Effect of Diode Laser on Human Cytomegalovirus and Epstein - Barr Virus in Chronic Periodontitis Patients – A Quantitative Real-Time Polymerase Chain Reaction Study

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Abstract— Background and aims: Studies have investigated the occurrence of Human cytomegalovirus (HCMV) and Epstein Barr Virus (EBV) in samples of both chronic and aggressive periodontitis lesions and their role in the pathogenesis of the disease have been suggested. It is important to determine the role and presence of these viruses in chronic generalized periodontitis patients after Scaling and root planing (SRP) alone and in conjunction with diode laser application. Thus the aim of the present study was to determine the prevalence of human cytomegalovirus (HCMV), Epstein-Barr virus (EBV) in gingival crevicular fluid (GCF) samples obtained from chronic generalised periodontitis patients.

Materials (Subjects) and Methods: A total of 30 patients were included in the study based on the inclusion and exclusion criteria as follows: Group I – 15 chronic generalized periodontitis patients who received treatment with scaling and root planing. Group II – 15 chronic generalized periodontitis patients who received treatment with scaling and root planing and Diode laser. In each patient GCF samples were collected at baseline, 7 days and 21 days after treatment. Clinical parameters like gingival index, probing pocket depth and clinical attachment level were recorded and as GCF samples collected.

Results: Correlation between quantification of EBV values with clinical parameters such as gingival index, probing pocket depth and clinical attachment loss in two groups showed statistical significance ($P < 0.001$). Relative quantification of EBV was statistically significant, whereas HCMV was significantly absent. The mean values of GI, PPD and CAL parameters at different time interval was found to be statistically significant in both group ($P < 0.001$)

Conclusions: Only EBV was detected more frequently in patients with CGP, but HCMV was not detected with similar frequency in the groups studied. We also found that group with application of diode laser showed statistical significant results.

Index Terms— Chronic generalised periodontitis, Diode laser, Epstein – Barr virus, Gingival crevicular fluid, Human cytomegalovirus, PCR, Scaling and Root planning.

I. INTRODUCTION

Periodontitis is a disease attributable to multiple infectious agents and interconnected cellular and humoral host immune responses. However; it has been difficult to unravel the precise role of various putative pathogens and host responses in the pathogenesis of periodontitis. It is not understood why, in hosts with comparable levels of risk factors, some periodontal infections result in loss of periodontal attachment and alveolar bone, while other infections are limited to inflammation of the gingiva with little or no discernible clinical consequences. Also, many periodontitis patients do not show a remarkable level of classical risk factors [1], [2].

Detection and quantification of periodontopathic bacterial species are useful for identifying subjects at elevated risk of periodontitis, but do not consistently predict clinical outcome. These uncertainties have galvanized efforts to find additional etiologic factors for periodontitis. Even though specific infectious agents are of key importance in the development of periodontitis, it is unlikely that a single agent or even a small group of pathogens are the sole cause or modulator of this heterogeneous disease [3].

Since the mid-1990s, herpes viruses have emerged as putative pathogens in various types of periodontal disease. 4,5,6) In particular, human cytomegalovirus (HCMV) and Epstein-Barr virus (EBV) seem to play important roles in the etiopathogenesis of severe types of periodontitis.

Genomes of the two herpesviruses occur at high frequency in disease progressive as seen in progressive periodontitis in adults, localized and generalized aggressive (juvenile) periodontitis, HIV-associated periodontitis, acute necrotizing ulcerative gingivitis, periodontal abscesses, and some rare types of advanced periodontitis associated with medical disorders. 7,8) HCMV infects periodontal monocytes/ macrophages and T-lymphocytes, and EBV infects periodontal B-lymphocytes [9]

According to Slots [10] periodontal destruction may be associated with the coexistence of periodontal herpes viruses, especially human cytomegalovirus (HCMV) and Epstein-Barr virus (EBV), periodontopathogenic bacteria, and the local impaired host immune response. AgP sites presenting HCMV, EBV-1, or herpes simplex virus (HSV) had a higher occurrence of *P. gingivalis* and *D. pneumosintes* than sites without the viruses [11]. Also, Aggressive periodontitis (AgP) sites infected with active HCMV harbored more *A. actinomycetemcomitans* than sites with latent HCMV infection [12].

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Herpesvirus-infected inflammatory cells elicit tissue-destroying cytokines and may exert diminished ability to defend against bacterial challenge [13]. Herpesvirus-associated periodontal sites also tend to harbor elevated levels of periodontopathic bacteria, including *Porphyromonas gingivalis*, *Tannerella forsythia*, *Dialister pneumosintes* / *Dialister invisus*, *Prevotella intermedia*, *Prevotella nigrescens*, *Treponema denticola*, *Campylobacter rectus* and *Actinobacillus actinomycetemcomitans* [14], [15]. Transcripts of HCMV and EBV have been identified in the great majority of symptomatic periapical lesions as well. In the light of the close statistical relationship between herpesviruses and periodontitis, it is reasonable to surmise that some cases of the disease have a herpesviral component.

