



### Graduados en Ingeniería de la Salud

### **Biological Databases**

**Nutritional Database** 

Made by Susana Rocío Fernández Giaccomassi Pablo Moreno García-Espina

Supervised by Ismael Navas Delgado

Department Lenguajes y Ciencias de la Computación

MÁLAGA, June 2023





## ESCUELA TÉCNICA SUPERIOR DE INGENIERÍA INFORMÁTICA GRADUADO EN INGENIERÍA DEL SOFTWARE

Bases de Datos Biológicas

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## **Abstract**

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Keywords: A, B, C

### Resumen

### 0.1 Initial project description

We are going to create a database which use is going to be directly related to aliments. The goal of our database is to provide a determined patient with a diet following its main characteristics (age, height, weight) Therefore, we will need a group of nutritionists. They are the ones who will be in charge of the development of the diets. We will storage data for aliments mainly, and we will use this data in order to create new diets for our patients in the nutritional center.

We will use a relational database, a type of database that stores and provides access to data points that are related to one another.

In order to generate data for our database, we will use csv files for some of our tables. For the rest of the tables, we will try to generate data automatically with the tools we will see in class (Python)

### 0.2 RDBMS Deployment

In the schema of our database, we will have the following tables: Food, Patient, Nutritionist, Diet and List Of Food. One Diet will consist of different kinds of Food, a Diet will be created by a Nutritionist (although a Nutrionist can create more than one) and a Nutritionist will take care of one or more Patients. The table List Of Food is created as a consequence of the relation between Diet and Food (a relation M:N creates a new table including the primary keys of both tables included in the relation).

Firstly, we have introduced the table Food, which has the food ID as the primary key. The rest of columns of the table are related with information such as the energy in kcal and KJ that provides the food, the quantity of Protein and Fiber in food in every 100 grams... Then the table to register the information about the Patient has been created and fulfilled with relevant information about it in order

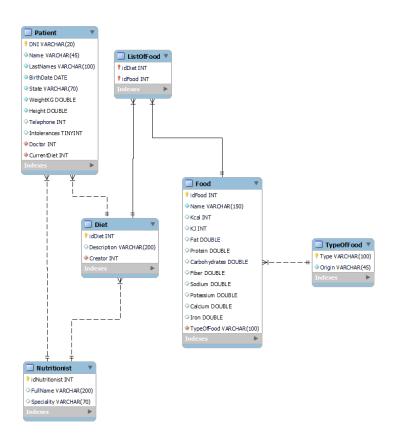


Figure 1: EER model

to organize the new diet. It also stores the current diet that the person is following and the intolerances, information that can be quite useful in order to design a new diet.

The table of the Nutritionist is created also with an ID as the primary key. It also contains information about the name of the nutritionist and the specialty that he or she has (obesity, for example)

For the diets we have another table. As an identifier, we have and ID for the diet (primary key) Also, a description and a Creator are columns for this table. The Description will contain the different steps to follow in order to do the diet. The data which has been introduced in the table Food has been obtained by a csv file in https://www.fao.org/infoods/infoods/tablas-y-bases-de-datos/es/

### 0.3 Query Design

· Food with more than 20 grams of protein (Fig. 2):

```
SELECT NutritionalDB.Food.Name
FROM NutritionalDB.Food WHERE Food.Protein > 20;
```

· All the names of patients treated by the doctor Pablo Moreno Garcia-Espina (Fig. 3):

```
SELECT NutritionalDB.Patient.Name
FROM NutritionalDB.Patient JOIN NutritionalDB.
    Nutritionist
ON Patient.Doctor = Nutritionist.idNutritionist
WHERE Nutritionist.FullName = 'Pablo Moreno Garcia-Espina';
```

· Description of the diets created by the doctor Susana Rocio Fernandez Giaccomassi (Fig. 4)

```
SELECT NutritionalDB.Diet.Description
```

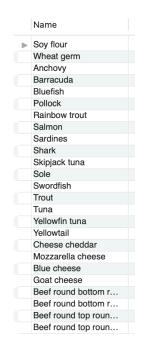


Figure 2: First query output

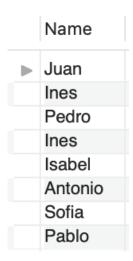


Figure 3: Second query output

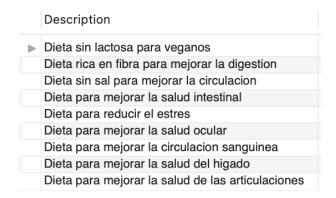


Figure 4: Third query output

```
FROM NutritionalDB.Diet JOIN NutritionalDB.
    Nutritionist
ON Diet.Creator = Nutritionist.idNutritionist
WHERE Nutritionist.FullName = 'Susana Rocio Fernandez
    Giaccomassi';
```

