

Detecting Customer Purchasing Patterns using Association Rule Mining

MSc Research Project Data Analytics

Rhutujit Uday Paradkar Student ID: 20187416

School of Computing National College of Ireland

Supervisor: Dr. Paul Stynes Dr. Anu Sahni

National College of Ireland Project Submission Sheet School of Computing



Student Name:	Rhutujit Uday Paradkar		
Student ID:	20187416		
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Configuration Manual Detecting Customer Purchasing Patterns using Association Rule Mining

Rhutujit Uday Paradkar 20187416

1 Introduction

The main aim of creation of this document is to give all the steps that are essential for reproducing the research in the future. The document acts as a technical manual for reproducing all the research related to the association rule mining , Kmeans clustering and the principal component analysis. This manual consists of the screenshots of the essential parts of the code so that the understanding of the code becomes easier.

2 Prequisites: Hardware and Software

The software and the hardware requirements that are needed for completing the research are mentioned below:

2.1 Hardware Requirements

- 1. **Operating System** : Windows 10 and above
- 2. Processor : Intel Core i7 7th Generation and above
- 3. **RAM** : 8 GigaBytes
- 4. **GPU** : Geforce GTX 1050 Ti
- 5. GPU Memory : 4 Gigabytes
- 6. Hard Disk : 1 Terra bytes
- 7. Solid state drive : 256 Gigabytes

3 Software Requirements

The Integrated Development environment or the IDE that was used for the research were:

- 1. Jupyter notebook for running the python code
- 2. R studio for running the R code

Anaconda software was used as a main controlling software for installing and controlling all the IDE's mentioned above.



3.1 Installing Anaconda Navigator

Figure 1: Website to download windows version of anaconda

The site www.anaconda.com has the links to install the anaconda for the following operating systems:

- 1. iOS
- 2. Windows
- 3. Linux

As the system that we are using is Windows we are using the windows version of anaconda.

3.2 Installing and Launching the Jupyter and the R notebooks from Anaconda

In this section, we will take a look at the process to install and run the Jupyter and the R notebooks. In some cases the Anaconda navigator comes preinstalled with Jupyter notebook and you just need to run it.

Home	Applications on base (root) Channels			Refres	
Candranamate	-				
- Cavironmencs	jupyterlab	notebook	qtconsole	spyder	
	An extensible environment for interactive	Web-based, interactive computing	PyQt GUI that supports inline figures,	Scientific PYthon Development	
Projects (beta)	and reproducible computing, based on the Jupyter Notebook and Architecture.	notebook environment. Edit and run human-readable docs while describing the	proper multiline editing with syntax highlighting, graphical calltips, and more.	EnviRonment. Powerful Python IDE with advanced editing, interactive testing,	
		data analysis.		debugging and introspection features	
Learning					
	Launch	Launch	Launch	Launch	
Community					
			•		
	l i	<u> </u>			
	d Ba	00	R		
	glueviz	orange3	rstudio		
Documentation	0.12.0	3.4.1	1.1.383		
	files. Explore relationships within and	Data visualization and data analysis for	you be more productive with R. Includes R		
Developer Blog	among related datasets.	novice and expert. Interactive workflows	essentials and notebooks.		
		with a brige coupon.			

Figure 2: Installation of R studio in the anaconda distribution

Once you click on the install button, it will take some time to install the R distribution and the progress will be shown in the bottom right of the main installation window. This process is same for the Jupyter notebook as well.



Figure 3: Installation of Jupyter in the anaconda distribution

4 Data Pre-processing

The libraries that are needed for the projects in this research paper are given below. To first implement various preprocessing techniques, it is first necessary to import the files into the code.

4.1 Experiment 1 - Market basket Analysis(Importing the libraries)

The file format of this dataset is csv. The pandas library is used for the preprocessing and the read_csv function is used to read the csv file and import the data into the jupyter notebook.



Figure 4: Code to read the csv and import the data

For applying the apriori algorithm to the dataset, the data needs to bee in the form of the list. That is why some preprocessing is needed to convert the data to a list . The figure 5 shows the code to convert the data to a list.



Figure 5: Converting the data to a list

4.2 Experiment 2 - Importing the dataset and the libraries

Figure 6: Importing data for the Glantus dataset

- 1. In this experiment the data is imported by using the inbuilt read_csv function of R
- 2. The nested json is converted to simplified version by using the jsonlite library of the R

```
library(arules)
library(arulesViz)
library(cluster)
library(cluster.datasets)
library(clValid)
library(clustree)
library(corrplot)
library(cowplot)
library(dendextend)
library(dplyr)
library(factoextra)
library(FactoMineR)
library(ggfortify)
library(GGally)
library(ggiraphExtra)
library(jsonlite)
library(knitr)
library(kableExtra)
library(lubridate)
library(magrittr)
library(NbClust)
library(RColorBrewer)
library(tidyverse)
```

Figure 7: Importing the necessary libraries in python for the instacart dataset

The figure 8 shows the necessary libraries that are need to perform the research on the glantus dataset. For installing the necessary libraries please follow the following command

1. install.packages(package name)

4.3 Experiment 3 - Importing the data and installing required packages

The datasets for the 3rd experiment are from the instacart dataset which is taken from kaggle. The dataset consists of multiple files combined together in a zip. Therefore, for importing the dataset the zipfile package from python is used which unzips the files so that we can get the individual csv files.

```
In [2]:
import pandas as pd
import numpy as np
from sklearn.cluster import KMeans
#For dimensionality reduction.
from sklearn.decomposition import PCA
from sklearn.neeprocessing import StandardScaler
from sklearn import decomposition
Xmatplotlib inline
import matplotlib.pyplot as plt
from pylab import rcParams
import seadorn as sb
rcParams['figure.figsize'] = 12, 4
sb.set_style('whitegrid')
np.random.seed(42)
```

Figure 8: Importing the necessary libraries in the R studio

```
import zipfile # Unzips the files
from subprocess import check_output
#Prior Dataset
with zipfile.ZipFile("instacart-market-basket-analysis/"+prior+".zip","r") as z:
   z.extractall("."
prior = pd.read_csv("order_products_prior.csv")
#Order_Train Dataset.
with zipfile.ZipFile("instacart-market-basket-analysis/"+order_train+".zip","r") as z:
   z.extractall(".")
order_train = pd.read_csv("order_products__train.csv")
#Orders Dataset.
with zipfile.ZipFile("instacart-market-basket-analysis/"+orders+".zip","r") as z:
   z.extractall(".")
orders = pd.read_csv("orders.csv")
#Products
with zipfile.ZipFile("instacart-market-basket-analysis/"+products+".zip","r") as z:
  z.extractall(".")
products = pd.read_csv("products.csv")
#Aisles
with zipfile.ZipFile("instacart-market-basket-analysis/"+aisles+".zip","r") as z:
   z.extractall(".")
aisles = pd.read_csv("aisles.csv")
#Departments
with zipfile.ZipFile("instacart-market-basket-analysis/"+departments+".zip","r") as z:
   z.extractall(".")
departments = pd.read_csv("departments.csv")
```

Figure 9: Code to unzip the files and get the individual csv files

The code in the figure 9 unzips all the files at the given path by passing the '.' parameter which means all files at the givenpath. The 'r' parameter of the ZipFile reads the raw string as it is.

```
In [5]:
combined_df_list = [products,orders, departments, aisles, prior, order_train]
In [6]:
#Check the size of the datasets.
for i in combined_df_list:
    print (i.shape)
del combined_df_list
    (49688, 4)
    (3421083, 7)
    (21, 2)
    (134, 2)
    (32434489, 4)
    (1384617, 4)
```



The code in the figure 10 is used to get an idea about the number of records in each

dataset.

4.4 Exploratory data analysis



Figure 11: Popular products in the dataset



Figure 12: Popular products in the dataset

The figures 12and 11 show the most popular items in the datasets.



Figure 13: Heatmap for the instacart data

The seaborn library is used to generate the heatmap shown in the fig.13.

Γ

pd.DataFra	pd.DataFrame(combined_df.groupby('user_id')['product_id'].count()).sort_values('product_id', ascending=False).head(2)			
#User_id =	#User_id = 142131			
product_id				
user_id				
142131	176			
169550	161			

Figure 14: Most popular user

The groupby clause is used here to get the user id of the customer who spends the most in the store.

In [21]:	<pre>pd.DataFrame(combined_df['aisle'].value_counts()).head(5</pre>		
Out[21]:		aisle	
	fresh fruits	56326	
	fresh vegetables	52073	
	packaged vegetables fruits	27347	
	yogurt	22485	
	packaged cheese	14960	

Figure 15: The aisle with the most products

The above code is used to get the aisle with the most number of products. The pandas library was used here.

In [20]:	: pd.DataFrame(combined_df['product_name'].value_counts()).he		
Out[20]:	pro	duct_name	
	Banana	7365	
	Bag of Organic Bananas	5920	
	Organic Strawberries	4023	
	Organic Baby Spinach	3797	
	Organic Hass Avocado	3317	

Figure 16: Most selling products

The code in the figure 16 is used to get the most frequent items in the dataset.

5 Implementation

The implementation section of the manual describes the different modelling techniques that are performed on the datasets in the research paper. The screenshots in the section show the code that was used to implement the modelling techniques.

```
In [5]:
             from apyori import apriori
             rules = apriori(transactions= transactions, min_support = .003, min_confidence = 0.25, min_lift = 2, min_length = 2, max_length
             Visualising the results
             Displaying the first results coming directly from the output of the apriori function
In [6]: results = list(rules)
In [7]: results
Out[7]: [RelationRecord(items=frozenset({'burgers', 'almonds'}), support=0.005199306759098787, ordered_statistics=[OrderedStatistic(ite
ms_base=frozenset({'almonds'}), items_add=frozenset({'burgers'}), confidence=0.25490196078431376, lift=2.923577382023146)]),
RelationRecord(items=frozenset({'bacon', 'spaghetti'}), support=0.003199573390214638, ordered_statistics=[OrderedStatistic(ite
             tic(items_base=frozenset({'blueberries'}), items_add=frozenset({'spaghetti'}), confidence=0.37681159420289856, lift=2.164214217
             546663)]),
RelationRecord(items=frozenset({'body spray', 'french fries'}), support=0.004266097853619517, ordered_statistics=[OrderedStati
              stic(items_base=frozenset({'body spray'}), items_add=frozenset({'french fries'}), confidence=0.37209302325581395, lift=2.177121
              50346479)]),
             Sobio4/9/j);
RelationRecord(items=frozenset({'cereals', 'milk'}), support=0.007065724570057326, ordered_statistics=[OrderedStatistic(items_
base=frozenset({'cereals'}), items_add=frozenset({'milk'}), confidence=0.2746113989637306, lift=2.119197637476279)]),
RelationRecord(items=frozenset({'light cream', 'chicken'}), support=0.004532728969470737, ordered_statistics=[OrderedStatistic
(items_base=frozenset({'light cream'}), items_add=frozenset({'chicken'}), confidence=0.29059829059829057, lift=4.8439506172839
             S)]),
RelationRecord(items=frozenset({'chocolate', 'tomato sauce'}), support=0.005065991201173177, ordered_statistics=[OrderedStatis
              tic(items_base=frozenset({'tomato sauce'}), items_add=frozenset({'chocolate'}), confidence=0.3584905660377358, lift=2.187988393
             6932925)]),
               RelationRecord(items=frozenset({'eggs', 'cider'}), support=0.004266097853619517, ordered_statistics=[OrderedStatistic(items_ba
             RelationRecond(items=trozenset({ eggs , cloer }), support=0.0042009705003917, ordered_statistics[orderedStatistics]orderedStatistics[clems_oa
se=frozenset({ cider '}), items_add=frozenset({ 'eggs '}), confidence=0.40506032911392405, lift=2.2539909101153137)]),
RelationRecond(items=frozenset({ 'milk', 'cider '}), support=0.00332288044140248, ordered_statistics=[OrderedStatistic(items_ba
se=frozenset({ 'cider '}), items_add=frozenset({ 'milk'}), confidence=0.31645569620253167, lift=2.4421133510444344)]),
RelationRecond(items=frozenset({ 'escalone'. 'mushroom cream sauce'}), supnort=0.005732568090801226, ordered statistics=[OrderedStatistics]),
```

Figure 17: Applying apriori on the dataset

The code in the figure 17 is used to generate the apriori rules and the package used for that is 'apyori'.

```
writing ... [1328 set(s)] done [0.01s].
Creating S4 object ... done [0.00s].
> eclat_rules <- ruleInduction(rules, marketdata, confidence = 0.2,lift=2)</p>
> inspect(sort(eclat_rules, by = 'lift')[1:10])
     1hs
                                                                            confidence lift
                                                rhs
                                                                support
                                                                                                itemset
                                             => {olive oil}
[1] {mineral water, whole wheat pasta}
                                                                0.003866151 0.4027778 6.115863 296
[2] {frozen vegetables, milk, mineral water} => {soup}
                                                                0.003066258 0.2771084 5.484407 615
[3] {fromage blanc}
                                                                0.003332889 0.2450980 5.164271 43
                                             => {honey}
[4] {spaghetti, tomato sauce}
                                             => {around beef}
                                                               0.003066258 0.4893617 4.980600 58
[5] {light cream}
                                             => {chicken}
                                                                0.004532729 0.2905983 4.843951 89
[6] {pasta}
                                             => {escalope}
                                                                0.005865885 0.3728814 4.700812 25
[7] {french fries, herb & pepper}
                                             => {around beef}
                                                               0.003199573 0.4615385 4.697422 540
[8] {cereals, spaghetti}
                                             => {ground beef}
                                                                0.003066258 0.4600000 4.681764 247
[9] {frozen vegetables, mineral water, soup} => {milk}
                                                                0.003066258 0.6052632 4.670863 615
[10] {french fries, ground beef}
                                             => {herb & pepper} 0.003199573 0.2307692 4.665768 540
×
```

Figure 18: Applying eclat on the mba dataset

The 'arules' package is used for applying the eclat algorithm on the mba dataset. The inspect function is used to sort the generated rules and then display it. The rules are sorted by the decreasing value of the lift.

5.1 Principal component analysis on the instacart dataset



Figure 19: Elbow method for the PCA

The figure shows the line plot and the code that is used to find the optimal number of clusters for the data.



Figure 20: Bar graph variance

This code is used to plot a bar graph that shows the variance that is explained by the number of selected features by performing the principal component analysis.

6 Conclusion

In the conclusion, the manual can be used to replicate the research that is done in the research. The steps and the code are explained in the manual by actual screenshots from the code.