in best models.items():
    print(f"Performing {k_folds}-fold cross-validation for {model_name} w
    # Initialize lists to store performance metrics for each fold
    fold accuracy = []
    fold_precision = []
    fold_recall = []
    fold_f1 = []
    fold_auc = []
    # For each fold
    for fold, (train idx, test idx) in enumerate(skf.split(X processed de
        # Split data
        X_train_fold, X_test_fold = X_processed_dense[train_idx], X_proce
        y_train_fold, y_test_fold = y_task1.iloc[train_idx], y_task1.iloc
        # Train model
        model.fit(X_train_fold, y_train_fold)
        # Make predictions
        y_pred_fold = model.predict(X_test_fold)
        # Calculate metrics
        acc = accuracy_score(y_test_fold, y_pred_fold)
        prec = precision_score(y_test_fold, y_pred_fold, zero_division=0)
        rec = recall_score(y_test_fold, y_pred_fold, zero_division=0)
        f1 = f1_score(y_test_fold, y_pred_fold, zero_division=0)
        # For AUC, we need probability estimates
        try:
            if hasattr(model, "predict proba"):
                y_prob = model.predict_proba(X_test_fold)[:, 1]
                auc_score = roc_auc_score(y_test_fold, y_prob)
                fold_auc.append(auc_score)
        except:
```

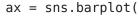
```
pass
        fold accuracy.append(acc)
        fold precision.append(prec)
        fold recall.append(rec)
        fold f1.append(f1)
    # Store average metrics and standard deviations
   best cv results[model name] = {
        'accuracy': {
            'mean': np.mean(fold accuracy),
            'std': np.std(fold accuracy)
        },
        'precision': {
            'mean': np.mean(fold precision),
            'std': np.std(fold_precision)
        },
        'recall': {
            'mean': np.mean(fold recall),
            'std': np.std(fold recall)
        },
        'f1': {
            'mean': np.mean(fold_f1),
            'std': np.std(fold f1)
        }
    }
    if fold auc:
        best cv results[model name]['auc'] = {
            'mean': np.mean(fold auc),
            'std': np.std(fold_auc)
        print(f" Average: AUC={best_cv_results[model_name]['auc']['mean'
    print(f"
              Average: Accuracy={best cv results[model name]['accuracy'][
              Average: F1 Score={best_cv_results[model_name]['f1']['mean'
    print(f"
    print()
# Create DataFrame for visualization
best_results_df = pd.DataFrame({
    'Model': [],
    'Metric': [],
    'Mean': [],
    'Std': []
})
for model_name in best_cv_results:
    for metric in ['accuracy', 'precision', 'recall', 'f1']:
        best_results_df = pd.concat([best_results_df, pd.DataFrame({
            'Model': [model name],
            'Metric': [metric.capitalize()],
            'Mean': [best_cv_results[model_name][metric]['mean']],
            'Std': [best cv results[model name][metric]['std']]
        })], ignore_index=True)
    if 'auc' in best_cv_results[model_name]:
        best_results_df = pd.concat([best_results_df, pd.DataFrame({
            'Model': [model_name],
            'Metric': ['AUC'],
            'Mean': [best_cv_results[model_name]['auc']['mean']],
            'Std': [best_cv_results[model_name]['auc']['std']]
```

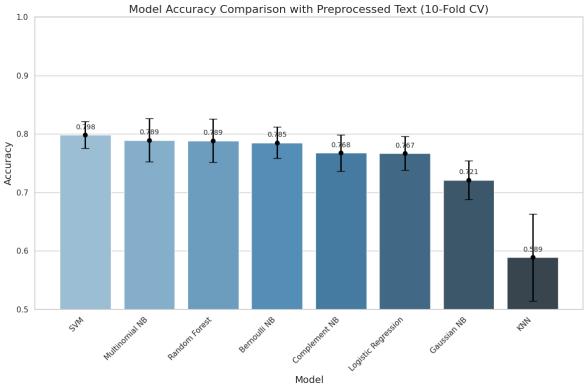
```
})], ignore_index=True)
# Sort models by accuracy
best model order = best results df[best results df['Metric'] == 'Accuracy'
# Create plots
plt.figure(figsize=(12, 8))
sns.set style("whitegrid")
# Create bar plot for accuracy
ax = sns.barplot(
   data=best results df[best results df['Metric'] == 'Accuracy'],
   x='Model',
   y='Mean',
   order=best model order,
   palette='Blues_d'
# Add error bars
for i, model in enumerate(best_model_order):
    row = best_results_df[(best_results_df['Model'] == model) & (best res
   ax.errorbar(
        i, row['Mean'], yerr=row['Std'],
        fmt='o', color='black', elinewidth=2, capsize=6
    )
# Add value labels on top of bars
for i, bar in enumerate(ax.patches):
    ax.text(
        bar.get x() + bar.get width()/2,
        bar.get_height() + 0.01,
        f"{bar.get_height():.3f}",
        ha='center',
        fontsize=10
    )
plt.title(f'Model Accuracy Comparison with Preprocessed Text ({k folds}-F
plt.xlabel('Model', fontsize=14)
plt.ylabel('Accuracy', fontsize=14)
plt.xticks(rotation=45, ha='right')
plt.ylim([0.5, 1.0])
plt.tight layout()
plt.show()
# Create a grouped bar chart for all metrics
plt.figure(figsize=(15, 10))
sns.set_style("whitegrid")
# Create grouped bar plot
ax = sns.barplot(
   data=best_results_df[best_results_df['Metric'] != 'AUC'],
   x='Model',
   y='Mean',
   hue='Metric',
   order=best_model_order,
   palette='Set2'
plt.title(f'Model Performance with Preprocessed Text ({k_folds}-Fold CV)'
plt.xlabel('Model', fontsize=14)
```

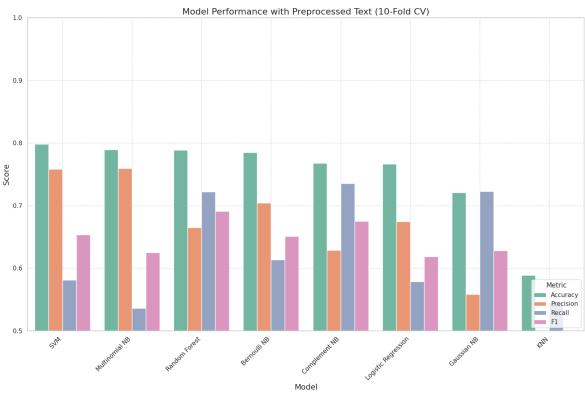
```
plt.ylabel('Score', fontsize=14)
 plt.xticks(rotation=45, ha='right')
 plt.legend(title='Metric', loc='lower right')
 plt.ylim([0.5, 1.0])
 plt.grid(True, linestyle='--', alpha=0.7)
 plt.tight layout()
 plt.show()
Performing 10-fold cross-validation for Gaussian NB with best parameters o
n preprocessed text...
 Average: AUC=0.7498 \pm 0.0386
 Average: Accuracy=0.7211 \pm 0.0332
 Average: F1 Score=0.6283 \pm 0.0560
Performing 10-fold cross-validation for Multinomial NB with best parameter
s on preprocessed text...
  Average: AUC=0.8442 \pm 0.0334
 Average: Accuracy=0.7895 \pm 0.0370
 Average: F1 Score=0.6251 \pm 0.0685
Performing 10-fold cross-validation for Complement NB with best parameters
on preprocessed text...
 Average: AUC=0.8550 \pm 0.0328
 Average: Accuracy=0.7675 \pm 0.0310
 Average: F1 Score=0.6750 \pm 0.0425
Performing 10-fold cross-validation for Bernoulli NB with best parameters
on preprocessed text...
 Average: AUC=0.8421 \pm 0.0246
 Average: Accuracy=0.7851 \pm 0.0270
 Average: F1 Score=0.6510 \pm 0.0474
Performing 10-fold cross-validation for KNN with best parameters on prepro
cessed text...
 Average: AUC=0.6058 \pm 0.0533
 Average: Accuracy=0.5886 \pm 0.0745
 Average: F1 Score=0.4461 \pm 0.1151
Performing 10-fold cross-validation for SVM with best parameters on prepro
cessed text...
 Average: AUC=0.8463 \pm 0.0323
 Average: Accuracy=0.7982 \pm 0.0229
 Average: F1 Score=0.6534 \pm 0.0402
Performing 10-fold cross-validation for Logistic Regression with best para
meters on preprocessed text...
 Average: AUC=0.8347 \pm 0.0318
 Average: Accuracy=0.7667 \pm 0.0289
 Average: F1 Score=0.6182 \pm 0.0508
Performing 10-fold cross-validation for Random Forest with best parameters
on preprocessed text...
 Average: AUC=0.8573 \pm 0.0307
 Average: Accuracy=0.7886 \pm 0.0373
 Average: F1 Score=0.6906 \pm 0.0609
```

/tmp/ipykernel_368067/2418365623.py:144: FutureWarning:

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







