Evaluation of SVM Kernels for Health Risks Assessment

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Abstract
As per the statistics of the National Family Health Survey – 4 (2015-16) only 28.7% of families in India have been covered under Health Insurance. There are many categories of the population that are not covered by the insurance companies as they are reluctant to insure them. The main reason is that they are not able to compute fitness level of the people who wants to get insured properly. In this paper, the preliminary analysis of data set is given equal importance as in constructing and fine-tuning machine learning model. The selection of features for machine learning models was done based on correlation as well as on the medical significance of the attributes. The features that are medically significant and has a minimum correlation among themselves were selected for constructing SVM kernel models. The selection of the most appropriate SVM Kernel was done by multiple evaluations of six SVM kernels. It was found that the Medium Radial kernel performs best in term of accuracy but linear kernel training time is least among all kernels. The values of C-statistics are consistent with the accuracy values in almost all cases and it shows that medium radial kernel is the best choice to automate the health risk assessment as it is faster as well as most accurate in prediction.

Keywords
Support Vector Machine, Kernels, Health Risk, Classification.

Introduction
Support Vector Machine [1] [2] [3] [4][5] (SVM) is an algorithm that distinguishes boundaries between the classes within a dataset [6]. The basis of the SVM is a regression, but it is more than just finding and separating borderlines between the classes [7]. The algorithm uses a ‘kernel trick’ [8] to transform the dataset into separate planes [9]. The word support vector represents the coordinates of each individual observations mapped on the vector space model [10]. For example, if \((x_1,y_1)\) corresponds to Class A and \((x_2,y_2)\) corresponds to the Class B, and if these data points are far apart, then they can be considered as ‘support vectors’ to differentiate between the Class A and Class B. In other words, higher the distance/difference, easier it is to identify particular class of 'support vector'. Hence by definition, the aim of the SVM is to identify the separating hyperplane (optimal) which maximizes the margin of the training dataset between respective classes [11]. Secondly, the logic of the SVM algorithm(s) is to maximize the distance between the two decision boundaries of the classes that it deals with. This means increasing the separation distance by choosing the best hyperplanes [12][13]. If the boundary is non-linear due to overlapping data points then the inner product method is used to transform the data into higher feature space. This is primarily called a kernelized machine learning models but Kernelization can only happen if the kernel function(s) satisfy the condition of symmetry \((k(x,y) = k(y,x))\) and positive semidefiniteness [14]. Checking symmetry is a straightforward method but checking semi-definiteness can be done by random simulation or experimentation with the feature data. Hence, selecting the right kernel function(s) is required and this can be done using an empirical approach and evaluations on the datasets. The current narrative in every field including insurance [15] [16] [17] [18] is to apply machine learning algorithms to automate the workflows [19] [20]. For, this there is always a need to evaluate the nature of the dataset as well.

Organization of the Paper
Section 3 gives a Review of the work done whereas Section 4 explains the Scope of work and objectives. In Section 5 implementation is done and in Section 6 data collection and dataset, characteristics are discussed. Section 7 shows performance evaluation and comparison and finally, Section 8 is a conclusion and Section 9 future scope.

Review
Many researchers have used support vector machine learning algorithms to check [16][15] [18] [17], insurance claims [21] [22] and classifying the nature subjects data for Health/disease risks [23] [24] [25] . Pieces of evidence can be found about the use of support vector machine algorithms to check the level of risks associated with the person. Study of all these research reveals that there is always the need to evaluate existing algorithms for new datasets before designing new algorithms or reinventing the wheel. There are many cases where fine-tuning of parameters [26] , appropriate selection of training data [27][28] [29] and optimization of the kernel functions [30] [31] [32] of support vector machines have yielded to highly accurate results.
Combinational approaches [33] [34] [35] have also been found to improve the accuracy of the Support vector machine algorithms.

Support Vector Machine learning models have been applied to a multitude of problems due to the effectiveness of Kernel functions. In the paper [36] the researchers have processed the structured as well as unstructured data patterns. The processing of both types of data is done to construct a machine learning model that can classify and predict cerebral infarction. The authors have evaluated multiple models (unit and multi models) to arrive at an optimal learning model that produces a high degree of accuracy. The algorithms employed methods that are variants of convolution models. Elaborate experimentation of text-based data and tabular form have been shown in the results section. The result claimed in the paper show that the approach in pre-processing and selection of the algorithm for the said data set works well to produce accuracy above 94%.

This research work [37] investigates the application of machine learning about predicting heart-related issues using clinical data, that shows machine learning models such as random forest, logistic regression, neural network and gradient boosting algorithm do help in improving the predictability of the data for classification of cardio-vascular issues. Out of all these models neural networks has the maximum ability to do so as it is most accurate in solving such problems. According to the authors, the result was validated using multiple performance parameters such as AUC.

In the works of [30], the Support Vector Machine (Correlation Kernel) has been revised to supports positive semi-definite values. The outcome of the modified classifier shows better performance achieved, as compared to the KNN and Decision Tree algorithm. These comparative conclusions were made on the basis of cancer data. The generic correlation kernel may produce negative values, as certain parameters may have negative casual relationships. In order to avoid the problem of non-positive values the kernel matrix, and diagonal matrix ‘P’ is transformed into P: where all values in the matrix are always positive in nature.

Wellness health insurance [38] [39] [40] is a method of rewarding and giving a bonus to the people, who maintain good health. It is a way by which the health insurance companies and improve their operating margins as a number of claims may go down. At the same time, people can maintain their fitness level to get rewarded. This idea has been applied to do health risk analysis [41] [25] [42] [43] applicable to insurance by many researchers. The adoption is a form of health insurance depends on the advances in medical sensor [41] technologies [44]. Many pieces of evidence can be found in the contemporary literature about the use of medical sensors, cloud and IoT technologies employed in the collection of biological data [44] [45] [38] [46]. Such advances in collection and monitoring [47] of biophysical data methods can help the health insurance industry immensely according to many researchers [48] [39] [40]. But, it also evident that use such technologies need careful evaluation/assessment of the algorithms, legal bindings, and platforms for it to be successful.

The Scope of Work and Objectives

It is abundantly clear that a preliminary analysis of the dataset can help in improving the classification tasks, especially when the datasets are not benchmarked. Computing fitness levels [49], health risks [50] and proneness to the disease are requisite in “wellness health insurance”. Therefore, based on a systematic review on the problems and issues of optimization of the support vector machine [13] [32] based functions for new datasets the scope and objectives of the work can be formulated as follows:

i. Data Collection for Health Risk Assessment

ii. Analysis of health risk attributes values for gaining apriori knowledge for selection of kernels of SVMs.

iii. Application and evaluation of Kernels of SVM (six) for automating the health risk assessment.

Implementation

This section explains the implementation steps taken to achieve the aforementioned objectives. Care has been taken to follow the well-established procedures for evaluating the six kernels of the SVM for automating the Health Risk Assessment. For maintaining the quality of the dataset, all procedures related to missing values treatment, data cleaning, data grouping, and data validation have been followed during data collection and analysis. The next section explains the characteristics of Health Risk Data.

Data Collection and Dataset Characteristics

The Health Risk dataset as shown in Table 1 consists of sixteen attributes or the indicators of the health risk. The values were collected using Omron body composition machine [51]. Demographically the dataset consists of Asians located in North India, it is available at [52]. The numbers of instances are 500 and dataset does not contain any missing or any duplicate values [53]. The key objective of data collection is to leverage the use of machine learning algorithm for conducting an automated health risk assessment. The description of the dataset is as follows:

| Attributes     | Description                                      | Notation |
|----------------|--------------------------------------------------|----------|
| Birth Age      | Age of a person                                  | BA       |
| Height         | The height of a person in centimeters            | H        |
| Gender         | Male or Female                                   | G        |
| Weight         | The weight of a person in Kg                     | WT       |
| Body fat       | It is body fat mass w.r.t total body weight.     | BF       |
| Visceral Fat   | Fat around the internal organs such as the liver, pancreas, intestines etc. | VF       |
Table 1: Health Risk Attributes and Notations

| Attribute      | Range value(s) | Fitness Level class |
|----------------|----------------|---------------------|
| BMI            | 18.5 – 24.9    | FL1                 |
|                | > 25           | FL2                 |
| BP Systolic    | 100 – 120      | FL1                 |
|                | > 120          | FL2                 |
| BP Diastolic   | 65 – 80        | FL1                 |
|                | > 80           | FL2                 |
| P              | <60 - 100      | FL1                 |
|                | >100           | FL2                 |
| BF (%age)      | Male           | FL1                 |
|                | Female         | FL2                 |
|                | 10.0 – 19.9    | FL1                 |
|                | > 20.0         | FL2                 |
|                | 20.9 – 29.9    | FL1                 |
|                | > 30.0         | FL2                 |
| VF (%age)      | 1 – 9 %        | FL1                 |
|                | >10 %          | FL2                 |
| SM (%age)      | Male           | FL1                 |
|                | Female         | FL2                 |
|                | 32.9 – 35.7    | FL1                 |
|                | > 35.8         | FL2                 |
|                | 25.9 – 27.9    | FL1                 |
|                | > 28           | FL2                 |
| BDA (Yrs.)     | ± 3            | FL1                 |
|                | > 3            | FL2                 |
| Blood Sugar (mg/dL) | Fasting | 80 – 120 | FL1 |
|                | > 120          | FL2                 |
|                | After Meal (PP)| 80 – 140 | FL1 |
|                | > 140          | FL2                 |
| Waist (cm)     | (Height (cm)/2) + (0.03 * Height) | FL1 |
|                | (Height (cm)/2) > (0.03 * Height) | FL2 |

Table 2: Medical Classification of Fitness Level of the Subject Under Observation

| FL1 – Fitness Level 1 (Healthy), FL2 – Fitness Level 2 (Unhealthy) |

Table 2 gives information on the medical conditions and the logic based on which the fitness data is grouped. Care has been taken to use universal medical standards, definitions, and ranges for each fitness factor recorded.

6.1 Preliminary Analysis of Dataset

The accuracy of the machine learning algorithm depends largely on the nature of the dataset [57]. If the dataset has large numbers of dimensions, it may lead to the “curse of dimensionality” [58] and may render the performance of the machine learning model in a reduced form. Secondly, some non-significant factors also play role in reducing the overall performance where missing values and correlation between the factors also contribute [57]. The selection of SVM Kernel also depends on the shape of the distribution and linearity between the various factors [58] [32] [59] [31]. Hence, this section 6.1 gives the initial statistical observations of the dataset that would help in constructing the machine learning model.

6.2 Non-linearity and Erratic value Analysis

It is statistically and visually clear from the graph Fig.1 (a) – Fig. (l) that the data is non–linear data. But, a minimal degree of linear trend can also be seen, if a trend line is created between the data points. The dataset shape shows that linear, as well as non-linear kernels, may work up to a certain level of accuracy. But, for a full evaluation, the speed at which the algorithm learns (training time), the empirical values of prediction speed also need to be evaluated for final selection. In general sense, most of the health parameters seem to be non-linear in nature. This fact can be validated by drawing mathematical relation equations [Table 3] from the dataset using curve fitting methods. It can be seen that most of the attributes have a polynomial curve (degree 3) equation in relation to the response variable ‘y’ (y ∈ fitness Level 1 | Fitness Level 2). If the dataset is inappropriately labeled or has erratic values, the accuracy of the learning model will suffer. Hence, it appears that there is a need to choose or construct those kernels that can handle non-linear data at first glance.
Fig. 1(a) – Fig. (l) shows the Line Graph for Different Fitness Attributes

| Attributes | Fitness Level 1 Dataset | Fitness Level 2 Dataset |
|------------|-------------------------|-------------------------|
|            | f(x) = p1*x^3 + p2*x^2 + p3*x + p4 | f(x) = p1*x^3 + p2*x^2 + p3*x + p4 |
|            | Goodness of fit          | Goodness of fit          |
|            | SSE: 3.479e+04, R-square: 0.2521, Adjusted R-square: 0.24, RMSE: 11.89 | SSE: 2.971e+04, R-square: 0.02105, Adjusted R-square: 0.009114, RMSE: 10.99 |

| BA         | p1 = 2.664 (0.775, 4.553) | p1 = 1.204 (-0.5497, 2.957) |
|           | p2 = 6.76 (5.102, 8.417) | p2 = -0.02769 (-1.564, 1.509) |
|           | p3 = -7.235 (-10.94, -3.528) | p3 = -3.453 (-6.883, -0.02384) |
|           | p4 = 22.46 (20.24, 24.68) | p4 = 38 (35.95, 40.06) |

| H          | p1 = -1.77 (-3.167, -0.3737) | p1 = 0.7202 (-0.4954, 1.936) |
|           | p2 = 0.1287 (-1.097, 1.354) | p2 = -0.04148 (-1.107, 1.024) |
|           | p3 = 4.766 (2.026, 7.506) | p3 = -1.001 (-3.378, 1.377) |
|           | p4 = 162.3 (160.7, 164) | p4 = 162.6 (161.2, 164.1) |
It can be seen from Table 3 that in both the cases polynomial degree 3 equation can be used to represent the behavior of each factor and in all cases, the values of x are normalized. The second thing that can be observed from these equations is that the value of R-square remains between 0.05 to 0.33 which shows that the data is quite high in degree of non-linearity, as it is unable to do data fitting. This clearly shows that for constructing a trend in this dataset, there is a need for doing some kind of data fitting.
smoothing and interpolation. This can also be reaffirmed from the values of the R-squared test and SSE (Sum of Square Error) metric. From all these statistical tests it appears that the machine learning algorithm will have difficulty in identifying instances of each fitness class. Those kernels or function that transform the data into linear form as a preprocessing step may be able to identify data points of these two classes.

6.3 Separation and Overlapping Analysis

Scatter plots [Fig. 2(a) – Fig. 2(l)] were constructed initially to understand the nature of attributes values distribution and it was found that the Fitness Level 1 and Fitness Level 2 class instance(s) values overlap and it shall be difficult to find hard margins and boundaries between the classes for most of the support vector machine kernel functions.

It can be seen that in case of body fat Fig. 2(b), skeleton muscle Fig. 2(d) and pulse Fig. 2(i) attributes the degree of overlapping is too high and in case of weight Fig. 2(a), visceral fat Fig. 2(c) body age Fig. 2(e), BMI Fig. 2(f), BP Fig. 2(g) - Fig. 2(h), sugar level Fig. 2(j) – Fig. 2(k) and waist Fig. 2(l) the overlapping is moderate. The scatter plot of the feature dataset also showed that some of the data points are forming non-linear trend lines. And apparently, the SVM algorithm might have to deal with many outliers and employ way to ignore these outliers. In Fig. 2 (a) to Fig (l), it can be seen that the SVM will not have a linear hyperplane, to resolve such a condition. the SVM might have to introduce a virtual or additional attribute such that the separation between the classes increases to some extent. All these conditions force us to run more evaluations with many combinations of parameters of each kernel.

6.4 Correlation between the Attributes
Theoretically, adding more features should help in improving the discrimination power of the dataset. But this is not always true, especially when the features have some degree of causal relationship computed with the help of correlation. Hence, there is a need to eliminate irrelevant features if required. Correlation is one of the reliable methods to eliminate irrelevant features, especially when the correlation among the attributes is complex and high in nature. However, in the context of the current problem, it is not surprising that the Birth Age and Skeleton Muscle are highly correlated with each other. In fact, the analysis shows that they have a strong relationship among themselves as shown in Table 4.

Pearson correlation coefficient \( [60] \)

\[
r = \frac{N \sum xy - (\sum x)(\sum y)}{\sqrt{N \sum x^2 - (\sum x)^2} \sqrt{N \sum y^2 - (\sum y)^2}}
\]

where

- \( N \) = number of pairs of scores.
- \( \sum xy \) = sum of the products of paired scores.
- \( \sum x \) = sum of x scores.
- \( \sum y \) = sum of y scores.
- \( \sum x^2 \) = sum of squared of x scores.
- \( \sum y^2 \) = sum of squared of y scores.

The relationship among the response \( y = \) Fitness Level 1 | Fitness Level 2 and predictor Birth Age and Skeleton Muscle (92%) are quite high as per the Pearson correlation method matrix \([61]\). Hence, depending on which predictor is included in the learning model, the results obtained will lead to different estimation of slope when the SVM kernel will compute the boundaries. After all, the SVM is basically a regression modeling. However, for supervised machine learning tasks, the accomplishment is faulty if there is a correlation between the indicators (Weight, Blood Sugar etc.) A valid feature matrix is the one, which contains a subset of features that are highly correlated with (predictive of) the class, yet has uncorrelated with (not predictive of) each other.

### Table 4: Correlation between Health Attributes

| Birth Age | Weight | Gender | Height | Body Fat | Visual Fat | Skeleton Muscle | Body Age | BM | BMZ | BPsys | BPdia | Pulse | Sugar F | Sugar FF | Weist | Group |
|-----------|--------|--------|--------|----------|------------|----------------|----------|----|-----|-------|-------|-------|---------|---------|-------|-------|
| Birth Age | 1      | -0.06  | 0.26   | 0.36     | 0.29       | 0.53          | 0.92      | -0.78| 0.94| 0.43  | 0.39  | 0.35  | -0.04   | 0.43    | 0.32  | 0.35  |
| Height    | -0.06  | 1      | -0.48  | 0.45     | -0.24      | 0.06          | 0.37      | 0    | 0.51| -0.03 | 0.07  | 0     | 0       | 0       | 0.04 | -0.03|
| Gender    | -0.26  | -0.48  | 1      | 0.29     | 0.44       | -0.16         | -0.53     | -0.15| -0.53| -0.06 | -0.14 | -0.13 | -0.19   | -0.03   | 0    | -0.09|
| Weight    | 0.36   | 0.45   | -0.29  | 1        | 0.48       | 0.76          | -0.3       | 0.73 | 0.85 | 0.65  | 0.51  | 0.27  | 0.63    | 0.95    | 0.28 | 0.51 |
| Body Fat  | 0.29   | -0.24  | 0.44   | 0.48     | 1          | 0.56          | -0.79      | 0.62 | 0.24| 0.67  | 0.45  | 0.39  | 0.25    | 0.56    | 0.52 | 0.51 |
| Visual Fat| 0.53   | 0.05   | -0.16  | 0.76     | 0.56       | 1             | -0.32      | 0.61 | 0.62| 0.82  | 0.71  | 0.58  | 0.29    | 0.77    | 0.7   | 0.63 |
| Skeleton Muscle | 0.92 | 0.39 | -0.55 | -0.2 | -0.79 | -0.32 | 1 | -0.4 | 0.02 | -0.42 | -0.22 | -0.22 | -0.24 | -0.37 | -0.23 | 0.28 |
| Body Age  | 0.78   | 0      | -0.15  | -0.73   | 0.62       | 0.81          | -0.4       | 1    | 0.57| 0.82  | 0.67  | 0.58  | 0.2     | 0.71    | 0.63 | 0.65 |
| BM        | 0.34   | 0.51   | -0.53  | 0.85     | 0.2        | 0.62          | 0.02       | 0.57| 1   | 0.67  | 0.47  | 0.38  | 0.11    | 0.52    | 0.5   | 0.58 |
| BMZ       | 0.63   | -0.03  | -0.06  | 0.87     | 0.67       | 0.92          | -0.62      | 0.02 | 0.67| 1     | 0.67  | 0.57  | 0.3     | 0.7     | 0.96 | 0.72 |
| BPsys     | 0.39   | 0.07   | -0.14  | 0.65     | 0.45       | 0.71          | -0.22      | 0.67 | 0.47| 0.69  | 1     | 0.79  | 0.47    | 0.67    | 0.62 | 0.67 |
| BPdia     | 0.35   | 0      | 0.13   | 0.51     | 0.39       | 0.36          | -0.22      | 0.58 | 0.38| 0.87  | 0.79  | 1     | 0.39    | 0.56    | 0.53 | 0.69 |
| Pulse     | -0.04  | 0      | 0.15   | 0.27     | 0.25       | 0.25          | -0.24      | 0.4    | 0.11| 0.3   | 0.47  | 0.39  | 0.1     | 0.35    | 0.29 | 0.28 |
| Sugar F   | 0.43   | 0      | -0.03  | 0.62     | 0.58       | 0.77          | -0.37      | 0.71 | 0.52| 0.7   | 0.67  | 0.56  | 0.25    | 0.91    | 0.59 | 0.67 |
| Sugar FF  | 0.32   | 0.03   | 0      | 0.62     | 0.57       | 0.73          | 0.63       | 0.56 | 0.66| 0.62  | 0.53  | 0.29  | 0.01    | 1       | 0.56 | 0.62 |
| Weist     | 0.35   | 0.12   | -0.09  | 0.71     | 0.51       | 0.63          | -0.23      | 0.65 | 0.58| 0.72  | 0.67  | 0.51  | 0.28    | 0.6     | 0.55 | 1 |
| Group     | -0.34  | -0.01  | 0.07   | -0.71    | -0.52      | -0.76         | 0.23       | -0.68| -0.51| -0.79 | -0.02 | -0.69 | -0.40   | -0.67   | -0.62 | -0.67 |

6.5 Further Inferences and Interpretations

It is always desired that all the variables must have a direct correlation with the response variable (\( f(y) = \) Fitness Level 1 \| Fitness Level 2), and must have an insignificant correlation among themselves. The observations show in table 4 that SVM kernels will face difficulty due to this to identify the data points of each class. The computation of the Pearson correlation allows the screening of irrelevant noisy attributes. This would ultimately helps in improving the classification accuracy with highly relevant features. According to this correlation matrix attributes that have a correlation of more than 90% may be eliminated from the current feature set, but at the same time, it raises the question of the elimination process. It may happen such that:

iv. Certain parameters that are significantly relevant may not be medically relevant or dominant in declaring a person/subject category as healthy or unhealthy (Fitness Level 1 \| Fitness Level 2) \([Table 4]\). Hence, this can be stated as

\[ P \Rightarrow \text{“All medically relevant health indicators are statistically not correlated to response variable } y = \text{“Fitness Level 1 \| Fitness Level 2”} \]

v. The converse is true i.e certain parameters may be critical in making a medical decision, but statistically, in this specific dataset, they are not relevant.

\[ Q \Rightarrow \text{“All not correlated health indicators are medically relevant to response variable } y = \text{“Fitness Level 1 \| Fitness Level 2 “} \]

Therefore, by considering these two statements (P, Q) or arguments, truth table \([Table 5]\) can be constructed to show that, \( P \Rightarrow Q \) (P implies Q) and \( Q \Rightarrow P \) (Q implies P) need to be evaluated as the selection of attributes for machine learning models based on some mathematical formula with a “fallacy”, if medical facts are not considered.

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Table 5: Truth Table

| P  | Q  | P→Q | Q→P (converse) |
|----|----|-----|----------------|
| T  | T  | T   | T              |
| T  | F  | F   | T              |
| F  | T  | T   | F              |
| F  | F  | T   | T              |

Let ‘STM’ be an argument for selecting an attribute of the form P implies Q (P → Q). Then the converse of ‘STM’ is the statement Q implies P (Q → P). Table 5 shows that the statements (P, Q) and converse of the statements (P, Q) are not logically equivalent unless both terms imply each other. Therefore, for selection of attributes both must be considered. The next section gives information on the selection process based on medical criteria using Delphi method [62].

6.6 Medical Relevance

A questionnaire [63] was prepared for ranking all the indicators of health on a scale of 10 from three medical experts as per their role and influence in classifying a person/subject to fall under Fitness Level 1 or Fitness Level 2. Table 6 gives the ranking given by each respondent. The modus operandi of collected responses for this questionnaire was based on the Delphi method.

![Medical Relevancy Ranking](image)

**Fig. 3: Medical Relevancy Ranking**

The average ranking matrix shows that attributes body age, gender, weight, visceral fat, BMI, BP, pulse, sugar and waist plays a major role in classifying a person/subject under Fitness Level 1 or Fitness Level 2 medically as their score is about 60% of threshold. Therefore, based on the correlation matrix and medical relevancy test of attributes the final dataset that can be subjected to SVM can be selected using set theory as follows.
A = \{ H, G, WT, BF, VF, BDA, RM, BMI, BPsys, BPdia, P, SF, SPP, W \}, where A ϵ feature set selected on the basis of correlation.

B = \{ G, WT, VF, BDA, BMI, BPsys, BPdia, P, SF, SPP, W \}, where B ϵ the feature set selected on the basis of medical relevancy test.

A ∩ B = \{ G, WT, VF, BDA, BMI, BPsys, BPdia, P, SF, SPP, W \}

6.7 Fine Tuning Learning Models

For classification and prediction tasks the dataset is divided into three parts for ensuring the quality of results. The work begins with a right balance of proportion (50:50) of the features rows of the classes (Fitness Level 1 | Fitness Level 2). This is done to give the machine algorithm a balanced dataset for learning. If a proportion of one class set of data is more, it is likely that the machine learning algorithm may develop a bias in the learning process, and this is done in all other phases (Testing and Validation) of the algorithm. Typically, there is a need to start with default values for each SVM kernel, especially in the context of the Radial/Gaussian Kernel. The values of gamma 'g' parameter play a major role in maintaining a good accuracy at all stages. There is normally a need to scale and standardized the dataset values before they become suitable for SVM algorithms. The default values of the parameters are given in Table 7.

| S. No | Parameters                  | Medium Radial | Polynomial Degree 3 | Linear | Polynomial Degree 2 | Coarse Radial | Fine Gaussian |
|-------|-----------------------------|---------------|---------------------|--------|---------------------|---------------|-------------|
| 1     | Kernel offset (Scale)       | 3.9           | 1                   |        | 1                   | 15            | 0.97        |
| 2     | Standardized Data           |               |                     |        |                     |               |             |
| 3     | Multiclass method           |               |                     |        | Yes                 |               |             |

Table 7: Parameters of SVM Algorithms

2. Performance Evaluation and Comparison

This section gives details of six kernel algorithms evaluations and their performances in the context of the problem of health risk classification and prediction.

7.1 Time and speed Analysis of Kernels

I. Training Time

It is time taken by the algorithm to learn from the existing dataset and groups/labels given to each data point. Lower training times is desired:

\[ \text{Training Time} = \frac{\Sigma t_t}{n} \quad (2) \]

where ‘tt’ is time taken by SVM kernel to learn from the given data set in seconds and n are the number of observation.

| S. No | SVM Classifier Kernel | Training Time (sec) |
|-------|-----------------------|---------------------|
| 1     | Linear                | 7.0458              |
| 2     | Polynomial Degree 2   | 7.0847              |
| 3     | Polynomial Degree 3   | 7.5249              |
| 4     | Fine Gaussian         | 7.9644              |
| 5     | Medium Radial         | 8.5103              |
| 6     | Coarse Radial         | 8.4147              |

Table 8: Time Analysis of SVM Classifier Kernels
It is clear from the tabular data [Table 8] and bars graph that Linear kernel is taking minimum time to learn and construct hyperplanes/decision boundaries. This can be attributed to the fact that SVM transforms the data into higher dimension linearly and finding the distance between the said classes is not that hard and time-consuming as compared to other kernels. The Polynomial degree 2 algorithm is performing second and Polynomial degree 3 algorithm is in the third position.

II. Prediction Speed

It is time taken by the algorithm to predict the given dataset with respect to the actual class

| S. No | SVM Classifier Kernel | Prediction speed (obs/sec) |
|-------|-----------------------|---------------------------|
| 1     | Linear                | 6600                      |
| 2     | Polynomial Degree 2   | 9500                      |
| 3     | Polynomial Degree 3   | 14000                     |
| 4     | Fine Radial/Gaussian  | 11000                     |
| 5     | Medium Radial         | 12000                     |
| 6     | Coarse Radial /       |                           |
|       | Gaussian              | 12000                     |

Table 9: Kernel Prediction Speed Analysis

Prediction time = $\sum_{t}^{p} t_i n$ 

where ‘pt’ is time taken to predict per observation in seconds and n are the number of observations.

Higher prediction time is always desired especially when the prediction is to be done in real time. The tabular data [Table 9] and a bar graph shows that Linear kernel takes 7.0458 seconds to perform 6600 number of observations while polynomial degree 2 takes 7.0847 seconds to perform 9500 numbers of observations and Medium Radial takes 8.5103 seconds to perform 12000 of observations.

From all these outcomes, it can be concluded that in terms of training time and prediction speed Medium Radial SVM algorithm performance best.

7.2 Classification Accuracy

It is measured as the fraction of sum of the true positive instances and true negatives instance by total instances in the dataset. Table 10 gives the accuracy of each classifier kernel.
### Table 10: SVM Classifier Kernel Accuracy

| S. No | SVM Classifier Kernel  | Accuracy Percentage |
|-------|------------------------|---------------------|
| 1     | Linear                 | 96.0                |
| 2     | Polynomial Degree 2    | 97.8                |
| 3     | Polynomial Degree 3    | 97.2                |
| 4     | Fine Gaussian          | 90.4                |
| 5     | Medium Radial          | 98.2                |
| 6     | Coarse Radial          | 96.0                |

### Interpretaion

The Medium Radial SVM is giving the maximum accuracy i.e. 98.2 %, which shows that this kernel is able to identify the boundary between the two classes more distinctively and has the lowest false alarm rate as compared to the other kernels. It further shows that the Medium Radial SVM is able to select and reject the instances as per the actual class more accurately as compared to other kernels. Hence, it can be safely said that the Medium Radial SVM algorithm has the maximum predictive power among Linear, Polynomial degree 2, Polynomial degree 3, Fine Gaussian and Coarse Radial.

### 7.3 Confusion Matrix

This matrix is also called the contingency matrix. It helps to understand the abilities of the classifier in terms of selecting as well as rejecting instances with respect to each class. If the classifier is able to match the fitness level 1 instance with fitness level 2 instances and predict healthy instances of healthy with healthy class, then the logic of implementing the SVM algorithm will be successful. The results are as follows:

| SVM Classifier Kernel | TP | TN | FP | FN |
|-----------------------|----|----|----|----|
| Linear SVM            | 2  | 45 | 5  | 1  |
| Polynomial Degree 2   | 2  | 47 | 3  | 8  |
| Polynomial Degree 3   | 2  | 46 | 4  | 1  |
| Fine SVM              | 2  | 49 | 1  | 0  |
| Gaussian SVM          | 4  | 29 | 1  | 0  |
| Medium Radial SVM     | 2  | 49 | 1  | 0  |
| Coarse Radial SVM     | 2  | 43 | 7  | 8  |

### Formulas

\[
TPR = \frac{TP}{TP + FN}
\]

\[
FPR = \frac{FP}{FP + TN}
\]

\[
ACC = \frac{TP + TN}{TP + TN + FP + FN}
\]

### Table 11: Performance Evaluation of Six SVM Kernel

Note: TP – True Positive, TN – True Negative, FP – False Positive, FN – False Negative

### Interpretation

It is clear from Table 11 that Medium Radial kernel and Fine Gaussian Kernel is producing the maximum number of true positives (>99%) and lowest false alarms. From this value, it can be safely interpreted that there are <1% chances that this classifier might go wrong, which is the lowest among all the other classifiers.

### 7.4 Area Under the Curve
The “Area under the curve” is also known as C-statistics. It gives the likelihood that the classifier in question will rank (or assign score) a randomly selected positive instance, higher than the randomly selected negative one. Higher the AUC value, higher is the probability of the positive instances and consequently higher is the chance of accuracy. [Table 12]

| S.No | SVM Classifier Kernel     | AUC  |
|------|----------------------------|------|
| 1    | Linear                     | 0.99 |
| 2    | Polynomial Degree 2        | 1.0  |
| 3    | Polynomial Degree 3        | 0.99 |
| 4    | Fine Gaussian              | 0.99 |
| 5    | Medium Radial              | 1.0  |
| 6    | Coarse Radial              | 0.99 |

Table 12: Area Under the Curve for SVM Classifier Kernel

Interpretation
It is clear from the AUC value of Polynomial Degree 2 SVM and Medium Radial SVM has the highest probability of getting positives and the lowest possibility of getting false alarm rate. The outcome of this metric is consistent with respect to the accuracy and values of the true positive rates. In this case, also, the Polynomial Degree 2 SVM and Medium Radial SVM is performing well in terms of Area Under the curve (1.0).

Conclusion
This paper work validates the latest finding in the current literature on the quality of dataset, feature sets and process of selection of machine learning algorithms i.e all aspects of the process are critical for the successful implementation and application of the machine learning algorithms for problems of classification and prediction. Since machine learning algorithm takes its inspiration from statistics as well as from the pattern finding algorithm. This paper focused on both the aspects. The evaluation strategy followed in feature selection in this paper can be operated independently of any learning algorithm as it helps to remove insignificant features before the learning process begins. The feature selection process goes through multiple revisions to arrive at the final feature set that leads to the high accuracy of the six algorithms.

The classification and the prediction process goes through a series of evaluations and reruns of the SVM with six kernels. The Medium Radial kernel of the SVM gave the maximum prediction power on the dataset. It performed well in almost all evaluations, whether it was the values of true positive rate or Area Under a Curve (AUC) is 0.1. It was empirically found that the main advantage of using SVM kernel functions is that it allows us to change Euclidean geometry so that it fits into the context of the problem and the dataset becomes workable for classification.

SVM classification algorithms are useful in cases where the regression method works well to identify the trend lines between the datasets. And it may fail in cases where there are noise and a lot of overlapping data points in the dataset. The initial statistical analysis of the feature set showed some degree of overlapping but the trend lines show a good degree of separation between the classes. Secondly, it was found that, for finding "best" separation hyper-plane, quantity (number of Instances of features) did not always convert into quality, the advantage of SVM is that it does not require a large number of instances to produce a fair percentage of accuracy as it is clear from the outcomes.

Future Scope
This paper gives computing model of Health Risk Assessment using learning models that are based on six kernel tricks. The extension of this work can be done by converting these dataset group into a multiclass risk assessment problem. The health risk classification may further be divided into three class problem (Fitness Level 1, Fitness Level 2 and Fitness Level 3). Second, for further direction, this dataset and its features may use for formulating health insurance policies. Insurance companies can leverage the use of machine learning algorithms for computing health plus insurance risks.
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