Saygun et al. reported that HCMV, EBV-1, and HSV-1 were positively associated with *P. gingivalis*, *P. intermedia*, *T. forsythia*, and *C. rectus* but not with *A. actinomycetemcomitans* in young adults with advanced periodontitis [16].

The herpesviral infection can stimulate the release of cytokines and chemokines from inflammatory and non-inflammatory cells and impair the periodontal immune defense, resulting in more virulent resident bacteria [13]. Periodontal bacterial infection elicits an inflammatory response in periodontal tissues, resulting in dense infiltration of the T- lymphocytes, B lymphocytes, and macrophages [13].

However, from our systematic literature review, there is limited data available on the effect of diode laser treatment on the reduction of viral load in South Indian population. Hence this study has been undertaken to evaluate the effect of diode laser on the prevalence of HCMV & EBV in chronic periodontitis using real time polymerase chain reaction technique. Thus the aim of the present study was to determine the prevalence of human cytomegalovirus (HCMV), Epstein-Barr virus (EBV) in gingival crevicular fluid (GCF) samples obtained from chronic periodontitis patients. In addition, the effect of diode laser in periodontal treatment was to examine the long-term effect of diode laser therapy on periodontal pockets with regard to the improvement of periodontal condition.

II. MATERIALS AND METHODS

A. Patients

Patients who visited outpatient department of Periodontology, Rajarajeswari Dental College and Hospital, Bangalore were selected for this study between June 2014 and August 2015. Inclusion criteria were: a) Patient having a minimum of 20 teeth, b) Sites having a probing depth ≥ 5 mm, c) Sites having clinical attachment loss ≥ 3 mm, d) Radiographic evidence of alveolar bone loss. Exclusion criteria were: a) Patients with history of systemic diseases affecting the periodontium, b) Smokers, c) Patients on any medication taken within the last 6 months which may alter the periodontal status, d) Pregnant and lactating mothers, e) Patients who have undergone periodontal treatment within a period of 1 year.

The study protocol was approved by the Research Ethics Committee of Rajarajeswari Dental college and hospital. All

patients were informed about the study and gave written informed consent to participate.

All the subjects underwent a full mouth periodontal probing and charting and were screened for suitability. A proforma was designed for the present study so as to have a systematic and methodical recording of all the observations and information. The relevant data were recorded in the proforma. Chronic periodontitis patients were diagnosed based on criteria discussed at the 1999 International workshop of periodontology.

A total of 30 patients included in the study and were grouped as follows:

Group I – 15 Chronic generalized Periodontitis patients were treated with scaling and root planing alone.

Group II – 15 chronic generalized periodontitis patients were treated with scaling and root planing and Diode laser.

B. Clinical parameters

All baseline and follow up, intrasurgical measurements were carried out by the same examiner. Measurements were made using a graduated William's periodontal probe and were rounded off to the nearest millimeter

Gingival index (Loe H & Silness P, 1963): Gingival index (GI) was assessed using a mouth mirror and probe. The teeth and gingiva were dried lightly with cotton rolls. The tissues surrounding each tooth were divided into 4 gingival scoring units: distal-facial papilla, facial margin, mesial-facial papilla and the entire lingual gingival margin. Each of the four gingival units was assessed.

Probing Pocket Depth (PPD): The graduated William's periodontal probe was inserted parallel to the long axis of the tooth to measure the pocket depth at 3 buccal (mesiobuccal, midbuccal and distobuccal) and 3 lingual (mesiolingual, midlingual and distolingual) sites for all maxillary and mandibular teeth. The pocket depth was evaluated by determining the distance between crest of the gingival margin to the base of the pocket. Mean pocket depth for a subject was calculated by adding measured pocket depth around all the teeth and divided by number of teeth examined.

Clinical attachment level (CAL): CAL was measured as the distance from the cemento-enamel junction (CEJ) to the base of the gingival sulcus using the periodontal probe.

Mean CAL for a subject was calculated by adding measured pocket depth around all the teeth and divided by number of teeth examined.

GI, PPD and CAL were recorded at baseline before any non-surgical periodontal therapy started and also 7 and 21 days after the therapy.

III. SAMPLE COLLECTION

All the subjects were examined and screened for the study. Patients were informed and written consent was obtained. GCF was collected using sterile paper point by inserting into deep periodontal pocket without applying pressure. GCF sample was collected 24 hours after periodontal examination in order to avoid blood contamination of samples. All samples were collected from the deepest periodontal pocket.

