

# **Deep Incremental Statistical Closeness Factor Based Algorithm**

## (DIS-CFBA) to assess Diabetes Mellitus

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**Abstract** - Diabetes Mellitus (DM) is potential epidemic in India. Millions of individual diagnosed with this disease. Different types and number of lab tests used to diagnose it. This paper presents a novel tactic to assess DM based on CFBA and statistical methods. DM parameters assessed and categorized into significant and insignificant parameters. DM dataset processed using cluster analysis and principal component analysis (PCA). Cluster analysis creates fifteen distinct clusters. PCA applied to find out variability and to do categorization of parameters. Thus, this paper illustrates utility of statistical clustering for effective DM management.

Kev Words: Deep Incremental; Statistical; CFBA; Diabetes Mellitus etc.

#### 1. INTRODUCTION

DM is an unending, enduring situation. It affects body's capacity to use the food energy. These day's it has treated as a major nuisance in health care industry. To develop evolved systems, incremental clustering acts as a key. It can help in effective knowledge augmentation. It in turn leads to effectual knowledge management [1, 2, 5 and 13-28]. Incremental learning can also be achieved through web based interactive data mining tools [4]. Data science is a booming field. Its innovative usage can reduce tests and trouble to patients during DM diagnosis. Data science coupled with Knowledge Management System (KMS) can pattern. alter traditional DM dealings Also recommendations presented through this coupled system can act as preventive measures [3]. In this paper, a novel DIS-CFBA proposed and applied on different parameter values of patients.

Multivariate statistical approach used for different purposes like water quality assessment of river, assessment of trace elements levels in patients with type 2 diabetes, diabetes classification, identification of cigarette design influencing features and name a few [6-12]. In a similar way, multivariate statistical analysis used here to identify significant DM parameters.

Section 2 presents methods used. Section 3 throws light on results and related discussion followed by outlook in section 4. The references referred are listed at the end of this paper. Table -1: DM ATTRIBUTES RANGE [6]

Sr. No.	Name of attribute	Range of attributes in mg/dl	
1	BLOOD GLUCOSE FASTING	115-210	
2	BLOOD GLUCOSE PP	140-250	
3	CHOLESTROL	140-250	
4	TRIGLYCERIDES	140-300	
5	HDL CHOLESTROL	40-60	
6	VLDL	20-60	
7	LDL CHOLESTROL	60-115	
8	NON HDL CHOLESTROL	120-170	

#### 2. METHODS USED

#### 2.1 Cluster Analysis through DIS-CFBA

Input: Raw Dataset, Data Series (DS)

Output: Parameters categorization as significant and insignificant one

#### **Closeness Factor Based Algorithm (CFBA)**

- 1. Initial count of clusters K = 0
- Calculate CF for DS (i) 2.
- 3. Calculate CF for DS (i++)
- 4. Clusters formed on the basis of calculated CF
- 5. If (~Processed \_Flag) then CF (added new cluster) = xi insert\_counter (added new cluster) = 1 & CFBA\_Clusters ← Clusters ∪ added new cluster
- 6. for all  $xi \in I$
- 7. As Processed Flag = False
- 8. For all clusters  $\in$  clusters do
- 9. if || xi || < = CF then
- 10. As Processed \_Flag = False
- 11. Update (cluster)
- 12. insert\_counter (cluster)
- 13. Processed \_Flag = True
- 14. Exit loop



- 15. end if
- 16. end for
- 17. end if

#### DIS - CFBA

- 18. if (~Processed \_Flag) then
- 19. if (Required\_Parameter\_Range (added new cluster))
   == (Required\_Parameter\_Range (CFBA\_Clusters))
- 20. Update Required\_Parameter\_Range (CFBA\_Clusters)
- 21. Else discard (added new cluster) & Processed\_Flag = True
- 22. end if
- 23. end if
- 24. Correlation, KMO & Bartlett's test & Factor Loadings on required\_Parameter\_Range (CFBA\_Clusters)
- 25. Parameters\_Categorization as significant and insignificant one
- 26. Exit

### 2.2 Sampling

The DM data set of working adults whose age is between 35 to 45 years considered for 2016 -17 [6]. Sample analysis takes place through below mentioned principal activities.

- 1. Table 1 shows DM parameters range based on pathology reports.
- 2. After application of DIS-CFBA fifteen distinct clusters obtained.
- 3. Required parameters range analyzed on the basis of obtained clusters.
- 4. Statistical methods applied on this analysis does categorization of parameters.

#### 2.3 Statistical Data Treatment

DM data generated through clustering normalized by log normal transformation. The fitness of data for PCA verified through Kaiser-Mayer-Olkin (KMO) and Barlett's tests. The DM management subjected to two major approaches viz., cluster analysis and PCA. All statistical calculations carried out through Minitab 17.0 software. KMO and Barlett's tests performed through R programming language.

#### 2.4 Principal Component Analysis (PCA)

Two Key aspects of PCA used viz. Data reduction and summarization. It analyzes interrelationships among attributes with their common causal dimensions, branded as parameters [9]. PCA performed on normalized parameters to get significant Principal Components (PCs). These PCs further lessen the contribution of parameters with minor significance. These PCs subjected to varimax rotation generates Vari-Factors (VFs) [9]. PCs defined as parameters when their variance is greater than 1 [9]. The standard test score of any parameter must be more than variance of any single parameter [9]. Hence, PCA coupled with varimax rotation along with Kaiser Normalization applied. The extracted Eigen values from correlation matrix, significant factors and variance percent presented.

#### 3. Results and Discussion

Statistical summary of extracted DM parameters through clustering observed at developed clusters. Table 2 illustrates the same. The values of Triglycerides vary from 80 to 530 which show its importance in relation to DM. Also, the values of Blood Glucose vary from 135 to 350 which is second important attribute in relation to DM. HDL Cholesterol, VLDL, LDL Cholesterol and Non HDL Cholesterol have lower ranges which implies their DM specific importance.

Correlation coefficient of DM parameters presented in Table 3. There is strong positive correlation between Blood Glucose Fasting and Triglycerides (r=0.41). Also, significant correlation has found between Blood Glucose and Triglycerides (r=0.36). Moderate correlation is there between Blood Glucose PP and Cholesterol (r=0.33).

**Table -2:** Summary of DM Parameters of Generated Clusters

Sr. No.	Parameters	Min	Max	Mean	Std. Dev.
I	Blood Glucose Fasting (mg/dl)		260	144.77	21.41
II	Blood Glucose PP (mg/dl)	135	350	200.6	24.82
III	Cholesterol (mg/dl)	130	240	179.66	25.03
IV	Triglycerides (mg/dl)	80	530	231.3	26.67
V	HDL Cholesterol (mg/dl)	30	59	49.49	5.58
VI	VLDL (mg/dl)	15	106	38.51	10.26
VII	UDL VII Cholesterol (mg/dl)		143	88.08	14.01
VIII	Non HDL Cholesterol (mg/dl)	90	185	141.56	12.74

Before PCA, dataset normalized using log normal transformation and Kolmogorove Smirnov (K-S) statistics test. These tests verify the goodness of fit of the data to log-normal distribution [8]. The results of K-S test shows that all attributes follows the log normal distribution. To investigate suitability of data for PCA, KMO and Barlett's test of sphericity performed [7]. The significance value of 0.62 (Table 4) indicates dataset is fit for PCA. Barlett's test of



sphericity test used to check relatedness of parameters. Its 0.57 Significance level indicates correlation matrix is an identity matrix. So, parameters are related to each other. PCA applied on obtained clusters using Minitab 17.0 software. PCA yields correlation matrix for parameters I to VIII (TABLE 3) and factors extracted through centroid method of varimax rotation. Eigen value indicates the significance of the parameter. Eigen value more than one treated as significant one [10, 11, 12].

Table -3: CORRELATION MATRIX (PEARSON (n))

	Ι	II	III	IV	V	VI	VII	VIII
Ι	1							
II	0.02	1						
III	0.2	0.33	1					
IV	0.41	0.36	0.28	1				
V	0	0.11	0.09	-0.02	1			
VI	0.09	0.05	0.03	0.11	0.06	1		
VII	0.08	0.14	0.01	0.24	0.02	0.09	1	
VIII	0.11	0.14	-0.03	0.2	0.01	0.08	0.03	1

Table -4: KMO AND BARLETT'S TEST

Kaiser-Meyer-Olkin M	0.62	
Adequ	0.02	
Approx. Ch	5.69	
Bartlett's Test of	df	7
Sphericity	Sig.	0.57



Figure -1: SCREE PLOT FOR PARAMETERS

PCA result shows first four Eigen values are higher than one and considered as significant. The scree plot (Fig. 1) shows greater part of variance in the original data covered by first four parameters. These four PCs accounts for 57.6% of the total variance associated with all 08 parameters. Table 5 shows variance of first four parameters for rotated and unrotated factor loadings. PC1 has 16% of total variance. It has strong positive loading of Triglycerides, negative loading of blood glucose. PC2 handles 14% of total variance, has strong positive loading of LDL Cholesterol. This factor thus acts as a reactive component for DM. Cholesterol reveals its DM specific influence in relation to obtained loadings. PC3 has a strong loading of blood glucose fasting and accounts for 13% of variance. PC4 exhibited again 13% of total variance within the data set. It has strong positive loading of VLDL and HDL Cholesterol. VLDL and HDL Cholesterol are indicative details of PC4. Table 6 represents the factor loadings of various parameters.

#### Table -5: VARIANCE

	Principal Components				
Parameters	1	2	3	4	
<b>Blood Glucose Fasting</b>	0.013	0.029	-0.906	0.016	
Blood Glucose PP	-0.812	0.091	0.139	-0.068	
Cholesterol	0.065	-0.769	0.319	0.027	
Triglycerides	0.811	0.070	0.134	-0.074	
HDL Cholesterol	0.044	-0.099	-0.044	0.406	
VLDL	-0.123	-0.142	-0.055	0.661	
LDL Cholesterol	0.027	0.546	0.284	-0.181	
Non HDL Cholesterol	0.077	0.422	0.172	0.654	

 Table -6: FACTOR LOADINGS

PCs	Sum of squ	ared loadings	Rotated sum of squared loadings		
	Variance	% of Variance	Variance	% of variance	
1	1.35	0.16	1.34	0.16	
2	1.11	0.14	1.10	0.14	
3	1.08	0.135	1.075	0.13	
4	1.05	0.132	1.072	0.12	

Hierarchical cluster analysis used to club DM attributes with certain similarity. The dendogram (Fig. 2) categorizes eight parameters of DM in two clusters on the basis of similarity between their characteristics. Cluster 1 consists of first five attributes while cluster 2 consists of remaining three. Cluster 1 and cluster 2 corresponded to high and low significant DM parameters. The patient's data among these clusters can attribute to difference in their influence to DM.

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Figure -2: DENDOGRAM FOR DM PARAMETERS

### 4. CONCLUSION

Statistical techniques powered by clustering and PCA applied on 5K dataset for 2016-2017. Blood Glucose Fasting, Blood Glucose PP, VLDL and Non HDL cholesterol are significant parameters. This illustrates utility of statistical techniques in DM assessment. So, cut down of these parameters can improve DM management. This information can also alter pathology test pattern for DM.

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