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Glucocorticoid-augmented efferocytosis inhibits pulmonary pneumococcal clearance in mice by reducing alveolar macrophage bactericidal function

Valerie R. Stolberg^{*,1}, Alexandra L. McCubbrey^{†,1}, Christine M. Freeman^{*,‡,1}, Jeanette P. Brown[‡], Sean W. Crudgington[‡], Sophina H. Taitano[†], Bridget L. Saxton[§], Peter Mancuso^{†,¶}, and Jeffrey L. Curtis^{†,‡,||}

*Research Service, VA Ann Arbor Healthcare System, Ann Arbor, MI 48105

[†]Graduate Program in Immunology, University of Michigan, Ann Arbor, MI 48109

[‡]Division of Pulmonary & Critical Care Medicine, Department of Internal Medicine, University of Michigan Health System, Ann Arbor, MI 48109

§Kalamazoo College, Kalamazoo, MI 49006

[¶]Department of Environmental Health Sciences, School of Public Health, University of Michigan, Ann Arbor, MI 48109

^{II}Medical Service, VA Ann Arbor Healthcare System, Ann Arbor, MI 48105

Abstract

Inhaled corticosteroid(s) (ICS) increase community-acquired pneumonia (CAP) incidence in patients with chronic obstructive pulmonary disease (COPD) by unknown mechanisms. Apoptosis is increased in the lungs of COPD patients. Uptake of apoptotic cells (AC) ("efferocytosis") by alveolar macrophages (AMø) reduces their ability to combat microbes, including *Streptococcus* pneumoniae, the most common cause of CAP in COPD patients. Having shown that ICS significantly increase AMø efferocytosis, we hypothesized that this process, termed glucocorticoid-augmented efferocytosis (GCAE), might explain the association of CAP with ICS therapy in COPD. To test this hypothesis, we studied the effects of fluticasone, AC or both on AMø of C57BL/6 mice in vitro and in an established model of pneumococcal pneumonia. Fluticasone plus AC significantly reduced TLR4-stimulated AMø IL-12 production, relative to either treatment alone, and decreased TNF-a, CCL3, CCL5 and KC, relative to AC. Mice treated with fluticasone plus AC before infection with viable pneumococci developed significantly more lung CFU at 48 h. However, none of the pretreatments altered inflammatory cell recruitment to the lungs at 48 h post-infection, and fluticasone plus AC less markedly reduced in vitro mediator production to heat-killed pneumococci. Fluticasone plus AC significantly reduced in vitro AMø killing of pneumococci, relative to other conditions, in part by delaying phagolysosome acidification without affecting production of reactive oxygen or nitrogen species. These results

Address correspondence and reprint requests to: Jeffrey L. Curtis, M.D., Pulmonary and Critical Care Medicine Section (506/111G), VA Ann Arbor Healthcare System, 2215 Fuller Road; Ann Arbor, MI 48105-2303, U.S.A., Phone: 734-845-3457; Fax: 734-845-3257; jlcurtis@umich.edu. ¹These authors contributed equally.

support GCAE as a potential explanation for the epidemiological association of ICS therapy of COPD patients with increased risk of CAP, and establish murine experimental models to dissect underlying molecular mechanisms.

Keywords

Streptococcus pneumoniae; pneumonia; bacterial; mice; inbred strain C57BL/6

INTRODUCTION

Therapy with inhaled corticosteroids (ICS) is central to chronic obstructive pulmonary disease (COPD) management, but is associated in this patient population with excess cases of community-acquired pneumonia (CAP) both in multi-center clinical trials and in analyses of administrative databases (1–10). Suggestions that this risk is balanced by reduced mortality in COPD patients admitted with CAP while using ICS (11, 12) have been contested (13–15). Defining the molecular basis of this epidemiological association could lead to more precisely personalized therapies and better outcomes in COPD, currently the third leading cause of death in the USA (16).

In COPD, as in the general population, for almost three decades the organism most commonly identified in CAP has been *Streptococcus pneumoniae*, also known colloquially as pneumococcus (17–19). It might seem intuitively obvious that the immunosuppressive properties of ICS should increase pneumonia frequency. However, in the sole study using a murine model, glucocorticoids (GC) alone actually reduced lung burden of pneumococcus (20). Additionally, ICS therapy is extremely prevalent in asthma, but most but not all studies have shown no similar increased risk of CAP in asthmatics (21–24). These findings suggest that additional factors may underlie the association of increased CAP risk and ICS therapy in COPD.

Apoptotic cells (AC) are increased in the lungs of COPD patients (25–29). Uptake of AC, also known as efferocytosis, is a complex, incompletely understood process that is relevant to many lung diseases (30). AC uptake by alveolar macrophages (AMø) is lower than by Mø from other organs, and is further reduced by smoking and in COPD (29, 31–37). This lung-specific suppression of AC uptake is mediated in part by interactions between the lung collectins surfactant proteins A (SP-A) & D (SP-D) and the inhibitory Mø receptor signal regulatory protein-alpha (SIRPa, CD172A) (38). The reduced basal efferocytic capacity of AMø is germane to increased CAP risk, as AC uptake induces a unique Mø activation state that favors resolution of inflammation (39), but which also blocks proinflammatory mediator production (40–42). This state differs both from classical and alternative Mø activation (43). The immunosuppressive effects of efferocytosis on host defense have been shown in vivo (39), including in a murine model of pneumococcal pneumonia (44).

We previously reported that clinically-relevant doses of the potent GC fluticasone increase in vitro AC uptake by murine AMø, via both a rapid mechanism dependent in part on downregulation of SIRP α and by a delayed yet sustained mechanism dependent on protein synthesis (45). We term this change GC-augmented efferocytosis (GCAE). The purpose of

this study was to determine whether GCAE alters host defense against pneumococcus. Early host responses to pneumococci depend crucially on AMø (46–48), yet relatively little is known about how AMø responses to pneumococcus are impacted by GC (49). Using murine AMø analyzed in vitro and an established mouse model of pneumococcal pneumonia, we demonstrate a potent detrimental effect on AMø antimicrobial function when GC therapy is used in the presence of alveolar AC.