The sample was isolated from saliva with cotton rolls and gently air dried. The supragingival plaque was removed by

sterile cotton pellets. Two paper points were inserted in each periodontal pocket until a mild resistance for 30 seconds. Paper point of ISO # 40 was used. Paper points contaminated with blood were not used in the analysis. These paper points were placed in sterile plastic tubes containing TE buffer and stored at -70° C until further analysis.

In Group 1, each patient received scaling and root planning using Gracey curettes until a hard, smooth and calculus-free root surface was achieved.

In Group 2, Diode laser ((SIROLaser®, Sirona, Bensheim, Germany) 980 nm at a power output of 2.5W in pulsed mode (30 Hz, pulse duration 60 s) was used as an adjunctive to scaling and root planing. The optic fiber of 400µm was moved from the coronal to apical side of the pocket in parallel paths with an inclination of approximately 20°. Each pocket of the test group was lasered for 30s twice, with a 60 s interval.

IV. TREATMENT PROCEDURE

All patients received full mouth disinfection i.e. scaling and root planing along with chlorhexidine applications within 24 hours.

A. Scaling and root planing:

Scaling and root planning was performed by the same investigator in both the groups using EMS scaler and standard periodontal curettes. Local anesthesia was used whenever needed. The time needed for the procedure was approximately 1 hour. Recall examinations were done at 7th and 21st day after the treatment for assessment of clinical parameters and collection of GCF samples for viral analysis.

B. For sample storage: -70° C refrigerator (New Brunswick Scientific)

C. DNA extraction

DNA extraction from the GCF samples was done using highly purified Invitrogen DNA isolation kit (Purelink™ DNA extraction kit) Applied Biosystems, India. Before DNA extraction, the paper point was completely submerged in TE buffer and kept in room temperature for 2hrs.

1. The vials containing TE buffer was centrifuged for 3000rpm for 2minutes.
2. About 200 µl of the solution was transferred into sterile micro centrifuge tube.
3. Then 20 µl of proteinase K was added to a sterile microcentrifuge tube.
4. The tubes were kept at 55°C for 2 hours in water bath with occasional vortexing unless and until its completed.
5. 20 µl of RNase-A was added to the lysate and mixed well by briefly vortexing and was incubated at room temperature for 2 minutes.
6. 200 µl of Purelink™ Genomic Lysis/Binding buffer was added and mixed well by vortexing to obtain a homogenous solution.
7. 200 µl of 96-100 % ethanol was then added and mixed well by vortexing for 5 seconds to obtain a homogeneous solution and was subjected to purification protocol immediately.

D. DNA Purification

1. The purification procedure is designed for purifying genomic DNA using a spin- column based centrifugation procedure on a total time of 10-15 minutes.
1. Purelink™ spin-column was removed from the package and placed in a clean collection tube.
2. The entire lysate prepared with Purelink™ Genomic Lysis/Binding buffer and ethanol was added to the spin-column.
3. The column was centrifuged at 10000 rpm for 1 minute at room temperature.
4. The spin-column was then placed into a clean Purelink™ collection tube supplied with the kit.
5. 500 µl of “Wash Buffer-1” prepared with ethanol was added to the column.
6. The column was centrifuged at maximum speed for 3 minutes at room temperature. The collection tube was discarded.
7. The spin-column was placed in a sterile 1.5 ml micro-centrifuge tube.
8. 40 µl of Purelink™ genomic elution buffer was added to the column and incubated at room temperature for 1 minute.
9. The column was then centrifuged at maximum speed for 1 minute at room temperature thus obtaining purified genomic DNA in the tube.
10. Purified DNA was stored at -20°C until further processing.

E. Primers and probes

Custom TaqMan inventoried assay for pathogen identification was used to identify HCMV. The details are as follows and available at www.lifetechnologies.com/order/genome-database/details/gene-expression/Pa03453400s1.

Assay ID for HCMV is Pa03453400_s1, the gene symbol is UL132 and the catalogue number is 4331182. Assay ID for EBV is Pa03453399_s1, the gene symbol is IR1 and the catalogue number is 4331182.

The sequences of HCMV and EBV used are as follows:

The assay is designed to a sequence that does not exhibit typical exon-intron-exon gene structure such as fusion transcript or artificial prokaryotic or mitochondrial sequences. As the assay procured from Applied Biosystem / Life technologies was inventoried, the individual HCMV and EBV primer and probe sequences are not provided or disclosed by the manufacturer.

F. PCR procedure

The PCR procedure was carried out in laboratory for molecular biology in Rajarajeswari Dental College and Hospital, Bangalore. Analysis of HCMV and EBV in the gingival crevicular fluid samples was done using quantitative real time polymerase chain reaction.

