Health Records Analysis to build Final Patient dataset for dementia disease

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Abstract

Big Data and data mining technologies are becoming an essential technology in everyday life. Banking uses these tools to monitor the financial market through network activity monitors to minimize fraudulent transition. Automotive uses these tools to reduce the implementation of opt having an underperforming market. The Healthcare sector, however, still has not benefitted from the wide-scale use of Big Data technologies. The reason for it is the Healthcare sector has been traditionally slow in adopting new ICT techniques because most clinical data was stored in paper form. The distribution of Electronic health record (EHR) in healthcare facilities, it was an important step towards digitalization of health.

Using the current hospital IT system is impossible to use data to make the clinical decisions because more of that has been collected in a wrong way or rather to achieve operational purposes. Dementia disease is one of the most widespread disease in the world among older people. Someone in the world develops dementia every 3 seconds. Around 50 million people have dementia, and there are nearly 10 million new cases every year. The most common form of dementia is Alzheimer disease that is associated with 60-70% of cases. The focus of this master thesis is dementia disease using anonymized data collecting various information during the disease progression. We present a data-driven approach to build the main dataset called Final Patient Data and with it define a set of three key performance indicators that it a preliminary step of a more complex machine learning analysis: 1) Multiple Filter Analysis 2) identification of strong correlation 3) rate of progression after the first diagnosis.

In order to achieve the goal this thesis presents a method to be able, first of all, to extract processes of the patient from unstructured data sets. To structure the project and make it replicable, CRISP-DM has been adopted as a methodology to fulfill the goals. The thesis also presents methods to analyze, clean and prepare data to obtain structured datasets from which the mentioned KPIs can be measured.
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1 Introduction and Motivation

1.1 Introduction

Data are one of the most useful resources for a huge variety of productive sectors. Data into computer science meaning is an elementary description of an information, an entity, a phenomenon or an event[4].

With the advent of technology, there has been an impressive increase of data and new branches of science were born with the goal to exploit it. An IDC report[5] said that global data traffic will reach 175 zettabytes by 2025. This uncontrollable data traffic increase is due to the spread of smart devices. In 2020 connected smart devices are 30 billion, they are estimated to be 75 billion by 2025 thanks to a trend in constant and exponential growth.

Another IDC report said that now all production sectors make full use of data to carry out analysis, statistical studies and so on. This report said that health care is growing the usage of data science tools, in fact it is assumed that there will be 36% growth in generated healthcare data by 2025. The growth in accumulated data is due to the integration of IT systems within hospitals and the use of the Electronic Health Record (EHR).

This increase in the healthcare sector has spread the belief of using it to extract new knowledge from the data. The idea is to exploit all the tools provided by big data, machine learning and deep learning techniques in order to provide a powerful mean to improve the diagnoses, the treatments provided and their effectiveness for identifying the risk factors associated to a specific type of illness or to translate the acquired knowledge into new hospital policies to reduce costs and waste of resources.

One of the many diseases that we want to study using the methods provided by data science is dementia that is an extremely common disease spread through the human population. In Italy in 2019 there are 1,271,000 people suffering from dementia which in 2050 it is expected to double and in the world it is estimated to have 48.8 million dementia sufferers, where more than 60% are affected by Alzheimer’s disease.

The promoter of this study is the Oxford Project to investigate Memory and Aging (OPTIMA)[6] that has the aim of identifying which are the still unknown causes of dementia and in particular Alzheimer.
The guidelines published by the World Health Organization (WHO)\[7\] on May 14 2019 emphasize how a change in lifestyles (such as smoking habits, excessive alcohol consumption, unbalanced nutrition) or the control of certain diseases (such as hypertension, diabetes, obesity, depression, hypercholesterolemia) and factors that are not strictly sanitary (such as social isolation and cognitive stimuli) may be implicated in the onset of dementia and, in general, of cognitive impairment.

This master thesis was developed as part of the OPTIMA project in order to build a reference dataset that have been called Patient Final Data and carry out a set of analysis on it.

### 1.2 State of The Art

In the past, several studies have been done about dementia and its influencing and risk factors. During this study, a different type of features that are perfectly associated with these past studies have been considered.

In the study from E.S Sharp e.t \[8\] has been studied the relationship between Dementia and education where has been identified that lower education is associated with a greater risk for dementia in many but not all patients. Another study from C.C. Liu e.t \[9\] showed that a specific type of gene, allele E4, provide clinical evidence of the relationship between it and the spread of Alzheimer's disease into the population. Although the presence of APOE E4 does not necessarily entail disease development, this genetic isoform probably accelerates the rate of disease conversion and progression.

Lastly, study from T.B. Chen e.t. \[10\] analyzed the relation between dementia and comorbidities. Regardless of the cognitive condition, people that have more than 60 years have at least one comorbidity disease. The study shows that people who have more than 60 years and at least three different types of comorbidities, have higher possibility to developing cognitive impairment problems. All the features used in these studies have been included in the Patients Final Data created in this project.

### 1.3 Goals

Starting from unstructured clinical data it will be used Data Science techniques to obtain the following goals:
Multiple Filter Analysis

Correlation Analysis on Patients Final Data

Rate of progression after first diagnosis

In order to fulfil these goals, the following steps must be followed:

- Exploration OPTIMA Data Source
- Extraction patients information from the relevant files into OPTIMA Data Source
- Doctor’s rules application.

1.4 Structure of Contents

Project has been divided into the following chapters:

- Chapter 2 Theoretical Framework: It describes the background of this study and explains the main tools used to develop it. Furthermore it describes the primary tools used to develop a Data Science Project and which is the main reason of the study.

- Chapter 3 Materials and Methods: A general overview of the raw OPTIMA Data Source and the process, called Data Pipeline, to build the Patient Final data is describe.

- Chapter 4 Proposed approach to extract pattern of dementia evolution: This is the main chapter on this project. This Chapter, starting from the business project analysis, will describe the primary Key Performance Indicators (KPI) developed. Then there is a complete description about Raw Data and the approach used to extract data from it to create Patients Final Data. Furthermore, the Doctor’s rules have been described and how to apply it and finally there is a phase of validation of results.

- Chapter 5 Discussion of Results: Chapter describe how the KPI defined into the previous chapter, are been computed and then a discussion of results obtained.
Chapter 6 Conclusions and Future Lines of Research: We presented the main conclusions and we have been proposed possible ways to improve the accuracy of the result obtained.
2 Theoretical Framework

In this chapter detailed description of dementia disease and what is Optima project have been presented to understand which are the main goals of this ambitious project. Furthermore, During the second part of the chapter, a theoretical explanation has been shown about the key performance indicators and how they have been defined and finally how a data science project is structured.

2.1 Dementia

Dementia is a descriptive term indicating an observable decline in mental abilities. It is an acquired clinical syndrome characterised by deterioration of mental functioning in its cognitive, emotional and conative aspects[11].

In the collective imagination, dementia is directly associated to Alzheimer’s disease but this idea is totally wrong. In fact there are over 100 diseases that may cause dementia each one associated with a set of specific causes. In general, it results from neurons degeneration due to disturbances of body systems

The following list shows which are the common causes of dementia:

- **Alzheimer’s disease**: Type of dementia that is totally irreversible. Alzheimer outlines a progressive brain disorder that destroys memory and thinking skills.

- **Parkinson’s disease**: The disease damages nerve cells producing dopamine into the black substance area of the brain. Dopamine has many different tasks to do, such as impacting thought processes, mental functions and memory management.

- **Vascular dementia**: It is a cognitive impairment problem caused by brain damage due to impaired blood flow to the brain. The main causes of vascular dementia are diseases like stroke that blocks an artery in the brain or conditions that damage blood vessels or blood circulation.

- **Frontotemporal dementia**: Type of dementia associated with a progressive nerve cell loss into frontal and temporal lobes. Normally FTD leads to loss of important cognitive functions.
• **Cortical Lewy Body dementia:** Type of dementia normally associated with a Lewy bodies disease.

Among the listed types of dementia, one is most common than the others, as the graph from the University of Queensland reports (figure 1).

![Figure 1: Distribution Dementia Types](image)

Among the listed types of dementia, one is most common than the others, as the graph from the University of Queensland reports (figure 1).

to understand if a patient has one of dementia diseases several different tests exist and they can be splitted into three macro categories: Neurological exams, Mental status tests and brain imagining exams. In mental status tests patients' thinking skills are tested during the resolution of simple problems. Two common types of mental status tests are the MMSE.
and Mini-Cog test. This project used MMSE test consisting into a 30-point questionnaire. This method is extensively used in clinical research to measure cognitive impairment through simple targeted questions and small graphic tasks to test the domain of different brain functions, such as orientation (autopsychic and outward), memory, attention and calculation, the ability to recall certain acquisitions and language.

2.2 Alzheimer and OPTIMA DATASET

As showed into the figure 2.1, Alzheimer’s disease is the most common cause of dementia. It is a disease that causes a progressive loss of memory and a lowering of cognitive abilities limiting the normal activities of daily life.

Alzheimer has no cure but just a set of treatments to slow down the disease are known. Scientists haven’t found Alzheimer’s causes but only a partial set of hypothesis. Some studies claim that the most common risk factors for dementia amenable to prevention are those associated with cerebrovascular disease[12].

Another study says that the most common cause of dementia is $\beta$-amyloid:it states neural death appears to be associated with intracellular calcium homeostasis and subsequent production of free radicals by calcium-sensitive enzymes[13].

This project focused on trying to extend research studies of the causes of Alzheimer’s and in general of all types of dementia. To developed it, the data has been provided by the Oxford Project to Investigate Memory and Ageing(OPTIMA) [14].

OPTIMA collect a huge amount of data on both cognitively impaired and cognitively normal elderly people during life.

Inside informations, can be splitted into the following subcategory:

- Medical history and physical examination.
- Neuropsychological assessments.
- Brain scans.
- Urine and csf samples.
- Histopathological information following brain donation.
- DNA
2.3 Key Performance Indicator (KPI)

A Key Performance Indicator (KPI) is a measurable value used to gauge the performance of a process and identify the best practices. It measures the trend of a business process and helps making decisions to improve the outcomes[15]. Each company can use a set of metrics to track the status of a specific process. With KPIs, the company focuses the attention to one or more metric to become a point of reference to identify progresses. The administration is in charge of defining business goals for its company, then, a so-called business analyst will analyze business goals to define KPIs. However, not all processes can be analyzed with KPIs and in general, the opportunity to analyze a process using them is measured with a robustness scale, which considers[16]:

- **Ease of understanding** - KPIs must explain what it is measuring at the minimum possible grain level
- **Cost of data** - Collecting data to enable analysis on a KPI costs. These costs can include collecting data, structuring data and experts on interpret results
- **Meaningfulness** - KPIs must reflect only critical goals i.e. the ones which give the company an improvement on quality of processes
- **Frequency of data change** - the more informative the indicator is with time, the more robust will be

During the development of this thesis some performance indicators have been defined to optimize the analysis on the dataset and identify the best practices.
2.4 Data Science

Data science is one of the scientific disciplines having an enormous growth during the 21st century. This field of science consists of using computer facilities, mathematics and statistics models to extract information and knowledge from unstructured and structured data. In its first application, data science has been used solely to building statistics but it has rapidly evolved becoming one of the pillars of today’s society. We find this discipline in areas as artificial intelligence, machine learning and Internet of things. With the advent of Big Data has been applied in more traditional fields like medicine, engineering and social science.

Many different dates can be used to track Data Science’s slow growth and its impact on the data management industry. In 1996, the term data mining became popular thank to an article called From Data Mining to Knowledge Discovery in Databases[17] where the overall process of discovering useful information from data has been explained.

As already mentioned, this multi-disciplinary field broke out during the early 2000s with the advent of the Internet era. In the age of Internet increased suddenly the amount of data and this phenomenon led to the creation of the term Big Data. The term Big data indicates the usage of large amount of information to perform analysis and extraction of new knowledge.

However, data are often stored without any planning for future analysis and leaving their potential knowledge hidden. Then if we want to extract this knowledge, we need to make a great effort to extract their real value. In the next sections, we will see that the most important operation during the data science project is the cleaning and standardization phases using the data wrangling procedure.
2.5 Data Science Project Development

Data Science Project has role to identify, analyze and extract the knowledge of a dataset. During this type of project, the main key is to have excellent communication among the client the company that commissiones the project and the data scientist team. To accomplish an excellent work the developer team need to understand perfectly the context, intermediate objectives and achieve the final goals.

