Recommender System on Nutritional Values for Different Food Items

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Abstract

The vast amount of data available today allows information enthusiasts with right skills to draw fruitful insights by inferences and patterns from data. These inferences and patterns give rise to the concept of Recommender Systems which play a major role in our day-to-day decisions in the form of various categories like "Recommended Items", "You may also Like" or "People you may know" in fields of ecommerce and social media. We intend to extend the capabilities of Recommender system to find similarities and diversity in various food nutritional values and present them as alternative to the food being looked for.

I. Introduction

COVID-19 is an unforeseen pandemic that struck the world on an unprecedented scale during the early months of 2020. With panic and lot of measures being placed, one of the most essential necessities is stocking up on food items as there are social restrictions placed in various different parts of the country with little to no access to all the food items that are otherwise available easily. With the growing need to find alternative foods with same nutritional values, it seems apt to design a recommender system that would recommend various food items with similar nutritional values in same or different food types with flexibility as desired by the user.

A recommender system, or a recommendation system is a subclass of information filtering systems that seeks to predict the "Rating" or "Nearness" of an item. The goals of a recommender system are relevance to the product being searched, novelty, serendipity, Increasing recommendation diversity. There are different types of recommender systems such as collaborative recommender system, content-based recommender system, knowledge-based recommender system, demographic recommender system, Utility based recommender system, Hybrid recommender system. In this project, we use item collaborative filtering with neighborhood-based methods to make recommendations. Diversity is a term sparsely concentrated in any recommender system. The objective of this project is providing the user with an option to diversify their recommendations where we recommend them a number of popular but less similar items while keeping in mind the nutritional requirement of the food item in question.

II. Conceptual Approach

Item-based collaborative filtering is a model-based algorithm for making recommendations. In the algorithm, the similarity between different items in the dataset is calculated by using a similarity measure, and then these similarity values are used to predict nearness to the searched food item. Here, we use the method of K-Nearest Neighbors to recommend the most similar food items based parametrically on the nutritional values. The KNN algorithm assumes that similar things exist in close proximity.

The k nearest neighbor classifier is based on the Euclidian distance between products. Let x_i be the input sample with p features $(x_{i^1}, x_{i^2}...x_{ip})$, n be the total samples (I = 1, 2, 3...n) and p be the total number of features (j = 1, 2, 3...p). Then, the Euclidian distance between x_i and x_i (I = 1, 2, 3...n) is defined as

$$d(xi, xl) = \sqrt{(x_{i1}-x_{l1})^2 + (x_{i2}-x_{l2})^2 + ... + (x_{ip}-x_{lp})^2}$$

The predictions of KNN are directly from the dataset. For new sample(x), the prediction is done by searching through the entire dataset for the nearest k-neighbors by using Euclidian distance. KNN is a lazy learning algorithm, which means, we do not need model fitting or a training step. The value of k in KNN determines the number of nearest neighbors to be selected.

In recommender systems, by informal definition, a longtail is divided into two separate parts: the head and the tail. The head part contains of most similar items (in this case, the items with nearest distance) which are often items or

recommendations of the same food type. And the tail of the curve is characterized by the remainder of the products. Computationally, the long tail model, F(x) simulates any heavy-tailed distribution. It models the cumulative distribution of long tail data F(x) which represents the share (%) of total volume covered by objects up to rank x:

$$F(x) = \frac{\beta}{(\frac{N_M}{r})^{\alpha} + 1}$$

Where α is the factor that defines S-Shape of the function. β is the total volume share and N_M is the median of the number of objects that cover half the total volume.

In this project, we diversify the items from the recommender system based on user selection by selecting the 'n' items from the tail end, to provide a different set of recommendations that are distinct in product type from remainder of items from the KNN.

III. Dataset and Preprocessing of Data

The dataset being used in "Composition of Foods, Raw, Processed, Prepared USDA National Nutrient Database for Standard Reference, Release 28" from U.S. Department of Agriculture. This dataset contains data for all food items in Standard Reference (SR-28), with nutrient values in 53 columns. These data are presented per 100 grams, edible portion. The Daily values of Nutrition and Supplement is obtained from the U.S. Food and Drug Administration.

