$\begin{array}{c} \textbf{IAML-INFR10069 (LEVEL 10):} \\ \textbf{Assignment} \ \#2 \\ \\ \$8141215 \end{array}$

PART A: 20-NEWSGROUPS [60 POINTS]

Question 1: (10 points) Exploratory Analysis

We will begin by exploring the Dataset to get some insight about it.

[1.1] (5 points) Focusing first on the training set, summarise the key features/observations in the data: focus on the dimensionality, data ranges, feature and class distribution and report anything out of the ordinary. What are the typical values of the features like?

Table Here		

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Your Answer Her	:e		

[1.2] (3 points) Looking now at the Testing set, how does it compare with the Training Set (in terms of sizes and feature-distributions) and what could be the repurcussions of

ust the frequency of times a word appears in a document as a feature?	
Your Answer Here	

[1.3] (2 points) Why do you think it is useful to consider TF-IDF weights as opposed to

Question 2: (24 points) Unsupervised Learning

We will now explore the documents in some detail by way of clustering.

[2.1] (2 points) The K-Means algorithm is non-deterministic. Explain why this is, and how the final model is selected in the SKLearn implementation of KMeans.

Your Answer Here			

[2.2] (1 point) One of the parameters we need to specify when using k-means is the number of clusters. What is a reasonable number for this problem and why?
Your Answer Here

[2.3] (5 points) We will use the Adjusted Mutual Information (AMI) i.e. adjusted_mutual _info_score between the clusters and the true (known) labels to quantify the performance of the clustering. Give an expression for the MI in terms of entropy. In short, describe what the MI measures about two variables, why this is applicable here and why it might be difficult to use in practice. Hint: MI is sometimes referred to as Information Gain: note that you are asked only about the standard way we defined MI and not the AMI which is adjusted for the size of the domain and for chance agreement.

Your Answer Here		

[2.4] (4 points) Fit K-Means objects with n_clusters ranging from 2 to 12. Set the random seed to 1000 and the number of initialisations to 50, but leave all other values at default. For each fit compute the adjusted mutual information (there is an SKLearn function for that). Set average_method='max'. Plot the AMI scores against the number of clusters (as a line plot).

Your Image Here

2.5] (3 points) Discuss any trends and interesting aspects which emerge from this follow from your expectations?	om the plot.
Your Answer Here	

[2.6] (6 points) Let us investigate the case with four (4) clusters in some more detail. Using seaborn's **countplot** function, plot a bar-chart of the number of data-points with a particular class (encoded by colour) assigned to each cluster centre (encoded by position on the plot's x-axis). As part of the cluster labels, include the total number of data-points assigned to that cluster.

Your Image Here	

2.7] (3 points) How doe looes it conform to your	_	- ()	with the true class l	abels
Your Answer Here				

Question 3: (26 points) Logistic Regression Classification

We will now try out supervised classification on this data. We will focus on Logistic Regression and measure performance in terms of the F1 score (familiarise yourself with this score which is related to the precision and recall scores that we learnt about in class).

[3.1] (3 points) What is the F1-score, and why is it preferable to accuracy in our problem? How does the macro-average work to extend the score to multi-class classification?

Your Answer Here			

average for the f1_score.					
Your Answer Here					

[3.2] (2 points) As always we start with a simple baseline classifier. Define such a classifier (indicating why you chose it) and report its performance on the **Test** set. Use the 'macro'

discussed in the lectures.			
Your Answer Here			

[3.3] (3 points) We will now train a LogisticRegression Classifier from SKLearn. By referring to the documentation, explain how the Logistic Regression model can be applied to classify multi-class labels as in our case. *Hint: Limit your explanation to methods we*

Comment on the result.			
Your Answer Here			

[3.4] (4 points) Train a Logistic Regressor on the training data. Set solver='lbfgs', multi_class='multinomial' and random_state=0. Use the Cross-Validation object you created and report the average validation-set F1-score as well as the standard deviation.