```
import seaborn as sns
import pandas as pd
import numpy as np
from sklearn.metrics import confusion_matrix, roc_curve, auc
```

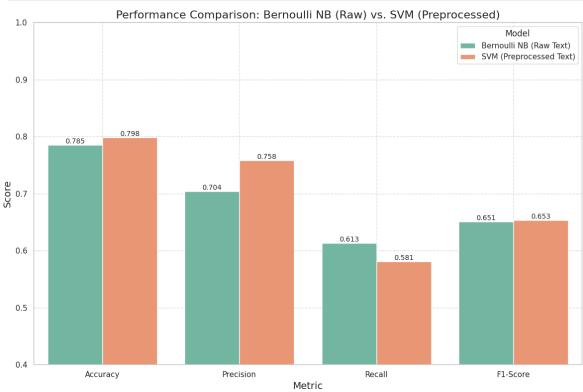
```
# Compare performance of SVM with preprocessed data and Bernoulli NB with
import matplotlib.pyplot as plt
# Create a DataFrame for comparison
comparison data = [
    {
        'Model': 'Bernoulli NB (Raw Text)',
        'Accuracy': best cv results['Bernoulli NB']['accuracy']['mean'],
        'Precision': best_cv_results['Bernoulli NB']['precision']['mean']
        'Recall': best_cv_results['Bernoulli NB']['recall']['mean'],
        'F1-Score': best cv results['Bernoulli NB']['f1']['mean']
    },
        'Model': 'SVM (Preprocessed Text)',
        'Accuracy': best_cv_results['SVM']['accuracy']['mean'],
        'Precision': best_cv_results['SVM']['precision']['mean'],
        'Recall': best_cv_results['SVM']['recall']['mean'],
        'F1-Score': best_cv_results['SVM']['f1']['mean']
    }
# Create DataFrame
compare df = pd.DataFrame(comparison data)
# Reshape data for visualization
compare_melted = pd.melt(compare_df, id_vars='Model',
                         value_vars=['Accuracy', 'Precision', 'Recall', '
                         var_name='Metric', value_name='Score')
# Plot comparison
plt.figure(figsize=(12, 8))
sns.set_style("whitegrid")
# Create grouped bar chart
ax = sns.barplot(data=compare melted, x='Metric', y='Score', hue='Model',
plt.title('Performance Comparison: Bernoulli NB (Raw) vs. SVM (Preprocess
plt.xlabel('Metric', fontsize=14)
plt.ylabel('Score', fontsize=14)
plt.ylim([0.4, 1.0])
plt.grid(True, linestyle='--', alpha=0.7)
plt.legend(title='Model')
# Add value labels on bars
for p in ax.patches:
    ax.annotate(f'{p.get_height():.3f}',
                (p.get_x() + p.get_width() / 2., p.get_height()),
                ha = 'center', va = 'bottom',
                fontsize=10)
plt.tight_layout()
plt.show()
# Statistical significance test (if applicable)
print("\nPerformance Summary:")
print(compare_df.set_index('Model'))
# Analyze differences
diff_accuracy = abs(comparison_data[0]['Accuracy'] - comparison_data[1]['
diff_f1 = abs(comparison_data[0]['F1-Score'] - comparison_data[1]['F1-Sco
```

```
print(f"\nDifference in Accuracy: {diff_accuracy:.3f}")
print(f"Difference in F1-Score: {diff_f1:.3f}")

# Print conclusions
if comparison_data[0]['Accuracy'] > comparison_data[1]['Accuracy']:
    winner = "Bernoulli NB with raw text"

else:
    winner = "SVM with preprocessed text"

print(f"\nConclusion: {winner} performs better in terms of accuracy.")
print("This suggests that " +
    ("text preprocessing is beneficial for this classification task."
    if winner == "SVM with preprocessed text"
    else "raw text features might contain important signals that are l
```



Performance Summary:

Accuracy Precision Recall F1-Score Model Bernoulli NB (Raw Text) 0.785088 0.704412 0.613158 0.651009

SVM (Preprocessed Text) 0.798246 0.758331 0.581081 0.653448

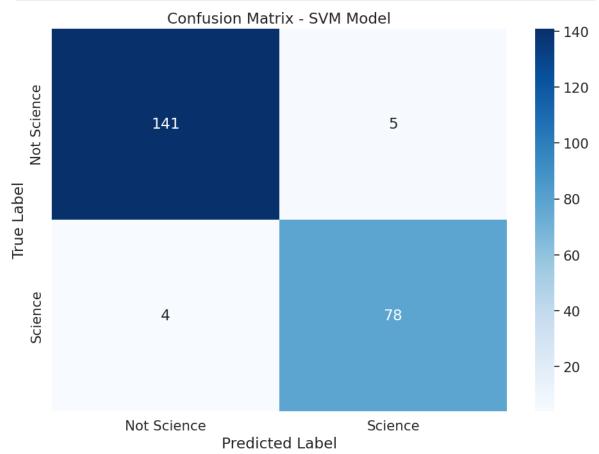
Difference in Accuracy: 0.013 Difference in F1-Score: 0.002

Conclusion: SVM with preprocessed text performs better in terms of accuracy.

This suggests that text preprocessing is beneficial for this classification task.

```
In [100... best_model = best_models['SVM']
In [101... from sklearn.metrics import confusion_matrix, classification_report import seaborn as sns import numpy as np
    import matplotlib.pyplot as plt
```

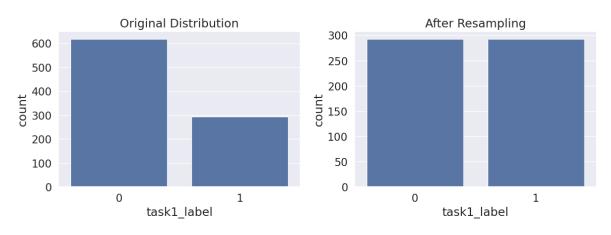
```
# Get predictions from the best model on the test set
y pred = best model.predict(X test task1)
# Calculate the confusion matrix
cm = confusion matrix(y test task1, y pred)
# Create a prettier visualization of the confusion matrix
plt.figure(figsize=(10, 8))
sns.set(font_scale=1.4)
ax = sns.heatmap(cm, annot=True, fmt='d', cmap='Blues',
                 xticklabels=['Not Science', 'Science'],
                 yticklabels=['Not Science', 'Science'])
# Add labels, title and ticks
plt.ylabel('True Label')
plt.xlabel('Predicted Label')
plt.title('Confusion Matrix - SVM Model')
# Calculate metrics
report = classification_report(y_test_task1, y_pred, output_dict=True)
accuracy = (cm[0, 0] + cm[1, 1]) / np.sum(cm)
precision = report['1']['precision']
recall = report['1']['recall']
f1 = report['1']['f1-score']
# Display metrics on the plot
plt.figtext(0.5, 0.01, f'Accuracy: {accuracy:.4f} | Precision: {precision
            ha='center', fontsize=12, bbox={'facecolor': 'lightblue', 'al
plt.tight layout(rect=[0, 0.03, 1, 0.95])
plt.show()
```



Accuracy: 0.9605 | Precision: 0.9398 | Recall: 0.9512 | F1-Score: 0.9455

```
In [102... %pip install imbalanced-learn
         # Import required modules
         from imblearn.under sampling import RandomUnderSampler
         from imblearn.over sampling import RandomOverSampler, SMOTE
         from collections import Counter
        Requirement already satisfied: imbalanced-learn in ./.venv/lib/python3.10/
        site-packages (0.13.0)
        Requirement already satisfied: numpy<3,>=1.24.3 in ./.venv/lib/python3.10/
        site-packages (from imbalanced-learn) (2.2.5)
        Requirement already satisfied: scipy<2,>=1.10.1 in ./.venv/lib/python3.10/
        site-packages (from imbalanced-learn) (1.15.2)
        Requirement already satisfied: scikit-learn<2,>=1.3.2 in ./.venv/lib/pytho
        n3.10/site-packages (from imbalanced-learn) (1.6.1)
        Requirement already satisfied: sklearn-compat<1,>=0.1 in ./.venv/lib/pytho
        n3.10/site-packages (from imbalanced-learn) (0.1.3)
        Requirement already satisfied: joblib<2,>=1.1.1 in ./.venv/lib/python3.10/
        site-packages (from imbalanced-learn) (1.4.2)
        Requirement already satisfied: threadpoolctl<4,>=2.0.0 in ./.venv/lib/pyth
        on3.10/site-packages (from imbalanced-learn) (3.6.0)
        Note: you may need to restart the kernel to use updated packages.
In [103... # For Task 1 (science related classification)
         def apply undersampling(X, y):
             undersampler = RandomUnderSampler(random_state=42)
             X_resampled, y_resampled = undersampler.fit_resample(X, y)
             print(f"Original distribution: {Counter(y)}")
             print(f"Distribution after undersampling: {Counter(y resampled)}")
             return X resampled, y resampled
         # Apply to Task 1
         X_train_task1_under, y_train_task1_under = apply_undersampling(X_train_ta
        Original distribution: Counter({0: 619, 1: 293})
        Distribution after undersampling: Counter({0: 293, 1: 293})
In [104... # Visualize effect of sampling
         def plot sampling comparison(original y, resampled y, title):
             plt.figure(figsize=(12, 5))
             plt.subplot(1, 2, 1)
             sns.countplot(x=original y)
             plt.title('Original Distribution')
             plt.subplot(1, 2, 2)
             sns.countplot(x=resampled y)
             plt.title('After Resampling')
             plt.suptitle(title)
             plt.tight layout()
             plt.show()
         # Example for Task 1
         plot sampling comparison(y train task1, y train task1 under, 'Effect of U
```

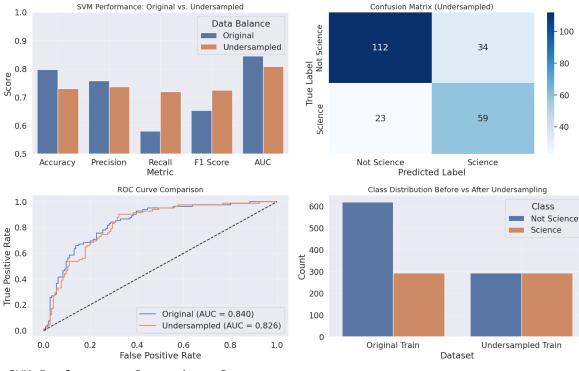
Effect of Undersampling on Task 1