Normally a Data Science project follows a common methodology called Cross-industry standard process for data mining (CISP-DM). CISP-DM is a technique conceived in 1996 defining a guideline to direct data mining efforts. It is composed of the steps shown in the figure below:

![CRISP_DM phases and relationship against its](image)

*Figure 3: CRISP_DM phases and relationship against its [3]*

A chief features of this technique is that it is no a linear process but a cycle process, it is as an idealised sequence of events. Tasks can be performed in a different order and it will often be necessary to backtrack to previous tasks and repeat certain steps again.
2.5.1 Business Understanding

The primary objective of this phase of the project is to translate business goals into data mining goals. Business goals is part of the planning process and normally describe what a company want to accomplish using business terminology, whereas data mining goals describe the same goals but in technical terms.

To realize this translation, we need to follow these steps:

1. Define final output
   - **Set Objectives** - Definition and description of primary objective in business terms.
   - **Project Plan Definition** - Definition and description of the ways to achieve final goals and the phases this procedure is divided into. Project Plan fixes the tools and techniques to be used.
   - **Business Success Criteria** - Definition of criteria to understand if the project plan reaches fulfil the business goals.

2. Evaluate the initial situation
   - **Inventory of Resources** - List the available resources for the project.
   - **Requirements, Assumptions and Constraints** - List of all the project requirements.
   - **Risks and Contingencies** - Identification of possible risks and delays associated to them.
   - **Terminology** - Glossary of terminology relevant to the project.
   - **Costs and Benefits** - Analysis based on the relationship between costs and benefits. The idea is to understand if we have a set of final benefits upper than costs associated to develop the project.

3. Define data mining goals
   - **Business success criteria** - Description of criteria to consider outputs consistent with the achievement of the business objectives.
Data mining success criteria - Description of criteria in order to consider outcomes successful for the projects in technical terms.

4. Produce project plan

- Project plan - List of the stages to be executed in the project, together with their duration, required resources, inputs, outputs, and dependencies.

- Initial assessment of tools and techniques - At the end of the first phase you should undertake an initial assessment of tools and techniques. Here, for example, you select a data mining tool that supports various methods for different stages of the process. It is important to assess tools and techniques early in the process since the selection of tools and techniques may influence the entire project.

2.5.2 Data Understanding

Data understanding phase includes three steps.

Data Collection listed data sources acquired. At the end of this phase it is necessary to develop a report for data description. At this scope, a preliminary descriptive analysis is needed, to extract information. In this project preliminary descriptive analysis is based on following the doctor’s guidelines.

During Data Exploration a deep analysis of acquired data is fulfilled. To make this, we used reporting techniques and data visualization tools that are provided by the programming language used for the project.

Into Report for Data Exploration, consists of listing the results of the exploration of your information. This includes the initial hypothesis and states how they impact the development of the entire project.

2.5.3 Data Preparation

The third stage of the project is one of the most important because we have to decide which information we are going to use in the next phases. In all data science projects the first three phases (business understanding, data understanding and data preparation) account for more than 80% of the entire project efforts. 76% of data scientists say that
data preparation is the worst part of their job[18].
During data preparation, the following steps will be executed:

- **Data Gathering** - It defines the ways how data are provided from the company’s data lake to perform the project. There are different ways to make this as using API, non-relational dataset or using a format like CSV. Into this project, data are provided CSV files. After getting the data, the goal is to identify the right columns in its. This happens using a data catalog or a data guideline where all tables of company’s data lake are listed and for each table a features description is reported to help data scientist to identify right columns. Furthermore, data guideline shows which are data type used and their meaning.

- **Data Cleaning** - It is the most important step in data preparation phase and it is crucial for removing faulty data and filling in gaps. Often data provided, are unstructured then have to apply on it, techniques to manage missing values, errors and to build conforming data to a standardized pattern.

  Data cleaning has to strong impact on the future phases of the project because it defines the quality of data.

- **Data Filtering** - Data filtering refers to removing unnecessary information. the general idea is to remove the "noise" into the data defining specific constraints to apply. The final goal is to obtain a dataset where can perform a better analysis.

- **Data Transformation** - It is a process where the main objective is to apply an updating the format where there are a converting on data types. Then, data transformation wants to achieve a well-defined result to obtain data more understood.

### 2.5.4 Modelling

The modelling phase is based on selecting the specific modelling technique to improve quality of data among the different available in the literature. Before we define a model we need to structure a scheduling procedure to test model’s quality and validity.

Into machine learning world there are many different tools and procedures. The choice is associated to the goal that we want to obtain : Regression, Classification, Clustering, Dimensionality Reduction, Ensemble Methods, Neural Nets and Deep Learning, Transfer
Learning and Reinforcement Learning. After having defined the objective and choosing the tool to use, the following iterative procedures will be followed:

- **Algorithm implementation.**
- **Training** - Data is fed into the algorithm to eventually learn patterns in data.
- **Validation** - The trained algorithm is checked against a validation data set to tune parameters that best fit our data.
- **Testing** - The algorithm is tested against new data to evaluate the abstraction power of the model.

*Figure 4: Modelling Steps Overview*
2.5.5 Evaluation

Evaluation phase checks the obtained results using factors as accurateness, precision and computational effort. The idea is to understand if the model is able to meet the business goals defined during the business understanding phase.

We will do an evaluation of the data mining result considering the final business success standard and create a report on it. Then we will perform the process of reviewing where a more comprehensive review for the data mining process, is developed checking if every task has been overlooked. The idea is to identify if some keys objective haven’t been sufficiently considered.

2.5.6 Deployment

During this phase is defined as the structures through which the data will be presented to the final client is built. The tools chosen to present the results are dashboard, API and web app.
3 Materials and Methods

In this chapter we explain every material and method used to develop this project. The chapter is splitted into two parts:

- The first one explains which data are used to create final dataset called Patient Final Data.

- The second one explains which programming languages them and packages are used to achieve the data science project proposing a general overview of them.

3.1 Raw Data

As mentioned in section 2.2, the data source used to develop this project is based on the OPTIMA data source that provides information on different fields: medical stuff, parent’s history, general information and personal information. All the documents provided by OPTIMA are totally anonymous so that there is no way to have personal data like name, surname or date of birth.

OPTIMA data source have stored patient information since 2009. Every data have been provided in CSV format and it have the following basic features:

- **ID** - It identifies a unique patient.

- **Study ID** - Numerical value that identifies which are study:OPTIMA or Challenge.

- **Gender** - Gender of the patient.

- **DOD** - It identifies the date when data was collected.

- **EPISODE DATE** - It identifies the Date when episode happens.

With OPTIMA data source has been provided an excel file called Variables Guide. Variables Guide is a descriptive file providing us a complete overview about tables presenting the data source and a deep description of each columns. This description contains data types, which are value used and what are meaning. About Variables Guidelines there is an overview into section 4.3.1.
3.2 Patients Final Data

Patients Final Data will be the final output obtained after the first three phases of the CRISP-DM process.

To build Patient Final Data rules have been followed provided by a team of doctors that define in deep the features to use and how to combine them. In general we can say that Patients Final data is the output of the Data preparation phase and can be splitted into the following sub-category:

- **General information** - Global patient id, gender, episode date, age at episode.

- **EHR information** - body max index, weight, comorbidity types, APOE.

- **Lifestyle information** - education level, smoking level, alcohol level.

- **Mini Mental Score information** - mmmse for episode, mmmse discretized, mmmse final.

- **Dementia type** - alzheimer, other dementia, no dementia.

A deep description of the rules and filter applying are shown into the following chapter.

![Figure 5: Final Patient Data Overview](image)

19
3.3 Python and R (Programming Languages)

the main programming language used to develop this project is Python. It is a multi-paradigm language that has among its main objectives: dynamism, simplicity and flexibility [19]. It was conceived in 1980 by Guido van Rossum at Centrum Wiskunde Informatica (CWI) and it is an interpreted, high-level, general-purpose programming language. It is the most powerful language because it is able to integrate a huge variety of programming paradigms and presents a comprehensive standard library.

Python is an open-source language and there is a global community called CPython that provides reference implementation, a complete explanation about the libraries and how to use it.

Python is a language that provides a huge set of tools in the area of statistics. Being a open-source system, many different packages containing a variety of tools, methods and techniques are freely available.

Also there is a huge number of packages available for the machine learning sector and for the data mining world. in this project, many different packages have been used to help with data visualization and data manipulation stage. These packages have been explained in deep in the following paragraphs. Furthermore, during this project R programming language has been used to develop survival analysis. R programming language is also an open-source language that provide good tools to made analysis into a dataset.

3.4 Visualization Packages in Python

Matplotlib.Pyplot

Matplotlib.pyplot is a collection of command style functions having the role to create graphs, figures like Matlab’s styles. Each pyplot function makes some changes to a figure such as creating a figure, creating a plotting area in a figure, figuring some lines in a plotting area, decorating the plot with labels and so on. It is much popular for the data visualization stage using elementary a graph’s style.

Seaborn

Seaborn is a library to achieve statistic graphs based on matplotlib with strong integration with pandas data structure. Pandas is the main library that can be used to perform anal-
ysis with dataframes, datalakes and in general to use data mining and machine learning tools. Seaborn aims to make visualization a primary part of exploration and to understanding data defining semantic mapping and statistical aggregation necessary to build informative graphs.

3.5 Data Manipulation Packages in Python

Pandas

Pandas is a software library to perform operations for data manipulation and analysis. Pandas was created by Wes McKinney in 2008 when he needed to perform quantitative analysis on financial data. It contains data structures for tables and series manipulation and provide the possibility to import a huge variety of file formats such as csv, excel, JSON, ZIP, images and so on. Pandas library allows us to use SQL tools like groupby, merge, concatenation and also tools to make data cleaning and data filtering. It is the main instrument for data manipulation phase in Python language.

NumPy

NumPy is a library to manage multidimensional array objects with a set of routines collection. Furthermore, it allow us to manage a large collection of high-level mathematical functions. The name Numpy is a combination of the two words : num = Numerical Py = Python and was created by Jim Hugunin.

Math

Library that provides a huge set of mathematical functions coded in the C standard. This type of function can be used with all types of existing numbers(complex, real, irrational, etc).

Datetime

Python DateTime, TimeDelta, Strftime(Format) with Examples. In Python, date, time and datetime classes provides a number of function to deal with dates, times and time intervals. Date and datetime are objects in Python, so when you manipulate them, you are actually manipulating objects and not string or timestamps.
4 Proposed approach to extract pattern of dementia evolution

The following chapter explains in detail the whole procedure followed to achieve the final objectives. The central element is to follow the CRISP-DM methodology for the realization of the project. The procedure used to translate the business objectives into data mining objectives, what are the KPI set and partial data exploration of raw data, will be explained. Furthermore, a complete explanation about features putting in my final dataset has been presented. Many of these features have been setted using the doctors 's rules that are explained into section data extraction.

In general we can say that the entire data science project procedure have been followed. Into the following section chapter have been splitted : Bussines understanding, Data understanding, Data exploration, Data preparation and Data extraction.

4.1 Description of the goals

During the development of a Data Science project, a common misperception is that the main part of the work is focused to apply statistics algorithms using tools like deep learning or machine learning but this is often only the last step. In practice a data scientist spends more than 60% of its time to discuss and understand the data after that data scientist starts to apply the prediction methods mentioned before.

This procedure, called data wrangling, represents much of the analyst’s professional process and it takes up most of its working day. Activities to apply are: to understand the informations are provided by the data, to choose the data to use, to understand how to combine multiple different data sources, to decide how to split the result in terms of size and shape that can drive the next steps of analysis.

The final goal of this thesis is to define methods and techniques that can been used to establish dementia patient information and how they evolve after the first episode to the last one. The idea is that a clinic will use this model to aggregate information and predict the likelihood of a rapid or slow disease progression. The rate of progression will be defined according to a specific metric derived from a measure of cognitive ability as mentioned in section 2.1, mini mental state examination(MMSE).
4.2 Business Understanding

In this project three different KPIs have been defined to understand and improve the knowledge about dementia and its causes: Multiple Filter Analysis, Identification of possible strong correlations between MMSE and patient information, How to evolve the rate of progression after the first diagnosis.

4.2.1 Multiple Filter Analysis

During this phase filter analysis have been performed on Patient Final Data based on different factors such as age, sex, comorbidities, lifestyle, family history (both maternal and paternal), age of onset, BMI and so on.

Process Outcome KP1 – Multiple Filter Analysis

This key indicator is composed by the following metrics:

- Main comorbidities associated to moderate level of dementia
- Main comorbidities associated to severe level of dementia.
- Number of patients that are strong smoker with at least one type of dementia.
- Number of patients that are strong drinker with at least one type of dementia.
- Number of patients getting worse.
- Number of patients remaining stable.
- Number of patients getting better.

4.2.2 Identification of possible strong correlations between MMSE and patient information

As stated before, this KPI is used to analyze any possible correlation between MMSE and patient information. In this section it has been tried to understand if some features have values above a fixed threshold using data analysis tools like heatmap and bar charts. Using the reference document, we can say that correlation coefficients are used in statistics to measure relationship between two variables. The formulas return a value between
-1 and 1, where 1 indicates a strong positive relationship, -1 indicates a strong negative relationship and zero indicates no relationship on its. [20].

