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CSE AI ML-B
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DECISION TREE
# Import necessary libraries
import pyspark
import os
import sys
from pyspark import SparkContext
from pyspark.sql import SparkSession
# Set up PySpark environment
os.environ['PYSPARK PYTHON'] = sys.executable # Setting Python executable
for PySpark
os.environ['PYSPARK DRIVER PYTHON'] = sys.executable # Setting driver
Python executable for PySpark
spark = SparkSession.builder.config("spark.driver.memory",
"16g").appName('chapter 4').getOrCreate() # Creating a SparkSession with
configuration
# Load the dataset without header
data without header = spark.read.option("inferSchema",
True).option("header", False).csv("data/covtype.data") # Loading data
without header
# Define the column names
colnames = ["Elevation", "Aspect", "Slope", # Defining column names for
the dataset
            "Horizontal Distance To Hydrology",
            "Vertical Distance To Hydrology",
"Horizontal Distance To Roadways",
            "Hillshade 9am", "Hillshade Noon", "Hillshade 3pm",
            "Horizontal_Distance_To_Fire_Points"] +
            [f"Wilderness Area {i}" for i in range(4)] +
            [f"Soil Type {i}" for i in range(40)] +
            ["Cover Type"]
# Create a DataFrame with column names and cast the label column to
DoubleType
data = data without header.toDF(*colnames).withColumn("Cover Type",
col("Cover Type").cast(DoubleType())) # Creating DataFrame with specified
column names and data types
# Split the data into train and test sets
(train data, test data) = data.randomSplit([0.9, 0.1]) # Splitting data
into train and test sets
train data.cache() # Caching train data for optimization
test data.cache() # Caching test data for optimization
# Feature engineering
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input_cols = colnames[:-1] # Defining input columns for feature
engineering
vector assembler = VectorAssembler(inputCols=input cols,
outputCol="featureVector") # Creating VectorAssembler for feature
engineering
assembled train data = vector assembler.transform(train data) #
Transforming train data using VectorAssembler
# Train a Decision Tree Classifier
classifier = DecisionTreeClassifier(seed=1234, labelCol="Cover Type",
featuresCol="featureVector", predictionCol="prediction") # Creating
DecisionTreeClassifier
model = classifier.fit(assembled train data) # Training the model
# Print feature importances
print(model.toDebugString) # Printing the debug string of the model
pd.DataFrame(model.featureImportances.toArray(), index=input cols,
columns=['importance']).sort values(by="importance", ascending=False) #
Displaying feature importances
# Make predictions and evaluate the model
predictions = model.transform(assembled train data) # Making predictions
on train data
evaluator = MulticlassClassificationEvaluator(labelCol="Cover Type",
predictionCol="prediction") # Creating evaluator for model evaluation
print(f"Accuracy:
{evaluator.setMetricName('accuracy').evaluate(predictions)}") #
Calculating and printing accuracy
print(f"F1-score: {evaluator.setMetricName('f1').evaluate(predictions)}")
# Calculating and printing F1-score
confusion matrix = predictions.groupBy("Cover Type").pivot("prediction",
range(1,8)).count().na.fill(0.0).orderBy("Cover Type") # Calculating
confusion matrix
confusion matrix.show() # Displaying confusion matrix
# Calculate class probabilities
def class probabilities(data):
   total = data.count()
    return
data.groupBy("Cover Type").count().orderBy("Cover Type").select(col("count"
).cast(DoubleType())).withColumn("count proportion",
col("count")/total).select("count proportion").collect()
train prior probabilities = class probabilities(train data) # Calculating
train prior probabilities
test_prior_probabilities = class_probabilities(test_data) # Calculating
test prior probabilities
train prior probabilities = [p[0] for p in train prior probabilities]
Extracting train prior probabilities
test prior probabilities = [p[0] for p in test prior probabilities] #
Extracting test prior probabilities
print(f"Sum of train and test prior probabilities: {sum([train p * cv p for
train_p, cv_p in zip(train_prior_probabilities,
test prior probabilities)]) }")  # Calculating and printing sum of train and
test prior probabilities
# Hyperparameter tuning
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assembler = VectorAssembler(inputCols=input cols,
outputCol="featureVector") # Creating VectorAssembler for hyperparameter
tuning
classifier = DecisionTreeClassifier(seed=1234, labelCol="Cover Type",
featuresCol="featureVector", predictionCol="prediction") # Creating
DecisionTreeClassifier for hyperparameter tuning
pipeline = Pipeline(stages=[assembler, classifier]) # Creating pipeline
for hyperparameter tuning
paramGrid = ParamGridBuilder().addGrid(classifier.impurity, ["gini",
"entropy"]).addGrid(classifier.maxDepth, [1,
20]).addGrid(classifier.maxBins, [40, 300]).addGrid(classifier.minInfoGain,
[0.0, 0.05]).build() # Creating parameter grid for hyperparameter tuning
multiclassEval =
MulticlassClassificationEvaluator().setLabelCol("Cover Type").setPrediction
Col("prediction").setMetricName("accuracy") # Creating evaluator for model
evaluation in hyperparameter tuning
validator = TrainValidationSplit(seed=1234, estimator=pipeline,
evaluator=multiclassEval, estimatorParamMaps=paramGrid, trainRatio=0.9) #
Creating TrainValidationSplit for hyperparameter tuning
validator model = validator.fit(train data) # Fitting TrainValidationSplit
on train data
best model = validator model.bestModel # Extracting best model from
TrainValidationSplit
print(best model.stages[1].extractParamMap()) # Printing best model
parameters
print(f"Best model accuracy on validation set: {metrics[0]}") # Printing
best model accuracy on validation set
print(f"Best model accuracy on test set:
{multiclassEval.evaluate(best model.transform(test data))}") # Calculating
and printing best model accuracy on test set
# One-hot encoding
def unencode one hot(data):
    wilderness cols = ['Wilderness Area ' + str(i) for i in range(4)] #
Defining wilderness area columns
    wilderness assembler =
VectorAssembler().setInputCols(wilderness cols).setOutputCol("wilderness")
# Creating VectorAssembler for wilderness area
    unhot udf = udf(lambda v: v.toArray().tolist().index(1)) # Creating
UDF for unencoding one-hot vectors
    with wilderness =
wilderness assembler.transform(data).drop(*wilderness cols).withColumn("wil
derness", unhot udf(col("wilderness")).cast(IntegerType())) # Unencoding
wilderness area
    soil_cols = ['Soil_Type_' + str(i) for i in range(40)] # Defining soil
type columns
    soil assembler =
VectorAssembler().setInputCols(soil cols).setOutputCol("soil") # Creating
VectorAssembler for soil type
    with soil =
soil assembler.transform(with wilderness).drop(*soil cols).withColumn("soil
", unhot udf(col("soil")).cast(IntegerType()))  # Unencoding soil type
    return with soil # Returning unencoded DataFrame
unenc train data = unencode one hot(train data) # Unencoding one-hot
vectors in train data
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unenc_test_data = unencode_one_hot(test_data) # Unencoding one-hot vectors
in test data
# Random Forest Classifier
cols = unenc train data.columns # Extracting columns from unencoded train
data
input cols = [c for c in cols if c!='Cover Type'] # Defining input columns
for Random Forest Classifier
assembler =
VectorAssembler().setInputCols(input cols).setOutputCol("featureVector") #
Creating VectorAssembler for Random Forest Classifier
indexer =
VectorIndexer().setMaxCategories(40).setInputCol("featureVector").setOutput
Col("indexedVector")  # Creating VectorIndexer for Random Forest Classifier
classifier = RandomForestClassifier(seed=1234, labelCol="Cover Type",
featuresCol="indexedVector", predictionCol="prediction") # Creating
RandomForestClassifier
pipeline = Pipeline().setStages([assembler, indexer, classifier]) #
Creating pipeline for Random Forest Classifier
paramGrid = ParamGridBuilder().addGrid(classifier.impurity, ["gini",
"entropy"]).addGrid(classifier.maxDepth, [1,
20]).addGrid(classifier.maxBins, [40, 300]).addGrid(classifier.minInfoGain,
[0.0, 0.05]).build() # Creating parameter grid for Random Forest
Classifier
multiclassEval =
MulticlassClassificationEvaluator().setLabelCol("Cover Type").setPrediction
Col("prediction").setMetricName("accuracy") # Creating evaluator for model
evaluation in Random Forest Classifier
validator = TrainValidationSplit(seed=1234, estimator=pipeline,
evaluator=multiclassEval, estimatorParamMaps=paramGrid, trainRatio=0.9) #
Creating TrainValidationSplit for Random Forest Classifier
validator model = validator.fit(unenc train data) # Fitting
TrainValidationSplit on unencoded train data
best model = validator model.bestModel # Extracting best model from
TrainValidationSplit
forest model = best model.stages[2] # Extracting Random Forest model from
best model
feature importance list = list(zip(input cols,
forest model.featureImportances.toArray())) # Creating list of feature
importances
feature importance list.sort(key=lambda x: x[1], reverse=True) # Sorting
feature importances
print("Feature importances:") # Printing feature importances
pprint(feature importance list) # Pretty-printing feature importances
# Make predictions on the test set using the best Random Forest model
best model.transform(unenc test data.drop("Cover Type")).select("prediction
").show(1) # Making predictions on test data using best Random Forest
model and displaying predictions
```

ENTITY RESOLUTION

```
import pyspark # Import PySpark library
import os # Import os module for operating system functionalities
import sys # Import sys module for system-specific parameters
from pyspark.sql import SparkSession # Import SparkSession from PySpark
# Set up PySpark environment
os.environ['PYSPARK PYTHON'] = sys.executable # Setting Python executable
for PySpark
os.environ['PYSPARK DRIVER PYTHON'] = sys.executable # Setting driver
Python executable for PySpark
spark = SparkSession.builder.config("spark.driver.memory",
"16g").appName('chapter 2').getOrCreate() # Creating a SparkSession with
configuration
# Read the CSV files
prev = spark.read.csv("data/linkage/donation/block 1/block 1.csv") #
Reading a CSV file without header
prev.show(2) # Displaying the first two rows of the DataFrame
# Read and parse CSV with options
parsed = spark.read.option("header", "true").option("nullValue",
"?").option("inferSchema",
"true").csv("data/linkage/donation/block 1/block 1.csv") # Reading and
parsing a CSV file with options
parsed.printSchema() # Printing the schema of the DataFrame
parsed.show(5) # Displaying the first five rows of the DataFrame
# Count rows and cache DataFrame
parsed.count() # Counting the number of rows in the DataFrame
parsed.cache() # Caching the DataFrame for optimization
# Group by 'is match' column and show counts
from pyspark.sql.functions import col # Importing col function from
PySpark
parsed.groupBy("is match").count().orderBy(col("count").desc()).show() #
Grouping by 'is match' column and showing counts in descending order
# Create temporary view for SQL queries
parsed.createOrReplaceTempView("linkage") # Creating a temporary view for
SQL queries
spark.sql("""
SELECT is match, COUNT(*) cnt
FROM linkage
GROUP BY is match
ORDER BY cnt DESC
""").show() # Executing a SQL query and displaying the result
```

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# Summary statistics and operations
summary = parsed.describe() # Generating summary statistics for the
DataFrame
summary.select("summary", "cmp fname c1", "cmp fname c2").show()
Selecting specific columns from the summary DataFrame and showing them
# Filtering and describing matches and misses
matches = parsed.where("is match = true") # Filtering rows where
'is match' is true
match summary = matches.describe() # Generating summary statistics for
matches
misses = parsed.filter(col("is match") == False) # Filtering rows where
'is match' is false
miss summary = misses.describe() # Generating summary statistics for
misses
# Converting summary to Pandas for operations
summary p = summary.toPandas() # Converting the summary DataFrame to
Pandas DataFrame
summary p.head() # Displaying the first few rows of the Pandas DataFrame
summary p.shape # Displaying the shape of the Pandas DataFrame
