```
In [32]: %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
e-packages (from pandas) (2.2.5)
Requirement already satisfied: python-dateutil>=2.8.2 in ./.venv/lib/pytho
n3.10/site-packages (from pandas) (2.9.0.post0)
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te-packages (from pandas) (2025.2)
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kages (from python-dateutil>=2.8.2->pandas) (1.17.0)
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ackages (3.10.1)
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e-packages (from matplotlib) (0.12.1)
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O/site-packages (from matplotlib) (1.4.8)
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Note: you may need to restart the kernel to use updated packages.
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ages (0.13.2)
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Requirement already satisfied: pandas>=1.2 in ./.venv/lib/python3.10/site-
packages (from seaborn) (2.2.3)
Requirement already satisfied: matplotlib!=3.6.1,>=3.4 in ./.venv/lib/pyth
on3.10/site-packages (from seaborn) (3.10.1)
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site-packages (from matplotlib!=3.6.1,>=3.4->seaborn) (1.3.2)
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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)
Requirement already satisfied: python-dateutil>=2.7 in ./.venv/lib/python
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
        e-packages (from pandas>=1.2->seaborn) (2025.2)
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        te-packages (from pandas>=1.2->seaborn) (2025.2)
        Requirement already satisfied: six>=1.5 in ./.venv/lib/python3.10/site-pac
        kages (from python-dateutil>=2.7->matplotlib!=3.6.1,>=3.4->seaborn) (1.1
        7.0)
        Note: you may need to restart the kernel to use updated packages.
        Requirement already satisfied: scikit-learn in ./.venv/lib/python3.10/sit
        e-packages (1.6.1)
        Requirement already satisfied: numpy>=1.19.5 in ./.venv/lib/python3.10/sit
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        e-packages (from scikit-learn) (1.4.2)
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        Note: you may need to restart the kernel to use updated packages.
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        s(3.9.1)
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        es (from nltk) (8.1.8)
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        ges (from nltk) (1.4.2)
        Requirement already satisfied: regex>=2021.8.3 in ./.venv/lib/python3.10/s
        ite-packages (from nltk) (2024.11.6)
        Requirement already satisfied: tqdm in ./.venv/lib/python3.10/site-package
        s (from nltk) (4.67.1)
        Note: you may need to restart the kernel to use updated packages.
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        e-packages (from wordcloud) (2.2.5)
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        ges (from wordcloud) (11.2.1)
        Requirement already satisfied: matplotlib in ./.venv/lib/python3.10/site-p
        ackages (from wordcloud) (3.10.1)
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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)
        Requirement already satisfied: python-dateutil>=2.7 in ./.venv/lib/python
        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 [33]: 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[33]:	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 [34]:	<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 [35]:	<pre># Compare different feature extraction methods from sklearn.feature_extraction.text import CountVectorizer from sklearn.feature_selection import SelectKBest, chi2, mutual_info_clas</pre>						
	<pre># Create label for Task 1 y_task1 = df['science_related']</pre>						
	<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,</pre>						
	<pre>ngram_range=(1, 2)) X_tfidf = tfidf_vec.fit_transform(df['text'])</pre>						
	# 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)
```

warnings.warn(msg, UserWarning)

/home/hurel/Documents/repo/projet-ml/.venv/lib/python3.10/site-packages/sk learn/metrics/cluster/_supervised.py:59: UserWarning: Clustering metrics e xpects discrete values but received continuous values for label, and binar y values for target

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/home/hurel/Documents/repo/projet-ml/.venv/lib/python3.10/site-packages/sk learn/metrics/cluster/_supervised.py:59: UserWarning: Clustering metrics e xpects discrete values but received continuous values for label, and binar y values for target

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/home/hurel/Documents/repo/projet-ml/.venv/lib/python3.10/site-packages/sk learn/metrics/cluster/_supervised.py:59: UserWarning: Clustering metrics e xpects discrete values but received continuous values for label, and binary values for target

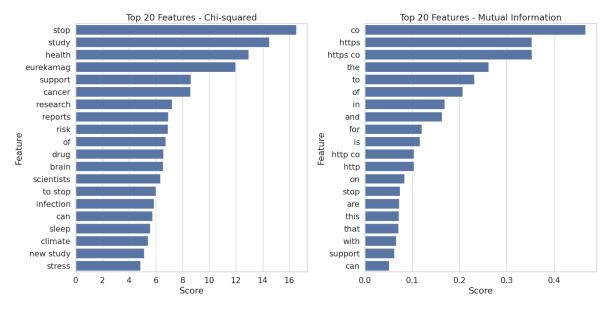
warnings.warn(msg, UserWarning)

/home/hurel/Documents/repo/projet-ml/.venv/lib/python3.10/site-packages/sk learn/metrics/cluster/_supervised.py:59: UserWarning: Clustering metrics e xpects discrete values but received continuous values for label, and binar y values for target

warnings.warn(msg, UserWarning)

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

Features after Mutual Information selection: (1140, 100)



```
In [36]: # 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['sdf_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 [37]: 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: Scier	ice Related	Classifica	tion		
Model: Gaussian NB Accuracy: 0.6711					
	precision	recall	f1-score	support	
0	0.79	0.66	0.72	146	
1	0.53	0.70	0.60	82	
accuracy			0.67	228	
macro avg weighted avg	0.66 0.70	0.68 0.67	0.66 0.68	228 228	
weighted avg	0.70	0.07	0.00	220	
Model: Multir Accuracy: 0.7					
Accuracy: 0.7	precision	recall	f1-score	support	
0	0.74	0.96	0.83	146	
1	0.84	0.39	0.53	82	
accuracy			0.75	228	
macro avg	0.79	0.67 0.75	0.68	228	
weighted avg	0.77	0.75	0.73	228	
Model: Complement NB Accuracy: 0.7851					
	precision	recall	f1-score	support	
0	0.84	0.82	0.83	146	
1	0.69	0.73	0.71	82	
accuracy			0.79	228	
macro avg weighted avg	0.77 0.79	0.77 0.79	0.77 0.79	228 228	
weighted avg	0.79	0.79	0.79	220	
Model: Bernoulli NB Accuracy: 0.7895					
	precision	recall	f1-score	support	
0	0.80	0.90	0.85	146	
1	0.77	0.59	0.67	82	
accuracy			0.79	228	
macro avg	0.78	0.74	0.76	228	
weighted avg	0.79	0.79	0.78	228	
Model: KNN					
Accuracy: 0.7	105 precision	recall	f1-score	support	
•					
0 1	0.80 0.58	0.73 0.68	0.76 0.63	146 82	
_				- -	

accuracy			0.71	228
macro avg	0.69	0.70	0.70	228
weighted avg	0.72	0.71	0.71	228

Model: SVM

Accuracy: 0.7895

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

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: 0.7544 precision recall f1-score support 0.74 0.95 0.83 146 0 1 0.81 0.41 0.55 82 0.75 228 accuracy macro avg 0.78 0.68 weighted avg 0.77 0.75 0.69 228 0.75 0.73 228

