Stimulated and Non-Stimulated Salivary Flows Should Be Tested for the Presence of HCV RNA in Saliva Samples from Patients with Chronic Hepatitis C

Grossmann SMC1,4*, De Oliveira GC2, Teixeira R3 and Vieira do Carmo MA1

1Department of Oral Surgery and Pathology, School of Dentistry, Universidade Federal de Minas Gerais, Belo Horizonte, Brazil
2René Rachou Fiocruz Research Centre, Belo Horizonte, Minas Gerais, Brazil
3Department of Medical Clinic, School of Medicine, Universidade Federal de Minas Gerais, Belo Horizonte, Brazil
4Department of Oral Diagnosis, School of Dentistry, Universidade Vale do Rio Verde, Belo Horizonte, Brazil

Corresponding author: Soraya de Mattos Camargo Grossmann, Universidade Vale do Rio Verde, Faculdade de Odontologia, Rua Gentios, 1420 – Luxemburgo, Belo Horizonte - CEP: 37410-000, Minas Gerais, Brazil, Tel: + 55-31-33441366; E-mail: sogrossmann@uol.com.br

Rec date: Feb 26, 2014, Acc date: Apr 18, 2014, Pub date: Apr 20, 2014

Copyright: © 2014 Grossmann SMC, et al. This is an open-access article distributed under the terms of the Creative Commons Attribution License, which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited.

Abstract

Objective: In most of the studies which analyzed the presence of HCV RNA in saliva from patients with chronic Hepatitis C only stimulated saliva samples have been used for viral detection. Thus, this study compared the prevalence of HCV RNA in non-stimulated and stimulated salivary flows in patients with chronic Hepatitis C.

Design: Saliva samples of non-stimulated and stimulated salivary flows from 24 patients were collected, and the HCV RNA was investigated by RT-nested PCR. Data regarding age, gender, risk factors for HCV infection, xerostomia and hyposalivation were also analyzed.

Results: The HCV RNA could be detected in 11 (45.8%) non-stimulated and in 14 (58.3%) stimulated saliva samples, without statistical significance (p=0.472). However, in 18 (75.0%) patients it was possible to detect the presence of the HCV RNA at least in one of the saliva samples. Six (25.0%) patients complained of xerostomia and nine (37.5%) presented hyposalivation, but in only 3 (12.5%) patients, these conditions could be observed, simultaneously. No significant correlation between the presence of HCV RNA in saliva and age, gender, risk factors for HCV infection, xerostomia and hyposalivation could be identified.

Conclusion: Both stimulated and non-stimulated saliva samples must be investigated for the presence of HCV RNA in patients with chronic Hepatitis C, to avoid underestimated prevalence of HCV in this group of patients.

Keywords: HCV RNA; Saliva; Chronic Hepatitis C; HCV

Introduction

Hepatitis C Virus (HCV) infection is a major cause of chronic liver disease and liver-related morbidity and mortality worldwide and it is the most prevalent reason for liver transplants in Europe and in the USA [1].

Up to 74% of the HCV-infected patients may develop at least one Extra Hepatic Manifestation (EHM) during the course of the infection, such as salivary gland disorders, being the HCV considered as a sialotrophic virus [2-6]. The reported prevalence of the HCV RNA in saliva ranges from 0 to 100% [7,8]. In a recent study conducted with Brazilian patients, Grossmann et al. [9] detected the HCV RNA in 39.0% of the non-stimulated saliva samples in patients with serum positive HCV RNA. These discrepancies may be due to the heterogeneity of the study design, population studied as well as different methodologies used as saliva sampling methods.

Xerostomia in patients with chronic Hepatitis C ranges from 10 to 20% [2,10] and its association with an objective evidence of salivary gland dysfunction like hyposalivation has been described in previous literature [11,12]. It has been suggested that whole saliva could be used as a screening test for identifying a suspected exocrine hypofunction and an abnormality in the salivary gland, such as the Sjögren Syndrome [13].

As the prevalence of HCV in saliva samples shows important differences among the studies, and most of them analyze only stimulated salivary flows [11,14], the aim of this study was to compare the prevalence of HCV RNA in saliva samples between non stimulated and stimulated salivary flows in a group of patients with chronic Hepatitis C. The possible association with xerostomia and presence of hyposalivation were also evaluated.

