



Roxithromycin attenuates inflammation via modulation of RAGE-influenced calprotectin expression in a neutrophilic asthma model

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Background: Roxithromycin (RXM), a macrolide antibiotic, exhibits anti-asthmatic effects, but its specific mechanism of action remains elusive. We evaluated the effects of RXM on airway inflammation, the expression of calprotectin, and the activity of the receptor of advanced glycation end products (RAGE) to determine whether RXM alleviates inflammation by regulating RAGE activation, and thereby calprotectin expression, in neutrophilic asthma.

Methods: Male Brown Norway rats were sensitized with ovalbumin (OVA) and Freund's complete adjuvant (FCA) mixture, followed by OVA challenge to induce neutrophilic asthma. RXM (30 mg/kg) or FPS-ZM1 (RAGE inhibitor, 1.5 mg/kg) was administered 30 min prior to each challenge. The infiltration of airway inflammatory cells and cytokines, as well as the expression of calprotectin and RAGE, was assessed.

Results: The expression of airway inflammatory cells and cytokines was found to be significantly elevated in OVA + FCA-induced rats. Increased expression of both calprotectin and RAGE was also detected in OVA + FCA-induced asthma [bronchoalveolar lavage fluid (BALF) calprotectin: 15.07±1.79 vs. 3.86±0.69 ng/mL; serum calprotectin: 20.47±1.64 vs. 9.29±1.31 ng/mL; lung tissue homogenates calprotectin: 28.82±1.01 vs. 12.02±1.38 ng/mg; BALF RAGE: 762.93±36.47 vs. 294.25±45.92 ng/mL; serum RAGE: 906.43±58.95 vs. 505.60±30.16 ng/mL; lung tissue homogenates RAGE: 1,585.24±177.59 vs. 461.53±63.40 ng/mg; all P<0.001]. However, all of these changes were interrupted by RXM and FPS-ZM1.

Conclusions: RXM exerted similar effects as the RAGE inhibitor FPS-ZM1 in terms of reducing airway inflammation and downregulating the expression of calprotectin and RAGE in a neutrophilic asthma model. Our findings provide novel insights into the mechanisms underlying the effect of RXM pretreatment on neutrophilic asthma. Furthermore, FPS-ZM1 may be useful as an intervention specific to the neutrophilic asthma phenotype.

Keywords: Calprotectin; receptor of advanced glycation end products (RAGE); roxithromycin (RXM); asthma; airway inflammation

Submitted Jan 12, 2021. Accepted for publication Mar 20, 2021.

doi: 10.21037/atm-21-859

View this article at: <http://dx.doi.org/10.21037/atm-21-859>

Introduction

Asthma is a heterogeneous disease with various phenotypes, and is mainly characterized by airway inflammation and airway hyperresponsiveness (1). Increasing evidence indicates that airway neutrophil infiltration in asthma is associated with disease severity and acute asthma exacerbation (2). Therefore, having safe and effective treatments for neutrophilic asthma is crucial.

MRP8 and MRP14, members of the S100 protein family, exist as the protein heterodimer calprotectin (also called S100A8/A9). Calprotectin is predominantly expressed by activated neutrophils, and comprises >30% of neutrophil cytoplasmic proteins (3). Recently, it was identified as an important damage-associated molecular pattern (DAMP) via the receptor of advanced glycation end products (RAGE) and Toll-like receptor 4 (TLR4) (4,5). Emerging evidence suggests that calprotectin is a useful biomarker of inflammatory processes, including rheumatoid arthritis, juvenile idiopathic arthritis, and inflammatory bowel disease (6). The recent study has shown that calprotectin levels in patients with asthma correlated negatively with forced expiratory volume in one second/forced vital capacity, body mass index, and positively with smoke amount and blood neutrophil percentage (7). In addition, calprotectin has a pivotal role in inflammation by inducing the secretion of inflammatory cytokines (8,9). Owing to its critical involvement in inflammatory reactions, calprotectin has received significant attention in studies of asthma (10,11).

RAGE, a multi-ligand receptor belonging to the immunoglobulin receptor superfamily, is an important transmembrane protein (12). There is a growing body of evidence implicating RAGE, via the binding of its ligands, including calprotectin, as a major driving mechanism in the pathogenesis of inflammatory states (13–15). Furthermore, one study showed that RAGE and its ligands may be potential hallmarks for inflammatory disorder progression and severity (16). Nevertheless, little has been revealed about RAGE's role in asthma to date.

It has been shown that macrolides appear to be beneficial in the prevention of myocardial remodeling via matrix metalloproteinase (MMP) and nitric oxide (NO) regulation, which indicates that macrolides may be alternative therapies for suppressing inflammatory remodeling (17,18). Roxithromycin (RXM), a broad-spectrum macrolide antibiotic, enhances phagocytosis and the bactericidal activities of neutrophils (19). Studies demonstrated that RXM ameliorates colitis by suppressing oxidative stress and downregulating NF- κ B-

mediated proinflammatory signaling (20,21). Furthermore, it has become increasingly evident that RXM has efficacy in reducing the risk of asthma exacerbations (22,23); however, the mechanisms underlying this effect are not yet fully understood.