```
In [38]: 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)

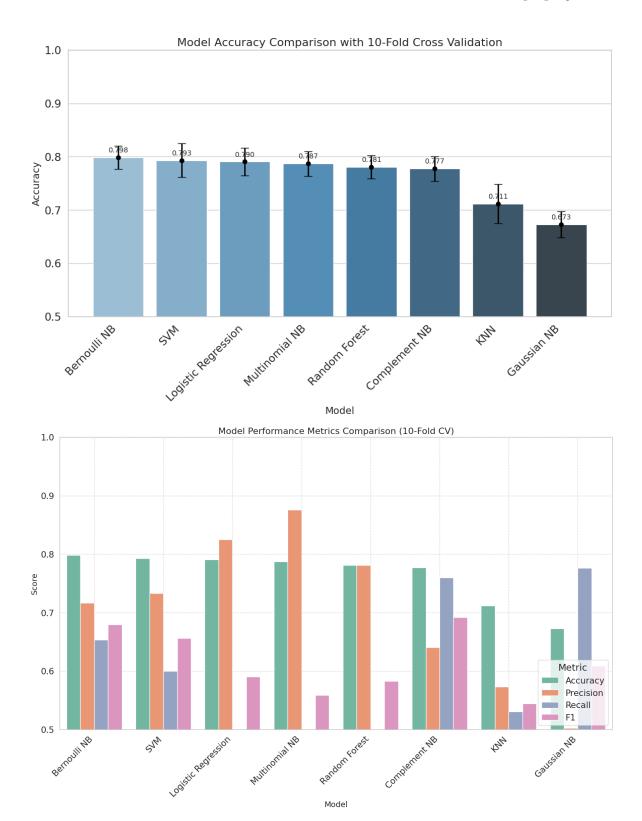
# Dictionary to store results
    cv_results = {}

# 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 0.0218
 Average: F1 Score=0.5826 \pm 0.0543
/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 [39]: from sklearn.model_selection import GridSearchCV

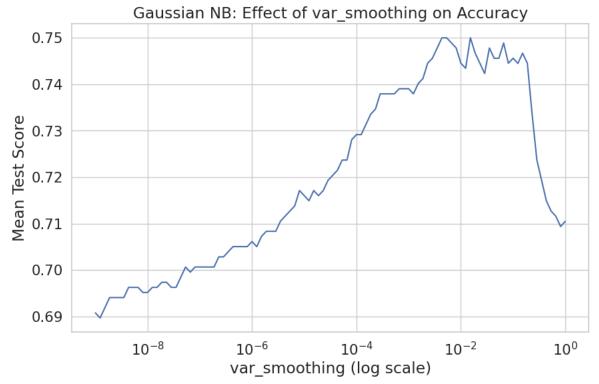
for model_name, model in nb_models.items():
```

```
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...

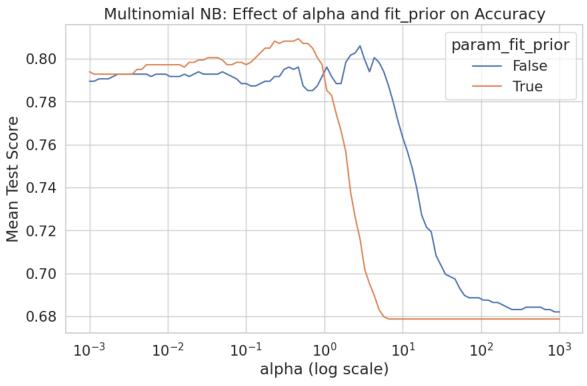
Fitting 10 folds for each of 200 candidates, totalling 2000 fits

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

ie}

Best Cross-Validation Score: 0.8093

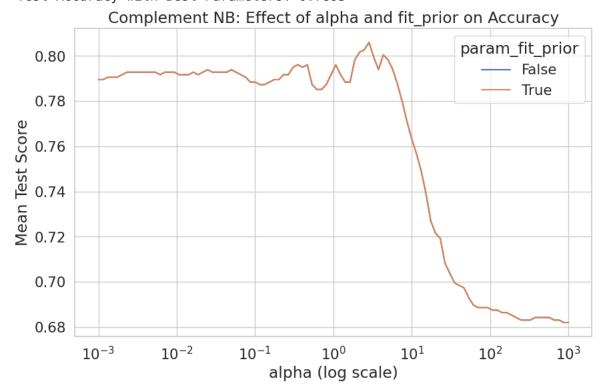
Test Accuracy with Best Parameters: 0.7982



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
e}

Best Cross-Validation Score: 0.8060

Test Accuracy with Best Parameters: 0.7895



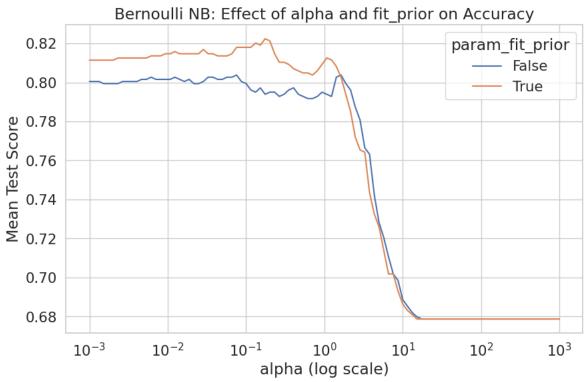
Performing Grid Search for Bernoulli NB...

