



Detection of Human Cytomegalovirus (Hcmv) Epstein Barr Virus (Ebv) and Herpes Simplex Virus (Hsv) in Periodontal Disease and Effect of Scaling and Root Planing (Srp) on the Presence of These Viruses.

KEYWORDS

human cytomegalovirus, EBV, herpes virus 1, polymerase chain reaction, scaling and root planing, periodontal disease.

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ABSTRACT Aim: To evaluate the presence of human cytomegalovirus (HCMV), Epstein-Barr virus (EBV) and herpes simplex virus type 1 (HSV-1) in subjects with chronic periodontitis and periodontally healthy subjects and effect of scaling and root planing (SRP) on the presence of these viruses.

Materials and methods: At baseline in the test group, subgingival plaque samples were collected from the periodontal pocket and in the control group, plaque samples were collected from the gingival sulcus. In the test group after the baseline plaque sampling was done, scaling and root planing was performed. Plaque was collected after 8 weeks of the SRP again. Hot STAURT multiplex PCR method was used to detect the viruses in the plaque sample.

Results: Overall the presence of HSV-1 in the test group was statistically high at base line compared to the control group. There was reduction in HSV-1, HCMV and EBV after SRP in the test group compared to the baseline.

Conclusion: Herpes viruses were found in higher frequency in chronic periodontitis patients than in healthy subjects. SRP reduced the percentage of these viruses after 8 weeks.

Introduction: Periodontal disease is a microbial infection involving a variety of microbes that trigger inflammation, loss of connective tissue attachment and alveolar bone around the teeth. The primary etiologic factor of periodontitis is bacterial plaque. In fact the propensity of periodontitis to proceed with periods of exacerbation and remission could suggest that the presence of other organisms contributes to the disease.¹ Various studies have shown that human viruses, especially human cytomegalovirus (HCMV), Epstein-Barr virus (EBV) and herpes simplex virus (HSV-1) seem to play a part in the pathogenesis of periodontal disease.^{2,3,4} The hypothesis of a correlation between HCMV and EBV infection and the pathogenesis and progression of aggressive periodontitis has been proposed by various studies.^{5,6,7} Periodontal destruction may be associated with the coexistence of periodontal herpes viruses, especially HCMV, EBV and periodontopathic bacteria. The herpes viral 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.⁸ Till date several techniques have been employed for the detection of the viruses, which include culture methods, DNA-DNA hybridization technique and various types of polymerase chain reactions (PCR) such as Hot STAURT PCR, multiplex PCR, nested PCR, reverse transcriptase and real time PCR.^{9, 10, 11, 12}

PCR is a rapid, accurate and sensitive technique for the detection of bacterial and viral DNA sequences. Scaling and root planing resulted in reduction of periodontal pathogens^{13,14,15,16} The aim of this study was to examine the presence of human cytomegalovirus (HCMV), Epstein-Barr virus (EBV) and herpes simplex virus (HSV) in subjects with healthy periodontium and in subjects having chronic periodontitis and to examine the effect of SRP on these viruses

in chronic periodontitis subjects.

Methods: It is a pilot study where in subjects with severe chronic periodontitis having pocket depth ≥ 7 mm with good general health, no history of periodontal therapy in past 6 months were included in the study. Smokers, pregnant women and lactating mothers, subjects not compliant with the terms of the study, subjects who were on antimicrobial drugs in the previous six months were excluded from the study. Informed consent was taken from all the subjects and the approval from the ethical committee was taken. Subjects having healthy periodontium and no signs of gingival inflammation and no probing pocket depth were sixteen. Ten subjects having chronic periodontitis were selected. There were 16 periodontal pockets measuring ≥ 7 mm in these subjects. Two groups were made for the study having 16 sites each. All clinical parameters were assessed in test group at baseline and 8 weeks after SRP. The clinical parameters which were measured by UNC 15 probe were, Probing pocket depth (PPD), Relative attachment level¹⁷ Plaque index,¹⁸ and Gingival index.¹⁸

