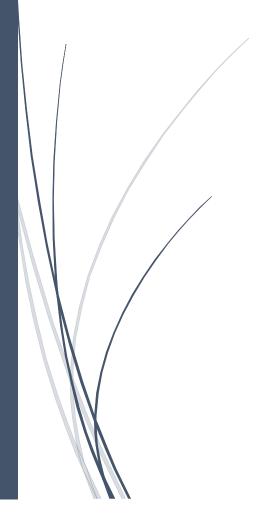
3/9/2021

Machine Learning and Content Analytics

MINI-PROJECT



VASILOPOULOU-CHONDROU ELESA P2822028 VLACHANTONI EVANGELIA P2922002 GKOTSI MARIA-ELENI P2822029

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1. Introduction

1.1. A few words for our "Machine Learning team project"

Our team was established in July 2021 for the needs of Machine Learning mini-project and consists of three members, all of them women, each of one aiming to contribute from her point of view and field of study in this accomplishment. More specifically, Mrs Vassilopoulou-Chondrou Elessa, as Statistician and Business Analyst Consultant, Mrs Vlachantoni Evangelia, Economist and Payroll manager and Mrs Maria-Eleni Gkotsi, Software Engineer and CRM Consultant.

1.2. What is the main idea

The main idea of this project is to predict structure or argument labels when examining abstracts of published articles of different scientific fields. Project's approach will focus first on creating a baseline intuition that will classify each line of abstract to a specific argument label and then by the use of neural networks will try to create a model that predicts argument and structure labels. There are many ways to proceed into this venture. Three main ways are Fasttext Approach (Version 1 and 2), Custom Approach and finally Transformer Network Approach one of which will be applied and analyzed in depth. The final step will be Abstract Clustering with the use of k-means method, trying to reveal main groups of scientific abstract's content.

1.3. What are our goals and expectations from our research

Through this research we expect to discover the power of neural networks more specific the application on text recognition, to enrich our skills in Python Programming language and to learn from each other. We expect to create a model that predicts structure and argument labels accurately with low level of error, to depict the groups of scientific abstracts and finally to compare our results with our baseline intuition created as first step.

2. Methodology

2.1 Data Collection

All abstracts were collected and uploaded in Label Studio (Figure 1) where we conducted annotation procedure in over 100 scientific abstracts without collaboration only by following tutor's instructions about main abstract's structure and basic rules for this kind of text annotation. This phase of methodology, was a necessary step in order to create our dataset in following steps.

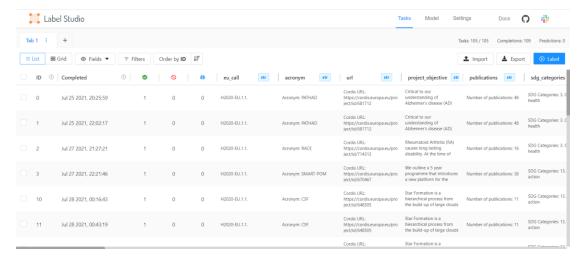


Figure 1 Data Collection in the Label Studio

2.2 Description of the data

Our data splits into three different datasets. The first one is containing argument labels (Figure 2), the second one structure labels (Figure 3) and the last one citations labels (Figure 4). All three datasets contain three columns named Document, Sentences and Labels. The first column of each dataset contains the doi number of each abstract, the second column contains the sentences and last column the labels of each category. For example, the labels of argument are divided into two subcategories: a) Claim, b) Evidence. Structure's labels are divided into five different subcategories: a) Background, b) Objective/Aim c) Method d) Result e) Conclusion. Depending on the keywords we choose to annotate each sentence with a specific structure label. The basic structure of a scientific abstract is the one mentioned above from a to e. As far as concerned argument labels

were combined either with conclusion or result label (claim and evidence respectively). If there were not any claim in the abstract it was difficult to find evidence.