**Process Outcome KP2 - Identification of strong correlations between MMSE and patient information**
The main metrics to achieve reasonable results are the following:

- Heatmap analysis.
- Identification of features with high correlation ratio.
- Level of multi correlation in the Patients Final Data episode by episode.
- Bivariate analysis episode by episode.

**4.2.3 How evolve the rate of progression after First diagnosis**
Survival Analysis is used to estimate the lifespan of a particular population under study. The goal is to estimate the time for an individual or a group of individuals to experience an event of interest [21]. Into this project the two events used to build survival analysis are the first episode analyzed and the episode when the patient becomes a severe one. Then, it have been computed "time to event" how week among it.

**Process Outcome KP3 - Rate of progression After First diagnosis**
Into this section, main key indicator are:

- Time for Mini Mental State Examination to become severe.
- Identification of features that have p_value less than a specific threshold.
- Hazard Ratio Analysis for a set of features.

**4.2.4 Technical Goals**
Once business goals have been defined, it's time to translate the upon goals into a more technical objectives using the data mining language. The first stuff needed to do is to apply programming tools to make data ready. In this phase we have applied the following Preparation Goals:
1. **General Preparation Goals**

- Preparation Goals 0.1 - Cleaning and standardization of available data.
- Preparation Goals 0.2 - Extraction of patient’s information from the OPTIMA documents.

2. **KPI 1 Main Goals** In order to obtain insights to fulfill the KPI 1, the following tasks are required:

- Preparation Goals 1.1 - Classification of patients into a fixed set of subgroups in order to enable more precise analysis.
- Preparation Goals 1.2 - Definition of filter analysis on comorbidities type
- Preparation Goals 1.3 - Definition of filter analysis based on APOE genotype to understand if there is a more important one.
- Preparation Goals 1.4 - a complete analysis of patient’s lifestyle trying to identify if exists a correlation between behaviour and mental state examination.
- Preparation Goals 1.5 - Definition of filter analysis on three type of dementia.
- Preparation Goals 1.6 - Setting a new feature called Mixed Status that is a combination of more than one features present into Patient Final Data.

3. **KPI 2 Main Goals** In order to obtain insights to fulfill the KPI 2 the following tasks are required:

- Preparation Goals 2.2 - Setting a constraint that limits patients in the dataset at only those than having at least 5 different episodes.
- Preparation Goals 2.2 - As explained in paragraph 5.3.3, Setting a threshold that defines the minimum number of patients for an episode.

4. **KPI 3 Main Goals** In order to obtain the outcome KP 3 the following steps are required:

- Preparation Goals 3.1 - Identification of features that have p-value less than 0.05.
• Preparation Goals 3.2 - Using features found into PG3.1, applying on it Cox Proportional Hazard Models.

4.3 Data Understanding

The intention of this section is to fulfill a first preliminary exploration on data. To develop this project the main dataset, called Final Patient Data has been created, that combine a subset of information taken of the OPTIMA.

4.3.1 Description of OPTIMA Data Source

Optima Data Source is a huge Data Source that contains many different Data Files used to take informations. Into this Data Source the principal file is Variable Guide-UPM.xls, a file containing guidelines for the variables. For each variable there is a brief description, data types and information about strange or incorrect values. For example, some tables presented 9999 values that means null value.

![Figure 6: Overview Variable Guide](image-url)
4.3.2 Description of the Data Files

To build Patients Final Data an huge different files taken from the OPTIMA project have been used. Files used are the following:

- **CAMDEXSCORES.xlsx**
  To obtain MMSE(Mini Mental State Examination) and gender for each patient episodes.

- **optimaDX.xlsx**
  To classify if the patient has Alzheimer, other type of dementia or no dementia.

- **Data Request Jan 2019 (Optima, Lead, Challenge) final history.xlsx and nart duration smoking.xlsx**
  To classify patients into one of the following categories: current smoker, former smoker and no smoker.

- **Data Request Jan 2019 (Optima, Lead, Challenge) final history.xlsx**
  To define Alcohol Consumption and Education Level of each patient.

- **Clinical background.xlsx**
  To obtain weight and BMI for each patient.

- **Data Request Jan 2019 (Optima, Lead, Challenge) final history.xlsx**
  To define Malignant Comorbidity, Physiatric Comorbidity, Vascular Comorbidity, Sistemic Comorbidity, Sistemic Comorbidity.

- **DiseasesORcomorbidities - comorbidities.xlsx and optima patients CUIs.xlsx**
  To identify if the patient has Chronic Infectin Comorbidity, Metabolic Comorbidity, Respiratory Comorbidity, K Comorbidity, Inflammatory Comorbidity.

- **APOE.xlsx**
  To obtain APOE for each patient.

Using these files a main database with 30 features has been build. An overview of the characteristics features is given in the table below:
<table>
<thead>
<tr>
<th>Variable Name</th>
<th>Variable Type</th>
<th>Column Description</th>
</tr>
</thead>
<tbody>
<tr>
<td>GLOBAL PATIENT DB ID</td>
<td>continuos</td>
<td>unique key dataset</td>
</tr>
<tr>
<td>GENDER</td>
<td>categorical</td>
<td>male or female</td>
</tr>
<tr>
<td>DEMENTIA</td>
<td>categorical</td>
<td>1=yes, 0=no or no value</td>
</tr>
<tr>
<td>OTHER DEMENTIA</td>
<td>categorical</td>
<td>1=yes, 0=no or no value</td>
</tr>
<tr>
<td>NO DEMENTIA</td>
<td>categorical</td>
<td>1=yes, 0=no or no value</td>
</tr>
<tr>
<td>PETERSON MCI</td>
<td>categorical</td>
<td>1=yes, 0=no or no value</td>
</tr>
<tr>
<td>SMOKER</td>
<td>categorical</td>
<td>current smoker, ex smoker, no smoker or no value</td>
</tr>
<tr>
<td>ALCOHOL</td>
<td>categorical</td>
<td>extreme drinking, mild drinking, no drinking or no value</td>
</tr>
<tr>
<td>EDUCATION</td>
<td>categorical</td>
<td>medium, basic, high, no value</td>
</tr>
<tr>
<td>BMI NUMERIC</td>
<td>continuos</td>
<td>body max index number</td>
</tr>
<tr>
<td>BMI DISCRETIZED</td>
<td>categorical</td>
<td>healthy weight, obese, overweight or underweight</td>
</tr>
<tr>
<td>WEIGHT</td>
<td>continuos</td>
<td>numeric value in kg</td>
</tr>
<tr>
<td>EPISODE DATE</td>
<td>date</td>
<td>standard gg/mm/aaaa</td>
</tr>
<tr>
<td>AGE AT EPISODE</td>
<td>continuos</td>
<td>numeric value</td>
</tr>
<tr>
<td>MMSE FOR EPISODE</td>
<td>continuos</td>
<td>score in numeric value</td>
</tr>
<tr>
<td>MMSE DISCRETIZED INIT</td>
<td>categorical</td>
<td>mild, moderate, normal, severe</td>
</tr>
<tr>
<td>EPISODE DATE FINAL</td>
<td>date</td>
<td>standard gg/mm/aaaa</td>
</tr>
<tr>
<td>AGE AT FINAL EPISODE</td>
<td>continuos</td>
<td>numeric value</td>
</tr>
<tr>
<td>MMSE FINAL</td>
<td>continuos</td>
<td>score in numeric value</td>
</tr>
<tr>
<td>MMSE DISCRETIZED FINAL</td>
<td>categorical</td>
<td>mild, moderate, normal, severe</td>
</tr>
<tr>
<td>MALIGNANT COMORBIDITY</td>
<td>categorical</td>
<td>1=yes, 0=no or no value</td>
</tr>
<tr>
<td>PSYSIATRIC COMORBIDITY</td>
<td>categorical</td>
<td>1=yes, 0=no or no value</td>
</tr>
<tr>
<td>VASCULAR COMORBIDITY</td>
<td>categorical</td>
<td>1=yes, 0=no or no value</td>
</tr>
<tr>
<td>CHRONIC INFECTION COMORBIDITY</td>
<td>categorical</td>
<td>1=yes, 0=no or no value</td>
</tr>
<tr>
<td>SISTEMIC COMORBIDITY</td>
<td>categorical</td>
<td>1=yes, 0=no or no value</td>
</tr>
<tr>
<td>METABOLIC COMORBIDITY</td>
<td>categorical</td>
<td>1=yes, 0=no or no value</td>
</tr>
<tr>
<td>RESPIRATORY COMORBIDITY</td>
<td>categorical</td>
<td>1=yes, 0=no or no value</td>
</tr>
<tr>
<td>K COMORBIDITY</td>
<td>categorical</td>
<td>1=yes, 0=no or no value</td>
</tr>
<tr>
<td>IMFLAMMATORY COMORBIDITY</td>
<td>categorical</td>
<td>1=yes, 0=no or no value</td>
</tr>
<tr>
<td>APOE</td>
<td>categorical</td>
<td>all types of APOE genotype</td>
</tr>
</tbody>
</table>

Table 1: Overview of Final Patient Data columns
4.4 Exploratory Analysis

In the first three paragraphs some information about the raw data and then some distribution graphs about the main features into Patients Final Data have been showed.

4.4.1 Raw Data Exploration

A general overview of my raw Data is shown. The general idea has been to make some exploration analysis into three specific dataset: CAMDEXSCORES.xlsx, APOE.xlsx and Data Request Jan 2019 (Optima, Lead, Challenge) final history.xlsx.

<table>
<thead>
<tr>
<th>Important Features CAMDEXSCORES file</th>
</tr>
</thead>
<tbody>
<tr>
<td>Variable Name</td>
</tr>
<tr>
<td>GLOBAL PATIENT DB ID</td>
</tr>
<tr>
<td>GENDER</td>
</tr>
<tr>
<td>EPISODE DATE</td>
</tr>
<tr>
<td>CAMDEX SCORES: MINI MENTAL SCORE</td>
</tr>
<tr>
<td>AGE AT EPISODE</td>
</tr>
<tr>
<td>Number of patients</td>
</tr>
<tr>
<td>1035</td>
</tr>
</tbody>
</table>

Table 2 : Overview of CAMDEXSCORES.xlsx

<table>
<thead>
<tr>
<th>Important Features Data Request Jan 2019 final history file</th>
</tr>
</thead>
<tbody>
<tr>
<td>Variable Name</td>
</tr>
<tr>
<td>GLOBAL PATIENT DB ID</td>
</tr>
<tr>
<td>GENDER</td>
</tr>
<tr>
<td>EPISODE DATE</td>
</tr>
<tr>
<td>AGE AT EPISODE</td>
</tr>
<tr>
<td>HISTORY PATIENT 74-119: (079) HISTORY OF HEAVY SMOKING</td>
</tr>
<tr>
<td>HISTORY OF PATIENT 74-119:(080): HIST. OF REGULAR DRINKING</td>
</tr>
<tr>
<td>HISTORY OF PATIENT 74-119:(081): HISTORY OF HEAVY DRINKING</td>
</tr>
<tr>
<td>HISTORY OF PATIENT 74-119:(081): HISTORY OF DRINKING PROBLEM</td>
</tr>
<tr>
<td>HISTORY PATIENT 74-119:(115)HISTORY CANCER</td>
</tr>
<tr>
<td>Number of patients</td>
</tr>
<tr>
<td>1035</td>
</tr>
</tbody>
</table>

Table 3 : Overview of Data Request Jan 2019 (Optima, Lead, Challenge) final history.xlsx
### Important Features APOE file

<table>
<thead>
<tr>
<th>Variable Name</th>
<th>Variable Type</th>
</tr>
</thead>
<tbody>
<tr>
<td>GLOBAL PATIENT DB ID</td>
<td>Continuous</td>
</tr>
<tr>
<td>APOE</td>
<td>Categorical</td>
</tr>
<tr>
<td>EPISODE DATE</td>
<td>Date</td>
</tr>
</tbody>
</table>

<table>
<thead>
<tr>
<th>Number of patients</th>
<th>total rows</th>
</tr>
</thead>
<tbody>
<tr>
<td>1035</td>
<td>9565</td>
</tr>
</tbody>
</table>

*Table 4: Overview of APOE.xlsx*

Following rules provided by doctor’s team patients having null value for CAMDEX SCORES: MINI MENTAL SCORE feature into CAMDEXSCORES file and null value for APOE feauture into APOE have been removed. During the first analysis into CAMDEXSCORES.XLSX has been seen that we have **9565** tuple splitting for **1035** different patients. If we apply filters to remove null value we obtain the following results:

<table>
<thead>
<tr>
<th>Number of Rows</th>
<th>Number of Patients</th>
</tr>
</thead>
<tbody>
<tr>
<td>9565</td>
<td>1035</td>
</tr>
</tbody>
</table>

<table>
<thead>
<tr>
<th>Number of Rows</th>
<th>Number of Patients</th>
</tr>
</thead>
<tbody>
<tr>
<td>8172</td>
<td>1013</td>
</tr>
</tbody>
</table>

*Table 5: Comparison between Raw Data and Clean Data*

Then 1393 rows and 22 patients. In order to obtain these results the join tools have been used merging the previous dataframes. The keys used to make inner join operation is the pair GLOBAL PATIENT DB ID, EPISODE DATE.
4.4.2 Patient Final Data Exploration

In this section, using Data Visualization tools, the distribution of the most important features in Patient Final Data has been shown to understand if dataset is homogeneous or not and if it is possible to see specific trend or something like that.

4.4.3 Distribution of Patients by AGE AT FIRST EPISODE

![Histogram of Age Distribution](image)

*Figure 7: Distribution of Patients by AGE*

**Data Source:** Patients File

In figures 7 the distribution of Patients into the OPTIMA Data Source based on Age has been shown. The histogram shows that more than 75% of the people are more than 65 years old. This results make sense because this type of disease is associated with older people. It is good to remember that this graph shows people during the first episode. The peak is in the range between 75 and 80 years old.
4.4.4 Distribution of Patients by MMSE SCORE

**Data Source:** Patients File

In figure 11 the distribution of MMSE SCORE has been shown for the first episode and for the last one.

![Figure 8: Distribution of Patient by MMSE SCORE](image)

It is possible to comprehend that normal status and severe status follow an inverse trend. The number of normal status decreases of about 33% (200 units less) while severe status increase of about 300% (300 units more).

These result are inline with scientific research that says risk of dementia diseases grows up in proportion of seniority. MMSE is provided to us as a numerical value then have been applied a code to categorize it to build a categorical analysis like the previous one.
The applied rules, are shown in the following code:

```python
list_mmse = MMSE_data['CAMDEX SCORES: MINI MENTAL SCORE'].tolist()
list_mmse_categorical = []
for i in range(len(list_patient)):
    if int(list_mmse[i]) > 24:
        list_mmse_categorical.append('normal')
    if int(list_mmse[i]) >= 19 and int(list_mmse[i]) <= 24:
        list_mmse_categorical.append('mild')
    if int(list_mmse[i]) >= 14 and int(list_mmse[i]) < 19:
        list_mmse_categorical.append('moderate')
    if int(list_mmse[i]) < 14:
        list_mmse_categorical.append('severe')
```

The Mini-Mental state examination is a cognitive test scoring on a scale of 0-30 where cognitive impairment is evaluated in this way:

- **Normal cognitive impairment**: >24 (It means no relevant problem with memory, language and in general no cognitive problem).

- **Mild cognitive impairment**: 19-24 (It defines problems with memory, language, thinking and judgment at a greater level if it is compared with normal age-related changes. In general, it does not interfere notably with activities of daily life [22]).

- **Moderate cognitive impairment**: 14-18 (During this stage are visible the first dementia symptoms, such as difficulties with language and problem-solving).

- **Severe cognitive impairment**: <14 (A severe cognitive impairment is defined as a deterioration or loss in intellectual capacity that places a person in jeopardy of harming him or herself or others and, therefore, the person requires substantial supervision by another person).
4.4.5 Distribution of Patients by ALLELE

Data Source: Patients File

In figure 8 the distribution of APOE has been shown.

What’s APOE?

"The APOE gene provides instructions for making a protein called apolipoprotein E. This protein combines with fats (lipids) in the body to form molecules called lipoproteins[23]".

Three different type of alleles of APOE gene exist: E2, E3, E4. Everyone has one pair of the gene and the possible combination are the following: E3E3, E3E4, E2E3, E2E4, E4E4, E2E2, E3E3. Many studies prove that APOE4 allele increase the risk for Alzheimer’s and
in general for dementia diseases. Having one copy of E4 (E3/E4) can increase your risk by 2 to 3 times while two copies (E4/E4) can increase the risk by 12 times.[9]

Into OPTIMA dataset patients are distributed in this way: APOE with at least one gene like E3 are more than 82% of samples. Furthermore the graph shows that APOE with at least one gene as E4 are more than 42%.

![Graph showing distribution of patients by MMSE](image)

*Figure 10: Distribution of Patient by MMSE only for patients that have at least one allele E4*

The figure above show how evolve the Mini Mental Score from the first test to the last one considering only patients having at least one gene like E4 (427 patients).

Using this filter, we noticed that figure 9 clearly shows that during the first episode more than 52% of patients have normal MMSE status and only 16% have severe status. While in the last episode more than 56% have a severe status and less than 30% have normal status. This can be considered a good indicator meaning the progression rate of Dementia’s disease has a reliable correlation with the E4 gene.
4.4.6 Distribution of Patients by GENDER

![Bar chart showing distribution of patients by gender.](image)

**Figure 11: Distribution of Patient by GENDER**

**Data Source:** Patients File

In figure above distribution of patients based on gender has been shown. The distribution based on gender is the following: number of women 510 and number of men 503. Then the distribution of patients based on gender is perfectly homogeneous.

4.4.7 Distribution of Patients by COMORBIDITIES

**Data Source:** Patients File

In this section the most common comorbidities for patients into the OPTIMA data source have been analyze. In this stage the goal is to make a preliminary analysis based only on the first episode. Into Patients Final Data there are 9 different types of comorbidity for this reason only a subset of these have been considered:

- **Malignant Comorbidity - Diseases list:** all types of cancer disease based on patient history.

- **Vascular Comorbidity - Diseases list:** diabetes, hypertension, heart disease, hypercholesterolemia, and peripheral vascular disease.

- **Psysiatric Comorbidity - Diseases list:** The most prevalent disease into this category are anxiety disorders, substance use disorders and other depressive disorders.
- **Chronic Infectin Comorbidity - Diseases list**: Burkitt lymphoma, cervical cancer disease, chronic Lyme arthritis disease, neoroborreliosis disease, kaposi sarcoma disease, bacillary angiomatoso disease.

![Graphs showing distribution of malignant and vascular comorbidities](image)

*Figure 12: Distribution patient by Comorbidities: Malignant and vascular comorbidities*

The first graph in figure 13 show that during the first episode, people having Malignant Comorbidity are about 40% of the samples. This trend is in line with some studies that claim cancer is more common in people that have more than 70 years where 46% of them being dead.[24]. The second graph show distribution patient based on vascular comorbidity during the first episode where only 16.8% of samples has the disease.
Graphs above show that about 80% of people doesn’t have chronic infectin comorbidity whereas about 66% of samples doesn’t have psysiatric comorbidity.

4.5 Data Preparation

OPTIMA Data Source is a repository filled by a Doctor’s Team using simple tools, like Excel file. During this phase, doctors doesn’t have a good knowledge of data analysis standard and then they made mistakes in filling the tables. The first task has been data cleaning operation.

In order to enable operations like aggregations, filtering and search on a data cleaning stage is needed.

Applying this task is also helpful for the future studies that will continue this project. Defining a rigorous data cleaning procedure is an essential task.
4.5.1 Section Data Cleaning

Into this section the data cleaning procedure used to obtain a clean data from a raw data has been explained.

Figure 14: data_cleaning

Documents File Cleaning

Goals: Cleaning subcategory fields:

- Removing special characters.
- Removing possible double spacing between words.
- Removing possible duplicated data that are not used.
- Removing rows that have PM(post mortem) wording.

Section File Cleaning

File Cleaning is a huge procedure having the aim to remove all the informations that are wrong or senseless. All the cleaning tasks that are explained below are applied on the
Input File: Section File

Goals: Cleaning fields

- Filtering sections by type, selecting only the ones used for our scope.
- Special characters removal excluding characters involved in the dates format.
- Fixing words spacing and leaving single space between words.
- Removing all value into column used for our scope that have value no consistent.
- Removing rows patient that have no value for mmse score.

Results:

All texts correctly cleaned and formatted. They are now ready to be used for information extraction.
4.5.2 Data Extraction

Data Extraction is a process that has the role of retrieval of data from various sources. This procedure is the most important process during the data analysis project called ETL(Extraction, Transform and Load). In this project, we used a Full Extraction approach. In full extraction, data is copied from the source system in its entirety.

The paragraphs below have been split into two parts: the former explains which is the final result that we want to obtain and the latted one explains in detail the algorithm used to achieve the above objective. To build a dataset that shows the evolution of the EHR(electronic health record) parameters, lifestyle parameters, MMSE and comorbidities episode by episode, episode date are used as a joining key so that we have a consistent dataset. This type of merge is applicable because all the tables that have been used, have an episode date column with consistent data.

To define raw columns that have been taken to create features for Final Patients Data, the idea is looking for related keywords inside Variable Guide-UPM.xls. For example, if we want to identify raw columns for the education level feature, we will seek keywords as school, education, university, degree and so on.

4.5.2.1 Smoking Level Extraction

**Final Target**: Building a column called SMOKING that has four different values: current smoker, former smoker, no smoker, no value.

To identify Smoking Level of each patient two different tables have been taken nart duration smoking.xlsx and Data Request Jan 2019 (Optima, Lead, Challenge) final-history.xlsx. Inside this two tables there is a subset of columns of interest: "SMOKING: SMOKING" and "SMOKING: SMOKING:" TEXT inside nart duration smoking.xlsx and "HISTORY PATIENT 74-119: (079) HISTORY OF HEAVY SMOKING" inside Data Request Jan 2019 (Optima, Lead, Challenge) final-history.xlsx.
Table 6: Overview of files using to Smoking Level

"SMOKING: SMOKING" has four distinct values: 1 indicates people that are current smoker, 2 indicates people that are former smoker, 0 indicates never smoker and 9 indicates that the information about this features are missing.

"SMOKING: SMOKING : TEXT" is a text field where the previous column has a text defining how long this patient has quitted smoking if it is equal to 2. Inside this field the text format is no unique how it is possible to see in the following tables.

| 40 years ago |
| 7 years ago("Never been a heavy smoker") |
| Regular smoker but stopped 18 years ago |
| Stopped 22 years ago |
| Stopped smoking 20 years ago |
| Stopped smoking regularly 14 years ago |

Table 7: Different text inside SMOKING: SMOKING : TEXT

Therefore it is necessary to define a rules to extract the information of interest from this column. Rules are explained into the algorithm section below.

"HISTORY OF HEAVY SMOKING" defines if a patient has smoked more than 20 cigarettes per day for a long time (more than 15 years). This column has 4 values: 1 is equal to YES, 0 is equal to NO, 9 is equal to NOT ASKED, 8 is equal to NOT KNOWN.

For my objective "HISTORY OF HEAVY SMOKING" is more useful than the other two columns because it has at least one value for 1035 patients.
Algorithm:

STEP 1: First of all, only the columns of interest have been taken from the two tables: GLOBAL PATIENT DB ID and EPISODE DATE that are the key of each tables and the three columns mentioned before.

STEP 2: On "SMOKING: SMOKING: TEXT" a substring of a text field has been extracted using a pandas tools: numeric value indicating how long patient has quitted smoking.

Therefore, before applying the extraction rules the result have been stored into a new column called num_year.

Then this algorithm has been applied:

```python
for index, row in list_smoker2.iterrows() :
    if SMOKING:SMOKING == 2 and num_year > 5 :
        patient_categorization_smoke.append('former_smoker')
        patient_id.append(GLOBAL_PATIENT_DB_ID)
    else if SMOKING:SMOKING == 2 and num_year <= 5 :
        patient_categorization_smoke.append('current_smoker')
        patient_id.append(GLOBAL_PATIENT_DB_ID)
```

The code explain that if "SMOKING:SMOKING" is equal to 2 a patient is a former smoker. But if he/she quitted smoking no more than five years ago he/she is labeled as a current smoker. In other case he/she is labeled as a former smoker. Therefore, with the application of this constraint we have increased the number of current smokers compared to those of former smokers.

STEP 3: Taken data from "SMOKING : SMOKING" column using the following algorithm:

```python
for index, row in list_smoker2.iterrows() :
    if SMOKING:SMOKING == 1 and GLOBAL_PATIENT_DB_ID not in patient_id :
        patient_categorization_smoke.append('current_smoker')
        patient_id.append(GLOBAL_PATIENT_DB_ID)
    if SMOKING:SMOKING == 2 and GLOBAL_PATIENT_DB_ID not in patient_id :
```
patient_categorization_smoke.append('former_smoker')
patient_id.append(GLOBAL_PATIENT_DB_ID)

if SMOKING == 0 and GLOBAL_PATIENT_DB_ID not in patient_id :
    patient_categorization_smoke.append('no_smoker')
    patient_id.append(GLOBAL_PATIENT_DB_ID)

The only patient that have been analyze are only patient that are no taken yet.

STEP 4: Last step, taken data from "HISTORY OF HEAVY SMOKING" column. It is the column that has more consistent data because it is has 9565 rows for 1035 patients.

for index,row in list_smoker.iterrows() :
    if HISTORY PATIENT 74-119:(081)HISTORY OF HEAVY SMOKER == 1 or
    GLOBAL_PATIENT_DB_ID not in patient_categorization_smoke:
        patient_categorization_smoke.append('current_smoker')
        patient_id.append(GLOBAL_PATIENT_DB_ID)
    if HISTORY PATIENT 74-119:(081)HISTORY OF HEAVY SMOKER == 0 or
    row['GLOBAL_PATIENT_DB_ID'] not in patient_categorization_smoke:
        patient_categorization_smoke.append('no_smoker')
        patient_id.append(GLOBAL_PATIENT_DB_ID)

The only patient that have been taken are only patient that are no taken yet.
4.5.2.2 Education Level Extraction

Final Target: It defines a column called EDUCATION that has four different values: basic, medium, high and no value. To build this column, the number of education years during each patient's life has been considered.

To identify education level of each patient, information has been taken from Data Request Jan 2019 (Optima, Lead, Challenge) final-present.xlsx. Inside this table, the subset of columns is the following: "Age Left School" and "Years in Further education".

Algorithm:

STEP 1: First of all, only the columns of interest have been taken from the two tables: GLOBAL PATIENT DB ID and EPISODE DATE are the key of each table, and the two columns above.

STEP 2: Using "Age Left School" and "Years in Further education" columns, the formula below has been applied:

\[
education = (AgeLeftSchool + YearsInFurtherEducation) - 5
\]

The idea is to create a new column called education that is a sum of the two previous columns minus the age when people start to go to school typically (5 years old).

STEP 3: During the last step, an if condition has been set to split patients' education level into one of the category defined into the target features. The following code explains how are set ranges:

```python
for i in range(len(education_patient)) :
    if education_patient <= 16 :
        education_categorization.append('basic')
        patient_id.append(GLOBAL_PATIENT_DB_ID);
    if education_patient > 16 and education_patient > 22 :
        education_categorization.append('medium')
        patient_id.append(GLOBAL_PATIENT_DB_ID);
    if education_patient >= 22 :
```
4.5.2.3 Drinking Level Extraction

**Final Target:** It defines a column called DRINKING that has four different values: no drinking, extreme drinking, mild drinking and no value.

To define Drinking level of each patient information on Data Request Jan 2019 (Optima, Lead, Challenge)_final-history.xlsx. have been used. This table contains 1030 different patient with at least one record.

For my objective, the subset of interest are the following: HISTORY PATIENT 74-119: (080) HIST. OF REGULAR DRINKING, HISTORY PATIENT 74-119: (081) HISTORY OF HEAVY DRINKING and HISTORY PATIENT 74-119: (082) HIST. OF PROBLEM DRINKING.

**Algorithm:**

**STEP 1:** First of all, only the columns of interest has been taken from the table: GLOBAL PATIENT DB ID and EPISODE DATE that are the key of each tables and the three columns above.

**STEP 2:** During this step the way informations are used inside the columns have been defined. Into the following table the possible values have been showed.

<table>
<thead>
<tr>
<th>Raw Column Name</th>
<th>Data Type</th>
</tr>
</thead>
<tbody>
<tr>
<td>(080) HIST. OF REGULAR DRINKING</td>
<td>9=NotAsked 8=NotKnown 0=No 1=Yes</td>
</tr>
<tr>
<td>(081) HISTORY OF HEAVY DRINKING</td>
<td>9=NotAsked 8=NotKnown 0=No 1=Yes</td>
</tr>
<tr>
<td>(082) HIST. OF PROBLEM DRINKING</td>
<td>9=NotAsked 8=NotKnown 0=No 1=Yes</td>
</tr>
</tbody>
</table>

*Table 8: Tables used to build DRINKING LEVEL columns*

**STEP 3:** An if condition has been set splitting information into one of the category define into the target feature. the following code explain how are setted ranges:
for index, row in list_alcohol.iterrows:
    if HISTORY PATIENT 74-119:(081)HISTORY OF HEAVY DRINKING == 1 or
        HISTORY PATIENT 74-119:(081)HIST. OF PROBLEM DRINKING == 1 :
        patient_categorization.append('extreme drinking')
    else if HISTORY PATIENT 74-119:(081)HIST. OF REGULAR DRINKING == 1 :
        patient_categorization.append('mild drinking')
    else :
        patient_categorization.append('no drinking')

The algorithm above shows the way data are split. We can notice that is set as extreme
drinking, patients that have value 1 for heavy drinking column and for problem drinking
column, as mild drinking, people that have value 1 for regular drinking column and no
drinking people that have 0 value for the three columns mentioned.
4.5.2.4 Comorbidity Extraction

**Final Targets:** It defines 9 columns that describe different comorbidities. Each comorbidity column has three different values: True, False, no value.

During this task has been used to compute these 9 columns three different tables: Data Request Jan 2019 (Optima, Lead, Challenge)_final-history.xlsx, diseasesORcomorbidities - comorbidities.xlsx, and optima_patients_CUIs.xlsx

If we want to use the information inside the last two tables we need to apply a specific procedure:

**Procedure comorbiditiy_identify:**

1. Definition of **comorbidity_type** variable using table diseasesORcomorbidities - comorbidities.xlsx. This excel file contains a list of different types of diseases and their classification in a specific column called TYPE. It is used to label the disease into one of the 9 comorbidities. For example, if we set TYPE = N that is linked to the physiatric comorbidity, we obtain the following result:

   | C0012546 |
   | C0191877 |
   | C3864035 |
   | C0152025 |
   | C0024437 |
   | C0041466 |
   | C0149958 |
   | C0014544 |

*Figure 15: Overview list of CUI for physiatric_comorbidity*
2. Definition of variable *comorbidity_patients_optima* based on *optima_patients_CUIs.xlsx* to obtain a list of CUI afflicting each patient.

<table>
<thead>
<tr>
<th>Index</th>
<th>Patient ID</th>
<th>Episode Date</th>
<th>CUIs List</th>
</tr>
</thead>
<tbody>
<tr>
<td>1928</td>
<td>2002-03-01</td>
<td>00:00:00</td>
<td>['0000768', '00008274', '00007383']</td>
</tr>
<tr>
<td>60</td>
<td>1994-03-01</td>
<td>00:00:00</td>
<td>['000001126', '00002708']</td>
</tr>
<tr>
<td>554</td>
<td>1992-07-01</td>
<td>00:00:00</td>
<td>['00001256', '000082866', '000206099', '00003631', '00161373', '04628976', '01321137', '00023603']</td>
</tr>
<tr>
<td>1893</td>
<td>2001-06-28</td>
<td>00:00:00</td>
<td>['00001883', '00264413', '00001562']</td>
</tr>
<tr>
<td>1693</td>
<td>1997-04-07</td>
<td>00:00:00</td>
<td>['00006292', '00026267', '00031315']</td>
</tr>
<tr>
<td>1693</td>
<td>1994-04-26</td>
<td>00:00:00</td>
<td>['00002052']</td>
</tr>
<tr>
<td>3550</td>
<td>2001-06-18</td>
<td>00:00:00</td>
<td>['00000761']</td>
</tr>
<tr>
<td>7116</td>
<td>1999-10-25</td>
<td>00:00:00</td>
<td>['00002892', '00136815']</td>
</tr>
<tr>
<td>1693</td>
<td>2001-05-15</td>
<td>00:00:00</td>
<td>['00002960']</td>
</tr>
<tr>
<td>1488</td>
<td>1991-10-07</td>
<td>00:00:00</td>
<td>['00002965']</td>
</tr>
<tr>
<td>8255</td>
<td>1995-11-21</td>
<td>00:00:00</td>
<td>['00005606', '00304129', '00423798', '00049751']</td>
</tr>
<tr>
<td>5350</td>
<td>1998-07-21</td>
<td>00:00:00</td>
<td>['00005864', '00008564', '00351339', '00085611', '0187779']</td>
</tr>
<tr>
<td>5350</td>
<td>1999-07-04</td>
<td>00:00:00</td>
<td>['00005864', '00099599', '00085611']</td>
</tr>
<tr>
<td>1753</td>
<td>1991-06-22</td>
<td>00:00:00</td>
<td>['00005867', '00318447', '00118459', '00013820', '0017081']</td>
</tr>
<tr>
<td>1162</td>
<td>2002-09-16</td>
<td>00:00:00</td>
<td>['00005867', '00304635', '01212121', '01840906']</td>
</tr>
<tr>
<td>2750</td>
<td>2000-10-31</td>
<td>00:00:00</td>
<td>['00005867']</td>
</tr>
<tr>
<td>6000</td>
<td>1991-05-21</td>
<td>00:00:00</td>
<td>['00005867']</td>
</tr>
<tr>
<td>1103</td>
<td>1992-01-28</td>
<td>00:00:00</td>
<td>['00005631', '00031117', '0002226']</td>
</tr>
<tr>
<td>1544</td>
<td>1990-07-09</td>
<td>00:00:00</td>
<td>['00005631']</td>
</tr>
<tr>
<td>4610</td>
<td>2001-05-04</td>
<td>00:00:00</td>
<td>['00005664', '00002871']</td>
</tr>
<tr>
<td>4669</td>
<td>2000-06-25</td>
<td>00:00:00</td>
<td>['00005664', '00136284', '00423653', '00031595']</td>
</tr>
<tr>
<td>4669</td>
<td>2000-06-11</td>
<td>00:00:00</td>
<td>['00005664', '00162478']</td>
</tr>
<tr>
<td>319</td>
<td>2001-07-18</td>
<td>00:00:00</td>
<td>['00005664', '00205158', '00156383']</td>
</tr>
</tbody>
</table>

Figure 16: CUI list overview for pair GLOBAL PATIENT DB ID and episode date

To retrieve comparable data with *comorbidity_type* we need to apply text cleaning operation to remove comma or special character like round bracket and square bracket.

3. At the end, using the two previous variables defined, a procedure code has been apply to obtain list of patients with that comorbidity under analysis. For example, the next code show how the procedure on psysiatric comorbidity has been carried out:

```python
for index, row in comorbidity_patients_optima:
    if row['diagnosis_CUIs'] in comorbidity_type_psyciatric
        patient_with_comorbidity.append(row['GLOBAL_PATIENT_DB_ID'])
```
Where comorbidity_patients_optima is a list containing GLOBAL PATIENT DB ID, episode date and diagnosis_CUI afflicting patient during episode date and comorbidity_type is a list of CUI for psychiatric comorbidity. Then if we find a match between diagnosis_CUI and CUI we can put True on psychiatric comorbidity for that patient on that episode date.

**Algorithm:**

**STEP 1:** First of all, from the three tables only the columns of interest have been taken: GLOBAL PATIENT DB ID and EPISODE DATE that are the key of each tables and the other columns need to compute all types of comorbidities.

**STEP 2:** In this step it has been explained which are the columns taken and how to use them to calculate features.

1. **MALIGNANT\_COMORBIDITY**
   - It uses variable HISTORY OF CANCER from Data Request Jan 2019 (Optima, Lead, Challenge)_\_final-history.xlsx file, we describe if there is incidence of cancer into family history patient.
   - Possible value: 0=no, 1=yes, 9=not known, 8=not asked

```
if HISTORY OF CANCER == 1 :
  MALIGNANT\_COMORBIDITY = TRUE;
if HISTORY OF CANCER == 0 :
  MALIGNANT\_COMORBIDITY = FALSE;
if HISTORY OF CANCER != 1 and HISTORY OF CANCER !=0 :
  MALIGNANT\_COMORBIDITY = no value;
```

2. **CHRONIC\_INFECTIN\_COMORBIDITY**
   - It takes information from diseases\_OR\_comorbidities - comorbidities.xlsx where TYPE = V and optima\_patients\_CUIs.xlsx using procedure comorbidity\_identify explained above.
3. PSYSIATRIC_COMORBIDITY

- It takes info from HISTORY OF PSYCH. ILLNESS From Data Request Jan 2019 (Optima, Lead, Challenge)_final-history.xlsx.
  Possible value: 0=no, 1=yes, 9=not known, 8=not asked
- It Holds information inside diseasesORcomorbidities - comorbidities.xlsx where TYPE = N and optima_patients_CUIs.xlsx using procedure comorbidity_identify explained above.

4. METABOLIC_COMORBIDITY

- It takes information from diseasesORcomorbidities - comorbidities.xlsx where TYPE = M and optima_patients_CUIs.xlsx using procedure comorbidity_identify explained above.