The viruses HCMV, EBV and HSV-1 were detected by Hot STAURT multiplex PCR*. In the test group, supra gingival plaque was gently removed with sterile cotton pellets and the sample site was isolated with cotton rolls to avoid contamination with the saliva and blood. Sterile periodontal curette was inserted into the bottom of the pocket and the plaque was removed by single stroke and collected into the vial containing phosphate buffered saline, in the same way plaque sample was collected from the gingival sulcus in the control group.⁶

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The primers[®] which were used in the study are as follows¹⁹

HSV-1 5'- F; 5'- CGTACTGCGGCTCGTGAA

R; 5'-AGCAGGGTGCTCGTGTATGGGC

HCMV- F; 5'-ACGTGTTACTGGCGGAGTCG-3'

R; 5'-TTGAGTTGTGGCCAGACTGAG-3'

EBV-F; 5'-AGCACTGGCCAGCTCATATC-3'

R; 5'TTGACGTCATGCCAAGCAA-3' 38

DNA extraction procedure from plaque sample: Plaque was transferred to the tube containing "TRIS-EDTA (T.E) buffer". It was centrifuged at 50,000 rpm for 2 minutes, supernatant was discarded. A fresh 200 micro liter T.E. buffer was added and then centrifuged for 3-4 minutes. Above procedure was repeated 3-4 times with fresh T.E. buffer, supernatant was discarded and 500 micro liter lysis buffer 1 was added and centrifuged at 5,000 rpm, supernatant was discarded and 50 micro liter lysis buffer 2 and 5 micro liter proteinase – K was added. It was kept in water bath over night then kept in boiling water bath for 10 minutes at 90°C. DNA was stored at -20°C. The tube containing DNA was taken and allowed to thaw. Then the master mix was prepared.

The contents of the master mix; For one reaction; Water – 14.25µL, dNTP mix - 2 µL, Enzymes – 0.25 µL, Primer- 0.5 µL. To a fresh tube, 22 µL of master mix was added and 3 µL of the DNA sample was added and it was inserted into the tube into the PCR cyclor.

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Statistical evaluation: The collected data was entered in the excel format. It was subjected to statistical analysis using SPSS software. The statistical tests applied were as below

1. Fisher exact test
2. Mann Whitney U test
3. Kruskal-Wallis test

For all statistical tests $p < 0.05$ was considered statistically significant.

Results: Decrease in plaque index and gingival index scores from baseline to 8 weeks in the test group which was statistically significant. There was a significant decrease in the probing pocket depth and gain in the attachment levels from baseline to 8 weeks in the test group.

Overall the presence of HSV-1 in the test group was significantly high compared to the control group at baseline. **(Table 1 & Graph1)** Presence of HCMV & EBV was high in the test group compared to the control group at baseline which was not statistically significant **(Table 1& Graph1)** there was reduction in HSV-1, HCMV and EBV after SRP in the test group compared to the baseline. Only HSV-1 reduction was statistically significant. **(Table 2 Graph 2.)**

Discussion: Viruses were found in patients with aggressive periodontitis and chronic periodontitis.¹ Viral DNA had been detected in gingival tissue, gingival crevicular

fluid (GCF) and subgingival plaque from periodontally diseased sites.²⁰ In particular, DNA from herpesviruses such as HCMV and EBV had been detected in high percentages of subgingival plaque samples from periodontitis patients. Periodontally healthy individuals had very low prevalence of such viruses.^{21, 22}