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

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

Best Cross-Validation Score: 0.8224

Test Accuracy with Best Parameters: 0.8070



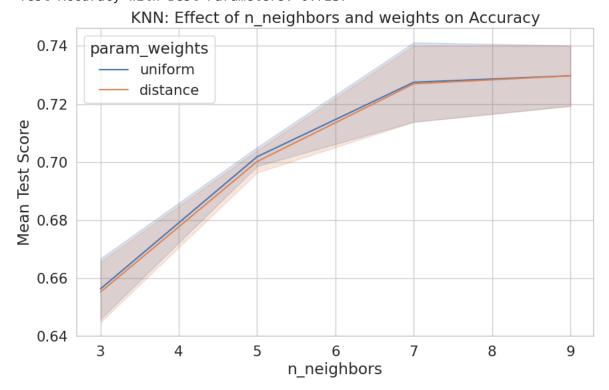
Performing Grid Search for KNN...

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...

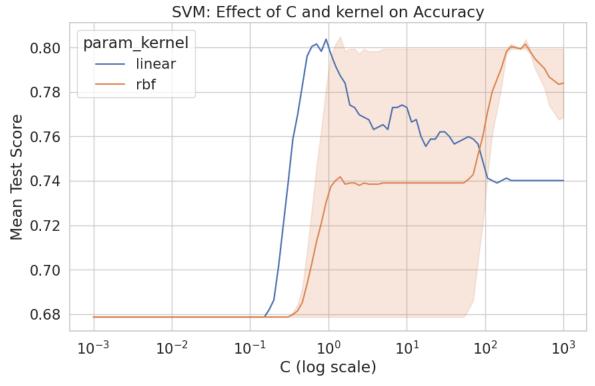
Fitting 10 folds for each of 400 candidates, totalling 4000 fits

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

```
/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
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/home/hurel/Documents/repo/projet-ml/.venv/lib/python3.10/site-packages/sk
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  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
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  warnings.warn(
/home/hurel/Documents/repo/projet-ml/.venv/lib/python3.10/site-packages/sk
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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
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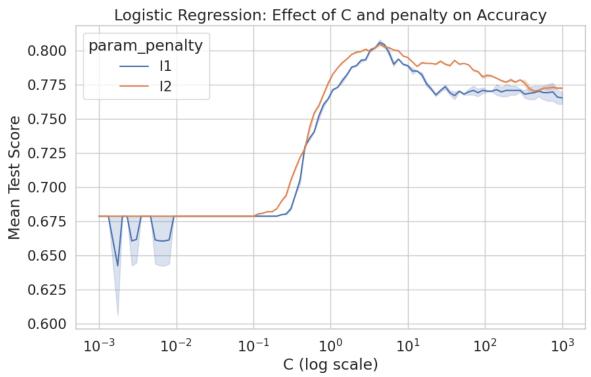
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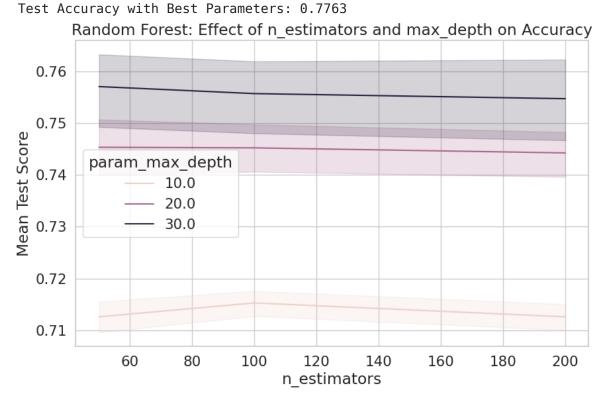
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Best Parameters: {'C': np.float64(4.328761281083062), 'penalty': 'l1', 'so
lver': 'saga'}
Best Cross-Validation Score: 0.8081
Test Accuracy with Best Parameters: 0.7939
```



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': 2, 'n_estimators': 200}
Best Cross-Validation Score: 0.7829



Training comparaison with best parameter

```
In [40]: 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)
```

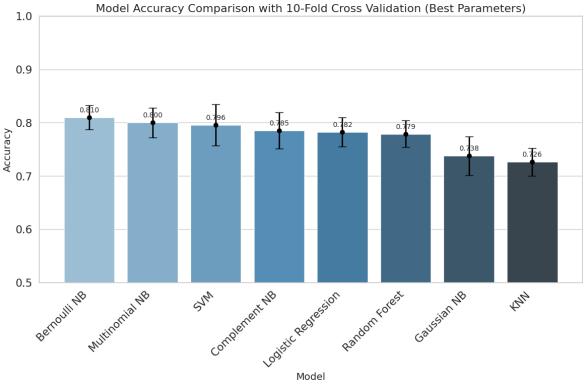
```
fold recall.append(rec)
        fold f1.append(f1)
    # Store average metrics and standard deviations
   best cv results[model name] = {
        'accuracy': {
            'mean': np.mean(fold accuracy),
            'std': np.std(fold accuracy)
        },
        'precision': {
            'mean': 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
```

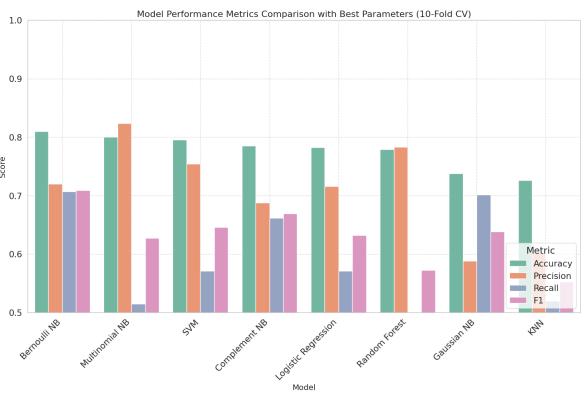
```
# Create plots
plt.figure(figsize=(12, 8))
sns.set style("whitegrid")
# Create bar plot for accuracy
ax = sns.barplot(
   data=best results df[best results df['Metric'] == 'Accuracy'],
   x='Model',
   y='Mean',
   order=best model order,
   palette='Blues d'
# Add error bars
for i, model in enumerate(best_model_order):
    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 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=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 Optimizatio
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 0.0212
  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=0.8542 \pm 0.0250
  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=0.8426 \pm 0.0310
  Average: Accuracy=0.7789 \pm 0.0251
  Average: F1 Score=0.5725 \pm 0.0594
/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(
```