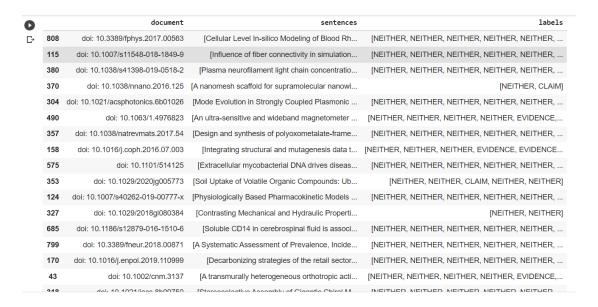


Figure 2 Argument dataset

	document	sentences	labe
330	doi: 10.1029/2018ms001327	[The Relative Influence of Atmospheric and Oce	[NEITHER, BACKGROUND, OBJECTIVE, OBJECTIVE, ME
79	doi: 10.1007/s00125-016-4184-0	[The heritable basis of gene-environment inter	[NEITHER, BACKGROUND, OBJECTIVE, METHOD, RESU
199	doi: 10.1016/j.jad.2019.04.080	[Reduced vascular endothelial growth factor le	[NEITHER, BACKGROUND, BACKGROUND, METHOD, RES
78	doi: 10.1007/s00125-016-4094-1	[Serum endotrophin identifies optimal responde	[NEITHER, BACKGROUND, BACKGROUND, OBJECTIVE,
448	doi: 10.1038/s41598-020-57790-5	[Unraveling the choice of the north Atlantic s	[NEITHER, BACKGROUND, BACKGROUND, METHOD, RES
650	doi: 10.1161/circimaging.118.007753	[Complex Congenital Heart Disease Associated W	[NEITHER, BACKGROUND, BACKGROUND
229	doi: 10.1016/j.redox.2018.04.018	[Protection against gamma-radiation injury by	[NEITHER, BACKGROUND, BACKGROUND, METHOD, MET
369	doi: 10.1038/nmat4652	[Tuning the energetics and tailoring the optic	[NEITHER, BACKGROUND, OBJECTIVE, METHOD, RESU
852	doi: 10.3762/bjoc.12.276	[The digital code driven autonomous synthesis	[NEITHER, OBJECTIVE, OBJECTIVE, METHOD, CONCL
651	doi: 10.1161/hypertensionaha.119.12412	[Vitamin K-Dependent Matrix Gla Protein as Mul	[NEITHER, OBJECT
89	doi: 10.1007/s00382-018-4521-8	[A first-of-its-kind multi-model convection pe	[NEITHER, BACKGROUND, OBJECTIVE, OBJECTIVE, M
364	doi: 10.1038/ncomms10144	[A simple and versatile design concept for flu	[NEITHER, BACKGROUND, BACKGROUND, OBJECTIVE,
67	doi: 10.1002/psp4.12473	[Open Systems Pharmacology Community—An Open A	[NEITHER, BACKGROUND, OBJECTIVE, OBJECTIVE, C
661	doi: 10.1175/bams-d-19-0068.1	[LongRunMIP: Motivation and Design for a Large	[NEITHER, OBJECTIVE, BACKGROUND, OBJECTIVE, C
835	doi: 10.3390/jcm9030814	[Assessing the Implementation of Pharmacogenom	[NEITHER, BACKGROUND, OBJECTIVE, METHOD, METH
880	doi: 10.5194/bg-15-3591-2018	[Rapid mineralization of biogenic volatile org	[NEITHER, OBJECTIVE, OBJECTIVE, OBJECTIVE, OF

Figure 3 Structure dataset

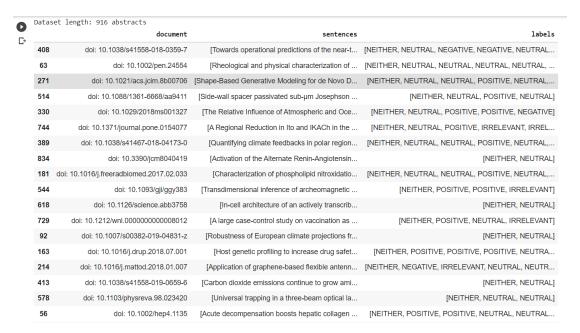


Figure 4 Citations dataset

Citation labels are divided in four different categories a) Positive b) Negative c) Neutral d) Irrelevant. The aim in this type of label is to characterize whether a citation supports (positive label) or not (negative label) the opinion or scientific analysis exposed in each abstract. With label irrelevant we characterized either citations that were not aligned with their appropriate citation number or citations written in different language or sentences containing symbols with no structure and similar sense.

2.3 Data processing

First step in our data process is to import json files with the corresponding python code. In order to start answering the first question regarding the baseline tuition, we had first to assign sentence column in separate object to create a dataframe easier to read and ready to use it for the first task. The primitive data type of each row of the column sentence was organized in list. Which means each row of sentence is a list. By using explode we created a data frame containing each sentence of type object with a correspondent document id. The same transformation happened to column labels and at the end a new data frame was prepared, named datargument, ready to be used.

2.4 Annotation

Before uploading our json dataset, in order to accomplish text recognition in upcoming assignment tasks we had to complete a previous important phase. Annotation. Among various scientific abstracts of different subjects, we had to append the appropriate label according three categories. Firstly, the structure label, whether a sentence belongs to background, previous knowledge acquired from other surveys, objective or aim of the scientific abstract, method, results and conclusion. After that, the argument label, whether a sentence already labeled with one of the previous structure could maintain a second label, specifically of evidence or claim depending the content of phrase and keywords. Finally, the last step in the annotation of each abstract, the citation's label, selecting whether a citation was positive, negative, neutral or irrelevant to the writer's analysis.

