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# Defining the role of *Porphyromonas gingivalis* peptidylarginine deiminase (PPAD) in rheumatoid arthritis through the study of PPAD biology

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# Abstract

**Background**—Antibodies to citrullinated proteins (ACPAs) are a hallmark of rheumatoid arthritis (RA). *Porphyromonas gingivalis* peptidylarginine deiminase (PPAD) has been implicated in the initiation of RA by generating citrullinated neoantigens and due to its ability to autocitrullinate.

**Objectives**—To define the citrullination status and biology of PPAD in *P gingivalis* and to characterize the anti-PPAD antibody response in RA and associated periodontal disease (PD).

**Methods**—PPAD in *P gingivalis* cells and culture supernatant was analyzed by immunoblotting and mass spectrometry to detect citrullination. Recombinant PPAD (rPPAD), inactive mutant PPAD (rPPAD<sup>C351S</sup>), and N-terminal truncated PPAD (rPPAD<sup>Ntx</sup>) were cloned and expressed in*E coli*. Patients with RA and healthy controls were assayed for IgG antibodies to citrullinated rPPAD and unmodified rPPAD<sup>C351S</sup> by ELISA. Anti-PPAD antibodies were correlated with anti-CCP3 antibody levels, RA disease activity, and PD status.

**Results**—PPAD from *P gingivalis* is truncated at the N- and C-terminal domains and not citrullinated. Only when artificially expressed in *E coli*, full-length rPPAD, but not truncated (fully active) rPPAD<sup>Ntx</sup>, is autocitrullinated. Anti-PPAD antibodies show no heightened reactivity to citrullinated rPPAD, but are exclusively directed against the unmodified enzyme. Antibodies against PPAD do not correlate with anti-CCP levels and disease activity in RA. By contrast, anti-PPAD antibody levels are significantly decreased in RA patients with PD.

**Conclusions**—PPAD autocitrullination is not the underlying mechanism linking PD and RA. N-terminal processing protects PPAD from autocitrullination and enhances enzyme activity. Anti-PPAD antibodies may have a protective role for the development of PD in RA patients.

Competing Interests None declared.

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# Keywords

Rheumatoid arthritis; periodontal disease; *Porphyromonas gingivalis* peptidylarginine deiminase; anti-PPAD antibodies; ACPA

# INTRODUCTION

Rheumatoid arthritis (RA) is a systemic autoimmune disease of unknown etiology characterized by synovial inflammation and joint destruction.<sup>12</sup> Loss of tolerance to citrullinated proteins is a hallmark of RA pathogenesis.<sup>3</sup> Autoantibodies to citrullinated proteins (ACPA) are highly specific for RA and precede the onset of clinical disease by years.<sup>4–6</sup> Peptidylarginine deiminases (PAD) are enzymes that catalyze the posttranslational modification reaction of arginine residues to citrulline.<sup>78</sup> By modifying autoantigens in synovial tissue and alternative sites of inflammation, PAD activity may be a *sine qua non* in initiating and sustaining autoimmunity in ACPA-positive RA. *Porphyromonas gingivalis* peptidylarginine deiminase (PPAD), a bacterial deiminase evolutionarily unrelated to human PAD,<sup>89</sup> has received considerable attention by investigators trying to identify a mechanism linking periodontal disease (PD), bacterial citrullination and RA.<sup>10–13</sup>

PD is a chronic inflammatory disease caused by infection of the supporting tissues of the teeth, ultimately resulting in alveolar bone destruction and tooth loss.<sup>14</sup> An increased prevalence of PD has repeatedly been reported in RA.<sup>15–19</sup> Among the periodontal pathogens, *P gingivalis*, an anaerobic bacterium strongly associated with chronic periodontitis,<sup>20–22</sup> is unique in its expression of a bacterial PAD.<sup>9</sup> PPAD is believed to be a major virulence factor of *P gingivalis* due to its capacity to generate ammonia in the deimination reaction of arginine to citrulline.<sup>8</sup> Ammonia may protect *P gingivalis* during acidic cleansing cycles in the mouth,<sup>92324</sup> and promote periodontal infection via inhibitory effects on neutrophil function.<sup>2526</sup>

