# An Univariate Feature Elimination Strategy for Clustering Based on Metafeatures

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Abstract—Feature selection plays a very important role in all pattern recognition tasks. It has several benefits in terms of reduced data collection effort, better interpretability of the models and reduced model building and execution time. A lot of problems in feature selection have been shown to be NP - Hard. There has been significant research in feature selection in last three decades. However, the problem of feature selection for clustering is still quite an open area. The main reason is unavailability of target variable as compared to supervised tasks. In this paper, five properties or metafeatures like entropy, skewness, kurtosis, coefficient of variation and average correlation of the features have been studied and analysed. An extensive study has been conducted over 21 publicly available datasets, to evaluate viability of feature elimination strategy based on the values of the metafeatures for feature selection in clustering. A strategy to select the most appropriate metafeatures for a particular dataset has also been outlined. The results indicate that the performance decrease is not statistically significant.

*Index Terms*—Feature Selection, Feature Elimination, Entropy, Skewness, Kurtosis, Coefficient of Variation, Correlation.

# I. INTRODUCTION

Feature selection is one of the most important preprocessing tasks in any data mining, machine learning, and pattern recognition process. This has several benefits like [1][2][3] – reduced data collection effort, reduced storage cost, lesser model building and execution time and better model interpretability. The interpretability of the model is a key requirement and this is one of the reasons why feature selection is often preferred over dimensionality reduction methods like Principal Component Analysis, Factor analysis etc. where the original features are transformed to generate new set of features and semantics of the features are lost [2][4]. The problem of feature selection is very relevant with the advent of Big-data, as the dimensionality of the datasets have increased significantly.

Feature selection for classification is relatively well defined, as the relevance of a feature can be estimated by its ability to predict the target or the class variable [24][25]. In case of clustering the problem is yet to be defined with equivalent clarity. So feature selection for clustering is still quite an open area of research[4]. Feature selection can be broadly categorized as filter and wrapper [29]. A filter strategy is generic and it depends on characteristics of the features or metafeatures. The wrapper on the other hand is hardwired with a learning algorithm and an optimal feature subset is obtained on the basis of algorithm's performance(Classification Accuracy, F-Score etc. for classification, DB Index, Mirkin index, rand index, Silhouette width, purity in case of clustering [26]). There has been extensive research in the domain of feature selection in last 30 years. The key motivations of the proposed work are,

- a) Feature selection, is often more important than the task itself. The practitioners, data mining and science professionals' use feature selection techniques as much as the research community. As a result, easy to interpret models are more successful and adopted, than theoretically robust complex models.
- b) Feature selection methods have been designed with a 'one size fit all' assumption. A need to analyze a dataset through its metafeatures is perceived by the authors for selecting or building an appropriate feature selection method. The appropriateness of a metafeature to be used for feature selection can be conjectured based on



generic characteristics of that metafeatures across all datasets.

Feature selection techniques are often quite involved and computationally complex. The methods for feature selection can be classified as either univariate or multivariate. A univariate method assumes the features to be independent and produces a ranked set of features. A multivariate method on the other hand employs, some goodness of a feature subset concept like Correlation Based Feature Selection (CFS) [5], minimum redundancy maximum relevance (mRMR) [6] etc. The multivariate methods are theoretically robust and they need high computational resources. Here a strategy has been discussed for feature elimination. The feature elimination is to be performed as a univariate preprocessing step before the feature selection. It is to be performed based on information theoretic and statistical properties of the features or metafeatures. Based on these metafeatures, the features are ranked and few features are eliminated. Now with these reduced set of features a multivariate method can be applied.

The methods have been examined for the unsupervised tasks and can be easily customized for supervised tasks. The different metrics that have been used are Pearson's correlation coefficient, Entropy, Skew, Kurtosis and Coefficient of variation. Reason of selecting the above metrics is that they are extensively used and well understood in the research community. It is to be noted that correlation coefficient, entropy, coefficient of variation has been found in the literature to be used for feature selection. However, no referential work could be found where skewness or kurtosis has been applied for the said task.

The intuitive guidelines for feature eliminations employing the metafeatures may be defined as follows: -

- I. Features which have low variance i.e. low coefficient of variation are candidates for elimination.
- II. Features which are relatively unrelated with other features i.e. low average correlation can be eliminated.
- III. Features which have lower entropy i.e. lesser information content can be eliminated.
- IV. Features which have highly asymmetric distribution measured by skewness are more suitable to be removed.
- V. Features with exhibit varying peaks measure in terms of kurtosis scan be eliminated

Apart from the above generic guidelines which can be applied for all datasets, an approach to select the most appropriate of the above five metafeatures have been outlined. This is arrived at, by comparing individual characteristics of a dataset, with overall characteristics of all datasets.

The organization of the rest of the paper is as follows: In Section II, a brief outline of the metrics has been given. In Section III, related works where these metafeatures have already been used is elaborated. Additionally, researches focusing on choosing a feature selection method based on characteristics of the data is also outlined. Section IV, details out the methods and materials used in the experiment. In Section V, the results of the experiments have been presented and critically discussed, with necessary statistical analysis of the results. Section VI contains conclusion with direction for future work.