```
In [105...
        # Retrain the best model (SVM) with undersampled data
         best model = best models['SVM']
         # Evaluate on the original test data for fair comparison
         best model.fit(X train task1 under, y train task1 under)
         y pred under = best model.predict(X test task1)
         accuracy under = accuracy score(y test task1, y pred under)
         report_under = classification_report(y_test_task1, y_pred_under, output_d
         cm under = confusion matrix(y test task1, y pred under)
         # Perform k-fold cross validation with undersampled data
         # Initialize StratifiedKFold
         skf = StratifiedKFold(n splits=k folds, shuffle=True, random state=42)
         # Store evaluation metrics
         fold accuracy under = []
         fold precision under = []
         fold recall under = []
         fold f1 under = []
         fold auc under = []
         # Get a balanced dataset
         X balanced = X train task1 under
         y balanced = y train task1 under
         # Perform cross-validation
         for fold, (train idx, test idx) in enumerate(skf.split(X balanced, y bala
             X_train_fold, X_test_fold = X_balanced[train_idx], X_balanced[test_id
             y train fold, y test fold = y balanced.iloc[train idx], y balanced.il
             best model.fit(X train fold, y train fold)
             y_pred_fold = best_model.predict(X_test_fold)
             # Calculate metrics
             acc = accuracy_score(y_test_fold, y_pred_fold)
             prec = precision score(y test fold, y pred fold, zero division=0)
             rec = recall score(y test fold, y pred fold, zero division=0)
             f1 = f1_score(y_test_fold, y_pred_fold, zero_division=0)
             # For AUC
             if hasattr(best model, "predict proba"):
                 y prob = best model.predict proba(X test fold)[:, 1]
                 auc_score = roc_auc_score(y_test_fold, y_prob)
                 fold auc under.append(auc score)
```

```
fold accuracy under.append(acc)
   fold_precision_under.append(prec)
    fold recall under.append(rec)
    fold f1 under.append(f1)
# Compare performance metrics before and after undersampling
comparison data = {
    'Metric': ['Accuracy', 'Precision', 'Recall', 'F1 Score', 'AUC'],
    'Original': [
        best cv results['SVM']['accuracy']['mean'],
        best cv results['SVM']['precision']['mean'],
       best cv results['SVM']['recall']['mean'],
       best_cv_results['SVM']['f1']['mean'],
        best_cv_results['SVM']['auc']['mean'] if 'auc' in best_cv_results
    ],
    'Undersampled': [
        np.mean(fold accuracy under),
        np.mean(fold_precision_under),
        np.mean(fold_recall_under),
       np.mean(fold_f1_under),
        np.mean(fold_auc_under) if fold_auc_under else None
    ]
comparison df = pd.DataFrame(comparison data)
# Create visualizations
plt.figure(figsize=(16, 10))
# Bar chart comparison
plt.subplot(2, 2, 1)
comparison_melted = pd.melt(comparison_df, id_vars=['Metric'],
                           value_vars=['Original', 'Undersampled'],
                           var name='Data Balance', value name='Score')
# Filter out any None values
comparison_melted = comparison_melted.dropna()
sns.barplot(x='Metric', y='Score', hue='Data Balance', data=comparison_me
plt.title('SVM Performance: Original vs. Undersampled', fontsize=14)
plt.ylim(0.5, 1.0)
plt.grid(True, linestyle='--', alpha=0.7)
# Confusion matrix for undersampled model
plt.subplot(2, 2, 2)
sns.heatmap(cm_under, annot=True, fmt='d', cmap='Blues',
            xticklabels=['Not Science', 'Science'],
            yticklabels=['Not Science', 'Science'])
plt.title('Confusion Matrix (Undersampled)', fontsize=14)
plt.ylabel('True Label')
plt.xlabel('Predicted Label')
# ROC Curve comparison if available
if hasattr(best_model, "predict_proba"):
   plt.subplot(2, 2, 3)
    # For original model
   y_prob_orig = best_model.fit(X_train_task1, y_train_task1).predict_pr
   fpr_orig, tpr_orig, _ = roc_curve(y_test_task1, y_prob_orig)
    auc_orig = auc(fpr_orig, tpr_orig)
```

```
# For undersampled model
   best_model.fit(X_train_task1_under, y_train_task1_under)
   y prob under = best model.predict proba(X test task1)[:, 1]
    fpr under, tpr under, = roc curve(y test task1, y prob under)
   auc_under = auc(fpr_under, tpr under)
    # Plot ROC curves
   plt.plot(fpr orig, tpr orig, label=f'Original (AUC = {auc orig:.3f})'
   plt.plot(fpr_under, tpr_under, label=f'Undersampled (AUC = {auc_under
   plt.plot([0, 1], [0, 1], 'k--')
   plt.title('ROC Curve Comparison', fontsize=14)
   plt.xlabel('False Positive Rate')
   plt.ylabel('True Positive Rate')
   plt.legend()
   plt.grid(True, linestyle='--', alpha=0.7)
# Class distribution before and after undersampling
plt.subplot(2, 2, 4)
class_dist = pd.DataFrame({
    'Dataset': ['Original Train'] * 2 + ['Undersampled Train'] * 2,
    'Class': ['Not Science', 'Science'] * 2,
    'Count': [
        sum(y train task1 == 0),
        sum(y_train_task1 == 1),
        sum(y train task1 under == 0),
        sum(y_train_task1_under == 1)
    ]
})
sns.barplot(x='Dataset', y='Count', hue='Class', data=class dist)
plt.title('Class Distribution Before vs After Undersampling', fontsize=14
plt.grid(True, linestyle='--', alpha=0.7)
plt.tight_layout()
plt.show()
# Print summary
print("\nSVM Performance Comparison Summary:")
print(comparison_df.set_index('Metric'))
print("\n0riginal Class Distribution:", Counter(y_train_task1))
print("Undersampled Class Distribution:", Counter(y train task1 under))
# Calculate improvement percentages
improvement_df = pd.DataFrame({
    'Metric': comparison_df['Metric'],
    'Improvement (%)': [
        ((comparison_df['Undersampled'][i] - comparison_df['Original'][i]
        if comparison df['Original'][i] is not None and comparison df['Un'
        else None
        for i in range(len(comparison df))
    ]
print("\nRelative Improvement After Undersampling:")
print(improvement df.set index('Metric'))
```



SVM Performance Comparison Summary: Original Undersampled

Metric
Accuracy 0.798246 0.730479
Precision 0.758331 0.737308
Recall 0.581081 0.719885
F1 Score 0.653448 0.725460
AUC 0.846289 0.809384

Original Class Distribution: Counter({0: 619, 1: 293})
Undersampled Class Distribution: Counter({0: 293, 1: 293})

Relative Improvement After Undersampling:

```
Improvement (%)
```

Metric
Accuracy -8.489412
Precision -2.772259
Recall 23.887196
F1 Score 11.020415
AUC -4.360885

```
In [106... # TASK 2: Scientific Claim/Reference Classification

# Since Task 2 is only applicable to science-related tweets, we need to u
print("Task 2: Scientific Claim/Reference Classification\n" + "="*50)

# Get a dense version of features for science-related tweets
# Convert the pandas Series to a numpy array with .values
X_dense_sci = X_selected[(df['science_related'] == 1).values].toarray()
y_task2 = df_sci['task2_label']

# Split data for Task 2
X_train_task2, X_test_task2, y_train_task2, y_test_task2 = train_test_spl
    X_dense_sci, y_task2, test_size=0.2, random_state=42
)

# Define the best models from Task 1 (you can modify these based on Task
task2_models = {
```

```
'Gaussian NB': GaussianNB(var_smoothing=0.0152),
'Multinomial NB': MultinomialNB(alpha=0.4641, fit_prior=True),
'Complement NB': ComplementNB(alpha=2.8480, fit_prior=True),
'Bernoulli NB': BernoulliNB(alpha=0.17475, fit_prior=True),
'KNN': KNeighborsClassifier(n_neighbors=7, weights='uniform', metric='SVM': SVC(C=1.4174742, kernel='rbf', gamma='scale', probability=True'Logistic Regression': LogisticRegression(C=np.float64(4.328761281083'Random Forest': RandomForestClassifier(n_estimators=200, max_depth=N)

# Train and evaluate models for Task 2
task2_results = {}
for name, model in task2_models.items():
    task2_results[name] = evaluate_model(
        model, X_train_task2, X_test_task2, y_train_task2, y_test_task2,
)
    print("\n")
```

Task 2: Scientific Claim/Reference Classification

533			
precision	recall	f1-score	support
0.30	0.43	0.35	7
0.94	0.90	0.92	68
		0.85	75
0.62	0.66		75
0.88	0.85	0.86	75
		6.3	
precision	recall	fl-score	support
0.00	0.00	0.00	7
0.91	1.00	0.95	68
		0.91	75
0.45	0.50	0.48	75
0.82	0.91	0.86	75
ment NB			
precision	recall	f1-score	support
0.18	0.57	0.28	7
0.94	0.74	0.83	68
		0.72	75
0.56	0.65	0.55	75
0.87	0.72	0.78	75
precision	recall	f1-score	support
0.00	0.00	0.00	7
0.90	0.94	0.92	68
		0.85	75
0.45	0.47	0.46	75
0.82	0.85	0.83	75
	an NB 533 precision	an NB 533 precision	precision recall f1-score 0.30