MATERIALS AND METHODS

Mice

C57BL/6 mice were purchased from Jackson Laboratories (Bar Harbor, ME). Mice were housed under specific pathogen-free conditions in the Animal Care Facility at the VA Ann Arbor Healthcare System, which is fully accredited by the American Association for Accreditation of Laboratory Animal Care. Mice were fed standard animal chow (rodent lab chow 5008; Purina, St. Louis, MO) and chlorinated tap water ad libitum and were used for experiments between 8 and 16 weeks of age. Animal care and experimentation were conducted in accordance with the *Guide for the Care and Use of Laboratory Animals (8th edition)* and were approved by the Ann Arbor VA Healthsystem Subcommittee on Animal Studies.

Pneumococcus cultivation

Streptococcus pneumoniae serotype 3 (clone 6303) stock culture was obtained from ATCC (Manassas, VA). Stocks of bacteria were immediately thawed for 30 seconds at 37°C and cultured in 5 mL Todd Hewitt (TH) broth supplemented with 0.5% yeast extract for 3 h at 37°C and 5% CO₂. Bacterial stock was frozen in 1 mL TH broth with 10% Glycerol (Sigma Aldrich, St. Louis, MO) and used for subsequent in vitro studies. Bacterial CFU counts were verified both by optical density (OD) at 600 nm and by quantitative culture on SRBC/ tryptose agar (Fisher Scientific, Pittsburgh, PA). To generate heat-killed *S. pneumoniae* for use in vitro, bacteria were incubated in a water bath at 56°C for 60 min. No live bacteria were detected after plating onto agar plates.

To maintain bacterial virulence for in vivo experiments, we first passaged *S. pneumoniae* serotype 3 in vivo, using our established murine pneumococcal pneumonia model (50). Untreated C57BL/6 mice received a intratracheal (IT) inoculum using the surgical technique described below, at a dose $(1 \times 10^6 \text{ CFU})$ designed to induce bacteremia. After 24 h, mice were euthanized; spleens were harvested aseptically and processed to isolate multiple individual pneumococcal clones on blood agar plates. These clones were expanded once in TH broth, and then frozen. In all subsequent in vivo experiments, these in vivo passaged pneumococcal clones were defrosted, expanded once in TH broth and used immediately without further passage on agar plates.

Induction of thymocyte apoptosis and quantification of efferocytosis

To induce apoptosis, we treated single cell suspensions of murine thymocytes with 10 μ M dexamethasone (Sigma) for 4 h at 37°C. These conditions consistently produced 50–60% Annexin⁺, PI⁻ thymocytes, as we have previously shown (51). Efferocytosis was quantified

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using a chamber slide-based microscopic assay, as previously described (32). Data are expressed as % efferocytosis, based on the number of AMø ingesting at least one AC; and as the efferocytic index, which was generated by dividing the total number of ingested AC cells by the total number of AMø counted.

AMø isolation and culture

Murine AMø were isolated by BAL using 10–15 mL PBS containing 0.5 mM EDTA in 1 mL aliquots (45). BAL cells were plated in lymphocyte culture media (LCM) (10% FBS, 1 mM sodium pyruvate, 0.5 mM 2-Mercaptoethanol, 1 mM HEPES, 100 u/ml penicillin, 100 u/ml streptomycin, 0.292 mg/ml L-Glutamine in RPMI) for 1.5 h at 37°C and 5% CO₂, and AMø were adhesion purified from this population, by discarding non-adherent cells. For in vitro stimulation studies, AMø were treated with one of four conditions: media alone; 2 μ M fluticasone for 3 h; AC (at a ratio of 10 AC/AMø) for 2 h; or 2 μ M fluticasone for 3 h followed by AC for 2h (Flu + AC). Without washing, LPS from *E. coli* K12 (InvivoGen, San Diego, CA) (1 ng/mL) or heat-killed *S. pneumonia*, at a multiplicity of infection (MOI) of 10 or 100 was added for an additional 24 h. Supernatants were collected and stored at –20 °C until assayed by Luminex.

Protein analysis of supernatants

We used the Luminex 200 system (Luminex Corporation, Austin, TX) running StarStation Software (Applied Cytometry, Dinnington, Sheffield, UK) according to manufacturer's instructions to determine protein levels for TNF- α , IL-1 β , IL-6, IL-12, CCL3, CCL5 and KC (Life Technologies, Grand Island, NY).

GCAE in vivo model

For the GCAE model, mice were administered saline, fluticasone, AC, or fluticasone plus AC via the intranasal (IN) route. All mice received two IN administrations, given 4 h apart, with one of the following: saline + saline; fluticasone + saline; saline + AC, or fluticasone + AC. The dose of fluticasone varied between experiments, ranging from 100 ng to 10 μ g; 1 × 10⁷ AC per mouse was used in all experiments. To deliver the reagents, mice were anesthetized with isoflurane via the open drop method and then were held with their heads elevated. Saline, fluticasone or AC were delivered via one nostril in a volume of 30 μ L. Mice were held in the upright position for an additional 60 seconds after IN administration before being returned to their cages.