StepOne™ (48-well) Real-time PCR Systems by Life Technologies™ and reagents from Applied Biosystems™ were used in this study. Custom TaqMan pathogen identification assay for HCMV and EBV was used in this study. TaqMan Polymerase Reaction was performed with a final volume of 20µl Polymerase Chain Reaction mixture containing (Table 1) 10µl TaqMan universal PCR mastermix (Applied biosystems), 1µl custom TaqMan assay containing probes and primers for specific virus (Human Cytomegalovirus and Epstein-Barr virus), 3µl of extracted DNA sample and 6µl RNase free water to make up to 20µl.

Each virus was identified in separate reaction for each sample.

TABLE 1: CONTENTS OF THE POLYMERASE CHAIN REACTION MIXTURE

Contents of PCR mixture	Volume in μ l
TaqMan universal PCR mastermix	10
Custom TaqMan assay containing probes and primers specific virus	1
Extracted DNA sample	3
RNase free water	6
Final volume of reaction mixture	20

The condition for Real Time Polymerase Chain Reaction was as follows. Pre-heating at 50°C for 2 minutes and at 95°C for 10 minutes, followed by 40 cycles of shuttle heating at 95°C for 15 seconds and at 60°C for 1 minute. 16S RNA was used as the endogenous control (TaqMan pre-developed assay reagents Applied Biosystems – India). The comparative threshold cycle or Ct method was used to assess the amplified transcripts.

V. STATISTICAL ANALYSIS

The following methods of statistical analysis have been used in this study. Data was entered in Microsoft excel and analysed using SPSS (Statistical Package for Social Science, Ver.10.0.5) package.

The results were averaged (mean + standard deviation) for continuous data and number and percentage for dichotomous data are presented in Table and Figure. Normality of data was tested using Shapiro-Wilk test.

Proportions were compared using Chi-square (χ^2) test of significance. Proportion of cases belonging to specific group of parameter or having a particular problem was expressed in absolute number and percentage. The student 't' test was used to determine whether there was a statistical difference between groups in the parameters measured if the data is normal. One way analyses of variance were used to test the difference between groups. If F value is significant there is a significant, difference between group means. To find out which of the two groups means is significantly difference post hoc test of Tukey test is used. In case of F value is not significant it indicates that there is no significant difference between the groups and stops the analysis at this stage and does not used Tukey test.

The Kruskal-Wallis H test was used to determine if there are statistically significant differences between two or more groups of an independent variable on a continuous or ordinal dependent variable. Mann-Whitney U test is used to compare differences between two independent groups when the dependent variable is either ordinal or continuous, but not normally distributed.

VI. RESULTS

A. Figures and Tables

The present randomized controlled study was undertaken to quantitatively analyse the levels of HCMV and EBV in GCF samples obtained from chronic generalized periodontitis patients who were treated with scaling and root planing and diode laser. We also assessed the effect of diode laser on the prevalence of HCMV and EBV counts in GCF of chronic generalized periodontitis patients using

quantitative polymerase chain reaction. The study population consisted of 30 subjects with 15 in each group.

Mean age of group 1 and group 2 was 43.93 ± 5.65 and 43.60 ± 5.48 respectively as shown in Table 2, 3 and Fig 1.

Fig 1: Age distribution of the study groups

TABLE 2: AGE DISTRIBUTION OF PATIENTS

Group	N	Mean age	SD	Median	Min.	Max.	't' value	'p' value
Group 1	15	43.9	5.650	43.0	36	54	0.027	0.871
Group 2	15	43.6	5.475	44.0	36	54		

TABLE 3: DISTRIBUTION OF PATIENTS AMONG TWO GROUPS

Group	N	Mean age	SD	Median	Min.	Max.	't' value	tABLE
Group 1	15	43.9	5.650	43.0	36	54	0.027	0.871
Group 2	15	43.6	5.475	44.0	36	54		

In both the groups, maximum subjects were belonging to 31- 60 years. Gender was almost equally distributed in subjects in both groups (Table 4, Fig 2). Gender distribution of the subjects in the two different groups is shown in table 4 and figure 3. Among 15 subjects in group 1, 8 males (53.3%) and 7 females (46.7%) were included in the study, and in group 2, it was 7 males (46.7%) and 8 females (53.3%).

TABLE 4: DISTRIBUTION OF PATIENTS AMONG TWO GROUPS BY GENDER

	Gender			χ^2 value	'p' value
	Male	Female			
Group 1	8	7	15	0.133	0.715
	53.3%	46.7%	100.0%		
Group 2	7	8	15		
	46.7%	53.3%	100.0%		
Total	15	15	30		
	50.0%	50.0%	100.0%		

TABLE 5: COMPARISON OF GINGIVAL INDEX BETWEEN TWO GROUPS AT BASELINE TO 7TH AND 21ST DAY.