In this study, we established a neutrophilic asthma model to investigate the effects of RXM and its underlying mechanisms. To this end, we hypothesized that RXM may alleviate airway inflammation via inhibition of RAGE, which modulates the expression of calprotectin.

We present the following article in accordance with the ARRIVE reporting checklist (available at <http://dx.doi.org/10.21037/atm-21-859>).

Methods

Animals and grouping

Male Brown Norway rats, aged 6–8 weeks and weighing 150–180 g, were obtained from the Beijing Vital River Laboratory Animal Center (Beijing, China) and allowed to acclimate to their environment for 7 days. The rats were kept in a pathogen-free environment at 25 °C with a 12-h light/dark cycle, and were allowed access to food and water ad libitum. Experiments were performed under a project license (protocol wyd2019-019) granted by the Animal Care and Protection Committee of Wenzhou Medical University, in compliance with the Animal Care and Protection Committee of Wenzhou Medical University guidelines for the care and use of animals. The rats were randomly assigned to four groups (each comprising six rats): control, asthma, asthma + FPS-ZM1, asthma + RXM.

Neutrophilic asthma model and treatments

Our neutrophilic asthma model was established according to the improved method of Dejager *et al.* (24). Briefly, the rats were sensitized with an intraperitoneal (i.p.) injection of 0.5 mg ovalbumin (OVA, Sigma-Aldrich, USA) emulsified in 0.5 mL Freund's complete adjuvant (FCA, Sigma-Aldrich, USA) in 0.5 mL normal saline (NS) on days 1 and 8. From days 15–19, the sensitized rats were challenged with 1% OVA aerosol for 30 min once daily. Rats in the asthma + RXM group had intragastric (i.g.) administration of RXM (MedChemExpress, USA; 30 mg/kg), and rats in the asthma + FPS-ZM1 group had i.g. administration of FPS-ZM1 (SelleckChem, USA; 1.5 mg/kg) 30 min before the challenge. Rats in the control group were sham-sensitized and challenged with NS (*Figure 1A*).