```
/home/hurel/Documents/repo/projet-ml/.venv/lib/python3.10/site-packages/sk
learn/metrics/ classification.py:1565: UndefinedMetricWarning: Precision i
s ill-defined and being set to 0.0 in labels with no predicted samples. Us
e `zero division` parameter to control this behavior.
  warn prf(average, modifier, f"{metric.capitalize()} is", len(result))
/home/hurel/Documents/repo/projet-ml/.venv/lib/python3.10/site-packages/sk
learn/metrics/ classification.py:1565: UndefinedMetricWarning: Precision i
s ill-defined and being set to 0.0 in labels with no predicted samples. Us
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e `zero_division` parameter to control this behavior.
  _warn_prf(average, modifier, f"{metric.capitalize()} is", len(result))
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s ill-defined and being set to 0.0 in labels with no predicted samples. Us
e `zero division` parameter to control this behavior.
  _warn_prf(average, modifier, f"{metric.capitalize()} is", len(result))
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s ill-defined and being set to 0.0 in labels with no predicted samples. Us
e `zero_division` parameter to control this behavior.
  _warn_prf(average, modifier, f"{metric.capitalize()} is", len(result))
/home/hurel/Documents/repo/projet-ml/.venv/lib/python3.10/site-packages/sk
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s ill-defined and being set to 0.0 in labels with no predicted samples. Us
e `zero_division` parameter to control this behavior.
  warn prf(average, modifier, f"{metric.capitalize()} is", len(result))
/home/hurel/Documents/repo/projet-ml/.venv/lib/python3.10/site-packages/sk
learn/metrics/_classification.py:1565: UndefinedMetricWarning: Precision i
s ill-defined and being set to 0.0 in labels with no predicted samples. Us
e `zero_division` parameter to control this behavior.
  _warn_prf(average, modifier, f"{metric.capitalize()} is", len(result))
/home/hurel/Documents/repo/projet-ml/.venv/lib/python3.10/site-packages/sk
learn/metrics/_classification.py:1565: UndefinedMetricWarning: Precision i
s ill-defined and being set to 0.0 in labels with no predicted samples. Us
e `zero_division` parameter to control this behavior.
  _warn_prf(average, modifier, f"{metric.capitalize()}            is", len(result))
/home/hurel/Documents/repo/projet-ml/.venv/lib/python3.10/site-packages/sk
learn/metrics/ classification.py:1565: UndefinedMetricWarning: Precision i
s ill-defined and being set to 0.0 in labels with no predicted samples. Us
e `zero_division` parameter to control this behavior.
  _warn_prf(average, modifier, f"{metric.capitalize()} is", len(result))
```

Model: KNN					
Accuracy: 0.9	precision	recall	f1-score	support	
0 1	0.00 0.91	0.00 1.00	0.00 0.95	7 68	
accuracy macro avg weighted avg	0.45 0.82	0.50 0.91	0.91 0.48 0.86	75 75 75	
Model: SVM Accuracy: 0.9	200 precision	recall	fl-score	support	
0 1	1.00 0.92	0.14 1.00	0.25 0.96	7 68	
accuracy macro avg weighted avg	0.96 0.93	0.57 0.92	0.92 0.60 0.89	75 75 75	
Model: Logist Accuracy: 0.9		n			
,	precision	recall	f1-score	support	
0 1	1.00 0.92	0.14 1.00	0.25 0.96	7 68	
accuracy macro avg weighted avg	0.96 0.93	0.57 0.92	0.92 0.60 0.89	75 75 75	
Model: Random Accuracy: 0.9		recall	f1-score	support	
0 1	1.00	0.14 1.00	0.25 0.96	7 68	
accuracy	0.06	0 57	0.92	75 75	

```
In [107... # Cross-validation for Task 2
print("Task 2: Cross-Validation\n" + "="*40)

# Initialize k-fold cross validation
k_folds = 10
skf = StratifiedKFold(n_splits=k_folds, shuffle=True, random_state=42)
```

0.60

0.89

75

75

macro avg 0.96 0.57 weighted avg 0.93 0.92

```
# Dictionary to store CV results
task2 cv results = {}
# Perform k-fold cross validation for each model
for model_name, model in task2_models.items():
    print(f"Performing {k folds}-fold cross-validation for {model name}...
    # Initialize lists to store performance metrics for each fold
   fold accuracy = []
   fold_precision = []
   fold recall = []
    fold_f1 = []
    fold auc = []
    # For each fold
    for fold, (train_idx, test_idx) in enumerate(skf.split(X_dense_sci, y)
        # Split data
        X train fold, X test fold = X dense sci[train idx], X dense sci[t
        y_train_fold, y_test_fold = y_task2.iloc[train_idx], y_task2.iloc
        # Train model
        model.fit(X_train_fold, y_train_fold)
        # Make predictions
        y_pred_fold = model.predict(X_test_fold)
        # Calculate metrics
        acc = accuracy_score(y_test_fold, y_pred_fold)
        prec = precision_score(y_test_fold, y_pred_fold, zero_division=0)
        rec = recall_score(y_test_fold, y_pred_fold, zero_division=0)
        f1 = f1_score(y_test_fold, y_pred_fold, zero_division=0)
        # For AUC, we need probability estimates
        try:
            if hasattr(model, "predict_proba"):
                y_prob = model.predict_proba(X_test_fold)[:, 1]
                auc score = roc_auc_score(y_test_fold, y_prob)
                fold_auc.append(auc_score)
        except:
            pass
        fold accuracy.append(acc)
        fold precision.append(prec)
        fold_recall.append(rec)
        fold_f1.append(f1)
    # Store average metrics and standard deviations
    task2 cv results[model name] = {
        'accuracy': {
            'mean': np.mean(fold accuracy),
            'std': np.std(fold_accuracy)
        },
        'precision': {
            'mean': np.mean(fold precision),
            'std': np.std(fold precision)
        },
        'recall': {
            'mean': np.mean(fold_recall),
            'std': np.std(fold_recall)
        },
```

```
'fl': {
        'mean': np.mean(fold_f1),
        'std': np.std(fold_f1)
}

if fold_auc:
    task2_cv_results[model_name]['auc'] = {
        'mean': np.mean(fold_auc),
        'std': np.std(fold_auc)
}