PPAD is almost exclusively detected in outer membrane (OM) fractions of *P gingivalis*,<sup>12</sup> and as a secreted enzyme.<sup>9</sup> Interestingly, OM-associated PPAD and secreted PPAD were only recently shown to be truncated both at the C- and N-terminal domains.<sup>927</sup> While C-terminal cleavage is required for cell surface translocation of PPAD,<sup>27–29</sup> N-terminal processing appears to maintain enzyme activity and stability.<sup>30</sup>

By citrullinating C-terminal peptidylarginine in the context of periodontal infection, *P* gingivalis has been hypothesized to play a primary role in RA pathogenesis.<sup>113132</sup> Moreover, the recent finding that recombinant PPAD (rPPAD) is autocitrullinated and preferentially recognized by antibodies in RA suggests that loss of tolerance to citrullinated proteins in RA may originate from an antimicrobial immune response directed against citrullinated PPAD.<sup>12</sup>

This study defines the structure and citrullination status of the cellular and secreted forms of PPAD (cPPAD and sPPAD, respectively) in *P gingivalis*. The relevance of the anti-PPAD antibody response for RA and associated PD is examined in the context of PPAD biology.

# MATERIAL AND METHODS

#### **Patients and controls**

Serum was obtained from 83 RA patients (1987 revised ARA criteria)<sup>33</sup> and 39 healthy controls recruited under the protocol of the Comprehensive Oral Health Assessment in Patients with Arthritis and Autoimmune Inflammatory Diseases (OHARA). The protocol was approved by the Johns Hopkins Medicine Institutional Review Board, and informed consent was obtained from all participants. Study participants underwent a comprehensive dental evaluation. PD was defined using the 2007 CDC/ AAP standard cases definitions for surveillance of moderate and severe periodontitis based on assessments of pocket depth and attachment loss;<sup>1434</sup> subjects not fulfilling these definitions were considered periodontally healthy.

### Bacterial samples and immunoblotting

*P gingivalis* strain W83 was obtained from the University of Maryland (courtesy of Mark A. Reynolds). Bacterial cells were pelleted by centrifugation, lysed in ice-cold NP-40 lysis buffer (20 mM Tris pH 7.4, 150 mM NaCl, 1 mM EDTA, 1% NP-40 Alternative), sonicated, and boiled in SDS sample buffer. *P gingivalis* culture supernatant was concentrated using Amicon Ultra-15 units (EMD Millipore) and buffer exchanged with 20 mM Tris pH 7.6, 150 mM NaCl, 10% glycerol. Samples were resolved by SDS-PAGE and analyzed by anti-modified citrulline (AMC) immunoblotting,<sup>35</sup> or using a polyclonal antibody raised in rabbits by immunization with recombinant PPAD (Covance).

# Cloning and purification of recombinant PPAD and alpha-enolase

The full-length PPAD coding sequence was amplified from *P gingivalis W83* DNA (ATCC) and cloned into pET-28a(+) (Novagen), generating a fusion protein with N-terminal His-tag. Enzymatically inactive PPAD was generated by site-directed mutagenesis to replace cysteine 351 at the active site of the protein with serine (PPAD<sup>C351S</sup>).<sup>36</sup> N-terminal truncated PPAD (rPPAD<sup>Ntx</sup>) was generated by amplifying the coding sequence for amino acids 44 to 556 of full-length PPAD with primers containing a 5' Kozak sequence. The PCR product was cloned into pET-28a(+) to encode for a C-terminal His-tagged fusion protein. Alpha-enolase encoding cDNA was cloned into pET-28a(+). Recombinant PPAD, PPAD<sup>C351S</sup>, PPAD<sup>Ntx</sup>, and enolase were expressed in *E coli* BL21 (DE3) (Agilent), and purified from the soluble fraction of cell lysates prepared in 20 mM Tris pH 7.6, 400 mM NaCl, 5 mM imidazole, 20 mM  $\beta$ -mercaptoethanol, 1% Triton X by Ni-NTA affinity chromatography (Qiagen). Purity of rPPAD and rPPAD<sup>C351S</sup> used in ELISA assays exceeded 95% (see online supplementary figure S1A). Enolase was citrullinated *in vitro* using human rPAD4 as previously described.<sup>37</sup>