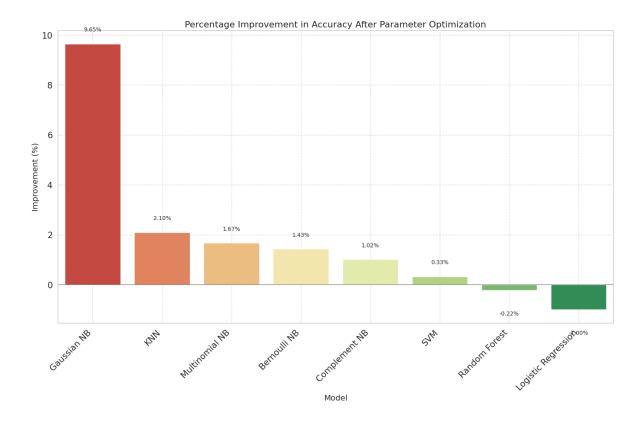




/tmp/ipykernel_368067/2330540877.py:224: FutureWarning:

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(

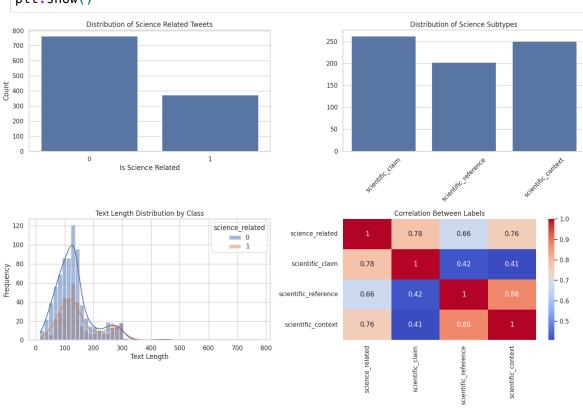


Preprocessing

```
In [41]: 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 [42]: 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)
```

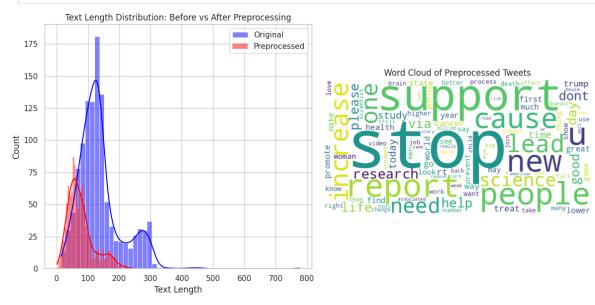
```
# Remove special characters and numbers
              text = re.sub(r'[^a-zA-Z\s]', '', text)
              # Tokenize
              tokens = nltk.word tokenize(text)
              # Remove stopwords
              stop words = set(stopwords.words('english'))
              tokens = [word for word 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[42]:
                                        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
             Does daily routine help prevent problems
                                                   daily routine help prevent problem bipolar
                                          with ...
                                                                                 dis...
In [43]: # 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 [44]: from sklearn.feature_extraction.text import TfidfVectorizer

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

```
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)
```

tfidf processed = TfidfVectorizer(max features=5000)

warnings.warn(msg, UserWarning)

/home/hurel/Documents/repo/projet-ml/.venv/lib/python3.10/site-packages/sk learn/metrics/cluster/_supervised.py:59: UserWarning: Clustering metrics e xpects discrete values but received continuous values for label, and binar y values for target

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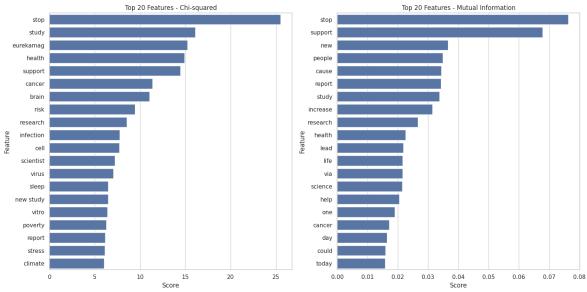
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Features after Chi-squared selection: (1140, 100)

Features after Mutual Information selection: (1140, 100)



```
In [46]: 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
             '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)

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: Scien	ce Related	Classifica	tion	
Model: Gaussi Accuracy: 0.6				
	precision	recall	f1-score	support
0 1	0.80 0.54	0.66 0.71	0.73 0.61	146 82
accuracy macro avg weighted avg	0.67 0.71	0.69 0.68	0.68 0.67 0.69	228 228 228
Model: Multin				
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 macro avg weighted avg	0.73 0.74	0.68 0.74	0.74 0.69 0.73	228 228 228
Model: Comple Accuracy: 0.7		recall	f1-score	support
0	0.82 0.64	0.78 0.70	0.80 0.67	146 82
accuracy macro avg weighted avg	0.73 0.76	0.74 0.75	0.75 0.73 0.75	228 228 228
Model: Bernou Accuracy: 0.7				
Accuracy: 0.7	precision	recall	f1-score	support
0 1	0.78 0.69	0.86 0.56	0.81 0.62	146 82
accuracy macro avg weighted avg	0.73 0.74	0.71 0.75	0.75 0.72 0.74	228 228 228
Model: KNN Accuracy: 0.4			61	
	precision	recall	f1-score	support
0 1	0.91 0.40	0.20 0.96	0.33 0.57	146 82

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accuracy macro avg weighted avg	0.65 0.73	0.58 0.47	0.47 0.45 0.41	228 228 228
Model: SVM Accuracy: 0.7	763 precision	recall	fl-score	support
0 1	0.78 0.75	0.90 0.56	0.84 0.64	146 82
accuracy macro avg weighted avg	0.77 0.77	0.73 0.78	0.78 0.74 0.77	228 228 228
Model: Logist Accuracy: 0.7	•	.on		

	precision	recall	f1-score	support
0	0.79	0.88	0.83	146
1	0.72	0.57	0.64	82
accuracy	0.7.	0.0.	0.77	228
macro avg	0.75	0.72	0.73	228
weighted avg	0.76	0.77	0.76	228