· Obtain the description and the name of their creator of all diets with more than 10 foods (Fig. 5):

```
SELECT NutritionalDB.Diet.Description as 'Diets with
   more than ten foods', NutritionalDB.Nutritionist.
   FullName as 'Creator'
FROM NutritionalDB.Diet JOIN NutritionalDB.
     Nutritionist
ON Diet.Creator = Nutritionist.idNutritionist
WHERE NutritionalDB.Diet.idDiet IN (
SELECT NutritionalDB.Diet.idDiet
FROM NutritionalDB.ListOfFood
GROUP BY idDiet HAVING COUNT(*) > 10
);
```

· Get the names and protein content of foods that have more protein than the average protein of all foods (Fig. 6):

	Diets with more than ten foods	Creator
▶	Dieta rica en proteinas para deportistas	Pablo Moreno Garcia-Espina
	Dieta para reducir el colesterol	Pablo Moreno Garcia-Espina
	Dieta para aumentar la masa muscular	Pablo Moreno Garcia-Espina
	Dieta para reducir la inflamacion	Pablo Moreno Garcia-Espina
	Dieta para mejorar la concentracion	Pablo Moreno Garcia-Espina
	Dieta para aliviar la artritis	Pablo Moreno Garcia-Espina
	Dieta para mejorar la digestion	Pablo Moreno Garcia-Espina
	Dieta para reducir la hinchazon abdominal	Pablo Moreno Garcia-Espina
	Dieta para reducir la retencion de liquidos	Pablo Moreno Garcia-Espina
	Dieta sin lactosa para veganos	Susana Rocio Fernandez Giaccomass
	Dieta rica en fibra para mejorar la digestion	Susana Rocio Fernandez Giaccomass
	Dieta sin sal para mejorar la circulacion	Susana Rocio Fernandez Giaccomass
	Dieta para mejorar la salud intestinal	Susana Rocio Fernandez Giaccomass
	Dieta para reducir el estres	Susana Rocio Fernandez Giaccomass
	Dieta para mejorar la salud ocular	Susana Rocio Fernandez Giaccomass
	Dieta para mejorar la circulacion sanguinea	Susana Rocio Fernandez Giaccomass
	Dieta para mejorar la salud del higado	Susana Rocio Fernandez Giaccomass
	Dieta para mejorar la salud de las articul	Susana Rocio Fernandez Giaccomass
	Dieta baja en grasas	Ismael Navas Delgado
	Dieta equilibrada para mantener el peso	Ismael Navas Delgado
	Dieta sin cafeina para mejorar el sueno	Ismael Navas Delgado

Figure 5: Fourth query ouput

```
SELECT NutritionalDB.Food.Name, NutritionalDB.Food.

Protein

FROM NutritionalDB.Food

WHERE NutritionalDB.Food.Protein > (

SELECT AVG(NutritionalDB.Food.Protein)

FROM NutritionalDB.Food

);
```

### 0.4 RDBMS Optimization

Firstly, we have changed the first query in order to make a deeper search. We can also search in the database according to the type of food, as we made a join between the table of Food and the table of Type of Food.

In order to make the search easier when it comes to diets related with sporty people, we have decided to design two indexes: one with Name and Kcal and another with Protein and Name. This index becomes useful with our first query, as we are looking firstly at the food with protein's value of more than 20 grams.

For the previous index, when we try to execute the first query apparently the

	Name	Protein
<b></b>	Bagel	11
	Buckwheat	13.3
	Einkorn wheat	16.5
	Kamut	14.3
	Millet	11
	Quinoa	14.1
	Soy flour	39.6
	Soybeans cooked	16.6
	Spelt raw	14.6
	Wheat bran	15.6
	Wheat germ	27.2
	Wheat whole grain	12.6
	Whole-grain bread	10.4
	Anchovy	20.3
	Barracuda	25.4
	Bluefish	20.4
	Catfish	18

Figure 6: Fifth query output

index is not used. That could be because the improvement when the index is used is not considerably high when you compare it with the performance without the index.

This picture (Fig. 7) shows the performance of the first query without the index.

This picture (Fig. 8) shows the performance of the first query with the index.

Furthermore, as we will frequently look for the different diets that one nutritionist has created, we will create an index for the column "Full Name" in the table Nutritionist. We already have an index for the id of a nutritionist, but we will typically get a diet of a nutritionist by his concrete name. With this index we will get an advantage in the second and third query, as they use the full name of the nutritionist in order to make the query.

This picture (Fig. 9) shows the time which lasts the second query without the index.

This picture (Fig. 10) shows the time which lasts the second query with the index.

This picture (Fig. 11) shows the performance of the first query without the index.

This picture (Fig. 12) shows the performance of the third query with the index.

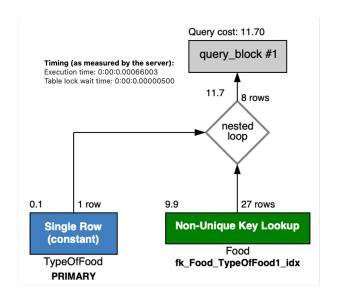


Figure 7: First query without index

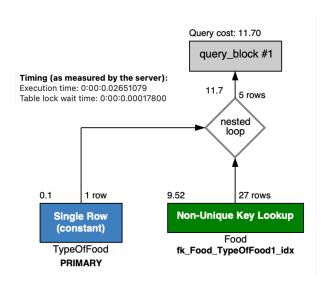


Figure 8: First query with index

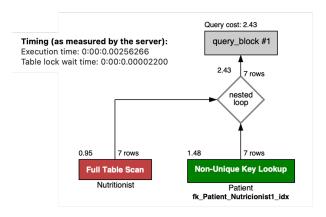


Figure 9: Second query without index

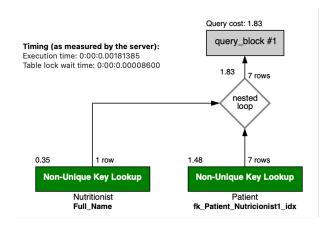


Figure 10: Second query with index

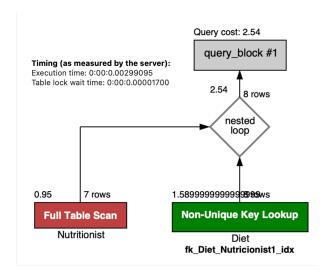


Figure 11: Third query without index

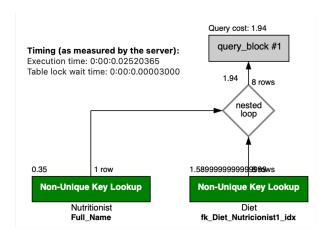


Figure 12: Third query with index

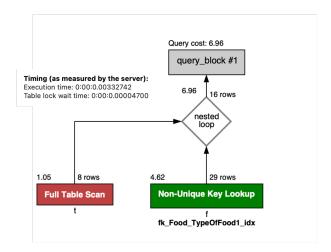


Figure 13: Extra query with index

We see how the time elapsed in the second time is lower than in the first time. The existence of the index makes the performance of the query higher.

We will also do queries which involve the name of the different patients we have in our database. Therefore, it would be suitable too to make an index with the name of the patients. Every time that we do a query in which the name of the patient is involved, we will get an advantage in terms of time, as we can see before.

We think that nutritionists may want to filter the aliments by the nutritional information and origin, so the query would be (Fig.9):

In this case, we have created an index for the table Food including the columns Protein and Iron, but it did not used it, neither the protein index created before.

### 0.5 Python

In order to develop the Python application to test the queries from an user interface, we have created a new project in PyCharm named NutritionalDB.

Specifically in the class NutritionalDB.py we have created three definitions (Fig. 9): one to initiate the connection to the database in SQL, one close the connection and another one to print the result of the queries we will implement next.

```
import pymysql
import pymysql.cursors
from pymysql import DatabaseError
class NutritionalQuerys(object):
    def __init__(self, host, user, password, database):
        self.connection = pymysql.connect(host=host,
                                            user=user.
                                            password=password,
                                            db=database,
                                            charset = 'utf8mb4',
                                            cursorclass =
   pymysql.cursors.DictCursor)
        print("Successfully Connected!")
    def close(self):
        self.connection.close()
        print("Connection Closed!")
    def print_query(self, query):
        cursor = self.exec_query(query)
        print()
        for row in cursor:
            print(row)
```

However, if we analyze the code for the "print query" definition we see that

another definition called "exec query" is called. This new definition is used in order to send the query to the cursor created with the connection (Fig. 10)

```
def exec_query(self, query):
    try:
        with self.connection.cursor() as cursor:
            cursor.execute(query)
        return cursor

except DatabaseError as e:
    if e.args[1] == '#42000Unknown Database':
        print(e)
    else:
        print(e)
```

After these first definitions, we have the definitions for each of the queries. When we made the queries in SQL we asked for concrete values of the columns (for example 20 grams of protein), however here we have parameterized the functions in order to make the query according to the user's goal.