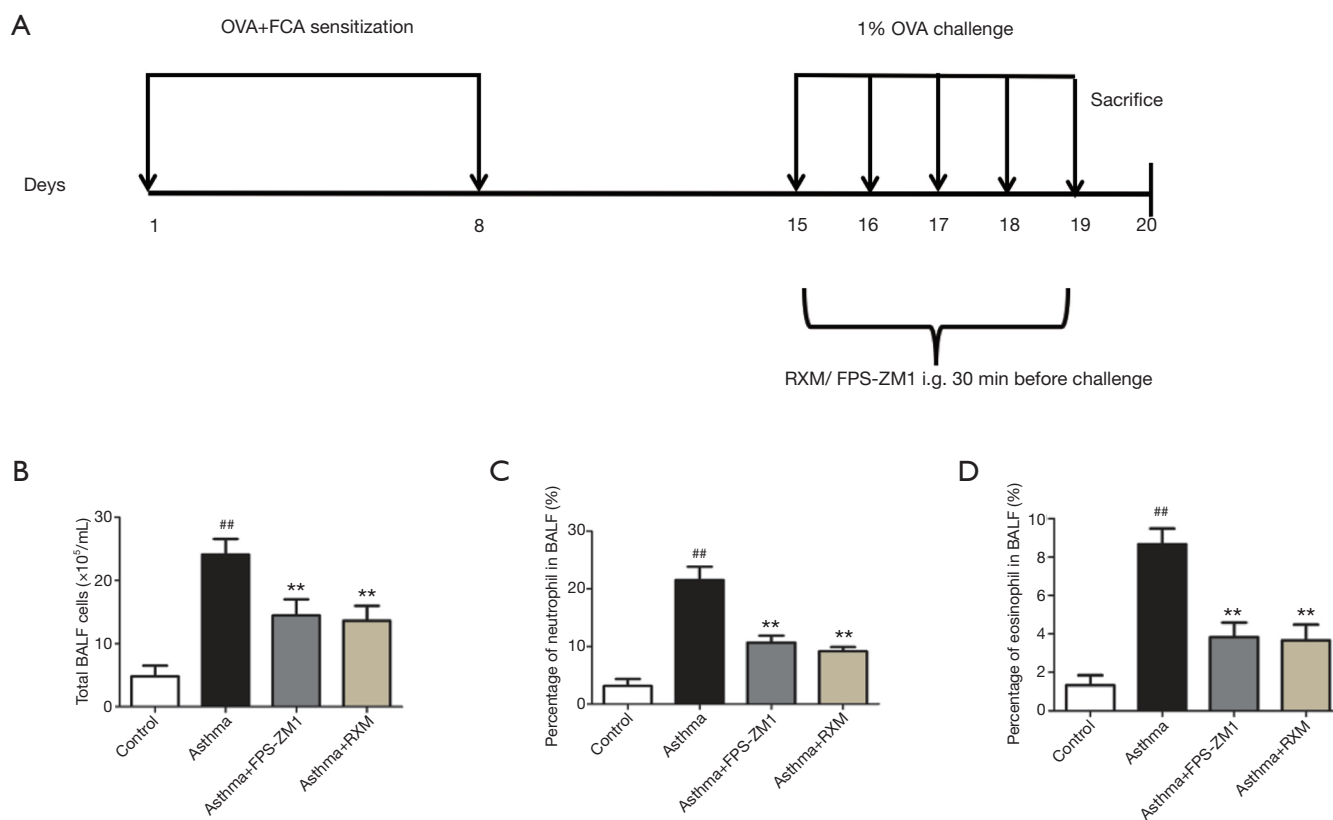


Figure 1 Effects of roxithromycin (RXM) on bronchoalveolar lavage fluid (BALF) cells of ovalbumin (OVA)+ Freund's complete adjuvant (FCA)-induced rats. (A) Experimental protocols for the neutrophilic asthma model and treatments. Total cells (B), percentage of neutrophil (C), and percentage of eosinophil (D) in BALF. Data are expressed as mean \pm standard deviation (SD) (n=4–6 per group). ^{##}, P<0.01 compared to the control group; ^{**}, P<0.01 compared to the asthma group.

Collection of serum and bronchoalveolar lavage fluid (BALF)

The rats were anesthetized and humanely killed 24 h after the last challenge. Blood samples were collected from the abdominal aorta and the supernatants were centrifuged at 1,006 g for 15 min and stored at -80°C . The rats' left lungs were washed with saline solution via a tracheal cannula, and the fluid collected was centrifuged at 112 g for 15 min at 4°C . The supernatants were kept as BALF and stored at -80°C . The cell pellet was resuspended in saline solution using a hemocytometer (Bio-Rad, USA) to count the total number of cells. Wright's staining was performed to obtain differential cell counts.

Histopathological analysis

The rats' right lungs were fixed in 4% paraformaldehyde

and stained with hematoxylin and eosin (H&E) for light microscopy (Leica, Germany). The inflammation score was evaluated by a sample-blinded pathologist using the H&E-stained sections (25).

Enzyme-linked immunosorbent assay

The concentrations of calprotectin, RAGE, interleukin (IL)-17, and IL-6 in BALF and serum, and the concentrations of calprotectin and RAGE in lung tissues, were quantified using enzyme-linked immunosorbent assay (ELISA) kits (Absin, China) according to the manufacturer's protocols.

Western blotting analysis

The isolated lungs were flash-frozen in liquid nitrogen, then homogenized and sonicated in ice-cold RIPA lysis buffer (Beyotime, China) containing protease and

phosphatase inhibitor (Applygen, China) to obtain protein extracts. A total of 50 µg of protein was loaded onto 10% or 12% sodium dodecyl sulfate (SDS)-polyacrylamide gels for electrophoresis (SDS-PAGE) and transferred onto polyvinylidene difluoride (PVDF) membranes (Millipore, USA). Membranes were blocked with Tris-buffered saline containing 5% non-fat dried milk for 2 h prior to incubation with primary antibodies against RAGE (Santa Cruz, USA) and GAPDH (CST, USA), respectively, overnight at 4 °C. The membranes were incubated with HRP-conjugated anti-mouse or anti-rabbit secondary antibodies (Biosharp, China) for 1.5 h at room temperature. After treatment with enhanced chemiluminescent reagents (Pierce, USA), the target protein was quantitatively analyzed using Image Lab software (Bio-Rad, USA) (target protein gray value/GAPDH gray value).

Statistical analysis

The data are expressed as mean ± standard deviation (SD). Comparisons among groups were performed using one-way analysis of variance followed by Tukey's (equal variances assumed) or Dunnett's (equal variances not assumed) post-hoc multiple comparisons. Correlation analysis was conducted using Pearson's correlation coefficient. SPSS 21.0 software (SPSS, USA) was used for statistical analyses, and $P < 0.05$ was considered statistically significant.

Results

Effect of RXM on airway inflammatory cells in BALF

Our results revealed a notable increase in the total number of inflammatory cells in OVA + FCA-induced rats relative to controls [(24.12 ± 2.49) vs. (4.82 ± 1.68) × 10⁵/mL, $P < 0.001$, *Figure 1B*], especially in the percentage of neutrophils (21.50% ± 2.35% vs. 3.17% ± 1.17%, $P < 0.001$, *Figure 1C*). The percentage of eosinophils was also increased in OVA + FCA-induced rats relative to controls (8.67% ± 0.82% vs. 1.33% ± 0.52%, $P < 0.001$, *Figure 1D*). Treatment with RXM significantly reduced these effects compared with the asthma group [(13.63 ± 2.36) vs. (24.12 ± 2.49) × 10⁵/mL; 9.17% ± 0.75% vs. 21.50% ± 2.35%; 3.67% ± 0.82% vs. 8.67% ± 0.82%; all $P < 0.001$, *Figure 1*].

Effect of RXM on infiltration of airway inflammatory cells

Relative to the controls, OVA + FCA-induced rats

developed evidently narrowed airways, increased mucosal wrinkles, and extensive inflammatory cell infiltration and aggregation in the bronchi, perivascular, and alveolar spaces (*Figure 2A*). The RXM treatment suppressed these histopathological responses, demonstrating more anti-inflammatory activity relative to the asthma group (*Figure 2*).