In vivo pneumococcal pneumonia model

At 24 h after the last IN treatment, mice were anesthetized with an intra-peritoneal injection of ketamine/xylazine at 90 mg/kg and 10 mg/kg, respectively. The plane of anesthesia was assessed by lack of response to toe pinch and mice were positioned supine on a surgical platform elevated to a 45 degree angle. To allow visualization of the trachea, a small midline skin incision was made and the neck muscles were retracted. Using a 26 gauge needle, *S. pneumoniae* were injected into the trachea (20 μ L PBS containing 50,000 CFU followed by 0.1 mL of air to assure deposition in the lungs). Mice were allowed to recover fully on a

water-jacketed heating pad and were returned to BSL2 housing until euthanasia 48 h later by exsanguination and induction of bilateral pneumothoraces under deep anesthesia.

Flow cytometry

The pulmonary vasculature was perfused via right heart injection of PBS until the effluent was clear, then the lungs were excised and mechanically disaggregated without enzyme treatments, which we have shown efficiently produces single-cell suspensions of high viability (52). After washing, lung cells were stained, fixed and run on an LSR II flow cytometer using FACSDiva software (version 6.1.3; BD Biosciences) with automatic compensation, and data were analyzed using FlowJo software (Tree Star, Ashland, OR) as previously described (45). We stained for the following antigens, using anti-murine antibodies (clone): CD1d (1B1), CD11b (M1/70), CD45 (30-F11), CD45R/B220 (RA3-6B2), CD103 (2E7), Ly6G (1A8), anti-GalCer:CD1d complex (L363) (Biolegend, San Diego, CA); CD3 (145-2C11), CD4 (GK1.5), CD11c (N418), MHC Class II (NIMR-4), NK1.1 (PK136) (eBioscience, San Diego, CA); and Ly6C (AL-21) (BD Biosciences, San Jose, CA).

We analyzed acidification of AMø phagolysosomes by measuring the pH-dependent change in fluorescence of pHrodo. AMø were plated at 40,000 AMø per well in LCM and adherence-purified for 1.5 h, then media was removed and fluticasone (2 μ M) in RPMI-5 (or RPMI-5 alone) was added and plates were cultured for 22 h at 37°C in 5% CO₂. Next, 400,000 AC (10:1 ratio of AC: AMø) in RPMI-5 (or RPMI-5 alone) were added and plates were incubated at 37°C in 5% CO₂ for another 2 h. Then, pHrodo Zymosan Bioparticles (Life Technologies) were added according to manufacturer's instructions and incubated at 37°C in 5% CO₂ for 90 minutes. As a control, one well in each condition received media alone and not Bioparticles. Samples were stained with CD45 and analyzed by flow cytometry. To exclude the possibility of treatment-induced differences in particle ingestion, we performed a similar experiment using FITC-Zymosan Bioparticles (Life Technologies).

Bacterial killing assay

AMø were plated in two different 96-well polystyrene tissue-culture plates (Corning Incorporated, Corning, NY) at 40,000 AMø per well in LCM. One plate was designated the time 0 min (T₀) control, to quantify live bacteria within AMø after phagocytosis but before killing could have time to occur; the other plate was designated the time 120 min (T₁₂₀) plate, to quantify ingested bacteria remaining viable 2 h after ingestion. AMø on both plates were adherence purified for 1.5 h, then media was removed by suctioning and fluticasone (2 μ M) in RPMI-5 (or RPMI-5 alone) was added and plates were cultured for 22 h at 37°C in 5% CO₂. Next, 400,000 AC (10:1 ratio of AC: AMø) in RPMI-5 (or RPMI-5 alone) were added and plates were incubated at 37°C in 5% CO₂ for another 2 h. Then, viable *Streptococcus pneumoniae*, which had been opsonized by incubation with normal rat serum (10% in Hank's Balanced Salt Solution) for 1 h at 37°C with constant shaking, were added to all wells at 2 × 10⁶ CFU per well. Plates were incubated for 20 min at 37°C and 5% CO₂ and then centrifuged for 5 min at 250 × *g*. Supernatants were aspirated and discarded, then 100 μ L of RPMI-5 was added to each well.

Plates were again centrifuged for 10 min at $250 \times g$, and at this point the two plates were treated differently. The T₀ plate received 20 µL of 5% saponin per well and was incubated for 1 min at room temperature. Then, 100 µL of Todd Hewitt broth was added; the plate was covered with parafilm and stored overnight at 4°C. By contrast, the T₁₂₀ plate was incubated at 37°C in 5% CO₂ for 2 h before addition of saponin and then Todd Hewitt broth, followed by storage overnight at 4°C.

The following day, both the T₀ and the T₁₂₀ plates were incubated at 37°C and 5% CO₂ for 2 h. After addition of 15 µL of 5 mg/ml 3-(4,5-dimethylthiazol-2-yl)2,5-diphenyltetrazolium bromide (MTT) (Sigma) and an additional 20 min incubation at 37°C in 5% CO₂, absorbance was measured at 570 nm using a microtiter plate reader. The CFU in the T₀ and T₁₂₀ plates were extrapolated from a standard curve. Bacterial killing was calculated as a percentage, using the following equation: *Bacterial killing* (%) = $((T_0 CFU - T_{120} CFU)/T_0 CFU) \times 100$.

Analysis of reactive oxygen species (ROS) and reactive nitrogen species (RNS)

To detect ROS and RNS, AMø were plated in 96-well polystyrene tissue-culture plates (Corning Incorporated, Corning, NY) at 40,000 AMø per well in LCM. AMø were adherence purified for 1.5 h, then media was removed by suctioning.

