Molecular Alterations of Parotid Saliva in Infantile Chronic Recurrent Parotitis

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ABSTRACT: Infantile chronic recurrent parotitis (ICRP) is an insidious disease whose etiopathogenesis remains an enigma. Alterations in the physical appearance of parotid saliva from ICRP patients have been frequently reported. However, sialochemical studies in regard to ICRP are very rare. The aim of this study was to determine whether saliva of ICRP patients presents major physicochemical and biochemical alterations compared with saliva from paired healthy controls. Parotid, whole, and submandibular/sublingual saliva was collected at an asymptomatic stage from 33 ICRP patients (5-16 y old, both sexes) and from 33 sex- and age-matched healthy controls. Saliva was analyzed for protein concentration, mode of protein diffusion on cellulose membranes, unidimensional sodium dodecylsulfate (SDS)-polyacrylamide gel electrophoresis protein profiles and zymographic profiles of metalloproteinase 2 (MMP-2) and metalloproteinase 9 (MMP-9). Parotid saliva of ICRP patients showed an increased protein concentration, altered mode of protein diffusion, a higher frequency of polypeptide bands of 43, 37, 33, 29, 26, 16, and 10 kD, higher asymmetry in the polypeptide profiles of both contralateral parotid saliva, and an increase in the frequency of MMP-2 and MMP-9. Parotid saliva of patients with ICRP is molecularly altered with respect to normal saliva. Some of the molecular differences could be related to the etiopathogenesis of the disease. (Pediatr Res 61: 203-208, 2007)

ICRP is an inflammatory disease associated with nonobstructive glandular sialectasia. It is characterized by recurrent episodes of swelling of one or both parotid glands that alternate with remission intervals lasting for weeks or months in which glands are clinically asymptomatic (1,2). The onset of the disease usually occurs around age 3 y. Symptoms may diminish and disappear spontaneously after puberty. Sometimes the disease may persist until adulthood (1–3).

Among the postulated etiological factors of ICRP are congenital ductal malformations, genetic factors, allergies, bacterial infections, local manifestations of autoimmune disease, and immunodeficiencies (1,2,4–9). However, none of them

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has been demonstrated to play a causative role in the disease, and to date its etiopathogenesis remains an enigma.

Likewise, consensus on the therapeutic management of ICRP has yet to be achieved. Different specialists have opted for different therapies. Among the therapeutic indications described in the specialized literature is a diversity of physical maneuvers to stimulate the salivary flux (1) and pharmacotherapy with nonsteroid anti-inflammatory agents (7) in combination with sialogogues and/or antibiotics (3,6,10). Recently, Nahlieli *et al.* (11) proposed an interventional approach using a sialendoscope. In cases in which irreversible damage has occurred, ligation of the excretory duct is indicated to produce glandular atrophy (1,2). In severe cases, partial or total parotidectomy is prescribed as a last resource (1,2,12).

Studies on glandular tissue obtained from patients with ICRP have shown lymphocytic infiltrates and lymphoid follicles around dilated interlobular ducts (sialectasia) (1,13,14).

Most of the studies on ICRP describe qualitative changes in parotid saliva during the active period of the disease (1,2,4,7). Some authors describe the salivary fluid either as mucopurulent (1), as a milky viscous secretion with the accumulation of semisolid material (2), as purulent and filamentous (4), or as a turbid and purulent material (7). If those physical changes of parotid saliva from ICRP patients could be associated with changes in its molecular composition, it might be expected to use parotid saliva as an indicator of the functional status of the gland. In that direction, the use of saliva for the diagnosis of disorders that affect salivary glands has been documented in regard to Sjögren syndrome, in therapeutic irradiation of the maxillofacial territory in cancer patients, and in cystic fibrosis (15–18). However, in the specialized literature, just a few sialochemical studies are referred to ICRP.