Effect of RXM on inflammatory cytokines in BALF and serum

The concentrations of IL-17 and IL-6 in the BALF (136.31 ± 6.18 vs. 32.65 ± 9.17 ng/mL; 69.60 ± 6.02 vs. 10.62 ± 3.83 ng/mL; all $P < 0.001$) and serum (185.76 ± 18.82 vs. 89.15 ± 15.24 ng/mL; 99.16 ± 9.60 vs. 46.65 ± 5.59 ng/mL; all $P < 0.001$) were markedly elevated in the asthma group compared with controls (*Figure 3A,B*). Of note, all of these elevated expression levels were reduced by RXM compared with the asthma group (108.12 ± 8.28 vs. 136.31 ± 6.18 ng/mL, $P = 0.002$; 39.07 ± 6.95 vs. 69.60 ± 6.02 ng/mL, $P < 0.001$; 138.42 ± 15.13 vs. 185.76 ± 18.82 ng/mL, $P = 0.007$; 64.81 ± 4.65 vs. 99.16 ± 9.60 ng/mL, $P < 0.001$, *Figure 3*).

Effect of RXM on expression of calprotectin in OVA + FCA-induced rats

Calprotectin plays a critical role in the inflammatory response (26), so we examined its expression in BALF, serum, and lung tissue. Our results revealed a notable increase in the concentration of both BALF and serum calprotectin in the asthma group relative to the control group (15.07 ± 1.79 vs. 3.86 ± 0.69 ng/mL; 20.47 ± 1.64 vs. 9.29 ± 1.31 ng/mL; all $P < 0.001$, *Figure 3C*). Similar results were found according to the ELISA results for the lung tissue homogenates (28.82 ± 1.01 vs. 12.02 ± 1.38 ng/mg, $P < 0.001$, *Figure 4A*). In contrast, RXM reversed the expression pattern of calprotectin compared with the asthma group (11.21 ± 0.76 vs. 15.07 ± 1.79 ng/mL, $P = 0.001$; 14.36 ± 0.78 vs. 20.47 ± 1.64 ng/mL, $P < 0.001$; 16.94 ± 1.40 vs. 28.82 ± 1.01 ng/mg, $P < 0.001$, *Figures 3C, 4A*).

Involvement of RAGE in the effect of RXM on OVA + FCA-induced changes in calprotectin expression

As illustrated in *Figure 4B,C*, RAGE was upregulated after OVA + FCA exposure (1.06 ± 0.05 vs. 0.18 ± 0.05, $P < 0.001$). ELISA analysis also showed high levels of RAGE in BALF, serum, and lung tissue compared with controls (762.93 ± 36.47 vs. 294.25 ± 45.92 ng/mL; 906.43 ± 58.95 vs.

