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import pandas as pd
import seaborn as sns
from sklearn.linear_model import LogisticRegression
from sklearn.model_selection import train_test_split
from sklearn.metrics import confusion matrix
from sklearn import datasets
cancer= datasets.load breast cancer()
X = cancer.data
y = cancer.target
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)
model = LogisticRegression()
model.fit(X_train, y_train)
y pred = model.predict(X test)
cm = confusion_matrix(y_test, y_pred)
sns.heatmap(cm, annot=True, cmap='Blues')
2
import numpy as np
import matplotlib.pyplot as plt
from scipy.optimize import minimize_scalar
def objective_function(x):
  return x**2 + 2*x +1
result = minimize_scalar(objective_function)
optima = result.x
x = np.linspace(-10, 10, 100)
y = objective_function(x)
plt.plot(x, y)
plt.scatter(optima, objective_function(optima), color='red', label='Optima')
plt.legend()
plt.xlabel('x')
plt.ylabel('f(x)')
plt.title('Optimizing Unconstrained Convex Univariate Function')
plt.show()
```

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3
import numpy as np
import matplotlib.pyplot as plt
from scipy.optimize import minimize
def objective(x):
  return x[0]**2 + x[1]**2
def constraint(x):
  return x[0] + x[1] - 1
x0 = np.array([0.5, 0.5])
result = minimize(objective, x0, method='SLSQP', constraints={'type': 'eq', 'fun': constraint})
print("Minimum found at:")
print("x:", result.x[0])
print("y:", result.x[1])
print("Objective value:", result.fun)
x1 grid, x2 grid = np.meshgrid(np.linspace(0, 1, 100), np.linspace(0, 1, 100))
Z = objective([x1\_grid, x2\_grid])
x_values = np.linspace(0, 1, 100)
y_values = 1 - x_values
plt.contourf(x1_grid, x2_grid, Z, levels=20, cmap='viridis')
plt.plot(x values, y values, label='Constraint: x + y = 1', color='red')
plt.scatter(result.x[0], result.x[1], color='black', marker='x', s=100, label='Minimum')
plt.xlabel('x')
plt.ylabel('y')
plt.title('Optimization with Constraint and Objective Function')
plt.colorbar(label='Objective Function Value')
plt.legend()
plt.grid(True)
plt.show()
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```
import numpy as np
from sklearn.metrics import mean squared error
def user based cf(user item matrix, target user):
  similarities = np.dot(user item matrix, user item matrix[target user]) / (
     np.linalg.norm(user_item_matrix, axis=1) * np.linalg.norm(user_item_matrix[target_user]))
  top items = np.argsort(similarities)[::-1][:3]
  recommended items = [item for item in top items if user item matrix[target user, item] == 0]
  unrated items = np.where(user item matrix[target user] == 0)[0]
  predicted_ratings = np.sum(similarities[:, np.newaxis] * user_item_matrix[:, unrated_items],
axis=0)
  actual_ratings = user_item_matrix[target_user, unrated_items]
  rmse = mean squared error(actual ratings, predicted ratings, squared=False)
  return recommended_items, rmse
user item matrix = np.array([[5, 4, 0],
                  [4, 0, 5],
                  [0, 2, 3]]
target user = 0
recommended_items, rmse = user_based_cf(user_item_matrix, target_user)
print("Recommended items for user", target user, ":", recommended items)
print("RMSE:", rmse)
import numpy as np
import math
def user based collaborative filtering(data):
  user means = np.mean(data, axis=1)
  centered_data = data - user_means[:, np.newaxis]
  similarity = np.dot(centered data, centered data.T) / (np.linalg.norm(centered data,
axis=1)[:, np.newaxis] * np.linalg.norm(centered_data.T, axis=0))
  np.fill diagonal(similarity, 0)
  predicted_ratings = user_means[:, np.newaxis] + np.dot(similarity, centered_data) /
np.sum(np.abs(similarity), axis=1)[:, np.newaxis]
  mse = np.mean((data - predicted_ratings) ** 2)
  return predicted ratings, mse
data = np.array([[5, 3, 0, 1],
          [4, 0, 0, 1],
          [1, 1, 0, 5],
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[1, 0, 0, 4],
          [0, 1, 5, 4]])
predicted_ratings, mse = user_based_collaborative_filtering(data)
pred=list(predicted ratings)
new=[]
for i in range(data.shape[0]):
  new.append([])
  for j in range(data.shape[1]):
    if data[i,j] == 0:
       replace_idx = i % len(pred)
       f=pred[replace idx]
       new[i].append(abs(round(f[j],2)))
     else:
       new[i].append(data[i][j])
print("Predicted Ratings:")
for i in new:
  print(i)
print("Mean Squared Error (MSE):", mse)
5
!pip install gensim
!pip install nltk
import nltk
nltk.download('punkt')
from gensim.models import Word2Vec, Doc2Vec
from gensim.models.doc2vec import TaggedDocument
from sklearn.metrics.pairwise import cosine_similarity
from nltk import word tokenize
document1 = "This is the first document."
document4 = "Is this the first document?"
tokenized doc1 = word tokenize(document1.lower())
tokenized_doc4 = word_tokenize(document4.lower())
word2vec_model = Word2Vec([tokenized_doc1,tokenized_doc4], min_count=1)
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tagged_documents = [TaggedDocument(words=word_tokenize(doc.lower()), tags=[str(i)]) for i, doc in enumerate([document1, document4])]
doc2vec_model = Doc2Vec(tagged_documents, vector_size=100, min_count=1, epochs=40)
word_embedding1 = word2vec_model.wv[tokenized_doc1]
word_embedding4 = word2vec_model.wv[tokenized_doc4]

doc_embedding4 = doc2vec_model.infer_vector(tokenized_doc4)

cosine_similarity_word = cosine_similarity([word_embedding1[:,0]], [word_embedding4[:,0]])
cosine_similarity_doc = cosine_similarity([doc_embedding1], [doc_embedding4])
jaccard_similarity = len(set(tokenized_doc1).intersection(tokenized_doc4)) /
len(set(tokenized_doc1).union(tokenized_doc4))

print("Cosine Similarity (Word Embedding):", cosine_similarity_word[0][0])
print("Cosine Similarity:", jaccard_similarity)
```