In 1978, Tabak *et al.* (19) analyzed parotid saliva of ICRP patients in which salivary concentrations of albumin and lactoferrin were increased during the acute episode and asymptomatic stage. Then in 1996, Ericson and Sjoback (20) analyzed samples of parotid saliva from 17 children with ICRP and observed a significant increase in the concentrations of albumin, IgA, lactoferrin, and kallikrein in relation to saliva

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Abbreviations: ICRP, infantile chronic recurrent parotitis; MMP-2, metalloproteinase 2; MMP-9, metalloproteinase 9

from healthy controls. Both studies, however, are referred to a small group of proteins. Furthermore, to date, no sialochemical study approaches the analysis of proteins that could be related to the damage of salivary glands.

The aim of this study was to determine whether saliva of ICRP patients presents, in relation to saliva from paired healthy controls, major physicochemical and biochemical alterations. To this end, we assayed that saliva for modes of protein diffusion on cellulose membranes, total protein content, electrophoretic unidimensional polypeptide profiles, and occurrence of MMP-2 and MMP-9. We postulate that parotid saliva of ICRP patients displays major physicochemical and biochemical alterations.

METHODS

Subjects. Thirty-three ICRP patients under current surveillance at the Pediatric Maxillofacial Surgery Division of San Juan de Dios Hospital were included. Inclusion criteria were (a) at least two reported acute inflammatory episodes affecting the same parotid gland, (b) bilateral sialographic study, (c) absence of any systemic disease, and (d) normal appearance of gingiva, teeth, and oral mucosa. Patients were recruited at an asymptomatic stage of the condition and at various stages of the treatment. The control population consisted of 33 healthy volunteers under preventive dental care at the same hospital, paired by gender and age, who fulfilled the following criteria: (a) absence of any oral, glandular, or systemic disease and (b) no medication in the past 3 mo. The study was conducted according to the guidelines of the Ethics Committee of the Faculty of Dentistry, University of Chile. Informed consent was obtained from the parents at the beginning of the study.

Saliva collection. Saliva samples were collected by a single trained operator after 2% citric acid stimulation at about 0.900–11.00 under standardized conditions (21). Whole saliva was expectorated into two collection vessels. Whole saliva collected during the first 30 s was excluded from protein analysis. Parotid and submandibular/sublingual saliva samples were collected by using adapted devices (22). Flow rates were estimated by scoring total saliva volumes and the corresponding collection times. Samples were maintained in ice-cooled containers during the collection and then were stored at -20° C until their analysis.

Quantification of salivary proteins. Proteins were measured according to Bramhall *et al.* (23) with previously described modifications (24). Aliquots of saliva or bovine serum albumin $(10-50 \ \mu g)$ were spotted on cellulose disks (Whatman 1, diameter 2.5 cm), fixed in 5% trichloracetic acid, washed in 80% ethanol and 3:1 ethanol/ether, stained with Coomassie blue R-250, and washed in 7% acetic acid until clear background. The disks were photographed to keep records of the mode of diffusion (monophasic or biphasic) of the salivary protein (25). The blue dye of each disk was eluted in 45% methanol/0.25% ammonia and subjected to spectrophotometry at 610 nm.

Unidimensional electrophoresis. Aliquots of saliva (8.5 μ g protein) were fractionated in 10% SDS-polyacrylamide gels ($20 \times 20 \times 0.15$ cm³) according to Laemmli (26) and developed with Coomassie blue. Gels were calibrated by using broad range protein M_r standards. Identification of individual bands was based on the apparent M_r . Samples of contralateral parotid glands of single subjects were run in neighbor tracks. Polypeptide bandings of parotid saliva samples from ICRP patients were compared with those of healthy controls. Also, the degree of concordance of polypeptide profiles (symmetry/asymmetry) between both parotid saliva samples of each subject