When the user in Python calls the definition in order to ask for a food whose protein content is greater than X (Fig. 11), he has to specify this X value in the call to the method.

```
def gtProtein(self, protein):
    sql = 'SELECT NutritionalDB.Food.Name FROM Food
WHERE Food.Protein > ' + str(protein) + '; '
    print()
    print('Food with more than ' + str(protein) + '
grams of protein:')
    self.print_query(sql)
```

As we can see (Fig. 12), all of the definitions we have implemented in Python have the same structure. Firstly we create a string called sql in which we write the proper query. We have to insert here the different parameters we have in the header of the definition. Secondly, we print an empty line in order to make a separation in the terminal between queries. Then we print a text in order to make

the user understand about the meaning of the query, and finally we execute the method "print query" in order to show in the terminal the results of the query.

We have created a method that receives a doctor as input and returns the patients treated by the input doctor.

```
def treatedBy(self, doctor):
    sql = "SELECT NutritionalDB.Patient.Name FROM
NutritionalDB.Patient JOIN Nutritionist " \
        "ON Patient.Doctor = Nutritionist.
idNutritionist " \
        "WHERE Nutritionist.FullName = '" + doctor + "
';"
    print()
    print('Patients treated by ' + str(doctor) + ':')
    self.print_query(sql)
```

The following definition receives an integer value and returns the diets with more than that number of food.

```
"" + str(quantity) + ");"
print()
print('Diets with more than ' + str(quantity) + '
foods:')
self.print_query(sql)
```

We can also get the food with more than the average of an input column. And the last definition receives two columns and two values, so we can filter by columns and grams.

```
def moreThanAvg(self, column):
     sql = "SELECT NutritionalDB.Food.Name, NutritionalDB
.Food." + column + \
           "FROM NutritionalDB.Food" \
           " WHERE NutritionalDB.Food." + column + " > (
SELECT AVG(NutritionalDB.Food." \
           + column + ") FROM NutritionalDB.Food);"
     print()
     print('Food with more ' + str(column) + ' than the
average: ')
     self.print_query(sql)
 def filterByInfoAndOrigin(self, info1, info2, value1,
value2, origin):
     sql = "SELECT NutritionalDB.Food.Name FROM
NutritionalDB.Food" \
           " JOIN NutritionalDB. TypeOfFood ON Food.
TypeOfFood = TypeOfFood.Type" \
           " WHERE TypeOfFood.Origin = '" + origin + \
           "' AND (Food." + info1 + " > " + str(value1) +
 " OR Food." + info2 + 
           " > " + str(value2) + ");"
     print()
```

Lastly, we have created a main class to test these definitions:

```
if __name__ == '__main__':
    password = 'Your_Password'
    ndb = NutritionalQuerys('localhost', 'root', password, '
    NutritionalDB')
    ndb.gtProtein(20)
    ndb.treatedBy('Pablo Moreno Garcia-Espina')
    ndb.dietsWithMoreFood(10)
    ndb.moreThanAvg('Protein')
    ndb.filterByInfoAndOrigin('Protein', 'Iron', 15, 5, '
    animal')
```

A preview of the outputs can be seen below in figures 15-18:

### 0.6 NoSQL Database Design

In this section we are going to create an alternative to our SQL database. This is going to be done in MongoDB, using JSON documents to create the collections. The structure that our NoSQL database is going to have is simple, having one collection for each table on the relational SQL database.

To export our tables in the JSON format, we used a powerful tool of MySQL Workbench. We selected each table and chose the option "export wizard table". Then, we selected the csv format, and changed it for JSON file when saving it into the repository folder.

In order to create the different collections in MongoDB, we opened a new connection in MongoDB Compass and started to create new collections in a folder. We created the collections with the same name as the tables in SQL and import the data from the JSON files we have just generated.

```
Food with more than 20 grams of protein:

{'Name': 'Pollock'}
{'Name': 'Slue'}
{'Name: 'Anchovy'}
{'Name: 'Inout'}
{'Name: 'Shuefish'}
{'Name: 'Shuefish'}
{'Name: 'Salmon'}
{'Name: 'Salmon'}
{'Name: 'Salmon'}
{'Name: 'Shoutder most'}
{'Name: 'Beef round bottom round roast separable lean and fat trimmed to 0" fat all grades raw'}
{'Name: 'Beef round bottom round roast separable lean and fat trimmed to 0" fat all grades raw'}
{'Name: 'Beef round bottom round roast separable lean and fat trimmed to 0" fat all grades raw'}
{'Name: 'Bout cheese'}
{'Name: 'Yellowtail'}
{'Name: 'Yellowtail'}
{'Name: 'Wild boar chop'}
{'Name: 'Wild boar chop'}
{'Name: 'Shoutder pot roast or steak boneless separable lean only trimmed to 0" fat all grades raw'}
{'Name: 'Shoutder pot roast or steak boneless separable lean only trimmed to 0" fat all grades raw'}
{'Name: 'Shoutder pot roast or steak boneless separable lean only trimmed to 0" fat all grades raw'}
{'Name: 'Shoutder pot roast or steak boneless separable lean only trimmed to 0" fat all grades raw'}
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{'Name: 'Shoutder pot roast or steak boneless separable lean only trimmed to 0" fat all grades raw'}
{'Name: 'Shoutder pot roast or steak boneless separable lean only trimmed to 0" fat all grades raw'}
{'Name: 'Shoutder pot roast or steak boneless separable lean only trimmed to 0" fat all grades raw'}
{'Name: 'Shoutder pot roast or steak boneless separable lean only trimmed to 0" fat all grades raw'}
{'Name: 'Shoutder pot roast or steak boneless separable lean only
```

Figure 14: gtProtein() output

```
Patients treated by Pablo Moreno Garcia-Espina:

{'Name': 'Juan'}

{'Name': 'Ines'}

{'Name': 'Pedro'}

{'Name': 'Ines'}

{'Name': 'Isabel'}

{'Name': 'Antonio'}

{'Name': 'Sofia'}

{'Name': 'Pablo'}
```

Figure 15: treatedBy() output

Figure 16: dietsWithMoreFood() output

```
Food with more Protein than the average:

{'Name': 'Whole-grain bread', 'Protein': 10.4}
{\name': "Reses's Peanut Butter Cups", 'Protein': 10.5}
{'Name': 'Bagel', 'Protein': 11.0}
{\name': 'Millet', 'Protein': 11.0}
{\name': 'Notage cheese', 'Protein': 11.1}
{\name': 'Sottage cheese', 'Protein': 11.3}
{\name': 'Soybeans', 'Protein': 12.0}
{\name': 'Soybeans', 'Protein': 12.0}
{\name': 'Buckwheat', 'Protein': 13.3}
{\name': 'Guinoa', 'Protein': 14.1}
{\name': 'Feta cheese', 'Protein': 14.2}
{\name': 'Feta cheese', 'Protein': 14.3}
{\name': 'Spelt ram', 'Protein': 14.6}
{\name': 'Squid', 'Protein': 15.6}
{\name': 'Squid', 'Protein': 15.6}
{\name': 'Whoat bran', 'Protein': 16.1}
{\name': 'Sibkoans cooked', 'Protein': 16.5}
{\name': 'Beef ground 85% lean meat 15% fat raw', 'Protein': 17.34}
{\name': 'Zebra fish', 'Protein': 17.5}
{\name': 'Carp', 'Protein': 17.5}
{\name': 'Carp', 'Protein': 17.5}
{\name': 'Name': 'Shok', 'Protein': 18.0}
{\name': 'Name': 'Shok', 'Protein': 18.0}
{\name': 'Hadioth', 'Protein': 18.0}
{\name': 'Hadioth', 'Protein': 18.0}
{\name': 'Hadioth', 'Protein': 18.0}
{\name': 'Thotein', 'Protein': 18.5}
{\name': 'Thotein', 'Protein': 18.5}
{\name': 'Thotein', 'Protein': 18.6}
{\name': 'Thote
```

Figure 17: moreThanAvg() output

```
{'Name': 'Whole-grain bread', 'Protein': 10.4}
{'Name': "Reese's Peanut Butter Cups", 'Protein': 10.5}
{'Name': 'Bagel', 'Protein': 11.0}
{'Name': 'Millet', 'Protein': 11.0}
{'Name': 'Millet', 'Protein': 11.0}
{'Name': 'Gottage cheese', 'Protein': 11.3}
{'Name': 'Soybeans', 'Protein': 12.0}
{'Name': 'Wheat whole grain', 'Protein': 12.6}
{'Name': 'Buckwheat', 'Protein': 13.3}
{'Name': 'Quinoa', 'Protein': 14.1}
{'Name': 'Guinoa', 'Protein': 14.3}
{'Name': 'Spelt raw', 'Protein': 14.6}
{'Name': 'Spelt raw', 'Protein': 15.6}
{'Name': 'Susage', 'Protein': 15.6}
{'Name': 'Wald boar sausage', 'Protein': 16.1}
{'Name': 'Wald boar sausage', 'Protein': 16.5}
{'Name': 'Beef ground 85% Lean meat 15% fat cooked pan-broiled', 'Protein': 17.34}
{'Name': 'Beef ground 85% Lean meat 15% fat raw', 'Protein': 17.34}
{'Name': 'Carp', 'Protein': 17.5}
{'Name': 'Carp', 'Protein': 17.5}
{'Name': 'Carp', 'Protein': 18.0}
{'Name': 'Shad', 'Protein': 18.0}
{'Name': 'Shad', 'Protein': 18.0}
{'Name': 'Halbut', 'Protein': 18.0}
{'Name': 'Halbut', 'Protein': 18.2}
{'Name': 'Halbut', 'Protein': 18.2}
{'Name': 'Halbut', 'Protein': 18.5}
{'Name': 'Lounder', 'Protein': 18.6}
{'Name': 'Cad', 'Protein': 18.6}
{'Name': 'Cader', 'Protein': 18.6}
{'Name': 'Cader', 'Protein': 18.6}
{'Name': 'Cader', 'Protein': 18.6}
{'Name': 'Cader', 'Protein': 18.6}
```

Figure 18: filterByInfoAndOrigin() output

```
_id: 1
Description: "Dieta baja en grasas"
Creator: 3

_id: 2
Description: "Dieta sin lactosa para veganos"
Creator: 2

_id: 3
Description: "Dieta rica en proteinas para deportistas"
Creator: 1

_id: 4
Description: "Dieta vegetariana equilibrada"
Creator: 4
```

Figure 19: Diet Collection in MongoDB

Before we add the data to the collections we have to do a change in the JSON files. As MongoDB always renames the primary key with "\_id", in order to relate the primary key in SQL with the primary key in MongoDB we have renamed the primary key in the JSON files as "\_id".

These is the appearance of the different collections in MongoDB: (Fig. 19-24)

### 0.7 NoSQL Database Queries and Pipelines

In this part of the project we are going to translate the different SQL queries we designed for our database into MongoDB language.

In the first MongoDB query we see the following:

```
_id: 1
Name: "Asparagus"
Kcal: 20
KJ: 84
Fat: 0.2
Protein: 2.2
Carbohydrates: 3.9
Fiber: 2.1
Sodium: 2
Potassium: 202
Calcium: 24
Iron: 1.1
TypeOfFood: "vegetables"
_id: 2
Name: "Bell Pepper"
Kcal: 20
KJ: 84
Fat: 0.2
Protein: 0.9
Carbohydrates: 4.6
Fiber: 1.7
Sodium: 3
Potassium: 211
```

Figure 20: Food Collection in MongoDB

```
_id: ObjectId('6438338635aa65b68feebd89')
idDiet: 26
idFood: 1

_id: ObjectId('6438338635aa65b68feebd8a')
idDiet: 46
idFood: 1

_id: ObjectId('6438338635aa65b68feebd8b')
idDiet: 30
idFood: 4

_id: ObjectId('6438338635aa65b68feebd8c')
idDiet: 1
idFood: 5
```

Figure 21: ListOfFood Collection in MongoDB

```
_id: 1
FullName: "Pablo Moreno Garcia-Espina"
Speciality: "Deporte"

_id: 2
FullName: "Susana Rocio Fernandez Giaccomassi"
Speciality: "Descenso"

_id: 3
FullName: "Ismael Navas Delgado"
Speciality: "Deporte"

_id: 4
FullName: "Antonio Rodriguez Horcas"
Speciality: "Ascenso"
```

Figure 22: Nutritionist Collection in MongoDB

```
_id: "02358124Z"
Name: "Carmen"
LastNames: "Ruiz Soto"
BirthDate: "1952-08-08"
State: "descenso"
WeightKG: 89.71
Height: 1.86
Telephone: 608765432
Intolerances: 0
Doctor: 3
CurrentDiet: 44
_id: "05622134W"
Name: "Pilar"
LastNames: "Ortega Jimenez"
BirthDate: "1959-12-21"
State: "ascenso"
WeightKG: 161.1
Height: 1.91
Telephone: 650123456
Intolerances: 0
Doctor: 7
CurrentDiet: 27
```

Figure 23: Patient Collection in MongoDB

```
_id: "cereals and bread"
Origin: "processed"

_id: "chocolates and sweets"
Origin: "processed"

_id: "dairy"
Origin: "animal"

_id: "fish"
Origin: "animal"

_id: "fruits"
Origin: "vegetal"

_id: "legumes"
Origin: "vegetal"
```

Figure 24: TypeOfFood Collection in MongoDB

```
},
},

{
    $project: {
        _id: 0,
        Name: 1,
        Protein: 1,
    },
},
```

Firstly we have extracted all of the food with protein levels higher than twenty grams, and then we have extracted out of them the "meat" food. In order to obtain the same query as in SQL, we have made a project in order to stay just with the fields name and protein.

In the second query we have:

```
[
  {
    $lookup: {
      from: "Patient",
      localField: "_id",
      foreignField: "Doctor",
      as: "patients",
    },
  },
  {
    $match: {
      FullName: "Pablo Moreno Garcia-Espina",
    },
  },
  {
    $project: {
      _id: 0,
```

```
"patients.Name": 1,
},
},
]
```

We make a join from the table Nutritionist to the table Patient relating the identifier field and the Doctor one. Then we filter the FullName avoiding all excepting "Pablo Moreno Garcia-Espina" and finally we project the patient's name.

The third query in MongoDB is:

```
[
  {
    $lookup: {
      from: "Nutritionist",
      localField: "Creator",
      foreignField: "_id",
      as: "nutritionist",
    },
  },
  {
    $match: {
      "nutritionist.FullName":
        "Susana Rocio Fernandez Giaccomassi",
    },
  },
  {
    $project: {
      _id: 0,
      Description: 1,
    },
  },
]
```

We join the table Diet with Nutritionist by relating the fields Creator in Diet and identifier in Nutritionist. Then we remain only with the diets made by the nutritionist "Susana Rocio Fernandez Giacomassi" and we project the diet's description.

The fourth query is:

```
client = MongoClient('mongodb+srv://susanfg:bS23eImV0TRer7Xe
  @cluster0.f34dy4x.mongodb.net/test')
result = client['NutritionalDB']['Diet'].aggregate([
    {
        '$lookup': {
            'from': 'Nutritionist',
            'localField': 'Creator',
            'foreignField': '_id',
            'as': 'nutritionistDoctor'
        }
    }, {
        '$lookup': {
            'from': 'ListOfFood',
            'localField': 'id',
            'foreignField': 'idDiet',
            'as': 'foods'
        }
    }, {
        '$match': {
            'foods': {
                '$gt': {
                    '$size': 10
            }
        }
    }, {
        '$project': {
            'Diets with more than ten foods': '$Description'
            'Creator': '$nutritionistDoctor.FullName'
```

```
}
])
```

Here we are doing an aggregation from the Diet collection, first we use a lookup to create an array with the nutritionists of each diet. Then we do another lookup with ListOfFood to have all the foods of a diet in the foods array. Then we use match to filter the arrays food with size greater than 10. Lastly, we project the description of the diets and their creator. Basically, we are searching the diets with more than ten foods.

### 0.8 Python Export to XML from Queries

In this section we are going to use a python script to convert the JSON files obtained for MongoDB to XML files. The code run for this is the following one:

```
if __name__ == '__main__':
    # get the xml from an URL that return json
    print("Reading ...")
    data = readfromjson("TypeOfFood.json")
    print("Translating ...")
    f = open("TypeOfFood.xml", "a")
    f.write(json2xml.Json2xml(data).to_xml())
    f.close()
```

In this program, we use the function readfromjson() that reads the .json file and returns an object that represents the data. Then we open the new file .xml with open and we use 'a' as second parameter, to append the data. The library json2xml is used to convert the JSON object into a XML, and we write the XML object in the new .xml file.

Once we have the six tables converted to .xml files, we can upload them into ExistDB. We have a docker container with ExistDB, so we access using local-host:8080. We have chosen exide to create a collection. The steps performed to create the collection and upload the files are:

- Select directory
- File > Manage > Log in > Create collection
- We named it "NutritionalDB"
- Double click in the collection
- Upload Files

Finally, we uploaded the files into the collection.