GROUPS AT BASELINE TO 7TH AND 21ST DAY.									
		N	Me an	SD	Medi an	Mi n.	Mi n.	Mann - Whitn ey U	'p' val ue
Baseli ne	Gro up 1	1 5	2.4 0	0.1 46	2.30	2.2	2.6	101.0 0	0.6 10
	Gro up 2	1 5	2.4 2	0.1 37	2.30	2.3	2.6		
7 days	Gro up 1	1 5	1.5 1	0.1 36	1.50	1.3	1.3	45.00 0	0.0 03
	Gro up 2	1 5	1.3 0	0.0 0	1.30	1.3	1.3		

	up 2	5	7	98					
21 days	Gro up 1	1 5	1.1 6	0.1 24	1.20	0.9	0.9	94.00 0	0.4 24
	Gro up 2	1 5	1.1 1	0.1 71	1.00	0.8	0.8

Fig 2: Gender distribution of study groups.

The difference in mean gingival index between group 1 and group 2 at various time intervals was statistically not significant (table 5, fig 3) whereas within each group the difference in mean gingival index was statistically significant at various time intervals. Pairwise comparison within group 1 and group 2 at various time intervals showed statistically significant difference (table 6, fig 4)

TABLE 6: COMPARISON GINGIVAL INDEX WITHIN EACH GROUP FROM BASELINE TO 7TH AND 21ST DAY.

			Me an	SD	Medi an	Mi n.	Mi n.	Chisqu re*	'p' valu e
Gro up 1	Basel ine	5	2.4 0	0.1 46	2.30	2.2	2.6	332.46 4	<0.0 01
	7 days	5	1.5 1	0.1 36	1.50	1.3	1.7		
	21 days	5	1.1 6	0.1 24	1.20	0.9	1.3		
Gro up 2	Basel ine	5	2.4 2	0.1 37	2.30	2.3	2.6	377.73 4	< 0.00 1
	7 days	5	1.3 7	0.0 98	1.30	1.3	1.5		
	21 days	5	1.1 1	0.1 71	1.00	0.8	1.3

Fig 3: Comparison of Gingival Index between Study groups at baseline to 7 and 21 days

Fig 4: Comparison of Gingival Index between different visit by Study Groups

The difference in mean pocket depth between group 1 and group 2 at various time intervals was not statistically significant (table 7, fig 5). Within each group the difference in mean probing pocket depth at various time intervals was statistically significant. Pairwise comparison within group 1 and group 2 at various time intervals showed statistically significant difference (table 8, fig 6)

TABLE 7-COMPARISON OF PROBING POCKET DEPTH BETWEEN TWO GROUPS FROM BASELINE TO 7TH AND 21ST DAY.

Visit	Gro up	N	Mea n	SD	Medi an	Mi n.	Mi n.	't' valu e	'p' valu e
Basel ine	Gro up 1	5	6.95	0.35 0	7.00	6.3	7.5	0.00 3	0.95 9
	Gro up 2	5	6.95	0.34 6	6.90	6.3	7.5		
7 days	Gro up 1	5	6.10	0.29 5	6.10	5.5	6.5	0.27 7	0.60 3
	Gro up 2	5	6.15	0.25 9	6.10	5.5	6.5		
21 days	Gro up 1	5	5.68	0.31 2	5.60	5.2	6.3	2.40 0	0.13 3
	Gro up 2	5	5.83	0.22 3	5.80	5.3	6.2

Fig 5: Comparison of Probing Pocket Depth between Study groups at baseline to 7 and 21 days

TABLE 8-COMPARISON OF PROBING POCKET DEPTH WITHIN EACH GROUP FROM BASELINE TO 7TH AND 21ST DAY.

Gro up	Vi sit	N	Me an	SD	Medi an	Mi n.	Mi n.	'F' valu e	'p' valu e
Gro up 1	Baseli ne	5	6.9 5	0.3 50	7.00	6.3	7.5	60.9 71	<0.0 01
	7 days	5	6.1 0	0.2 95	6.10	5.5	6.5		
	21 days	5	5.6 8	0.3 12	5.60	5.2	6.3		
Gro up 2	Baseli ne	5	6.9 5	0.3 46	6.90	6.3	7.5	63.3 81	<0.0 01

7 days	5	6.1	0.2	6.10	5.5	6.5	
21 days	5	5.8	0.2	5.80	5.3	6.2

Fig 6: Comparison of Probing Pocket Depth between different visit by Study Groups

The difference in the mean clinical attachment loss in between group 1 and group 2 was statistically not significant. It is found that in group 1 the mean CAL at various time intervals was not statistically significant (table 9, fig 7). Intra group analysis of group 2 showed that the difference in mean CAL at various time intervals was statistically significant. Pairwise analysis in group 2 showed between baseline and 21st day was statistically significant, whereas group1 at various time intervals was not significant (table 10, fig 8).

TABLE 9-COMPARISON OF CLINICAL ATTACHMENT LEVEL BETWEEN TWO GROUPS FROM BASELINE TO 7TH AND 21ST DAY.