Zymography. Aliquots of saliva (5 µg protein) were mixed with 60 mmol/L Tris-HCl pH 6.8, 5% SDS, 20% glycerol, and 0.03% bromophenol

blue and electrophoresed at constant 125 V according to Laemmli (26) in 10% SDS-polyacrylamide gels containing 1 mg/mL gelatin. After fractionation, gels were incubated twice at room temperature for 20 min each in 2.5% Triton X-100 and then incubated overnight at 37°C in activation buffer (10 mmol/L Tris-HCl (pH 8.0)/1.25% Triton X-100/5 mmol/L CaCl₂/1 μ mol/L ZnCl₂). Gels were stained in 0.25% Coomassie blue for 6 h and destained in 10% methanol/15% acetic acid until clear bands were resolved. Gels were calibrated using M_r standards. Nonmetalloproteinase proteases were discarded by incubating salivary samples in 50 mmol/L *N*-ethylmaleimide/50 mmol/L phenylmethylsulfonylfluoride Metalloproteinase activity was confirmed by incubating gels in activation buffer containing 5 mmol/L ethylenediamine tetraacetic acid, a metalloproteinase inhibitor. Further, metalloproteinase activities were contrasted against MMP-2 and MMP-9 standards.

Statistics. For age, flow rate, and protein concentration in saliva, averages and standard deviations were calculated. Statistical analyses were performed by a *t* test or the Wilcoxon Mann-Whitney test as appropriate. For the mode of protein diffusion, presence of electrophoretic bands, symmetry/asymmetry, and presence of zymographic bands, frequencies were determined. In these cases, statistical analyses were performed by a χ^2 or Fisher exact test as appropriate. Association between variables was assessed by logistic regression analysis. Stata Version 7.0 statistical software was used. Statistical significance was set at p < 0.05.

RESULTS

Patients. A total of 66 subjects, 33 ICRP patients (12 female, 21 male) with an average age of 10.3 ± 3.3 y (range, 5-16) and 33 age- and sex-matched controls were studied (Table 1). Fourteen patients were affected bilaterally, and 19 expressed either right (n = 6) or left (n = 13) clinical manifestations of the disease (Table 2). Sialographic findings but not gender or age were associated with the laterality of the disease [p < 0.001; odds ratio (OR) = 69.3] [95% confidence interval (CI) = 12.9-372.7]. Thus, bilateral parotid sialographic alterations were observed in 13 of 14 bilateral patients and ipsilateral alterations were detected in 18 of 19 unilaterally affected patients. The only exceptions were a bilateral patient displaying sialographic alterations only in the left parotid gland, two unilaterally affected patients displaying sialographic alterations also in the contralateral parotid gland, and a unilateral patient presenting normal sialographic images on both sides (Table 2).

Salivary samples. Whole saliva, submandibular/sublingual saliva, and both samples of parotid saliva from ICRP patients and healthy controls were pus free and blood free and displayed a translucid appearance. In the group of ICRP patients, 16 samples of parotid saliva had a whitish particulate material and eight had a viscous appearance. Mean value of flow rates of parotid saliva among the ICRP patients (0.136 mL/min/gland) was significantly lower (p < 0.001) than the one corresponding to the control group (0.384 mL/min/gland). Protein concentration in parotid saliva of ICRP patients showed statistically significant increases compared with healthy controls (Table 1).

Table 1. Age and sex distribution and protein concentration of parotid saliva among ICRP patients and controls

	S	ex		Protein (mg/mL	Δ),* mean \pm SD
Group	F	М	Age (y), mean \pm SD (range)	Right parotid	Left parotid
Control $(n = 33)$	12	21	$9.4 \pm 2.4(4-14)$	0.554 ± 0.186	0.552 ± 0.204
ICRP $(n = 33)$	12	21	$10.3 \pm 3.3(5-16)$	$0.659 \pm 0.292 \ddagger$	$0.693 \pm 0.304 \dagger \$$

* According to the method of Bramhall et al. (22) and using bovine seroalbumin as standard.

† Statistical significances of ICRP versus the corresponding control, Wilcoxon Mann-Whitney test.

 $\ddagger p = 0.034; \$ p = 0.010.$

 Table 2. Distribution of ICRP patients according to laterality of the disease and sialographic findings

	Right parot	id sialography	Left parotid sialography		
Laterality*	Normal	Sialectasis	Normal	Sialectasis	
Right $(n = 6)$	0	6	5	1	
Left $(n = 13)$	12	1	1	12	
Bilateral $(n = 14)$	1	13	0	14	

* Symptoms affecting either (left or right) or both parotid glands.