To assay ROS, fluticasone (2 μ M) in RPMI-5 (or RPMI-5 alone) was added and plates were cultured for 22 h at 37°C in 5% CO₂. Next, 400,000 AC (10:1 ratio of AC: AMø) in RPMI-5 (or RPMI-5 alone) were added and plates were incubated at 37°C in 5% CO₂ for another 2 h. Media was suctioned off and cells were washed with warm HBSS. A 50 μ m solution of 2', 7'-dichlorodihydrofluorescein diacetate (H₂DCFDA) (Life Technologies) was prepared in RPMI without phenol red and then added to each well. Cells were incubated at 37°C in 5% CO₂ for 1 h. The plate was washed with HBSS, then read at an excitation of 492 nm and emission of 522 nm using an FLx800 fluorescent plate reader (BioTek, Winooski, VT). Then, viable *Streptococcus pneumoniae* were added at 2 × 10⁶ CFU per well and incubated at 37°C in 5% CO₂. The plate was read every 15 minutes for a total of 90 minutes and returned to the incubator after each reading.

To assay RNS, media was removed and fluticasone (2 μ M) was prepared in a fluorescein amine methyl ester (FA-OMe) staining solution, as directed by the manufacturer (Cayman Chemical, Ann Arbor, MI). Wells that were not treated with fluticasone received the fluorescein amine methyl ester staining solution alone. Plates were cultured for 22 h at 37°C in 5% CO₂. Next, 400,000 AC in the staining solution (or staining solution alone) were added and plates were incubated at 37°C in 5% CO₂ for another 2 h. Then, viable *Streptococcus pneumoniae* were added at 2 × 10⁶ CFU per well and incubated at 37°C in 5% CO₂ for 2 h. Media was suctioned off and Hoechst Dye staining solution was prepared and added to each well, as directed by the manufacturer (Cayman Chemical). Plates were incubated for 10 minutes at 37°C. After a final wash, cells were analyzed using an FLx800 fluorescent plate reader. Nitric oxide staining intensity was measured with excitation and emission wavelengths of 485 and 535 nm, respectively. Cell number/density, indicated by the Hoechst Dye, was detectable at excitation and emission wavelengths of 355 and 465 nm, respectively.

Statistics

Statistical analyses were performed using GraphPad Prism 6.0.1 (GraphPad Software, Inc., La Jolla, CA) on a Macintosh Quad-Core Intel Xeon computer running OS X 10.10.3 (Apple; Cupertino, CA). To test for significant differences between groups, we used either ANOVA with appropriate post-hoc testing (Dunnett for comparison to a single control condition, Fisher LSD testing for multiple comparisons) or the analogous non-parametric Kruskal-Wallis test with appropriate post-hoc testing (Dunn for comparison to single control group, Holm-Sídák for multiple comparisons). A p value of < 0.05 was considered to indicate significance.

RESULTS

GCAE reduced production of inflammatory mediators by murine AMø

To test for suppression of host defenses by GCAE, we first studied its effect on stimulated production of inflammatory cytokines by murine AMø after in vitro pre-treatment with either fluticasone, AC, or fluticasone followed by AC. AMø activation by pneumococci depends chiefly on TLR4 recognition of pneumolysin (53), and to a lesser degree on recognition of peptidoglycan by NOD2 (53). Accordingly, we initially stimulated murine AMø in vitro via TLR4, using its prototypic agonist, purified LPS. As a read-out, we measured production of the inflammatory cytokines TNF- α , IL-6 & IL-12, which have all been shown to be essential to combat pneumococci in vivo (54–56), and the chemokines CCL3 (MIP-1 α), CCL5 (RANTES) (57) and KC, a murine functional homologue of human CXCL1.

Exposure of AMø to fluticasone plus AC significantly reduced AMø elaboration of IL-12 relative to the other three conditions, and also significantly decreased secretion of TNF- α relative to AMø pretreated with AC alone or with medium alone (Fig. 1, A–C). Fluticasone plus AC treatment also led to significantly reduced secretion of CCL3 and KC (Fig. 1, D–F) in comparison to the un-pretreated AMø or those that received AC alone. Fluticasone alone strongly inhibited AMø production of all three chemokines and especially of IL-6, so that IL-12 was the only analyte that was significantly lower in AMø pretreated with fluticasone plus AC than in those pretreated with fluticasone alone, although for TNF- α , CCL5 and KC, there were non-significant trends towards lower production in response to the combined stimuli. We also measured IL-1 β in these experiments, but secretion was near or below the limit of detection for all conditions (data not shown). These results imply that GCAE could reduce the ability of murine AMø to secrete inflammatory mediators crucial for recruitment of other leukocyte subsets during early pneumococcal pneumonia. Accordingly, we next set about to test that possibility in vivo.

GCAE inhibited in vivo clearance of S. pneumoniae in a murine model

To establish a murine model of GCAE, we first needed to verify that a physiological dose of fluticasone altered murine AMø efferocytosis in vivo. Mice were pretreated with fluticasone by IN inoculation for 6 h, then received AC by the IN route. After an additional 2 h, we harvested AMø by BAL and quantified ingested AC. Results showed that fluticasone

pretreatment significantly increased uptake of AC by murine AMø in vivo in a dosedependent manner (Fig. 2).

In the next set of experiments, mice were pretreated with saline, fluticasone, AC, or fluticasone followed by AC (all by IN route), and then all groups received *Streptococcus pneumoniae* (50,000 CFU/mouse by the IT route). After 48 h, lungs were harvested and total CFU per lung were calculated (Fig. 3A). Pretreatment by fluticasone followed by AC significantly reduced pneumococcal clearance from the lungs, relative to all other treatment groups (Fig. 3B). Thus, short-term treatment leading to GCAE in vivo had an adverse effect during bacterial pneumonia.

GCAE did not alter inflammatory cell recruitment to the lungs during pneumococcal pneumonia

As a first step toward determining the mechanism by which GCAE had this adverse effect on lung host defenses, we used a separate cohort of mice to analyze inflammatory cell recruitment to the lungs during pneumonia. Mice were treated exactly as in the previous experiment, but at 48 h, lungs were harvested without previous bronchoalveolar lavage and total cells were stained for flow cytometric analysis.