#### PPAD ELISA

Polystyrene plates (Costar) were coated with 100ng/well citrullinated rPPAD (cit-rPPAD), rPPAD<sup>C351S</sup> or PBS pH 7.4 alone (used as a negative control). Coated plates were washed with PBS 0.05% Tween 20 (PBS–T), and unoccupied binding sites blocked with PBS–T 3% non-fat dry milk (PBS–TM). Sera were diluted at 1:1000 in PBS–TM 1% and assayed in

duplicate. HRP-conjugated anti-human IgG (Jackson ImmunoResearch) was used as a secondary antibody. A serial dilution of rabbit anti-PPAD was used as a standard. Arbitrary units (AU) were calculated from standard dilutions, and individual values corrected for background by subtracting the reactivity of PBS-coated wells.

# RESULTS

### PPAD from P gingivalis is truncated and not citrullinated

Cell lysates and supernatant from *P gingivalis* were initially studied by immunoblotting to characterize bacterial PPAD. Two distinct patterns of PPAD were identified. In bacterial cells, anti-PPAD immunoblotting detected bands of approximately 75-85 kDa (cPPAD), while a single band of 47 kDa (sPPAD) was identified in *P gingivalis* supernatant (figure 1A, middle panel). The observed size of sPPAD is consistent with the truncated form of PPAD.<sup>927</sup> In contrast to a previous report,<sup>11</sup> only low levels of protein citrullination were detected by AMC immunoblotting in bacteria lysed in NP-40 buffer (figure 1A, right panel), and no citrullination was found when *P gingivalis* was directly lysed and boiled in SDS sample buffer (see online supplementary figure S1B). Bacterial citrullination may therefore occur *in vitro* during cell lysis. Importantly, cPPAD and sPPAD are not citrullinated in bacterial samples (figure 1A, right panel and online supplementary figure S1B).

Mass spectrometry (MS) analysis of cPPAD and sPPAD invariably identified a truncated protein missing the first 43 N-terminal and 97 C-terminal amino acids of full-length PPAD (figure 1B and 1C, respectively). Peptides outside these terminal domains showed sequence coverage of 82.9% (cPPAD) and 92.5% (sPPAD). In contrast to a previous study of full-length rPPAD expressed in *E coli*,<sup>12</sup> we found that PPAD in *P gingivalis* is not citrullinated (figure 1B,C). Lack of autocitrullination was not attributable to loss of enzymatic function as confirmed using the BAEE assay (data not shown).

#### N-terminal truncation prevents PPAD autocitrullination and amplifies enzyme function

Previous studies of recombinant full-length PPAD and N-terminal truncated PPAD have shown conflicting results with regard to enzyme autocitrullination.<sup>1230</sup> While full-length PPAD is strongly citrullinated during expression in *E coli*,<sup>12</sup> the truncated enzyme is not autocitrullinated.<sup>30</sup> We hypothesized that N-terminal processing of PPAD may protect the enzyme from autocitrullination in *P gingivalis*. Indeed, when we expressed full-length PPAD (rPPAD) and N-terminal truncated PPAD (rPPAD<sup>Ntx</sup>) in *E coli*, only the full-length enzyme was citrullinated (figure 2A). Analogously, enzymatically inactive full-length PPAD (PPAD<sup>C351S</sup>) was not citrullinated when expressed in *E coli* (figure 2A, lower panel), confirming that PPAD autocitrullination is dependent on both the catalytic activity of the enzyme and its N-terminal domain. Importantly, we confirmed that the absence of autocitrullination in rPPAD<sup>Ntx</sup> was not due to loss of enzymatic function. Indeed, N-terminal truncated PPAD showed a significant increase in enzyme activity as compared to full-length rPPAD (mean citrulline production in  $\mu$ M: 191.9 vs. 88.9; \*p<0.0001) (figure 2B).