Figure 1. Modes of protein diffusion of salivary protein on cellulose paper membranes. Aliquots of parotid saliva (A) or submandibular/sublingual saliva (B) were placed on the center of a disk, fixed, and stained. Monophasic (A) and biphasic (B) modes of protein diffusion.

Diffusion assay of salivary proteins. Protein of different types of saliva display different modes of diffusion when placed punctually on cellulose laminar matrices (25). Thus, parotid saliva from healthy glands presents a monophasic mode of diffusion that is characterized by a homogeneous radial distribution of the protein on the cellulose lamina (Fig. 1A). In contrast, the samples of submandibular/sublingual and whole saliva displayed biphasic diffusion modes in which a nondiffusible central protein fraction becomes surrounded by a peripheral, radially diffusible protein fraction (Fig. 1B). In the present study, those observations including the analysis of various types of saliva from 33 healthy subjects were mostly corroborated. Thus, among these subjects, 30 right parotid saliva samples and 32 left parotid saliva samples showed a monophasic diffusion mode. Whole and submandibular/ sublingual saliva samples from all the control subjects (n =33) displayed a biphasic mode of diffusion. When saliva samples from the ICRP patients were subjected to this anal-

 Table 4. Mode of diffusion of parotid saliva in relation to the laterality of the disease among ICRP patients

	Mode of protein diffusion of parotid saliva						
Laterality	Biphasic right	Biphasic left	Both biphasic	Monophasic			
Right $(n = 6)$	0	0	6	0			
Left $(n = 13)$	1	1	8	3			
Bilateral $(n = 14)$	1	1	11	1			
Total $(n = 33)$	2	2	25	4			

Results in this table show the number of patients displaying either monophasic or biphasic modes of protein diffusion in either parotid saliva sample in relation to the laterality of the disease.

ysis, some striking differences were observed. On the one hand, both whole and submandibular/sublingual saliva samples from those patients expressed the biphasic mode of diffusion as it normally occurs among healthy subjects. In contrast, 27 of 33 ICRP patients (82%) displayed a biphasic mode of diffusion in at least one of their parotid saliva samples. Clearly, the frequencies of parotid saliva showing the biphasic mode of diffusion in both groups under study were statistically different (Fisher exact test, p < 0.001, Table 3). Furthermore, regardless the laterality of the ICRP disease in the population under study, 25 of the 33 ICRP patients displayed a biphasic mode of diffusion both in their left and right parotid saliva samples (Table 4).

Electrophoretic profiles of salivary proteins in ICRP patients. A direct comparison of the unidimensional electrophoretic protein profiles of parotid saliva samples in our healthy and ICRP groups showed marked differences as to the frequency of a few polypeptide bands. Those were the ones whose apparent molecular weights were 43, 37, 33, 29, 26, 16, and 10 kD. All of them were more frequent among the ICRP patients compared with healthy controls (Fisher exact test, p < 0.05, Table 5). No single change in polypeptide bands was identified in all the ICRP patients (Fig. 2, Table 5). However, 23 of the 31 ICRP patients displaying the 10-kD polypeptide band showed that polypeptide in the saliva produced by the healthy gland. That observation sustained an association between that polypeptide band and the clinic commitment of the gland (p = 0.011; OR = 0.22) (95% CI = 0.68–0.71).

On the other hand, a band-by-band comparative analysis of the protein profiles displayed by the right and the left parotid saliva samples from the same subject allowed us to define the occurrence or absence of symmetry. From an operational point

Table 3. Mode of protein diffusion of both whole and glandular saliva samples from ICRP patients and healthy controls

				Saliva	1			
	Right p	arotid	Left parotid		SBM/SL		Whole	
Group	Mono	Bi	Mono	Bi	Mono	Bi	Mono	Bi
Control $(n = 33)$ ICRP $(n = 33)$	30 6*	3 27*	32 6*	1 27*	0 0†	33 33†	0 0†	33 33†

Mono, monophasic; Bi, biphasic; SBM/SL, submandibular/sublingual.

Results in this table show the number of saliva samples displaying either a monophasic or a biphasic mode of protein diffusion. Statistical significance of ICRP versus the corresponding control was assessed by the Fisher exact probability test.