5. SISTEMIC_COMORBIDITY

- It Uses information inside Data Request Jan 2019 (Optima, Lead, Challenge)_final-history.xlsx. The Data are inside columns called (075) HISTORY OF RAISED BP and HISTORY OF DIABETES.
  Possible value: 0=no, 1=yes, 9=not known, 8=not asked.

6. RESPIRATORY_COMORBIDITY

- It takes information from diseasesORcomorbidities - comorbidities.xlsx where TYPE = R and optima_patients_CUIs.xlsx using procedure comorbidity_identify explained above.

7. VASCULAR_COMORBIDITY

- It Uses information inside Data Request Jan 2019 (Optima, Lead, Challenge)_final-history.xlsx. The taken columns, are inside columns called (075) HISTORY OF HEART ATTACK and HISTORY OF STROKE
  Possible value: 0=no, 1=yes, 9=not known, 8=not asked
• It holds information inside diseasesORcomorbidities - comorbidities.xlsx where TYPE = V and optima_patients_CUIs.xlsx using procedure comorbidity_identify explain above.

8. IMFLAMMATORY_COMORBIDITY

• It takes information from diseasesORcomorbidities - comorbidities.xlsx where TYPE = V and optima_patients_CUIs.xlsx using procedure comorbidity_identify explain above.

9. K_COMORBIDITY

• It takes information from diseasesORcomorbidities - comorbidities.xlsx where TYPE = K and optima_patients_CUIs.xlsx using procedure comorbidity_identify explain above.
4.5.2.5 APOE Extraction

Final Target: It builds categorical features that show what is the APOE genotype for each patient.

Before setting APOE’s column we need to remove all the wrong values inside the original raw column.

![Distribution Patient by APOE](image)

*Figure 17: Distribution Patient by APOE*

<table>
<thead>
<tr>
<th>APOE data</th>
<th></th>
</tr>
</thead>
<tbody>
<tr>
<td>Raw Data</td>
<td>Clean data</td>
</tr>
<tr>
<td>number of distinct patient</td>
<td>number of distinct patient</td>
</tr>
<tr>
<td>1023</td>
<td>1013</td>
</tr>
</tbody>
</table>

*Table 9: Comparison between APOE Raw Data and Clean Data*

**Algorithm:**

**STEP 1:** First of all, from the two tables only the columns of interest have been taken: GLOBAL PATIENT DB ID and EPISODE DATE that are the key of each tables and the APOE column above.

**STEP 2:** During this step, the only thing that has been done is to take the values found in the original table after the data cleaning operation.
4.5.2.6 Dementia Type Extraction

Final Targets: It defines 3 columns that describe if the patient has Dementia, Other Dementia or no Dementia; the first one includes only Alzheimer’s disease while other dementia is a huge group of diseases like vascular disease, Parkinson disease, Lewy body dementia and so on.

To define these three columns data inside optimaDX.xlsx have been manipulated, using 13 different columns. To correctly use the information in this table, a set of rules has been provided by the team of doctors. Doctor's rules define a set of constraints to categorize patients into the correct group.

dementia (Alzheimer’s disease), other dementia, no dementia into the table below are respectively D, OD, ND. Rules defined by doctor’s team will be explain into Step 2 of algorithm section.

Algorithm:

STEP 1: First of all, only the columns of interest have been taken from the two tables: GLOBAL PATIENT DB ID and EPISODE DATE that are the key of each tables that are considered and the other columns need to compute all types of comorbidities.

STEP 2: On this section, doctor’s rules have been applied to build dementia, other dementia and no value columns. In the following table every constraints used to define dementia type are explain:

<table>
<thead>
<tr>
<th>Doctor’s rules to define dementia</th>
</tr>
</thead>
<tbody>
<tr>
<td>Variable Name</td>
</tr>
<tr>
<td>COGNITIVE IMPAIRMENT</td>
</tr>
<tr>
<td></td>
</tr>
<tr>
<td></td>
</tr>
<tr>
<td>PETERSEN MCI</td>
</tr>
<tr>
<td></td>
</tr>
<tr>
<td></td>
</tr>
</tbody>
</table>

Table 10: Overview of columns used to build dementia, other dementia, no dementia columns 1/2
<table>
<thead>
<tr>
<th>Variable Name</th>
<th>possible values</th>
<th>doctor’s rule</th>
</tr>
</thead>
<tbody>
<tr>
<td>VCI</td>
<td>1=Yes 0=No 8=Not Known 9=Not Asked</td>
<td>if 1 then D=0, OD=1, ND=0</td>
</tr>
<tr>
<td>DEMENTIA PRESENT</td>
<td>0=No 1=Mild 2=Moderate 3=Severe</td>
<td>if 1,2,3 or 4 then D=1, OD=1, ND=0</td>
</tr>
<tr>
<td>AD (NINCDS-ADSDA)</td>
<td>0=Negative 1=Possible 2=Probable 8=Not Asked</td>
<td>if 1,2,3 or 4 then D=1, OD=0, ND=0</td>
</tr>
<tr>
<td>MIXED DEMENTIA TYPE</td>
<td>1=Yes 0=No 9=Not Asked</td>
<td>if 1,3 then D=1, OD=1, ND=0</td>
</tr>
<tr>
<td>VASCULAR DEMENTIA</td>
<td>0=Negative 1=Possible 2=Probable 8=Not Asked</td>
<td>if 1,2,3 or 4 then D=0, OD=1, ND=0</td>
</tr>
<tr>
<td>DEMENTIA OTHER</td>
<td>1=Yes 0=No</td>
<td>if 1 then D=0, OD=1, ND=0</td>
</tr>
<tr>
<td>PARKINSON DISEASE</td>
<td>1=Yes 0=No</td>
<td>if 1 then D=0, OD=0, ND=1</td>
</tr>
<tr>
<td>PROGRESSIVE SUPRA-NUCLEAR PALSY</td>
<td>1=Yes 0=No</td>
<td>if 1 then D=0, OD=1, ND=0</td>
</tr>
<tr>
<td>FRONTO-TEMPORAL DEMENTIA</td>
<td>0=Negative 1=Possible 2=Probable 8=Not Asked</td>
<td>if 1,2 then D=0, OD=1, ND=0</td>
</tr>
<tr>
<td>CORTICO-BASAL DEGENERATION</td>
<td>0=Negative 1=Possible 2=Probable 8=Not Asked</td>
<td>if 1,2 then D=0, OD=1, ND=0</td>
</tr>
<tr>
<td>LEWY-BODY DISEASE</td>
<td>1=Yes 0=No 8=Not Known 9=Not Asked</td>
<td>if 1 then D=0, OD=1, ND=0</td>
</tr>
</tbody>
</table>

*Table 11: Overview of columns used to build dementia, other dementia, no dementia columns 2/2*
The table optimaDX has 9565 rows and at least one row for 1030 patients but if we do more accurate analysis, we can notice that there are many patients that have only values like 9, 8 and nan for each episode. For these subgroup of patients is impossible to used doctor's rules "no_value" string has been used to point out patients that have no consistent values.

The following figure shows the way patients distribute between consistent and unconsistent data:

![Figure 18: Overview of data distribution on optimaDX](image)

The graph shows that more than 25% of patients has no data for either column for all episode under analysis, then for these patients, it is impossible to have information about Alzheimer or another type of dementia.
STEP 3: During this stage doctor’s rules have been used to build the columns mentioned above.

The Final result into the Final Patient Data is the following:

<table>
<thead>
<tr>
<th>GLOBAL_PATIENT_DB_ID</th>
<th>GENDER</th>
<th>DEMENTIA</th>
<th>OTHER_DEMENTIA</th>
<th>NO_DEMENTIA</th>
</tr>
</thead>
<tbody>
<tr>
<td>8743</td>
<td>Female</td>
<td>1</td>
<td>1</td>
<td>0</td>
</tr>
<tr>
<td>8742</td>
<td>Female</td>
<td>1</td>
<td>1</td>
<td>0</td>
</tr>
<tr>
<td>8741</td>
<td>Male</td>
<td>1</td>
<td>1</td>
<td>0</td>
</tr>
<tr>
<td>8740</td>
<td>Male</td>
<td>1</td>
<td>1</td>
<td>0</td>
</tr>
<tr>
<td>8739</td>
<td>Male</td>
<td>0</td>
<td>1</td>
<td>0</td>
</tr>
<tr>
<td>8738</td>
<td>Female</td>
<td>1</td>
<td>1</td>
<td>0</td>
</tr>
<tr>
<td>8737</td>
<td>Male</td>
<td>1</td>
<td>1</td>
<td>0</td>
</tr>
<tr>
<td>8736</td>
<td>Female</td>
<td>no_value</td>
<td>no_value</td>
<td>no_value</td>
</tr>
<tr>
<td>8735</td>
<td>Male</td>
<td>1</td>
<td>1</td>
<td>0</td>
</tr>
<tr>
<td>8734</td>
<td>Male</td>
<td>no_value</td>
<td>no_value</td>
<td>no_value</td>
</tr>
<tr>
<td>8733</td>
<td>Female</td>
<td>1</td>
<td>1</td>
<td>0</td>
</tr>
<tr>
<td>8732</td>
<td>Female</td>
<td>1</td>
<td>1</td>
<td>0</td>
</tr>
<tr>
<td>8731</td>
<td>Male</td>
<td>1</td>
<td>1</td>
<td>0</td>
</tr>
<tr>
<td>8730</td>
<td>Male</td>
<td>no_value</td>
<td>no_value</td>
<td>no_value</td>
</tr>
<tr>
<td>8729</td>
<td>Male</td>
<td>no_value</td>
<td>no_value</td>
<td>no_value</td>
</tr>
<tr>
<td>8727</td>
<td>Male</td>
<td>1</td>
<td>1</td>
<td>0</td>
</tr>
<tr>
<td>8726</td>
<td>Male</td>
<td>no_value</td>
<td>no_value</td>
<td>no_value</td>
</tr>
<tr>
<td>8725</td>
<td>Male</td>
<td>1</td>
<td>1</td>
<td>0</td>
</tr>
</tbody>
</table>

*Figure 20: Overview of DEMENTIA,OTHER,DEMENTIA,NO DEMENTIA into Final Patient Data*
5 Discussion of Results

Chapter can be splitted into two part:

1. An exploration on the final dataset Final Data Patients.

2. The development of the KPI defined during the previous chapter. During this stage many techniques have been applied (heatmap, stacked bar chart, survival analysis, distribution analysis) to achieve a complete and consistent analysis of them.

5.1 Exploratory Analysis of Output Data

Once we processed the raw data to extract information from Optima Data Source, it is time to develop the key performance indicator mentioned in section 4.2.

5.1.1 Exploring Final Data Patients

Table Structure: in this table there are 8134 rows associated to 1013 distinct patients. As already discussed in section 4.3.2, dataset contains 30 columns that we can split into the following subcategories: general information, dementia information, episode data, lifestyle patient information and comorbidity information.

5.1.1.1 Number of Episode per Patients

![Figure 21: Number of episode for patients](image)
The figure 21 shows that on average, each patient has an average value of 8 episodes with minimum number of episode equal to 1 and maximum number of episode equal to 35. Graphs below show the chief statistics about episode per Patients

<table>
<thead>
<tr>
<th>Statistics Type</th>
<th>Episode Per Patients</th>
</tr>
</thead>
<tbody>
<tr>
<td>count</td>
<td>1013</td>
</tr>
<tr>
<td>mean</td>
<td>8.02962</td>
</tr>
<tr>
<td>std</td>
<td>5.33049</td>
</tr>
<tr>
<td>min</td>
<td>1</td>
</tr>
<tr>
<td>25%</td>
<td>4</td>
</tr>
<tr>
<td>50%</td>
<td>6</td>
</tr>
<tr>
<td>75%</td>
<td>11</td>
</tr>
<tr>
<td>max</td>
<td>35</td>
</tr>
</tbody>
</table>

*Table 12: Final Patient Data Statistics Episode Number*

5.1.1.2 Distribution of Patients by Dementia Type

Using Doctor’s rules, as discussed on section 4.5.3.6, these four subcategories have been setted. During the first episode patients that have no value are more than 50% whereas in the last episode patients that has no value are less than 50% . An important consideration we come out from the previous chart: every patients that has
1 value about dementia column also have 1 value in the other dementia column. This means that the constraints defined by the doctors to filter any patient with only dementia (alzheimer disease) don’t work maybe because they have nan or not asked value. Using the variations of the graph from the first to the last episode we can define a starting point to perform Multiple Filter Analysis operations.