print(f" Average: AUC={task2_cv_results[model_name]['auc']['mean

print(f" Average: Accuracy={task2_cv_results[model_name]['accuracy']
    print(f" Average: F1 Score={task2_cv_results[model_name]['f1']['mean print()
```

Task 2: Cross-Validation _____ Performing 10-fold cross-validation for Gaussian NB... Average: $AUC=0.5660 \pm 0.1187$ Average: Accuracy= 0.8217 ± 0.0635 Average: F1 Score= 0.8988 ± 0.0375 Performing 10-fold cross-validation for Multinomial NB... Average: AUC=0.7806 \pm 0.0932 Average: Accuracy= 0.9148 ± 0.0152 Average: F1 Score= 0.9554 ± 0.0081 Performing 10-fold cross-validation for Complement NB... Average: $AUC=0.7468 \pm 0.0979$ Average: Accuracy= 0.7151 ± 0.0768 Average: F1 Score= 0.8205 ± 0.0532 Performing 10-fold cross-validation for Bernoulli NB... Average: $AUC=0.7860 \pm 0.0991$ Average: Accuracy= 0.8935 ± 0.0332 Average: F1 Score= 0.9423 ± 0.0184 Performing 10-fold cross-validation for KNN... Average: $AUC=0.5324 \pm 0.0660$ Average: Accuracy= 0.9094 ± 0.0126 Average: F1 Score= 0.9525 ± 0.0069 Performing 10-fold cross-validation for SVM... Average: $AUC=0.7151 \pm 0.1219$ Average: Accuracy= 0.9175 ± 0.0179 Average: F1 Score= 0.9567 ± 0.0095 Performing 10-fold cross-validation for Logistic Regression... Average: AUC=0.7019 \pm 0.1686 Average: Accuracy= 0.9068 ± 0.0242 Average: F1 Score= 0.9503 ± 0.0131 Performing 10-fold cross-validation for Random Forest... Average: $AUC=0.7275 \pm 0.1402$ Average: Accuracy= 0.9148 ± 0.0152 Average: F1 Score= 0.9554 ± 0.0081

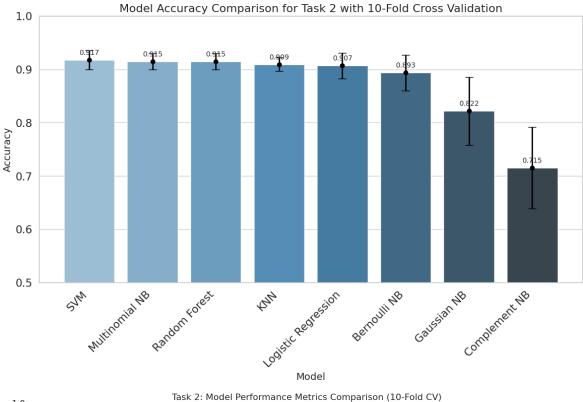
```
In [108... # Visualize Task 2 cross-validation results
         # Create DataFrame for visualization
         task2 results df = pd.DataFrame({
              'Model': [],
              'Metric': [],
              'Mean': [],
              'Std': []
         })
         for model name in task2 cv results:
             for metric in ['accuracy', 'precision', 'recall', 'f1']:
                  task2 results df = pd.concat([task2 results df, pd.DataFrame({
                      'Model': [model_name],
                      'Metric': [metric.capitalize()],
                      'Mean': [task2 cv results[model name][metric]['mean']],
                      'Std': [task2 cv results[model name][metric]['std']]
                 })], ignore index=True)
             if 'auc' in task2_cv_results[model_name]:
                 task2 results df = pd.concat([task2 results df, pd.DataFrame({
                      'Model': [model name],
                      'Metric': ['AUC'],
                      'Mean': [task2 cv results[model name]['auc']['mean']],
                      'Std': [task2 cv results[model name]['auc']['std']]
                 })], ignore index=True)
         # Sort models by accuracy
         task2 model order = task2 results df[task2 results df['Metric'] == 'Accur
         # Create accuracy bar plot
         plt.figure(figsize=(12, 8))
         sns.set style("whitegrid")
         # Create bar plot for accuracy
         ax = sns.barplot(
             data=task2_results_df[task2_results_df['Metric'] == 'Accuracy'],
             x='Model',
             y='Mean',
             order=task2 model order,
             palette='Blues d'
         )
         # Add error bars
         for i, model in enumerate(task2 model order):
              row = task2_results_df[(task2_results_df['Model'] == model) & (task2_
             ax.errorbar(
                 i, row['Mean'], yerr=row['Std'],
                 fmt='o', color='black', elinewidth=2, capsize=6
             )
         # Add value labels on top of bars
         for i, bar in enumerate(ax.patches):
             ax.text(
                 bar.get x() + bar.get width()/2,
                 bar.get height() + 0.01,
                 f"{bar.get_height():.3f}",
                 ha='center',
                 fontsize=10
         plt.title(f'Model Accuracy Comparison for Task 2 with {k folds}-Fold Cros
```

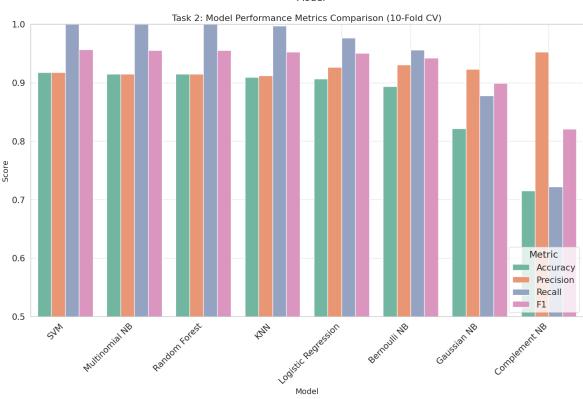
```
plt.xlabel('Model', fontsize=14)
plt.ylabel('Accuracy', fontsize=14)
plt.xticks(rotation=45, ha='right')
plt.ylim([0.5, 1.0])
plt.tight layout()
plt.show()
# Create a grouped bar chart for all metrics
plt.figure(figsize=(15, 10))
sns.set_style("whitegrid")
# Create grouped bar plot
ax = sns.barplot(
    data=task2_results_df[task2_results_df['Metric'] != 'AUC'],
   x='Model',
   y='Mean',
   hue='Metric',
    order=task2 model order,
    palette='Set2'
plt.title(f'Task 2: Model Performance Metrics Comparison ({k_folds}-Fold
plt.xlabel('Model', fontsize=14)
plt.ylabel('Score', fontsize=14)
plt.xticks(rotation=45, ha='right')
plt.legend(title='Metric', loc='lower right')
plt.ylim([0.5, 1.0])
plt.grid(True, linestyle='--', alpha=0.7)
plt.tight layout()
plt.show()
```

/tmp/ipykernel 368067/1744879752.py:34: FutureWarning:

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

```
ax = sns.barplot(
```





```
In [109... # Hyperparameter optimization for the best model on Task 2
# Let's assume SVM was the best model (update based on your results)
best_task2_model_name = 'SVM' # Replace with the actual best model from
best_task2_model = task2_models[best_task2_model_name]

print(f"Performing Grid Search for {best_task2_model_name} on Task 2...")

# Define the parameter grid for the best model
if best_task2_model_name == 'SVM':
    param_grid = {
        'C': np.logspace(-3, 3, 10),
        'gamma': ['scale', 'auto'] + list(np.logspace(-3, 3, 5)),
        'kernel': ['rbf', 'linear']
```

```
}
elif best task2 model name == 'Random Forest':
    param grid = {
        'n estimators': [50, 100, 200],
        'max depth': [None, 10, 20, 30],
        'min_samples_split': [2, 5, 10],
        'min samples leaf': [1, 2, 4]
elif best task2 model name.endswith('NB'):
    if best_task2_model_name == 'Gaussian NB':
        param grid = {
            'var smoothing': np.logspace(-10, 0, 11)
        }
    else:
        param grid = {
            'alpha': np.logspace(-3, 3, 10),
            'fit prior': [True, False]
else:
    # Default grid for other models
    param_grid = {
        'C': np.logspace(-3, 3, 10)
    }
# Create and fit GridSearchCV
grid search task2 = GridSearchCV(
    estimator=best_task2_model,
    param_grid=param_grid,
    cv=5,
    scoring='accuracy',
    n jobs=-1,
    verbose=1
grid search task2.fit(X train task2, y train task2)
# Display the best parameters and score
print(f"Best Parameters: {grid_search_task2.best_params_}")
print(f"Best Cross-Validation Score: {grid_search_task2.best_score_:.4f}"
# Evaluate on test set
best model task2 = grid search task2.best estimator
y_pred_task2 = best_model_task2.predict(X_test_task2)
accuracy_task2 = accuracy_score(y_test_task2, y_pred_task2)
print(f"Test Accuracy with Best Parameters: {accuracy_task2:.4f}")
print("\nClassification Report:")
print(classification_report(y_test_task2, y_pred_task2))
# Plot confusion matrix
plt.figure(figsize=(10, 8))
cm = confusion_matrix(y_test_task2, y_pred_task2)
sns.heatmap(cm, annot=True, fmt='d', cmap='Blues',
            xticklabels=['No Claim/Ref', 'Claim/Ref'],
            yticklabels=['No Claim/Ref', 'Claim/Ref'])
plt.title(f'Confusion Matrix - Task 2 ({best_task2_model_name})')
plt.ylabel('True Label')
plt.xlabel('Predicted Label')
plt.tight_layout()
plt.show()
```

75

75

75

accuracy

macro avg

weighted avg

0.96

0.93

```
Performing Grid Search for SVM on Task 2...
Fitting 5 folds for each of 140 candidates, totalling 700 fits
Best Parameters: {'C': np.float64(215.44346900318823), 'gamma': 'auto', 'k
ernel': 'rbf'}
Best Cross-Validation Score: 0.9167
Test Accuracy with Best Parameters: 0.9200
Classification Report:
              precision
                           recall f1-score
                                             support
           0
                   1.00
                             0.14
                                       0.25
                                                    7
           1
                   0.92
                             1.00
                                       0.96
                                                   68
```

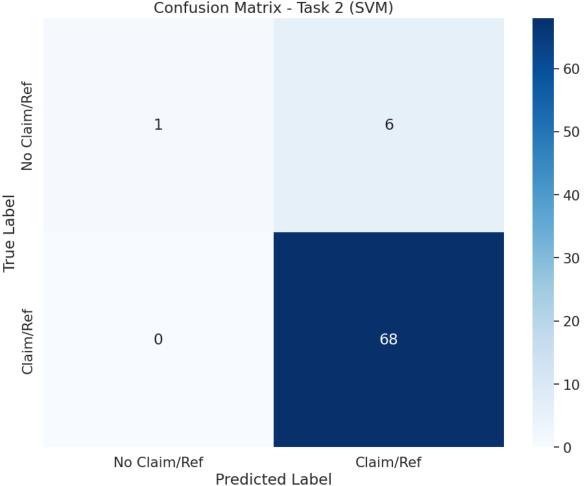
0.92

0.60

0.89

0.57

0.92



```
In [110... # Check for class imbalance in Task 2
         print("Task 2 class distribution:")
         print(y_task2.value_counts())
         print(y_task2.value_counts(normalize=True).round(3) * 100, '%')
         # Apply class balancing for Task 2 if needed
         from imblearn.over_sampling import SMOTE
         from imblearn.under_sampling import RandomUnderSampler
         from sklearn.base import clone # Add this import
         # Let's try SMOTE for handling imbalance
```