Mass spectrometry of full-length rPPAD achieved 96% protein sequence coverage, which included 17 of the 18 arginines found in the enzyme. Among these, 8 arginine residues were citrullinated (figure 2C). The probability of correct citrullination site localization approached 100% for all eight sites (figure 2D). These findings are in general accordance with the citrullination sites previously reported for rPPAD,<sup>12</sup> with two additional sites identified at R129 and R286. In peptides covering R63 and R70, citrullination was located to R63, but not confirmed at R70 as previously reported (figure 2D).<sup>12</sup> While this data confirms that rPPAD is autocitrullinated in *E coli*, this process appears to be unique to artificially expressed PPAD in the absence of physiological enzyme processing and regulators found in *P gingivalis*.

#### Anti-PPAD antibodies in RA are not directed against citrullinated PPAD

While we found no evidence that PPAD is citrullinated in *P gingivalis*, antibodies specific for citrullinated rPPAD have previously been reported in patients with RA.<sup>12</sup> To dissect the role of citrullination for anti-PPAD antibody recognition, we assayed serum samples of RA patients and healthy controls for antibodies to cit-rPPAD and uncitrullinated rPPAD<sup>C351S</sup> by ELISA. IgG anti-PPAD antibodies were common in patients with RA and in healthy controls (figure 3A). Strikingly, citrullinated rPPAD<sup>C351S</sup> did not differ significantly among RA patients (median 0.069 vs. 0.070; median of differences: 0.0; p=0.14) or controls (median 0.165 vs. 0.170; median of differences: -0.001; p=0.87) (figure 3A and online supplementary figure S2). Indeed, anti-cit-rPPAD and anti-rPPAD<sup>C351S</sup> antibody levels in RA were highly correlated (r=0.99, \*p<0.0001) (figure 3B). Control subjects demonstrated a similar degree of correlation (r=0.99, \*p<0.0001).

Importantly, levels of anti-cit-rPPAD were not significantly different between patients with RA and controls (median 0.069 vs. 0.165; p=0.085). Although the highest antibody reactivity was observed in the RA group (5/83 above the 95<sup>th</sup> percentile) compared to controls (1/39), statistical significance to support this association was not attained (Fisher's exact test; p=0.66). In fact, the overall trend observed was towards higher anti-PPAD antibody levels in the control group (figure 3A). To confirm the specificity of the ELISA assay for anti-PPAD, antibodies were additionally demonstrated by immunoprecipitation (IP) using radiolabeled PPAD generated by *in vitro* transcription/translation (figure 3C). In agreement with the ELISA assay, RA patient sera showed similar reactivity against PPAD and mutant PPAD<sup>C351S</sup> by IP (figure 3C, top and middle panel, respectively). Importantly, only sera reactive by ELISA immunoprecipitated radiolabeled PPAD (figure 3C, anti-PPAD + vs. anti-PPAD-). Similarly, radiolabeled truncated PPAD<sup>Ntx</sup> (figure 3C, bottom panel), a protein that better mimics PPAD in *P gingivalis*, was only precipitated by sera positive by ELISA. These data support the specificity of the ELISA assay and strongly suggest that anti-PPAD antibody binding in RA is independent of PPAD citrullination.

To confirm that citrullination is not a determinant in the antibody recognition of rPPAD, we further performed competition experiments. Antibody binding to cit-rPPAD was completely abrogated by preincubation of anti-cit-rPPAD positive RA sera with uncitrullinated rPPAD<sup>C351S</sup>, but not by preincubation with unmodified or citrullinated enolase (figure

4A,B). This demonstrates that anti-PPAD antibodies in RA are exclusively directed against unmodified PPAD.

#### Anti-PPAD antibodies do not correlate with anti-CCP levels or RA disease activity

To further characterize their relevance in RA, we inquired how anti-PPAD antibodies relate to anti-CCP levels and disease activity measures. Anti-CCP3 antibodies in RA did not correlate with anti-cit-rPPAD or anti-rPPAD<sup>C351S</sup> levels (r=0.07, p=0.56; r=0.09, p=0.44; respectively) (figure 5A). Moreover, we found no significant association between anti-rPPAD antibody levels and RA disease activity as measured using DAS28-CRP (r=-0.11, p=0.30) (figure 5B).

#### Anti-PPAD antibody levels are decreased in RA patients with PD

To elucidate the role of anti-PPAD antibodies in RA-associated PD, we investigated the PD status in RA and healthy controls (clinical characteristics summarized in Table 1). PD was more common among patients with RA compared to age and sex-matched controls (60/83; 72.3% vs. 9/39; 23.1%). Interestingly, anti-rPPAD antibody levels were significantly lower in RA patients with PD compared to periodontally healthy controls (median 0.058 vs. 0.162; \*p=0.046). Analogously, anti-rPPAD levels were decreased in PD-positive RA when compared to PD-negative RA patients (median 0.058 vs. 0.162). However, this analysis was underpowered and did not reach statistical significance (p=0.073) (figure 5C). Moreover, a significant decline in antibody levels with RA disease duration was observed in PD-positive RA (r=-0.28, \*p=0.03) (figure 5D), but not in the PD-negative RA group (r=-0.01, p=0.97; data not shown). Although the duration of PD in these groups is unknown, the negative association with RA-associated PD may suggest a protective role of anti-PPAD antibodies for periodontal health.

# DISCUSSION

PPAD autocitrullination has provided an intriguing framework to explain the loss of tolerance to citrullinated proteins in RA. In this study, we show that PPAD expressed by *P gingivalis* is not citrullinated, and demonstrate that PPAD citrullination is not recognized by anti-PPAD antibodies in RA. These findings have critical implications for the biology of PPAD and redefine the relevance of the anti-PPAD response in RA.

Despite the difference in molecular weight, cellular PPAD (75-85 kDa) and secreted PPAD (47 kDa) were indistinguishable in protein sequence. Mass spectrometry identified both as a truncated form of PPAD missing the first 43 N-terminal and 97 C-terminal amino acids. Full-length PPAD was not detected in our analysis, suggesting that similar to proforms of other *P gingivalis* virulence factors, this pro-enzyme is rapidly processed by truncation and concomitantly translocated to the bacterial cell surface.<sup>273940</sup> As such, glycosylation of cPPAD for membrane attachment may be sufficient to explain the striking difference in molecular weight to sPPAD,<sup>29</sup> as previously shown for arginine gingipain.<sup>41</sup> The truncation sites identified in this study may indicate specific processing by *P gingivalis* envelope-associated cysteine proteases with arginine-X (at R43) and lysine-X activity (at K459).<sup>42</sup> Truncation at R43 has previously been shown for sPPAD by N-terminal sequencing,<sup>9</sup> and

MS.<sup>27</sup> While we cannot exclude that truncation at K459 is an artifact of tryptic digestion, semitryptic peptides indicative of alternative cleavage sites were not detected.