### II. METRICS USED FOR FEATURE ELIMINATION

For both the filter and wrapper methods, it is important to reduce the search space of feature subsets. The different measures or meta features used for feature elimination are namely, Shanon's Entropy, Pearson's product moment correlation coefficient, Coefficient of variation, Skew and Kurtosis.

## Shanon's Entropy

For a finite sample, Shanon's Entropy is taken as  $\sum_i p(x_i) \log_b p(x_i)$ , where  $x_i$  are the values taken by random variables, and b is the logarithmic base, taken as 2 generally. The continuous variables have been appropriately discretized.

## Pearson's product moment correlation coefficient:

Pearson's product moment correlation coefficient between two variables x and y, is given by the following equation (1),

$$\rho(\mathbf{x}, \mathbf{y}) = \frac{cov(x, y)}{\sqrt{var(x) * var(y)}}$$
(1)

Few underlying assumptions are a) the relationship between x and y is linear. b) x and y are normally distributed. c) The residuals in the scatter plot are homoscedastic i.e. they are random. Correlation coefficient, has a value between -1 and +1, higher the absolute value, higher the strength of the relationship. It is also symmetric, i.e., the correlation coefficient between x and y and correlation coefficient between y and x are same. Another important property of Correlation Coefficient is it is scale invariant.

Some other measures which can be used are in place of correlation coefficient are Mutual Information, Normalized mutual information [7], Maximal Information Coefficient [8] etc.

### Coefficient of Variation:

Coefficient of variation is a measure of dispersion for any frequency distribution or probability distribution. It is given as  $Cov(x) = \mu / \sigma$ , where  $\mu$  is the arithmetic mean and  $\sigma$  is the standard deviation of the distribution. The advantage of this measure is it is expressed as a ratio to mean, however it loses significance when the variables take negative values.

## Skew:

Skew is a measure of asymmetry of a probability

distribution. For a unimodal distribution, negative skew indicates the left hand side tail is longer, while positive skew indicates the converse. It is denoted by  $\gamma 1$  and defined as  $E\left[\left(\frac{x-\mu}{\sigma}\right)^3\right]$ .

## Kurtosis:

Kurtosis is a measure of peakedness of a probability distribution. It is denoted by  $\gamma 2$  and defined as  $E\left[\left(\frac{x-\mu}{\sigma}\right)^4\right]$ . High kurtosis means sharp peak and fatter tails while low kurtosis means rounder peak and thinner tails. There are many other univariate measures. However, for keeping the discussion focused, the scope has been confined to above five popular measures.

## III. RELATED WORK

In paper [9], authors have discussed effectiveness of measures like Skewness and correlation for feature selection in a pattern recognition task dealing with statistical process control data. In paper [10], authors have discussed a feature selection technique based on clustering the coefficient of variations. As observed in paper [11], SPSS, which is a leading commercial tool for data mining by IBM recommends screening those features which has a low coefficient of variation. Another commercial tool SQL Server Analysis Service from Microsoft outlines the importance of entropy in finding interestingness or importance of an attribute [12]. There are numerous papers using correlation coefficient and Mutual Information for feature selection, however very rarely they have been utilized for feature selection in clustering [27] [28][30].

In paper [13] authors propose that, the characteristics of dataset play a role in choosing the feature selection method for classification. The different attributes which are considered for the dataset are mean correlation coefficient, mean skew, mean kurtosis and mean entropy. As a measure of central tendency, median have been used as it is more robust to outliers. Coefficient of variation has been used as a measure of dispersion in the said work.

Table 1. Classification of datasets based on MVS

| Category of Dataset | MVS Range  |
|---------------------|------------|
| Strong Independent  | < 20       |
| Weak Independent    | 20 – 72. 5 |
| Weak Correlated     | 72.5 - 150 |
| Strong Correlated   | > 150      |

In paper [14], a measure has been proposed named as 'MVS' (Multi Variate Score), which quantifies the strength of association between the variables in a dataset derived from its correlation matrix. MVS (Multivariate Score) is defined as  $MVS = \sum_{i=1}^{10} w_{1i} * w_{2i} * d_i$ , where the absolute value of all the possible pair wise correlation coefficients are picked up and then distributed in 10

buckets(0 - 0.1, 0.1 - 0.2, 0.2 - 0.3, 0.3 - 0.4, 0.4 - 0.5, 0.5 - 0.6, 0.6 - 0.7, 0.7 - 0.8, 0.8 - 0.9, 0.9 - 1.0). For further details, the said paper can be referred.

The paper advocates choosing feature selection strategies based on the dataset characteristics. Some other popular univariate measures for feature selection are Laplacian Score[15] and Spec [16] respectively. However, as these are neighborhood based methods they are computationally more expensive.