Visit	Group	N	Mean	SD	Median	Min.	Max.	t' value	p' value
Baseline	Group 1	5	7.07	0.403	7.20	6.3	7.5	1.005	0.325
	Group 2	5	7.20	0.321	7.30	6.6	7.6		
7 days	Group 1	5	6.80	0.434	7.00	6.0	7.3	0.395	0.535
	Group 2	5	6.89	0.377	7.00	6.3	7.3		
21 days	Group 1	5	6.67	0.398	7.00	6.0	7.0	0.810	0.376
	Group 2	5	6.55	0.283	6.50	6.0	7.0

Fig 7: Comparison of Clinical Attachment Level between Study groups at baseline to 7 and 21 days

TABLE 10-COMPARISON OF CLINICAL ATTACHMENT LEVEL WITHIN EACH GROUP FROM BASELINE TO 7TH AND 21ST DAY.

Group	Visit	N	Mean	SD	Median	Min.	Max.	F' value	p' value
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Group 1	Baseline	5	7.07	0.403	7.20	6.3	7.5	3.667	0.034
	7 days	5	6.80	0.434	7.00	6.0	7.3		
	21 days	5	6.67	0.398	7.00	6.0	7.0		
Group 2	Baseline	5	7.20	0.321	7.30	6.6	7.6	14.499	<0.001
	7 days	5	6.89	0.377	7.00	6.3	7.3		
	21 days	5	6.55	0.283	6.50	6.0	7.0

Fig 8: Comparison of Clinical Attachment level between different visit by Study Groups

The difference in the mean RQ of EBV value in group 1 and group 2 at various time intervals was not statistically significant (table 11, fig 9). Within group 1 the mean RQ value of EBV at baseline, 7 and 21 days were 28.53, 19.52 and 0.19 respectively. The difference in mean RQ values in the above said intervals was statistically significant ($p < 0.001$). Within group 2 the mean RQ value of EBV at baseline, 7th and 21st days were 21.88, 3.36 and 0.05 respectively. The difference in mean RQ values in the above said intervals was statistically significant ($p < 0.001$) (table 12, fig 10).

TABLE 11 - COMPARISON OF RELATIVE QUANTIFICATION OF EBV SCORES BY MANN- WHITNEY U TEST BETWEEN TWO GROUPS FROM BASELINE TO 7TH AND 21ST DAY.

Visit	Group	N	Mean	SD	Median	Min.	Max.	Mann - Whitney U	p' value
Baseline	Group 1	5	28.531	25.521	26.182	3.97	79.84	32.000	0.470
	Group 2	5	21.886	24.098	13.432	1.00	79.84		
7 days	Group 1	5	19.526	28.526	7.688	1.06	61.67	4.500	0.176
	Group 2	5	3.367	5.108	1.843	0.03	12.37		
21 days	Group 1	5	0.199	0.335	0.073	0.00	1.06	21.000	0.266
	Group 2	5	0.050	0.054	0.018	0.00	0.14

Fig 9: Comparison of EBV Values between Study groups at baseline to 7 and 21 days

TABLE 12 - COMPARISON OF RELATIVE QUANTIFICATION OF EBV SCORES BY CHISQUARE TEST IN EACH GROUP FROM BASELINE TO 7TH AND 21ST DAY.

Gro up	Visit	N	Mea n	SD	Med ian	Mi n.	Ma x.	Chisq ure*	'p' valu e
Gro up 1	Basel ine	5	28.5 31	25.5 21	26.1 82	3.9 7	79. 84	14.797	<0.0 01
	7 days	5	19.5 26	28.5 26	7.68 8	1.0 6	61. 67		
	21 days	5	0.19 9	0.33 5	0.07 3	0.0 0	1.0 6		
Gro up 2	Basel ine	5	21.8 86	24.0 98	13.4 32	1.0 0	79. 84	14.652	<0.0 01
	7 days	5	3.36 7	5.10 8	1.84 3	0.0 3	12. 37		
	21 days	5	0.05 0	0.05 4	0.01 8	0.0 0	0.1 4

Fig 10: Comparison of EBV Values between different visit by Study Groups.

It was found that the effect of diode laser as an adjunct to SRP showed greater reduction of EBV levels in concentration in chronic generalized periodontitis patients compared to SRP.

Out of 90 samples of GCF taken from both the groups, EBV was present in most of the samples, whereas HCMV was not detected in any of the samples across the group. The prevalence of EBV was found higher in both the groups and also the quantity increased in line with higher clinical parameters.

VII. DISCUSSION

The present randomized controlled trial was undertaken to determine the prevalence of human cytomegalovirus (HCMV) and Epstein-Barr virus (EBV) in gingival crevicular fluid (GCF) of chronic generalized periodontitis patients. We also evaluated the effect of scaling and root planing (SRP) and diode laser on the quantity of HCMV and

EBV in chronic generalized periodontitis patients using quantitative real time polymerase chain reaction. According to our knowledge this is the first study conducted in this regard.