Considering full-length rPPAD is readily autocitrullinated when expressed in  $E \ coli$ ,<sup>12</sup> the lack of citrullinated PPAD in P gingivalis is astounding. Unique strategies, which are absent in *E coli*, may have evolved in *P gingivalis* to prevent PPAD autocitrullination. In this study, we provide evidence that N-terminal processing may be the primary mechanism by which PPAD is protected from autocitrullination. Indeed, PPAD citrullination may be biologically unfavorable for P gingivalis. Rodríguez et al reported that PPAD activity declines with enzyme autocitrullination,<sup>30</sup> and citrullination of R352 adjacent to the active site of PPAD (C351) has been hypothesized to explain this reduction in enzyme activity.<sup>123036</sup> Mechanisms that block PPAD autocitrullination in *P gingivalis* may therefore conserve PPAD activity. The molecular basis by which N-terminal truncation abrogates PPAD autocitrullination is unclear. While PPAD is generally restricted to modify only free arginine and C-terminal arginine residues in cleaved proteins,<sup>91112</sup> PPAD autocitrullination involves citrullination of internal arginine residues in the intact enzyme. The N-terminal domain may therefore facilitate peptidylarginine autocitrullination by uniquely expanding the catalytic function of full-length PPAD. Rapid removal of the N-terminal domain by P gingivalis may render autocitrullination insufficient and aid bacterial survival in the periodontal pocket.

The finding that anti-PPAD antibodies in RA do not target citrullinated PPAD is consistent with the biology of PPAD identified in this study. However, our findings markedly contrast the conclusions reached by another study in which antibodies specific for citrullinated fulllength rPPAD were identified in 38% of RA patients.<sup>12</sup> The discrepancy between the two studies may be explained by differences in ELISA methodology, in patient cohorts studied, or in the rPPAD fusion proteins used. We tried to minimize error and exclude plate variability by measuring antibodies to cit-rPPAD, rPPAD<sup>C351S</sup> and background reactivity on the same plate. Using competition assays, we show that anti-PPAD antibodies in patients with RA are exclusively directed against the unmodified enzyme. We cannot fully exclude major differences in the populations studied. Lastly, there is a significant difference in the recombinant proteins used in our study. Previously, GST-His-tagged rPPAD was used to screen for anti-PPAD antibodies.<sup>12</sup> The GST tag (25 kDa) adds 9 additional arginine residues to the fusion protein, which may potentially be citrullinated in the process of GST-His-rPPAD autocitrullination. By contrast, rPPAD in our study was designed without GST tag and only contains one additional arginine in the His-tag used (2.5 kDa). Thus, we cannot exclude that the antibodies identified in the previous study may in fact target citrullinated GST in the fusion protein. Such antibodies would not be identified when screening against enzymatically inactive GST-His-rPPAD. Nonetheless, the existence of citrullinated-PPADspecific antibodies in RA cannot be supported, and appears implausible in the absence of PPAD citrullination in P gingivalis.

While anti-PPAD antibodies are common in patients with RA (and healthy controls), the significance of the antibody response against this bacterial virulence factor for RA-associated PD is not clear. In our study, anti-PPAD antibodies did not correlate with disease activity or anti-CCP levels, reinforcing that anti-PPAD antibodies are not part of the ACPA-response. Antibodies to *P gingivalis* cell extracts have been reported to correlate with anti-

CCP antibodies and disease activity measures in RA,<sup>1943–45</sup> suggesting that anti-PPAD antibodies may represent a unique antibody population. Despite this association, anti-*P* gingivalis antibodies do not distinguish RA from controls cases,<sup>45</sup> and have no predictive value for the development of RA in seropositive arthralgia patients.<sup>46</sup> In this context, it is compelling that anti-PPAD levels are significantly lower in RA with PD compared to periodontally healthy controls. Anti-PPAD antibodies may have a protective role for PD in RA, and clearance of PPAD may decrease survival of *P* gingivalis in the periodontal pocket by limiting ammonia production. The finding that antibodies against major *P* gingivalis virulence factors confer protection in murine models of periodontal infection and potentially human disease further supports this possibility.<sup>1847–50</sup>

Our findings suggest that PPAD autocitrullination is not the underlying mechanism linking *P gingivalis*-associated PD and RA. Anti-PPAD antibodies may be useful markers to predict the risk of PD progression in patients with RA.