All cell populations were initially gated on viable CD45⁺ cells. We defined AMø as high autofluorescent, CD11c⁺ cells that were negative for CD3, CD19 & Ly6G. Neutrophils were identified as high side scatter cells abundantly expressing Ly6G. We defined dendritic cells (DC) as low autofluorescent, MHC II⁺, CD11c^{dim} cells that were negative for CD3, CD19 and Ly6G; two DC subsets were distinguished, CD11b⁺ CD103⁻ (which we and others have shown are located predominately in parenchymal lung interstitium) and CD11b⁻ CD103⁺ (which have been shown to reside largely in airways) (58, 59). Similarly, exudate Mø were identified as low autofluorescent, CD11b^{high} cells, among which we distinguished two subsets by their expression of Ly6C. We also identified CD4+ T cells, which have recently been recognized to contribute importantly and acutely to host defense during pneumococcal pneumonia (60, 61). Finally, we searched for NK cells and NKT cells, using staining for NK1.1, CD1d, CD3 and anti-GalCer.

There were no significant differences between the four in vivo treatments for any of these inflammatory cell populations at this time post-infection, whether expressed as absolute number of cells per mouse lung (Fig. 4) or as percentage of each cell type among all CD45⁺ lung cells (data not shown). Although we identified lung NK cells (NK1.1⁺, CD3⁻, CD1d⁻, anti-GalCer⁻), we found no evidence of classical (type I) NKT cells (NK1.1⁺, CD1d⁺ anti-GalCer⁺) or non-classical (type II) NKT cells (CD1d⁺ anti-GalCer⁻). Interestingly, for all cell types except the CD11b⁻, CD103⁺ DC subset, there was a non-significant trend towards higher absolute numbers in the fluticasone-treated group (Fig. 4). The reason for this trend is not apparent. Thus, the observed significant differences in lung pneumococcal CFU associated with GCAE (Fig. 3) could not be attributed to disparity in numbers or relative composition of lung inflammatory cells.

GCAE minimally decreases production of cytokine and chemokine murine AMø in response to heat-killed pneumococcus

This unanticipated result for inflammatory cell accumulation led us to repeat our initial in vitro stimulation of murine AMø pre-treated with fluticasone, AC or both, but this time using heat-killed pneumococci to induce inflammatory mediator production. Results were strikingly different than with LPS, as the combined treatment induced a significant decrease, relative to the other three conditions, only in IL-6 production (Fig. 5). At this ratio of heat-killed pneumococci to AMø (MOI 100), the combined stimulus of fluticasone followed by AC significantly decreased production of TNF- α and CCL3, relative to no pretreatment or AC alone, but the change was not statistically significant for fluticasone pretreatment alone. Interestingly, there were no differences between groups in response to heat-killed organisms for IL-12 or KC, which had shown the greatest difference in response to LPS.

We considered that the ratio of organisms to AMø, the MOI, might be too great to see an effect of GCAE, so in one experiment performed in parallel we analyzed an MOI of 10. However, at that lower MOI, the only analyte detected in any condition was CCL-3 (data not shown). Collectively, these findings are consistent with the lack of differences in inflammatory cell recruitment in vivo, and imply that the effect of GCAE varies with the stimulus, being much stronger with stimulation of TLR4 than with the heat-killed organism. Nevertheless, these results do not provide an explanation for the significant differences observed in lung CFU.

GCAE significantly reduced in vitro killing of pneumococcus by murine AMø

To test whether bacterial killing by AMø was inhibited by GCAE and could contribute to the observed increase in bacterial burden in vivo, we performed an in vitro bacterial killing assay. AMø from normal C57BL/6 mice were adherence-purified before pre-treatment with either saline, fluticasone, AC, or fluticasone followed by AC. Pre-treatment with fluticasone plus AC significantly reduced AMø-mediated bacterial killing, relative to each of the three other conditions (Fig. 6A). Fluticasone pre-treatment also resulted in a modest decrease in bacterial killing, which was significantly different from the saline control. Hence, an additive reduction in AMø killing of pneumococcus is one means by which GCAE impairs host defense against pneumonia. Numbers of CFU at T_0 were significantly increased in AMø pretreated with fluticasone plus AC, relative to the untreated and AC only groups (Fig. 6B). This result implies that the observed GCAE-induced deficit in killing by T_{120} mins did not result from reduced initial phagocytosis of pneumococcus.

An effect of AC on in vitro killing of *K. pneumoniae* by rat AMø was previously shown in experiments using opsonization by specific IgG (44). Accordingly, we investigated whether the GCAE-induced decrement in killing of pneumococcus by murine AMø that we showed using complement-opsonized bacteria would also be seen with specific IgG opsonization. We compared killing of pneumococci opsonized by the two methods by untreated murine AMø or AMø treated in vitro with AC plus fluticasone. Results showed a similar reduction in bacterial killing by GCAE regardless of the method of opsonization (Supplemental Figure 1), although the difference from untreated AMø did not attain statistical significance in this single experiment.

GCAE and fluticasone impaired AMø killing of S. pneumoniae in part by reducing phagolysosome acidification

We performed several types of experiments to begin to define the possible mechanisms by which GCAE impairs in vitro killing of *S. pneumoniae* by murine AMø. Bacterial killing by murine Mø often involves production of ROS by the NADPH oxidase complex and of RNS by inducible nitric oxide. However, neither exposure of murine AMø to fluticasone, AC or both significantly affected ROS production in response to ingestion of viable pneumococci (Supplemental Fig. 2A). Similarly, none of these treatments impacted production of RNS (Supplemental Fig. 2B).