Patients and Methods

Patients

A cross-sectional survey was carried out on 24 HCV patients with confirmed diagnosis of chronic Hepatitis C (anti-HCV positive, Elisa III, HCV RNA qualitative test positive) admitted to the Viral Hepatitis Reference Center of the Alfa Institute of Gastroenterology at the Clinical Hospital of Universidade Federal de Minas Gerais (AHEV/ IAG-HC/UFMG), Belo Horizonte, Brazil. The study was approved by the UFMG Ethical Committee for Surveys, and all volunteers signed an informed consent form. All patients were HIV (human immunodeficiency virus) and HBV (Hepatitis B virus) negative and had no other concomitant liver diseases. Patients receiving antiviral
treatment for Hepatitis C were also excluded from the study. Demographical data as well as risk factors for the acquisition of the infection were assessed from medical records. Intra-oral examination was performed on all patients.

Xerostomia

Patients were inquired in relation to symptom of dry mouth (xerostomia). If present, xerostomia was classified as mild, moderate, or severe, as previously described [4] and the patients received treatment with artificial saliva gel (Oral Balance- Biotene) to relieve their symptom.

Saliva samples

Patients refrained from oral hygiene, eating, drinking, and smoking 60 min before non-stimulated (NSSF) and stimulated (SSF) salivary collection in sterile 50 mL Falcon tubes. The entire NSSF produced during three minutes was collected. Rates ≤ 0.1 ml/min was considered hyposalivation, as previously established [15]. Subsequently, patients were asked to chew a sterile rubber cylinder for three minutes before spitting saliva to measure SSF. Rates of ≤ 0.7 ml/min were considered hyposalivation, as previously established [15]. All saliva samples were immediately stored at – 80°C.

Detection of HCV RNA in the saliva

HCV RNA was extracted from 200 μl of undiluted saliva using a commercial viral RNA isolation kit (Qiamp, Qiagen, Hildem Germany), according to manufacturer’s instructions. Nested reverse transcription-polymerase chain reaction (RT-PCR) was performed, by amplification of a 251 bp fragment from the 5’ UTR region of HCV with a minor adaptation described by Oliveira et al. [16]. Briefly, a single-strand cDNA was synthesized from 18 μL of the RNA sample at 42°C for 50 minutes with 200 U of reverse transcriptase (Superscript II, Gibco, BRL, Rockville, MD, USA) in 30 μL of a manufacturer supplied buffer containing 10 pmole of primer 209 (ATACTCGAGGTGCACGGGTCTACGAGACCT), 10 mM of each dNTP, and 10 nM of dithiothreitol. For the first-round of PCR, 2 μL of cDNA were added to a mixture containing 2 μl of 10x of the supplied PCR buffer, 5 mM of MgCl2, 10 mM of each dNTP, 10 pmole of primers 939 (CTGTGAGGAACTACTGTCTT) and 211 (TTCACGCAGAAAGGGTCTAG) and 211 single-strand cDNA was synthesized from 18 μL of the RNA sample at 42°C for 50 minutes with 200 U of reverse transcriptase (Superscript II, Gibco, BRL). The mixture was cycled 30x in a thermocycler (Perkin Elmer Gene Amp PCR System 2400) at 95°C for 5 minutes, 95°C for 1 minute, 50°C for 1 minute, 72°C for 1 minute, and again at 72°C for 1 minute. The second-round of PCR (nested) was carried out as above, using primers 940 (TTCACGCGAAGGGTCTAG) and 211 (CAGCTTCCGAGACCCTATCAACGGCAGT) as well as 1.5U of Taq DNA polymerase (GIBCO/ BRL). The PCR products were electrophoresed in a 6% polyacrylamide gel, using a vertical electrophoresis apparatus (Model S2 Life Technologies Inc.) and stained with silver nitrate. The banding pattern was visualized and analyzed as described by McOmish [17]. A known HCV RNA positive saliva sample was used as a positive control and, to a negative control, viral RNA was omitted.

Statistical analysis

Chi-square and Fisher’s tests were used for univariate analysis. p value ≤ 0.05 was considered significant.

Results

The group studied consisted of 13 (54.2%) females and 11 (45.8%) males, with a mean age of 52.4 years and median of 52 years (range 29-73 years of age). Main sources of infection included blood transfusions in 10 (41.7%) cases and intravenous drug use in 4 (16.7%) cases. Risk factors could not be identified in 4 (16.7%) patients (Table 1).