# Transpose and convert back to Spark DataFrame
summary p = summary p.set index('summary').transpose().reset index()
Transposing the Pandas DataFrame and resetting the index
summary p = summary p.rename(columns={'index':'field'}) # Renaming columns
of the DataFrame
summary p = summary p.rename axis(None, axis=1) # Removing the axis name
from the DataFrame
summary p.shape # Displaying the shape of the DataFrame
summaryT = spark.createDataFrame(summary p) # Creating a Spark DataFrame
from the Pandas DataFrame
summaryT.printSchema() # Printing the schema of the Spark DataFrame
# Convert metric columns to DoubleType
from pyspark.sql.types import DoubleType # Importing DoubleType from
PySpark
for c in summaryT.columns: # Looping through columns of the DataFrame
    if c == 'field': # Skipping the 'field' column
    summaryT = summaryT.withColumn(c, summaryT[c].cast(DoubleType())) #
Converting columns to DoubleType
summaryT.printSchema() # Printing the schema of the DataFrame
# Define a function for pivoting and converting summary
def pivot summary(desc): # Defining a function for pivoting and converting
summary
   desc p = desc.toPandas() # Converting Spark DataFrame to Pandas
DataFrame
   desc p = desc p.set index('summary').transpose().reset index() #
Transposing and resetting index of the Pandas DataFrame
   desc p = desc p.rename(columns={'index':'field'}) # Renaming columns
of the DataFrame
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desc p = desc p.rename axis(None, axis=1) # Removing the axis name
from the DataFrame
   descT = spark.createDataFrame(desc p) # Creating a Spark DataFrame
from the Pandas DataFrame
    for c in descT.columns: # Looping through columns of the DataFrame
        if c == 'field': # Skipping the 'field' column
        else:
           descT = descT.withColumn(c, descT[c].cast(DoubleType())) #
Converting columns to DoubleType
   return descT # Returning the converted DataFrame
match summaryT = pivot summary(match summary) # Pivoting and converting
match summary
miss summaryT = pivot summary(miss summary) # Pivoting and converting miss
summary
match summaryT.createOrReplaceTempView("match desc") # Creating a
temporary view for match summary
miss summaryT.createOrReplaceTempView("miss desc") # Creating a temporary
view for miss summary
# SQL query for comparing matches and misses
spark.sql("""
SELECT a.field, a.count + b.count total, a.mean - b.mean delta
FROM match desc a INNER JOIN miss desc b ON a.field = b.field
WHERE a.field NOT IN ("id 1", "id 2")
ORDER BY delta DESC, total DESC
""")  # Executing a SQL query for comparing matches and misses
# Good features and scoring
good features = ["cmp lname c1", "cmp plz", "cmp by", "cmp bd", "cmp bm"]
# Defining good features for scoring
sum expression = " + ".join(good features) # Joining good features with a
sum expression
from pyspark.sql.functions import expr # Importing expr function from
scored = parsed.fillna(0, subset=good features).withColumn('score',
expr(sum expression)).select('score', 'is match') # Filling NA values,
scoring, and selecting columns
scored.show() # Displaying the scored DataFrame
# Cross-tabulation function
def crossTabs(scored: DataFrame, t: DoubleType) -> DataFrame: # Defining a
function for cross-tabulation
    return scored.selectExpr(f"score >= {t} as above",
"is match").groupBy("above").pivot("is match", ("true", "false")).count()
# Performing cross-tabulation
crossTabs(scored, 4.0).show() # Performing cross-tabulation with threshold
4.0
crossTabs(scored, 2.0).show() # Performing cross-tabulation with threshold
2.0
```

K-Means

```
# Import necessary libraries
import pyspark # Importing PySpark for distributed computing
import os # Operating System module for environment setup
import sys # System-specific parameters and functions
from pyspark import SparkContext # SparkContext for controlling Spark
functionality
from pyspark.sql import SparkSession # SparkSession for working with
structured data
from pyspark.sql.functions import col # Column functions for data
manipulation
from pyspark.ml.feature import VectorAssembler, StandardScaler,
OneHotEncoder, StringIndexer # ML feature engineering tools
from pyspark.ml.clustering import KMeans # KMeans clustering algorithm
from MLlib
from pyspark.ml import Pipeline # Pipeline for ML workflow management
from random import randint # Random number generator
from math import log # Logarithmic function for entropy calculation
from pyspark.sql import functions as fun # Alias for SQL functions in
from pyspark.sql import Window # Window functions for data aggregation and
manipulation
# Set up Spark session
os.environ['PYSPARK PYTHON'] = sys.executable # Set Python executable for
PySpark
os.environ['PYSPARK DRIVER PYTHON'] = sys.executable # Set Python
executable for PySpark driver
spark = SparkSession.builder.config("spark.driver.memory",
"16g").appName('chapter 5').getOrCreate() # Configure Spark session
# Read data and define column names
data without header = spark.read.option("inferSchema",
True).option("header", False).csv("data/kddcup.data 10 percent corrected")
# Read CSV data without inferring schema
column names = [ "duration", "protocol type", "service", "flag",
"src bytes", "dst bytes", "land", "wrong fragment", "urgent", "hot",
"num_failed_logins", "logged_in", "num_compromised", "root_shell",
"su attempted", "num root", "num file creations", "num shells",
"num access files", "num outbound cmds", "is host login", "is guest login",
"count", "srv_count", "serror_rate", "srv_serror_rate", "rerror_rate",
"srv_rerror_rate", "same_srv_rate", "diff_srv_rate", "srv_diff_host_rate",
"dst_host_count", "dst_host_srv count", "dst host same srv rate",
"dst host diff srv rate", "dst host same src port rate",
"dst host srv diff host rate", "dst host serror rate",
"dst_host_srv_serror_rate", "dst_host_rerror_rate",
"dst host srv rerror rate", "label"] # Define column names for the data
data = data_without_header.toDF(*column_names) # Create DataFrame with
specified column names
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# Display count of each label
data.select("label").groupBy("label").count().orderBy(col("count").desc()).