5.2 KPI 1 - Multiple Filter Analysis

In this section different and various Multiple Filter Analysis have been done. The general scope of this task is to apply some filtering to find new type of information and to identify possible new ways to read the data. To do this, some advice was provided by the doctors after their analysis of Final Patient Data contents. The following Multiple Filter Analysis have been developed:

1. Main comorbidities associated with moderate level of MMSE

2. Main comorbidities associated with severe level of MMSE.

3. Number of patients that are strong smoker when have at least one types of dementia.

4. Number of patients that are strong drinker when have at least one types of dementia.

5. Number of patients getting worse.

6. Number of patients remaining stable.

7. Number of patients getting better.

8. Mixed Status Column Creation.

9. Remove patients that no change their MMSE status from initial episode to final episode.

10. Remove all patients that no have at least five episodes.

5.2.1 Main comorbidities associated to moderate or severe level of MMSE

Moderate level of Mini Mental State Examination

Patients Number on the last episode: 282
Table 13: Distribution moderate MMSE Comorbidities

Analyzing patients having moderate level of MMSE on the last episode, it is possible to notice that the more common comorbidities are malignant and psychiatric. Malignant comorbidity is present on 19% of the samples whereas systemic comorbidity on 15% of the samples.

Severe level of Mini Mental State Examination

Patients Number on the last episode: 567

Table 14: Distribution severe MMSE Comorbidities

The table below shows that patients reaching severe status during the last episode have no specific comorbidities more common than others.
5.2.2 Number of patients that are strong smoker or strong drinker when have at least one types of dementia.

Starting from Final Patient Data, a filter has been applied to remove patients that no have any type of dementia and then on it an analysis to identify strong smokers and strong drinkers have been developed. The goal is to understand if smoke or alcohol influence the development of dementia.

<table>
<thead>
<tr>
<th>Original Final Patient Data</th>
<th>Filtering Final Patient Data</th>
</tr>
</thead>
<tbody>
<tr>
<td>Number of Rows</td>
<td>Number of Rows</td>
</tr>
<tr>
<td>8595</td>
<td>2398</td>
</tr>
<tr>
<td>Number of patient</td>
<td>Number of patient</td>
</tr>
<tr>
<td>1013</td>
<td>556</td>
</tr>
</tbody>
</table>

*Table 15: Comparison number of rows and patient between the original dataset and the filtering dataset*

The two graphs below shows only patient that during the last episode have flag Alzheimer disease and other dementia diseases equal to 1.

*Figure 22: Distribution of Patients by SMOKING level*
SMOKING level graphs shows that more than 65% of the patients never smoked regularly in their lives. Furthermore more than 25% are current or former smoker. Relying on statistics just obtained, no meaningful conclusions could be drawn from the analysis provided.

Instead figure 27 show that more than 75% of the patient are no drinking then meaningful conclusions could be drawn: high consumption of alcohol haven’t strong correlation with dementia’s disease.
5.2.3 Mixed Status Column and Patients Evolution

After an overview of the doctor’s team, has been defined a new column combining two column present into Final Patients Data, DEMENTIA and MMSE_DISCRETIZED column. In this way we obtained a more complex analysis incorporate different but correlate features.

```python
if DEMENTIA == 1 and MMSE_DISCRETIZED == 'mild':
    mixed_status = 'mild'
elif DEMENTIA == 1 and MMSE_DISCRETIZED == 'moderate':
    mixed_status = 'moderate'
elif DEMENTIA == 1 and MMSE_DISCRETIZED == 'severe':
    mixed_status = 'severe'
elif PETERSEN_MCI == 1:
    mixed_status = 'mci_value'
else:
    mixed_status = 'no_value'
```

<table>
<thead>
<tr>
<th>Distribution Mixed_status during first episode</th>
</tr>
</thead>
<tbody>
<tr>
<td>mild</td>
</tr>
<tr>
<td>44</td>
</tr>
</tbody>
</table>

<table>
<thead>
<tr>
<th>Distribution Mixed_status during last episode</th>
</tr>
</thead>
<tbody>
<tr>
<td>56</td>
</tr>
</tbody>
</table>

Table 16: Comparison mix status between first and last episode

Table below shows the distribution of patients during the first episode and the last episode. We can observe that number of patients in each category is growing. The goal is to identify patients that change status and define if is correlative to some other features.
The following table show change status:

<table>
<thead>
<tr>
<th>From no_value to Severe on Mixed Status</th>
<th>14</th>
</tr>
</thead>
<tbody>
<tr>
<td>From MCI_value to Moderate</td>
<td>6</td>
</tr>
<tr>
<td>From no_value to Mild</td>
<td>34</td>
</tr>
<tr>
<td>From no_value to MCI_value</td>
<td>57</td>
</tr>
<tr>
<td>From no_value to Severe</td>
<td>41</td>
</tr>
<tr>
<td>From no_value to Moderate</td>
<td>6</td>
</tr>
<tr>
<td>From MCI_value to Mild</td>
<td>8</td>
</tr>
<tr>
<td>From Mild to Moderate</td>
<td>9</td>
</tr>
<tr>
<td>From Mild to Severe</td>
<td>7</td>
</tr>
</tbody>
</table>

*Table 17: Overview of the mix status value from the first episode to the last*

**From MCI_value to Severe on Mixed Status**

Analyzing this subcategory associated with 14 different patients that have an average of 13 episodes each one, we obtain the following results from which subsequent considerations for future studies can be made.

1. 72% of the patients have at least one genotype E4.

2. 75% of the patients have at least one of the comorbidities defined into Final Patient Data.

3. 92% of the patients are not a former smoker or current smoker.

**From no_value to Severe on Mixed Status**

Patients inside this subcategory are 41 with an average of 12 episodes each one.

1. 41% of the patients inside this subcategory are current smoker while the remaining part 59% are no smoker.

2. The following graph show distribution based on APOE type:
Figure 24: Overview of APOE for sub category from no_value to Severe

It is possible to note that, also in this sub category, the percentage of patients who have at least one e4 genotype is high, equal to 63%.

From no_value to MCI_value on Mixed Status

Number of patients equal to 57 with an average of 7 episodes per patient.

1. 55% of patients in this subgroup suffered from malignant comorbidity.

2. 19 patients on 57 have no dementia flag equal to 1 during the first episode and have dementia (Alzheimer’s disease) and other dementia flag equal to 1 during the last episode.

5.2.4 Patients that changes your MMSE status and with at least 5 episode

On Final Patient Data have been applied a filter considering only patient that changes their mini mental state examination score from the first to the last episode and furthermore all the patients that no have at least 5 episode have been removed.

```python
list_patient_5 = final_list[4]["GLOBAL_PATIENT_DB_ID"]
list_patient_5 = list_patient_5.tolist()
for i in range(len(final_list)) :
    for index, row in final_list[i].iterrows() :
        if row["GLOBAL_PATIENT_DB_ID"] not in list_patient_5 :
            final_list[i] = final_list[i].drop(index)
```
With filter just shown, the number of total patients has fallen by 284 units, becoming equal to 724 and number of total rows becomes equal to 7276.

```python
patient_status_not_change = []
patient_status_change = []
for index, row in final_list[0].iterrows():
    if row['MMSE_DISCRETIZED_INIT'] == 'normal' and row['MMSE_DISCRETIZED_FINAL'] == 'normal':
        patient_status_not_change.append(row['GLOBAL_PATIENT_DB_ID'])
    else:
        patient_status_change.append(row['GLOBAL_PATIENT_DB_ID'])
for i in range(len(final_list)):
    for index, row in final_list[i].iterrows():
        if row['GLOBAL_PATIENT_DB_ID'] not in patient_status_change:
            final_list[i] = final_list[i].drop(index)
```

Further, the application of this new filter above reduces the number of patients in Final Patient Data: 449 patients have changed status from the first to the last episode and the total lines are 4935.

### 5.3 KPI 2 - Correlation Analysis on Patient Final Data

In this section graphical representations have been carried out to analyze the correlations between Mini Mental State Examination and EHR patients, Mini Mental State Examination and general information. The tools used are Stacked bar plots and heatmap which give the possibility to obtain a complete view of the possible correlations. To compute correlation matrix Pearson’s correlation has been used.

Correlation analysis is a statistical method used to evaluate the strength of relationship between two quantitative variables. A high correlation means that variables have a strong relationship with each other, while a weak correlation means that the variables are hardly related [25].

For data scientists, checking correlations is an important part of the exploratory data analysis process. This analysis is used to decide which features affect the target variable.
the most. To do that, heatmap is used because visualization is generally easier to understand than reading tabular data, heatmaps are typically used to visualize correlation matrices. All the proposed analyzes were carried out on Final Patient Data after applying of KPI 1.

5.3.1 Correlation Analysis Between MMSE and general information

![HeatMap MMSE and general information](image)

*Figure 25: HeatMap MMSE and general information*

Into the above figure, have been showed correlation between Mini Mental State Examination and general information (bmi, age, weight).

Looking HeatMap graph we can notice that there are no good positive or negative correlation between mmse and these feautures. Obviously the only strong correlations present are between features directly related to each other, such as body mass index and weight.
5.3.2 Correlation Analysis Between MMSE and comorbidity information

Figure 26: HeatMap MMSE and comorbidity information

Figure 26, show correlation between comorbidity information and Mini Mental State Examination. The two previous analyzes show us that using the pearson coefficient in search of strong correlations for our case study is totally useless.

5.3.3 Multicorrelation Analysis episode by episode

Multi correlation analysis is the most helpful tools to define a sensible study of this KPI. It has been used to show the progression of some features episode by episode, after applying the filters mentioned in section 5.2.2. The following schema shows how much patients have values for a episode. For example, the first five episodes have data for 449 different patients whereas the sixth episode have data for 389 different patients.
After the fourteenth episode it possible to notice that number of patients decreases considerably. Then, for the following stacked bar have been considering only the first 15 episodes. In the next bar graphs, achievement progression of MMSE and some comorbidities types are being compared. The left (vertical) axis represents patients in percentage, and the right (horizontal) axis represents achievement episodes. About the following stacked bar graphs we have to point out that number of patients isn’t constant from episode 1 to episode 15, as mentioned in figure 28. To obtain a more clear understanding of the following stack bar graphs is presented for each of them a table showing:

- number of patients for each patient.
- the distribution of MMSE episode by episode.
- the distribution of feature under analysis episode by episode.
5.3.3.1 Mini Mental Score Examination and Malignant Comorbidity

As explained in the previous section, patients are lost from episode to episode because the number of the episode that has been recorded is no unique for all the patients. About the first five episodes where the number of patients under analysis is homogeneous, the following line chart with normalized data has been defined to obtain more clear representation.

Figure 28: Stack Bar MMSE and Malignant Comorbidity
We can see that as people with severe cognitive impairment increase, the number of people with malignant comorbidity increases but is not enough to say that they are related. For the other episodes, the following table has been used to exhibit some statistics and analysis:

<table>
<thead>
<tr>
<th>Number of Patient</th>
<th>Distribution Data</th>
<th>Episode</th>
</tr>
</thead>
<tbody>
<tr>
<td>389</td>
<td>malignant comorbidity: true=108 false=234, mmse: severe=195 mod=33 mild=55 normal=59</td>
<td>episode6</td>
</tr>
<tr>
<td>342</td>
<td>malignant comorbidity: true=95 false=202, mmse: severe=187 mod=46 mild=41 normal=23</td>
<td>episode7</td>
</tr>
<tr>
<td>297</td>
<td>malignant comorbidity: true=85 false=174, mmse: severe=152 mod=43 mild=43 normal=21</td>
<td>episode8</td>
</tr>
<tr>
<td>259</td>
<td>malignant comorbidity: true=74 false=153, mmse: severe=147 mod=32 mild=30 normal=18</td>
<td>episode9</td>
</tr>
<tr>
<td>227</td>
<td>malignant comorbidity: true=67 false=134, mmse: severe=119 mod=29 mild=29 normal=24</td>
<td>episode10</td>
</tr>
</tbody>
</table>

Table 18: Stack bar malignant-mmse statistics 1/2
Table 19: Stack bar malignant-mmse statistics 2/2

5.3.3.2 Mini Mental Score Examination and Vascular Comorbidity

Figure 30: Stack Bar MMSE and Vascular Comorbidity

Figure 31 showing the progression of vascular comorbidity and mini mental score examination where we can notice that during the first five episode progression is not proportional between them.
## Vascular Comorbidity Evolution

<table>
<thead>
<tr>
<th>Number of Patient</th>
<th>Distribution Data</th>
<th>Episode</th>
</tr>
</thead>
<tbody>
<tr>
<td>389</td>
<td>vascular comorbidity: true=62 false=280</td>
<td>episode6</td>
</tr>
<tr>
<td></td>
<td>mmse: severe=195 mod=33 mild=55 normal=59</td>
<td></td>
</tr>
<tr>
<td>342</td>
<td>vascular comorbidity: true=59 false=238</td>
<td>episode7</td>
</tr>
<tr>
<td></td>
<td>mmse: severe=187 mod=46 mild=41 normal=23</td>
<td></td>
</tr>
<tr>
<td>297</td>
<td>vascular comorbidity: true=53 false=20</td>
<td>episode8</td>
</tr>
<tr>
<td></td>
<td>mmse: severe=152 mod=43 mild=43 normal=21</td>
<td></td>
</tr>
<tr>
<td>259</td>
<td>vascular comorbidity: true=43 false=184</td>
<td>episode9</td>
</tr>
<tr>
<td></td>
<td>mmse: severe=147 mod=32 mild=30 normal=18</td>
<td></td>
</tr>
<tr>
<td>227</td>
<td>vascular comorbidity: true=40 false=161</td>
<td>episode10</td>
</tr>
<tr>
<td></td>
<td>mmse: severe=119 mod=29 mild=29 normal=24</td>
<td></td>
</tr>
</tbody>
</table>