In this study the viruses were isolated from subgingival plaque sample as all the microorganisms are colonized in the subgingival plaque. It is the simplest method of collection and less time consuming. In the test group, plaque sample was taken at baseline and 8 weeks after SRP as there will be recolonization and reestablishment of subgingival micro biota within 8 weeks after SRP.²³ The subgingival plaque was collected from pockets having ≥ 7 mm as the HCMV, EBV and HSV were detected with higher frequency in deep than in shallow periodontal pockets.²⁴ Full mouth plaque index and gingival index were taken to ascertain the patient compliance towards periodontal therapy. All patients had shown improvements in the indices scores after SRP compared to baseline which could be attributed to constant reinforcement of oral hygiene instruction and Hawthorne effect as given by Knowles et al., in 1979 and Ramfjord et al., 1987.²⁵ There was significant reduction in the probing depth and gain in relative attachment level which was in accordance to the study done by Shibata Y 1989 where in there was decrease in the probing depth and gain in attachment level after SRP.^{26,27} Patients with chronic periodontitis showed higher percentage of HSV-1 (81.25 %), HCMV (50%), and EBV (37.50 %) than patient having healthy periodontium who had HSV-1 (31.25 %), HCMV (31.25%), and EBV (25 %). The lesser percentage of viruses in healthy sites may be because of the presence of virus in latent form.²⁰ The frequency of viruses in periodontal disease increases as these viruses may get reactivated. Contreras et al., 1999 studied the presence of herpes viruses in adult periodontitis patients. He showed the higher percentage of HCMV than EBV and HSV by using nested PCR method.³ Li Jane Ling et al., found the higher prevalence of HCMV than HSV and EBV-1 which was determined by nested PCR method.²⁴ Higher percentage of HCMV was found by Kamma JJ et al., when compared with EBV-1 and HSV.²⁸ Our study showed higher percentage of HSV-1 than HCMV and EBV, the variation could be due to demographic and geographic difference.

EBV and HCMV were rarely found in chronic periodontitis patients by Dolphus R et al., by using real time PCR.¹ Similar results were found by Nibali L et al., in plaque samples of chronic periodontitis, generalized aggressive periodontitis and localized aggressive periodontitis.²⁹ Our study showed similar results as the study done by Imbronito et al., who found higher percentage of HSV-1 than HCMV in subjects with generalized aggressive periodontitis, chronic periodontitis and gingivitis by nested PCR method.³⁰ These studies suggest that Herpes viruses were found in higher frequency in chronic periodontitis patients than in healthy subjects.

The variation in the results of the above mentioned studies and our study might be due to the differences in the methodology, method of collection of samples, the type of PCR used and the periodontal disease activity at the time of sample collection. Nested PCR which was used in most of the previous studies is very sensitive and susceptible to contamination and may give rise to false positive results (Burkard, 2000).²⁰ The multiplex Hot STAURT PCR which was used in our study is a qualitative PCR method with a detection range of 50-100 subgingival viruses. The

main advantage of multiplex PCR is that large scale amplification of viral DNA can be done with minimum cost and time (Shin C et al., 2003).³¹

The mixed herpes viral infections might promote subgingival colonization of pathogenic organisms by a multiplicity of mechanisms. A gingival herpes viral infection may impair the local host defense by infecting and altering the function of polymorphonuclear leukocytes,^{32, 33} lymphocytes^{34, 35} and macrophages.^{36, 37} A virally induced neutrophil dysfunction may serve to potentiate the overgrowth and virulence of *P. gingivalis* and other members of the subgingival microbiota.⁴ In addition the herpesvirus infection can lyse or affect oral epithelial cells. The disruption of the protective epithelial barriers of the periodontium may facilitate access of periodontopathic bacteria to deeper tissues of the periodontium and create additional sites for bacterial binding.^{38, 39} The frequency of viruses was higher in active than in inactive sites (Ling L et al.).¹³ Virus-infected periodontal sites tend to exhibit more breakdown than herpes virus free sites and a herpes viral active infection was associated with an elevated risk of progressive periodontal disease.¹⁴

After SRP, there was reduction in the HSV-1, HCMV and EBV score compared to the baseline score in chronic periodontitis group which was significantly higher in the HSV-1. This indicates that SRP was effective in reducing the amount of viral load. The above results are in accordance to the study done by Grenier et al 2009 who found reduction in the scores of HCMV, EBV and HSV after SRP,²⁶ this suggests that periodontal therapy was effective in reducing the viral load.