3. Experiments in Argument/Structure Prediction

3.1 Intuitive Baseline

Digging into our venture we will try to build a simple baseline based on exploratory analysis that will examine the data, for example label distribution in abstracts by using lexicon as a mean of classification that assigns labels to sentences. Our baseline intuition will be applied in argument labels forming an iterative repetition in each sentence which will examine whether there is or not a keyword justifying whether the argument label belongs to claim or evidence. Before this, an important step to this approach is to create a bug of keywords recently appeared in each argument label category. For this, with function str.split(" ") we create a dataframe with one column containing all abstract's words eliminate the factor of lower case or upper case letters, delete special characters such as commas, points etc. dropping out all duplicates, and create the list that the for loop will use to check if the sentence of the abstract is containing one of them. In the case that one word included in the list exists also in the sentence the corresponding label will be assigned.

```
[] # what we found with for loop
    cross['LABEL'].value_counts()

NONE 511
    claim 443
    Name: LABEL, dtype: int64

[] # what professor found
    cross['label'].value_counts()

CLAIM 954
    Name: label, dtype: int64
```

Figure 5 Cross check Argument Label Claim

After counting what our for loop found as claim label, we saw that from 954 sentences labeled as CLAIM we detected 443 (Figure 5), which means 46,4% of claim labels were assigned correctly, while the rest was not detected by the for loop. The total number of appended label claims using the for loop was 3997 which means 3554 labels were wrongly appended (not very encouraging).

Another approach for the labeling of Claim was to label the last sentences of an Abstract as Claim, because the majority of the last sentences of the Abstracts were Conclusion as Structure and Claim as Argument.

```
# another method to detect claim label

## LAST SENTENCES CLAIM
datatest=dataargument
datatest=datatest.drop_duplicates(subset='doc_id_y',keep='last')
datatest
```

The result here is that we have correctly appended 473 Claim Labels from the total 954 and the rest represent another label.

```
wordsevidence_df['LABEL'].value_counts()

evidence 5521

NONE 3864

Name: LABEL, dtype: int64

cross2['LABEL'].value_counts()

evidence 1191

NONE 387

Name: LABEL, dtype: int64
```

As far as evidence label is concerned, after counting what our for loop found as evidence label, we saw that from 1578 sentences labeled as EVIDENCE we detected 1017 (Figure 6), which means 64,4% of evidence labels where assigned correctly, while the rest was not detected by the for loop. The total of evidence labels obtained from the for loop were 5521 which means that 4330 were wrongly appended (not good as well).

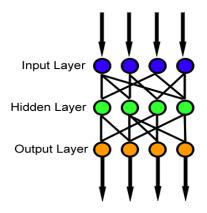
Figure 6 Cross check Argument Label Evidence

Another approach for the labeling of the Evidence was to create a list with all the symbols which could be referred to evidence and find the sentences in which these symbols are included.

```
sym = ["%","*","<",">","+","β","=","±","‰","Ξ2","Ξ","2"]
```

The result here was that from the 1578 Evidences, this method found 921 Evidence sentences and from them the 405 were assigned correctly. It seems that this method could be more effective.

3.2 Feed Forward Network Method



A feedforward neural network is an artificial neural network wherein connections between the nodes do not form a cycle. As such, it is different from its descendant: recurrent neural networks. The feedforward neural network was the first and simplest type of artificial neural network devised. In this network, the information moves in only one direction—forward—from the input nodes, through the hidden nodes (if any) and to the output nodes. There are no cycles or loops in the network.

Figure 7 Feed Forward Network Schema

3.2.1. Argument Prediction

After loading our dataset (dataset_aueb_argument_v1) and proceeding with basic descriptive statistics in order to better understand data's content and we isolated sentences and labels columns into two different variables. Next important step is one of the most frequent steps on a machine learning pipeline is splitting data into training, validation and test sets. The splitting process requires a random shuffle of the data followed by a partition using a preset threshold. On classification variants, you may want to use stratification to ensure the same distribution of classes on both sets. Splitting procedure occurred in two phases. First, we created train-validation and test dataset, and afterwards we continued with splitting train-validation into two separated datasets, the train and the validation one. In order to make the algorithm work, we had to encode our y variable in both three sets by using OneHotEncoder. The idea is to create arrays with 0-1 values representing whether a sentence is characterized as claim, evidence or neither. If one of the three previous options existed as label in a sentence then it was represented in the array as one. In other case 0. The procedure which is known as embeddings is a method used to represent discrete variables as continuous vectors. In the context of neural networks, embeddings are low-dimensional, learned continuous vector representations of discrete variables. Neural network embeddings are useful because they can reduce the dimensionality of categorical variables and meaningfully represent categories in the transformed space. Next step in our pre-process itinerary is the creation of tokens. As creation of tokens or Tokenization, is a way of separating a piece of text into smaller units called tokens. Here, tokens can be either words, characters, or subwords. Hence, tokenization can be broadly classified

into 3 types – word, character, and subword (n-gram characters) tokenization. Creating Vocabulary is the ultimate goal of Tokenization. One of the simplest hacks to boost the performance of the NLP model is to create a vocabulary out of top K frequently occurring words. Now, let's understand the usage of the vocabulary in Traditional and Advanced Deep Learning-based NLP methods. Traditional NLP approaches such as Count Vectorizer and TF-IDF use vocabulary as features. Each word in the vocabulary is treated as a unique feature. In Advanced Deep Learning-based NLP architectures, vocabulary is used to create the tokenized input sentences. Finally, the tokens of these sentences are passed as inputs to the model.