Our results showed a high prevalence of EBV in GCF samples (46% in group 1 and 60% in group 2) from deep periodontal pockets similar to studies conducted by Watanabe SA et al. [65] where they found a significantly higher occurrence of EBV in periodontal sites than in gingivitis sites. Watanabe et al [17] discussed that higher occurrence of EBV in deep periodontal sites is consistent with the herpes viruses playing a role in the pathogenesis of periodontitis.

Contreras A et al [18] reported that EBV was the second most prevalent virus detected occurring in 50% of tissue biopsy specimens and in 45% of B-lymphocyte fractions from the adult periodontitis lesions. EBV is a potent polyclonal B-lymphocyte activator, capable of inducing proliferation and differentiation of immunoglobulin secreting cells.

EBV affects over 90% of humans and is usually transmitted by oral secretions or blood. The virus replicates in epithelial cells or B-cells of the oropharynx. Resting memory B cells are the main site of persistence of EBV in the body. The number of latently infected cells in a person remains stable for over years [19] Latent EBV infection can be reactivated, leading to viral shedding into oral mucosa [20].

Konstantinidis et al [6] showed EBV to be present in 11 out of 12 chronic periodontitis patients and only one in healthy controls. Similarly, in aggressive periodontitis patients, EBV was found to be significant (32%) compared with healthy controls (8%). However, no difference was found between chronic periodontitis and aggressive periodontitis patients in this study. This is contradictory to the study results obtained by Kubar et al [21] where EBV was seen to be present in 89% sites in aggressive periodontitis patients compared with 46% in chronic periodontitis patients, which was statistically significant.

Some investigators have suggested that once EBV is present (or active), its herpes virus mediated impairment of the host defense may give rise to subgingival growth of periodontal pathogens and subsequent periodontal attachment loss [17], [22]. Saygun et al [23] found EBV in 44.3% of sites from patients with chronic periodontitis and only 14.3% of the sites obtained from healthy controls.

The results of the present study revealed an association between the presence of EBV and chronic periodontitis lesions (PD =5 mm). Our results correlated with that of previous studies that showed statistically significant levels of EBV in patients with chronic periodontitis. Slots et al. discovered more EBV in the gingival crevicular fluid and saliva of patients with periodontal diseases than in the saliva of an otherwise healthy control group [4], [24]. The same group demonstrated a correlation between EBV prevalence in periodontal patients and periodontal pocket depth [21], [25], [26]. Because EBV detection using nested PCR is a qualitative test. The results showed that real-time PCR data was consistent with nested PCR data. Real-time PCR did detect the presence of EBV in those sites in which nested PCR had detected the presence of the DNA. In addition,

least one order or more of EBV was detected in deeper probing depth sites compared with shallow probing depth sites.

Contreras et al [9] have detected HCMV in monocytes / macrophages and T lymphocytes, EBV in B lymphocytes, and HSV in T lymphocytes and monocytes / macrophages. Infected immune cells may not mount a proper immune response against periodontopathogenic bacteria predisposing to microbial superinfection. Herpes virus induced defective polymorphonuclear neutrophils can increase the risk for destructive periodontal disease. The above said study found herpes viruses infecting various types of inflammatory cells in periodontitis lesions. It was not known whether the infected cells carry functional or latent herpes viruses. Active herpes virus infections would potentially be more detrimental to the periodontium than latent herpes virus infections.

Previous studies have shown the association of HCMV to periodontitis sites, [4], [18], [21], [27], [23] which are in contrast with our results as HCMV was not detected in chronic periodontitis patient. This discrepancy may also be due to sample size, selection of the subjects evaluated, or ethnic differences. The absence of HCMV could denote the stability of periodontitis sites in the sample, since a previous study observed a higher frequency of HCMV in progressing periodontitis sites compared to that of stable sites [28].

In a study conducted by Ting et al [29] it was hypothesized that active HCMV infection could be associated with the initiation and progression of localized juvenile periodontitis. In another study conducted by Contreras and Slots [30] results suggested that active HCMV replication could occur in periodontitis sites, but it remained unclear if HCMV reactivation was related to the initiation or the progression of destructive periodontal disease.

The present findings are consistent with the premise that high levels of EBV in GCF samples reflect increased likelihood of several subgingival and periodontal sites having high EBV burden. In contrast, the low prevalence of HCMV in GCF samples suggests that this virus is an infrequent participant of periodontal disease in immunocompetent individuals. Interestingly, we did not observe a relationship between the presence and amount of HCMV in chronic generalized periodontitis patients.