#### IV. PROPOSED METHOD

In this section, the methods that have been used for feature elimination (FEE) have been elaborated. The five metafeatures as discussed in Section II, have been computed for all the features in dataset. The features are then ranked by the value of metafeatures and then the lower ranked features are eliminated based on the elimination threshold level  $(\alpha)$ . As explained features with lower values of coefficient of variation, average correlation and entropy and higher values of skewness and kurtosis have been eliminated. At step 1, a max-min normalization to scale the feature values within the range [0, 1] has been performed. This method is preferred to other normalization techniques like z-score as it retains partial information about standard deviation [17]. The method produces 15 subsets of features for 5 metafeatures and three elimination levels respectively.

**Procedure:** Feature Elimination Exhaustive (FEE) **Input:** Dataset D **Parameter:** Elimination level α (0.1,0.2, 0.25) **Output:** FS [15][]

*Step 1:The features (F) are scaled using max-min normalization.* 

**Step 2:**Calculate Entropy, Skewness, Kurtosis, and Coefficient of Variation and average correlation of the attributes.

**Step 3:** Using the above five measures,  $\alpha$  % features are eliminated as appropriate

- For Entropy, Coefficient of Variation and Average Correlation the features with lower values are eliminated.
- For Skew and Kurtosis, features with higher values are eliminated.

*Step 4:* for each of the 15 combinations, the feature subsets are added to FS.

The notations used are as follows,

F indicates the complete feature set.

 $F_{0.1}^{En}$  indicates the feature subset produced by eliminating 10% of the features using entropy as the metric. The general form of the feature subset notation is  $F_P^M$ , where M can be any one of the five metrics, Entropy (En), Skew (Sk), Kurtosis (Kt), Coefficient of Variation (Cv) and Average Correlation Coefficient (Ac). The different levels of elimination ( $\alpha$ ) used are 0.1,

0.2 and 0.25 in the current setup.

Next a study has been conducted by computing metafeatures of all the datasets. To better analyzing datasets rather than working with individual values of the meta features, they are grouped based on the quartile values based on a concept similar to quartile clustering [18]. This technique has been applied to the first four metrics namely (Entropy (EN), Coefficient of Variation (CV), Skewness (SK) and Kurtosis (KT)). The representation scheme is elaborate d in Table 2. 'V' is the value of the metafeatures for that particular dataset and Q1, Q2, Q3 denotes quartile 1, median and quartile 3 values respectively.

Table 2. Coding strategy for datasets based on metafeatures

| Range of Value  | Code | Description    |
|---|------|----------------|
| V<= Q1  | LL   | Low low        |
| Q1 <v<=q2< td=""><td>LM</td><td>Low medium</td></v<=q2<>      | LM   | Low medium     |
| Q2 <v<=q3< td=""><td>HM</td><td>High<br/>Medium</td></v<=q3<> | HM   | High<br>Medium |
| V>Q3  | HH   | High high      |

For all the metafeatures, median values of the metafeatures for that particular datasets have been used for the comparison, with the exception of average correlation. For average correlation, MVS (Multivariate Score) of the dataset has been used and as this is already grouped the above grouping is not required for MVS.

**Procedure**: Feature Elimination Greedy (FEG) **Input**: Dataset D **Parameter**: Elimination level α **Output**: Feature Subsets [K][]

*Step 1:The features (F) are scaled using max-min normalization.* 

**Step 2:** Calculate Entropy, Skew, Kurtosis, and Coefficient of Variation and average correlation of the attributes in 'D'.

**Step 3:** The median values of all the metrics for 'D' is computed for four metafeatures and MVS value is calculated for average correlation. **Step 4:** 

- These values are coded to 'LL', 'LM', 'HM', 'HH' for Entropy, skewness, Kurtosis and Coefficient of Variation.
- For MVS, the dataset is coded as 'SI', 'WI', 'WC' and 'SC' respectively

**Step 5:** Identify the metric/metrics, which is/are either encoded as 'HH' or 'SC'

*Step 6:* Using the selected measures, 10%, 20% and 25% features are eliminated respectively

• For Entropy, Coefficient of Variation and Average Correlation the features with lower values are eliminated.

• For Skew and Kurtosis, features with higher values are eliminated

Step 7: If the criteria in step 4, results in any dataset which does not have 'HH' or 'SC' then metrics having value as 'HM' or 'WC' is chosen next. Step 9: If criteria at step5 and step 7 generate empty set then Feature Elimination Exhaustive (FEE)is performed.

