

WILSON BAMBIRRA JÚNIOR

AVALIAÇÃO DA EXPRESSÃO DE INTEGRINA ALFA2, PROTEÍNA DE  
CHOQUE TÉRMICO 47, E MEDIADORES PRO-INFLAMATÓRIOS E  
IMUNORREGULATÓRIOS EM RESPOSTA À INFECÇÃO ENDODÔNTICA.

FACULDADE DE ODONTOLOGIA  
UNIVERSIDADE FEDERAL DE MINAS GERAIS  
BELO HORIZONTE

2014

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Dissertação apresentada ao Colegiado do programa de Pós-Graduação  
da Faculdade de Odontologia da Universidade Federal de  
Minas Gerais, como requisito parcial para a obtenção  
do grau de Mestre em Odontologia - área de  
concentração em Endodontia

Orientador: Prof. Dr. Antônio Paulino Ribeiro Sobrinho

Co-orientador (a): Prof (a). Dr (a). Leda Quercia Vieira

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UNIVERSIDADE FEDERAL DE MINAS GERAIS

PROGRAMA DE PÓS-GRADUAÇÃO EM ODONTOLOGIA



## FOLHA DE APROVAÇÃO

**Avaliação da expressão de integrina alfa 2, proteína de Choque Térmico 47, e mediadores proinflamatórios e imunorregulatórios em resposta a infecção endodôntica.**

### WILSON BAMBIRRA JÚNIOR

Dissertação submetida à Banca Examinadora designada pelo Colegiado do Programa de Pós-Graduação em ODONTOLOGIA, como requisito para obtenção do grau de Mestre em ODONTOLOGIA, área de concentração ENDODONTIA.

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Prof(a). Evandro Neves Abdo  
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Belo Horizonte, 28 de agosto de 2014.



UNIVERSIDADE FEDERAL DE MINAS GERAIS

PROGRAMA DE PÓS-GRADUAÇÃO EM ODONTOLOGIA



## ATA DA DEFESA DA DISSERTAÇÃO DO ALUNO WILSON BAMBIRRA JÚNIOR

Realizou-se, no dia 28 de agosto de 2014, às 08:30 horas, Sala 3403, Faculdade de Odontologia da Universidade Federal de Minas Gerais, a defesa de dissertação, intitulada *Avaliação da expressão de integrina alfa 2, proteína de Choque Térmico 47, e mediadores pró-inflamatórios e imunorregulatórios em resposta a infecção endodôntica.*, apresentada por WILSON BAMBIRRA JÚNIOR, número de registro 2012733063, graduado no curso de ODONTOLOGIA, como requisito parcial para a obtenção do grau de Mestre em ODONTOLOGIA, à seguinte Comissão Examinadora: Prof(a). Antonio Paulino Ribeiro Sobrinho - Orientador (UFMG), Prof(a). Leda Quercia Vieira (UFMG), Prof(a). Luciana Carla Neves de Brito (FOUI), Prof(a). Evandro Neves Abdo (UFMG).

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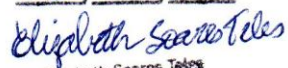
  
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*“Tenha coragem de seguir o seu próprio coração e a sua intuição. Eles de alguma maneira já sabem o que você quer se tornar. Todo o resto é secundário.”*

**STEVE JOBS.**

## RESUMO

**Objetivo:** Analisar a expressão de integrina alfa2, mediadores moleculares, citocinas e quimiocinas, a partir de células presentes no líquido intersticial periapical adjacente a dentes portadores de infecção dos canais radiculares.

**Metodologia:** Os 13 pacientes incluídos no estudo foram encaminhados à Faculdade de Odontologia da Universidade Federal de Minas Gerais (Belo Horizonte, MG, Brasil). As amostras foram retiradas de dentes com necrose pulpar e nenhum paciente apresentou sintomas periapicais agudos no momento das coletas. Após a limpeza e formatação dos sistemas de canais de radiculares (SCR), 3 cones de papel absorventes foram introduzidos no SCR. De forma passiva, o cone de papel ultrapassou o ápice radicular em 2 mm e permaneceu por 1 minuto. As amostras foram coletadas imediatamente após a limpeza do canal radicular e 7 dias após, para caracterizar as expressões dos genes ITGA2, Hsp47, FAK, OPN, IL-1 $\beta$ , TGF- $\beta$ , IL-17A, IL-10, IFN- $\gamma$ , IL-8, CCL2/MCP-1, CCL5, utilizando-se a PCR em Tempo Real.

**Resultados:** Observaram-se níveis significativamente mais baixos de TNF- $\alpha$ , CCL5, CCL2 / MCP-1 e IL-8 em dentes com cargas bacterianas reduzidas (segunda coleta), quando comparada à primeira coleta. Do mesmo modo, a expressão do mRNA das proteínas SSP1 / OPN e FAK diminuiu nas amostras coletadas na segunda amostragem (7 dias). A expressão gênica da IL-10 foi significativamente superior nas amostras coletadas 7 dias após a limpeza e formatação dos SCR quando comparada àquela coletada no dia 0. As expressões gênicas da IL-1 $\beta$ , IL-17A, IFN- $\gamma$ , ITGA2 e Hsp47 / SERPINH1 foram semelhantes nos dois momentos avaliados.

**Conclusão:** Estes resultados sugerem que após a redução da carga bacteriana do SCR uma resposta anti-inflamatória se instala no periápice, aparentemente mediada pela IL-10.

**Palavras-chave:** integrina, SSP1 / OPN, FAK, ITGA2, Hsp47 / SERPINH1, periodontite apical, citocinas e quimiocinas.

## **Abstract**

**Aim:** To examine alpha 2 integrin, molecular mediators, cytokines, and chemokines from cells in periapical interstitial fluid from root canal infections.

**Methodology:** Subjects included 13 patients referred to the College of Dentistry at the Universidade Federal de Minas Gerais (Belo Horizonte, MG, Brazil). Clinical samples were taken from teeth with pulp necrosis and no patients had acute periapical symptoms at the time of the appointments. After cleaning and drying, 3 paper points were introduced into the root canal, passing passively through the root apex (2 mm) into the periapical tissues for 1 minute. The samples were collected immediately after root canal cleaning and 7 days later (restrained root canal bacterial load) to characterize ITGA2, Hsp47, FAK, OPN, IL-1 $\beta$ , TGF- $\beta$ , IL-17A, IL-10, IFN- $\gamma$ , IL-8, CCL2/MCP-1, CCL5 gene expressions using Real Time PCR.