<table>
<thead>
<tr>
<th>Risk Factors</th>
<th>n (%)</th>
<th>Positive n (%)</th>
<th>Negative n (%)</th>
<th>p</th>
</tr>
</thead>
<tbody>
<tr>
<td>Blood transfusion</td>
<td>11</td>
<td>5 (35.7)</td>
<td>5 (50.0)</td>
<td>0.241</td>
</tr>
<tr>
<td>i.v. drug use</td>
<td>11</td>
<td>3 (27.3)</td>
<td>8 (72.7)</td>
<td>0.472</td>
</tr>
<tr>
<td>Percutaneous exposure</td>
<td>11</td>
<td>1 (9.1)</td>
<td>10 (90.9)</td>
<td>0.076</td>
</tr>
<tr>
<td>Others</td>
<td>11</td>
<td>0 (0.0)</td>
<td>11 (100.0)</td>
<td>0.004</td>
</tr>
<tr>
<td>Unknown</td>
<td>11</td>
<td>2 (18.2)</td>
<td>9 (81.8)</td>
<td>0.115</td>
</tr>
</tbody>
</table>

Fisher’s exact test

Legend: i.v. – intravenous; n – absolute number; (%) – relative number

Table 1: Prevalence of HCV RNA in saliva, according to gender and risk factors

Treatment for Hepatitis C were also excluded from the study. Demographical data as well as risk factors for the acquisition of the infection were assessed from medical records. Intra-oral examination was performed on all patients.

<table>
<thead>
<tr>
<th>HCV RNA in stimulated salivary flow</th>
<th>Positive n (%)</th>
<th>Negative n (%)</th>
<th>Total n</th>
<th>p</th>
</tr>
</thead>
<tbody>
<tr>
<td>Positive</td>
<td>7 (63.6)</td>
<td>4 (36.4)</td>
<td>11</td>
<td>0.472</td>
</tr>
<tr>
<td>Negative</td>
<td>3 (27.3)</td>
<td>10 (72.7)</td>
<td>13</td>
<td></td>
</tr>
<tr>
<td>Fisher’s exact test</td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
</tbody>
</table>

Legend: n – absolute number; (%) – relative number

HCV RNA was detected in non-stimulated saliva samples of 11 (45.8%) patients (4 women and 7 men) and in stimulated saliva samples of 14 (58.3%) patients (6 women and 8 men), with no...
statistical difference (p=0.472) (Table 2). In only 7 (29.2%) patients the HCV RNA could be detected in both, NSSF and SSF, simultaneously. However, in 18 (75.0%) patients it was possible to detect the presence of the HCV RNA at least in one of the saliva samples.

<table>
<thead>
<tr>
<th>Xerostomia</th>
<th>Non stimulated salivary flow</th>
<th>Stimulated salivary flow</th>
<th>p</th>
<th>p</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Normal</td>
<td>Hyposalivation</td>
<td>Normal</td>
<td>Hyposalivation</td>
</tr>
<tr>
<td>Present</td>
<td>5</td>
<td>1 (11.1)</td>
<td>4</td>
<td>2 (25.0)</td>
</tr>
<tr>
<td>Absent</td>
<td>10</td>
<td>8 (88.9)</td>
<td>12</td>
<td>6 (75.0)</td>
</tr>
<tr>
<td>Total n (%)</td>
<td>15</td>
<td>9 (100.0)</td>
<td>18</td>
<td>8 (100.0)</td>
</tr>
</tbody>
</table>

Fisher’s exact test
Legend: n – absolute number; (%) – relative number

Table 3: Correlation between the prevalence of xerostomia and hyposalivation and salivary flows

Six (25.0%) patients complained of oral dryness (xerostomia) which was mild in 4 (16.7%) patients, and moderate in 2 (8.3%) cases. Considering the entire sample, hyposalivation could be observed in 9 (37.5%) cases of non-stimulated salivary flow and in 8 (33.3%) cases of stimulated salivary flow. However, in only 3 (12.5%) cases the xerostomia and hyposalivation could be observed, simultaneously (Table 3).

Among the patients with xerostomia and hyposalivation, in both different flows, the HCV could be detected more frequently in saliva sample from non-stimulated salivary flow (6 cases) (Table 4).

Oral mucosal alterations, including variations of normality and mucosal lesions were observed in 20 (83.3%) patients. The most common alterations were traumatic lesions in 8 (33.4%) cases, followed by Fordyce’s spots in 6 (25.0%) cases, and candidiasis in 4 (16.7%) cases.