Model fitting is a measure of how well a machine learning model generalizes to similar data to that on which it was trained. A model that is well-fitted produces more accurate outcomes. A model that is overfitted matches the data too closely. A model that is underfitted doesn't match closely enough. In our case, regarding the pairs of loss- accuracy for train and validation dataset (Figure 9) we see that while in the train set our accuracy is constantly rising to 0.9983 on the contrary in the validation dataset is starting from 0.7293_keeping a downward trend that stops in 0.6554.

Figure 8 Count the frequency of top words

```
Epoch 1/30
200/200 - 4s - loss: 0.7555 - accuracy: 0.7336 - val loss: 0.9013 - val accuracy: 0.7293
Epoch 2/30
200/200 - 1s - loss: 0.4627 - accuracy: 0.8295 - val loss: 1.1879 - val accuracy: 0.6823
Epoch 3/30
200/200 - 1s - loss: 0.1908 - accuracy: 0.9379 - val_loss: 2.0530 - val_accuracy: 0.6886
Epoch 4/30
200/200 - 1s - loss: 0.0675 - accuracy: 0.9806 - val_loss: 2.4452 - val_accuracy: 0.6523
Epoch 5/30
200/200 - 1s - loss: 0.0365 - accuracy: 0.9920 - val loss: 2.9868 - val accuracy: 0.6729
Epoch 6/30
200/200 - 1s - loss: 0.0241 - accuracy: 0.9942 - val loss: 3.2253 - val accuracy: 0.6723
Epoch 7/30
200/200 - 1s - loss: 0.0173 - accuracy: 0.9969 - val loss: 3.4835 - val accuracy: 0.6648
Epoch 8/30
200/200 - 1s - loss: 0.0136 - accuracy: 0.9964 - val_loss: 3.7976 - val_accuracy: 0.6667
Epoch 9/30
200/200 - 1s - loss: 0.0120 - accuracy: 0.9975 - val_loss: 4.0088 - val_accuracy: 0.6830
Epoch 10/30
200/200 - 1s - loss: 0.0103 - accuracy: 0.9980 - val_loss: 3.9365 - val_accuracy: 0.6604
Epoch 11/30
200/200 - 1s - loss: 0.0091 - accuracy: 0.9980 - val_loss: 4.2149 - val_accuracy: 0.6642
Fnoch 12/30
200/200 - 1s - loss: 0.0099 - accuracy: 0.9975 - val loss: 4.0115 - val accuracy: 0.6529
Epoch 13/30
200/200 - 1s - loss: 0.0104 - accuracy: 0.9975 - val_loss: 4.4636 - val_accuracy: 0.6679
Epoch 14/30
200/200 - 1s - loss: 0.0073 - accuracy: 0.9983 - val_loss: 4.6180 - val_accuracy: 0.6698
```

Figure 9 Fit argument prediction model over 32 epochs

Loss is a value that represents the summation of errors in our model. It measures how well (or bad) our model is doing. If the errors are high, the loss will be high, which means that the model does not fit well. Otherwise, the lower it is, the better our model works. In our case, for the train set the loss is always below 0 which means that errors are not high and our model works well, while in the validation set is starting 0.9 keeping upward trend that stops in 5.8760 depicting a moderate performance (Figure 10).

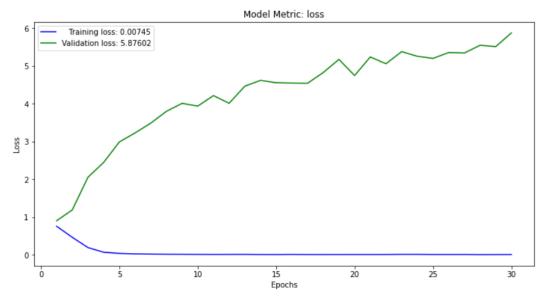


Figure 10 Model metric loss

Accuracy is more straightforward. It measures how well our model predicts by comparing the model predictions with the true values in terms of percentage which means that 99,83% our model predicts well on the train set but on the contrary in the validation dataset it predicts with 65,54% success (Figure 11). Now, if we analyze these two measurements together, we can infer more information about how is our model working.

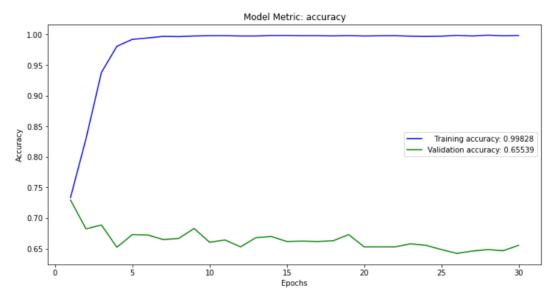


Figure 11 Model metric accuracy

Having a low accuracy but a high loss would mean that the model makes big errors in most of the data. But, if both loss and accuracy are low, it means the model makes small errors in most of the data. However, if they're both high, it makes big errors in some of the data. Finally, if the accuracy is high and the loss is low, then the model makes small errors on just some of the data, which would be the ideal case and is the category that we belong.