**Results:** Significantly lower levels of TNF- $\alpha$ , CCL5, CCL2/MCP-1, and IL-8 in teeth with restrained bacterial loads (second collection) compared to the first collection were observed. Similarly, the mRNA expression of the proteins SSP1/OPN and FAK decreased in samples from the second collection. The regulatory cytokine IL-10 mRNA was significantly higher in the samples from the second collection compared to the first collection. mRNA expression of IL-1 $\beta$ , IL-17A, IFN- $\gamma$ , ITGA2 and Hsp47/SERPINH1 were similar at both time points.

**Conclusion:** These findings suggest that after reducing the root canal bacterial load an anti-inflammatory response takes place in the periapical area regulated by the Treg IL-10 cytokine.

**Key words:** Integrin, SSP1/OPN, FAK, ITGA2, Hsp47/SERPINH1, apical periodontitis, cytokine and chemokine.

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## LISTA DE ABREVIATURAS E SIGLAS

1. CCL2/MCP-1: *monocyte chemotactic protein 1*
2. CCL5: *Chemokine ligant 5*
3. FAK: *focal adhesion kinases*
4. HSP47: *heat shock protein 47*
5. IFN: *interferon gamma*
6. IL: *interleukin*
7. ITGA2: *integrin alpha 2*
8. OPN: *osteopontin*
9. pH: potencial hidrogênionico
10. RANK: *receptor activator of nuclear factor Kappa B*
11. RANKL: *receptor activator of nuclear factor Kappa B ligant*
12. SCR: sistema de canais radiculares
13. TGF: *transforming growth factor*
14. Th: *T helper*
15. TNF: *tumor necrosis factor*

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*Introdução e relevância*

## 1 INTRODUÇÃO E RELEVÂNCIA

As patologias pulpares e perirradiculares iniciam-se no momento em que microorganismos oportunistas violam a estrutura dental através de lesões cariosas, traumas, restaurações mal adaptadas e defeitos estruturais, como trincas e fissuras produzidas pelo estresse mastigatório (37). Isso permite que agentes patogênicos penetrem na estrutura dental provocando uma resposta inflamatória no hospedeiro frente aos agentes agressores que infectam e colonizam o Sistema de Canais Radiculares (SCR).

A resposta inflamatória contra os patógenos provenientes do sistema de canais radiculares (SCR) infectados induz a liberação de mediadores inflamatórios que, ao conter a invasão dos agentes agressores aos tecidos perirradiculares, promovem destruição localizada no próprio tecido do hospedeiro (50). Mediadores químicos como a Interleucina-1 $\beta$  (IL-1 $\beta$ ), Fator de Necrose Tumoral alfa (TNF- $\alpha$ ), Interferon gama (IFN- $\gamma$ ) promovem interações entre o receptor ativador do fator nuclear Kappa B (RANK) e o seu ligante (RANKL) e consequente ativação e diferenciação dos pré-osteoclastos em osteoclastos. Essa interação provocará uma reabsorção óssea periapical proporcionando a instalação de um tecido imunocompetente que visa impedir a disseminação de microorganismos para tecidos periapicais e para os demais tecidos do hospedeiro. (7,10,13).

No decorrer entre a penetração e a colonização dos SCR, inter-relações ecológicas microbianas ocorrem mudando o perfil e a patogenicidade microbiana. Fatores como: ambiente específico; pressões seletivas existentes nos canais radiculares (48); disponibilidade de nutrientes; o pH; temperatura e baixa tensão de oxigênio

(potencial redox) (43,48) podem determinar estas interações como sinérgicas ou antagônicas (33,34,45,46,48). Essas relações influenciam de forma decisiva na colonização e no estabelecimento da microbiota infectante do SCR (3, 48), culminando na instalação da infecção endodôntica composta principalmente por espécies anaeróbias (2,7,11,21,31,34,43,44,52), e por conseguinte podem produzir respostas agudas ou crônicas aos tecidos periapicais do hospedeiro. Assim a virulência e o número de espécies bacterianas determinarão os efeitos patogênicos microbianos que atuam de forma direta, através da produção de enzimas e de toxinas sobre os tecidos que circundam o dente; e indireta, via estimulação e liberação de mediadores solúveis derivados do hospedeiro, como as citocinas e quimiocinas (42).

A patogenicidade da microbiota infectante determinará o recrutamento de células inflamatórias, liberação de citocinas e quimiocinas pró-inflamatórias e imunorregulatórias responsáveis pela polarização da resposta imune. De acordo com o conhecimento atual, os Linfócitos T *helper* (Th) ativados se subdividem em subgrupos Th1, Th2, Th17 e Treg que se caracterizam principalmente pelo perfil de citocinas produzidas e seus efeitos sobre o organismo (25). A resposta imune do tipo 1 (Th1) caracteriza-se principalmente pela produção de TNF- $\alpha$ , IFN- $\gamma$  e IL-1 $\beta$  que induzirão a diferenciação e ativação dos pré-osteoclastos pelo ligante do receptor ativador do fator nuclear Kappa B (RANKL) e consequente progressão da infecção (10,51). Os osteoclastos se ligam a hidroxiapatita, através da Osteopontina, que é uma proteína fosfoproteica biosintetizada por células do sistema imune, como macrófagos, células T e B, neutrófilos e células dendríticas. A OPN atua como imunomodulador ao promover o recrutamento de células inflamatórias ao sítio da inflamação, influenciando a imunidade celular com funções de citocinas pró-inflamatórias ao estimular a produção de IFN- $\gamma$ , IL-12 e IL-17 e inibir a produção de IL-10 (35,41). A OPN age também como proteína

adesiva, na ligação entre células e átomos de cálcio, disponíveis em tecidos ósseos e dentes, através do fosfato de cálcio atuando como remodelador ósseo e inibidor de crescimento de cristais de cálcio (26,56).

A resposta celular do tipo 2 (Th2) produz IL-4, IL-5, IL-6, IL-10 e IL-13 que estão relacionadas aos processos de cura e reparo da área afetada (19,53). Esses dois subgrupos, Th1 e Th2, apresentam regulação cruzada, através de liberação de citocinas antagônicas como a IL-10 e o fator de crescimento tumoral (TGF- $\beta$ ). Ou seja, as células Th1 inibem a resposta Th2 pela produção de IFN- $\gamma$  e IL-12, enquanto as células Th2 inibem a geração de uma resposta Th1 produzindo citocinas como a IL-4 e IL-10 (28).