Finally, we examined the effect of GCAE on phagolysosome acidification. We assayed AMø phagolysosome acidification by quantifying the change in fluorescence of the pH-sensitive dye pHrodo using flow cytometry. Pretreatment with either fluticasone or fluticasone followed by AC significantly impaired phagosome acidification in vitro (Fig. 7A, B). To exclude the possibility that this difference was artifactual, and instead reflected treatment-induced differences in particle ingestion, we performed a similar experiment using FITC-Zymosan Bioparticles. That control experiment showed that the combination of fluticasone followed by AC slightly but significantly increased AMø ingestion, which was avid for all treatments (Fig. 7C).

DISCUSSION

The results of this study demonstrate that by the process we term GCAE, the combination of the potent GC fluticasone and subsequent AC exposure impairs the ability of resident AMø to defend against pneumococcus more greatly than either stimulus alone. GC pre-treatment in vitro significantly reduced pro-inflammatory production by murine AMø in response to TLR4-stimulation, although the effect was less marked using heat-killed pneumococci under the conditions tested. The combination of steroid pre-treatment and intra-alveolar AC led to significantly greater lung bacterial burdens in a murine model of pneumococcal pneumonia, without altering inflammatory cell recruitment to the lungs. GCAE also decreased in vitro killing of pneumococci by murine AMø, but not their uptake, in association with a decrease in phagolysosome acidification. These data provide one mechanism for the increased susceptibility to CAP when COPD patients are treated with potent ICS therapy.

Streptococcus pneumoniae is a near-obligate human pathogen that causes more deaths globally than any other organism, principally due to infections in children under the age of five years in the developing world (62). However, pneumococcal pneumonia also continues to be a very significant health problem in industrialized nations, despite vaccines that have efficacy even in elderly patients with COPD (63). Moreover, the lethality of pneumococcal pneumonia remains considerable even when appropriate antibiotic therapy is initiated promptly (64). Providing appropriate antibiotic therapy against pneumococcus is becoming more difficult, as high-grade resistance to multiple antibiotics is already prevalent in many regions, especially Asia. Although antibiotic stewardship may delay the global spread of resistance, the profound ability of pneumococcus to undergo DNA transformation virtually assures that the trend will continue. Hence, investigating the immunological basis of susceptibility to pneumococcal pneumonia will remain important.

Our studies employed mice, a well-accepted model of human pneumococcal pneumonia, provided appropriate attention is given to pneumococcal serotype, murine strain and anesthesia method (46, 65, 66). We used the encapsulated serotype 3, which in humans remains associated both with common nasopharyngeal carriage and frequent pneumonia with relatively high mortality (67–69), perhaps recently in part due to its lower immunogenicity relative to other strains in polyvalent vaccines (70). We (50, 71, 72) and others (44, 73–85) have used variations of this murine model extensively in wild-type and transgenic mice to define the molecular mechanisms of lung host defense against pneumococcus. Together with established murine models mimicking the pathogenic changes of COPD, either employing or independent of cigarette smoke-exposure (86, 87), the tools are now available to determine whether the beneficial effects of ICS can be dissociated from this and other adverse effects.

The demonstration that GCAE was associated with both significantly greater lung bacterial burdens and reduced killing in vitro is important because resident AMø are so crucial to defend against pneumococcal pneumonia (88). Part of their role depends on killing pneumococci, which for AMø occurs almost entirely intracellularly. AMø require opsonization to ingest encapsulated strains, which are associated with most episodes of CAP in humans. Accordingly, we used the encapsulated serotype 3 and performed serum opsonization in all experiments. Ingestion depends on several Mø receptors, including $Fc\gamma R$ and the scavenger receptors SR-A and MARCO (89) and (primarily for unopsonized pneumococci) mannose receptor. Our data indicate GCAE and fluticasone alone increased ingestion of pneumococcus by murine AMø in vitro, which literature searching leads us to conclude is a novel finding. The reason for this increase is uncertain. Our previous demonstration that fluticasone induced downregulation by murine AMø of the inhibitory receptor SIRPa (45) together with the finding that alveolar lining fluid [which contain surfactant proteins (SP–) A and SP-D, the collectin ligands for SIRPa] impaired phagocytosis by murine AMø (90) provide a possible explanation. However, increased phagocytosis of pneumococcus was induced in human neutrophils by both SP-A and SP-D (91), and in murine AMø by SP-A (via surface localization of SR-A) but not by SP-D (92). Hence, further study will be needed to establish that mechanism. Importantly, the calculation of percentage bacterial killing takes into account difference in bacterial uptake, so this effect does not explain the GCAE-induced killing defect we show.

Unlike the better studied neutrophil, which eliminate pneumococci using multiple extracellular and intracellular mechanisms, precisely which elements are essential for AMø to kill ingested pneumococci remains incompletely defined. Our finding that neither GCAE nor fluticasone alone affected ROS production extends results of Marriott and colleagues, who used gp91phox–/– mice to show that ROS production was dispensable in a subclinical pneumonia model (93). Glucocorticoids do reduce phagocyte generation of superoxide at least in part via reduced eicosanoid signaling (94), but that effect is seen at micromolar concentrations irrelevant to GCAE, which occurs at nanomolar concentration identical to those achieved clinically during ICS therapy. Further investigation is needed to define the roles of ROS and RNS in AMø killing of pneumococcus, but our results show that changes in these mediators cannot explain the defect induced by GCAE.

To our knowledge, corticosteroids have not previously been shown in any mammalian phagocyte to affect phagolysosome acidification, a particularly crucial step in the killing process (95). An acidic intraluminal pH is necessary for optimal activity of cathepsins, which contribute to pneumococcal clearance in murine models both by inducing AMø apoptosis linked to bacterial killing (83), and at least in neutrophils, by direct anti-bacterial activity (96). Additionally, acidification of the maturing phagolysosome is important to counteract three potentially detrimental effects of the oxidative burst that would otherwise increase pH: (a) consumption of protons as superoxide undergoes dismutation to hydrogen peroxide; (b) leakage of H+ into the cytoplasm due to membrane oxidation; and (c) impaired recruitment to the phagosome of V-ATPase, which is essential to complete acidification (97). Because the effect on acidification was equivalent in AMø treated with fluticasone without or with subsequent AC, this mechanism cannot explain the even greater reduction in killing induced by GCAE, relative to fluticasone alone. Considerable additional investigation will be required to define the molecular basis for this GCAE-induced defect, but these results advance the field by excluding defective ROS or RNS generation.