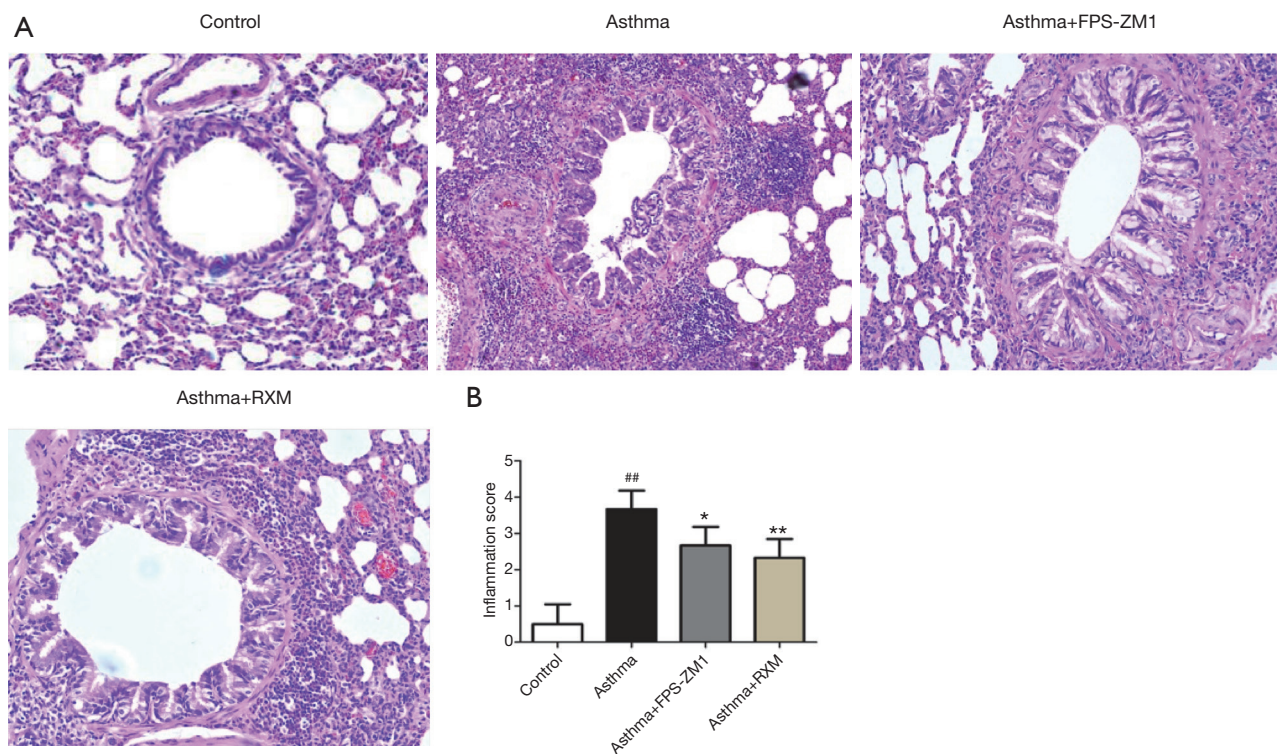


Figure 2 Effects of roxithromycin (RXM) on lung histopathology of ovalbumin (OVA)+ Freund's complete adjuvant (FCA)-induced rats. Hematoxylin and eosin (H&E) staining of lung tissues at 200 × magnification (A). The inflammation score of each group (B). The data are expressed as mean ± standard deviation (SD) (n=4–6 per group). ##, P<0.01 compared to the control group; *, P<0.05 compared to the asthma group; **, P<0.01 compared to the asthma group.

505.60±30.16 ng/mL; 1,585.24±177.59 vs. 461.53±63.40 ng/mg; all P<0.001, *Figures 3D,4D*); RXM treatment significantly inhibited this elevation (559.70±64.32 vs. 762.93±36.47 ng/mL; 649.85±61.33 vs. 906.43±58.95 ng/mL; 860.30±134.56 vs. 1,585.24±177.59 ng/mg; all P<0.001). To further investigate the mechanisms of RXM intervention in calprotectin expression and neutrophilic asthma, we used the FPS-ZM1 group for comparison. As expected, FPS-ZM1 suppressed calprotectin expression and reduced airway inflammation, further supporting a preeminent role for RAGE in the pathogenesis of neutrophilic asthma.

Correlative analysis of various parameters

The ELISA revealed that the concentration of calprotectin correlated positively to the concentration of RAGE (r=0.9611, 0.9063, 0.9353; all P<0.001; *Figures 4E,5A,B*). The concentration of calprotectin correlated positively to IL-17 in both serum and BALF (r=0.8674, 0.9670; all

P<0.001; *Figure 5C,D*), and it also correlated positively to IL-6 in both serum and BALF (r=0.9037, 0.9362; all P<0.001; *Figure 5E,F*). Moreover, there was a positive correlation between calprotectin and both total BALF cells and total neutrophils (r=0.9096, 0.8760; all P<0.001; *Figure 5G,H*).

Discussion

Neutrophils and their harmful secretions are well recognized as one of the reliable clinical biomarkers of lung disease progression (27). Pronounced airway neutrophil infiltration is a central feature of neutrophilic asthma, which is a major cause of both severe and refractory asthma (1,2). Based on our earlier findings, we developed a neutrophilic asthma model induced by OVA and FCA. Our results revealed airway hyperresponsiveness and a notable increase in neutrophil-predominant inflammatory cells in the OVA + FCA-induced rats. Neutrophil infiltration is known to involve a vast array of chemokines and cytokines. IL-17

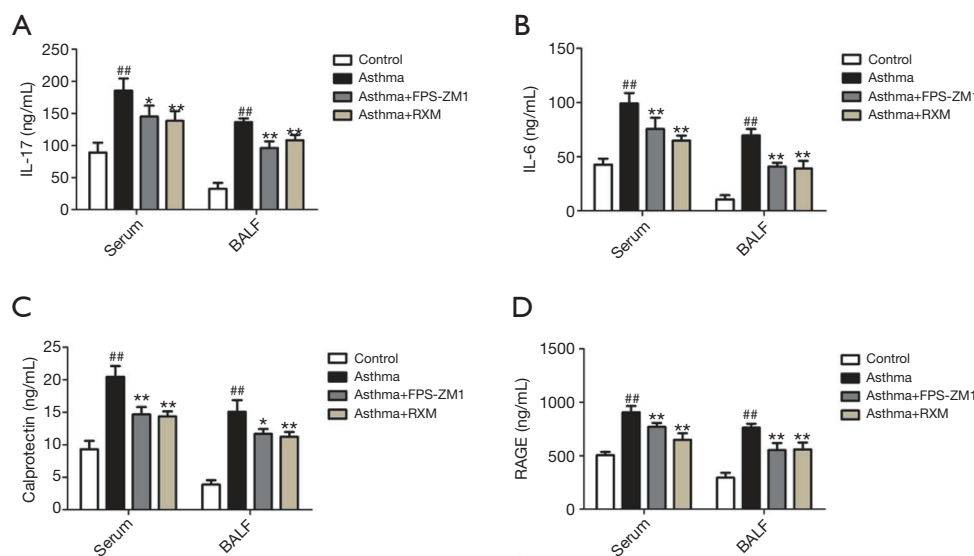


Figure 3 Effects of roxithromycin (RXM) on inflammatory cytokines, calprotectin and the receptor of advanced glycation end products (RAGE) in serum and bronchoalveolar lavage fluid (BALF). The concentrations of interleukin (IL)-17 (A), IL-6 (B), calprotectin (C), and RAGE (D) in serum and BALF by Enzyme-linked immunosorbent assay (ELISA). Results were obtained from three independent experiments. The data are expressed as mean \pm standard deviation (SD) (n=4–6 per group). ##, P<0.01 compared to the control group; *, P<0.05 compared to the asthma group; **, P<0.01 compared to the asthma group.