# **Supplementary Material**

Refer to Web version on PubMed Central for supplementary material.

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#### Figure 1.

Cellular PPAD (cPPAD) and secreted PPAD (sPPAD) from *P gingivalis* are truncated and not citrullinated. (A) *P gingivalis* cell lysate (CL) and culture supernatant (CS) were analyzed by SDS-PAGE. Ponceau staining before antibody probing is shown to visualize loading (left panel). Anti-PPAD immunoblotting shows cPPAD (black arrowhead) and sPPAD (white arrowhead) in *P gingivalis* CL and CS, respectively (middle panel). AMC immunoblotting was used to detect citrullination in bacterial samples (right panel). (B and C) Mass spectrometry sequence coverage map of *P gingivalis* cPPAD (B) and sPPAD (C). Peptide sequences identified are shown in blue; the N-terminal and C-terminal domains missing in cPPAD and sPPAD are shown in grey. Arginine residues are underlined. Numbered arrows indicate possible PPAD truncation sites at R43 and K459.



#### Figure 2.

Citrullination of rPPAD expressed in *E coli* is dependent on the N-terminal domain. (A) Purified rPPAD, mutant rPPAD<sup>C351S</sup> (C351S) and truncated rPPAD missing the N-terminal domain (Ntx) were analyzed by SDS-PAGE and visualized by anti-PPAD immunoblotting (top panel). AMC immunoblotting (bottom panel) was used to detect citrullination. (B) Enzymatic activity of rPPAD, rPPAD<sup>C351S</sup> and rPPAD<sup>Ntx</sup> was measured in the BAEE assay. Average citrulline concentrations (in  $\mu$ M) from two independent experiments are shown (error bars indicate standard deviation). Enzyme activities were analyzed by two-way ANOVA (rPPAD<sup>Ntx</sup> vs. rPPAD; \*p<0.0001). (C) Mass spectrometry sequence coverage map of full-length rPPAD expressed in *E coli*. Peptide sequences identified by MS are shown in blue; confirmed citrullination sites in pink. Arginine residues are underlined. (D) Localization probabilities for individual arginine deimination sites in rPPAD were calculated using the Ascore algorithm. An Ascore of 20 (99% certainty; p=0.01) was considered significant.<sup>38</sup>



# Figure 3.

Antibodies to citrullinated rPPAD (cit-rPPAD) and uncitrullinated PPAD (rPPAD<sup>C351S</sup>) in serum of patients with RA and controls. (A) RA patients (n=83) and controls (n=39) were assayed for antibodies to cit-rPPAD (full circles) and rPPAD<sup>C351S</sup> (empty circles) by ELISA. Antibody reactivity is expressed as arbitrary units (AU). Statistical analysis was performed using the Wilcoxon matched-pairs signed-rank test (anti-citPPAD vs. anti-PPAD<sup>C351S</sup>) and Mann-Whitney test (RA vs. control subjects). The red line represents the median reactivity for individual groups. ns: not significant. (B) Correlation of anti-cit-rPPAD and anti-rPPAD<sup>C351S</sup> antibody levels in patients with RA (r=0.99, p<0.0001). (C) <sup>35</sup>S-methionine-labeled PPAD (top panel), PPAD<sup>C351S</sup> (middle panel) and truncated PPAD<sup>Ntx</sup> (bottom panel) generated by IVTT were immunoprecipitated using RA patient serum. Immunoprecipitates were electrophoresed on SDS-PAGE and visualized by fluorography. Representative results for anti-rPPAD positive (anti-PPAD+) and anti-rPPAD negative (anti-PPAD-) patients, as determined by ELISA, are shown.