show(25) # Group by label, count occurrences, and show top 25 counts
# Define functions for clustering and evaluation
# Function to calculate clustering score based on training cost
def clustering score(input data, k):
    # Prepare data
   input numeric only = input data.drop("protocol type", "service",
"flag")  # Exclude non-numeric columns
    assembler =
VectorAssembler().setInputCols(input numeric only.columns[:-
1]).setOutputCol("featureVector")  # Assemble feature vector
KMeans().setSeed(randint(100,100000)).setK(k).setPredictionCol("cluster").s
etFeaturesCol("featureVector") # Define KMeans algorithm
    pipeline = Pipeline().setStages([assembler, kmeans]) # Create ML
pipeline
    # Fit pipeline and get training cost
    pipeline model = pipeline.fit(input numeric only)
    kmeans_model = pipeline model.stages[-1]
    training cost = kmeans model.summary.trainingCost
   return training cost
# Function to calculate clustering score with additional parameters
def clustering score 1(input data, k):
    # Prepare data
    input numeric only = input data.drop("protocol type", "service",
"flag")  # Exclude non-numeric columns
    assembler =
VectorAssembler().setInputCols(input numeric only.columns[:-
1]).setOutputCol("featureVector")  # Assemble feature vector
    kmeans =
KMeans().setSeed(randint(100,100000)).setK(k).setMaxIter(40).setTol(1.0e-
5).setPredictionCol("cluster").setFeatureSCol("featureVector") # Define
KMeans algorithm with additional parameters
   pipeline = Pipeline().setStages([assembler, kmeans]) # Create ML
pipeline
    # Fit pipeline and get training cost
    pipeline model = pipeline.fit(input numeric only)
    kmeans_model = pipeline_model.stages[-1]
    training cost = kmeans model.summary.trainingCost
    return training_cost
# Function to calculate clustering score with standard scaling
def clustering score 2(input data, k):
    # Prepare data
    input numeric only = input data.drop("protocol type", "service",
"flag") # Exclude non-numeric columns
    assembler =
VectorAssembler().setInputCols(input numeric only.columns[:-
1]).setOutputCol("featureVector")  # Assemble feature vector
    scaler =
StandardScaler().setInputCol("featureVector").setOutputCol("scaledFeatureVe
ctor").setWithStd(True).setWithMean(False) # Scale features
```

```
kmeans =
KMeans().setSeed(randint(100,100000)).setK(k).setMaxIter(40).setTol(1.0e-
5).setPredictionCol("cluster").setFeaturesCol("scaledFeatureVector")
Define KMeans algorithm with scaled features
    pipeline = Pipeline().setStages([assembler, scaler, kmeans]) # Create
ML pipeline
    # Fit pipeline and get training cost
    pipeline model = pipeline.fit(input numeric only)
    kmeans model = pipeline model.stages[-1]
    training cost = kmeans model.summary.trainingCost
    return training cost
from pyspark.ml.feature import OneHotEncoder, StringIndexer
def one hot pipeline(input col):
# Function to calculate clustering score with one-hot encoding
def clustering score 3(input_data, k):
    # Prepare data
    proto type pipeline, proto type vec col =
one_hot_pipeline("protocol_type") # One-hot encode protocol_type column
    service_pipeline, service_vec_col = one_hot_pipeline("service") # One-
hot encode service column
    flag pipeline, flag vec col = one hot pipeline("flag") # One-hot
encode flag column
    assemble cols = set(input data.columns) - {"label", "protocol type",
"service", "flag"} | {proto type vec col, service vec col, flag vec col} #
Combine columns
    assembler =
VectorAssembler().setInputCols(list(assemble cols)).setOutputCol("featureVe
ctor") # Assemble feature vector
    scaler =
StandardScaler().setInputCol("featureVector").setOutputCol("scaledFeatureVe
ctor").setWithStd(True).setWithMean(False) # Scale features
KMeans().setSeed(randint(100,100000)).setK(k).setMaxIter(40).setTol(1.0e-
5).setPredictionCol("cluster").setFeaturesCol("scaledFeatureVector") #
Define KMeans algorithm with scaled features
    pipeline = Pipeline().setStages([proto_type_pipeline, service_pipeline,
flag pipeline, assembler, scaler, kmeans]) # Create ML pipeline
    # Fit pipeline and get training cost
    pipeline model = pipeline.fit(input data)
    kmeans_model = pipeline_model.stages[-1]
    training cost = kmeans model.summary.trainingCost
    return training cost
# Function to calculate entropy
def entropy(counts):
    values = [c for c in counts if (c > 0)] # Filter non-zero counts
    n = sum(values) # Calculate total count
    p = [v/n for v in values] # Calculate probabilities
    return sum([-1*(p v) * log(p_v) for p_v in p]) # Calculate entropy
# Function to fit pipeline for clustering with one-hot encoding
def fit pipeline 4(data, k):
    (proto type pipeline, proto type vec col) =