and IL-6, among other cytokines, are crucial mediators of neutrophil recruitment and activation (28,29). Previous research has demonstrated that the airway tissues of asthmatic subjects with severe and persistent symptoms are characterized by preferential neutrophilic infiltration with elevated levels of both IL-17 and IL-6 (28–30). In addition to the increased airway neutrophils, our current study detected upregulation of IL-17 and IL-6 in both the serum and BALF from the asthmatic rats, which corroborated our model of neutrophil-mediated inflammation. Our primary finding was that pretreatment with RXM alleviated airway inflammation and modulated the expression of calprotectin and RAGE in neutrophilic asthmatic rats. Further, we found that blocking RAGE activation suppressed calprotectin expression and weakened the inflammatory response.

Calprotectin, as one of the DAMPs belonging to the family of Ca^{2+} -binding proteins, has been identified as a critical component of inflammatory diseases, including asthma (8,10). Differential proteomic analysis of BALF from asthmatic subjects has revealed that calprotectin is strongly linked to the pathogenesis of airway inflammation (31). The expression of calprotectin in lung tissue is significantly elevated during the early stage of OVA-induced asthma, so calprotectin may be involved in the initiation of the early

inflammatory response in asthma (9). Moreover, proteomic identification of $\text{MMP-2}^{-/-}/\text{MMP-9}^{-/-}$ asthmatic mice demonstrated that calprotectin was upregulated and that function-blocking antibodies to calprotectin meaningfully inhibited the migration of inflammatory cells into the alveolar space. This finding indicates that calprotectin induces chemotaxis and migration of airway inflammatory cells in asthma (32).

Recent evidence has shown that calprotectin exerts its proinflammatory functions mainly by binding to the pattern recognition receptor (PRR) RAGE and thus activating inflammation-related downstream signaling pathways and upregulating the expression of proinflammatory cytokines (33). Another study showed that calprotectin could be upregulated with a high AGE burden, thereby aggravating proinflammatory conditions via the activation of RAGE; calprotectin expression and proinflammatory conditions were significantly suppressed by RAGE antagonism (34). In our previous experiment, we found that the concentration of S100A8 in asthmatic rats was notably increased, with relatively higher levels of RAGE (35). In this study, we added the FPS-ZM1 group, to further verify that the significant elevation of calprotectin in our neutrophilic asthma model positively correlated with the concentration of RAGE and airway inflammation. Most importantly, all of these