Haffajee AD et al. had done a study to examine the effect of SRP on clinical and microbiological parameters in 57 subjects with adult periodontitis. Subjects were monitored clinically and microbiologically prior to and 3, 6 and 9 months after full-mouth SRP under local anaesthesia. They observed that mean prevalences and levels of *P. gingivalis*, *T. denticola* and *B. forsythus* were significantly reduced after SRP, while *A. viscosus* showed a significant increase in mean levels. The mean decrease in prevalence of *P. gingivalis* was similar at all pocket depth categories, while *B. forsythus* decreased more at shallow and intermediate pockets and *A. viscosus* increased most at deep sites. *P. gingivalis*, *B. forsythus* and *T. denticola* were equally prevalent. Clinical improvement post-SRP was accompanied by a modest change in the subgingival microbiota, primarily a reduction in *P. gingivalis*, *B. forsythus* and *T. denticola*, suggesting potential targets for therapy and indicating that

radical alterations in the subgingival microbiota.

Grenier G et al [31] determined the prevalence of human cytomegalovirus (HCMV), Epstein-Barr virus (EBV) and herpes simplex virus (HSV) in gingival crevicular fluid (GCF) samples obtained from periodontally healthy, gingivitis and periodontitis patients. In addition, the effect of periodontal treatment (scaling and root planing) on the persistence of herpetic viruses was evaluated in a sub-group of patients suffering from chronic periodontitis. They observed that there was a higher prevalence of HCMV was associated with deep periodontal pockets in subjects suffering from periodontitis. This said study showed that the prevalence of HCMV and HSV viruses in GCF is higher in patients suffering from periodontitis compared to periodontally healthy subjects, and that the prevalence of HCMV is higher in deep periodontal pockets. It also brought evidences that periodontal therapy may be associated with virus elimination in diseased sites.

Laser therapy has been proposed as an alternative or adjunctive treatment to conventional periodontal therapy. A variety of lasers have been proposed, including the CO₂; neodymium doped:yttrium, aluminum, and garnet (Nd:YAG); diode; and erbium-doped: YAG (Er:YAG) lasers.

The bactericidal and detoxifying effects of the diode laser during non-surgical periodontal treatment have been documented [32], [33]. For patients with periodontitis, the adjunctive use of a diode laser with scaling and root planing (SRP) has shown a superior effect over SRP or the laser alone for certain microbial and clinical parameters [34].

Diode lasers (810-980 nm range) emit laser light in the near infra-red spectrum of the electromagnetic radiation which are highly absorbed in haemoglobin and other pigments. One of the main benefits of using diode lasers is the ability to selectively and precisely interact with diseased tissues. Lasers also allow the clinician to reduce the amount of bacteria and other pathogens in the surgical field and in the case of soft-tissue procedures, achieve good haemostasis without sutures [35], [17].

The purported advantages of diode lasers versus conventional surgery include increased coagulation that yields a dry surgical field and better visualization; the ability to negotiate curvatures and folds within tissue contours; tissue surface sterilization and therefore, reduction in bacteraemia; decreased swelling, edema, and scarring; decreased pain; faster healing response; and increased patient acceptance [17].

When laser cutting is in progress, small blood and lymphatic vessels are sealed due to the generated heat, thereby reducing or eliminating bleeding and edema. Denatured proteins within tissue and plasma are the source of the layer termed "coagulum", which is formed because of laser action and serves to protect the wound from bacterial or frictional action. Also the diode laser did not produce any deleterious effect on the root surface. Therefore, diode laser surgery can be performed safely in close proximity to dental hard tissue. During SRP procedure, there was no bleeding and postoperatively no pain was experienced by the patient and no swelling or any other signs of infection were noticed, whereas conventional procedures require the administration of antibiotics and analgesics to minimize postoperative

infection and pain [35].

In the present study a significant reduction of mean relative quantification (RQ) values of EBV in relation to 16sRNA in GCF samples of chronic generalized periodontitis patients was observed. The mean RQ values of EBV at baseline (28.53) was reduced to 0.19 on 21st day in group 1 and in group 2 it was reduced to 0.05 from 21.86.

VIII. CONCLUSION

It was found that the effect of diode laser as an adjunct to Scaling and root planing (SRP) showed greater reduction of EBV levels concentration in chronic generalized periodontitis patients compared to SRP and we also observed marked improvement of clinical parameters (GI, PPD and CAL) from baseline to 7th and 21st day.

The higher prevalence of EBV in the GCF of chronic generalised periodontitis patients suggest that EBV may play an important role in the pathogenesis of chronic periodontitis.

Within the limitations, we suggest that diode laser provides improvement in clinical and biochemical benefits over conventional mechanical treatment during medium term of observation.

IX. ACKNOWLEDGMENT

No commercial relationship or conflict of interest with any of the products used in this investigation, and the study is designed by our own initiative. Our study was self-funded by the authors and the institution.

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