one hot pipeline("protocol type") # One-hot encode protocol type column
    (service pipeline, service vec col) = one hot pipeline("service")
One-hot encode service column
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(flag pipeline, flag vec col) = one hot pipeline("flag") # One-hot
encode flag column
    assemble cols = set(data.columns) - {"label", "protocol type",
"service", "flag"} | {proto type vec col, service vec col, flag vec col} #
Combine columns
    assembler = VectorAssembler(inputCols=list(assemble cols),
outputCol="featureVector") # Assemble feature vector
    scaler = StandardScaler(inputCol="featureVector",
outputCol="scaledFeatureVector", withStd=True, withMean=False) # Scale
features
    kmeans model = KMeans(seed=randint(100,100000), k=k, maxIter=40,
tol=1.0e-5, predictionCol="cluster", featureSCol="scaledFeatureVector")
Define KMeans algorithm with scaled features
    pipeline = Pipeline(stages=[proto type pipeline, service pipeline,
flag pipeline, assembler, scaler, kmeans model]) # Create ML pipeline
    pipeline model = pipeline.fit(data) # Fit pipeline
    return pipeline model
# Define function to calculate clustering score based on training cost
def clustering score 4(input data, k):
    # Fit the ML pipeline for clustering
    pipeline model = fit pipeline 4(input data, k)
    # Perform clustering and label analysis
    cluster label = pipeline model.transform(input data).select("cluster",
"label") # Transform data and select relevant columns
    df = cluster label.groupBy("cluster",
"label").count().orderBy("cluster")  # Group by cluster and label, count
occurrences, and order by cluster
    # Calculate probabilities and entropy for each cluster
    w = Window.partitionBy("cluster") # Define window for partitioning
    p col = df['count'] / fun.sum(df['count']).over(w) # Calculate
probabilities within each cluster
    with_p_col = df.withColumn("p_col", p_col) # Add probability column to
DataFrame
   result = with p col.groupBy("cluster").agg(-fun.sum(col("p col") *
fun.log2(col("p col"))).alias("entropy"), # Calculate entropy for each
fun.sum(col("count")).alias("cluster size")) # Calculate cluster size
    result = result.withColumn('weightedClusterEntropy', col('entropy') *
col('cluster_size')) # Calculate weighted cluster entropy
    weighted cluster entropy avg =
result.agg(fun.sum(col('weightedClusterEntropy'))).collect() # Calculate
average weighted cluster entropy
    return weighted_cluster_entropy_avg[0][0] / input_data.count() #
Return normalized clustering score
# Fit the ML pipeline for clustering with k=180
pipeline model = fit pipeline 4(data, 180)
# Perform clustering and label analysis
count by cluster label = pipeline model.transform(data).select("cluster",
"label").groupBy("cluster", "label").count().orderBy("cluster", "label")
count by cluster label.show()
# Display the result of clustering and label analysis
```

MONTE-CARLO

```
# Import necessary libraries
import pyspark
import os
import sys
from pyspark import SparkContext
# Set environment variables for PySpark
os.environ['PYSPARK PYTHON'] = sys.executable
os.environ['PYSPARK DRIVER PYTHON'] = sys.executable
# Import SparkSession from PySpark SQL module
from pyspark.sql import SparkSession
# Create a SparkSession with specified configuration
spark = SparkSession.builder.config("spark.driver.memory",
"16g").appName('chapter 8').getOrCreate()
# Read CSV files into a PySpark DataFrame
stocks =
spark.read.csv(["data/stocksA/ABAX.csv","data/stocksA/AAME.csv","data/stock
sA/AEPI.csv"], header='true', inferSchema='true')
# Show the first two rows of the DataFrame
stocks.show(2)
# Import SQL functions from PySpark
from pyspark.sql import functions as fun
# Add a column 'Symbol' to the DataFrame based on the file name
stocks = stocks.withColumn("Symbol", fun.input file name()).
withColumn("Symbol", fun.element at(fun.split("Symbol", "/"), -1)).
withColumn("Symbol", fun.element_at(fun.split("Symbol", "."), 1))
# Show the first two rows of the updated DataFrame
stocks.show(2)
# Read the same CSV files into another DataFrame
factors =
spark.read.csv(["data/stocksA/ABAX.csv","data/stocksA/AAME.csv","data/stock
sA/AEPI.csv"], header='true', inferSchema='true')
# Add a column 'Symbol' to the DataFrame based on the file name
factors = factors.withColumn("Symbol", fun.input file name()).
withColumn("Symbol", fun.element_at(fun.split("Symbol", "/"), -1)).
withColumn("Symbol", fun.element at(fun.split("Symbol", "."), 1))
# Import Window function from PySpark
```

```
from pyspark.sql import Window
# Filter the stocks DataFrame based on a condition
stocks = stocks.withColumn('count', fun.count('Symbol').
over(Window.partitionBy('Symbol'))).