#### V. METHODS AND MATERIALS

Twenty-one public datasets have been used from publicly available sources [19][20]. The computing environment that is used is 'R' [21]. Few 'R' libraries have been used for different computations. [21][22][23]. The datasets used are enclosed in Table 3a,

| Dataset  | # Records | # Features | # Class |
|----------|-----------|------------|---------|
| bands    | 365       | 19         | 2       |
| btissue  | 106       | 9          | 6       |
| CTG      | 2126      | 34         | 10      |
| Darma    | 358       | 34         | 6       |
| Dow      | 995       | 12         | 10      |
| Heart    | 270       | 13         | 2       |
| hepa     | 80        | 19         | 2       |
| Leaf     | 340       | 15         | 36      |
| magic    | 19020     | 9          | 2       |
| mdlon    | 2000      | 500        | 2       |
| optdgt   | 5620      | 62         | 10      |
| Pen      | 10992     | 16         | 10      |
| Saeheart | 462       | 9          | 2       |
| satimg   | 1166      | 18         | 7       |
| satt     | 4435      | 36         | 6       |
| Sonar    | 208       | 60         | 2       |
| Veichle  | 846       | 18         | 4       |
| waveform | 5000      | 21         | 3       |
| wbdc     | 569       | 31         | 2       |
| Wine     | 178       | 13         | 3       |
| wqwhite  | 4898      | 11         | 7       |

Table 3a. Datasets characteristic

The reason for selecting classified data is that, though there is several cluster validity measures like Silhouette Coefficient, SSE, entropy to name a few, different indices give varying amounts of emphasis on cohesion and separability and hence are subjective and difficult to compare. An external measure like purity is more objective and intuitive. Purity is defined as below,

**Purity**:  $p_{ij}$  is defined as the probability of a member of the cluster *i* belongs to the class *j*, given by  $m_{ij} / m_i$ , where  $m_{ij}$  and  $m_i$  are counts as appropriate. Now purity of a cluster *i* is by  $p_i = max_jp_{ij}$ . The overall purity is given by. $\sum_{k=1}^{i} \frac{m_i}{m} * p_i$ .

Table 3b enlists median value of the metafeatures for each dataset.

| Dataset  | Median<br>Skew | Median<br>Kurtosis | Median<br>Coefficient<br>of Variation | Median<br>Entropy | MVS    |
|----------|----------------|--------------------|---------------------------------------|-------------------|--------|
| Bands    | 0.8            | 1.68               | 0.37                                  | 1.95              | 12.46  |
| btissue  | 1.73           | 3.49               | 1.89                                  | 1.55              | 522.13 |
| CTG      | 1.66           | 3.05               | 3.85                                  | 2.02              | 49.67  |
| darma    | 1.34           | 1.3                | 1.74                                  | 0.9               | 118.47 |
| dow      | 0.19           | 1.07               | 0.31                                  | 2.96              | 673.74 |
| heart    | 0.72           | 1.43               | 0.5                                   | 1.08              | 8.4    |
| hepa     | 1.06           | 1.72               | 0.49                                  | 0.67              | 13.72  |
| Leaf     | 1.45           | 1.86               | 0.81                                  | 2.28              | 408.35 |
| mdlon    | 0.06           | 0.15               | 0.29                                  | 3.13              | 0.32   |
| mgc      | 0.86           | 2.7                | 0.55                                  | 4.04              | 200.36 |
| optdgt   | 5.82           | 169.53             | 5.12                                  | 1.73              | 15.82  |
| pen      | 0.41           | 0.98               | 0.65                                  | 4.1               | 74.84  |
| saehart  | 0.9            | 1.94               | 0.49                                  | 2.37              | 36.68  |
| sat      | 0.39           | 0.82               | 0.39                                  | 3.61              | 680.4  |
| satimg   | 1.3            | 0.84               | 1.7                                   | 2.93              | 324.69 |
| sonar    | 0.93           | 1.06               | 0.78                                  | 2.28              | 57.48  |
| veichle  | 0.5            | 0.61               | 0.56                                  | 2.95              | 538.16 |
| waveform | 0.15           | 0.45               | 0.3                                   | 3.81              | 81.31  |
| wbdc     | 1.41           | 2.96               | 0.72                                  | 2.47              | 312.69 |
| wine     | 0.3            | 0.68               | 0.5                                   | 2.23              | 74.09  |
| wqwhite  | 0.98           | 3.46               | 0.46                                  | 3.05              | 54.2   |

Table 3b. Metafeatures of datasets

From figure 1a to 1f, the distribution of the five metrics has been displayed using histogram. The histogram with kurtosis has been repeated with eliminating of very high outlying value in 1e.



Fig.1a. Histogram showing median entropy of datasets



Fig.1b. Histogram showing median Coefficient of Variations of datasets



Fig.1c. Histogram showing median Skewness of datasets



Fig.1d. Histogram showing median kurtosis of datasets



Fig.1e. Histogram showing median kurtosis of datasets after outlier removal



Fig.1f. Histogram showing MVS of datasets

The observations from the histograms and table 3b are as follows: -

- 'Optdgt' dataset, seems to be an outlier with very high value of skew, kurtosis and coefficient of variation
- 'btissue', 'dow', 'leaf', 'mgc', 'sat', 'satimg', 'wbdc' and 'vehicle' are identified as strongly correlated datasets as per the MVS score
- 'mdlon' has very low values for skew, kurtosis and coefficient of variation
- 'CTG also has a relatively high coefficient of variation.

In table 4, the datasets have been coded as per the proposed scheme in table 2.

| Dataset  | SK | КТ | CV | EN | MVS |
|----------|----|----|----|----|-----|
| Bands    | LM | HM | LL | LL | SI  |
| btissue  | HH | HH | HH | LL | SC  |
| CTG      | HH | HH | HH | LM | WI  |
| darma    | HM | LM | HH | LL | WC  |
| dow      | LL | LM | LL | HM | SC  |
| heart    | LM | LM | LM | LL | SI  |
| hepa     | HM | HM | LM | LL | SI  |
| Leaf     | HH | HM | HM | LM | SC  |
| mdlon    | LL | LL | LL | HH | SI  |
| mgc      | LM | HM | LM | HH | SC  |
| optdgt   | HH | HH | HH | LL | SI  |
| pen      | LL | LM | HM | HH | WC  |
| saehart  | LM | HM | LM | LM | WI  |
| sat      | LL | LL | LL | HH | SC  |
| satimg   | HM | LL | HH | HM | SC  |
| sonar    | HM | LM | HM | LM | WI  |
| veichle  | LM | LL | HM | HM | SC  |
| waveform | LL | LL | LL | HH | WC  |
| wbdc     | HH | HH | HM | HM | SC  |
| wine     | LL | LL | LM | LM | WC  |
| wqwhite  | HM | HH | LL | HM | WI  |

Table 4. Codified datasets

## VI. RESULTS AND DISUCSSION

This section has two parts. Initially the result using FFE has been presented for all the 5 metafeatures and 3 elimination levels. These are represented in Table 5a to Table 5e. All the tables, contain result obtained using all features in the last column.

#### Results with Entropy:

In the below table, the results using entropy for feature elimination is presented. The  $2^{nd}$  to  $4^{th}$  columns indicate purity achieved using different feature elimination level.

Table 4a. Results with Entropy

| Dataset  | a = 0.1 $a = 0.2$ $a = 25$ |         | a = 25  | All      |
|----------|----------------------------|---------|---------|----------|
| Dataset  | u – 0.1                    | u – 0.2 | u – 25  | features |
| bands    | 0.63014                    | 0.63014 | 0.63014 | 0.63014  |
| btissue  | 0.56132                    | 0.54726 | 0.5533  | 0.55623  |
| CTG      | 0.85956                    | 0.8142  | 0.77717 | 0.95912  |
| Darma    | 0.88307                    | 0.86721 | 0.86648 | 0.86763  |
| Dow      | 0.5664                     | 0.5599  | 0.55968 | 0.57545  |
| Heart    | 0.81852                    | 0.7963  | 0.81111 | 0.84433  |
| hepa     | 0.8375                     | 0.8375  | 0.8375  | 0.8375   |
| Leaf     | 0.54309                    | 0.55276 | 0.55432 | 0.54915  |
| magic    | 0.64837                    | 0.64837 | 0.64837 | 0.64837  |
| mdlon    | 0.92267                    | 0.91564 | 0.90861 | 0.91037  |
| optdgt   | 0.65835                    | 0.69756 | 0.72975 | 0.65516  |
| Pen      | 0.74327                    | 0.68377 | 0.68378 | 0.71778  |
| Saeheart | 0.65368                    | 0.65368 | 0.65368 | 0.65368  |
| satimg   | 0.6015                     | 0.65212 | 0.64066 | 0.58433  |
| satt     | 0.74679                    | 0.74453 | 0.747   | 0.74611  |
| Sonar    | 0.53365                    | 0.53365 | 0.53365 | 0.53365  |
| Veichle  | 0.38967                    | 0.38142 | 0.38771 | 0.36921  |
| waveform | 0.5264                     | 0.5268  | 0.5268  | 0.5316   |
| wbdc     | 0.92267                    | 0.91564 | 0.90861 | 0.91037  |
| Wine     | 0.96067                    | 0.95506 | 0.93258 | 0.96629  |
| wqwhite  | 0.48685                    | 0.47344 | 0.47349 | 0.47863  |

As per the above dataset, purity at all the three levels are equivalent to purity achieved with full feature set, with the exception of the dataset 'CTG' and 'Heart', where there is a drop in purity, by more than a percentage point.

Table 4b. Results with average correlation coefficient

| Datasets | α = 0.1 | $\alpha = 0.2$ | α = 25  | All     |
|----------|---------|----------------|---------|---------|
| bands    | 0.63014 | 0.63014        | 0.63014 | 0.63014 |
| btissue  | 0.58642 | 0.57358        | 0.56208 | 0.55623 |
| CTG      | 0.88315 | 0.88096        | 0.85448 | 0.95912 |
| Darma    | 0.86201 | 0.86684        | 0.86612 | 0.86763 |
| Dow      | 0.58095 | 0.56606        | 0.56579 | 0.57545 |
| Heart    | 0.83704 | 0.82593        | 0.80741 | 0.84433 |
| hepa     | 0.8375  | 0.8375         | 0.8375  | 0.8375  |
| Leaf     | 0.51944 | 0.53           | 0.50515 | 0.54915 |
| magic    | 0.64837 | 0.64837        | 0.64837 | 0.64837 |
| mdlon    | 0.91037 | 0.91037        | 0.90861 | 0.91037 |
| optdgt   | 0.65111 | 0.71644        | 0.71735 | 0.65516 |
| Pen      | 0.71241 | 0.68018        | 0.68046 | 0.71778 |
| Saeheart | 0.65368 | 0.65368        | 0.65368 | 0.65368 |
| satimg   | 0.61026 | 0.62405        | 0.65948 | 0.58433 |
| satt     | 0.74547 | 0.74611        | 0.74566 | 0.74611 |
| Sonar    | 0.54327 | 0.53365        | 0.53365 | 0.53365 |
| Veichle  | 0.37194 | 0.3885         | 0.38014 | 0.36921 |
| waveform | 0.5278  | 0.531          | 0.533   | 0.5316  |
| wbdc     | 0.91037 | 0.91037        | 0.90861 | 0.91037 |
| Wine     | 0.96067 | 0.91011        | 0.88213 | 0.96629 |
| wqwhite  | 0.45767 | 0.45532        | 0.45529 | 0.47863 |

With average correlation coefficient too, the reduction in purity is very marginal for all the three levels, so it can be said, they produce equivalent results. In fact, for datasets with high MVS, the reduced feature subsets seem to give a marginally better result on average. Only for 'wqwhite' and 'CTG' dataset, there is % drop in accuracy by more than a percentage point.

Table 4c. Results with average coefficient of variation

| Datasets | $\alpha = 0.1$ | α = 0.2 | <i>α</i> = 25 | Full    |
|----------|----------------|---------|---------------|---------|
| bands    | 0.63014        | 0.63014 | 0.63014       | 0.63014 |
| btissue  | 0.58642        | 0.57443 | 0.56208       | 0.55623 |
| CTG      | 0.98159        | 0.98188 | 0.9766        | 0.95912 |
| Darma    | 0.86388        | 0.86249 | 0.85997       | 0.86763 |
| Dow      | 0.56626        | 0.55987 | 0.55983       | 0.57545 |
| Heart    | 0.83333        | 0.82963 | 0.8037        | 0.84433 |
| hepa     | 0.8375         | 0.8375  | 0.8375        | 0.8375  |
| Leaf     | 0.54115        | 0.51718 | 0.49341       | 0.54915 |
| magic    | 0.64837        | 0.64837 | 0.64837       | 0.64837 |
| mdlon    | 0.91916        | 0.90334 | 0.89807       | 0.91037 |
| optdgt   | 0.6371         | 0.59911 | 0.551         | 0.65516 |
| Pen      | 0.70702        | 0.72491 | 0.72536       | 0.71778 |
| Saeheart | 0.65368        | 0.65368 | 0.65368       | 0.65368 |
| satimg   | 0.57155        | 0.58788 | 0.58066       | 0.58433 |
| satt     | 0.74994        | 0.74858 | 0.74837       | 0.74611 |
| Sonar    | 0.53365        | 0.53365 | 0.55288       | 0.53365 |
| Veichle  | 0.36725        | 0.37323 | 0.37096       | 0.36921 |
| waveform | 0.5284         | 0.5286  | 0.5264        | 0.5316  |
| wbdc     | 0.91916        | 0.90334 | 0.89807       | 0.91037 |
| Wine     | 0.93258        | 0.91011 | 0.92697       | 0.96629 |
| wqwhite  | 0.4791         | 0.48244 | 0.48244       | 0.47863 |

With coefficient of variation, also the results are more or less similar with results obtained from all features. More than 1% performance degradation is observed in few of the datasets. The results obtained with skew as the feature elimination metric yields equivalent purity.

Table 4d. Results with average Skew

| Datasets | $\alpha = 0.1$ | $\alpha = 0.2$ | a = 25  | Full    |
|----------|----------------|----------------|---------|---------|
| bands    | 0.63014        | 0.63014        | 0.63014 | 0.63014 |
| btissue  | 0.56509        | 0.54811        | 0.55    | 0.55623 |
| CTG      | 0.87427        | 0.86265        | 0.84915 | 0.95912 |
| Darma    | 0.85922        | 0.86564        | 0.87346 | 0.86763 |
| Dow      | 0.56221        | 0.56864        | 0.55608 | 0.57545 |
| Heart    | 0.77407        | 0.83333        | 0.76667 | 0.84433 |
| hepa     | 0.8375         | 0.8375         | 0.8375  | 0.8375  |
| Leaf     | 0.55853        | 0.54471        | 0.56118 | 0.54915 |
| magic    | 0.64837        | 0.64837        | 0.64837 | 0.64837 |
| mdlon    | 0.5748         | 0.5736         | 0.57745 | 0.91037 |
| optdgt   | 0.63954        | 0.71443        | 0.72295 | 0.65516 |
| Pen      | 0.70663        | 0.69376        | 0.66302 | 0.71778 |
| Saeheart | 0.65368        | 0.65368        | 0.65368 | 0.65368 |
| satimg   | 0.63053        | 0.62367        | 0.60309 | 0.58433 |
| satt     | 0.74561        | 0.74656        | 0.74703 | 0.74611 |
| Sonar    | 0.53365        | 0.55769        | 0.53365 | 0.53365 |
| Veichle  | 0.38369        | 0.38972        | 0.38652 | 0.36921 |
| waveform | 0.5342         | 0.53           | 0.5292  | 0.5316  |
| wbdc     | 0.92267        | 0.91564        | 0.92794 | 0.91037 |
| Wine     | 0.94382        | 0.9382         | 0.9044  | 0.96629 |
| wqwhite  | 0.45788        | 0.46419        | 0.46331 | 0.47863 |

One dataset, which has a close to 30% difference in accuracy is mdlon, which is the dataset with lowest average skew.

| Datasets | α = 0.1 | $\alpha = 0.2$ | α = 25  | Full    |
|----------|---------|----------------|---------|---------|
| bands    | 0.63014 | 0.63014        | 0.63014 | 0.63014 |
| btissue  | 0.55849 | 0.54717        | 0.5566  | 0.55623 |
| CTG      | 0.96195 | 0.86769        | 0.85884 | 0.95912 |
| Darma    | 0.8676  | 0.87793        | 0.87542 | 0.86763 |
| Dow      | 0.57709 | 0.5804         | 0.5804  | 0.57545 |
| Heart    | 0.77037 | 0.84444        | 0.84815 | 0.84433 |
| hepa     | 0.8375  | 0.8375         | 0.8375  | 0.8375  |
| Leaf     | 0.55118 | 0.55882        | 0.55206 | 0.54915 |
| magic    | 0.64837 | 0.64837        | 0.64837 | 0.64837 |
| mdlon    | 0.5628  | 0.54985        | 0.5502  | 0.91037 |
| optdgt   | 0.65187 | 0.70918        | 0.72925 | 0.65516 |
| Pen      | 0.69376 | 0.69333        | 0.69504 | 0.71778 |
| Saeheart | 0.65368 | 0.65368        | 0.65368 | 0.65368 |
| satimg   | 0.62744 | 0.59537        | 0.58293 | 0.58433 |
| satt     | 0.74927 | 0.74656        | 0.74744 | 0.74611 |
| Sonar    | 0.53365 | 0.53365        | 0.53365 | 0.53365 |
| Veichle  | 0.38771 | 0.38995        | 0.38002 | 0.36921 |
| waveform | 0.5186  | 0.5142         | 0.5152  | 0.5316  |
| wbdc     | 0.92267 | 0.91564        | 0.91564 | 0.91037 |
| Wine     | 0.96067 | 0.9606         | 0.9438  | 0.96629 |
| wqwhite  | 0.48979 | 0.48032        | 0.47997 | 0.47863 |

Table 4e. Results with average kurtosis

One dataset, which has a close to 30% difference in accuracy, is mdlon, which is the dataset with lowest average skew. The five methods are compared in the below figure, the red line indicates purity achieved by using all the features



Fig.2. Comparing performance of different feature elimination strategies.

At a summary level, methods based on Coefficient of variation and Entropy is closest to purity achieved with all features. In table 5, a paired t-test has been performed between results with all features and that obtained with the 15 different feature subsets. It can be seen, for none of the 15 settings the Null hypothesis can be rejected, at 99% significance level. Hence the feature elimination strategies do not result in any statistically significant performance degradation, which was indeed one of the objectives of the study. Entropy followed by average correlation has the highest 'p' values for hypothesis testing. The t-statistics and p-values have been listed in table 5.

| Metric   | α = 0.1                           | $\alpha = 0.2$                    | $\alpha = 0.25$                |
|----------|-----------------------------------|-----------------------------------|--------------------------------|
| Cov      | t = 0.6197<br>p-value =<br>0.5425 | t = 1.4702<br>p-value =<br>0.1571 | t = 1.8483<br>p-value = 0.0794 |
| Acor     | t = 0.8931<br>p-value =<br>0.3824 | t = 0.7817<br>p-value =<br>0.4436 | t = 1.0477<br>p-value = 0.3073 |
| Entropy  | t = 0.2678<br>p-value =<br>0.7916 | t = 0.7698<br>p-value =<br>0.4504 | t = 0.7418<br>p-value = 0.4668 |
| Skew     | t = 1.3917<br>p-value =<br>0.1793 | t = 1.0684<br>p-value =<br>0.2981 | t = 1.4722<br>p-value = 0.1565 |
| Kurtosis | t = 1.0308<br>p-value =<br>0.3149 | t = 1.0335<br>p-value =<br>0.3137 | t = 1.0761<br>p-value = 0.2947 |

Table 5. Statistical Significance

The result is illustrated, further with each individual dataset. An improvement in purity is indicated by 'W', a tie is indicated by 'D' and a loss is indicated by 'L'. The cases where, in majority equivalent or better results are obtained are marked in bold.

From table 6, it can be observed that -

- Entropy and Kurtosis has given better or equal result at 71.42% cases at 10% elimination level.
- Using covariance and average correlation coefficient the same ratio is 57.14%, at 25% level,
- metrics which gives equivalent or better results in more than 50% case are Average Correlation, Skew and Kurtosis respectively across all the three levels

Average ranks of each of the methods are computed and compared in Figure 3.

| Metric   | $\alpha = 0.1$        | $\alpha = 0.2$       | $\alpha = 25$        |
|----------|-----------------------|----------------------|----------------------|
| Cov      | W-9, D-3, L-9         | W - 7, D - 3, L - 11 | W - 7, D - 3, L - 11 |
| Acor     | W- 6, D-6, L-9        | W-5, D-6, L-10       | W-5, D-6, L-10       |
| Entropy  | W- 10, D-5, L-6       | W-6, D-5, L-10       | W-5, D-5, L-11       |
| Skew     | W - 5, D - 5 , L - 11 | W - 6, D - 4, L - 11 | W- 7, D - 5, L - 9   |
| Kurtosis | W - 10, D - 5 , L - 6 | W - 8, D - 5, L - 8  | W - 8, D - 5, L - 8  |

Table 6. W-D-L by mete features



Fig.3. Comparing average rank of different feature elimination strategies.

The red line indicates rank achieved with all features. Kurtosis and Entropy achieves the best ranks as per the analysis.

In the table below, there is one column corresponding to each metrics and the encoded values as shown in table 4 are used. The last column indicates which of the feature elimination strategies give an equivalent or better result in average over the performance achieved with full feature set. The background color of 6<sup>th</sup> column is colored in green if FEG, correctly identifies the meta feature, Amber if fails to identify and no color if FEG can't come to a decision and FEE needs to be applied.

| Dataset  | SK | KT | CV | EN | MVS | Better or Equivalent |
|----------|----|----|----|----|-----|----------------------|
| Bands    | LM | HM | LL | LL | SI  | All                  |
| btissue  | HH | HH | HH | LL | SC  | CV, AC               |
| CTG      | HH | HH | HH | LM | WI  | CV                   |
| darma    | HM | LM | HH | LL | WC  | EN                   |
| dow      | LL | LM | LL | HM | SC  | KT, AC               |
| heart    | LM | LM | LM | LL | SI  | None                 |
| hepa     | HM | HM | LM | LL | SI  | ALL                  |
| Leaf     | HH | HM | HM | LM | SC  | SK , KT              |
| mdlon    | LL | LL | LL | HH | SI  | EN                   |
| mgc      | LM | HM | LM | HH | SC  | All                  |
| optdgt   | HH | HH | HH | LL | SI  | ENT, ACOR, SK, KT    |
| pen      | LL | LM | HM | HH | WC  | CV                   |
| saehart  | LM | HM | LM | LM | WI  | All                  |
| sat      | LL | LL | LL | HH | SC  | CV, KT, SK           |
| satimg   | HM | LL | HH | HM | SC  | AC, EN, SK, KT       |
| sonar    | HM | LM | HM | LM | WI  | SK, CV               |
| veichle  | LM | LL | HM | HM | SC  | ALL                  |
| waveform | LL | LL | LL | HH | WC  | None                 |
| wbdc     | HH | HH | HM | HM | SC  | SK,KT, ENT           |
| wine     | LL | LL | LM | LM | WC  | None                 |
| wqwhite  | HM | HH | LL | HM | WI  | KT,COV               |

Table 7. Meta feature selection strategy

The below is the result of applying FEG strategy

- In 16 of the datasets has at least one measure as 'HH' or 'SC' in case of MVS metric. These 16 datasets have been color coded and among them in 12 of them, this is seen to be good strategy i.e. a 75% success rate.
- In 3 of the datasets there is a presence of 'HM' or 'WC', and in all three of them strategy suggested by FEG, gives correct result.
- For the rest 2 datasets, FEE needs to be applied.

## VII. CONCLUSION

Feature selection for clustering is quite an open problem to solve. Here a feature elimination strategy(FEE), based on ranking the features using the value of metafeatures (Statistical and Information theoretic properties of the features) has been used. From the above 15 strategies, in 9 of them, the reduced features sets have a better or equivalent result. Additionally, all these methods have a linear computational time, making them computationally less expensive. It is to be noted that in none of the methods, there is a reduction in performance which is statistically significant. Α subsequent study reveals, these metafeatures, can be in turn compared with metafeatures from all datasets, to select an appropriate metafeatures for feature elimination (FEG). It is observed that, this strategy can be successfully applied to improve results in 79% cases (15 out of 19 datasets). Overall, use of metafeatures to understand the characteristics of the dataset as well as a

feature elimination strategy seems to be quite effective. The strategies (FEE & FEG) need to be tested on more datasets and with different levels of feature elimination.

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