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import pandas as pd
import numpy as np
import re
from bs4 import BeautifulSoup
from sklearn.model selection import train test split, KFold
from sklearn.preprocessing import LabelEncoder
from lime.lime text import LimeTextExplainer
import matplotlib.pyplot as plt
import tensorflow as tf
from tensorflow.keras.preprocessing.text import Tokenizer
from tensorflow.keras.preprocessing.sequence import pad sequences
from tensorflow.keras.models import Sequential
from tensorflow.keras.layers import Embedding, Conv1D, MaxPooling1D, Flatten, Dropout, Dense
import html
# Load and preprocess your data
df = pd.read csv('/kaggle/input/datset8/requirment8.csv', encoding='latin1')
# Preprocessing function
def clean text(text):
  text = BeautifulSoup(text, "lxml").get text() # Using BeautifulSoup to remove HTML content
  text = text.lower()
  text = re.sub(r'[^a-zA-Z0-9\s]', ", text)
  return text
df['Base Reviews'] = df['Base Reviews'].astype(str).apply(clean text)
# Encode the labels (categories)
label encoder = LabelEncoder()
df['category'] = label_encoder.fit_transform(df['category'])
class names = label encoder.classes .tolist()
# Tokenizer and Pad Sequences for CNN
tokenizer = Tokenizer(num words=1000)
tokenizer.fit on texts(df['Base Reviews'])
X = tokenizer.texts to sequences(df['Base Reviews'])
X = pad sequences(X, maxlen=100)
vocab size = len(tokenizer.word index) + 1
# Mapping categories to numerical labels
y = df['category'].values
# Split the data
X train, X test, y train, y test = train_test_split(X, y, test_size=0.2, random_state=123)
# Convert labels to categorical for classification
y train cat = tf.keras.utils.to categorical(y train, num classes=len(class names))
y test cat = tf.keras.utils.to categorical(y test, num classes=len(class names))
# Define the CNN model with learning rate 0.001
def create cnn model():
  model = Sequential([
    Embedding(input dim=vocab size, output dim=100, input length=100),
    Conv1D(128, 5, activation='relu'),
    MaxPooling1D(pool size=2),
    Flatten(),
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Dropout(0.5),
    Dense(len(class names), activation='softmax')
  optimizer = tf.keras.optimizers.Adam(learning rate=0.001)
  model.compile(optimizer=optimizer, loss='categorical crossentropy', metrics=['accuracy'])
  return model
# Implementing K-Fold Cross Validation with K=10
kfold = KFold(n splits=10, shuffle=True, random state=1)
for fold, (train idx, val idx) in enumerate(kfold.split(X train)):
  print(f"Training fold {fold + 1}")
  # Split the training data into training and validation sets
  X fold train, X fold val = X train[train idx], X train[val idx]
  y fold train, y fold val = y train cat[train idx], y train cat[val idx]
  # Create and train the CNN model for each fold
  cnn model = create cnn model()
  cnn model.fit(X fold train, y fold train, epochs=10, batch size=32, validation data=(X fold val, y fold val),
verbose=1)
  # Optionally evaluate on validation data
  score = cnn model.evaluate(X fold val, y fold val, verbose=0)
  print(f"Fold {fold + 1} validation accuracy: {score[1]}")
# Train the CNN model on the entire training set after cross-validation
cnn model = create cnn model()
cnn model.fit(X train, y train cat, epochs=10, batch size=32, validation data=(X test, y test cat), verbose=1)
# Initialize LIME Text Explainer
explainer = LimeTextExplainer(class_names=class_names)
# Function for CNN prediction probability
def cnn predict proba(texts):
  vec texts = tokenizer.texts to sequences(texts)
  vec texts = pad sequences(vec texts, maxlen=100)
  return cnn model.predict(vec texts)
# Generate LIME explanations for the first annotated review in each class from the full dataset
for class name in class names:
  class id = label encoder.transform([class name])[0]
  class_indices = df[df['category'] == class_id].index # Find indices of all reviews in this class
  if class indices. size > 0:
    first review index = class indices[0] # Get the index of the first review for this class
    text instance = df['Base Reviews'].iloc[first review index]
    # Generate the LIME explanation
    exp = explainer.explain instance(text instance, cnn predict proba, num features=10, labels=[class id])
    # Save explanation as HTML
    html file name = f'Z LIME explanation {class name}.html'
    exp.save to file(html file name)
    print(f'Saved LIME explanation for class "{class name}" to {html file name}')
    exp.show in notebook(text=True) # Optional: Display in notebook
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# Create and save the feature importance plot
num samples = 20
aggregate explanations = []
for i in range(num samples):
  text instance = df['Base Reviews'].iloc[i] # Get the review text
  exp = explainer.explain_instance(text_instance, cnn_predict_proba, num_features=20)
  aggregate explanations.extend(exp.as list())
# Calculate and plot the aggregate feature importance
feature importances = {}
for feature, importance in aggregate explanations:
  if feature not in feature importances:
     feature importances[feature] = 0
  feature importances[feature] += importance
sorted features = sorted(feature importances.items(), key=lambda x: abs(x[1]), reverse=True)[:20]
features, importances = zip(*sorted features)
colors = ['green' if x > 0 else 'red' for x in importances]
pos = np.arange(len(features)) + .5
# Plot feature importance
fig, ax = plt.subplots(figsize=(14, 10))
bars = ax.barh(pos, importances, align='center', color=colors)
ax.set yticks(pos)
ax.set yticklabels(features, fontsize=14)
ax.set xlabel('Aggregate Importance', fontsize=16)
ax.set title('Overall LIME Feature Importances (Top 20 Features)', fontsize=18)
plt.tight layout()
plt.savefig('z_Overall_LIME_feature_importances_CNN.png', dpi=300) # Save at 300 DPI
plt.show()
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