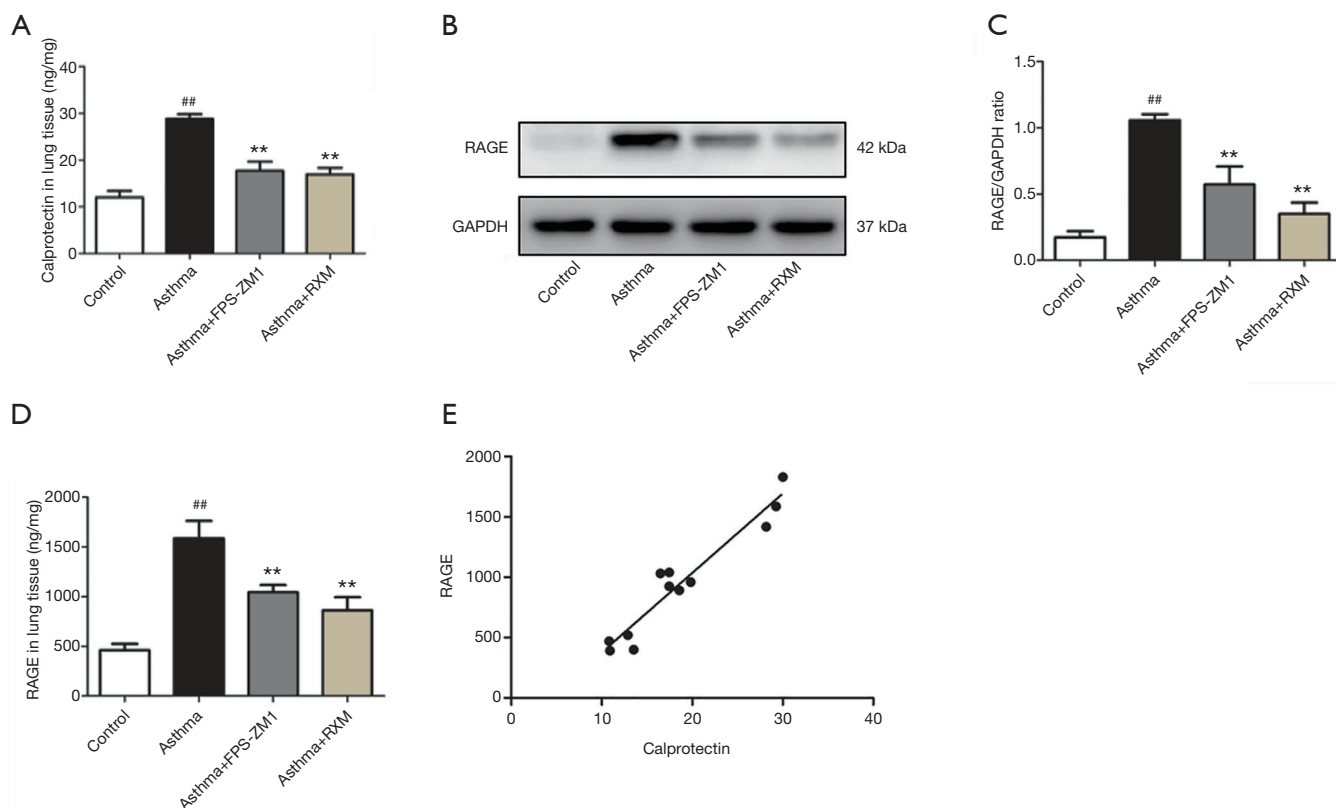


Figure 4 Effects of roxithromycin (RXM) on calprotectin and the receptor of advanced glycation end products (RAGE) in lung tissue. The concentration of lung tissue calprotectin (A) by Enzyme-linked immunosorbent assay (ELISA). The expression of RAGE in lung tissue was analyzed by Western blotting (B) and normalized to the levels of glyceraldehyde-3-phosphate dehydrogenase (GAPDH) (C). The concentration of lung tissue RAGE (D) by ELISA. The correlation between RAGE and calprotectin in lung tissue ($r=0.9611$; E). Results were obtained from three independent experiments. Data are expressed as mean \pm standard deviation (SD) ($n=4-6$ per group). ^{##}, $P<0.01$ compared to the control group; ^{**}, $P<0.01$ compared to the asthma group.

changes were alleviated by FPS-ZM1. These findings are in accordance with the results from the other studies discussed above.

As a PRR, RAGE has emerged as a key regulator of inflammatory processes (36), and it is strongly expressed in the lungs (12). Genome-wide association studies have identified RAGE as an important player in the pathogenesis of human asthma (37). Clinical studies have suggested an increased level of RAGE is present in sputum samples from asthmatic subjects; such levels correlate with disease severity (38). In a house dust mite asthmatic model, RAGE-knockout mice were devoid of most of the pathological features, including airway hypersensitivity, airway inflammation, and airway remodeling, that are found in wild-type mice. Furthermore, these findings were reproduced in an OVA-induced asthmatic model, which

demonstrated the fundamental importance of RAGE in asthma (39). Additional analysis revealed that RAGE promoted IL-33 expression and mediated its downstream inflammatory signaling effects in response to asthma; the absence of RAGE impeded this reaction (40). In the present study, OVA + FCA-induced rats showed elevated RAGE expression levels, and RAGE inhibition attenuated OVA + FCA-induced airway inflammation, which confirms the pronounced significance of RAGE in neutrophilic asthma.

Given the inefficacy of corticosteroids in neutrophilic asthma, considerable interest has been given to potential alternative therapeutics. Because infections with bacteria and viruses are moderately associated with asthma severity (41), antibiotics, especially macrolides, have shown promise, as presented in decreased sputum neutrophil counts and alleviation of symptoms (42-44). Our previous study found