	Low Loss	High Loss
Low Accuracy	A lot of small errors	A lot of big errors
High Accuracy	A few small errors	A few big errors

Figure 12 Loss-Accuracy interpration

In order to evaluate our model, we are going to implement two methods, the first is confusion matrix and second one classification report. Regarding the confusion matrix, it is a performance measurement for machine learning classification problem where output can be two or more classes. It is a table with different combinations of predicted and actual values.

	CLAIM	EVIDENCE	NEITHER
CLAIM	5	10	128
EVIDENCE	5	20	212
NEITHER	20	54	954

On the vertical column we have the predicted and on horizontal we have the actual values. True positive means that we predicted positive and it is true. Here we predict 5 sentences as claim and all

Figure 13 Confusion Matrix Argument Prediction evaluation

five are true. True negative means we predicted negative and it is true. False positive (Type error 1), we predicted positive and it is false. Here we predicted 5 sentences of argument evidence that belonged to argument claim. Finally, false negative (Type 2 Error) we predicted negative and it is false.

$\stackrel{\longrightarrow}{\Box}$	precision	recall	f1-score	support
CLAIM EVIDENCE NEITHER	0.17 0.24 0.74	0.03 0.08 0.93	0.06 0.12 0.82	143 237 1028
accuracy macro avg weighted avg	0.38 0.60	0.35 0.70	0.70 0.33 0.63	1408 1408 1408

The classification report visualizer displays the precision, recall, f1-score, and support scores for the model. Precision — What percent of your predictions were correct

Figure 14 Classification Report Argument Prediction evaluation

Precision-What percent of your predictions were correct? Precision is the ability of a classifier not to label an instance positive that is actually negative. For each class, it is defined as the ratio of true positives to the sum of a true positive and false positive. Precision: Accuracy of positive predictions. Here the precision for claim is 0.17, for evidence, 0.24 and for neither 0.74 which means that the model predicts better the label "neither". **Recall -What percent of** the positive cases did you catch? Recall is the ability of a classifier to find all positive instances. For each class it is defined as the ratio of true positives to the sum of true positives and false negatives. Recall: Fraction of positives that were correctly identified. Here the recall for claim is 0.03, for evidence 0.08 and for neither 0.93 which means that 93% were correct found positive. F1 score -What percent of positive predictions were correct? The F1 score is a weighted harmonic mean of precision and recall such that the best score is 1.0 and the worst is 0.0. F1 scores are lower than accuracy measures as they embed precision and recall into their computation. As a rule of thumb, the weighted average of F1 should be used to compare classifier models, not global accuracy. Here f1 for claim is 0.06, for evidence 0.12 and for neither 0.82. Support is the number of actual occurrences of the class in the specified dataset. Imbalanced support in the training data may indicate structural weaknesses in the reported scores of the classifier and could indicate the need for stratified sampling or rebalancing. Support doesn't change between models but instead diagnoses the evaluation process.

3.2.2 Structure Prediction

Further in our analysis, we will examine the model of structure prediction concerning the labels background, objective, method, result conclusion by loading the dataset "dataset_aueb_structure_v1". The pre-process is the same with the Argument Prediction described in the previous section. After carefully integrate all above steps we tried to fit our model to see the loss-accuracy pair in train and validation dataset.

```
Epoch 1/30
    200/200 - 4s - loss: 1.5446 - accuracy: 0.3963 - val_loss: 1.8361 - val_accuracy: 0.2619
   Epoch 2/30
    200/200 - 4s - loss: 0.8905 - accuracy: 0.6922 - val loss: 2.2798 - val accuracy: 0.2594
    Fnoch 3/30
    200/200 - 3s - loss: 0.3862 - accuracy: 0.8784 - val loss: 3.1901 - val accuracy: 0.2412
    200/200 - 3s - loss: 0.1511 - accuracy: 0.9604 - val loss: 4.1090 - val accuracy: 0.2475
    Epoch 5/30
    200/200 - 3s - loss: 0.0724 - accuracy: 0.9840 - val_loss: 4.6646 - val_accuracy: 0.2475
    Epoch 6/30
    200/200 - 4s - loss: 0.0600 - accuracy: 0.9876 - val_loss: 5.0079 - val_accuracy: 0.2607
    Epoch 7/30
    200/200 - 3s - loss: 0.0421 - accuracy: 0.9917 - val_loss: 5.2915 - val_accuracy: 0.2513
    Epoch 8/30
    200/200 - 3s - loss: 0.0379 - accuracy: 0.9929 - val_loss: 5.4980 - val_accuracy: 0.2419
    Epoch 9/30
    200/200 - 3s - loss: 0.0354 - accuracy: 0.9926 - val loss: 5.6905 - val accuracy: 0.2581
    Epoch 10/30
    200/200 - 4s - loss: 0.0310 - accuracy: 0.9925 - val_loss: 5.8596 - val_accuracy: 0.2362
    Epoch 11/30
    200/200 - 3s - loss: 0.0267 - accuracy: 0.9934 - val_loss: 6.0078 - val_accuracy: 0.2400
    Epoch 12/30
    200/200 - 3s - loss: 0.0227 - accuracy: 0.9947 - val loss: 6.3011 - val accuracy: 0.2393
    Epoch 13/30
    200/200 - 4s - loss: 0.0231 - accuracy: 0.9953 - val loss: 6.4256 - val accuracy: 0.2362
    Epoch 14/30
    200/200 - 4s - loss: 0.0275 - accuracy: 0.9936 - val_loss: 6.4774 - val_accuracy: 0.2425
    Epoch 15/30
    200/200 - 3s - loss: 0.0295 - accuracy: 0.9928 - val_loss: 6.5884 - val_accuracy: 0.2456
```