```
smote = SMOTE(random state=42)
X train task2 balanced, y train task2 balanced = smote.fit resample(X train
print("\nClass distribution after SMOTE balancing:")
print(pd.Series(y train task2 balanced).value counts())
# Train the best model with balanced data
best model balanced = clone(best model task2)
best_model_balanced.fit(X_train_task2_balanced, y_train_task2_balanced)
# Evaluate on the test set
y pred balanced = best model balanced.predict(X test task2)
accuracy_balanced = accuracy_score(y_test_task2, y_pred_balanced)
print(f"\nTest Accuracy with Balanced Training Data: {accuracy_balanced:.
print("\nClassification Report with Balanced Training Data:")
print(classification_report(y_test_task2, y_pred_balanced))
# Compare balanced vs unbalanced training
plt.figure(figsize=(12, 6))
metrics = ['Accuracy', 'Precision', 'Recall', 'F1-Score']
unbalanced_scores = [
    accuracy_task2,
    precision_score(y_test_task2, y_pred_task2),
    recall_score(y_test_task2, y_pred_task2),
    f1_score(y_test_task2, y_pred_task2)
balanced_scores = [
    accuracy_balanced,
    precision_score(y_test_task2, y_pred_balanced),
    recall_score(y_test_task2, y_pred_balanced),
    f1_score(y_test_task2, y_pred_balanced)
1
x = np.arange(len(metrics))
width = 0.35
plt.bar(x - width/2, unbalanced scores, width, label='Unbalanced Training
plt.bar(x + width/2, balanced_scores, width, label='Balanced Training')
plt.xticks(x, metrics)
plt.ylabel('Score')
plt.title('Impact of Class Balancing on Task 2 Performance')
plt.legend()
plt.ylim(0, 1)
plt.tight layout()
plt.show()
```

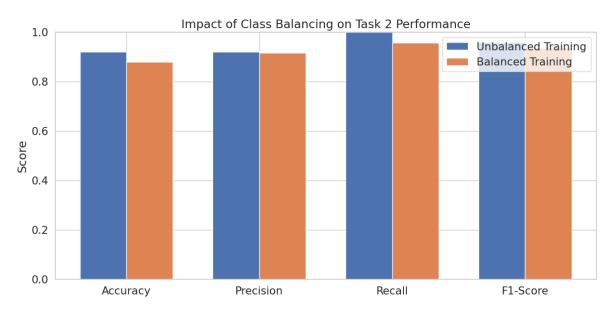
```
Task 2 class distribution:
task2_label
1    342
0    33
Name: count, dtype: int64
task2_label
1    91.2
0    8.8
Name: proportion, dtype: float64 %

Class distribution after SMOTE balancing:
task2_label
1    274
0    274
Name: count, dtype: int64
```

Test Accuracy with Balanced Training Data: 0.8800

Classification Report with Balanced Training Data:

	precision	recall	f1-score	support
0	0.25	0.14	0.18	7
1	0.92	0.96	0.94	68
accuracy			0.88	75
macro avg weighted avg	0.58 0.85	0.55 0.88	0.56 0.86	75 75



```
In [111... # TASK 3: Scientific Type Classification
print("Task 3: Scientific Type Classification\n" + "="*50)

# We already have the preprocessed features for science-related tweets (X # And we have the labels in df_sci['task3_label'] (with values 0, 1, 2 fo y_task3 = df_sci['task3_label']

# Split data for Task 3
X_train_task3, X_test_task3, y_train_task3, y_test_task3 = train_test_spl X_dense_sci, y_task3, test_size=0.2, random_state=42
)

# Use the same models as in Task 2
task3_models = {
```

```
'Gaussian NB': GaussianNB(var_smoothing=0.0152),
    'Multinomial NB': MultinomialNB(alpha=0.4641, fit_prior=True),
    'Complement NB': ComplementNB(alpha=2.8480, fit_prior=True),
    'Bernoulli NB': BernoulliNB(alpha=0.17475, fit_prior=True),
    'KNN': KNeighborsClassifier(n_neighbors=7, weights='uniform', metric='SVM': SVC(C=1.4174742, kernel='rbf', gamma='scale', probability=True'Logistic Regression': LogisticRegression(C=np.float64(4.328761281083'Random Forest': RandomForestClassifier(n_estimators=200, max_depth=N)

# Train and evaluate models for Task 3
task3_results = {}
for name, model in task3_models.items():
    task3_results[name] = evaluate_model(
        model, X_train_task3, X_test_task3, y_train_task3, y_test_task3,
)
    print("\n")
```

Task 3: Scientific Type Classification

Model , Caussian NP	

Model: Gaussia		======	=======	====
·	precision	recall	f1-score	support
0	0.86	0.89	0.87	54
1 2	0.67 0.30	0.43 0.43	0.52 0.35	14 7
2	0.50	0.45	0.55	,
accuracy	0.61	0 50	0.76	75 75
macro avg weighted avg	0.61 0.77	0.58 0.76	0.58 0.76	75 75
	• • • • • • • • • • • • • • • • • • • •			
Model: Multino Accuracy: 0.76				
Accuracy: 0.70	precision	recall	f1-score	support
0	0.76	1.00	0.86	54
1	0.75	0.21	0.33	14
2	0.00	0.00	0.00	7
accuracy			0.76	75
macro avg weighted avg	0.50 0.69	0.40 0.76	0.40 0.68	75 75
Model: Complem	ent NB			
Accuracy: 0.60	precision	recall	f1-score	support
0 1	0.78 0.31	0.72 0.29	0.75 0.30	54 14
2	0.31	0.29	0.30	7
accuracy	0 42	0 42	0.60	75 75
macro avg weighted avg	0.42 0.63	0.43 0.60	0.42 0.61	75 75
weighted dvg	0.03	0.00	0.01	,,
Model: Bernoul Accuracy: 0.70				
	precision	recall	f1-score	support
0	0.78	0.91	0.84	54
1	0.43	0.21	0.29	14
2	0.20	0.14	0.17	7
accuracy	0.47	0 42	0.71	75 75

Model: KNN

weighted avg

Accuracy: 0.7200

macro avg 0.47 ighted avg 0.66

84 sur 96 01/05/2025 17:27

0.42

0.71

0.43

0.67

75

75

	precision	recall	f1-score	support
0 1 2	0.73 0.50 0.00	0.96 0.14 0.00	0.83 0.22 0.00	54 14 7
accuracy macro avg weighted avg	0.41 0.62	0.37 0.72	0.72 0.35 0.64	75 75 75
Model: SVM Accuracy: 0.7	733 precision	recall	f1-score	support
0 1 2	0.79 0.60 0.50	1.00 0.21 0.14	0.89 0.32 0.22	54 14 7
accuracy macro avg weighted avg	0.63 0.73	0.45 0.77	0.77 0.47 0.72	75 75 75

```
/home/hurel/Documents/repo/projet-ml/.venv/lib/python3.10/site-packages/sk
learn/metrics/ classification.py:1565: UndefinedMetricWarning: Precision i
s ill-defined and being set to 0.0 in labels with no predicted samples. Us
e `zero division` parameter to control this behavior.
  warn prf(average, modifier, f"{metric.capitalize()} is", len(result))
/home/hurel/Documents/repo/projet-ml/.venv/lib/python3.10/site-packages/sk
learn/metrics/ classification.py:1565: UndefinedMetricWarning: Precision i
s ill-defined and being set to 0.0 in labels with no predicted samples. Us
e `zero division` parameter to control this behavior.
  warn prf(average, modifier, f"{metric.capitalize()} is", len(result))
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s ill-defined and being set to 0.0 in labels with no predicted samples. Us
e `zero_division` parameter to control this behavior.
  _warn_prf(average, modifier, f"{metric.capitalize()} is", len(result))
/home/hurel/Documents/repo/projet-ml/.venv/lib/python3.10/site-packages/sk
learn/metrics/_classification.py:1565: UndefinedMetricWarning: Precision i
s ill-defined and being set to 0.0 in labels with no predicted samples. Us
e `zero_division` parameter to control this behavior.
  _warn_prf(average, modifier, f"{metric.capitalize()} is", len(result))
/home/hurel/Documents/repo/projet-ml/.venv/lib/python3.10/site-packages/sk
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s ill-defined and being set to 0.0 in labels with no predicted samples. Us
e `zero_division` parameter to control this behavior.
  _warn_prf(average, modifier, f"{metric.capitalize()} is", len(result))
/home/hurel/Documents/repo/projet-ml/.venv/lib/python3.10/site-packages/sk
learn/metrics/ classification.py:1565: UndefinedMetricWarning: Precision i
s ill-defined and being set to 0.0 in labels with no predicted samples. Us
e `zero division` parameter to control this behavior.
  _warn_prf(average, modifier, f"{metric.capitalize()} is", len(result))
/home/hurel/Documents/repo/projet-ml/.venv/lib/python3.10/site-packages/sk
learn/metrics/_classification.py:1565: UndefinedMetricWarning: Precision i
s ill-defined and being set to 0.0 in labels with no predicted samples. Us
e `zero_division` parameter to control this behavior.
  _warn_prf(average, modifier, f"{metric.capitalize()} is", len(result))
/home/hurel/Documents/repo/projet-ml/.venv/lib/python3.10/site-packages/sk
learn/metrics/_classification.py:1565: UndefinedMetricWarning: Precision i
s ill-defined and being set to 0.0 in labels with no predicted samples. Us
e `zero division` parameter to control this behavior.
  _warn_prf(average, modifier, f"{metric.capitalize()} is", len(result))
/home/hurel/Documents/repo/projet-ml/.venv/lib/python3.10/site-packages/sk
learn/metrics/ classification.py:1565: UndefinedMetricWarning: Precision i
s ill-defined and being set to 0.0 in labels with no predicted samples. Us
e `zero_division` parameter to control this behavior.
  warn prf(average, modifier, f"{metric.capitalize()} is", len(result))
/home/hurel/Documents/repo/projet-ml/.venv/lib/python3.10/site-packages/sk
learn/metrics/_classification.py:1565: UndefinedMetricWarning: Precision i
s ill-defined and being set to 0.0 in labels with no predicted samples. Us
e `zero_division` parameter to control this behavior.
  _warn_prf(average, modifier, f"{metric.capitalize()} is", len(result))
/home/hurel/Documents/repo/projet-ml/.venv/lib/python3.10/site-packages/sk
learn/metrics/_classification.py:1565: UndefinedMetricWarning: Precision i
s ill-defined and being set to 0.0 in labels with no predicted samples. Us
e `zero_division` parameter to control this behavior.
  _warn_prf(average, modifier, f"{metric.capitalize()}            is", len(result))
/home/hurel/Documents/repo/projet-ml/.venv/lib/python3.10/site-packages/sk
learn/metrics/ classification.py:1565: UndefinedMetricWarning: Precision i
s ill-defined and being set to 0.0 in labels with no predicted samples. Us
e `zero_division` parameter to control this behavior.
  _warn_prf(average, modifier, f"{metric.capitalize()} is", len(result))
```

Model: Logistic Regression

Accuracy: 0.7067

	precision	recall	f1-score	support
0	0.77	0.89	0.83	54
1	0.33	0.29	0.31	14
2	1.00	0.14	0.25	7
accuracy			0.71	75
macro avg	0.70	0.44	0.46	75
weighted avg	0.71	0.71	0.68	75

Model: Random Forest Accuracy: 0.6667

Accuracy. 0		,0,7			
		precision	recall	f1-score	support
	0	0.82	0.78	0.80	54
	1	0.29	0.43	0.34	14
	2	0.67	0.29	0.40	7
accurac	у			0.67	75
macro av	′g	0.59	0.50	0.51	75
weighted av	'g	0.71	0.67	0.68	75

