



# Analysis of Immune Responses in *Acinetobacter baumannii*-Infected Klotho Knockout Mice: A Mouse Model of *Acinetobacter baumannii* Infection in Aged Hosts

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*Acinetobacter baumannii* is an important opportunistic pathogen that primarily afflicts elderly people. To clarify the pathogenicity of *A. baumannii* in the elderly, we investigated immune responses to *A. baumannii* ATCC 19606 infection in klotho knockout (KO) mice, the mouse model of aging. Following intravenous inoculation, the mice seldom displayed severe symptoms. However, the survival rate was 56% at 7 days post-infection. Bacteria were detected in the lungs of klotho KO mice but not klotho wildtype (WT) mice at 7 days post-infection. Neutrophils, eosinophils, interstitial macrophages, and monocyte/dendritic cell subset in the lungs of klotho KO mice were transiently induced after infection with *A. baumannii*. The number of alveolar macrophages in klotho KO mice was lower than that in klotho WT mice, except for 1 day post-infection. CD11b expression on neutrophils and alveolar macrophages in the lungs of klotho KO mice was seldom upregulated by the infection. These results suggested that immune functions eliminating bacteria in the lungs of klotho KO mice were insufficient. CD11b<sup>low</sup> conventional DC cells hardly increased in klotho KO mice infected with *A. baumannii*. Additionally, the production of interleukin (IL)-10 in the sera of klotho KO mice was significantly higher than that in klotho WT mice, whereas that production of interferon-gamma was not detected in the sera of klotho KO mice. These results suggested that acquired immune responses were hardly induced in klotho KO mice. *IL-1 $\beta$* , *CXCL1*, *CXCL2*, and *CCL2* expression was significantly higher in the lungs of klotho KO mice infected with *A. baumannii* than in those of klotho WT mice at 1 day post-infection. These results suggested that pulmonary inflammation was elicited in klotho KO mice during early infection. The expression levels of proinflammatory cytokines significantly correlated with *TLR9* expression in the lungs of klotho KO mice. The collective results demonstrate an *A. baumannii* infection state in aged hosts and suggest that pulmonary inflammation and bacterial burden should be noted in aged hosts even in the absence of severe symptoms of *A. baumannii* infection.

**Keywords:** *Acinetobacter baumannii*, klotho knockout mice, immune responses, pulmonary inflammation, toll-like receptors

## INTRODUCTION

*Acinetobacter baumannii* is an important opportunistic pathogen that is associated with nosocomial infections that include bacteremia, pneumonia, meningitis, urinary tract infections, and wound infections (1, 2). The recent increase in multidrug-resistant *A. baumannii* (MDRAB) outbreaks worldwide has become a concern (3–5). Additionally, *A. baumannii* is among the six nosocomial pathogens—*Enterococcus faecium*, *Staphylococcus aureus*, *Klebsiella pneumoniae*, *A. baumannii*, *Pseudomonas aeruginosa*, and *Enterobacter* spp.—that comprise the ESKAPE group that have acquired multidrug resistance and virulence (6, 7). *A. baumannii* is also an emerging pathogen among elderly people in community hospitals and nursing homes (8). Given the accelerating pace of aging of the global population, *A. baumannii* is an important global human pathogen.

Although *A. baumannii* is regarded as a low-virulence pathogen (9), it possesses several mechanisms of pathogenicity, including biofilm formation, adherence, and invasion of lung epithelial cells (10–14), host cell death (15–17), and iron acquisition (18). The pathogenicity of *A. baumannii* depends on various virulence factors, especially the outer membrane proteins (Omps) (19). