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Genetic Epidemiological Studies of Hearing Loss in Faisalabad City, Pakistan

By Sumaira Iqbal, Fazeela Yaqoob & Tayyaba Sultana

Government College University, Pakistan

Summary- Basic aim of the study was to investigate the incidence, prevalence, disease pattern and the etiology of the non-syndromic congenital deafness in a selected population from different areas and schools for special education of Faisalabad District, Pakistan. An epidemiological study was carried out through a questionnaire including different standard parameters like surname, parental marriage age, parental relationship of marriage, their family history including information about number of offspring, sex, birth order, number of normal and diseased individuals and age of onset. Information about economic status and educational qualification of parents, their locality, food habits and paternal smoking or non-smoking habits were also recorded. In a total of 436 deaf cases, 55.0% were males and 44.95% were females while 22 families were selected for pedigree analysis. Urban population (79.58%) was affected more than rural population (20.41%). Mean maternal and paternal marriage age was 20.63 ± 0.163 and 24.989 ± 0.217 , respectively with smaller (0-4 years) marriage age difference.

Keywords: epidemiology, deafness, inheritance, pedigree.

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Genetic Epidemiological Studies of Hearing Loss in Faisalabad City, Pakistan

Sumaira Iqbal ^α, Fazeela Yaqoob ^ο & Tayyaba Sultana ^ρ

Summary- Basic aim of the study was to investigate the incidence, prevalence, disease pattern and the etiology of the non-syndromic congenital deafness in a selected population from different areas and schools for special education of Faisalabad District, Pakistan. An epidemiological study was carried out through a questionnaire including different standard parameters like surname, parental marriage age, parental relationship of marriage, their family history including information about number of offspring, sex, birth order, number of normal and diseased individuals and age of onset. Information about economic status and educational qualification of parents, their locality, food habits and paternal smoking or non-smoking habits were also recorded. In a total of 436 deaf cases, 55.0% were males and 44.95% were females while 22 families were selected for pedigree analysis. Urban population (79.58%) was affected more than rural population (20.41%). Mean maternal and paternal marriage age was 20.63 ± 0.163 and 24.989 ± 0.217 , respectively with smaller (0-4 years) marriage age difference. First (24.54%) and second (22.24%) birth order showed maximum deaf patients. Offspring of first cousin marriages (67.66%) had significantly higher congenital hearing loss affliction than of unrelated couples as demonstrated by consanguineous studies. Coefficient of inbreeding was ($F=0.051$) high than in general population ($F=0.0271$). Autosomal recessive mode of inheritance was observed in twenty-two affected families.

Keywords: epidemiology, deafness, inheritance, pedigree.

I. INTRODUCTION

Deafness, a sensory defect, affecting 1 / 500-650 newborn children and 250 million people worldwide (Olusanya et al., 2007), of whom two thirds are in developing countries. It is considered to be second major cause of disability. Frequency of hearing loss in Southeast Asia varies from 4.6% to 8.8% (Garg et al., 2009). There are approximately 63 million (India) 28 million (USA) and 9 million (UK) hearing impaired people according to WHO survey. Deafness can be categorized as genetic or non-genetic, prelingual or postlingual, and syndromic or non-syndromic.

Approximately 60% cases of deafness have genetic basis (Mehl and Thomson, 2002). Environmental factors are responsible for upto 40% cases and few are due to unknown causes (Willems, 2000; Reddy et al., 2006). In genetic cases, 85% of the hearing impairment is described as non-syndromic (Kenneson et al., 2002).

Comparatively to advanced countries, people from developing countries are three to four times more susceptible to deafness largely because of unfavourable environmental factors. Asian, African and Latin American population practice consanguineous marriages frequently, so siblings of these marriages are at high risk of autosomal recessive disorder. Consanguinity basically disturbs the normal genetic pathway and paves the way for the development of hearing loss (Reddy et al., 2006). Autosomal recessive non-syndromic hearing impairment (ARNSHI) can be traced easily in Pakistani population due to the availability of large extended and highly consanguineous pedigrees (Santos et al., 2005). The prevalence of bilateral deafness in Pakistani population is as high as 1.6 per 1000, 60% higher than the world average (Elahi et al., 1998). In Pakistani population 80% marriages are first-cousin and 60% are consanguineous (Hussain, 1998). Neonatal, infant and childhood periods face the highest death rates because of this consanguinity (Bittles, 2001).

For present epidemiological study deaf population was selected at district level to collect data for determination of prevalence rates. Effect of consanguinity on congenital deafness, social and economic status impacts and mode of inheritance of genetic cases were studied.

II. MATERIALS AND METHODS

The families for genetic epidemiological studies were collected from different schools for special education of district Faisalabad, Total twenty-two families were scored by collecting data of about 436 deaf patients from special schools and different areas of Faisalabad city. These families provided necessary relevant information for genetic epidemiological studies. Family consisted of 4 or 5 generations was considered in which at least 4 or 5 individuals were afflicted with congenital deafness, to get better results.

The details of the family were traced back through the proband. The information was collected from subjects with the help of a data sheet including their surname, parental marriage age, parental relationship of marriage, their family history including information about number of offsprings (sex, birth order), similar and other disease in family, number of normal and diseased individuals in family and the age of onset of disease.

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a) *Data analysis*

The data was analyzed in three ways. Firstly, the sample was analyzed as a whole to get information of population and second in relation of consanguinity. Third pedigrees were draw in order to evaluate the mode of inheritance of disease. Genetic relationships in marriages were classified into first cousins (1C), first cousin once removed (11/2C), second cousins (2C), distant relatives (DR), bradari (B) and unrelated (U) (Shami and Iqbal, 1983).

b) *Statistical analysis*

The statistical analysis carried out for this study included percentage (%), mean (M), standard error (S.E) and chi-square test (χ^2). Mean coefficient of inbreeding (F-value) was calculated by following the method of Wright (1992).

65.165% in rural and 52.44% in urban areas whereas females were 34.83% and 47.55%, respectively.

b) *Parental marriage age*

In case of female patients mean maternal marriage age was 20.61 ± 0.262 years whereas mean paternal marriage age was 25.010 ± 0.335 years. Similarly in case of male deaf patients mean maternal and paternal marriage age was 20.654 ± 0.207 and 24.971 ± 0.284 , respectively. It was seen that second age group (18-22 years) of maternal marriage was more commonly distributed in deaf male and females than any other age group (Table-1). Children of couples with age difference ranged from 0-4 years had high percentage (59.63 %) of disease. Difference of 2, 3, 4 and 5 years at marriage have high percentages 20.87%, 13.76%, 10.55% and 11.46% of deaf patients, respectively.

III. RESULTS

a) *General population study*

Out of 436 deaf patients, 347 belonged to urban areas and 89 belonged to rural. Male patients were

Table 1 : Distribution of mean maternal marriage age (MA) of deaf patients in different age groups

Age groups at marriage (Years)	Maternal age of male patients (X ± S.E)	Maternal age of female patients (X ± S.E)	Sexes Combined (X ± S.E)
13-17	16.20 ± 0.208 (25)	16.136 ± 0.136 (22)	16.170 ± 0.127 (47)
18-22	19.940 ± 0.123 (166)	19.609 ± 0.127 (133)	19.793 ± 0.089 (299)
23-27	24.341 ± 0.155 (41)	24.645 ± 0.189 (31)	24.470 ± 0.120 (72)
28-32	29.571 ± 0.481 (7)	29.571 ± 0.297 (7)	29.571 ± 0.272 (14)
33-37	37.0 ± 0.00 (1)	35.333 ± 0.333 (3)	35.750 ± 0.479 (4)

$\chi^2 = 223.96; d.f=9, P < 0.001$

The number mentioned within parenthesis () is the number of deaf patients.

c) *Birth order*

Different birth orders of 436 deaf patients were recorded up to 12th birth order. Percentage distribution of combined sex in different birth orders shows that 1st (24.54%), 2nd (22.24%) and 3rd (17.20%) birth order had the highest representation while 9th (0.45%), 10th (0.68%) and 12th (0.229%) birth order had the lowest representation. In male deaf patients 1st (29.16%) and 2nd (22.08%) birth order and in female deaf patients 2nd (22.44%) and 4th (21.42%) birth order showed maximum percentages. Tenth and 12th birth order (0.416%) in males and 8th birth order (0.51%) in females were least suffered (Table-2).

Table 2 : Percentage distribution of deaf males and females in different birth orders

Sex		1 st order	2 nd order	3 rd order	4 th order	5 th order	6 th order	7 th order	8 th order	9 th order	10 th order	11 th order	12 th order
Males	No	70	53	36	31	20	14	6	6	2	1	-	1
	%	29.16	22.08	15.0	12.91	8.34	5.83	2.5	2.5	0.834	0.416	-	0.416
Females	No	37	44	39	42	18	8	5	1	-	2	-	-
	%	18.87	22.44	19.89	21.42	9.18	4.08	2.55	0.51	-	1.02	-	-
Combined Sex	No	107	97	75	73	38	22	11	7	2	3	-	1
	%	24.54	22.24	17.20	16.74	8.715	5.04	2.52	1.60	0.45	0.68	-	0.229

d) Socio- economic status, life style and education

Highest percentage representation of deaf patients was seen in skilled manual occupation (43.34%), followed by professional-managerial (19.03%) and unskilled categories (14.90%), respectively. Distribution of deaf patients according to their educational level was in primary (49.77%), Middle (14.90%), Matric (13.30%) and college education (2.52%), respectively. Ratio of uneducated smokers (53.45%) and non-smokers (34.29%) were highest in representation as compared to educated non-smokers (4.33%) and smokers (0.628%), respectively.

e) Consanguinity and deafness

Deaf patients were explored on the parental consanguinity basis, showed first cousin (67.66%), first

cousin once removed (0.458%), second cousin (0.68%), distant relations (11.23%), bradari (5.27%) and unrelated spouses (14.67%), respectively. First cousin marriage couples had maximum number of affected children as compared to unrelated. Table-3 shows analysis of goodness of fit for father's occupation and different genetic relationships of deaf patients, which indicated that father's occupation and genetic relationships had highly significant effect on the development of deafness ($\chi^2 = 68.774$; d.f = 25; $p < 0.01$). Educational contributions and different genetic relationships of parents were also compared in deaf patients that indicated a highly significant effect of education and genetic relationships on deafness (Table-4) ($\chi^2 = 349.41$; d.f = 40; $p < 0.01$).

Table 3 : Test of goodness of fit between father's occupation and different genetic relationships of deaf patients

Genetic relationships	Father's occupation						Total
	I	II	IV	V	VI	VII	
1C	57 (56.16)	39 (32.48)	125 (127.88)	2 (2.71)	42 (43.98)	30 (31.80)	295 (295.0)
1 ^{1/2} C	- (0.38)	- (0.22)	- (0.87)	1 (0.02)	1 (0.30)	- (0.22)	2 (2.00)
2C	1 (0.57)	- (0.33)	2 (1.30)	- (0.03)	- (0.45)	- (0.32)	3 (3.00)
B	4 (4.38)	2 (2.53)	11 (9.97)	1 (0.21)	3 (3.43)	2 (2.48)	23 (23.00)
DR	9 (9.33)	2 (5.39)	24 (21.24)	- (0.45)	9 (7.31)	5 (5.28)	49 (49.00)
U	12 (12.18)	5 (7.05)	27 (27.74)	- (0.59)	10 (9.54)	10 (6.90)	64 (64.00)
Total	83	48	189	4	65	47	436

$\chi^2 = 68.774$; d.f = 25 $P < 0.01$. The value mentioned within parenthesis () is the expected value 1C= First cousin. 1^{1/2}= First cousin once removed. 2C= Second cousin. B = Bradari

Table 4 : Test of goodness of fit between parental educational combinations and genetic relationships

Genetic Relationships	FATHER / MOTHER'S EDUCATION								
	N/N	N/S	S/N	S/S	S/C	C/N	C/S	C/C	U/U
1C	106 (106.23)	10 (15.56)	61 (52.10)	85 (87.96)	- (1.35)	2 (2.30)	19 (16.92)	6 (6.09)	6 (6.77)
1 ^{1/2} C	- (0.72)	- (0.11)	- (0.35)	1 (0.60)	- (0.01)	- (0.01)	- (0.11)	0 (0.04)	1 (0.05)
2C	- (1.08)	- (0.16)	- (0.53)	1 (0.89)	2 (0.01)	- (0.02)	- (0.17)	- (0.06)	- (0.07)
DR	17 (17.64)	6 (2.58)	4 (8.65)	17 (14.61)	- (0.22)	- (0.34)	2 (2.81)	- (1.01)	3 (1.12)
B	7 (8.28)	5 (1.21)	3 (4.06)	8 (6.86)	- (0.11)	- (0.16)	- (1.32)	- (0.47)	- (0.53)
U	27 (23.05)	2 (3.38)	9 (11.30)	18 (19.08)	- (0.29)	1 (0.44)	4 (3.67)	3 (1.32)	- (1.47)
Total	157 (157.0)	23 (23.00)	77 (77.00)	130 (130.0)	2 (2.00)	3 (3.00)	25 (25.00)	9 (9.00)	10 (10.0)

$\chi^2 = 349.412; d.f = 40, P < 0.01$

The value mentioned within parenthesis () is the expected value

N = Nil

S = School

C = College/University

U = Unknown

f) Disease in family

In 436 deaf patients, 282 (64.67%) cases had this disease in their family. Out of which 148 (52.48%) males and 134 (47.51%) females were those who had the same disease in their family, whereas 92 male (59.74%) and 62 female (40.25%) cases did not have family history for the disease. Table-5 represents the distribution of deaf males and their relatives diagnosed for the same disease. One forty eight affected males had 284 (47.17%) affected relatives. Among those the close blood relatives such as father, mother, daughter, son, sister and brother were 181 (63.73%) followed by 13 paternal (4.57%) and 8 maternal relatives (2.8%). 82 affected relatives (28.87%) were kept in category of others. One thirty four affected females had higher affected relative number (318) than that of males (284) and these affected relatives consisted of 188 close blood relation (59.11%), 15 maternal (4.71%), 14 paternal relatives (4.4%) and 101 (31.76%) included all these three relations (Table-6).

Table 5 : Distribution of deaf males and their relatives diagnosed for the same disease

Type of Relation	Affected Relatives	No of Affected Relatives	No of cases
Only Close Blood Relation Relatives	Only Sister	19	18
	Only Brother	51	42
	Only Mother	1	1
	Sister + Brother	88	34
	F+ M+ S+ B+ D+ Son	22	7
Only Maternal Relatives	M.G.F+ M.G.M+ M.A+	8	7
	M.U+ M.C		
Only Paternal Relatives	P.G.F + P.G.M + P.A +	13	11
	P.U + P.C		
Close blood relation Relatives & Maternal-Paternal Relatives	F+ M+ S+ B+ D+ Son+	82	8
	P.G.F+ P.G.M+ P.U+ P.A+		
	P.C+ M.G.F+ M.G.M+		
	M.A+ M.C+ M.U+ M.N+W		
Total		284	148

F = Father P.G.F = Paternal-grandfather, M = Mother P.G.M = Paternal-grandmother
 S = Sister P.U = Paternal uncle B = Brother P.A = Paternal Aunt
 D = Daughter P.C = Paternal Cousin W = Wife P.N = Paternal nephew
 H = Husband M.G.F = Maternal grandfather M.G.M = Maternal grandmother
 M.A = Maternal Ant M.U = Maternal uncle M.C = Maternal cousin M.N = Maternal nephew

Table 6 : Distribution of deaf females and their relatives diagnosed for the same disease

Type of Relation	Affected Relatives	No of Affected Relatives	No of cases
Only Close Blood Relation Relatives	Only Sister	45	31
	Only Brother	23	21
	Only Son	3	1
	Sister + Brother	97	34
	F+ M+ S+ B+ D+ Son	20	5
Only Maternal Relatives	M.G.F+ M.G.M+ M.A+	15	8
	M.U+ M.C		
Only Paternal Relatives	P.G.F + P.G.M + P.A +	14	7
	P.U + P.C		
Close blood relation Relatives & Maternal-Paternal Relatives	F+ M+ S+ B+ D+ Son+	101	27
	P.G.F + P.G.M + P.U		
	+ P.A+P.N+P.C+M.G.F		
	+ M.G.M +M.A+ M.C+ M.U+ M.N+H		
Total		318	134

F = Father, P.G.F = Paternal-grandfather, M = Mother, P.G.M = Paternal-grandmother
 S = Sister, P.U = Paternal uncle, B = Brother, P.A = Paternal Aunt
 D = Daughter, P.C = Paternal Cousin, W = Wife P.N = Paternal nephew
 H = Husband M.G.F = Maternal grandfather M.G.M = Maternal grandmother
 M.A = Maternal Ant M. U = Maternal uncle M.C = Maternal cousin M. N = Maternal nephew

g) Pedigree analysis

After studying the genetic cases of disease, twenty-two patients with family disease history were selected in total scored cases. Among Mendelian modes of inheritance only autosomal recessive mode was observed.

IV. PEDIGREE

a) Epidemiology

Pedigree (Fig-1) is an illustration of a Rajput family who is settled in Ali Hassan colony of Faisalabad. Eleven members including six males and five females were affected with congenital deafness. Individual III-5

was proband through which disease was traced back in his family. In first three generations, no affected individual was found except one affected male (III-5) produced in third generation. In this generation, first consanguineous marriage produced two affected sons (IV-1 and IV-3) and two affected daughters (IV-4, IV-6). But the second first cousin marriage of an affected male and normal female (III-5 and III-4) produced one affected daughter (IV-10), two affected sons (IV-7 and IV-8) and one normal son (IV-12). In fourth generation, two marriages were practiced, first was a

consanguineous marriage between two congenitally deaf persons (IV-6 and IV-7) that produced all affected offsprings including one male (V-3) and two females (V-1 and V-2). Whereas second consanguineous marriage between an affected female and normal male (IV-10 and IV-11) produced three normal daughters (V-4, V-5 and V-6).

b) Result

This pedigree also shows an autosomal recessive mode of inheritance (Fig-1).

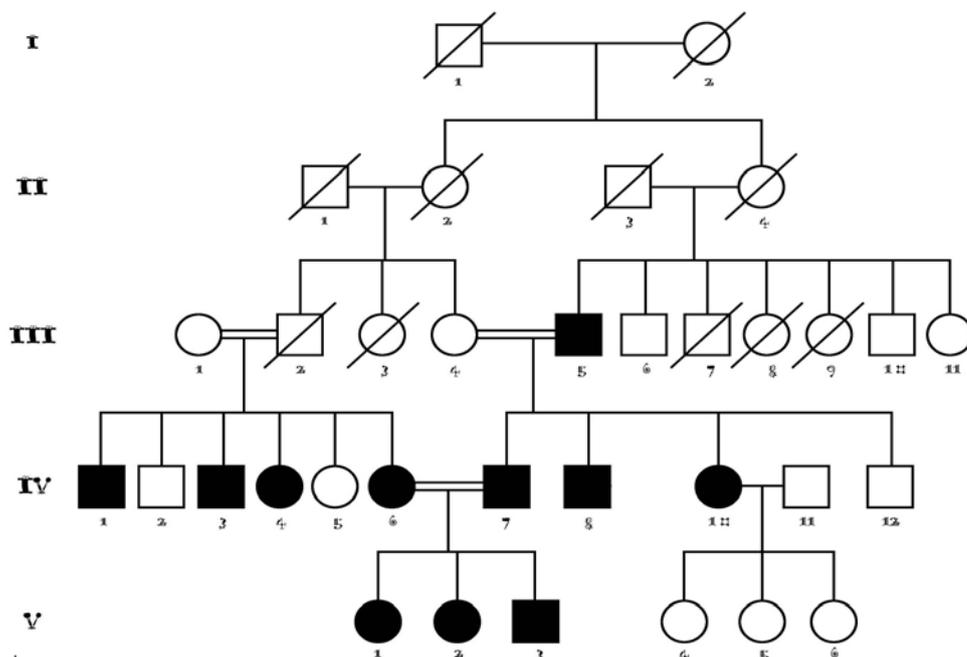


Figure 1 : pedigree reflecting an autosomal recessive mode of inheritance

V. DISCUSSION

This study showed that males were more liable to congenital deafness as compared to females. Same results were found by Liu et al. (1999). Mean maternal marriage age and mean paternal age was 20.63 ± 0.163 and 24.989 ± 0.217 recorded. Early age parental marriages could be a strong risky factor but it needs further investigation.

According to birth order study, 1st (24.54%), 2nd (22.24%) and 3rd (17.20%) birth orders were mostly affected with disease as compared to other birth orders irrespective of sex (Table-5). No supporting evidence is encountered on the importance of birth order because detailed literature review is silent on this aspect. Present study revealed that disease was common in lower social class (Category-IV), which was the case of skilled manual. This finding is comparable with the work of Bafaquee et al. (1994).

Highest percentage of deaf offspring was observed in uneducated people (49.77%) sector as compared to educated ones (2.52%). These investigations clearly show that lack of education increases the

risk of disease that was also reported by Bafaquee et al. (1994).

In Pakistan first cousin marriages are more common and rates of inbreeding ranges from 37.8% to 48.9%. Calculated coefficient of inbreeding (F) for general population ranged from 0.0236 to 0.0286 (Shami et al., 1990).

Das (2006) examined that there was an increase in genetic disorders with an increase in inbreeding in almost all populations. Consanguineous children had more familial aggregation of disease. The consanguinity rate was too high which is in agreement with Ansari et al. (2004). In Iran, the frequency of consanguinity of parents was 59.7% in congenitally malformed population and 31.5% in normal population (Khushki and Zeyghami, 2005).

It was found that most common parental relationship was 1st cousin marriages (67.66%) than unrelated marriages (14.67%). These findings are in accordance with those of Bafaquee et al. (1994), Zakzouk (2002), Bener and Hussain (2006) and Reddy et al. (2006).

Socio economic status showed significant ($P < 0.01$) influence on occurrence of congenital deafness among different marriage relationships. Education of parents also had significant effects ($P < 0.01$) on the occurrence of disease in their children. (Table-4). Similar arguments are reported by Bittles (2001) and Aber et al. (2005) for Muslim populations.

Only Mendelian mode of autosomal recessive inheritance pattern was observed in this study. Marazita et al. (1993) found that 62.8% causes of early onset deafness were genetic in which 47.1% were recessive and 15.7% were with dominant cases. Same analysis was revealed by Longtiano and Brunoni (2000). Park et al. (2003) concluded that recessive mutations are common cause of deafness due to consanguineous matings in Pakistani and Indian populations. Calapoglu et al. (2005), Zolotogora and Barges (2003) also reported the same findings in Muslim Israeli population.

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Classification Model for the Heart Disease Diagnosis

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Keywords: *heart disease, data mining techniques, classification rules, k-means clustering, and part.*

GJMR-F Classification : *NLMC Code: WG 200, WG 205*



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Classification Model for the Heart Disease Diagnosis

Atul Kumar Pandey ^α Prabhat Pandey ^σ & K.L. Jaiswal ^ρ

Abstract- Medical science industry has huge amount of data, but unfortunately most of this data is not mined to find out hidden information in data. Advanced data mining techniques can be used to discover hidden pattern in data. Models developed from these techniques will be useful for medical practitioners to take effective decision. In this research work, we have analyzed the performance of the classification rule algorithms namely PART based on K-Means Clustering algorithms. The k-means is the simplest, most commonly and good behavior clustering algorithm used in many applications. Firstly the preprocessed heart disease dataset is grouped using the K-means algorithm with the K =2 values on classes to cluster evaluation testing mode. After that data mining classification rule algorithms namely Projective Adaptive Resonance Theory are analyzed on clustered relevant dataset. In our studies 10-fold cross validation method was used to measure the unbiased estimate of the prediction model. Accuracy of K-Means Clustering, PART and PART based on K-Means Clustering are 81.08%, 79.05% and 84.12% respectively. Our analysis shows that out of these three classification models Classification based on Clustering predicts cardiovascular disease with improved accuracy.

Keywords: heart disease, data mining techniques, classification rules, k-means clustering, and part.

I. INTRODUCTION

Accurate and error-free of diagnosis and treatment given to patients has been a major issue highlighted in medical service nowadays. Quality service in health care field implies diagnosing patients correctly and administering treatments that are effective [11]. Hospitals can also minimize the cost of clinical tests by employing appropriate computer-based information and/or decision support systems. Most hospitals today use some sort of hospital information systems to manage their healthcare or patient data [10]. These systems generate huge amounts of data which take the form of numbers, text, charts and images.

Data mining is the process of extracting hidden patterns from large data sets. Data mining is a searching process done automatically for hidden patterns present in a large database [2]. Data mining is

an iterative process. Its progress is defined by discovery, through either automatic or manual methods. Data mining is reflected in its wide range of methodologies and techniques [8]. These techniques can be applied to a connection of problem sets. Classification deals in generating rules that partition the data into disjoint groups. Classification is a data mining (machine learning) technique used to predict group membership for data instances [4]. The goal of the classification is to assign a class to find previously unseen records as accurately as possible. Classification process consists of training set that are analyzed by a classification algorithms and the classifier or learner model is represented in the form of classification rules [9].

There are various kinds of classification method including decision tree induction, Bayesian networks, k-nearest neighbor classifier, case-based reasoning, genetic algorithm and fuzzy logic techniques. Systems that construct classifiers are one of the commonly used tools in data mining. Such systems take as input a collection of cases, each belonging to one of a small number of classes and described by its values for a fixed set of attributes, and output a classifier that can accurately predict the class to which a new case belongs [7].

Our goal is to use the publicly available dataset heart disease, and use PART and K-Means data mining algorithms to predict about heart disease, analyses the results and use the rules generated by these algorithms for further predictions. The rest of this paper is organized as following. Section II provides a review of literature. The problem definition is given in Section III. Subsequently, our proposed approach is discussed in Section IV. The experimental results are given in Section V. Finally, Section VI gives the conclusion and future work.

II. RELATED WORKS

A classification rule or classifier is a function that can be evaluated for any possible value specifically given the data it will yield a similar classification. In a binary classification, the elements that are not correctly classified are named false positives and false negatives [12]. Some classification rules are static functions. There are various classification rule algorithms namely OneR, Ridor, Conjunctive Rule etc. There are two types in extracting classification rules namely direct method and

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indirect method. In direct method the rules are extracted from data [5]. In indirect method the rules are extracted from other classification models. The classification rules are also known as if then rules.

In [1], the author proposed enhanced K-Means clustering algorithm for predicting coronary heart disease. There are two strategies are used for enhancing K-means clustering algorithm. First the author proposed weighted ranking algorithm to overcome the problem of random selection of initial centroids. Second the attributes associated with weights concerned by the physicians are taken into account in both ranking and the K-means algorithm instead of assigning unit weight to all the attributes. The heart dataset was collected from UCI machine learning repository. Moreover 35 conditions are carried out to assign weights to attributes. This paper describes about the rule based classification algorithm namely Part and Simple K-Means clustering algorithm. In this paper we review about the role of those two algorithms in various concepts.

III. PROBLEM DEFINITION

Given a dataset D , a set of classes C , a set of classification rules R over D through the algorithms K-Means, Part and Part based on K-Means, find the best algorithm using some the performance factors.

IV. PROPOSED SYSTEM

In the proposed system a clear view of the two algorithms is given. This section discusses a brief description of the two data mining algorithms.

a) K-Means Clustering Algorithm

Clustering the medical data into small with meaningful data can aid in the discovery of forms by supporting the abstraction of several suitable features from each of the collections thereby introducing party into the data and helping the application of orthodox data mining techniques. The k-means is the simplest, most commonly and good behavior clustering algorithm used in many applications [3, 6]. The simplicity is due to the use of squared error as the stopping criteria, which tends to work well with isolated and compact clusters. Its time complexity depends on the number of data points to be clustered and the number of iteration. The K mean algorithm works on the Euclidian Distance Method, is initialized from some random or approximate solution.

K-means groups the data in accord with their individual values into k distinct collections. Data categorized into the identical cluster have a like feature values. K , the positive number representing the number of collections, needs to be delivered in advance. The phases convoluted in a k-means algorithm are given consequently:

Prophecy of heart disease using K – Means clustering techniques

- K points denoting the data to be bunched are positioned into the space. These points signify the primary collection centroids.
- The data are consigned to the group that is nearby to the centroids.
- The points of all the K centroids are again calculated as swiftly as all the data are allotted.
- Steps 2 and 3 are repeated until the centroids stop affecting any further. This results in the isolation of data into groups from which the metric to be diminished can be reflected.

The preprocessed heart illness data is grouped using the K-means algorithm with the K values. Clustering is a type of multivariate statistical examination also known as cluster analysis, unsupervised classification analysis, or numerical taxonomy. K-Means clustering produces a definite number of separate, flat (non-hierarchical) clusters.

b) Classification Rule Based PART Algorithm

Classification is a concept or process of finding a model which finds the class of unknown objects. It basically maps the data items into one of the some predefined classes. Classification model generate a set of rules based on the features of the data in the training dataset. Further these rules can be use for classification of future unknown data items. Classification is the one of the most important data mining technique. Medical diagnosis is an important application of classification for example; diagnosis of new patients based on their symptoms by using the classification rules about diseases from known cases.

PART stands for Projective Adaptive Resonance Theory. The input for PART algorithm is the vigilance and distance parameters [13].

i. Initialization

Number m of nodes in F1 layer:=number of dimensions in the input vector. Number m of nodes in F layer: =expected maximum number of clusters that can be formed at each clustering level.

Initialize parameters L , ρ_0 , ρ_h , σ , α , θ , and e .

1. Set $\rho = \rho_0$.
2. Repeat steps 3 – 7 until the stopping condition is satisfied.
3. Set all F2 nodes as being non-committed.
4. For each input vector in dataset S , do steps 4.1-4.6.
 - Compute h_{ij} for all F1 nodes v_i and committed F2 nodes v_j . If all F2 nodes are non committed, go to step 4.3.
 - a. Compute T_j for all committed F2 nodes V_j .
 - b. Select the winning F2 node V_j . If no F2 node can be selected, put the input data into outlier 0 & then continue to do step 4.
 - c. If the winner is a committed node, compute r_j , otherwise go to step 4.6.
 - d. If $r_j \geq \rho$, go to step 4.6, otherwise reset the winner V_j and go back to step 4.3.

- e. Set the winner VJ as the committed and update the bottom-up and top-down weights for winner node VJ.
- 5. Repeat step 4 N times until stable clusters are formed (i.e. until the difference of output clusters at N-th and (N-1)-th time becomes sufficiently small).
- 6. For each cluster C_j in F2 layer, compute the associated dimension set D_j . Then, set $S = C_j$ and set $\rho = \rho + \rho_h$ (or $\rho = |D| = \rho_h$), go back to step 2.
- 7. For the outlier O, set $S = 0$, go back to step 2.

Fig. 1 : Algorithm for PART

V. EXPERIMENTAL RESULTS

The above two algorithms are combined using dataset namely Heart Disease. These dataset are collected from UCI Repository in the website www.ucirepository.com. The heart disease dataset contains 303 instances and 14 selected attributes. Initially dataset contained some fields, in which some value in the records was missing. These were identified and replaced with most appropriate values using ReplaceMissingValues filter from Weka 3.7. This process is known as Data Preprocessing. After pre-processing the data, data mining clustering and classification techniques namely Simple K-Means Clustering and PART were applied.

To measure the stability of the performance of the proposed model the data is divided into training and testing data with 10-fold cross validation. A confusion matrix shows how many instances have been assigned to each class. In our experiment we have two classes or clusters, and therefore we have a 2x2 confusion matrix. The entries of this matrix are used to explain the performance measures. The following charts and figure are based on the combined made of two algorithms namely K-Means and PART for heart disease dataset.

We are evaluating the performance of Simple K-Means algorithm Clustering using the mode of classes to clusters evaluation with the prediction attribute nom. Table 1, Table 2, Table 3 and Table 4 illustrates the confusion matrix of Simple k-means, PART, PART via Simple K-means (Classification via Clustering) and Accuracy of algorithm respectively. Results shows that 169 (56%) records are grouped into cluster 0 and 134 (44%) to cluster 1. Cluster 1 those who have heart disease and cluster 0 has no heart disease.

Table 1 : Confusion Matrix of K-Means

Predicted Class	Actual Class	
	1	0
Cluster 1 <-- 0	27	138
Cluster 0 <-- 1	107	31

Table 2 : Confusion Matrix of PART

Predicted Class	Actual Class	
	1	0
a=1	131	28
b=0	34	110

Table 3 : PART via Simple K-Means clustering

Predicted Class	Actual Class	
	b	a
b=cluster 1	125	9
a=cluster 0	12	157

Table 4 : Comparison of Data Mining Techniques

Classification Techniques	Time(seconds)	Accuracy %
Simple K-Means	0.02	80.858
PART	0.06	79.538
PART via K-Means	0.02	93.0693

Table 5 illustrates the number of rules created by PART algorithm without K-Means, PART based on K-Means. Figure 2 & 3 illustrates the rules generated by Part and Part with cluster relevant data where class value 0 & cluster value 1 has heart disease.

Table 5 : No. of Rules generated by Algorithm

Classification Techniques	No. of Rules
PART	26
PART via Simple K-Means Clustering	11

- PART DECISION LIST**
- 1) ca > 0.674497 AND cp = asympt AND sex = male: 0 (62.0/2.0)
 - 2) thal = normal AND ca <= 1 AND slope = up AND fbs = f AND exang = no AND age <= 56: 1 (54.0/1.0)
 - 3) oldpeak > 2.4 AND thal = reversable_defect: 0 (16.0/1.0)
 - 4) thal = normal AND ca <= 1 AND slope = up AND fbs = t: 1 (10.0)
 - 5) thal = normal AND age <= 45: 1 (16.0)
 - 6) exang = no AND sex = female AND fbs = f AND thal = normal: 1 (32.0/3.0)
 - 7) slope = flat AND sex = female: 0 (17.0/2.0)
 - 8) sex = female: 1 (6.0/1.0)
 - 9) slope = up AND fbs = t: 1 (5.0)
 - 10) ca <= 1 AND thal = normal AND exang = yes: 1 (7.0/1.0)
 - 11) exang = yes AND chol > 243: 0 (10.0)
 - 12) fbs = t AND ca <= 0: 1 (5.0)
 - 13) oldpeak > 0.7 AND slope = flat: 0 (19.0/3.0)
 - 14) exang = yes: 1 (6.0/1.0)
 - 15) ca <= 1 AND thal = fixed_defect: 1 (4.0)
 - 16) ca <= 1 AND thal = normal AND slope = up AND chol <= 271: 1 (4.0/1.0)
 - 17) ca <= 1 AND thal = normal AND slope = flat: 1 (3.0)
 - 18) restecg = left_vent_hyper AND cp = typ_angina: 0 (3.0/1.0)
 - 19) cp = atyp_angina AND slope = up AND trestbps > 122: 0 (3.0/1.0)
 - 20) cp = atyp_angina: 1 (5.0/1.0)
 - 21) ca <= 1 AND slope = flat AND ca <= 0: 1 (3.0/1.0)
 - 22) slope = up AND thal = normal: 0 (3.0)
 - 23) slope = up AND cp = asympt AND restecg = normal: 1 (3.0/1.0)
 - 24) cp = asympt: 0 (2.0)
 - 25) slope = up: 1 (2.0)
 - 26) : 0 (3.0/1.0)

Fig. 2 : Generated Rules by PART

- PART DECISION LIST ON CLUSTERED RELEVANT DATA**
- 1) exang = yes AND num = 0: cluster1 (76.0)
 - 2) thal = normal AND exang = no AND slope = up: cluster0 (92.0)
 - 3) restecg = normal AND cp = non_anginal: cluster0 (23.0)
 - 4) restecg = normal AND cp = atyp_angina: cluster0 (11.0)
 - 5) restecg = normal AND sex = female: cluster0 (7.0)
 - 6) cp = atyp_angina AND age <= 60: cluster0 (9.0)
 - 7) restecg = left_vent_hyper AND cp = asympt: cluster1 (29.0)
 - 8) age <= 53: cluster0 (17.0/3.0)
 - 9) exang = no AND slope = flat AND thal = reversable_defect: cluster1 (15.0)
 - 10) exang = no AND fbs = f AND oldpeak <= 3.6: cluster0 (13.0)
 - 11) : cluster1 (11.0)

Fig. 3 : Rule Generation by PART via K-Means

Figure 4 & 5 illustrates the threshold curve of PART algorithm for class 1 & 0. We can say that the area under ROC= 0.831.

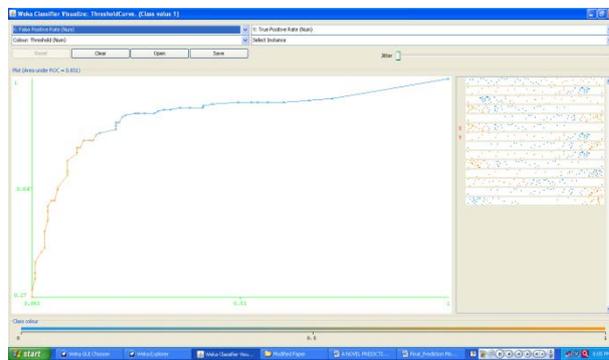


Fig. 4 : Threshold Curve of PART for Class 1

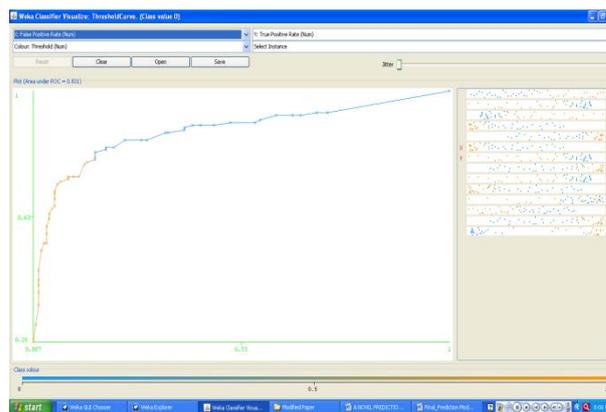


Fig. 5 : Threshold Curve of PART for Class 0

Figure 6 & 7 illustrates the threshold curve of PART algorithm for class value cluster1 & cluster0. We can say that the area under ROC= 0.959.

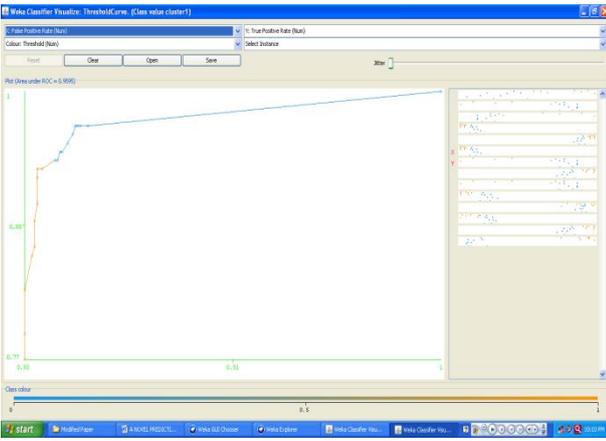


Fig. 6 : Threshold Curve of PART for Class value cluster 0

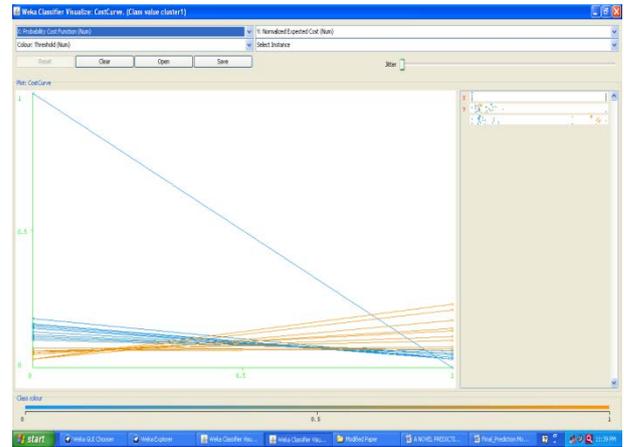


Figure 9 : Cost Curve for Class value Cluster1

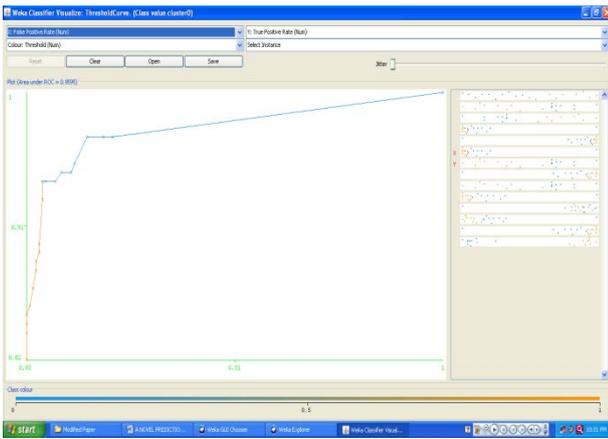


Fig. 7 : Threshold Curve of PART for Class value cluster 0

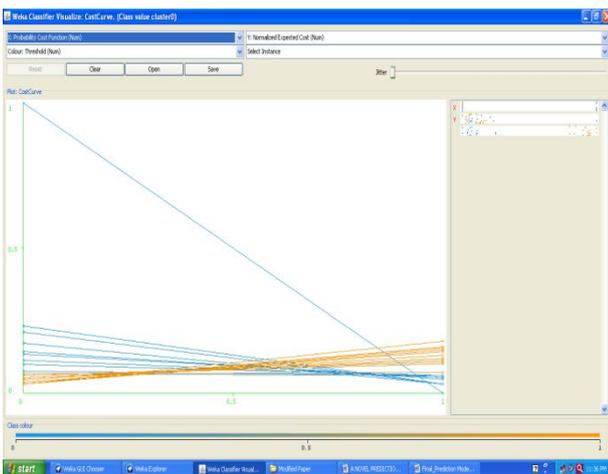


Figure 8 & 9 : Cost Curve for Class value Cluster0

VI. CONCLUSION AND FUTURE WORK

Around 18 million people 7% of the Indians are affected by heart disease. Heart disease is mostly affected the person under the age of 65. In this paper, we have compared PART and PART based on K-Means Clustering algorithms which are very suitable for generating rules in classification technique. The classification rule generation algorithms generates classification rules which is both sensitive and non sensitive. There are different data mining techniques that can be used for the identification and prevention of cardiovascular disease among patients. Our studies showed that Part based on K-Means Clustering turned out to be best classifier for cardiovascular disease prediction.

In our future work, we have planned to design and develop an efficient heart attack prediction system with Patient Prescription Support using the web mining and data warehouse techniques. New algorithms and techniques are to be developed which overcome the drawbacks of the existing system. In future some privacy preserving technique can be induced for the rule generation in the classification technique. We intend to improve performance of these basic classification techniques by creating Meta model which will be used to predict cardiovascular disease in patients.

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Risk Factors Associated with Transmission of Hepatitis B and Hepatitis C Virus in Pakistan

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Abstract- In Pakistan about 7 to 9 million people are living with HBV and 10 million people are living with HCV, with higher morbidity and mortality. This article reviews prevalence of Hepatitis B and Hepatitis C virus in Pakistan and risk factors associated with transmission of viral hepatitis. A literature search was conducted on research articles from Pubmed, PakMediNet and Google scholar. Prevalence of HBV and HCV infection varies in different regions due to inadequate knowledge and non implementation of international health standards. Majority of Pakistani population lives in rural areas as compared to urban areas. But unfortunately, up to date, no prevalence study has been reported from rural areas of Pakistan depicting HBV and HCV infection. Prevalence of HBV in general population, peds, pregnant women and IDUs was reported 4.5%, 1.8%, 0.34% to 12.62% and 22.4% respectively.

Keywords: *hepatitis B virus, hepatitis C virus, risk factors, awareness, needle stick injuries, barbers, blood transfusions.*

GJMR-F Classification : *NLMC Code: WI 715, QW 170*



RISKFACTORSASSOCIATEDWITHTRANSMISSIONOFHEPATITISBANDHEPATITISCVIRUSINPAKISTAN

Strictly as per the compliance and regulations of:



Risk Factors Associated with Transmission of Hepatitis B and Hepatitis C Virus in Pakistan

Umar Saeed ^α & Sobia Manzoor ^σ

Abstract- In Pakistan about 7 to 9 million people are living with HBV and 10 million people are living with HCV, with higher morbidity and mortality. This article reviews prevalence of Hepatitis B and Hepatitis C virus in Pakistan and risk factors associated with transmission of viral hepatitis. A literature search was conducted on research articles from Pubmed, PakMediNet and Google scholar. Prevalence of HBV and HCV infection varies in different regions due to inadequate knowledge and non implementation of international health standards. Majority of Pakistani population lives in rural areas as compared to urban areas. But unfortunately, up to date, no prevalence study has been reported from rural areas of Pakistan depicting HBV and HCV infection. Prevalence of HBV in general population, peds, pregnant women and IDUs was reported 4.5%, 1.8%, 0.34% to 12.62% and 22.4% respectively. Prevalence of HCV in general adult population, pediatric population, young population applying for recruitment, injecting drug users and multitransfused population was reported 4.95%, 1.72%, 3.64%, 57% and 48.67% respectively. Due to rapidly increasing threat of viral prevalence in Pakistani societies, it is hypothesized that Pakistan neither need enemies nor nuclear war; instead its rapidly increasing hepatitis epidemics can potentially wipeout the entire nation.

Keywords: hepatitis B virus, hepatitis C virus, risk factors, awareness, needle stick injuries, barbers, blood transfusions.

I. INTRODUCTION

The HBV and HCV are blood borne pathogens, frequently causing deaths among general populations and various high risk populations. HBV and HCV were discovered in 1963 and 1975 respectively. HBV infected 2 billion people worldwide and about 400 million of them were chronically infected (Ali et al., 2011; Li, et al., 2010; Zhu, et al., 2008; Alam, et al., 2007). The HBV and HCV infection constitutes approximately 530 million of 6 billion world population (Hwang, et al., 2006; WHO, 2000; National Foundation for Infectious Diseases, 2012). WHO has declared South East Asia a high risk area for HCV with prevalence rate of 2.15% (WHO, 2000). Liver cirrhosis develops in 20% of HCV infected patients after 10-20 years, while liver cancer usually appears after 20-40 years post infection (Denson, 2005). In Pakistan, 7 to 9 million people are living with HBV with an approximate carrier rate of 3 to 5% (Ali, et al., 2011). While about 10 million people are living with HCV in Pakistan. Asymptomatic spread of

chronic infection is major dilemma, which significantly contributes toward dissemination of lethal viral infection to others in society (Farooqi, et al., 2007). Horizontal transmission in early childhood is major contributing factor for chronic viral hepatitis infection in Pakistan (Mujeeb, et al., 1997). The perinatal transmission is leading cause of high rate of chronic infection in developing countries of Asia and Africa, whereas parenteral transmission is frequent cause of cases reported from developed industrialized countries (Maddrey, et al., 2000). Among injection drug, paid blood donors and multitransfused thalassemia populations, there is an increased transmission risk of transfusion transmitted HBV and HCV infections. In developing countries, increased transmission of HBV and HCV is due to non implementation of international standards on blood transfusions, perinatal transmission, intranasal cocaine use, shared shaving equipments, intravenous drug users, tattoos, sexual activity, needle stick injuries, body piercing, hemodialysis, hemophilia and transplants.

II. LITERATURE SEARCH

A literature search was performed via accessing research articles of previous two decade from PakMediNet, Pubmed and Google Scholar with key words of Hepatitis B virus and Hepatitis C virus associated risk factors, barbers, needle stick injuries, blood transfusion. The valued information was subjected for review.

III. HEPATITIS B VIRUS IN PAKISTAN

There exists a limited knowledge in general population about epidemiological patterns of hepatitis B prevalence in different communities of Pakistan (Waheed, et al., 2010). Noorali et al. reported prevalence of HBV in general population of Karachi to be of 4.5% (Noorali, et al., 2008). According to Chaudhary et al. the prevalence of HBV in individuals from Rawalpindi was approximately 2.3% (Chaudhary, et al., 2007). It has been reported that the prevalence of HBV in peds was 1.8% (Jafri, et al., 2006). In a study conducted on recruitment individuals from different areas of Pakistan, the prevalence of HBV was reported to be of 3.2% (Mirza, et al., 2006). Several seroprevalence studies have been conducted on blood transfusion populations. Prevalence of HBV in pregnant women is significantly

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different in different regions of Pakistan, depending upon availability or absence of hygienic environment. According to various studies conducted on pregnant women, it has been reported that HBV prevalence range between 0.34% (minimum reported rate) to 12.62% (maximum reported rate) (Sheikh, et al., 2009; Yousfani, et al., 2006). Sometimes health care workers are accidentally exposed to blood borne pathogens due to various reasons such as needle prick injuries and contact to patients body fluids. It has been reported that the prevalence of HBV among health care workers from Abbottabad was 2.4% (Sarwar, et al., 2008). Similar kind of study was reported from Karachi, with HBV prevalence of 2.4% (Aziz, et al., 2006). Injection drug users and multitransfused population can be considered as populations at risk for bloodborne pathogenic infection.

Unscreened blood supply could be a major threat for transfusion transmitted infections. According to study conducted from Karachi on IDUs, it has been reported that HBV prevalence was 22.4% (Alam, et al., 2007). Although there exist an increased threat of viral infection among multitransfused population, yet due to limited resources and awareness; only few studies have been reported from Pakistan. Majority of Pakistani population lives in rural areas as compared to urban areas. But unfortunately, up to date, no study has been reported from rural areas of Pakistan. According to HBV screening studies conducted in Islamabad, Lahore and Peshawar the prevalence of HBV was reported as 12.4%, 4% and 5% respectively (Hussain, et al., 2003; Khokhar, et al., 2004; Malik and Hussain, 2006). In order to prevent future epidemic of HBV in Pakistan, many efforts are required from both government and public sector authorities. In Pakistan, various vaccination campaigns are being conducted with major emphasis on prevention of HBV infection in neonates. It has been reported by UNICEF-WHO that approximately 73% of Pakistani neonates were vaccinated against hepatitis B virus (WHO/UNICEF, 2008). Frequent vaccination programs should be conducted at national level for people from all age groups in order to decrease the future burden of disease.

IV. HEPATITIS C VIRUS IN PAKISTAN

In 2008 Hakim et al. conducted a study on prevalence of HCV in Karachi with population size of 3820. According to the study the prevalence of HCV was 5.20 among general population (Hakim et al., 2008). It has been reported by Aziz et al. that prevalence of HCV is 1.40% in pediatric population of Karachi, which depicts increased threat of early chronic infection in such individuals (Aziz et al., 2007). In recruitment individuals from Sargodha, HCV prevalence was reported to be 4.41% (Alam et al., 2006). In Hyderabad, Yousfani et al. reported prevalence of HCV to be 16.50%; which describes HCV to be an increasingly

health problem in pregnant women (Yousfani et al., 2006). Sultan et al. reported prevalence of HCV to be 4.99% in 41 498 blood donors from different areas of Pakistan. IDUs are considered as high risk population for HCV transmission. Several studies have been conducted on seroprevalence of HCV among IDUs from different areas of Pakistan, with extremely increased percentage prevalence of viral hepatitis. It has been reported by Kuo et al. that HCV prevalence in IDUs of Lahore was 88% which describes an alarming situation. Similarly another study conducted in Karachi, it was reported that 94% of IDUs were HCV positive (Kuo et al., 2006; Altaf et al., 2009). Among health care workers HCV prevalence was reported to be 5.60% (Aziz et al., 2002). Limited studies have been conducted on multitransfused populations like thalassemia, dialysis patients and hemophilia patients. According to literature review the prevalence of HCV in thalassemia and hemophilia patients is approximately 57% and 56% respectively (Shah et al., 2005; Malik et al., 2006). Multitransfused populations are at increased risk of developing viral hepatitis due to limited screening facilities, prior to blood transfusion, at various blood transfusion setups of rural areas and towns. In majority of urban areas absence of healthy blood transfusion facilities and scarcity of quality assurance procedure drag the poor patients towards more serious consequences in term of lethal viral infections. Hepatitis treatment is very expensive and it creates huge burden on Pakistan economy. Government should raise awareness among the nation by frequent use of electronic media and by modifying syllabus of schools and colleges.

V. RISK FACTORS ASSOCIATED WITH TRANSMISSION OF HBV AND HCV

a) *Reuse of needles and needle stick injuries*

There exist an increased evidence of hepatitis B and hepatitis C virus transmission due to frequent reuse of needles and syringes. It has been reported that there are many group of individuals who are actively involved in repacking and recycling of used needles and syringes. These products are later on supplied at many drug stores. It was further reported that because of refine packaging it becomes very difficult for general public to distinguish between new disposable sterilized needles and repacked unsterilized syringes (Khan, et al., 2000; Simonsen, et al., 1999; Abdul Mujeeb, et al., 2003). It has been reported that during 2002 to 2007, 1382 needle stick injuries occurred at Aga Khan University hospital Pakistan with higher incidence associated with young doctors (28.5%), nurses (20.4%). It has been further reported that approximately 19% of overall injuries occurred during blood collection (Waheed, et al., 2010; WHO unsafe inj, 2000). In order to avoid further burden of disease, there is an urgent demand for alteration in behavior of both patients and

doctors. According to Janjua et al approximately 68% of individuals received injection in previous 3 months and out of those injections only 54% were provided from new syringes (Janjua, et al., 2005). It has been reported that estimated number of injections varies from 8.2 to 13.6 per person per year, which are highest as compared to other developing countries. Among these almost 94.2% were further categorized as unnecessary (Altaf, et al., 2007). It has been reported by Khan et al that 44% of Pakistani population would prefer injections (as compared to oral medicine) as therapeutic options against various ailments (Khan, et al., 2000). Due to poor sanitary conditions in various health sectors, it has been reported that almost 60% of used syringes are not properly destroyed, instead are majorly dumped into general public waste. Persons who are involved in garbage collection are more prone to acquire viral infections due to needle stick injuries (Abdul Mujeeb, et al., 2003; Waheed, et al., 2009).

b) *Barbers*

In Pakistan, most of HBV and HCV infected individuals have history of shaving from barbers. In 2010, a survey was conducted on 508 barber shops from capital twin cities of Pakistan. It was reported that 99.8% of barbers were washing razor with water. It was further reported that only 39.6% knew about HBV and HCV as bloodborne viral pathogens. There exist moderate awareness about different transmission routes of hepatitis and majority of them were unaware of vaccination (Waheed, et al., 2010). Most of barbers are involved in disposing of used blades in general public waste, hence posing higher risk for scavengers and sweepers during garbage handling. Injuries caused by infected blades, could bring havoc in lives of healthy individuals. It has been reported that approximately 80% of barbers applied potash alum stone (also known as Phatkari) on facial cuts in order to prevent bleeding. Using single stone on multiple customers could be a threat for transmission of viral hepatitis B virus, hepatitis C virus and human immunodeficiency virus. Some barbers have high burden of work especially at weekends and holidays. On special occasions (such as Eid days and festivals), the potash alum stone is consumed in just couple of days. It is the property of hepatitis B virus that it can easily survive for more than seven days on solid surfaces and instruments (Downey, 2008). In Muslim community, circumcision is universal. This procedure is mostly performed by barbers in rural and urban areas. Due to lack of awareness and knowledge about transmission of viral hepatitis, most of barbers, during circumcision, use contaminated instruments on multiple clients. This unhealthy practice makes infants prone to bloodborne viral pathogens.

c) *Blood Transfusions*

In Pakistan, most of the individuals have requirement of healthy blood donation for life saving

purpose. Under such circumstances, unscreened pathogen carrying blood could be a potential source of transfusion transmitted infections and future complications. It has been stated by WHO that annual blood transfusions carried out in Pakistan are approximately 1.2-1.5 million (WHO country office in Pakistan, Blood safety, 2009). It has been reported that during blood transfusions carried out in Karachi, only 25% donations were obtained from volunteer donors and 23% of those donations were screened for hepatitis C virus (Luby, et al., 2000). Almost sixty six percent of Pakistani population belongs to rural areas. In these sectors, the blood transfusion facilities are not satisfactory due to absence of organized infrastructure and electricity problem. Replacement donors including friends, family members and relatives are among the major source for blood donations. In order to strengthen the good relations, most of donors hide their health status and thus become willing to support their lives. Identification of most appropriate donors and ensuring proper blood screening prior to donations are the key factors associated with safe blood donation. For safe blood donation, the donors must possess repeated negative results for bloodborne pathogens. Blood requirements are on its peak during pregnancy related issues, traumatic injuries, thalassemia, dialysis, hemophilia. In order to cope with such emergency situations; organized, fair and safe blood transfusion setups must be developed at national level with easy accessibility. There is an urgent demand for implementation of international blood safety standards at various blood transfusion setups to ensure limited risk for transmission of hepatitis B and hepatitis C virus.

d) *Conclusion*

The prevalence of Hepatitis B virus Hepatitis C viruses is rapidly increasing in different regions of Pakistan due to non implementation of international health standards and limited awareness of HBV and HCV prevention among general and high risk populations. Although public health authorities are raising awareness among general and high risk population by the assistance of electronic and print media, yet unfortunately these efforts are not practiced at majority of Pakistani rural areas. In order to decrease future burden of disease preventive measures should be adopted, these includes; proper sterilization of health care instruments, use of disposable syringes and new razor blades and screening of blood against transfusion transmitted infectious agents.

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How to Evaluate the Risk of Malnutrition in Patients with Copd?

By Chandra Selvi. E, Saikumar. P, & Naveen Kumar

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Abstract- Background: The risk of malnutrition in patients with COPD increases the length of hospital stay, early readmission rates and poorer prognosis. Malnutrition is a significant problem in COPD and often goes undetected and often untreated. Many patients with COPD suffer with poor dietary intake and consequently reduced muscle mass.

Aim & Objective: To evaluate the effect of duration of disease and malnutrition in patients with COPD.

Material And Methods: Ten patients with COPD (both male and female) aged 30-50 yrs were recruited in this study. After getting informed consent the subjects were instructed to fill the questionnaire for the history related to our study. The subjects Body Mass Index (BMI Kg/m²), Body surface area, Duration of Disease, Skinfold thickness (Caliper), Calories intake, and Pulmonary function tests (RMS Polyrite) were evaluated.

Keywords: malnutrition, COPD, body mass index, skinfold thickness, calories intake, body surface area.

GJMR-F Classification : NLMC Code: WG 420



Strictly as per the compliance and regulations of:



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Aim & Objective: To evaluate the effect of duration of disease and malnutrition in patients with COPD.

Material And Methods: Ten patients with COPD both male and female) aged 30-50 yrs were recruited in this study. After getting informed consent the subjects were instructed to fill the questionnaire for the history related to our study. The subjects Body Mass Index (BMI Kg/m²), Body surface area, Duration of Disease, Skinfold thickness (Caliper), Calories intake, and Pulmonary function tests (RMS Polyrite) were evaluated.

Results: A significant individual patients variation was observed in our study. The mean BMI of the patients was (23.45±6.34) found to be reduced than normal. Duration of the disease (2yrs-28yrs) was negatively correlated with BMI, Free Fat Mass, Calorie intake and Expiratory Flow Rates (55%-65% pred). By using skinfold thickness (Biceps, Triceps, Sub scapular, Waist, Knee, Calf) Free Fat Mass was calculated.

Conclusion: Being an inflammatory disease COPD involves with lungs and affect other body tissues like bones and muscles, these are known as co morbidities. Diet and nutritional intake are important in COPD because they help to combat some of these co morbidities. So nutrition is an important therapy in the management of patients with COPD.

Keywords: malnutrition, COPD body mass index, skinfold thickness, calories intake, body surface area.

I. INTRODUCTION

From 1950s useful screening tools such as body weight and body mass index have been used to evaluate the nutritional status. Patients who are underweight or losing weight voluntarily associated with severity of airflow obstruction are the poor prognostic sign in chronic obstructive pulmonary disease (COPD). The causes of weight lose in patients with COPD are multifactorial including decreased oral intake as malnutrition, the effect of increased work of breathing due to abnormal respiratory mechanics and the effect of chronic systemic inflammation. Malnutrition can be defined as weight less than 90% of the predicted value as given by the Metropolitan Insurance Company or body mass index (BMI) of less than 18.4 kg/m².

The quality of life and survival limitation of chronic obstructive pulmonary disease (COPD) could be due to exercise intolerance and alterations in skeletal muscle like muscle wasting, muscle weakness and muscle fatigue rather than pulmonary problems. (Rob) Patients with COPD are commonly characterized by thin, breathlessness and voluntary weight loss. Long term use of medications such as bronchodilators (malabsorption), Corticosteroids (peripheral myopathy), and antibiotics (Gastrointestinal disturbances) can indirectly affect the nutritional status of COPD patients. (Macklem 2001). Studies have been proved that reduced maximal expiratory flow(Faulkner et al 2006), FEV1 in COPD correlates poorly with exercise capacity (Lencer et al 2003). Hence this study was designed to evaluate the baseline parameters to assess the nutritional status in patients with COPD.

II. MATERIALS AND METHODS

a) Study population

Ten COPD male patients aged 30-50 yrs with clinically stable were recruited from chest & TB department of Sree Balaji Medical College and hospitals. The study design was explained to the subjects and their informed consent was obtained. The COPD subjects were diagnosed according to the criteria given by Global Initiative for Chronic Obstructive lung Disease (COLD) Patients history like duration of disease, diet intake were obtained by questionnaire. Study was approved by the institutional medical Ethics committee of Sree Balaji Medical College, Chennai.

b) Parameters measured

Body weight (Kg) and height (cm) were measured with subjects wearing indoor clothing and BMI was calculated as by weight and height². Pulmonary function test: Flow rates and lung volumes were determined using computerized spirometer (Medispiror). Forced inspiratory and expiratory maneuvers were performed three times and the best values obtained from the maximum inspiratory and expiratory flow-volume curves were used for comparison. Body surface area was calculated in m². Skin fold thickness was taken in six sites of the body like biceps, triceps, subscapular, waist, knee and calf muscles by using digital skinfold thickness calipers. With the Skinfold Thickness and Body Surface Area, Total Body Fat was calculated.

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c) *Statistical analysis*

Statistical analysis will be performed by using statistical package for social sciences (SPSS). Data will be expressed as mean ± standard deviation. The correlation between the parameters will be analyzed by using Pearson's moment product correlation analysis. Any p value <0.05 will be considered significant

BMI from 16.90 to 30.37. Only thirty percent of the study population had less than normal BMI. In Table II duration of disease was compared with pulmonary function tests of study population. Values of FEV% and FEF 25-75% proved the obstructive pattern of lung disease and severity of diseases. Moreover as the duration of disease progresses reduction in pulmonary function tests parameters were observed.

III. RESULTS

In our results Table-1 showed duration of disease of the study population from 2-40 years and the

a) *Results*

Characteristic features of study population

Table 1

Subject	Age (yrs)	Sex	Disease Duration	Weight (Kg)	Height(M)	BMI	BSA
1	45	F	20	46	1.65	19.90	1.70
2	40	F	2	57	1.39	29.50	1.57
3	52	M	8	77	1.67	27.61	2.14
4	38	F	2	58	1.5	25.78	1.71
5	47	F	15	47	1.37	25.04	1.42
6	37	F	25	45	1.43	22.01	1.46
7	57	M	2	65	1.6	25.39	1.91
8	65	M	40	46	1.5	20.44	1.55
9	54	M	16	65	1.6	25.39	1.91
10	50	F	5	57	1.37	30.37	1.55

BMI- Body Mass Index, BSA-Body Surface Area

Table 2 : Comparison of duration of disease with pulmonary function tests

Subject	Disease Duration	PFT				
	Years	FEV ₁	FVC	FEV ₁ %	FEF ₅₀	FEF ₂₅₋₇₅
1	20	44	58	76	55	51
2	2	38	47	82	75	77
3	8	41	50	69	55	67
4	2	35	45	70	77	76
5	15	35	49	78	69	57
6	25	35	42	75	51	50
7	2	46	55	84	74	79
8	40	37	47	66	46	47
9	16	47	57	74	58	54
10	5	40	50	79	73	71

PFT- pulmonary function tests, FVC- Forced vital capacity, FEV1- Forced Expiratory volume

Table 3 : Comparison of duration of disease with Nutritional status

Subject	Disease Duration	Cal Intake	Calorie Deficiency	Sum(SFT)	TBF
1	20	1450	850	91.2	3.08
2	2	1165	1135	102.09	8.43
3	8	1400	900	190.1	11.65
4	2	1850	450	102.6	3.77
5	15	1450	850	99.2	2.98
6	25	735	1565	95	2.83

7	2	2045	255	164.33	8.58
8	40	1300	1000	72.3	1.71
9	16	1400	900	189	10.32
10	5	1500	800	159.56	6.66

SFT- Skin Fold Thickness, TBF- Total Body Fat

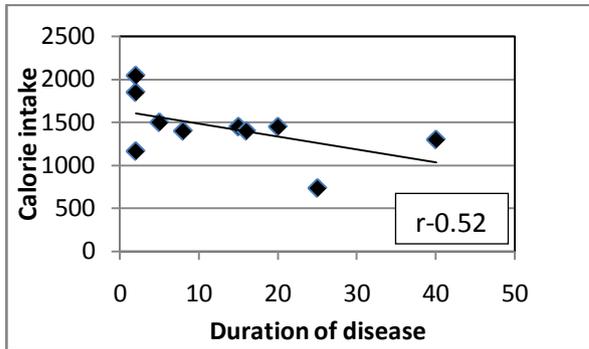


Figure 1

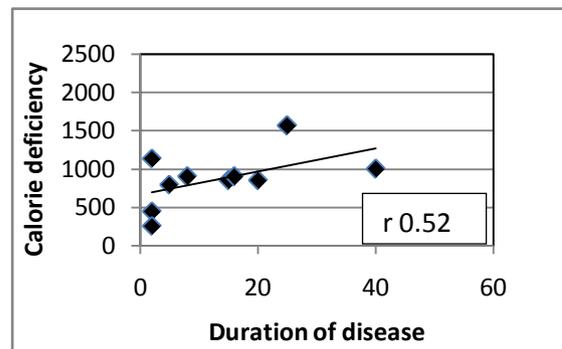


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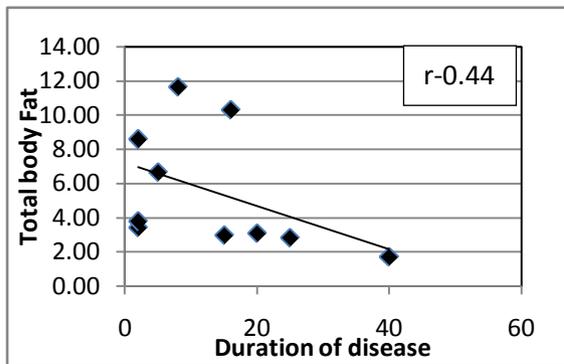


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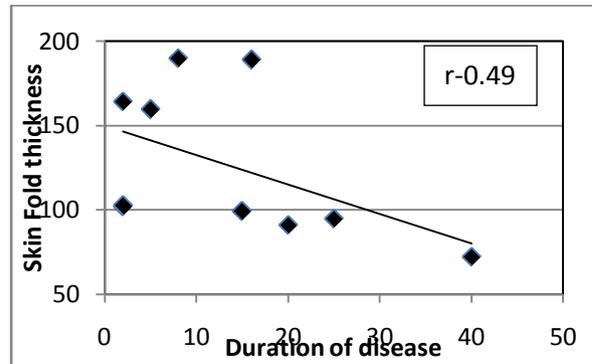


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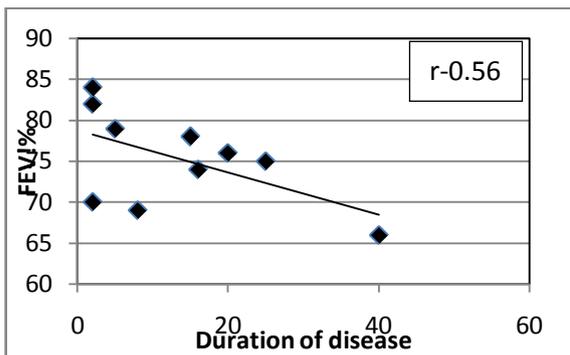


Figure 5

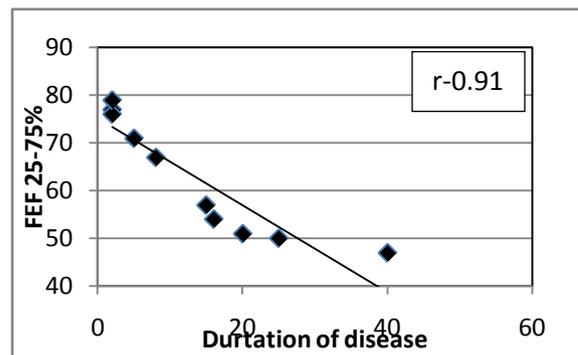


Figure 6

Duration of disease and pulmonary function disease parameters were negatively correlated which was shown in Fig-V and FIG-VI. Eighty percent of the study population were malnourished based on the calorie intake and calorie deficiency which was focused in Table III. In our study Calorie intake was significantly negatively correlated and calorie deficiency was statistically positively correlated which are shown in Fig-I and Fig-II and also statistically significant. This study showed that both SFT and TBF were very much reduced in all the subjects irrespective of BMI. Our study

showed significant negative correlation between duration of disease and SFT, TBF (Fig III, IV).

IV. DISCUSSION

Seventy percent of the COPD patients of our study with mild to severe disease had normal Body weight and BMI, this could be due to depletion of lean tissues (De Benedetto et al 2000).

Recent studies revealed that the regenerative capacity of skeletal muscle is impaired in mice with elevated circulating tumor necrosis factors (TNF) levels

(Langen et al 2006), lower testosterone (Vliet et al 2005), due to chronic hypoxia and corticosteroid therapy (Kamischke et al 1998).

In our study BMI of patients with COPD were negatively correlated with disease duration. This prevalence of malnutrition may be due to systemic inflammation, Low dietary intake (chronic mouth breathing, aerophagia, Dyspnea, old age), bronchodilators, corticosteroids, antibiotics. Elevated circulating leptin level in COPD patients may affects dietary intake and consequently muscle mass and function (Schols 2003).

Expiratory air flow limitation is the key to diagnose the severity of disease and traditional physiological changes in patients with COPD. This could be due to both small and peripheral airway obstruction and consequent increase in airway resistance. Loss of small airway patency due to destruction of alveolar tissues may play important role. Low FEV1, FEF 25-75% and FEV1% indicate the severity of disease of COPD patients. The airflow obstruction may increase the cost of breathing (Aliverti and Macklem 2001) which cause structural changes in the respiratory muscles due to the continuous overload (Orozco-Levi et al 2001)

The energy requirements of a healthy person vary depending on a number of factors including: age; gender; body composition; current and past nutritional status; and basal metabolic rate (BMR). BMR may be defined as the metabolic activity required for the maintenance of life including respiration, heartbeat and body temperature. When people experience illness, injury or surgery, their BMR increases. This causes metabolic stress, which, if uncontrolled, can lead to weight loss and eventually malnutrition. Without sufficient energy, protein stores in the body are mobilised from skeletal muscle, resulting in loss of lean body mass. This protein is broken down via biochemical oxidation to meet the body's increased energy needs. If the person's diet does not contain enough protein, this will lead to a negative nitrogen balance (Bongers et al, 2007). A positive nitrogen balance is essential for tissue repair after illness or major trauma (Soeters et al, 2004)

V. CONCLUSION

Being an inflammatory disease COPD involves with lungs and affect other body tissues like bones and muscles, these are known as co morbidities. Diet and nutritional intake are important in COPD because they help to combat some of these co morbidities. So nutrition is an important therapy in the management of patients with COPD.

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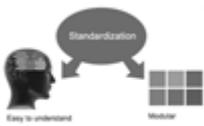
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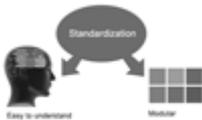
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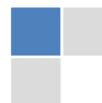
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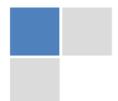
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Content

- Sum up your conclusion in text and demonstrate them, if suitable, with figures and tables.
- In manuscript, explain each of your consequences, point the reader to remarks that are most appropriate.
- Present a background, such as by describing the question that was addressed by creation an exacting study.
- Explain results of control experiments and comprise remarks that are not accessible in a prescribed figure or table, if appropriate.
- Examine your data, then prepare the analyzed (transformed) data in the form of a figure (graph), table, or in manuscript form.

What to stay away from

- Do not discuss or infer your outcome, report surroundings information, or try to explain anything.
- Not at all, take in raw data or intermediate calculations in a research manuscript.
- Do not present the similar data more than once.
- Manuscript should complement any figures or tables, not duplicate the identical information.
- Never confuse figures with tables - there is a difference.

Approach

- As forever, use past tense when you submit to your results, and put the whole thing in a reasonable order.
- Put figures and tables, appropriately numbered, in order at the end of the report
- If you desire, you may place your figures and tables properly within the text of your results part.

Figures and tables

- If you put figures and tables at the end of the details, make certain that they are visibly distinguished from any attach appendix materials, such as raw facts
- Despite of position, each figure must be numbered one after the other and complete with subtitle
- In spite of position, each table must be titled, numbered one after the other and complete with heading
- All figure and table must be adequately complete that it could situate on its own, divide from text

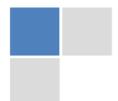
Discussion:

The Discussion is expected the trickiest segment to write and describe. A lot of papers submitted for journal are discarded based on problems with the Discussion. There is no head of state for how long a argument should be. Position your understanding of the outcome visibly to lead the reviewer through your conclusions, and then finish the paper with a summing up of the implication of the study. The purpose here is to offer an understanding of your results and hold up for all of your conclusions, using facts from your research and generally accepted information, if suitable. The implication of result should be visibly described. Infer your data in the conversation in suitable depth. This means that when you clarify an observable fact you must explain mechanisms that may account for the observation. If your results vary from your prospect, make clear why that may have happened. If your results agree, then explain the theory that the proof supported. It is never suitable to just state that the data approved with prospect, and let it drop at that.

- Make a decision if each premise is supported, discarded, or if you cannot make a conclusion with assurance. Do not just dismiss a study or part of a study as "uncertain."
- Research papers are not acknowledged if the work is imperfect. Draw what conclusions you can based upon the results that you have, and take care of the study as a finished work
- You may propose future guidelines, such as how the experiment might be personalized to accomplish a new idea.
- Give details all of your remarks as much as possible, focus on mechanisms.
- Make a decision if the tentative design sufficiently addressed the theory, and whether or not it was correctly restricted.
- Try to present substitute explanations if sensible alternatives be present.
- One research will not counter an overall question, so maintain the large picture in mind, where do you go next? The best studies unlock new avenues of study. What questions remain?
- Recommendations for detailed papers will offer supplementary suggestions.

Approach:

- When you refer to information, differentiate data generated by your own studies from available information
- Submit to work done by specific persons (including you) in past tense.
- Submit to generally acknowledged facts and main beliefs in present tense.



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<i>References</i>	Complete and correct format, well organized	Beside the point, Incomplete	Wrong format and structuring



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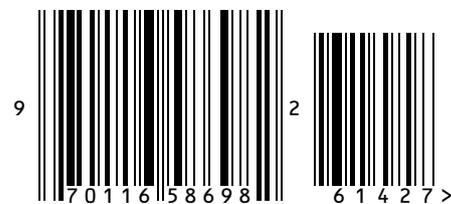
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