[3.5] (5 points) We will now optimise the Regularisation parameter C using cross-validation. Train a logistic regressor for different values of C: in each case, evaluate the F1 score on the training and validation portion of the fold. That is, for each value of C you must provide the training set and validation-set scores per fold and then compute (and store) the average of both over all folds. Finally plot the (average) training and validation-set scores as a function of C. Hint: Use a logarithmic scale for C, spanning 19 samples between 10^{-4} to 10^{5} .

Your Image Here	

-	 pening in your plot from Question 3: pgisticRegression page on SKLearn.	(e) Hin
Your Answer Here		

[3.6] (7 points) What is the optimal value of C (and the corresponding score)? How did you choose this value? By making reference to the effect of the regularisation parameter

on the entire training set (that is drop the folds). <i>H</i> = 200. Comment briefly on the result.	Hint:	You may n	eed to se	et max_iter
Your Answer Here				

[3.7] (2 points) Finally, report the score of the best model on the test-set, after retraining

PART B: BRISTOL AIR-QUALITY [90 POINTS]

Question 4: (30 Points) Exploratory Analysis

We will begin by exploring the Dataset to familiarise ourselves with it.

[4.1] (6 points) Summarise the key features/observations in the data: describe the purpose of each column and report (briefly) also on the dimensionality/ranges (ballpark figures only, and how they compare across features) and number of sites, and identify anything out of the ordinary/problematic: i.e. look out for missing data and negative values. Why are the latter unreasonable in such a dataset? *Hint: Refer to the documentation for how to interpret the pollutant values*.

Your Answer Here		

[4.2] (6 points) Repeat the same analysis but this time on a per-site basis. Provide a
table with the number of samples and percentage of problematic samples (negative and
missing) in each site. To report numbers, count a row which has at least one missing
entry as having missing data, and similarly for negative entries. Hint: Pandas has a
handy method, to_latex(), for generating a latex table from a dataframe.

Your Table Here		

4.3] (4 points) Briefly summarise how the sites compare in terms of number of san amount of problematic samples.	nples
Your Answer Here	

[4.4] (3 points) Given that the columns are all oxides of nitrogen and hence we expect them to be related, we will now look at correlations in our data. This will also be useful in determining how well we can predict any one of the readings from the other two. Remove the data from sites 3 and 15 and compute the **Pearson** correlation coefficient between each of the three pollutant columns on the remaining data. Visualise the coefficients between each pair of columns in a table.

Your Table Here		

[4.5]	(2 points) Comment on the level of correlation between each pair of pollutants.
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[4.6] (5 points) For each of the three pollutants, compute the Pearson correlation between sites. *Hint: You will need to remove the 'Date Time' column and then group by the first level of the columns.* Then plot these as three heatmaps: show the values within the

in the other ga	ses or if there is so	omething differen	nt.	
Your Answe	er Here			

[4.7] (4 points) Comment briefly on your observations from Question 4:(f): start by summarising the results from the NO gas and then comment on whether the same is observed

Question 5: (19 Points) Principal Component Analysis

One aspect which we have not yet explored is the temporal nature of the data. That is, we need to keep in mind that the readings have a temporal aspect to them which can provide some interesting insight. We will explore this next.

[5.1] (1 point) Plot the first 5 lines of data (plot each row as a single line-plot).

Your Image Here	

[5.2] (5 points) We will focus first on data solely from Site 1. Extract the data from this site, and run PCA with the number of components set to 72 for now. Set the random_state=0. On a single graph plot: (i) the percentage of the variance explained by each principal component (as a bar-chart), (ii) the cumulative variance (line-plot) explained by the first n components: (Hint: you should use twinx() to make the plot fit), and, (iii) mark the point at which the number of components collectively explain at least 95% of the variance (using a vertical line). Hint: Number components starting from 1.

Your Image Here	

[5.3] (2 points) Interpret and summarise the above plot.	
Your Answer Here	

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[5.4] (5 points) Generate three figures, one for the mean and one for each of the first 2 principal components: in each, plot the mean/component as three lines, one for each pollutant

[5.5] (6 points) Focusing on the mean and first principal component, are there any signifi-
cant patterns which emerge throughout the day? Hint: Think about car usage throughout
the day. What is different when interpreting the mean versus the first component? Hint:
Do peaks signify the same thing in both cases? Looking at the principal components
only, are there any significant differences between the pollutants? Why could this be
happening? Hint: You can refer to one of the limitations of PCA.