* p < 0.001; † NS.

Table 5. Distribution of individual polypeptide bands in parotid saliva from ICRP patients and healthy controls

		Polypeptide band							
Parotid saliva	43 kD	37 kD	33 kD	29 kD	26 kD	16 kD	10 kD		
Control $(n = 66)$ ICRP $(n = 66)$	38 (58%) 56 (85%)*	1 (1.5%) 11 (17%)†	3 (4.5%) 19 (29%)‡	12 (18%) 33 (50%)‡	10 (15%) 40 (61%)‡	9 (14%) 32 (49%)‡	16 (24%) 31 (47%)§		

Results in this table show the number of samples in which each selected polypeptide band was observed (percentage in brackets). Statistical significance was assessed by the Fisher exact probability test.

* p = 0.001; † p = 0.004; ‡ p < 0.001; § p = 0.011.



Figure 2. Representative gel of the electrophoretic fractionation of parotid saliva from healthy controls and ICRP patients. Both samples from each subject were run in neighbor lanes. Odd lanes correspond to right parotid saliva samples and even lanes to left parotid saliva samples. Lanes 1–6 show the polypeptide profiles of three ICRP patients and lanes 7–12 the profiles corresponding to three healthy controls. Lane 13 shows M_r standards. On the left, polypeptide bands differing in frequency in the ICRP and control groups are shown.

of view, we considered symmetry when the number of polypeptide bands distinguishing both parotid saliva samples was between 0 and 3 and asymmetry when the number of differences was four or more. Under these considerations, most of the healthy subjects included in the study (25 of 33) exhibited symmetry, whereas most of the ICRP patients (23 of 33) exhibited asymmetry (χ^2 test, p < 0.001) (Fig. 2, Table 6).

MMP-2 and MMP-9 in saliva of ICRP patients. The presence or absence of clear bands having identical mobility as pure metalloproteinases MMP-2 and MMP-9 (92 and 72 kD, respectively) was determined. In the control group, MMP-9 was observed in the right and left parotid saliva samples of only two of 33 subjects, whereas it was present in the whole saliva of 21 subjects and in the submandibular/ sublingual saliva of 13 subjects. In contrast, in the group of 33 ICRP patients, MMP-9 was observed in the right parotid saliva of 16 subjects and in the left parotid saliva of 18

Table 6. Subjects expressing symmetry or asymmetry	in the
polypeptide profile of both contralateral parotid saliva	samples
among healthy controls and ICRP patients	

Group	Symmetry	Asymmetry
Control $(n = 33)$	25	8
ICRP $(n = 33)$	10*	23*

Results in this table show the number of subjects in the control and ICRP groups displaying symmetry or asymmetry. Statistical significance was assessed by the χ^2 test.

* p < 0.001.

subjects, whereas it was detected in whole saliva of 25 subjects and in the submandibular/sublingual saliva of 18 subjects. The frequency of ICRP subjects presenting MMP-9 either in the right or left parotid saliva samples was statistically different from the control group (Fisher exact test, p <0.05) (Table 7, Fig. 3). Likewise, among the healthy subjects, MMP-2 was observed in the right and left parotid saliva samples of only two subjects, whereas it was observed in whole saliva and submandibular/sublingual saliva samples of 17 and eight subjects, respectively. When saliva from 33 ICRP patients was studied, MMP-2 was detected in the right and left parotid saliva samples of eight and 13 subjects, respectively, and in whole saliva and submandibular/sublingual saliva of 14 and eight subjects, respectively. The frequency of ICRP subjects presenting MMP-2 in the left parotid saliva was significantly different from that corresponding to the control group (Fisher exact test, p = 0.003). Comparison of right parotid saliva samples from ICRP and control subjects resulted in no statistical difference (p = 0.082) (Table 7, Fig. 3). Among ICRP patients, MMP-2 but not MMP-9 was significantly associated with the clinic commitment of the parotid gland (p = 0.031; OR = 3.8; 95% CI = 1.1-13.0).