In experimental models, AMø also contribute to defense against pneumococcal pneumonia by production of cytokines and chemokines, which are particularly crucial to activate and recruit other cell types in response to larger inocula (98). The disparity in results of stimulating AMø in vitro with purified LPS versus with heat-killed pneumococci does not necessarily detract from the relevance of GCAE shown by our lung CFU and in vitro killing data because neither experiment fully simulates the indirect contribution of AMø to host defense during pneumococcal pneumonia. For example, production of CCL5 by murine Mø in response to pneumococcus requires pneumolysin-dependent escape from phagolysosomes that likely does not occur efficiently using heat-killed organisms (99, 100). However, the absence of any significant difference between treatment groups in recruitment of inflammatory cell subsets implies that in vivo, either AMø production of recruitment signals was not essential under the conditions tested, or that other factors compensated. We suspect that the disparity in our results regarding neutrophil recruitment from the decreases seen by Medeiros and colleagues in response to AC alone (44) relate to differences from our experiments in pneumococcal inoculum (10⁶ CFU, versus 5×10^4 CFU in this study) (98). Moreover, it is likely that GCAE-induced failure of early pneumococcal eradication in our study led to compensatory changes by other cell populations that culminated in increased neutrophil recruitment to the lungs by the time-point we studied.

In adults, COPD is one of the strongest risk factors for development of CAP. COPD is also a pervasive and rapidly increasing problem worldwide, due to the juxtaposition in much of the developing world of rapidly increasing cigarette consumption and air pollution, particularly by indoor use of biomass fuel or coal. Although COPD is also increasing as a directly attributable cause of death in industrialized nations, its true burden is likely vastly underestimated. COPD is under-diagnosed, especially in women (101). Hence, defining precisely why there is an association between ICS therapy, which clearly has beneficial effects in COPD, and potentially lethal hospitalizations for CAP should be of significant public health interest. Our data imply that the risk of GCAE contributing to CAP might be greatest in those with emphysema, which among the several pathological processes contributing to COPD phenotypes is the one most closely associated with excessive

apoptosis of parenchymal lung cells. To date, none of the studies on the association of CAP and ICS usage have included the imaging data necessary to address that possibility. Indirect support for that possibility comes from the recent identification of the emphysematous subset as being at elevated risk for acute exacerbations of COPD (102), a largely infectious syndrome that blends clinically with CAP.

Based on our in vitro findings that budesonide also induced GCAE in murine AMø (45), we suspect that the effect we have shown here using fluticasone is a class-effect of all potent ICS medications. However, we recognize that there is controversy on this important clinical point. A retrospective observational trial using administrative health data found lower pneumonia event rates in COPD patients using budesonide compared to fluticasone (10). An individual-subject data meta-analysis relying heavily on adverse event reporting and including studies of relatively short duration found no difference in incidence of CAP from placebo among users of budesonide (103). However, this conclusion has also been contested (104, 105). Verifying whether specific ICS agents have different risks of infection or other adverse effects is a highly significant question, but one which will likely require large clinical trials or very careful epidemiological approaches. Importantly for our hypothesis that GCAE might explain the increased risk of CAP among COPD patients using ICS therapy, asthma is not associated with apoptosis of alveolar epithelial and endothelial cells that can interact with AMø.

In summary, we have extended our previous identification of GCAE as a property of resident tissue Mø (45) (i.e., not only a process relevant during maturation of blood monocytes) by showing its impact on a clinically relevant bacterial pathogen in vitro and in vivo. Our results support GCAE as a potential explanation for the epidemiological association between ICS therapy of COPD patients and an increased risk of CAP, although further support for that possibility is required, including studies using human AMø that we have underway. These findings also establish murine in vitro and in vivo experimental models to dissect the underlying molecular mechanisms by which GCAE impacts lung host defense.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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LIST OF ABBREVIATIONS USED

AC	apoptotic cell(s)
AMø	alveolar macrophages(s)
BAL	bronchoalveolar lavage
CAP	community acquired pneumonia
COPD	chronic obstructive pulmonary disease
GC	glucocorticoid
GCAE	glucocorticoid-augmented efferocytosis
ICS	inhaled corticosteroid(s)
IN	intranasal
IT	intratracheal
ТН	Todd Hewitt (broth)

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Figure 1. GCAE reduced inflammatory cytokine production by murine AMø in response to stimulation via TLR4

Adherence-purified AMø from normal C57BL/6 mice were pre-treated with media alone (none); 2 µM fluticasone for 3 h (Flu), AC (at a ratio of 10 AC/AMø) for 2 h (AC); or 2 µM fluticasone for 3 h followed by AC for 2h (Flu + AC). Next, LPS at 1 ng/mL was added to all wells for an additional 24 h. Supernatants were collected and assayed by Luminex for protein concentrations of (A) TNF- α ; (B) IL-6; (C) IL-12; (D) CCL3; (E) CCL5; and (F) KC. Results are mean ± SEM of four independent experiments, each using pooled AMø from two mice. *, *p* < 0.05; **, *p* < 0.01; ***, p<0.001; ****, p<0.0001; NS, not significant by ANOVA with Fisher LSD post-hoc testing.