Your Answer Here		

Question 6: (41 points) Regression

Given our understanding of the correlation between signals and sites, we will now attempt to predict the NOx level for Site 17 given the value at the other sites. We will evaluate our models using the Root Mean Squared Error (RMSE) i.e. the square root of the mean squared error score by sklearn.

[6.1] (2 points) First things first: since we are dealing with a supervised task, we will need to split our data into a training and testing set. Furthermore, since some of our regressors will involve hyper-parameter tuning, we will also need a validation set. Use the multi_way_split() method from mpctools.extensions.skext to split the data into a Training (60%), Validation (15%) and Testing (25%) set: use the ShuffleSplit object from sklearn for the splitter. Set the random state to 0. Hint: The method gives you the indices of the split for each set, which can then be applied to multiple matrices. Report the sizes of each dataset.

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YOUR	Answer	нere

he training and validation sets. Interpret this relative to the statistics of the data.
Your Answer Here

[6.2] (4 points) Let us start with a baseline. By using only the y-values, what baseline regressor can you define (indicate what it does)? Implement it and report the RMSE on

with LinearRegression. Train the regressor on the training data and the training and validation set, and comment on the relative performance.	*
Your Answer Here	

[6.3] (3 points) Let us now try a more interesting algorithm: specifically, we will start

[6.4] (5 points) We want to explore further what the model is learning. Explain why in Linear Regression, we cannot just blindly use the weights of the regression coefficients to evaluate the relative importance of each feature, but rather we have to normalise the features. By referring to the documentation for the LinearRegression implementation in SKLearn, explain what the normalisation does and how it helps in comparing features. Will this affect the performance of the Linear Regressor?

Your Answer Here		

4:(f), and why do you think that is?		-	
Your Answer Here			

[6.5] (5 points) Retrain the regressor, setting normalize=True and report (in a table) the ratio of the relative importance of each feature. Which is the most/least important site? How do they compare with the correlation coefficients for Site 17 as computed in Question

[6.6] (5 points) It might be that with non-linear models, we may get better performance. Let us try to use K-Nearest-Neighbours. Train a KNN regressor with default parameters on the training set and report performance on the training and validation set. Hint: it might be beneficial to set $n_{jobs=-1}$ to improve performance. How does it compare with Linear Regression in terms of performance on both sets? What is a limitation of the KNN algorithm for our dataset?

Your Answer Here			

will optimise only one: the number of neighbours to use. By using the validation set, find the optimal value for the n_neighbours parameter out of the values [2, 4, 8, 16, 32]. Plot the training/validation RMSE and indicate (for example with a line) the best value for n_neighbours. Your Image Here

[6.7] (4 points) The KNN regression allows setting a number of hyper-parameters. We

$\left[6.8\right]$ (1 points) What is the best-case RMSE performance	on the validation set for KNN?
Your Answer Here	

[6.9] (4 points) Let us try one last regression algorithm: we will now use DecisionTreeRegressor. Again, the algorithm contains a number of hyper-parameters, and we will optimise the depth of the tree. Train a series of Decision Tree Regressors, optimising (over the validation set) the max_depth over the values [2, 4, 8, 16, 32, 64]. Set random_state=0. Plot the training/validation RMSE and indicate (as before) the best value for max_depth.

Your Image Here

do you notice from the plot about the performance of the Decision Tree Regressor?
Your Answer Here

[6.10] (3 points) What is the best-case RMSE performance on the validation set? What

[6.11] (5 points) To conclude let us now compare all the models on the testing set. Com-
bine the training and validation sets and retrain the model from each family on it: in
cases where we optimised hyper-parameters, set this to the best-case value. Report the
testing-set performance of each model in a table Hint: You should have 4 values.

Your Answer Here			