Figure 15 Fit structure prediction model over 32 epochs

In comparison to the previous fit of argument prediction here we see that while loss-accuracy pair in train dataset is going well, in the validation loss accuracy pair we do not meet the same results. In other words, in our case above which number the error is considered to be low or big. Here the 6,5 comparing with other kind of errors in statistics we would say that is a small error accompanied with small accuracy, thus in this case our model makes small errors.

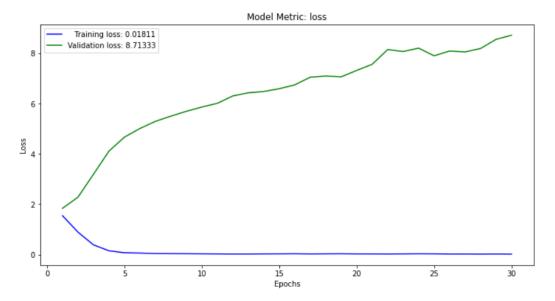


Figure 16 Loss metric structure prediction

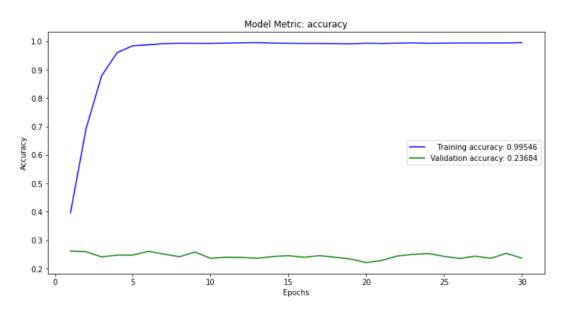


Figure 17 Accuracy structure prediction

₽		BACKGROUND	CONCLUSION	METHOD	NEITHER	OBJECTIVE	RESULT
	BACKGROUND	63	22	33	23	40	104
	CONCLUSION	36	16	14	9	25	60
	METHOD	36	16	25	11	39	87
	NEITHER	25	9	14	20	21	48
	OBJECTIVE	29	16	31	17	81	86
	RESULT	54	25	39	21	47	166

Figure 18 Confusion Matrix for Structure Prediction

₽	precision	recall	f1-score	support
BACKGROUND	0.26	0.22	0.24	285
CONCLUSION	0.15	0.10	0.12	160
METHOD	0.16	0.12	0.14	214
NEITHER	0.20	0.15	0.17	137
OBJECTIVE	0.32	0.31	0.32	260
RESULT	0.30	0.47	0.37	352
accuracy			0.26	1408
macro avg	0.23	0.23	0.22	1408
weighted avg	0.25	0.26	0.25	1408

Structure prediction model provided slightly better results in the prediction of result objective and result label in comparison to other labels but still with many errors. This claim it is identified also by the recall ratio where the 31% for objective is predicted correctly and the 47% for result is predicted correctly as well.

Figure 19 Classification Report for structure prediction

It is important to mention that when we change the epochs from 32 to 60, the performance of the model was improved significantly.

3.3 Abstract Clustering

3.3 K-means

In this part of the assignment, we will use the dataset named "dataset_aueb_argument_v2" which it contains two more columns, the eu_call and the project objective. Our aim is to use any of the embeddings created for Abstract, Project objective (each abstract belongs to a project), EU Call (each project belongs to an EU call), Claim only or Claim and Evidence to create clusters using embeddings from the abstract or a combination of features of any of the above mentioned categories.

3.3.1a Elbow method

The KElbowVisualizer implements the "elbow" method to help data scientists select the optimal number of clusters by fitting the model with a range of values for K. If the line chart resembles an arm, then the "elbow" (the point of inflection on the curve) is a good indication that the underlying model fits best at that point. In our case, we do not meet this point, thus we do not have clusters among abstracts or their combinations with aforementioned features.