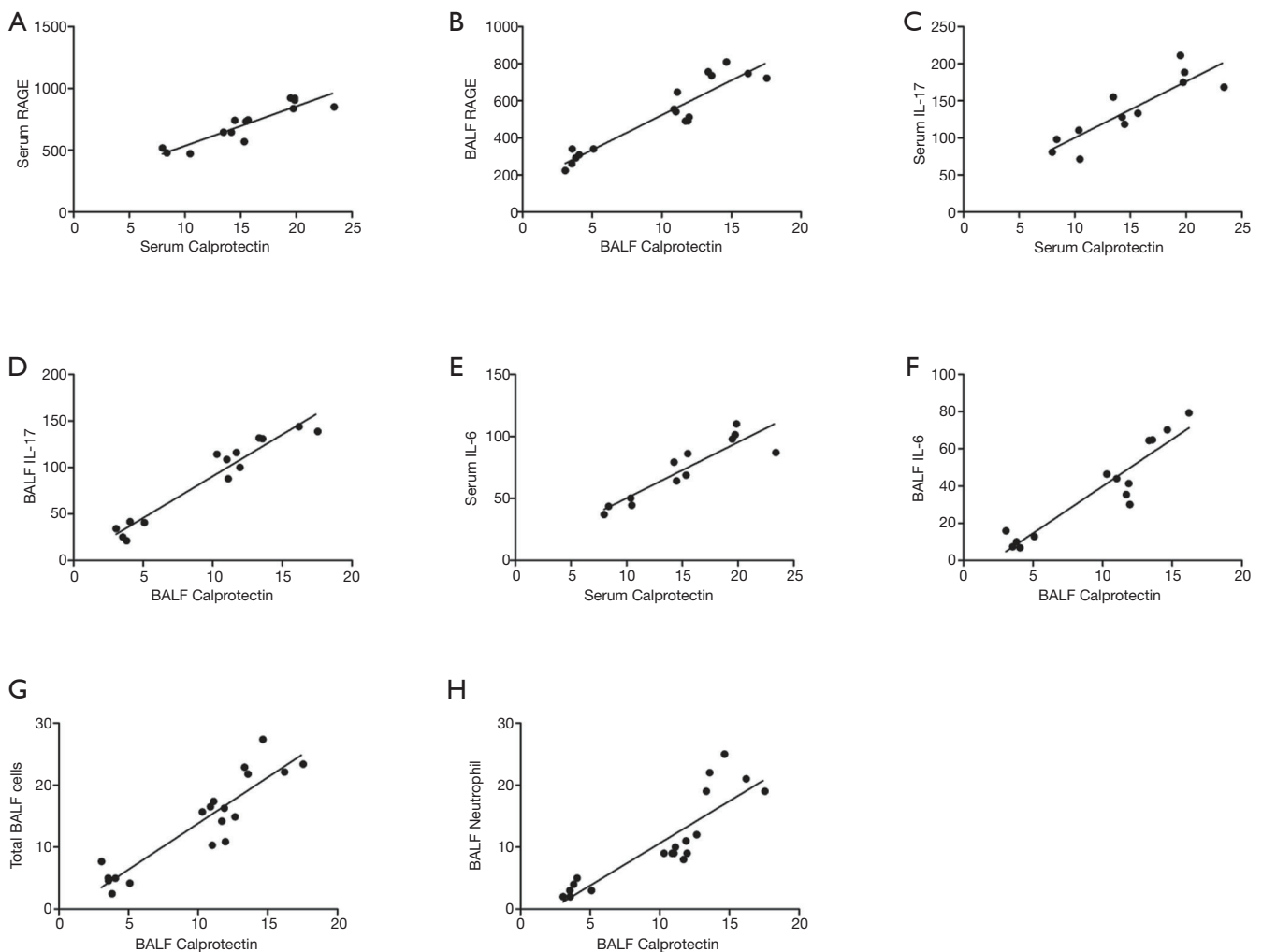


Figure 5 The correlation among different parameters. The correlation between the receptor of advanced glycation end products (RAGE) and calprotectin in serum ($r=0.9063$, A) and bronchoalveolar lavage fluid (BALF) ($r=0.9353$, B). The correlation between interleukin (IL)-17 and calprotectin in serum ($r=0.8674$, C) and BALF ($r=0.9670$, D). The correlation between IL-6 and calprotectin in serum ($r=0.9037$, E) and BALF ($r=0.9362$, F). The correlation between total inflammatory cells and calprotectin in BALF ($r=0.9096$, G). The correlation between neutrophils and calprotectin in BALF ($r=0.8760$, H). $n=4-6$ per group, all $P<0.01$.

that RXM could reduce airway inflammation by inhibiting the PI3K- δ /Akt signaling pathway and upregulating histone deacetylase 2 (HDAC2) expression (22). In the present study, RXM treatment reduced the number of airway inflammatory cells, neutrophils, and the concentration of inflammatory cytokines. RXM was also associated with downregulation of calprotectin and RAGE.

In conclusion, pretreatment with RXM exerted similar effects as the RAGE inhibitor FPS-ZM1, and RXM may suppress calprotectin expression via inhibition of RAGE activation, further suppressing the secretion of inflammatory

cytokines and airway inflammation, which improves the neutrophilic asthma. Our findings provide novel insights into the mechanism of RXM in neutrophilic asthma; namely, that a RAGE inhibitor may be a potential candidate for neutrophilic asthma therapy. Additional studies are underway that are exploring the promising new therapeutic options for neutrophilic asthma of inhibiting calprotectin and RAGE.

Acknowledgments

Funding: This work was supported by the National Natural

Science Foundation of China (81570027); Zhejiang Natural Science Funding (LY16H010007); Jinhua Science and Technology Bureau (2017-3-010).

Footnote

Reporting Checklist: The authors have completed the ARRIVE reporting checklist. Available at <http://dx.doi.org/10.21037/atm-21-859>

Data Sharing Statement: Available at <http://dx.doi.org/10.21037/atm-21-859>

Conflicts of Interest: All authors have completed the ICMJE uniform disclosure form (available at <http://dx.doi.org/10.21037/atm-21-859>). The authors have no conflicts of interest to declare.

Ethical Statement: The authors are accountable for all aspects of the work in ensuring that questions related to the accuracy or integrity of any part of the work are appropriately investigated and resolved. Experiments were performed under a project license (protocol wyd2019-019) granted by the Animal Care and Protection Committee of Wenzhou Medical University, in compliance with the Animal Care and Protection Committee of Wenzhou Medical University guidelines for the care and use of animals.

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(English Language Editor: K. Brown)

Cite this article as: Gu X, Shu D, Ying S, Dai Y, Zhang Q, Chen X, Chen H, Dai W. Roxithromycin attenuates inflammation via modulation of RAGE-influenced calprotectin expression in a neutrophilic asthma model. *Ann Transl Med* 2021;9(6):494. doi: 10.21037/atm-21-859