Os linfócitos T reguladores (Treg) atuam como agentes imunossupressores e reestabelecem o equilíbrio entre a progressão da infecção e a resposta imunológica do hospedeiro restringindo uma reação inflamatória mais intensa e danosa ao próprio organismo (10,38,58).

Células Th17 são caracterizadas por expressarem a IL-17 e IL-22 e estão relacionadas à exacerbação de processos inflamatórios em lesões periradiculares, através da indução de secreção de mediadores pró-inflamatórios como a IL-8, TNF- $\alpha$ , IL-1 $\beta$ , e IL-6 que promovem rapidamente o recrutamento de neutrófilos para o local da infecção (6), sendo consideradas pontes entre a resposta adaptativa e inata (59), através de estímulo para produzir outras citocinas pró-inflamatórias que atuam na progressão de lesões periapicais (6,57).

Quimiocinas possuem um papel de destaque no processo inflamatório uma vez que funcionam como agentes quimiotáticos e promovem uma resposta imune inicial para diferentes tipos de patógenos, através do recrutamento de células efetoras apropriadas para o sítio inflamatório, incluindo o recrutamento de diferentes células Th1 e Th2. CCL2/MCP-1 tem sido associado ao recrutamento de células inflamatórias e tem

ação sobre a diferenciação de células T efetoras, uma vez que leva a diminuição da produção da IL-12 pelos macrófagos e a supressão da resposta Th1. CCL-5 possui funções quimiotáticas para células T, eosinófilos, e basófilos, desempenhando um papel ativo no recrutamento de leucócitos, além de induzir a expressão de moléculas de adesão celular (7,15).

Os granulomas e cistos radiculares são lesões perirradiculares crônicas, consideradas como sequelas de processos infecciosos resultantes da necrose pulpar. Os linfócitos T CD4+, LT CD8+, neutrófilos, macrófagos, mastócitos e eosinófilos são as principais células presentes nestes sítios e, juntos, formam um intenso infiltrado inflamatório circundado por uma cápsula de tecido conectivo (32).

De todas as células presentes na lesão perirradicular os macrófagos tem merecido destaque na função fagocitária, desempenhando o papel de células apresentadoras de antígenos e secretando fatores de crescimento que estimularão o recrutamento de fibroblastos, proliferação e deposição de colágeno. Com isso tem papel fundamental na coordenação da reparação tecidual (1,9,23).

O processo de reparação da área afetada é ativado por fatores de crescimento, moléculas de adesão celulares, citocinas, quimiocinas, enzimas remodeladoras e componentes da matriz extracelular (MEC) (5,20).

As células produtoras de colágenos sempre são acompanhadas pela expressão de proteínas do choque térmico 47 (Hsp47) que são essenciais para a síntese e processamento de vários tipos de colágenos em vários tecidos e em algumas condições patofisiológicas (29,30). Hsp47 são chaperones moleculares localizados no retículo endoplasmático que se ligam transitoriamente ao pró-colágeno durante seu dobramento, montagem e transporte para o meio extracelular, assegurando a correta maturação de

vários tipos de moléculas de colágeno e impedindo a formação de agregados celulares (30,49).

Integrinas são glicoproteínas localizadas na membrana plasmática e estão envolvidas em interações entre células, células-matriz extracelular e célula-patógeno, por meio de ligações a ligantes distintos ou a combinação de ligantes (47). Essas atuam como mediadores no processo inflamatório, uma vez que exercem papel importante nas etapas de adesão leucocitária, quimiotaxia, fagocitose, adesão endotelial e, por fim, na agregação celular (4).

A classificação das integrinas é feita de acordo com a subunidade  $\beta$  em comum (17,40). Sua afinidade se dá pelo ligante na matriz extracelular que é conferida pela cadeia  $\alpha$  (54).

As principais integrinas que se ligam ao colágeno na célula são  $\alpha 1\beta 1$  e  $\alpha 2\beta 1$ . Essas integrinas se ligam ao colágeno tipo I, muito presente no ambiente perirradicular e também em lesões periapicais (8).

A matriz extracelular é constituída por diferentes combinações de colágenos, proteoglicanos, ácido hialurônico e várias glicoproteínas como a laminina e fibronectina, que atuam de forma a preencher os espaços intercelulares (14,39). A integrina  $\alpha 2\beta 1$  interage com o colágeno e a laminina, atuando não apenas como receptor para essas proteínas, mas também como organizadora da matriz extracelular, moduladora da migração celular e organizadoras do citoesqueleto e da produção de metaloproteinases (16,60).

A subunidade  $\alpha 2$  (ITGA2) é expressa tanto por células epiteliais quanto por fibroblastos, células endoteliais, plaquetas e células T ativadas. Tem como principal função estimular a expressão gênica de colágeno e colagenase em duas e três

dimensões, e assim reparar o arcabouço da matriz de colágeno na região afetada pela infecção (18,22,36).

Os sinais iniciados pela interação das proteínas de matriz com a integrina devem ser transduzidos no interior da célula por proteínas citosólicas denominadas quinase de adesão focal (FAK). Essas proteínas formam um elo entre a matriz extracelular, o citoesqueleto de actina e integrinas (12,27,55). Essa interação atua diretamente na plasticidade da estrutura do citoesqueleto e confere a habilidade da célula em migrar e ou evadir para os sítios de reparo em resposta aos fatores de crescimento.

Os tecidos possuem a capacidade de restaurar sua estrutura original após a remoção do agente agressor por meio de processos reparativos, regulados através de sinalizações e liberação de mediadores químicos. Essas sinalizações ocorrem entre célula-célula e célula-matriz com uma variedade de citocinas e fatores de crescimento que promoverão mitoses celulares e apoptoses, que restabelecem os tecidos afetados pela infecção (24). O conhecimento desses processos de inter-relações celular e tecidual vem desvendar, cada vez mais, a patogênese e a cura das doenças endodônticas, e, tem ainda, a capacidade de desenvolver novos métodos para a prevenção, diagnósticos e tratamento da periodontite apical.



## ***OBJETIVOS***

## **2 OBJETIVOS**

### **OBJETIVO GERAL:**

Avaliar a expressão de integrinas, chaperona molecular, citocinas e quimiocinas em dentes portadores de necrose pulpar, por meio do PCR em tempo real.

### **OBJETIVO ESPECÍFICO:**

Avaliar a expressão de ITGA2, Hsp47, OPN, FAK, IL-1 $\beta$ , IFN- $\gamma$ , TNF- $\alpha$ , CCL2/MCP-1, CCL5, IL-17A, IL-10 e IL-8 nos tecidos perirradiculares, imediatamente após a limpeza e formatação dos SCR (na presença de infecção) e 7 dias após (na ausência de infecção).

## ***TRABALHO CIENTÍFICO***

# **Assessment of apical expression of alpha2 integrin, heat shock protein, proinflammatory and immunoregulatory cytokines in response to endodontic infection.**

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## **Abstract**

**Aim:** To examine alpha 2 integrin, molecular mediators, cytokines, and chemokines from cells in periapical interstitial fluid from root canal infections before and after the reduction of the bacterial load by cleaning procedure.

**Methodology:** Subjects included 13 patients referred to the School of Dentistry at the Universidade Federal de Minas Gerais (Belo Horizonte, MG, Brazil). Clinical samples were taken from teeth with pulp necrosis and no patients had acute periapical symptoms at the time of the appointments. After cleaning and drying, three paper points were introduced into the root canal, passing passively through the root apex (2 mm) into the periapical tissues for 1 minute. The samples were collected immediately after root canal cleaning and seven days later (restrained root canal bacterial load) to characterize those gene expressions using Real Time PCR.

**Results:** Significantly lower levels of TNF- $\alpha$ , CCL5, CCL2/MCP-1, and IL-8 in teeth with restrained bacterial loads (second collection) compared to the first collection were observed. Similarly, the mRNA expression of the integrins SSP1/OPN and FAK decreased in samples from the second collection. The mRNA for the regulatory cytokine IL-10 was significant higher in samples from the second collection compared to the first collection. mRNA expression of IL-1 $\beta$ , IL-17A, IFN- $\gamma$ , ITGA2 and Hsp47/SERPINH1 were similar at both time points.

**Conclusion:** These findings suggest that after reducing the root canal bacterial load an decrease in the inflammatory response took place in the periapical area.

**Key words:** Integrin, SSP1/OPN, FAK, ITGA2, Hsp47/SERPINH1, apical periodontitis, cytokine and chemokine

## **Introduction**

Apical periodontitis is an inflammatory disease of periradicular tissues caused by the host immune response to root canal infection, characterized by localized inflammation concomitant with bone resorption. The pathogenesis of periapical disease involves degradation of extracellular matrix components, including laminin, fibronectin, collagen and proteoglycans (13, 21, 24, 27).

Lymphocytes are the most prevalent cells in the periapical inflammatory infiltrate (46). These cells are associated with the production of cytokines and chemokines that restrain the root canal infection (31, 38), but also lead to alveolar bone destruction.

The antigen-presenting cells, especially dendritic cells and macrophage, are responsible, at least, for the polarization of four different Th subsets (28). Inflammatory response is related to Th1 subset which produces cytokines like IFN- $\gamma$ , TNF- $\alpha$  and IL-1, which are involved in the progression, bone destruction and remodeling of periapical lesions (39). Conversely, the healing process is related to the Th2 subset (21). Th17 subset may play a role in exacerbating inflammation by stimulating the secretion of pro-inflammatory mediators, such as IL-8, TNF- $\alpha$ , IL-6 (20). Regulatory T (Treg) cells maintain normal homeostasis and reduce the over-activity of Th1, Th2 and Th17 response. IL-10 exhibits strong anti-inflammatory properties and are produced by Th1, Th17, Th2 and Treg cells (4).

Cleaning and shaping procedures play a role in the reduction of root canal microbiota and its byproducts, allowing the proper healing of apical tissues (14). This process is tightly regulated through cell-cell interactions and cell-matrix signaling,

promoted by integrins, a variety of cytokines and growth factors, as well as by cell mitosis and apoptosis (6,16).

Integrins are a large family of heterodimeric membrane glycoproteins involved in cellular processes mediated by cell to cell and cell to extra-cellular matrix interactions. Alpha-2 integrin mediates induction of collagenase MMP-1 in fibroblasts within collagen gels (22).

Heat Shock proteins (HSPs) exert a protective role against harmful environmental conditions and pathogens. They are characteristically induced by stress signals such as elevated temperature, infectious agents and inflammatory mediators participating in the folding and translocation of polypeptides across the cell membrane (11). Hsp47 is the major collagen-binding heat inducible glycoprotein expressed in fibroblast (30,43).

The aim of this study was to quantitatively assay the mRNA expression of the alpha 2 integrin (ITGA2), heat shock protein 47 (Hsp47/SERPINH1), osteopontin (OPN/SSP1), focal adhesion kinase FAK/PTK2, as well as TNF- $\alpha$ , IFN- $\gamma$ , IL-1 $\beta$ , IL-17, IL-10, CCL2/MCP-1, CCL5, and IL-8 in samples collected from interstitial fluid adjacent to human root canal infections. In addition, we detected the same integrins and cytokines seven days later, after the root canal cleaning procedures, when root canal bacterial load was strongly reduced, providing conditions for healing process. Finally, we statistically compared and analyzed both sets of data.

## **Materials and Methods**

### ***Study Participants***

Subjects were 13 patients referred to the Dental School at the Federal University of Minas Gerais (UFMG), Belo Horizonte, Brazil to receive endodontic care, 18 to 80 years old. The exclusion criterion for this study was anti-inflammatory and antibiotic therapy up to 3 months before starting endodontic therapy. All participants signed the informed consent formulary. This study was approved by the Ethics Committee of the Federal University of Minas Gerais (CAAE 20383914.0.0000.5149).

### ***Sample Collection***

Clinical samples were taken from 13 teeth (single and multirooted) presenting pulp necrosis and apical periodontitis, diagnosed by clinical and radiographic analyses and pulp sensibility tests. All selected patients failed to present acute periapical symptoms. Teeth were isolated by using a rubber dam followed by complete asepsis of the isolated crown with 30% hydrogen peroxide, followed by 5% iodine and by inactivation with 5% sodium thiosulfate as proposed by Möller. Cleaning and shaping of the root canals were completed by using ProTaper nickel-titanium files (Dentsply Maillefer, Ballaigues, Switzerland) in conjunction with 5.2% sodium hypochlorite. Collections were performed as previously described (19). Briefly, the samples were collected using paper points immediately after root canal cleaning to characterize the mRNA expression profiles of cytokines, chemokines, and integrins. After cleaning and drying, three paper points were introduced into the root canal through the root apex (2 mm) into the periapical tissues, for 1 minute (day 0). The 4 mm tip of the paper points were dropped into a microcentrifuge tube, and the samples were stored at -70°C. By



using this procedure, RNA was collected from the periapical interstitial fluid. No endodontic dressing was inserted into root canals. The coronal accesses of the teeth were restored with eugenol-based cement. Seven days later (day 7), the teeth were reopened, and the periapical interstitial fluid was sampled again to characterize the expressions of cytokines and integrins in teeth after instrumentation. In teeth with multiple canals, the first (day 0) and second (day 7) samples were collected from the same canal. No clinical signs or symptoms were present at the time of the second collection, and root canals were filled by lateral condensation technique.

### ***Sample Preparation***

Total RNA was extracted from each sample with TRIZOL reagent (GIBCO/BRL Laboratories, Grand Island, NY). Briefly, chloroform was added to the tubes containing the paper points, and the mixture was centrifuged at 12,000g at 4°C for 15 minutes; the aqueous phase was collected, and RNA was precipitated by isopropanol, 75% cold ethanol was added and samples were centrifuged again at 12,000g at 4°C for 15 minutes. Ethanol was discarded, samples were dried, dissolved in RNase-free water, and then incubated at 55°C for 10 minutes. The RNA was then stored at -70°C.

### ***Real-time Polymerase Chain Reaction***

Complementary DNA was synthesized by using 1 µg of RNA through a reverse transcription reaction as described by Barbosa Silva *et al.* Polymerase chain reaction (PCR) was carried out under standard conditions: a holding stage of 95°C (10 minutes), a cycling stage with 40 cycles of 95°C (15 seconds), followed by 60°C (1minute) and a melt curve stage of 95°C (15 seconds), 60°C (1 minute), and 95°C

(15 seconds). The primer sequences used for quantitative PCR analysis of IFN- $\gamma$ , TNF- $\alpha$ , IL-1 $\beta$ , IL-17, IL-10, CCL2/MCP-1, IL-8, CCL5, ITGA2, Hsp47, OPN, FAK, and GAPDH mRNA expression are shown in Table 1. The sequences of human primers were designed by using PRIMEREXPRESS software (Applied Biosystems, Foster City, CA) based on nucleotide sequences available in the GenBank database. The real-time PCR assay was performed by using the StepOne Real-time PCR System (Applied Biosystems). A Syber-Green detection system (Applied Biosystems) was used to assay primer amplification. Glyceraldehyde-3-phosphate dehydrogenase (GAPDH) was used as a housekeeping gene for normalization. All samples were run in duplicate. Sequence Detection System Software version v 2.4.1 (Applied Biosystems ®) was used to analyze data after amplification. Results were obtained as threshold cycle ( $C_T$ ) values. Expression levels were calculated by using the comparative  $2^{-\Delta\Delta C_T}$  method (25). The values were calculated as the mean value of duplicates for each patient, and the expression levels of mRNAs in all samples were expressed as the ratio of each specific primer to GAPDH expression.

### ***Statistical Analysis***

Data analysis was performed by using SPSS for Windows (version 15.0; SPSS, Chicago, IL, USA). Data were subjected to the Shapiro-Wilk test to characterize their normality. Because data did not present a normal distribution, the Wilcoxon test was used to determine statistical differences in samples from the same groups ( $P < 0.05$ ).

## Results

mRNA expression was determined by real-time PCR and quantified by comparison with the internal control gene GAPDH. Assessment revealed significant decreases in TNF- $\alpha$ , CCL5, CCL2/MCP-1, and IL-8 mRNA levels on day 7 when compared to day 0 ( $p < 0.05$ ) (Fig. 1). Similarly, the mRNA expression of the integrins SSP1/OPN and FAK decreased in samples from the second collection ( $p < 0.05$ ) (Fig. 2). IL-1 $\beta$ , IL-17A, IFN- $\gamma$  and Hsp47/SERPINH1 and ITGA2 gene expression was similar at both time points ( $p > 0.05$ , Fig 1 and 2). mRNA expression of IL-10 significantly increased at day 7 ( $P < 0.05$  Fig. 1).

## Discussion

The presence of microorganisms and their byproducts inside infected root canals evokes host immune response aimed at avoiding dissemination through the periapical foramen. Cleaning and shaping procedures strongly diminish the root canal bacterial load and, afterwards, periapical healing starts. This process involves overlapping phases of inflammation, proliferation and remodeling, each characterized by dynamic interactions among components of the extra-cellular matrix, growth factors, and cells (2, 33). In this study, expressions of inflammatory and regulatory genes were investigated in periapical lesions immediately after root canal cleaning procedures and 7 days later.

The identification of novel targets that restrain the progression of periapical lesions might improve diagnostic procedures and the development of individualized treatment plans and regenerative procedures. In this regard, this study investigated ITGA2 and Hsp47 gene expression in active lesions and seven days after root canal cleaning, since several periapical cellular processes involve cell-extra-cellular matrix and cell-cell interactions (43). Integrins are critical components of the cell attachment

machinery. Most normal vertebrate cells cannot survive unless they are anchored to the extra-cellular matrix. ITGA2 and Hsp47 are major collagen-binding heat-inducible glycoproteins in fibroblast (13), which are expressed associated with collagen in various tissues during pathophysiological conditions (30). In this study, ITGA2 and Hsp47 mRNA were present at similar levels in both conditions evaluated. This result suggests that the healing process is still in course seven days after root canal instrumentation. Recently, it was shown that ITGA4 mRNA expression was significantly higher in inactive compared with active periapical granulomas (16). However, no data for ITGA2 were found in the literature.

Hsp44 genes were significantly increased in a pool of periapical granulomas compared with a pool of healthy periodontal ligaments (17). These authors indicated that HSP40 gene family plays a crucial role in the maintenance of infection and tissue destruction in apical periodontitis. Here, we investigated the levels of Hsp47 in endodontic lesions. We found that Hsp47 was expressed in active lesions and seven days after cleaning, at similar levels, again indicating that seven days after root canal instrumentation tissue re-shaping was still in course.

Recently, it was shown that FAK expression is associated with cell motility, invasiveness, proliferation and apoptosis (18, 23, 47). In the present study, after cleaning procedures FAK mRNA expression was significantly down-regulated ( $p < 0.05$ ) indicating that microbial infection stimulates its expression, as demonstrated elsewhere (32).

OPN regulates adhesion and recruitment of neutrophils and macrophages, and T cell binding to integrins (34). It has been associated with Th1 responses by increasing IL-12 and diminishing IL-10 production (5, 45). In this study, OPN mRNA expression was significantly decreased at day 7 after cleaning procedures ( $p < 0.05$ ), which

mirrored the decrease of type 1 cytokines at the same time, as demonstrated by others (10, 40, 41).

The inflammatory cytokines IL-1 $\beta$ , IL-17, IFN- $\gamma$ , IL-8, and TNF- $\alpha$  were detected in periapical interstitial fluid, as observed elsewhere (10, 15, 35, 40, 42). In this study, mRNA expression of the inflammatory cytokines TNF- $\alpha$ , CCL5, CCL2/MCP-1, and IL-8 decreased at day 7 after cleaning and shaping procedures, in agreement with the findings previously observed (10, 40, 41).

IL-1  $\beta$  stimulates bone resorption by osteoclasts, by inducing IL-17 synthesis (28). Moreover, IL-17 induces the secretion of IL-8, TNF- $\alpha$  and IL-1 $\beta$  by monocytes. These cytokines promote quick neutrophil movement to the site of the infection (8, 20). In this study mRNA expression of IL-1 $\beta$  and IL-17 were similar at both times analyzed. Recently, similar results were found concerning IL-17 but not in regards to IL-1 $\beta$ , which was expressed in similar levels during the observation time (10, 40, 41). In an animal model, it was demonstrated that both cytokines (IL-1 $\beta$  and IL-17) were detected in higher levels during the active phase when compared with chronic phase of lesion development (21, 44). Thus, it is possible that, despite the cleaning procedure, some inflammation was still in course in our patients, as suggested by these cytokines. However, mRNA expression of TNF- $\alpha$  and IL-8 were significantly lower seven days after root canal cleaning ( $p < 0.05$ ), suggesting resorption and neutrophil recruitment decrease after this procedure. Other researchers did not observe any significant differences in mRNA TNF- $\alpha$  expressions, neither in HIV negative (10, 40, 41) nor in HIV positive individuals (9).

CCL2/MCP-1 and CCL5 are chemotactic for monocytes, T cells, eosinophils, and basophils, and play an active role in recruiting leukocytes into inflammatory sites (12, 26, 36). Association between CCL5 and Th1-type cellular responses was suggested

by the finding that neutralization of CCL5 reduced type 1 granuloma formation (7). In this study, CCL2 and CCL5 mRNA expressions were significantly decreased at day 7 after cleaning procedures ( $p < 0.05$ ). Previously it was demonstrated similar results concerning CCL5, but not in regards to CCL2, which was found at similar levels on days zero and seven after root canal cleaning (10). These contradictory results may be due to differences in the efficiency of the cleaning procedures. We suggest that the process of control of inflammation has started at this time point and if investigated later would reveal a more clear-cut non-inflammatory scenario. In addition, different initial infections might account for the variability encountered. Further evidence of the healing and immunoregulatory process at day 7 in our patients is the higher levels of IL-10 found seven days after root canal cleaning.

In conclusion, our results demonstrate a decrease in mRNA expression of proinflammatory mediators such as TNF- $\alpha$ , IL-8, OPN, FAK, CCL5 and an increase in the mRNA expression of immune regulator cytokine (IL-10) when the root canal bacteria load was reduced. Fibroblast and endothelial cells are expected to migrate into the healing tissue. Alpha 2 integrin and heat shock protein 47 had similar expression before and after cleaning procedures, probably indicative of the healing process. Further comprehensive clinical studies should be conducted in the periapical area, given the fact that several other players are involved in the inflammatory and healing processes. However, expression of the genes described here suggested an on-going healing process after reduction of bacterial load by root canal cleaning.

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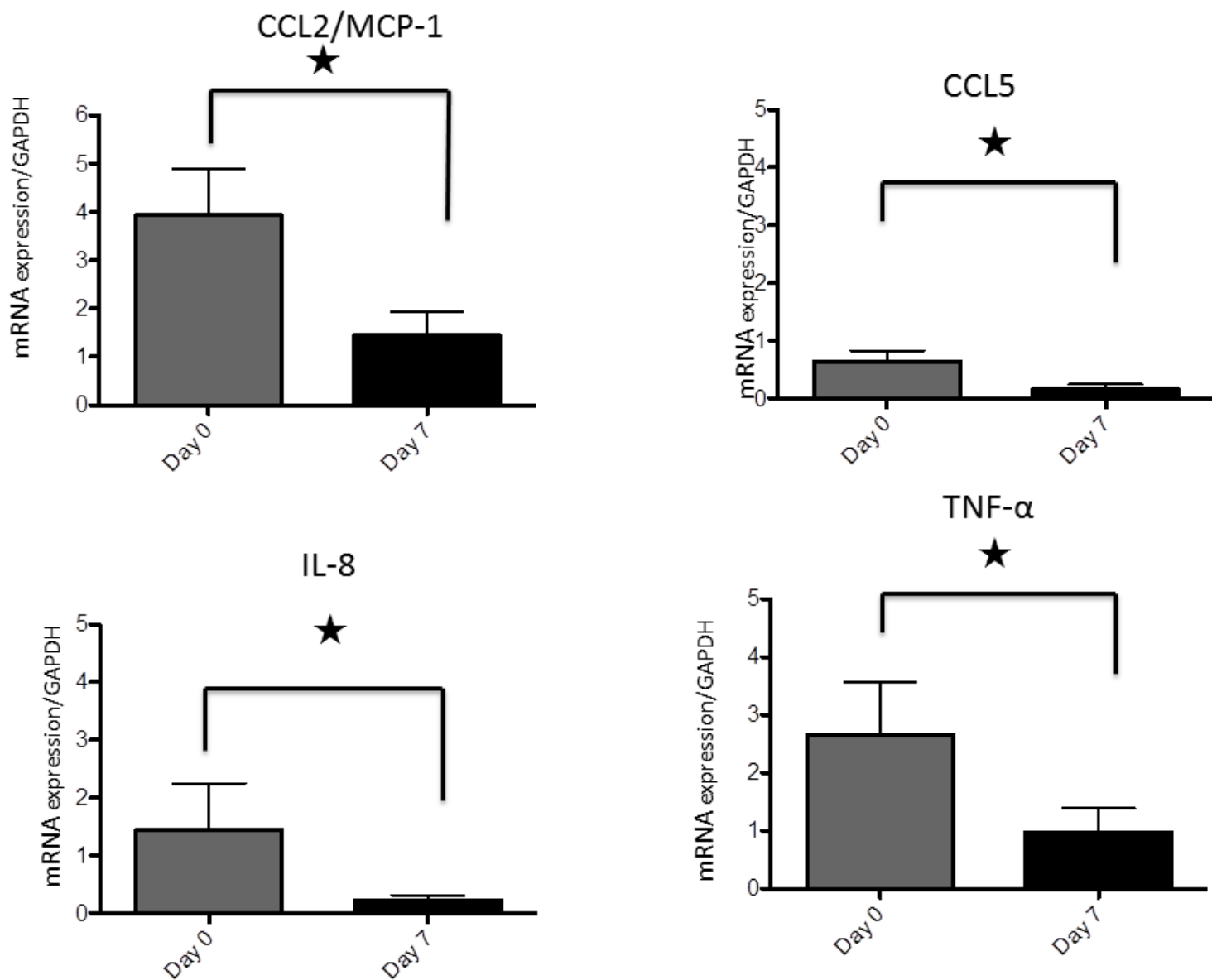
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Gene	Sense and antisense	Mt* (°C)	bp*
GAPDH	5'-GCA CCA CCA ACT GCT TAG CA- 3'	80	96
	5'-TGG CAG TGA TGG CAT GGA GGA- 3'		
TNF- $\alpha$	5'-TTC TGG CTC AAA AAG AGA ATT G- 3'	76	73
	5'-TGG TGG TCT TGT TGC TTA AGG- 3'		
IL-1 $\beta$	5'-TGG CAG AAA GGG AAC AGA A- 3'	73	59
	5'-ACA ACA GGA AAG TCC AGG CTA- 3'		
IL-10	5'-GGT TGC CAA GCC TTG TCT GA- 3'	81	107
	5'-TCC CCC AGG GAG TTC ACA T- 3'		
IL-17	5'-CAA TGA CCT GGA ATT ACC CAA- 3'	70	52
	5'-TGA AGG CAT GTG AAA TCG AGA- 3'		
IFN- $\gamma$	5'-ACT GAC TTG AAT GTC CAA CGC A- 3'	61	101
	5'-ATC TGA CTC CTT TTT CGC TTC CC- 3'		
ITGA2	5'-GCA GAT GGA CCA CAC TTT GA- 3'	60	115
	5'-TGT CTG TGC CCT TTT CCT CT- 3'		
HSP47	5'-TGC TGA GCC CGG AAA CTC- 3'	62	63
	5'-TTC AGG GCA GGC AGA ATG- 3'		
IL-8	5'- GAA TGG GTT TGC TAG AAT GTG ATA- 3'	60	129
	5'- CAG ACT AGG GTT GCC AG ATT TAA C - 3'		
CCL5	5'-CGT GCC CAC ATC AAG GAG TA- 3'	80	91
	5'-CAC ACA CTT GGC GGT TCT TTC- 3'		
OPN	5'-GCC GAG GTG ATA GTG TGG TT- 3'	62	101
	5'-TGA GGT GAT GTC CTC GTC TG- 3'		
FAK	5'-CAA CAG GTG AAG AGC GAT TA- 3'	58	99
	5'-CCA GTA TGA TCG CCG TAT TT- 3'		
CCL2	5'-AGG ACC ATT GTG GCC AAG GA-3'	81	93
	5'-CGG AGT TTG GGT TTG CTT GT-3'		

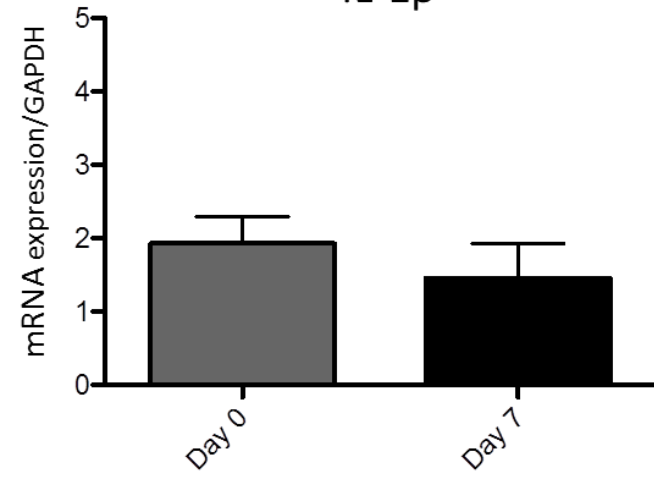
Table 1. Primer sequences

\*Mt: melting temperature; bp: base pairs of amplicon size

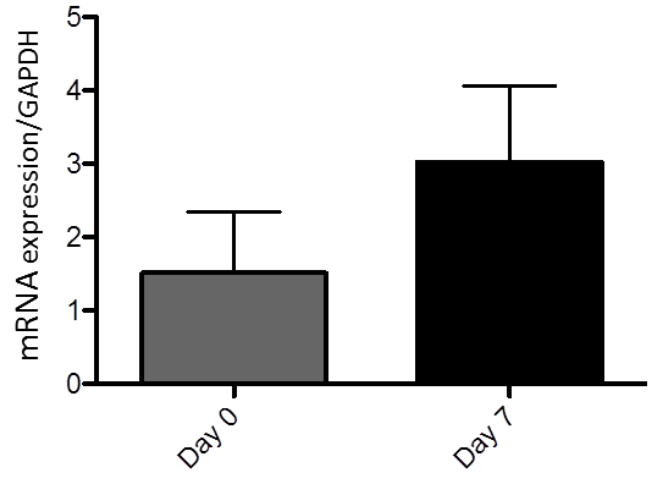
**Figure 1.** Expression of CCL2/MCP-1, CCL5/RANTES, IL-8, TNF, IL-1 $\beta$ , IFN- $\gamma$  IL-17A and IL-10 in infected root canal periradicular tissues. Expression was determined by real-time PCR and quantified by comparison with internal control (GAPDH). Bars represent mean values of samples recovered from 13 patients; lines represent standard error of mean. \* indicates  $p < 0.05$ .



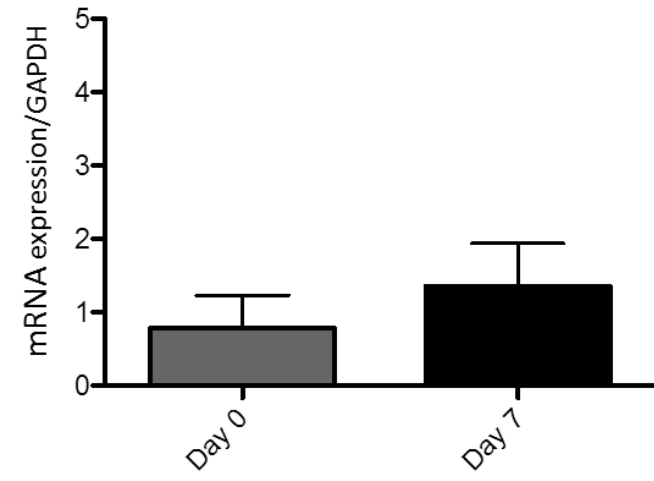
IL-1 $\beta$



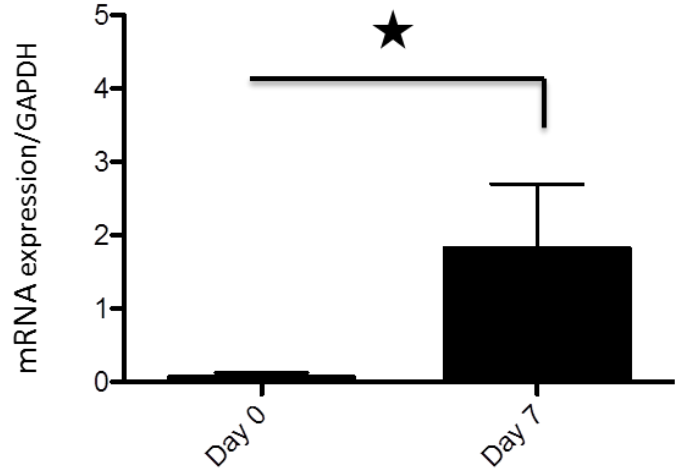
IFN- $\gamma$



IL-17A

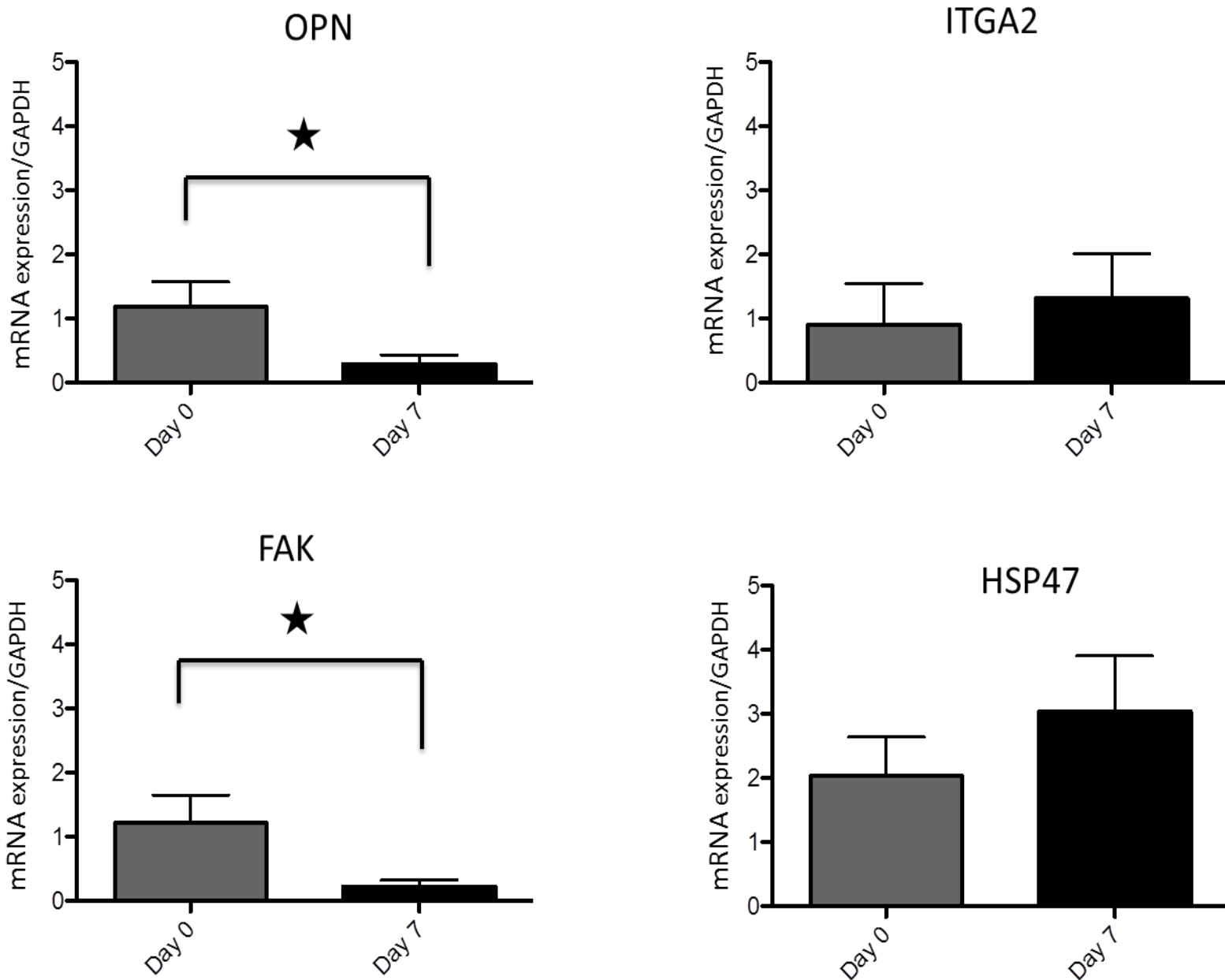


IL-10





**Figure 2.** Expression of SSP1/OPN, ITGA2, FAK and Hsp47/SERPINH1 in infected root canal periradicular tissues. Expression was determined by real-time PCR and quantified by comparison with internal control (GAPDH). Bars represent mean values of samples recovered from 13 patients; lines represent standard error of mean. \* indicates  $p < 0.05$ .



## ***CONCLUSÃO***

## 4 Conclusão

Conclui-se que a expressão de integrina alfa2 (ITGA2), proteína do choque térmico 47 (Hsp47), das proteínas citosólicas OPN e FAK, bem como dos mediadores pro-inflamatórios IL-1 $\beta$ , IFN- $\gamma$ , TNF- $\alpha$  e IL-17, das quimiocinas CCL2/MCP-1, CCL5 e IL-8, e da citocina reguladora IL-10 sofrem o efeito da presença ou da ausência de uma carga microbiana efetiva no interior do SCR. Ou seja, que ocorre um predomínio dos mediadores pró-inflamatórios na presença da infecção endodôntica e uma redução dos mesmos, em concomitância com o aumento da expressão da citocina IL-10, após a redução da carga bacteriana.

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