*Table 20: Stack bar vascular-mmse statistics 1/2*

<table>
<thead>
<tr>
<th>Number of Patient</th>
<th>Distribution Data</th>
<th>Episode</th>
</tr>
</thead>
<tbody>
<tr>
<td>201</td>
<td>vascular comorbidity: true=32 false=137</td>
<td>episode11</td>
</tr>
<tr>
<td></td>
<td>mmse: severe=110 mod=28 mild=19 normal=12</td>
<td></td>
</tr>
<tr>
<td>169</td>
<td>vascular comorbidity: true=119 false=26</td>
<td>episode12</td>
</tr>
<tr>
<td></td>
<td>mmse: severe=86 mod=29 mild=15 normal=15</td>
<td></td>
</tr>
<tr>
<td>145</td>
<td>vascular comorbidity: true=23 false=103</td>
<td>episode13</td>
</tr>
<tr>
<td></td>
<td>mmse: severe=80 mod=10 mild=20 normal=16</td>
<td></td>
</tr>
<tr>
<td>126</td>
<td>vascular comorbidity: true=22 false=82</td>
<td>episode14</td>
</tr>
<tr>
<td></td>
<td>mmse: severe=73 mod=9 mild=13 normal=9</td>
<td></td>
</tr>
<tr>
<td>104</td>
<td>vascular comorbidity: true=16 false=66</td>
<td>episode15</td>
</tr>
<tr>
<td></td>
<td>mmse: severe=53 mod=9 mild=11 normal=9</td>
<td></td>
</tr>
</tbody>
</table>

*Table 21: Stack bar vascular-mmse statistics 2/2*
5.3.3.3 Mini Mental Score Examination and Sistemic Comorbidity

Figure 31: Stack Bar MMSE and sistemic comorbidity

<table>
<thead>
<tr>
<th>Number of Patient</th>
<th>Distribution Data</th>
<th>Episode</th>
</tr>
</thead>
<tbody>
<tr>
<td>389</td>
<td>sistemic comorbidity: true=112 false=230 mmse: severe=195 mod=33 mild=55 normal=59</td>
<td>episode6</td>
</tr>
<tr>
<td>342</td>
<td>sistemic comorbidity: true=101 false=196 mmse: severe=187 mod=46 mild=41 normal=23</td>
<td>episode7</td>
</tr>
<tr>
<td>297</td>
<td>sistemic comorbidity: true=91 false=168 mmse: severe=152 mod=43 mild=43 normal=21</td>
<td>episode8</td>
</tr>
<tr>
<td>259</td>
<td>sistemic comorbidity: true=82 false=145 mmse: severe=147 mod=32 mild=30 normal=18</td>
<td>episode9</td>
</tr>
<tr>
<td>227</td>
<td>sistemic comorbidity: true=77 false=124 mmse: severe=119 mod=29 mild=29 normal=24</td>
<td>episode10</td>
</tr>
</tbody>
</table>

Table 22: Stack bar sistemic-mmse statistics 1/2
### Sistemic Comorbidity Evolution

<table>
<thead>
<tr>
<th>Number of Patient</th>
<th>Distribution Data</th>
<th>Episode</th>
</tr>
</thead>
<tbody>
<tr>
<td>201</td>
<td>vascular sistemic: true=66 false=103</td>
<td>episode11</td>
</tr>
<tr>
<td></td>
<td>mmse: severe=110 mod=28 mild=19 normal=12</td>
<td></td>
</tr>
<tr>
<td>169</td>
<td>sistemic comorbidity: true=56 false=89</td>
<td>episode12</td>
</tr>
<tr>
<td></td>
<td>mmse: severe=86 mod=29 mild=15 normal=15</td>
<td></td>
</tr>
<tr>
<td>145</td>
<td>sistemic comorbidity: true=74 false=52</td>
<td>episode13</td>
</tr>
<tr>
<td></td>
<td>mmse: severe= mod= mild= normal=</td>
<td></td>
</tr>
<tr>
<td>126</td>
<td>sistemic comorbidity: true=42 false=62</td>
<td>episode14</td>
</tr>
<tr>
<td></td>
<td>mmse: severe=73 mod=9 mild=13 normal=9</td>
<td></td>
</tr>
<tr>
<td>104</td>
<td>sistemic comorbidity: true=34 false =48</td>
<td>episode15</td>
</tr>
<tr>
<td></td>
<td>mmse: severe=53 mod=9 mild=11 normal=9</td>
<td></td>
</tr>
</tbody>
</table>

*Table 23: Stack bar sistemic-mmse statistics 2/2*
5.4 KPI 3 - Rate of progression after first diagnosis

As mentioned in section 4.3, survival analysis is a "time to event" technique. The general idea is to study the progress of a dataset in relation to time and an event of interest that is seen as our target features. In my case the event of interest is a variable that will be created for the purpose in this section.

Into the following step have been explain how are identify event of interest that is called several status and time.

```python
for i in range(len(patient_group)):
    for index, row in patient_group[i].iterrows():
        if row['MMSE_DISCRETIZED_INIT'] == 'severe':
            patient_that_became_severe.append(row['GLOBAL_PATIENT_DB_ID'])
            date_initial.append(row['EPISODE_DATE'])
            date_when_became_severe.append(row['EPISODE_DATE_FINAL'])
            several_status.append(True)
```

The code below show how has been computed several status. The objective is to locate date when the patient becomes severe, if there is, and then create a variable called severe status that is 1 if the patient becomes severe otherwise 0.

Furthermore, if a severe status has been found the respective date is saved otherwise if a severe status is not identified for that patient, the last episode is taken as the date.

```python
psysiatric_type = psysiatric_type[psysiatric_type['TYPE'] == 'N']
for n in range(len(psysiatric_type)):
    if another_patient_with_psysiatric_problem[n] not in list_patient_with_psyatric_problem:
```

To the temporal division, it has been decided to evaluate the difference between the first episode and the event where the patient becomes severe. The difference between the two dates is calculated as weeks rooting.

Set these two variables a typical nonparametric survival analysis tools was used: Kaplan-Meier Estimator.

**Kaplan-Meier Estimator:** Kaplan-Meier estimate is one of the best options to be used
to measure the fraction of subjects living for a certain amount of time after treatment. In clinical trials or community trials, the effect of an intervention is assessed by measuring the number of subjects survived or saved after that intervention over a period of time. The time starting from a defined point to the occurrence of a given event, for example death is called as survival time and the analysis of group data as survival analysis. This can be affected by subjects under study that are uncooperative and refused to be remained in the study or when some of the subjects may not experience the event or death before the end of the study, although they would have experienced or died if observation continued, or we lose touch with them midway in the study. We label these situations as censored observations. The Kaplan-Meier estimate is the simplest way of computing the survival over time in spite of all these difficulties associated with subjects or situations. The survival curve can be created assuming various situations. It involves computing of probabilities of occurrence of event at a certain point of time and multiplying these successive probabilities by any earlier computed probabilities to get the final estimate. This can be calculated for two groups of subjects and also their statistical difference in the survivals[26].

All the graphs associated with the Kaplan-Meier Estimator are composed of three sub-graphs: the first describes the Kaplan-Meier curve where on the x-axis we have the time intervals and on the y-axis we have the probability that a patient will become severe and into the graph is present p_value.

In statistics, the p-value is the probability of obtaining results as extreme as the observed results of a statistical hypothesis test, assuming that the null hypothesis is correct. The p-value is used as an alternative to rejection points to provide the smallest level of significance at which the null hypothesis would be rejected. A smaller p-value means that there is stronger evidence in favor of the alternative hypothesis[27].

The second graph divides the patients into sub-groups in relation to the feature used and analyzes the trend throughout the time interval, documenting with a percentage the number of patients who did not register the event or who were censored compared it to the initial total. Finally, the third graph shows the number of censured for each subgroup during the whole time interval considered.
5.4.1 APOE survival analysis

Analyzing the previous graph we see that the patients who have APOE e2e4 are those who have 50% likelihood of becoming severe after 440 weeks on average (8.5 years). We also note that patients who have APOE e2e3 and e4e4 are those who after only 260 weeks (5 years) go to severe status. We also note that patients who become severe or who have been censored between 900 and 1500 weeks belong to APOE e3e4 or e3e3.

In the following chart we will consider a survival analysis associated only with patients who have Alzheimer’s disease.

The graph presented confirms the studies mentioned in section 4.4.4. individuals with at
least one E4 genotype have a higher risk of going to a severe status than other patients. From this graph, the patients who go to severe status as late as possible are those with the E3E3 genotype.
5.4.2 ALCOHOL survival analysis

![ALCOHOL survival analysis graph](image)

Figure 34: ALCOHOL survival analysis

Figure 41 shows survival analysis for patients into Final Patient Data associated to alcohol consumption.

In particular:

- **ALCOHOL = 0** represent patients that are no drinking
- **ALCOHOL = 1** represent patients that are mild drinking
- **ALCOHOL = 2** represent patients that are extreme drinking

The graph clearly shows that patients who have a history of extreme drinking have a significantly better likelihood of moving to severe status in a short time. In fact, the graph shows that the probability of becoming severe is 50% already after 250 weeks (4.7 years) and 25% after 300 weeks (5.7 years).
5.4.3 VASCULAR COMORBIDITY survival analysis

Figure 35: VASCULAR comorbidity survival analysis

In the graph above, VASCULAR_COMORBIDITY = 0 indicates patients who do not have that disease while VASCULAR_COMORBIDITY = 1 indicates patients who present it. It is noted that in relation to this comorbidity there is no big difference between the two categories of patients.

This maker can also be seen in the percentage of number at risk where the ratio remains unchanged from one interval to another.
The graph above shows us that 5% of patients have inflammatory comorbidity. We also note that the probability of becoming severe is 50% after 290 weeks while it drops sharply to 25% after 350 weeks.

In relation to the data present in Final Patient Data we can say that the likelihood of passing to severe status is greater for patients who have inflammatory comorbidity.
6 Conclusions and Perspectives

This master thesis work makes it possible to transform raw data into a structured form that can be used to extract knowledge that can help to improve Key Performance Indicators and a set of methods and techniques have been applied in order to discover pattern in a medical data. To achieve this goals a main dataset called Patients Final Data has been developed containing all the information needed.

The main challenge that have been faced can be splitted into:

1. **preparation stage** - Noise removal in medical data, filtering section by type, understanding medical data to define which are suitable data for the proposal purposes, classification of patients into a fixed set of subgroups in order to enable more precise analysis.

2. **business understanding** - Translation of business requirement on a data mining terms, the definition of which are columns that the clinical want on Final Patient Data, identify KPIs and the relative metrics.

3. **data extraction** - the way to use raw data to obtain final features, how raw data have been combined to obtain features and an overview about the algorithm used to compute comorbidity features, drinking feature, education feature, smoking feature, APOE feature and dementia type features.

4. **several approach to extract new information** - Development of the Key Performance Indicators defined and identification of a new points of view usable in future studies.

To developed challenge proposed several tools have been used: histogram, bar chart, stacked bar chart, heatmap, survival analysis techniques, Kaplan-Meir Estimator.

The analysis of Patients Final Data corroborates some knowledge presenting on several studies about alzheimer's and dementia disease but, at the same time, develops new perspectives of analysis.

The project shows us that the likelihood to develop a dementia disease with severe status is more common for patients that have at least one e4 allele. Patients having e3e3 APOE develop the disease slowly and they often don't pass to the severe status.
Furthermore, the project shows us that smoking is not to be considered as a possible cause of disease while it is possible to say that strong drinking problem have a good relationship with the likelihood to become severe and then to developed dementia disease. Finally, no strong correlation was identified between one of the comorbidity analyzed in this project and the result provided by the cognitive tests MMSE. These results obtained aren’t consistent because raw data used to build Final Patient Data comorbidities features have a lot of missing or unknown data. To have a more clear understanding of these relationships we could increase the number of samples and data robustness using the final Patient data structure as a start point. We can consider this as a perspective for future studies because this project is the first part of a more ambitious project that wants to identify causes and possible solutions to dementia disease using machine learning techniques. In conclusion, for each KPIs, the best methods and techniques have been applied to obtain it despite result of the clinical point of view has no been validated.
Bibliography


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