filter(fun.col('count') > 260*5 + 10)
# Set a configuration for parsing date/time strings
spark.sql("set spark.sql.legacy.timeParserPolicy=LEGACY")
# Convert the 'Date' column to date format
stocks = stocks.withColumn('Date',
fun.to date(fun.to timestamp(fun.col('Date'), 'dd-MMM-yy')))
# Print the schema of the DataFrame
stocks.printSchema()
# Import datetime module
from datetime import datetime
# Filter the stocks DataFrame based on date range
stocks = stocks.filter(fun.col('Date') >= datetime(2009, 10, 23)).
filter(fun.col('Date') <= datetime(2014, 10, 23))</pre>
# Convert the 'Date' column to date format and filter based on date range
factors = factors.withColumn('Date',
fun.to date(fun.to timestamp(fun.col('Date'), 'dd-MMM-yy')))
factors = factors.filter(fun.col('Date') >= datetime(2009, 10, 23)).
filter(fun.col('Date') <= datetime(2014, 10, 23))</pre>
# Convert PySpark DataFrames to pandas DataFrames
stocks pd df = stocks.toPandas()
factors pd df = factors.toPandas()
# Print the first 5 rows of the factors pd df DataFrame
factors pd df.head(5)
# Set the number of steps for rolling window calculation
n \text{ steps} = 10
# Define a function to calculate returns
def my fun(x):
    return ((x.iloc[-1] - x.iloc[0]) / x.iloc[0])
# Calculate stock returns and factor returns using rolling window
stock returns =
stocks pd df.groupby('Symbol').Close.rolling(window=n steps).apply(my fun)
factors returns =
factors pd df.groupby('Symbol').Close.rolling(window=n steps).apply(my fun)
# Reset and sort the index of the returns DataFrames
stock returns =
stock returns.reset index().sort values('level 1').reset index()
\hbox{factors returns} \; = \;
factors returns.reset index().sort values('level 1').reset index()
# Add stock returns to the stocks DataFrame
```

```
stocks pd df with returns =
stocks pd df.assign(stock returns=stock returns['Close'])
# Add factor returns and squared factor returns to the factors DataFrame
factors pd df with returns =
factors pd df.assign(factors returns=factors returns['Close'],
factors returns squared=factors returns['Close']**2)
# Pivot the factors DataFrame to have factors and squared factors as
columns
factors pd df with returns = factors pd df with returns.pivot(index='Date',
columns='Symbol', values=['factors returns', 'factors returns squared'])
# Modify the column names of the pivoted DataFrame
factors pd df with returns.columns =
factors pd df with returns.columns.to series().str.join(' ').reset index()[
01
factors pd df with returns = factors pd df with returns.reset index()
# Print the first row and column names of the pivoted DataFrame
print(factors pd df with returns.head(1))
print(factors pd df with returns.columns)
# Import necessary libraries
import pandas as pd
from sklearn.linear model import LinearRegression
# Merge the stocks and factors DataFrames
stocks factors combined df = pd.merge(stocks pd df with returns,
factors pd df with returns, how="left", on="Date")
# Get the feature column names
feature columns = list(stocks factors combined df.columns[-6:])
# Drop rows with NaN values in feature columns and stock returns
with pd.option context('mode.use inf as na', True):
    stocks factors combined df =
stocks factors combined df.dropna(subset=feature columns +
['stock returns'])
# Function to find OLS coefficients
def find ols coef(df):
    y = df[['stock returns']].values
    X = df[feature columns]
    regr = LinearRegression()
    regr_output = regr.fit(X, y)
    return list(df[['Symbol']].values[0]) + list(regr output.coef [0])
# Calculate OLS coefficients for each stock
coefs per stock =
stocks factors combined df.groupby('Symbol').apply(find ols coef)
coefs per stock = pd.DataFrame(coefs per stock).reset index()
coefs_per_stock.columns = ['symbol', 'factor_coef_list']
coefs per stock = pd.DataFrame(coefs per stock.factor coef list.tolist(),
index=coefs per stock.index, columns=['Symbol'] + feature columns)
# Print the coefficients DataFrame
```

```
coefs per stock
# Select a sample of factor returns
samples = factors returns.loc[factors returns.Symbol ==
factors returns.Symbol.unique()[0]]['Close']
# Plot the kernel density estimate for the sample
samples.plot.kde()
# Select factor returns for the first three unique symbols
f 1 = factors returns.loc[factors returns.Symbol ==
factors returns.Symbol.unique()[0]]['Close']
f 2 = factors returns.loc[factors returns.Symbol ==
factors returns.Symbol.unique()[1]]['Close']
f 3 = factors returns.loc[factors returns.Symbol ==
factors returns.Symbol.unique()[2]]['Close']
# Print the sizes of the factor return series
print(f_1.size, len(f_2), f_3.size)
# Calculate correlation between the factor return series
pd.DataFrame({'f1': list(f 1)[1:1040], 'f2': list(f 2)[1:1040], 'f3':
list(f 3) }).corr()
# Calculate covariance matrix for factor returns
factors returns cov = pd.DataFrame({'f1': list(f 1)[1:1040], 'f2':
list(f 2)[1:1040], 'f3': list(f 3)}).cov().to numpy()
# Calculate mean for factor returns
factors returns mean = pd.DataFrame({'f1': list(f 1)[1:1040], 'f2':
list(f 2)[1:1040], 'f3': list(f 3)}).mean()
# Import multivariate normal function from numpy.random
from numpy.random import multivariate normal