Figure 2. Fluticasone increased in vivo uptake of AC by resident AMø

A. C57BL/6 mice received two IN administrations, of either various doses of fluticasone (100-10,000 ng/mL) or saline control, followed 6 h later by an IN administration of 1×10^7 AC. One h later, AMø were collected by BAL; cytospins were stained with H&E and ingested AC were counted under oil at $1000 \times$ final magnification. A. Representative cytospins showing in vivo AC uptake following in vivo fluticasone treatment with either saline (top panel) or 1000 ng fluticasone. Arrows point to ingested AC. Top panel, percentage of AMø ingesting at least one AC; lower panel, efferocytic index. B. Data are mean \pm SEM from three mice in a single experiment, and are representative of results of three independent experiments. Statistical testing using one-way ANOVA with Dunnett's post-hoc testing for multiple comparisons relative to saline-only control group.

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Figure 3. GCAE specifically reduced clearance of viable *S. pneumoniae* from the lungs in a murine model

C57BL/6 mice received two IN administrations, given 4 h apart, of either saline (indicated by minus symbol), fluticasone followed by saline, saline followed by AC, or fluticasone followed by AC. All mice were infected via the IT route 24 h after the final IN treatment using 50,000 CFU *S. pneumoniae* serotype 3. Lungs were collected 48 h later to assay total CFU by serial dilution on blood agar plates. Data are derived from 2–3 mice per condition assayed individually in each of three independent experiments (total n = 34). A. Log lung CFU of individual mice; symbols denote mice from different experiments. B. Fold-change in lung CFU, relative to the group pre-treated twice with saline before infection, the

geometric mean of which was set to 1. Data are shown as median, 25% & 75% (box) and 5%, 95% CI (whiskers), with outliers shown individually. *, p<0.5; **, p<0.01 by Kruskal-Wallis non-parametric ANOVA with Dunn's post-hoc testing for multiple comparisons to saline-only control group.



Figure 4. GCAE did not alter inflammatory cell recruitment to the lungs during pneumococcal pneumonia

Mice were pre-treated by the IN route and infected by the IT route with 50,000 CFU/mouse *S. pneumoniae* serotype 3, exactly as described in the Legend to Figure 3, except that 48 h post-infection, lungs were harvested and processed individually for flow cytometry. Hematopoietic cells were gated using light-scatter parameters and CD45 staining as described in the **Results**. Data are expressed as the absolute number of cells per lung for each cell type on the vertical axis (note differences in scales), as mean \pm SEM of 2–3 mice per condition assayed individually in each of two independent experiments (total *n* = 21). There were no statistically significant differences between treatment groups by ANOVA with Fisher's LSD post-hoc testing.



Figure 5. GCAE reduced inflammatory cytokine production by murine AMø in response to stimulation by heat-killed pneumococci

Adherence-purified murine AMø from C57BL/6 mice were pre-treated with media alone (none); 2 μ M fluticasone for 22 h (Flu), AC (at a ratio of 10 AC/AMø) for 2 h (AC); or 2 μ M fluticasone for 22 h followed by AC for 2h (Flu + AC). All conditions were incubated for and additional 24 h with heat-killed *S. pneumoniae* serotype 3 at a multiplicity of infection (MOI) of 100 (except for CCL3, for which MOI = 10). Supernatants were collected and assayed by Luminex for protein concentrations of TNF- α , IL-6, IL-12 (top row) and CCL3, CCL5 and KC (bottom row). Results are mean ± SEM of three independent experiments, each using pooled AMø from two mice. *, p < 0.05; **, p < 0.01; ***, p < 0.001; NS, not significant by ANOVA with Fisher LSD post-hoc testing.



Figure 6. GCAE significantly inhibited in vitro killing of *S. pneumoniae* **by murine AMø** Adherence-purified AMø from normal C57BL/6 mice were treated with one of four regimens: saline (indicated by minus sign) alone twice; 2 μ M fluticasone for 22 h followed by saline for 2 h; saline for 22 h followed by AC for 2 h; or 2 μ M fluticasone for 22 h followed by AC for 2 h. AC were added at a ratio of 10:1 relative to AMø. Next, viable pneumococci (2 × 10⁶ CFU) were added, and then bacterial killing was assayed as described in **Material & Methods**. Data are expressed as (A) the percentage of bacterial killing at T₁₂₀ mins; and (B) lung CFU (in millions) at T₀ mins; both are derived from four independent experiments each containing at least three mice per group. Results are depicted

as a box & whiskers plot, indicating median, 25^{th} & 75^{th} percentiles and minimum, maximum values; *, p < 0.05; **, p < 0.01, by one-way ANOVA with Holm-Sídák post-hoc testing for multiple comparisons.



Figure 7. Fluticasone alone and GCAE significantly inhibited phagolysosome acidification Adherence-purified AMø from normal C57BL/6 mice were treated with one of four regimens: saline alone twice; 2 μ M fluticasone for 22 h followed by saline for 2 h; saline for 22 h followed by AC for 2 h; or 2 μ M fluticasone for 22 h followed by AC for 2 h. Next, either pHrodo Bioparticles (A, B) or FITC Zymosan Bioparticles (C) were added and plates were cultured for an additional 90 minutes. Finally, AM were washed, harvested and analyzed by flow cytometry. Corrected MFI was calculated individually for each condition by subtracting the MFI of the well without added particles from the well with added particles in the same experiment. (A) Representative staining for pHrodo Bioparticles (FITC +) for each condition. (B) Aggregated corrected MFI for pHrodo. Results are mean \pm SEM of three independent experiments (4–5 mice per experiment); each symbol represents an individual well. (C) Aggregated corrected MFI for FITC-Zymosan; note compressed range. Results are mean \pm SEM from one experiment using pooled AMø from five mice; each symbol represents an individual well. *, p<0.05; **, p < 0.01, by one-way ANOVA with Holm-Sídák post-hoc testing for multiple comparisons; NS, not significant.