#### Figure 4.

Anti-PPAD antibodies in RA are exclusively directed against uncitrullinated PPAD. (A) Antibody reactivity against cit-rPPAD was measured after preincubation of anti-cit-rPPAD positive RA sera (n=8) with buffer alone (column 1), purified enolase (column 2), citrullinated enolase (cit-enolase) (column 3) or uncitrullinated rPPAD<sup>C351S</sup> (column 4; \*\*\*\*p<0.001). Comparisons were made using Friedman's test/ Dunn's post test for matched groups. Column height shows mean antibody reactivity expressed as arbitrary units (AU); bars indicate SEM. ns: not significant. (B) AMC immunoblotting to demonstrate the presence or absence of citrullination in proteins used in A. Purified cit-rPPAD, rPPAD<sup>C351S</sup>, enolase and cit-enolase were resolved by SDS-PAGE, transferred onto nitrocellulose and visualized by Ponceau staining as a loading control (top panel). Protein citrullination in the same membrane was detected by AMC immunoblotting (bottom panel).



#### Figure 5.

Anti-PPAD levels do not correlate with anti-CCP antibodies or RA disease activity, but are negatively associated with periodontal disease in RA. (A) Correlation of anti-PPAD antibodies with anti-CCP3 levels in RA. Antibody levels to CCP3 were measured by ELISA (QUANTA Lite IgG, INOVA). (B) Correlation of anti-PPAD levels with RA disease activity as measured using DAS28-CRP. ns: not significant. (C) Anti-PPAD antibody levels in RA patients with and without periodontal disease (*RA PD* and *RA No PD*, respectively; full circles) compared to periodontally healthy controls (empty circles). Antibody levels are expressed in arbitrary units (red line shows median reactivity; \*p<0.05). (D) Antibody levels in PD-positive RA patients as a function of RA disease duration in years.

#### Table 1

Demographic and clinical characteristics of rheumatoid arthritis patients and controls

| Parameter                                    | RA (83)       | Controls (39) | <b>p</b> * |
|--|---------------|---------------|------------|
| Female (%)                                   | 79.5 (66)     | 82.1 (32)     | 0.74       |
| Mean age in years (± SD)                     | 50.7 (± 12.9) | 48.1 (± 13.3) | 0.31       |
| Race / ethnicity                             |               |               | 0.00       |
| Caucasian (%)                                | 74.7 (62)     | 33.3 (13)     |            |
| African-American (%)                         | 16.9 (14)     | 35.9 (14)     |            |
| Smoking (%) $^{\dot{\tau}}$                  | 16.9 (14)     | 12.8 (5)      | 0.56       |
| Periodontal disease (%)                      | 72.3 (60)     | 23.1 (9)      | 0.00       |
| Mean RA disease duration in years $(\pm SD)$ | 6.3 (± 7.9)   |               |            |
| Seropositive for IgM RF (%)                  | 68.5 (50)     |               |            |
| Seropositive for IgG anti-CCP3 (%)           | 75.9 (63)     |               |            |
| Mean CRP in mg/L (± SD)                      | 6.9 (± 11.3)  |               |            |
| Mean DAS28-CRP (± SD)                        | 3.5 (± 1.4)   |               |            |
| Mean CDAI (± SD)                             | 15.6 (± 14.0) |               |            |

RF, rheumatoid factor, cut-off level for positivity 20 IU; anti-CCP3, anti-cyclic citrullinated peptide antibody (third-generation), cut-off level for positivity 20 units; CRP, C-reactive protein; DAS28-CRP, Disease Activity Score 28-CRP; CDAI, Clinical Disease Activity Index.

\*Comparison of parameters reflected in percentages: chi-square test for proportions, comparison of means: independent-samples t-test; SD, standard deviation.

 $^{\dagger}$ Current cigarette smoking