# Generate random samples from the multivariate normal distribution
multivariate normal(factors_returns_mean, factors_returns_cov)
# Broadcast the coefficients DataFrame and other variables across the
cluster
b coefs per stock = spark.sparkContext.broadcast(coefs per stock)
b feature columns = spark.sparkContext.broadcast(feature columns)
b factors returns mean = spark.sparkContext.broadcast(factors returns mean)
b factors returns cov = spark.sparkContext.broadcast(factors returns cov)
# Define parallelism and number of trials
from pyspark.sql.types import IntegerType
parallelism = 1000
num trials = 1000000
base seed = 1496
# Create a PySpark DataFrame with seeds
seeds = [b for b in range(base seed, base seed + parallelism)]
seedsDF = spark.createDataFrame(seeds, IntegerType())
seedsDF = seedsDF.repartition(parallelism)
# Import necessary libraries
import random
```

```
from numpy.random import seed
from pyspark.sql.types import LongType, ArrayType, DoubleType
from pyspark.sql.functions import udf
# Define a function to calculate trial returns
def calculate trial return(x):
    trial return list = []
    for i in range(int(num trials/parallelism)):
        random int = random.randint(0, num trials*num trials)
        seed(x)
        random factors = multivariate normal(b factors returns mean.value,
b factors returns cov.value)
        coefs per stock df = b coefs per stock.value
        returns per stock = (coefs per stock df[b feature columns.value] *
(list(random factors) + list(random factors**2)))
        trial return list.append(float(returns per stock.sum(axis=1).sum()
/ b coefs_per_stock.value.size))
    return trial return list
# Create a User Defined Function (UDF) from the calculate trial return
function
udf return = udf(calculate trial return, ArrayType(DoubleType()))
# Apply the UDF to a PySpark DataFrame to calculate trial returns
from pyspark.sql.functions import col, explode
trials = seedsDF.withColumn("trial return", udf return(col("value")))
trials = trials.select('value',
explode('trial return').alias('trial return'))
# Cache the trials DataFrame for faster access
trials.cache()
# Find the 5th percentile of trial returns
trials.approxQuantile('trial return', [0.05], 0.0)
# Calculate the average of the lowest 5% trial returns
trials.orderBy(col('trial return').asc()).
limit(int(trials.count()/20)).
agg(fun.avg(col("trial return"))).show()
# Convert trials DataFrame to pandas DataFrame and plot the distribution of
returns
import pandas
mytrials=trials.toPandas()
mytrials.plot.line()
```

MOVIE RECOMMENDOR

```
# Import necessary libraries
import os
import sys
import pyspark as ps
import warnings
from pyspark.sql import SQLContext
# Set environment variables for PySpark
os.environ['PYSPARK PYTHON'] = sys.executable
os.environ['PYSPARK DRIVER PYTHON'] = sys.executable
# Try to create a SparkContext, or print a warning if it already exists
    # Create SparkContext using all available CPUs
    sc = ps.SparkContext('local[*]')
    # Uncomment the line below if you also need SQLContext
    # sqlContext = SQLContext(sc)
   print("Just created a SparkContext")
except ValueError:
    warnings.warn("SparkContext already exists in this scope")
# Import unittest library for writing and running tests
import unittest
import sys
# Define a test case class for testing RDD operations
class TestRdd(unittest.TestCase):
    # Define a test method for the 'take' operation on RDDs
    def test take(self):
        # Create an RDD containing the numbers 1, 2, 3, 4
        input = sc.parallelize([1, 2, 3, 4])
        # Test if calling take(4) on the RDD returns [1, 2, 3, 4]
        self.assertEqual([1, 2, 3, 4], input.take(4))
# Define a function to run the tests
def run tests():
    # Load the test case suite from the TestRdd class
    suite = unittest.TestLoader().loadTestsFromTestCase(TestRdd)
    # Run the tests and display results with verbosity level 1
    unittest.TextTestRunner(verbosity=1, stream=sys.stderr).run(suite)
# Call the run tests function to execute the tests
run tests()
import json
# Define the fields for different types of entries in the JSON data
fields = ['product id', 'user id', 'score', 'time']
fields2 = ['product id', 'user id', 'review', 'profile name',
'helpfulness', 'score', 'time']
fields3 = ['product_id', 'user_id', 'time']
```

```
fields4 = ['user id', 'score', 'time']
# Function to validate if all required fields are present in an entry
def validate(line):
    for field in fields2: # Check against fields2 as it seems to be the
most detailed set
        if field not in line:
            return False
    return True
# Load the raw reviews data from a file and filter it based on validation
reviews raw = sc.textFile('data/movies.json')
reviews = reviews raw.map(lambda line: json.loads(line)).filter(validate)
reviews.cache() # Cache the RDD for efficiency
# Count the number of unique movies, users, and total entries in the
dataset
num movies = reviews.groupBy(lambda entry: entry['product id']).count()
num users = reviews.groupBy(lambda entry: entry['user id']).count()
num entries = reviews.count()
print(str(num entries) + " reviews of " + str(num movies) + " movies by " +
str(num users) + " different people.")