Our study showed the presence of viruses in healthy periodontium and chronic periodontitis. The percentage of viruses was more in subjects with chronic periodontitis. SRP was effective to reduce the viral load after 8 weeks in chronic periodontitis subjects. Very few studies are done where in nonsurgical periodontal therapy is used to reduce the viral load, and viruses were detected by Hot STAURT PCR, where in most of the studies viruses were detected by nested PCR.

Future studies are required with more sample size to identify the oral sites of herpes viral latency and characterization of viral gene expression during latency, to determine whether herpes viral replication, host immune response or direct effect of herpes viruses on immune cells act as determinants of periodontal disease.

Acknowledgements: I would like to thank Dr. Kishore Bhat (Microbiologist) for helping me in my study.

TABLES

Table 1: Comparison of HSV-1, HCMV & EBV-1 in the test and the control group at baseline

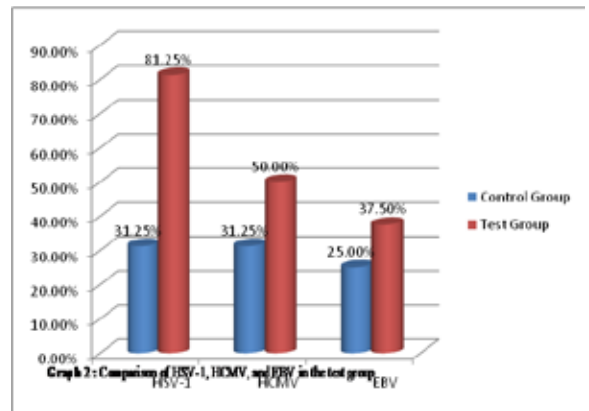
Viruses	Group	Negative	%	Positive	%	Total
HSV-1	Test	3	18.75	13	81.25	16
	Control	11	68.75	5	31.25	16
	Total	14	43.75	18	56.25	32
	Fisher exact test	p=0.00567				
	Z-value	-2.4121				
	p-value	0.0159				

HCMV	Test	8	50.00	8	50.00	16
	Control	11	68.75	5	31.25	16
	Total	19	59.38	13	40.63	32
	Fisher exact test	p=.23				
	Z-value	-0.90				
EBV-1	Test	10	62.5	6	37.50	16
	Control	12	75.00	4	25.00	16
	Total	22	68.75	10	31.25	32
	Fisher exact test	p=.35				
	Z-value	-0.60				
	p-value	0.54				

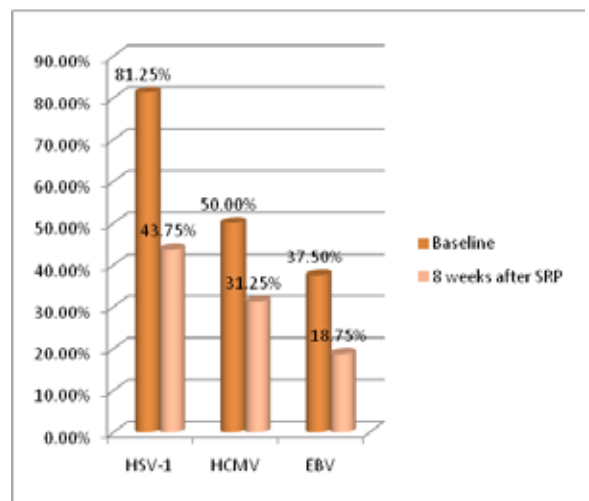
Table 2: Comparison of HSV-1, HCMV, and EBV in the test group.

Viruses	Time	Mean	SD	T-value	Z-value	p-level	Signi
HSV-1	BL-8weeks	0.81-0.44	0.40-0.51	0.00	2.20	0.02	S
HCMV	BL-8weeks	0.50-0.31	0.52-0.48	3.00	1.21	0.22	NS
EBV	BL-8weeks	0.38-0.19	0.50-0.40	0.00	1.60	0.10	NS

Graph 1: Comparison of the HSV-1, HCMV, and EBV in the test and control at baseline.



Graph 2 : Comparison of HSV-1, HCMV, and EBV in the test group



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