Model: Random Forest

Accuracy:	0.7	precision	recall	fl-score	support
	0	0.82	0.80	0.81	146
	1	0.66	0.70	0.68	82
accur	асу			0.76	228
macro	avg	0.74	0.75	0.75	228
weighted	avg	0.77	0.76	0.76	228

```
In [47]: # Using preprocessed text for feature extraction
         from sklearn.model_selection import cross_val_score, StratifiedKFold
         from sklearn.metrics import accuracy_score, precision_score, recall_score
         # Create TF-IDF features from preprocessed text
         tfidf_vec = TfidfVectorizer(max_features=5000,
                                    min df=5,
                                    max df=0.8,
                                    ngram range=(1, 2))
         X_tfidf_processed = tfidf_vec.fit_transform(df['processed_text'])
         # Feature selection using Chi-squared
         selector chi2 = SelectKBest(chi2, k=100)
```

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```
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:
```

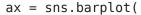
```
pass
        fold accuracy.append(acc)
        fold precision.append(prec)
        fold recall.append(rec)
        fold f1.append(f1)
    # Store average metrics and standard deviations
   best cv results[model name] = {
        'accuracy': {
            'mean': np.mean(fold accuracy),
            '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']]
```

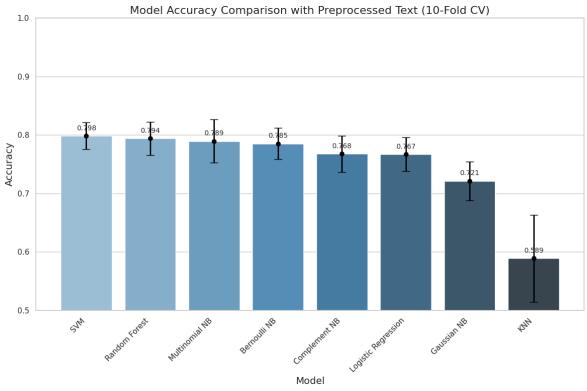
```
})], ignore_index=True)
# Sort models by accuracy
best model order = best results df[best results df['Metric'] == 'Accuracy'
# Create plots
plt.figure(figsize=(12, 8))
sns.set style("whitegrid")
# Create bar plot for accuracy
ax = sns.barplot(
   data=best results df[best results df['Metric'] == '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)
```

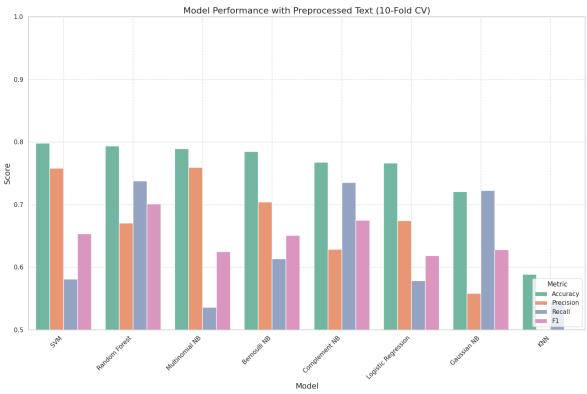
```
plt.ylabel('Score', fontsize=14)
 plt.xticks(rotation=45, ha='right')
 plt.legend(title='Metric', loc='lower right')
 plt.ylim([0.5, 1.0])
 plt.grid(True, linestyle='--', alpha=0.7)
 plt.tight layout()
 plt.show()
Performing 10-fold cross-validation for Gaussian NB with best parameters o
n preprocessed 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=0.8346 \pm 0.0317
  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=0.8594 \pm 0.0294
  Average: Accuracy=0.7939 \pm 0.0286
  Average: F1 Score=0.7011 \pm 0.0459
```

/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

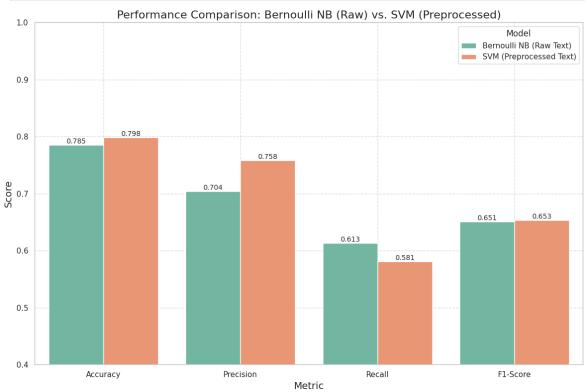
```
# Compare performance of SVM with preprocessed data and Bernoulli NB with
import matplotlib.pyplot as plt
# Create a DataFrame for comparison
comparison data = [
    {
        'Model': 'Bernoulli NB (Raw Text)',
        'Accuracy': best cv results['Bernoulli NB']['accuracy']['mean'],
        'Precision': 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

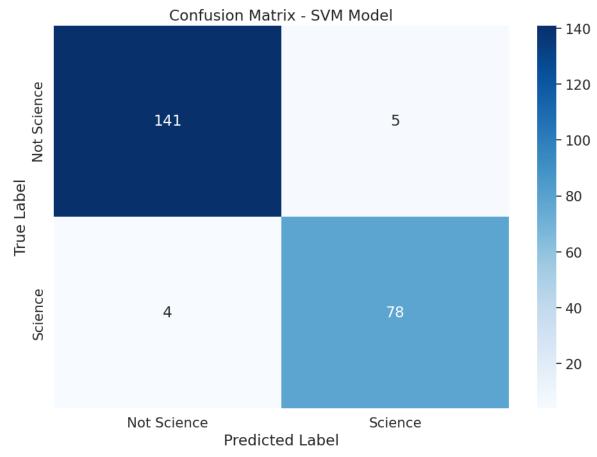
Difference in Accuracy: 0.013 Difference in F1-Score: 0.002

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

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

```
In [49]: best_model = best_models['SVM']
In [50]: from sklearn.metrics import confusion_matrix, classification_report import seaborn as sns import numpy as np
import matplotlib.pyplot as plt
```

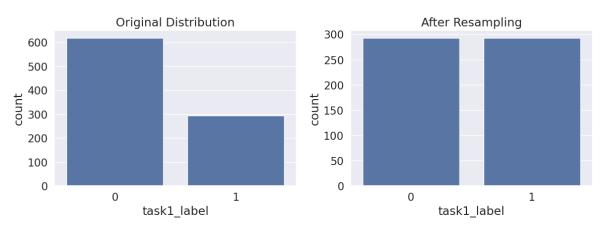
```
# Get predictions from the best model on the test set
y pred = best model.predict(X test task1)
# Calculate the confusion matrix
cm = confusion matrix(y test task1, y pred)
# Create a prettier visualization of the confusion matrix
plt.figure(figsize=(10, 8))
sns.set(font_scale=1.4)
ax = sns.heatmap(cm, 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 [51]: %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 [52]: # 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 [53]: # 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
```