For data preprocessing, we first scale down the number of different nutrients to most essential ones, viz. 'Energ_Kcal', 'Lipid_Tot_(g)', 'FA_Sat_(g)', 'Cholestrl_(mg)', 'Sodium_(mg)', 'Carbohydrt_(g)', 'Fiber_TD_(g)', 'Sugar_Tot_(g)', 'Protein_(g)', 'Vit_D_µg', 'Calcium_(mg)', 'Iron_(mg)', 'Potassium_(mg)'.

Then, we clear the rows with missing values of data. Since, each of the above nutrients have different measurement scales, scaling of data becomes essential. We use a Stadardscaler from python package 'sklearn' to attain scaling.

This data is then saved to a excel spreadsheet as "Preprocessed Data.csv" for predictions.

```
#Reading the dataset
datafile= pd.read excel("sr28abxl\ABBREV.xlsx")
 #Extraction of features
=
| features=['Energ_Kcal'
           'Lipid_Tot_(g)',
           'FA_Sat_(g)'
           'Cholestrl (mg)',
           'Sodium (mg)',
           'Carbohydrt_(g)',
           'Fiber_TD_(g)',
'Sugar_Tot_(g)'
           'Protein (g)',
           'Vit D µg',
           'Calcium_(mg)',
           'Iron (mg)',
           'Potassium (mg)']
 #Filtering only the 'Features' columns from the dataset
attributes = datafile[features]
 #Drop missing values
 attributes = attributes.dropna()
 dataset = attributes
 #Scaling the cleaned data
scaled = StandardScaler()
 dataset scaled = scaled.fit transform(attributes)
dataset["Itemname"] = datafile["Shrt Desc"]
dataset["NDB No"] = datafile["NDB_No"]
dataset = dataset.dropna()
 #Saving Preprocessed data
dataset.to_csv("Preprocessed_dataset.csv")
np.savetxt("Scaled dataset.csv", dataset scaled, delimiter=",")
```

IV. KNN Model, Front-End design and Computational Approach

We used the package 'Scikit Learn' for the computation of K-Nearest Neighnbors. Scikit-learn is an open source machine learning library that supports supervised and unsupervised learning. It also provides various tools for model fitting, data preprocessing, model selection and evaluation, and many other utilities. Sklearn.neighbors provides functionality for unsupervised and supervised neighbors-based learning methods. The principle behind nearest neighbor methods is to find a predefined number of training samples closest in distance to the new point, and predict the label from these. The number of samples can be a user-defined constant (k-nearest neighbor learning).

In scikit learn, KneighborsClassifier implements learning based on the nearest neighbors of each query point, where is an integer value specified by the user. Here, we are specifying the value of k = 100, such that first 100 similar products are listed. The output of KNN is stored in file 'finalized_model.sav' and the Euclidian distances in file 'finalized model dist.sav'.

```
#Building the model using K-NN with K = 100
recommendations = NearestNeighbors(n_neighbors=100, algorithm='auto').fit(dataset_scaled)
#Euclidian Distance Calculation
distances,item indices = recommendations.kneighbors(dataset scaled)
```

The distance matrix generated is as follows:

```
In [2]: runcell(0, 'C:/RecommenderSystem/CodeBase/recomimpl.py')
             0.61759664 1.04453146 ... 7.23189118 7.24119297 7.25147386]
[[0.
 [0.
             1.04453146 1.04938804 ... 6.45908489 6.46093493 6.47629927]
             2.47958752 2.5172238 ... 9.24217831 9.24846557 9.30596899]
 [0.
             0.6272778   0.63303357   ...   2.46431651   2.47899462   2.48355696]
 [0.
 [0.
             0.40336762 0.43586258 ... 0.76842859 0.7684736 0.76869717]
 [0.
             0.53225514 0.53313326 ... 0.73063537 0.73118318 0.73243512]] [[
                                                                                    107
       582 2275
                  56]
          0 107 ... 4673 3445 2312]
     1
             107 ... 3503 4823 4673]
     2
 [4840 4825 3587 ... 157 3443
 [4841 2804 4837 ... 4183 4266 4318]
  4842
        781
             806 ... 4152 1575 3961]]
```

We use the package "flask" to build a WebApp framework for the front-end UI of the recommender. The web-page is scripted using HTML and has the following outlook:

← → O 0 127.0.0.1:5000	7/4	3711	田	
Food Recommender Recommender System For Alternate Food With Similar Nutritional Values				
Food Recommender				
Product: Choose diversity over accuracy: Yes v Predict				

With the K-Nearest Neighbors calculated and Distance Matrix calculated, we use the following libraries:

Flask	Flask is a popular, extensible web microframework for
	building web applications with python
JobLib	JobLib is a set of tools that provide lightweigh
	pipelining in python
FuzzyWuzzy	FuzzyWuzzy is a library of python which is used for
	string matching by calculating the difference between
	sequences
Pandas	Pandas is an open-source data analysis and manipulation
	tool built on top of Python

In this program, we use the FuzzyWuzzy to introduce diversity to the system. FuzzyWuzzy package matches the string accuracy in a given sequence. When diversity is True, we use the output from FuzzyWuzzy to select the items that do not belong to a particular food type.

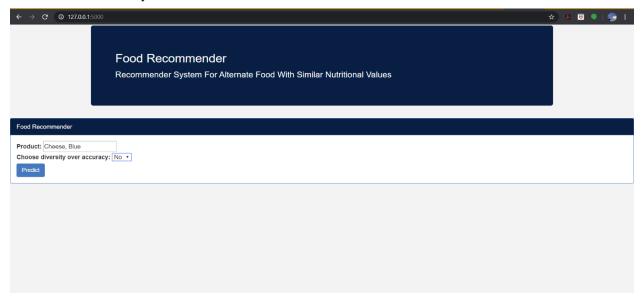
For ex. If searched for food type 'Cheese,Blue' and diversity is False, then the top 8 results belong to the food type 'cheese'. But if diversity is 'True', the program outputs list of food items from the top 100 nearest neighbors, whose string match is less than 40%.

With the following libraries invoked, we create a function predict () in python file app.py, which follows the flowas below:

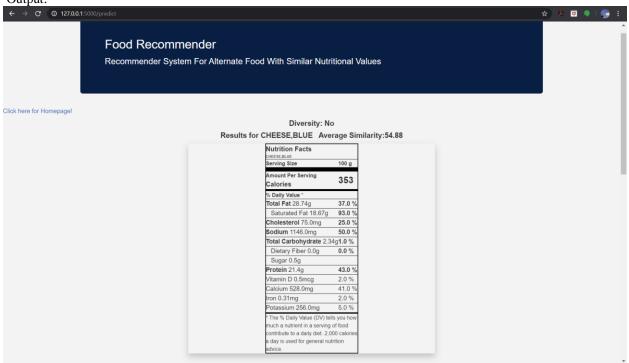
- 1. Load Welcome page 'welcome.html' to the browser
- 2. The values of daily nutrients are preloaded (Changes optional, if any changed)
- 3. Load files finalized_model.sav and finalized_model_dist.sav
- 4. Read data from file 'Preprocessed_data.csv'
- 5. When Food item is input on the 'welcome.html' page, initialize empty lists for all nutrient columns
- 6. Read Inputs 'Food Item' and 'Diversity'
- 7. Check if the given food name is present in the Preprocessed_dataset.csv
- 8. If present, Fetch the values of various nutrients from the dataset
- 9. If daily value is a non-zero entity, compare each of the nutrients with daily values and calculate the percentage
- 10. Calculate the similarity percentage
- 11. If input for Diversity = 'YES', then show the items which have accuracy of less than 40%
- 12. If input for Diversity = 'NO', show the top 8 nearest neighbor food items
- 13. Zip each of the nutrient and values under food items
- 14. Return the list in a formatted template on the 'results.html' page

The output page for a search for food item 'Cheese,Blue' is as displayed:

1. With Diversity = No



Output:



Recommended alternate foods which are similar to CHEESE,BLUE in nutritional values:

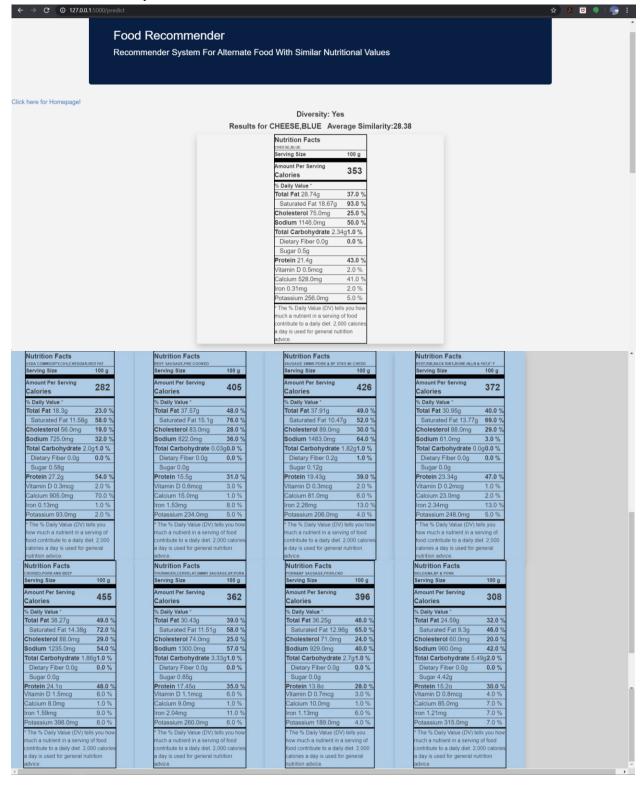
CHEESE, PAST PROCESS, PIMENTO Serving Size	100 g
Amount Per Serving	075
Calories	375
% Daily Value *	
Total Fat 31.2g	40.0 %
Saturated Fat 19.66g	98.0 %
Cholesterol 94.0mg	31.0 %
Sodium 915.0mg	40.0 %
Total Carbohydrate 1.73	g 1.0 %
Dietary Fiber 0.1g	0.0 %
Sugar 0.62g	
Protein 22.13g	44.0 %
Vitamin D 0.5mcg	2.0 %
Calcium 614.0mg	47.0 %
Iron 0.42mg	2.0 %
Potassium 162.0mg	3.0 %
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Nutrition Facts CHEESE,LIMBURGER	
Serving Size	100 g
Amount Per Serving	007
Calories	327
% Daily Value *	
Total Fat 27.25g	35.0 %
Saturated Fat 16.75g	84.0 %
Cholesterol 90.0mg	30.0 %
Sodium 800.0mg	35.0 %
Total Carbohydrate 0.49	
Dietary Fiber 0.0g	0.0 %
Sugar 0.49g	
Protein 20.05g	40.0 %
Vitamin D 0.5mcg	2.0 %
Calcium 497.0mg	38.0 %
Iron 0.13mg	1.0 %
Potassium 128.0mg * The % Daily Value (DV) tell:	3.0 %
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Nutrition Facts CHEESE, FONTINA	
Serving Size	100 g
Amount Per Serving	389
Calories	309
% Daily Value *	
Total Fat 31.14g	40.0 %
Saturated Fat 19.2g	96.0 %
Cholesterol 116.0mg	39.0 %
Sodium 800.0mg	35.0 %
Total Carbohydrate 1.5	5g 1.0 %
Dietary Fiber 0.0g	0.0 %
Sugar 1.55g	
Protein 25.6g	51.0 %
Vitamin D 0.6mcg	3.0 %
Calcium 550.0mg	42.0 %
Iron 0.23mg	1.0 %
Potassium 64.0mg	1.0 %
* The % Daily Value (DV) tel	ls you how
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2. With Diversity = Yes



V. Conclusion and Future Scope

From the above example, it is evident that when we focus on diversity, because of the lack of knowledge-discovery in the nutrition domain, there is a loss of accuracy on using string matching. However, with diversity = No, this project illustrates that with a little deviation on the nutrient values, we can model a recommender system to find alternate food items with similar nutritional values, that can substitute the food item in question. This system can be employed to find alternatives to food which may cause intolerance, for ex. A lactose intolerant individual can replace cheese with tofu which is similar in nutritional value.

With a lot of research in the field of recommender systems, we can further look to use a more powerful type of recommenders, such as a hybrid recommender system, which would take not just the similarity distance, but multiple parameters into account, thereby increasing diversity while maintaining the accuracy of the system.

VI. References

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