```
In [112... # Cross-validation for Task 3
         print("Task 3: Cross-Validation\n" + "="*40)
         # Initialize k-fold cross validation
         k folds = 10
         skf = StratifiedKFold(n_splits=k_folds, shuffle=True, random_state=42)
         # Dictionary to store CV results
         task3_cv_results = {}
         # Perform k-fold cross validation for each model
         for model_name, model in task3_models.items():
             print(f"Performing {k folds}-fold cross-validation for {model name}...
             # Initialize lists to store performance metrics for each fold
             fold accuracy = []
             fold_precision = []
             fold recall = []
             fold f1 = []
             # For each fold
             for fold, (train idx, test idx) in enumerate(skf.split(X dense sci, y
                 # Split data
                 X_train_fold, X_test_fold = X_dense_sci[train_idx], X_dense_sci[t
                 y train fold, y test fold = y task3.iloc[train idx], y task3.iloc
                 # Train model
                 model.fit(X_train_fold, y_train_fold)
                 # Make predictions
                 y pred fold = model.predict(X test fold)
```

```
# Calculate metrics
    acc = accuracy_score(y_test_fold, y_pred_fold)
    # For multi-class classification, use macro averaging
    prec = precision_score(y_test_fold, y_pred_fold, average='macro',
    rec = recall_score(y_test_fold, y_pred_fold, average='macro', zer
    f1 = f1 score(y test fold, y pred fold, average='macro', zero div
    fold_accuracy.append(acc)
    fold precision.append(prec)
    fold recall.append(rec)
    fold f1.append(f1)
# Store average metrics and standard deviations
task3_cv_results[model_name] = {
    'accuracy': {
        'mean': np.mean(fold accuracy),
        'std': np.std(fold accuracy)
    },
    'precision': {
        'mean': np.mean(fold_precision),
        'std': np.std(fold_precision)
    },
    'recall': {
        'mean': np.mean(fold_recall),
        'std': np.std(fold recall)
    'f1': {
        'mean': np.mean(fold f1),
        'std': np.std(fold f1)
    }
}
print(f"
         Average: Accuracy={task3_cv_results[model_name]['accuracy']
print(f" Average: F1 Score={task3 cv results[model name]['f1']['mean
print()
```

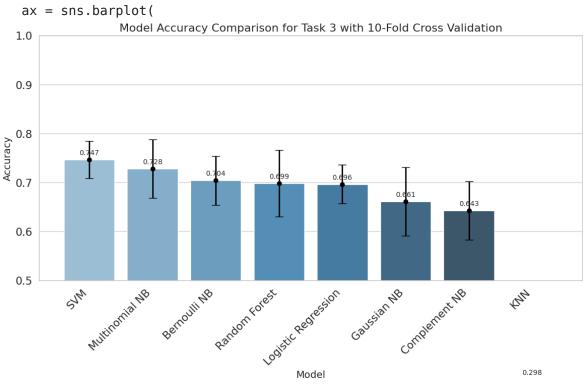
```
Task 3: Cross-Validation
        Performing 10-fold cross-validation for Gaussian NB...
          Average: Accuracy=0.6612 \pm 0.0701
          Average: F1 Score=0.4518 \pm 0.0643
        Performing 10-fold cross-validation for Multinomial NB...
          Average: Accuracy=0.7280 \pm 0.0599
          Average: F1 Score=0.4004 \pm 0.0922
        Performing 10-fold cross-validation for Complement NB...
          Average: Accuracy=0.6428 \pm 0.0595
          Average: F1 Score=0.4886 \pm 0.0616
        Performing 10-fold cross-validation for Bernoulli NB...
          Average: Accuracy=0.6428 \pm 0.0595
          Average: F1 Score=0.4886 \pm 0.0616
        Performing 10-fold cross-validation for Bernoulli NB...
          Average: Accuracy=0.7042 \pm 0.0498
          Average: F1 Score=0.4997 \pm 0.0956
        Performing 10-fold cross-validation for KNN...
          Average: Accuracy=0.2983 \pm 0.0671
          Average: F1 Score=0.1982 \pm 0.0542
        Performing 10-fold cross-validation for SVM...
          Average: Accuracy=0.7467 \pm 0.0379
          Average: F1 Score=0.4304 \pm 0.0634
        Performing 10-fold cross-validation for Logistic Regression...
          Average: Accuracy=0.6962 \pm 0.0395
          Average: F1 Score=0.4445 \pm 0.0834
        Performing 10-fold cross-validation for Random Forest...
          Average: Accuracy=0.6985 \pm 0.0678
          Average: F1 Score=0.4920 \pm 0.1049
In [113... | # Visualize Task 3 cross-validation results
         # Create DataFrame for visualization
         task3 results df = pd.DataFrame({
              'Model': [],
              'Metric': [],
              'Mean': [],
              'Std': []
         })
         for model name in task3 cv results:
             for metric in ['accuracy', 'precision', 'recall', 'f1']:
                  task3_results_df = pd.concat([task3_results_df, pd.DataFrame({
                      'Model': [model name],
                      'Metric': [metric.capitalize()],
                      'Mean': [task3 cv results[model name][metric]['mean']],
                      'Std': [task3_cv_results[model_name][metric]['std']]
                  })], ignore index=True)
         # Sort models by accuracy
         task3_model_order = task3_results_df[task3_results_df['Metric'] == 'Accur
```

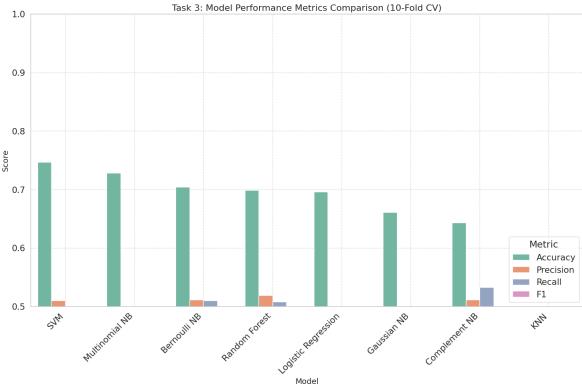
```
# Create accuracy bar plot
plt.figure(figsize=(12, 8))
sns.set_style("whitegrid")
# Create bar plot for accuracy
ax = sns.barplot(
   data=task3 results df[task3 results df['Metric'] == 'Accuracy'],
   x='Model',
   y='Mean',
   order=task3 model order,
   palette='Blues d'
)
# Add error bars
for i, model in enumerate(task3 model order):
    row = task3_results_df[(task3_results_df['Model'] == model) & (task3_
   ax.errorbar(
        i, row['Mean'], yerr=row['Std'],
        fmt='o', color='black', elinewidth=2, capsize=6
    )
# Add value labels on top of bars
for i, bar in enumerate(ax.patches):
    ax.text(
        bar.get_x() + bar.get_width()/2,
        bar.get height() + 0.01,
        f"{bar.get_height():.3f}",
        ha='center',
        fontsize=10
    )
plt.title(f'Model Accuracy Comparison for Task 3 with {k_folds}-Fold Cros
plt.xlabel('Model', fontsize=14)
plt.ylabel('Accuracy', fontsize=14)
plt.xticks(rotation=45, ha='right')
plt.ylim([0.5, 1.0])
plt.tight layout()
plt.show()
# Create a grouped bar chart for all metrics
plt.figure(figsize=(15, 10))
sns.set_style("whitegrid")
# Create grouped bar plot
ax = sns.barplot(
   data=task3_results_df,
   x='Model',
   y='Mean',
   hue='Metric',
   order=task3 model order,
   palette='Set2'
plt.title(f'Task 3: Model Performance Metrics Comparison ({k folds}-Fold
plt.xlabel('Model', fontsize=14)
plt.ylabel('Score', fontsize=14)
plt.xticks(rotation=45, ha='right')
plt.legend(title='Metric', loc='lower right')
plt.ylim([0.5, 1.0])
plt.grid(True, linestyle='--', alpha=0.7)
```

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

/tmp/ipykernel_368067/1263538886.py:27: FutureWarning:

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