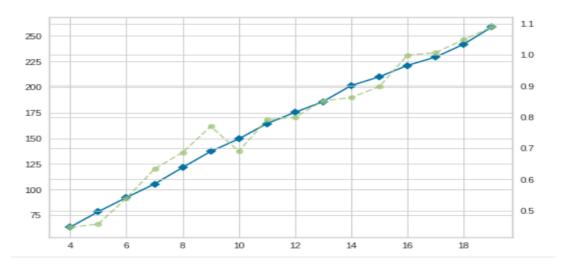


Figure 20 Elbow method for abstract embeddings

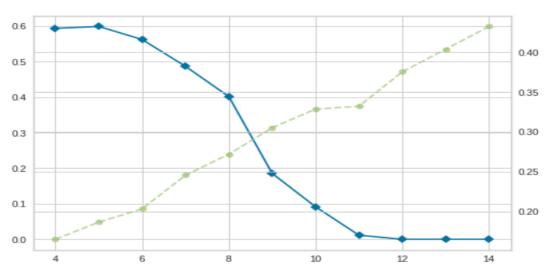


Figure 21 Elbow method clustering with eu_call embendings

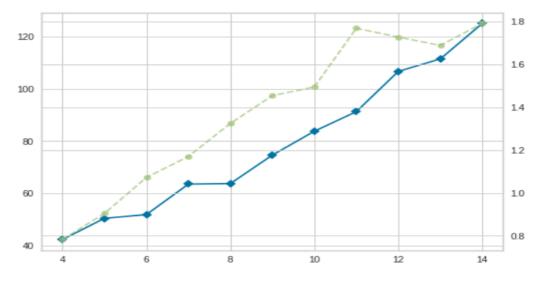


Figure 22 Elbow method with all document embeddings and claim

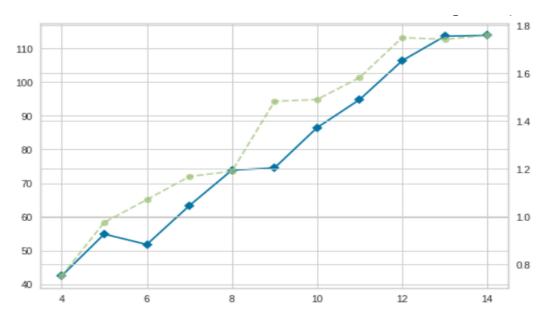


Figure 23 Elbow method with all document embeddings and claim evidence

3.3.1b Silhouette method

Silhouette analysis can be used to study the separation distance between the resulting clusters. The silhouette plot displays a measure of how close each point of one cluster is, related to the points of the neighboring clusters and thus provides a way to assess parameters, like number of clusters visually. This measure has a range of [-1, 1]. A value of 0 indicates that the sample is on or very close to the decision boundary between two neighboring clusters. Negative values indicate that those samples might have been assigned to the wrong cluster. The silhouette score falls within the range [-1, 1]. In other words, the silhouette score of 1 means that the clusters are very dense and nicely separated. The score of 0 means that clusters are overlapping. The score of less than 0 means that data belonging to clusters may be wrong/incorrect.

Our result is that all cases belong to the category that all abstracts' clusters are overlapping, thus it is difficult to organize scientific subject by examining their sentences.