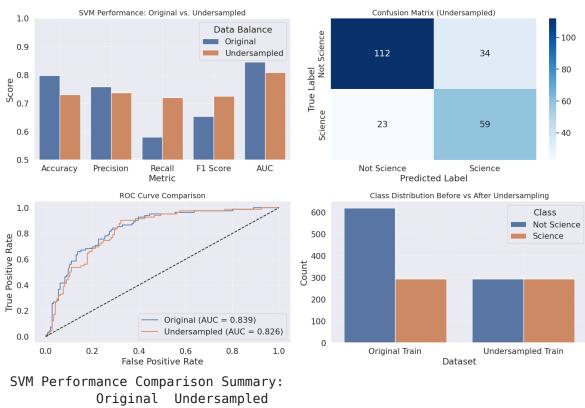
Effect of Undersampling on Task 1



```
In [54]: # 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'))
```



Metric
Accuracy 0.798246 0.730479
Precision 0.758331 0.737308
Recall 0.581081 0.719885
F1 Score 0.653448 0.725460
AUC 0.846290 0.809265

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 -4.374963

```
In [56]: # 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

Task 2: Scien					
Model: Gaussian NB Accuracy: 0.8533					
	precision	recall	f1-score	support	
0 1	0.30 0.94	0.43 0.90	0.35 0.92	7 68	
accuracy			0.85	75	
macro avg weighted avg	0.62 0.88	0.66 0.85	0.64 0.86	75 75	
Model: Multin 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			0.91	75	
macro avg weighted avg	0.45 0.82	0.50 0.91	0.48 0.86	75 75	
Model: Comple Accuracy: 0.7					
	precision	recall	f1-score	support	
0 1	0.18 0.94	0.57 0.74	0.28 0.83	7 68	
accuracy	0.50	0.05	0.72	75	
macro avg weighted avg	0.56 0.87	0.65 0.72	0.55 0.78	75 75	
Model: Bernou Accuracy: 0.8					
	precision	recall	f1-score	support	
0 1	0.00 0.90	0.00 0.94	0.00 0.92	7 68	
accuracy			0.85	75	
macro avg weighted avg	0.45 0.82	0.47 0.85	0.46 0.83	75 75	
Model: KNN Accuracy: 0.9	067				
curucy: 019	precision	recall	f1-score	support	

	precision	recall	f1-score	support
0	0.00	0.00	0.00	7
1	0.91	1.00	0.95	68

accuracy			0.91	75
macro avg	0.45	0.50	0.48	75
weighted avg	0.82	0.91	0.86	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
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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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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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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))
```

Model: SVM

Accuracy: 0.9200

	precision	recall	f1-score	support
Θ	1.00	0.14	0.25	7
1	0.92	1.00	0.96	68
accuracy			0.92	75
macro avg weighted avg	0.96 0.93	0.57 0.92	0.60 0.89	75 75

Model: Logistic Regression

Accuracy: 0.9200

	precision	recall	f1-score	support
0	1.00	0.14	0.25	7
1	0.92	1.00	0.96	68
accuracy			0.92	75
macro avg weighted avg	0.96 0.93	0.57 0.92	0.60 0.89	75 75

Model: Random Forest

Accuracy: 0.9200

	precision	recall	f1-score	support
0	1.00	0.14	0.25	7
1	0.92	1.00	0.96	68
accuracy			0.92	75
macro avg	0.96	0.57	0.60	75
weighted avg	0.93	0.92	0.89	75

```
In [57]: # 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)
         # 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)
    },
    'f1': {
        '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
```

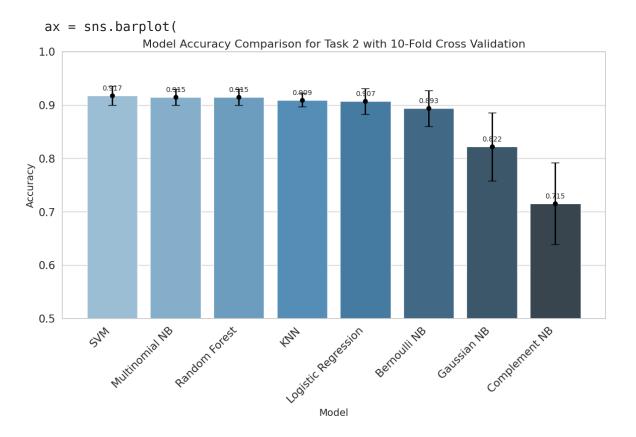
```
print(f" Average: Accuracy={task2 cv results[model name]['accuracy']
             print(f" Average: F1 Score={task2 cv results[model name]['f1']['mean
             print()
        Task 2: Cross-Validation
        Performing 10-fold cross-validation for Gaussian NB...
          Average: AUC=0.5660 \pm 0.1187
          Average: Accuracy=0.8217 \pm 0.0635
          Average: F1 Score=0.8988 \pm 0.0375
        Performing 10-fold cross-validation for Multinomial NB...
          Average: AUC=0.7806 \pm 0.0932
          Average: Accuracy=0.9148 \pm 0.0152
          Average: F1 Score=0.9554 \pm 0.0081
        Performing 10-fold cross-validation for Complement NB...
          Average: AUC=0.7468 \pm 0.0979
          Average: Accuracy=0.7151 \pm 0.0768
          Average: F1 Score=0.8205 \pm 0.0532
        Performing 10-fold cross-validation for Bernoulli NB...
          Average: AUC=0.7860 \pm 0.0991
          Average: Accuracy=0.8935 \pm 0.0332
          Average: F1 Score=0.9423 \pm 0.0184
        Performing 10-fold cross-validation for KNN...
          Average: AUC=0.5324 \pm 0.0660
          Average: Accuracy=0.9094 \pm 0.0126
          Average: F1 Score=0.9525 \pm 0.0069
        Performing 10-fold cross-validation for SVM...
          Average: AUC=0.7151 \pm 0.1219
          Average: Accuracy=0.9175 \pm 0.0179
          Average: F1 Score=0.9567 \pm 0.0095
        Performing 10-fold cross-validation for Logistic Regression...
        /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(
          Average: AUC=0.7019 \pm 0.1686
          Average: Accuracy=0.9068 \pm 0.0242
          Average: F1 Score=0.9503 \pm 0.0131
        Performing 10-fold cross-validation for Random Forest...
          Average: AUC=0.7260 \pm 0.1391
          Average: Accuracy=0.9148 \pm 0.0152
          Average: F1 Score=0.9554 \pm 0.0081
In [58]: # Visualize Task 2 cross-validation results
         # Create DataFrame for visualization
         task2 results df = pd.DataFrame({
              'Model': [],
              'Metric': [],
              'Mean': [],
              'Std': []
         })
```