# Calculate average number of movies watched per user and top movies by
watch count
r1 = reviews.map(lambda r: ((r['product id'],), 1)) # Map each movie to
avg3 = r1.mapValues(lambda x: (x, 1)).reduceByKey(lambda x, y: (x[0] + reduceByKey))
y[0], x[1] + y[1])) # Reduce to sum and count
avg3 = avg3.filter(lambda x: x[1][1] > 20) # Filter movies with less than
20 views
avg3 = avg3.map(lambda x: ((x[1][0] + x[1][1],),
x[0])).sortByKey(ascending=False)  # Sort by view count
# Print top movies by watch count
for movie in avg3.take(10):
    print("http://www.amazon.com/dp/" + movie[1][0] + " WATCHED BY : " +
str(movie[0][0]) + " PEOPLE")
# Calculate average number of movies watched by each user and top users by
movie count
r2 = reviews.map(lambda ru: ((ru['user id'],), 1)) # Map each user to
count 1
avg2 = r2.mapValues(lambda x: (x, 1)).reduceByKey(lambda x, y: (x[0] +
y[0], x[1] + y[1])) # Reduce to sum and count
avg2 = avg2.filter(lambda x: x[1][1] > 20) # Filter users with less than
20 reviews
avg2 = avg2.map(lambda x: ((x[1][0] + x[1][1],),
x[0])).sortByKey(ascending=False)  # Sort by review count
# Print top users by movie count
for user in avg2.take(10):
    print("User " + user[1][0] + " WATCHED : " + str(user[0][0]) + "
MOVIES")
# Filter entries with "George" in the profile name and print details
filtered = reviews.filter(lambda entry: "George" in entry['profile name'])
```

```
print("Found " + str(filtered.count()) + " entries.n")
for review in filtered.collect():
    print("Rating: " + str(review['score']) + " and helpfulness: " +
review['helpfulness'])
    print("http://www.amazon.com/dp/" + review['product id'])
    print(review['summary'])
    print(review['review'])
   print("n")
# Get best and worst rated movies
reviews by movie = reviews.map(lambda r: ((r['product id'],), r['score']))
# Calculate average rating per movie
avg = reviews by movie.mapValues(lambda x: (x, 1))
    .reduceByKey(lambda x, y: (x[0] + y[0], x[1] + y[1]))
avg = avg.filter(lambda x: x[1][1] > 20)
avg = avg.map(lambda x: ((x[1][0] / x[1][1],), x[0]))
    .sortByKey(ascending=True)
# Print top movies by watch count
for movie in avg.take(10):
    print("http://www.amazon.com/dp/" + movie[1][0] + " Rating: " +
str(movie[0][0]))
# Convert RDD to Pandas DataFrame for time series analysis
from datetime import datetime
timeseries rdd = reviews.map(lambda entry: {'score': entry['score'],
datetime.fromtimestamp(entry['time'])})
# Sample the data for visualization
sample = timeseries rdd.sample(withReplacement=False, fraction=20000.0 /
num entries, seed=1134)
timeseries = pd.DataFrame(sample.collect(), columns=['score', 'time'])
print(timeseries.head(3))
# Set up time series analysis and plot
timeseries.score.astype('float64')
timeseries.set index('time', inplace=True)
Rsample = timeseries.score.resample('Y').count()
Rsample.plot()
Rsample2 = timeseries.score.resample('M').count()
Rsample2.plot()
Rsample3 = timeseries.score.resample('Q').count()
Rsample3.plot()
# Visualize histograms for average rating of movies, number of movies
reviewed by users, and movies reviewed by number of users
for movie in avg.take(4):
    plt.bar(movie[1][0], movie[0][0])
    plt.title('Histogram of 'AVERAGE RATING OF MOVIE'')
    plt.xlabel('MOVIE')
    plt.ylabel('AVGRATING')
for movie in avg2.take(3):
    plt.bar(movie[1][0], movie[0][0])
   plt.title('Histogram of 'NUMBER OF MOVIES REVIEWED BY USER'')
plt.xlabel('USER')
```

```
plt.ylabel('MOVIE COUNT')
for movie in avg3.take(4):
   plt.bar(movie[1][0], movie[0][0])
    plt.title('Histogram of 'MOVIES REVIEWED BY NUMBER OF USERS'')
    plt.xlabel('MOVIE')
    plt.ylabel('USER COUNT')
# Build recommendation model using ALS
from pyspark.mllib.recommendation import ALS
from numpy import array
import hashlib
import math
# Function to hash strings
def get hash(s):
    return int(hashlib.sha1(s).hexdigest(), 16) % (10 ** 8)
# Prepare ratings data
ratings = reviews.map(lambda entry:
tuple([get hash(entry['user id'].encode('utf-8')),
get hash(entry['product id'].encode('utf-8')),
                                           int(entry['score'])]))
# Train ALS model
rank = 20
numIterations = 20
model = ALS.train(train data, rank, numIterations)
# Evaluate the model on test data
unknown = test data.map(lambda entry: (int(entry[0]), int(entry[1])))
predictions = model.predictAll(unknown).map(lambda r: ((int(r[0]),
int(r[1])), r[2]))
true and predictions = test data.map(lambda r: ((int(r[0]), int(r[1])),
r[2])).join(predictions)
# Analyze word frequencies for sentiment analysis
min occurrences = 10
good reviews = reviews.filter(lambda line: line['score'] == 5.0)
bad reviews = reviews.filter(lambda line: line['score'] == 1.0)
# Extract words and calculate frequencies
good words = good reviews.flatMap(lambda line: line['review'].split(' '))
num good words = good words.count()
good_words = good_words.map(lambda word: (word.strip(), 1))
    .reduceByKey(lambda a, b: a + b)
    .filter(lambda word count: word count[1] > min occurrences)
# Calculate word frequencies
frequency_good = good_words.map(lambda word: ((word[0],), float(word[1]) /
num good words))
# Join frequencies and analyze relative differences
joined frequencies = frequency good.join(frequency bad)
result = joined frequencies.map(lambda f: ((relative difference(f[1][0],
f[1][1]),), f[0][0]))
```

```
.sortByKey(ascending=False)

# Print top expressions for positive and negative reviews
for movie in result.take(50):
   plt.bar(movie[1], movie[0][0])
   plt.title('Histogram of 'SENTIMENT ANALYSIS'')
   plt.xlabel('WORD')
   plt.ylabel('NUMBER OF OCCURRENCES')
```