The format of Diet.xml is:

```
<all >
<item type="dict">
  <_id type="int">1</_id>
  <Description type="str">Dieta baja en grasas </

Description >
  <Creator type="int">3</Creator>
</item>
  ...
  <\all >
```

### Food.xml structure:

```
<TypeOfFood type="str">vegetables </TypeOfFood>
</item>
...
...
<\all>
```

ListOfFood.xml is structured as follows:

Nutritionist.xml lists the doctors in the following structure:

```
<all >
<item type="dict">
    <_id type="int">1</_id>
    <FullName type="str">Pablo Moreno Garcia-Espina</
FullName>
    <Speciality type="str">Deporte</Speciality>
</item>
    ...
    ...
    <\all >
```

For the table Patient, we have the Patient.xml file:

```
<all >
<item type="dict">
<_id type="str">02358124Z</_id>
<Name type="str">Carmen</Name>
```

And lastly, the TypeOfFood.xml file has the different types of food:

```
<all >
<item type="dict">
  <_id type="str">cereals and bread </_id >
  <Origin type="str">processed </Origin >
  </item >
    ...
  <\all >
```

#### 0.9 XML Database

Now that we have the tables in XML, we can translate our queries into XQuery language. The first query would be:

Figure 25: First XQuery output

```
<Protein > { data ( $food / Protein ) } < / Protein > </Food >
```

We are returning the result in the xml structure, and we obtain 22 Food items, as it was expected (Figure 25).

The following query asks for the patients treated by a specific doctor:

```
for $patient in doc("/db/NutritionalDB/Patient.xml")/all/
  item
for $nutritionist in doc("/db/NutritionalDB/Nutritionist.xml
  ")/all/item
where $patient/Doctor = $nutritionist/_id and $nutritionist/
  FullName = 'Pablo Moreno Garcia-Espina'
return data($patient/Name)
```

And it gives us the 8 patients treated by the doctor (Figure 26).

The third query is used in order to determine the diets which have been created by one nutritionist:

```
for $diet in doc("/db/NutritionalDB/Diet.xml")/all/item
for $nutritionist in doc("/db/NutritionalDB/Nutritionist.xml
   ")/all/item
where $diet/Creator = $nutritionist/_id and $nutritionist/
   FullName = "Susana Rocio Fernandez Giaccomassi"
return data($diet/Description)
```

It return 27 different diet descriptions (Figure 27)

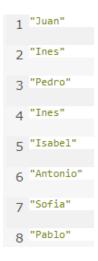


Figure 26: Second XQuery output

```
[] description (que-tra-distra sin taltes per subject descriptions
[] description (que-tra-distra exist on these persons a distribution-transpirate
[] description (que-tra-distra exist on the persons of a distribution-transpirate
[] description (que-tra-distra exist of persons)
[] description (que-tra-distra exist of persons)
[] description (que-tra-distra exist taltes)
[] description (que-tra-distra exist)
[] description (qu
```

Figure 27: Third XQuery output

The fourth query returns the diet descriptions and the creator of the diet which has more than ten food in their diet.

The last query searches the food with more protein than the average:

We calculate the average value of protein for the entire file and then we search the items with more than the average, returning an xml structure (Figure 29).

Figure 28: Last XQuery output

## Contents

	0.1	Initial project description	3			
	0.2	RDBMS Deployment	3			
	0.3	Query Design	5			
	0.4	RDBMS Optimization	8			
	0.5	Python	13			
	0.6	NoSQL Database Design	17			
	0.7	NoSQL Database Queries and Pipelines	21			
	0.8	Python Export to XML from Queries	28			
	0.9	XML Database	31			
1	Intr	oduction	37			
	1.1	Motivation	37			
	1.2	Objectives	38			
	1.3	Structure of the document	39			
	1.4	Technologies used	39			
2	Con	clusions and Futures Lines of Research	41			
	2.1	Conclusions	41			
	2.2	Future lines of Research	41			
3	Con	iclusiones y Líneas Futuras	43			
	3.1	Conclusiones	43			
	3.2	Líneas Futuras	43			
Aj	ppen	lix A Installation				
	Manual					

## 1

### Introduction

### 1.1 Motivation

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### 1.2 Objectives

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### 1.3 Structure of the document

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### 1.4 Technologies used

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## 2

# Conclusions and Futures Lines of Research

### 2.1 Conclusions

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### 2.2 Future lines of Research

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## Conclusiones y Líneas Futuras

### 3.1 Conclusiones

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#### 3.2 Líneas Futuras

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## Appendix A Installation Manual

Requirements:



E.T.S de Ingeniería Informática Bulevar Louis Pasteur, 35 Campus de Teatinos 29071 Málaga