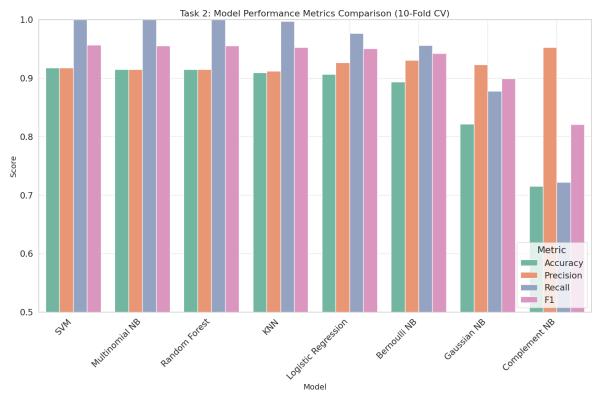
```
for model name in task2 cv results:
    for metric in ['accuracy', 'precision', 'recall', 'f1']:
        task2 results df = pd.concat([task2 results df, pd.DataFrame({
            'Model': [model name],
            'Metric': [metric.capitalize()],
            'Mean': [task2 cv results[model name][metric]['mean']],
            '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:

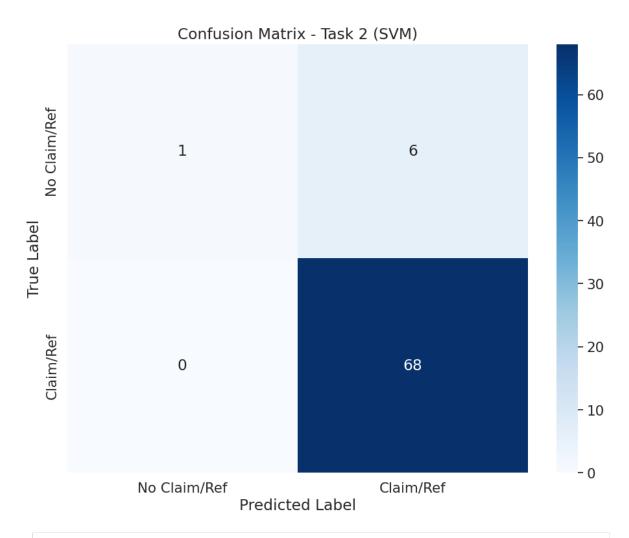
Passing `palette` without assigning `hue` is deprecated and will be remove d in v0.14.0. Assign the `x` variable to `hue` and set `legend=False` for the same effect.





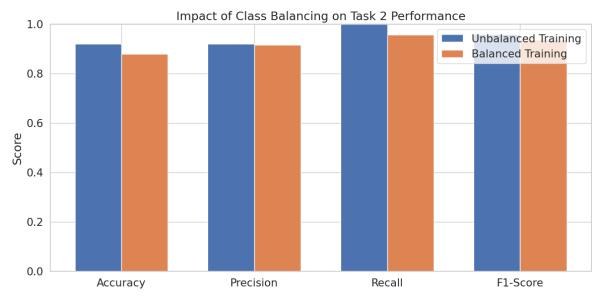
```
# Hyperparameter optimization for the best model on Task 2
# Let's assume SVM was the best model (update based on your results)
best_task2_model_name = 'SVM' # Replace with the actual best model from
best_task2_model = task2_models[best_task2_model_name]
print(f"Performing Grid Search for {best_task2_model_name} 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()
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
                                                   75
    accuracy
                   0.96
                            0.57
                                       0.60
                                                   75
   macro avg
weighted avg
                   0.93
                             0.92
                                       0.89
                                                   75
```



```
In [64]: # 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_tra
         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)
 1
 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
     342
1
     33
Name: count, dtype: int64
task2 label
   91.2
1
      8.8
Name: proportion, dtype: float64 %
Class distribution after SMOTE balancing:
task2 label
     274
1
     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
                   0.92
                             0.96
                                       0.94
                                                    68
                                                   75
                                       0.88
    accuracy
                  0.58
                             0.55
                                       0.56
                                                   75
   macro avg
weighted avg
                  0.85
                             0.88
                                       0.86
                                                   75
```



```
In [65]: # 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

======================================	an NB	======	:=======	=====
Accuracy: 0.76			6.3	
	precision	recall	f1-score	support
0	0.86	0.89	0.87	54
1	0.67	0.43	0.52	14
2	0.30	0.43	0.35	7
accuracy			0.76	75
macro avg	0.61	0.58	0.58	75 75
weighted avg	0.77	0.76	0.76	75
Model: Multino				
Accuracy. 0.70	precision	recall	f1-score	support
	p. 002020		000.0	0 a p p 0 . 1
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	0.50	0.40	0.40	75
weighted avg	0.69	0.76	0.68	75
Model: Complem				
	precision	recall	f1-score	support
0	0.78	0.72	0.75	54
1	0.31	0.29	0.30	14
2	0.17	0.29	0.21	7
accuracy			0.60	75
macro avg	0.42	0.43	0.42	75
weighted avg	0.63	0.60	0.61	75
Madal Dansaul	14 ND			
Model: Bernoul Accuracy: 0.70				
need dey! of re	precision	recall	f1-score	support
Θ	0.78	0.91	0.84	54
1	0.43	0.21	0.29	14
2	0.20	0.14	0.17	7
accuracy			0.71	75
macro avg	0.47	0.42	0.43	75
weighted avg	0.66	0.71	0.67	75