DISCUSSION

ICRP is a pathology of unknown etiology. Direct study of the affected parotid tissue is restricted due to its close imbrication with the facial nerve. Thus, molecular analysis of parotid secretion could offer a valuable complement to the study of the disease as well as to the noninvasive assessment of the glandular function. These aims would need to demonstrate first that parotid saliva is in fact affected in ICRP patients when compared with parotid saliva from healthy subjects. In the present study, some major physicochemical and biochemical alterations in the parotid saliva of ICRP patients were shown. First, parotid saliva of ICRP patients was shown to present a higher protein concentration compared with saliva from a control group. These results differ from

 Table 7. Subjects expressing metalloproteinases in whole and glandular saliva samples among ICRP patients and healthy controls

			Presen	ce of salivary m	etalloproteinase a	activity		
	Right parotid		Left parotid		SBM/SL		Whole	
Group	MMP-9	MMP-2	MMP-9	MMP-2	MMP-9	MMP-2	MMP-9	MMP-2
Control $(n = 33)$ ICRP $(n = 33)$	2 16*	2 8†	2 18*	2 13‡	13 18§	8 8§	21 25§	17 14§

Results in this table show the number of control and ICRP subjects expressing each of those proteases in the various types of saliva. Statistical significance of ICRP versus the corresponding control was assessed by the Fisher exact probability test.

* p < 0.001; † p = 0.082; ‡ p = 0.003; § NS.



Figure 3. Representative zymography gels for metalloproteinases MMP-2 and MMP-9 in whole and glandular saliva samples from healthy controls and ICRP patients. (*A*) Lanes 1–4 and 6–9 correspond, respectively, to the right and left parotid, submandibular/sublingual, and whole saliva samples from an ICRP patient and from a healthy subject. (*B*) Samples corresponding to a different ICRP patient and to a different healthy subject. Clear bands corresponding to metalloproteinases MMP-2 and MMP-9 are shown.

those of Ericson and Sjoback (20) who found no statistical differences in the total protein concentration but higher concentrations of salivary albumin, IgA, lactoferrin, and kallikrein between both groups under study. Higher protein concentration of parotid saliva could be due to an increase in the secretory protein that is synthesized by the gland, to leakage of plasmatic proteins, or to a decrease in the water content of saliva (17,27,28). Although additional molecular analyses are necessary to explain such an increase in protein concentration, the alteration in this biochemical parameter may well be indicative of a functional failure of the gland.

In the present study, we adapted a physicochemical approach to characterize qualitatively the salivary fluids. Saliva diffuses radially from the spotting site when placed punctually on cellulose membranes. After staining, the mode of diffusion displayed by the salivary protein allowed us to make a clear distinction between different types of saliva (25). Parotid saliva is a serous fluid whose protein diffuses homogeneously. On the contrary, submandibular/sublingual saliva and whole saliva exhibit viscoelastic properties that have been related to the presence of the MG1 and MG2 mucins and to intermolecular interactions between them and with other salivary macromolecules (29,30). Both salivary subtypes display a biphasic mode of diffusion (25). In our study, 32 left parotid saliva samples and 30 right parotid saliva samples from 33 healthy subjects (about 94%) showed normal monophasic mode of diffusion. Strikingly, about 82% of parotid saliva samples from ICRP patients showed a biphasic mode of protein diffusion (Fig. 1, Table 3). That observation was in accordance with the altered physical characteristics of parotid saliva that we have observed in these patients and also previously described by various laboratories (1,2,5,9). Moreover, that biphasic mode of diffusion displayed by this saliva was found to be sensitive to disulfide-bridge disrupters (unpublished observations). Our results suggest that at least a fraction of the protein components of parotid saliva from ICRP patients displays a more complex supramolecular nondiffusible protein organization. Again, that new organization may have different origins that would vary from the massive occurrence of new protein components to the occurrence of just a few molecular changes in the protein composition that may lead to a cascade of new intermolecular interactions. More surprisingly in this part of the study was finding that the mode of diffusion of parotid salivary protein of most of the ICRP patients (25 of 33) was affected bilaterally regardless of laterality of the clinical condition (Table 4). Although we have no definite explanation for this observation, our results tempt us to speculate that the healthy contralateral parotid gland of unilaterally affected ICRP patients is also subclinically affected. Studies in that direction are currently under development in our laboratory.