Discussion
As Hepatitis C is often asymptomatic or shows no specific manifestation in the acute phase, the WHO estimates that there are millions of undiagnosed HCV-infected people, constituting an important link in the chain of HCV transmission [18]. Thus, the recognition of extra Hepatic manifestations of this infection is considered of great importance regarding the establishment of an early diagnosis.

Table 4: Prevalence of HCV RNA in relation to the presence of xerostomia and hyposalivation in stimulated and non-stimulated salivary flows

As expected, the main source of transmission of HCV in the present study was blood transfusion (41.7%), which is in accordance with that reported in the literature [1]. Besides, in 4 patients (16.7%), the source of infection was unknown, in accordance with our previous study [9,19,20].

Considering the reported sialotropism of HCV, with different findings, and taking into account that most of the studies used only stimulated salivary flow, the present study aimed to compare the prevalence of HCV RNA in saliva samples from non-stimulated and stimulated salivary flows. Our results showed the presence of HCV RNA in non-stimulated and stimulated saliva samples in 11 patients (45.8%) and in 14 patients (58.3%), respectively, with no statistically significant difference (p=0.472). These findings suggest that both non-stimulated and stimulated saliva samples are equally suitable for the investigation of the presence of salivary HCV RNA. In only 7 (29.2%)
patients the HCV RNA could be detected in both, NSSF and SSF, simultaneously. However, a very higher prevalence could be observed in 18 (75.0%) patients in whom it was possible to detect the presence of the HCV RNA at least in one of the saliva samples. We consider this as an important finding that points toward the necessity of investigation the presence of HCV RNA in both salivary flows, taking into account that the mechanism of viral shedding is not well-understood. It is also necessary to emphasize that a more representative sample to determine the real prevalence of HCV RNA in saliva is warranted.

Some authors reported that there is no correlation between the presence of anti-HCV antibodies in saliva and the detection of HCV RNA in saliva and salivary glands in patients with chronic Hepatitis C [21]. However, others suggested that the salivary HCV viral load is significantly lower than the viral load in the serum16, and this could explain the controversial results that investigate the HCV in saliva in patients with chronic Hepatitis C.

No significant correlations were found between the presence of HCV RNA in non-stimulated or stimulated saliva samples and age, gender, risk factors, xerostomia, or hyposativation in accordance with previous study [9]. We consider that the xerostomia and hyposativation in Hepatitis C patients is not association with direct detection of HCV in saliva. We results suggest that an indirect effect as more inflammatory cells in salivary gland (PATRICIA) could be responsible for xerostomia and hyposativation in this group of patients.

In the present study, six (25.0%) patients complained of oral dryness (xerostomia) and hyposativation could be identified in 9 (37.5%) patients, with no statistical significance. Dry mouth (xerostomia) has been reported in 0 to 35% patients with chronic Hepatitis C [2,5,11,14]. Some authors have the HCV as a virus with a potential association between oral lichen planus and chronic Hepatitis C [20]. Due to the high global prevalence of chronic Hepatitis C, and the necessity of viral shedding is not well-understood. It is also necessary to emphasize that a more representative sample to determine the real prevalence of HCV RNA in saliva is warranted.

Further research to elucidate this point is needed.

Oral mucosal alterations were observed in almost all evaluated patients with chronic Hepatitis C (83.3%) as in our previous study [20]. Due to the high global prevalence of chronic Hepatitis C, and the potential association between oral lichen planus and chronic Hepatitis C [19], we consider important periodic oral examination in order to promote the suitable and full health assistance to these patients.

Although in stimulated and non-stimulated salivary flows it is possible the detection of HCV RNA, and regardless the small sample, our findings strongly suggest that the investigation should be performed in both samples, to avoid bias regarding the real prevalence of this virus in saliva of patients with chronic Hepatitis C. Moreover, the detection of HCV RNA in saliva cannot determine definitively a direct effect of this virus in presence of xerostomia and hyposativation in Hepatitis C patients. Thus, investigations that try elucidating the possible sialotropism of HCV are necessary.

Acknowledgements

This study was supported by grants from CAPES (Coordenação de Aperfeiçoamento de Pessoal de Nível Superior), FAPEMIG (Fundação de Amparo à Pesquisa do Estado de Minas Gerais) and CNPq (Conselho Nacional de Desenvolvimento Científico e Tecnológico - National Counsel of Technological and Scientific Development). Dra. Teixeira R and Dr. Oliveira GO are research fellows of CNPq.

References