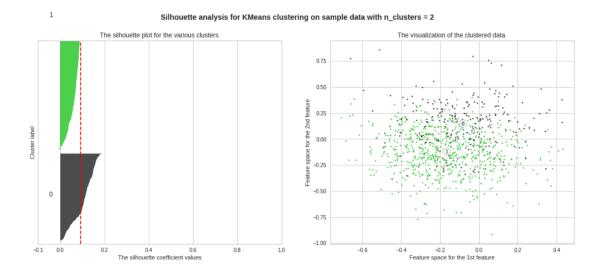


Figure 24 Silhouette method for abstract embeddings k=2

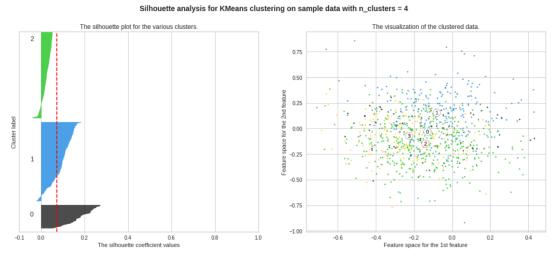


Figure 25 Silhouette method for abstract embeddings k=4

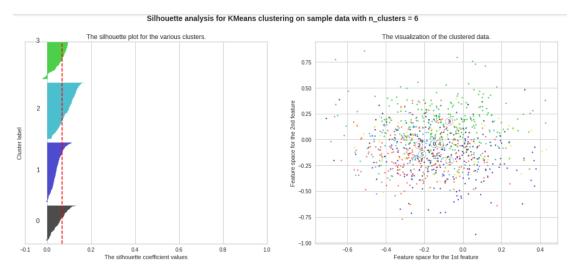


Figure 26 Silhouette method for abstract embeddings k=6

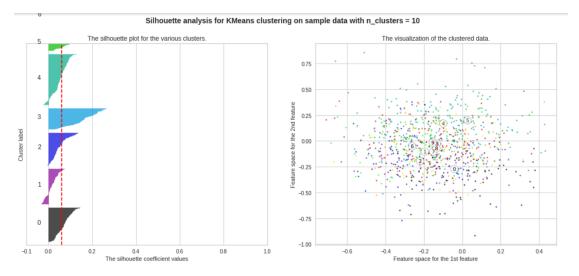


Figure 27 Silhouette method for abstract embeddings k=10

Then, we repeat the same process for the abstracts and features. Here, while the silhouette_score seems to be more acceptable (Figure 28) the clusters either are overlapping again or the size of clusters is totally different. As result, it is difficult again to examine the data.

```
For n_clusters = 2 The average silhouette_score is : 0.42654485

For n_clusters = 3 The average silhouette_score is : 0.3364397

For n_clusters = 4 The average silhouette_score is : 0.33232093

For n_clusters = 5 The average silhouette_score is : 0.32937774

For n_clusters = 6 The average silhouette_score is : 0.3360351

For n_clusters = 7 The average silhouette_score is : 0.34025684

For n_clusters = 8 The average silhouette_score is : 0.2812209

For n_clusters = 9 The average silhouette_score is : 0.29609248

For n_clusters = 10 The average silhouette_score is : 0.29856828
```

Figure 28 Silhouette Scores for the method for abstract embeddings and features



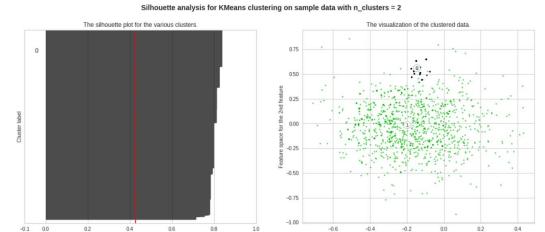


Figure 29 Silhouette method for abstract embeddings and features k=2

Figure 30 Silhouette method for abstract embeddings and features k=4

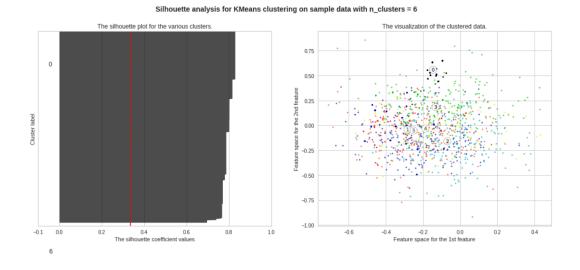


Figure 31 Silhouette method for abstract embeddings and features k=6

Silhouette analysis for KMeans clustering on sample data with n_clusters = 8

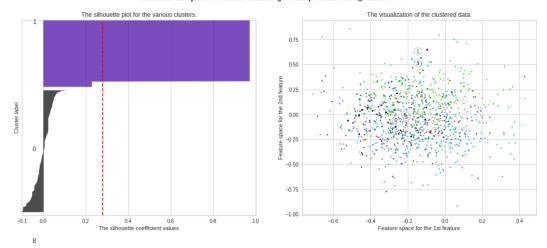


Figure 32 Silhouette method for abstract embeddings and features k=8

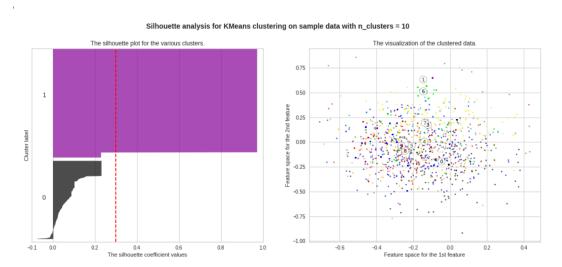


Figure 33 Silhouette method for abstract embeddings and features k=10

4. Results and Quantitave Analysis and Errors

4.1 Visualizations

All project's visualizations have been respectively incorporated in previous sections. (Figure 1- Figure 33)

4.2 Quantitave Analysis and Errors

As we show in Classification Report, the model for argument prediction could not predict well claim and evidence, in comparison to the neither label, the two first were predicted with errors. Perhaps, a reason that this event occurs, is because of the way the words were tokenized or

combined with the previous and the words after. In other words, maybe there were key-phrases all over 105 abstracts that were repeated declaring claim or evidence and our model due to its performance on keywords could not increase his success rate. The same explanation we give for structure prediction as well. Regarding errors in Clustering, the common line in both methods was that using sentences' embeddings is difficult to cluster abstracts. A reason would be that scientific abstracts utilize common words to form paragraphs, structures, methodologies etc. For this reason, when clustering method is looking to organize a bag of words in groups, it is difficult to discern which word goes in the correspondent bin. This project gave us the chance to meet and understand the power of neural networks and clustering capabilities, but as always there is space for ameliorate a prediction's model.

5. Discussion and Future goals

Machine learning journey was a great experience, from the annotation phase to the implementation of code and interpretation of result. Our feature goals as team certainly would be to investigate other algorithms of Machine learning and expand our knowledge in other kinds of recognition like image or voice.

6. Members and Roles

As we said in previous section, our team consists of three members all of them are women. All effort was deducted in team spirit, everyone did anything, which means there were not any role assigned. The result of this assignment is a collective effort, aiming to return knowledge and useful information to his members and the scientific community as well.

Figure 3x Time Plan

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