Unidimensional electrophoretic analysis of parotid saliva from ICRP patients showed that bands of 43, 37, 33, 29, 26, 16, and 10 kD were expressed in a frequency significantly higher than among healthy controls. It remains to be seen whether some of those polypeptides may account for the alterations in the physical properties of parotid saliva in ICRP patients. In the same regard, it would be interesting to get more insight into the 10-kD band given its higher frequency in saliva produced by the healthy parotid gland of ICRP patients. Conversely, at variance with the marked bilateral symmetry of polypeptide bandings of parotid saliva samples among healthy controls, parotid saliva samples from ICRP patients were found to be highly asymmetric. This was another main characteristic of parotid saliva in ICRP patients. As in the mode of diffusion of parotid salivary proteins, the increased asymmetry was unrelated to the laterality of the pathologic condition.

The study of more specific protein molecules and the expected damage of the parotid parenchyma in ICRP led us to study MMP-2 and MMP-9 in parotid saliva of ICRP patients (31). Those activities had been previously described in whole saliva of healthy subjects (15). Studies on metalloproteinases involving glandular saliva are rather scarce and have been restricted to low numbers of subjects (32,33). In agreement with previous reports, we found that more than 50% of the whole saliva samples from healthy subjects presented MMP-2 and MMP-9 activity. However, only two healthy subjects expressed that activity in parotid saliva, thus suggesting that either these proteins are in fact absent in this glandular fluid or they are present in parotid saliva of a small percentage of the normal population. We are unaware of whether both MMP-2- and MMP-9-positive subjects were subclinical carriers of a glandular disease. Contamination of parotid saliva of both positive healthy subjects by total saliva was ruled out since MMPs zymography studies of their total saliva were markedly different from those of their parotid saliva (not shown). By contrast, when the presence of MMP-2 and MMP-9 was studied in parotid saliva from ICRP patients, about a third of them (21 of 66) demonstrated MMP-2 activity and about a half (34 of 66) demonstrated MMP-9 activity. MMP-2 was found to be associated with the clinical commitment of the parotid gland since in 14 of the 21 ICRP patients displaying that enzyme, it was detected in the saliva produced by the altered gland. By contrast, despite its higher frequency, MMP-9 was not associated with the clinical condition of the gland since it was observed indistinctly both in healthy and altered parotid glands of ICRP patients. Previous studies had reported the presence of both enzymes in mixed whole saliva of patients affected by conditions that alter salivary glands (15,16). Our study reports for the first time that the frequency of subjects displaying MMP-2 and MMP-9 activities in parotid saliva is much higher among ICRP patients than among healthy subjects. A leading study in parotid saliva of Sjögren syndrome patients had also shown a marked increase in MMP-2 and MMP-9 expression (32). These metalloproteinases have been associated with tissue destruction in periodontitis, dentinal caries lesion progression, and a variety of other oral diseases (31). Thus, MMP-2 and MMP-9 may be involved in the etiopathology of the parotid damage in ICRP patients. On this basis, complementary studies on the quantification of MMP-2 and MMP-9 enzyme activities, the assessment of active and latent molecular forms of these enzymes and the MMPs/ TIMPs (tissue inhibitors of metalloproteinases) balance are currently being addressed. On the other hand, previous reports have shown that in vitro stimulation of a human salivary gland cell line by interferon- γ and tumor necrosis factor α provokes an increase in the expression and secretion of MMP-2 and MMP-9 (32). Thus, studies on the presence of particular cytokines and growth factors in parotid saliva of ICRP patients may also help us to gain valuable insight into the disease (32,34). Also, the possibility of inhibition of metalloproteinases by synthetic inhibitors with the aim of limiting parotid tissue destruction in ICRP patients might be evaluated (31). Thus, our results have clearly demonstrated that parotid saliva in ICRP patients is altered with respect to normal saliva and that some of the molecular differences could be related to the etiopathogenesis of the medical condition and may be of uppermost relevance to the management of ICRP.

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