Model: KNN

Accuracy: 0.7200

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	precision	recall	f1-score	support
0	0.73	0.96	0.83	54
1	0.50	0.14	0.22	14
2	0.00	0.00	0.00	7
accuracy			0.72	75
macro avg	0.41	0.37	0.35	75
weighted avg	0.62	0.72	0.64	75
Model: SVM Accuracy: 0.7				
	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 75
WCIGHTCU UVG	0.75	0.77	0.72	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
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s ill-defined and being set to 0.0 in labels with no predicted samples. Us
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  _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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learn/metrics/_classification.py:1565: UndefinedMetricWarning: Precision i
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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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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: 0.6800

,	precision	recall	f1-score	support
0	0.84	0.80	0.82	54
1	0.32	0.50	0.39	14
2	0.50	0.14	0.22	7
accuracy			0.68	75
macro avg	0.55	0.48	0.48	75
weighted avg	0.71	0.68	0.68	75

```
In [66]: # 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 ± 0.0701 Average: F1 Score= 0.4518 ± 0.0643 Performing 10-fold cross-validation for Multinomial NB... Average: Accuracy= 0.7280 ± 0.0599 Average: F1 Score= 0.4004 ± 0.0922 Performing 10-fold cross-validation for Complement NB... Average: Accuracy= 0.6428 ± 0.0595 Average: F1 Score= 0.4886 ± 0.0616 Performing 10-fold cross-validation for Bernoulli NB... Average: Accuracy= 0.7042 ± 0.0498 Average: F1 Score= 0.4997 ± 0.0956 Performing 10-fold cross-validation for KNN... Average: Accuracy= 0.2983 ± 0.0671 Average: F1 Score= 0.1982 ± 0.0542 Performing 10-fold cross-validation for SVM... Average: Accuracy= 0.7467 ± 0.0379 Average: F1 Score= 0.4304 ± 0.0634 Performing 10-fold cross-validation for Logistic Regression... /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(Average: Accuracy= 0.6962 ± 0.0395 Average: F1 Score= 0.4445 ± 0.0834 Performing 10-fold cross-validation for Random Forest... Average: Accuracy= 0.7011 ± 0.0700 Average: F1 Score= 0.5048 ± 0.1258 In [67]: # 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

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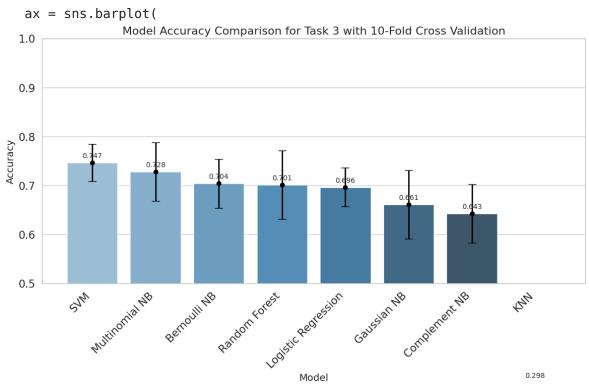
task3 model order = task3 results df[task3 results df['Metric'] == 'Accur

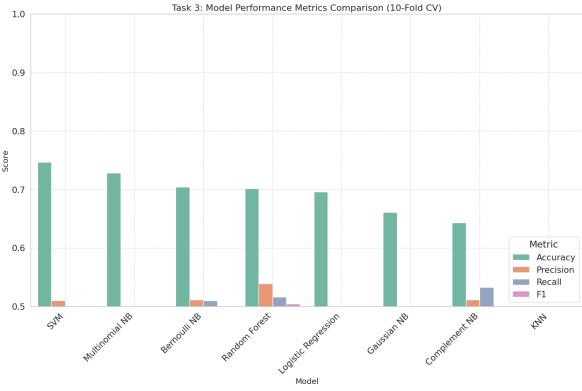
```
# Create accuracy bar plot
plt.figure(figsize=(12, 8))
sns.set_style("whitegrid")
# Create bar plot for accuracy
ax = sns.barplot(
   data=task3 results df[task3 results df['Metric'] == 'Accuracy'],
   x='Model',
   y='Mean',
   order=task3 model order,
   palette='Blues d'
)
# Add error bars
for 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 [68]: # 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'}

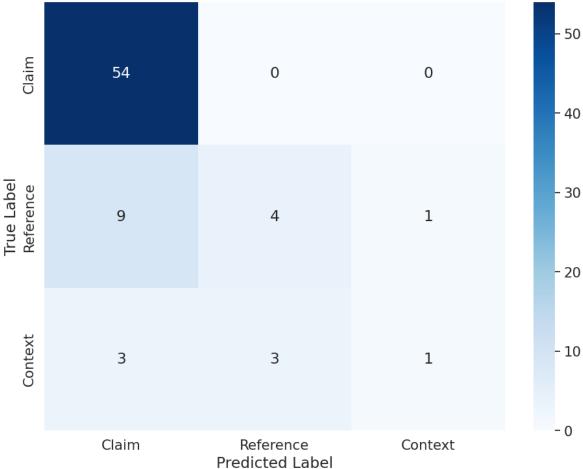
Best Cross-Validation Score: 0.7433

Test Accuracy with Best Parameters: 0.7867

Classification Report:

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





```
In [69]: # 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

8.8

2

Name: proportion, dtype: float64 %

Class distribution after SMOTE balancing:

task3_label 0 209 1 209 2 209

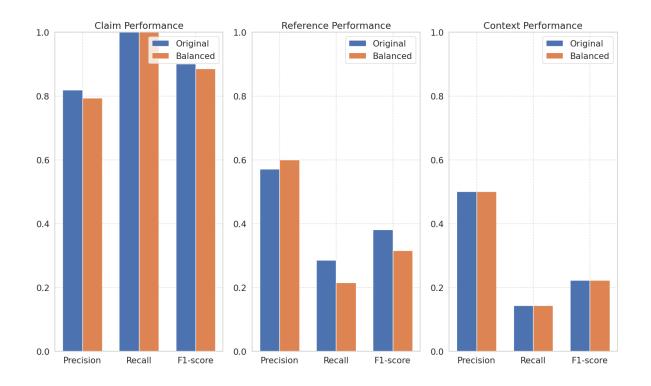
Name: count, dtype: int64

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 [70]: # 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()
```