```
In [114... # Choose the best model from cross-validation for Task 3
best_task3_model_name = task3_model_order[0] # The model with highest ac
best_task3_model = task3_models[best_task3_model_name]
```

```
# Hyperparameter optimization for the best model on Task 3
print(f"Performing Grid Search for {best task3 model name} on Task 3...")
# Define the parameter grid for the best model
if best task3 model name == 'SVM':
    param grid = {
        'C': np.logspace(-3, 3, 7),
        'gamma': ['scale', 'auto'] + list(np.logspace(-3, 3, 5)),
        'kernel': ['rbf', 'linear']
elif best task3 model name == 'Random Forest':
    param grid = {
        'n estimators': [50, 100, 200],
        'max_depth': [None, 10, 20, 30],
        'min_samples_split': [2, 5, 10],
        'min_samples_leaf': [1, 2, 4]
elif best task3 model name.endswith('NB'):
    if best task3 model name == 'Gaussian NB':
        param_grid = {
            'var_smoothing': np.logspace(-10, 0, 11)
    else:
        param_grid = {
            'alpha': np.logspace(-3, 3, 7),
            'fit prior': [True, False]
else:
    # Default grid for other models
    param grid = {
        'C': np.logspace(-3, 3, 7)
# Create and fit GridSearchCV
grid search task3 = GridSearchCV(
    estimator=best_task3_model,
    param_grid=param_grid,
    cv=5,
    scoring='accuracy',
    n_{jobs=-1}
    verbose=1
)
grid_search_task3.fit(X_train_task3, y_train_task3)
# Display the best parameters and score
print(f"Best Parameters: {grid_search_task3.best_params_}")
print(f"Best Cross-Validation Score: {grid search task3.best score :.4f}"
# Evaluate on test set
best_model_task3 = grid_search_task3.best_estimator_
y_pred_task3 = best_model_task3.predict(X_test_task3)
accuracy_task3 = accuracy_score(y_test_task3, y_pred_task3)
print(f"Test Accuracy with Best Parameters: {accuracy_task3:.4f}")
print("\nClassification Report:")
print(classification_report(y_test_task3, y_pred_task3))
# Plot confusion matrix
plt.figure(figsize=(10, 8))
cm = confusion_matrix(y_test_task3, y_pred_task3)
```

Performing Grid Search for SVM on Task 3...
Fitting 5 folds for each of 98 candidates, totalling 490 fits
Best Parameters: {'C': np.float64(10.0), 'gamma': 'scale', 'kernel': 'rb
f'}

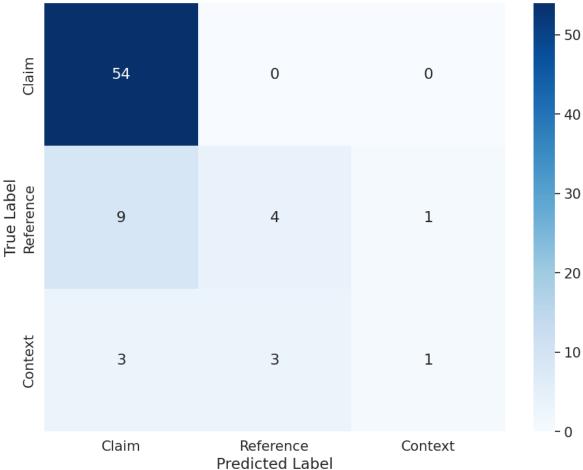
Best Cross-Validation Score: 0.7433

Test Accuracy with Best Parameters: 0.7867

Classification Report:

	precision	recall	f1-score	support
0 1 2	0.82 0.57 0.50	1.00 0.29 0.14	0.90 0.38 0.22	54 14 7
accuracy macro avg weighted avg	0.63 0.74	0.48 0.79	0.79 0.50 0.74	75 75 75





```
In [115... # Check for class imbalance in Task 3
print("Task 3 class distribution:")
print(y_task3.value_counts())
```

```
print(y task3.value counts(normalize=True).round(3) * 100, '%')
# Apply SMOTE for handling class imbalance in Task 3
smote = SMOTE(random state=42)
X train task3 balanced, y train task3 balanced = smote.fit resample(X tra
print("\nClass distribution after SMOTE balancing:")
print(pd.Series(y train task3 balanced).value counts())
# Train the best model with balanced data
best model balanced = clone(best model task3)
best model balanced fit(X train task3 balanced, y train task3 balanced)
# Evaluate on the test set
y pred balanced = best model balanced.predict(X test task3)
accuracy_balanced = accuracy_score(y_test_task3, y_pred_balanced)
print(f"\nTest Accuracy with Balanced Training Data: {accuracy balanced:.
print("\nClassification Report with Balanced Training Data:")
print(classification_report(y_test_task3, y_pred_balanced))
# Compare class-wise performance before and after balancing
original_report = classification_report(y_test_task3, y_pred_task3, outpu
balanced_report = classification_report(y_test_task3, y_pred_balanced, ou
# Create a DataFrame to compare per-class metrics
class comparison = []
for class_label in ['0', '1', '2']: # Claim, Reference, Context
    for metric in ['precision', 'recall', 'f1-score']:
        class comparison.append({
            'Class': ['Claim', 'Reference', 'Context'][int(class label)],
            'Metric': metric.capitalize(),
            'Original': original_report[class_label][metric],
            'Balanced': balanced_report[class_label][metric],
            'Difference': balanced_report[class_label][metric] - original
        })
class comp df = pd.DataFrame(class comparison)
# Plot class-wise performance comparison
plt.figure(figsize=(15, 10))
for i, cls in enumerate(['Claim', 'Reference', 'Context']):
    plt.subplot(1, 3, i+1)
    df_class = class_comp_df[class_comp_df['Class'] == cls]
    x = np.arange(len(df_class['Metric'].unique()))
    width = 0.35
    orig scores = df_class['Original'].values
    bal_scores = df_class['Balanced'].values
    plt.bar(x - width/2, orig scores, width, label='Original')
    plt.bar(x + width/2, bal scores, width, label='Balanced')
    plt.title(f'{cls} Performance')
    plt.xticks(x, df class['Metric'].unique())
   plt.ylim(0, 1)
    plt.legend()
    plt.grid(True, linestyle='--', alpha=0.7)
```

```
plt.suptitle('Impact of Class Balancing on Task 3 Performance by Class',
plt.tight_layout(rect=[0, 0, 1, 0.95])
plt.show()
```

Task 3 class distribution:

task3_label

0 263

1 79

2 33 Name: count, dtype: int64

task3_label 0 70.1

1 21.1 2 8.8

Name: proportion, dtype: float64 %

Class distribution after SMOTE balancing:

task3_label

0 209

1 209

2 209

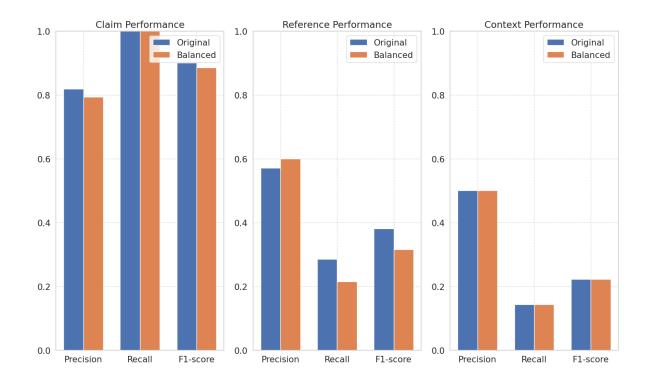
Name: count, dtype: int64

Test Accuracy with Balanced Training Data: 0.7733

Classification Report with Balanced Training Data:

	precision	recall	fl-score	support
0	0.79	1.00	0.89	54
1	0.60	0.21	0.32	14
2	0.50	0.14	0.22	7
accuracy			0.77	75
macro avg	0.63	0.45	0.47	75
weighted avg	0.73	0.77	0.72	75

Impact of Class Balancing on Task 3 Performance by Class



```
In [116... # Visualize confusion matrices for balanced vs unbalanced
          plt.figure(figsize=(16, 7))
          # Original model confusion matrix
          plt.subplot(1, 2, 1)
          cm_orig = confusion_matrix(y_test_task3, y_pred_task3)
          sns.heatmap(cm_orig, annot=True, fmt='d', cmap='Blues',
                      xticklabels=['Claim', 'Reference', 'Context'],
                       yticklabels=['Claim', 'Reference', 'Context'])
          plt.title('Confusion Matrix - Original Model')
          plt.ylabel('True Label')
          plt.xlabel('Predicted Label')
          # Balanced model confusion matrix
          plt.subplot(1, 2, 2)
          cm_balanced = confusion_matrix(y_test_task3, y_pred_balanced)
          sns.heatmap(cm_balanced, annot=True, fmt='d', cmap='Blues',
                       xticklabels=['Claim', 'Reference', 'Context'],
yticklabels=['Claim', 'Reference', 'Context'])
          plt.title('Confusion Matrix - Balanced Model')
          plt.ylabel('True Label')
          plt.xlabel('Predicted Label')
          plt.tight_layout()
          plt.show()
```

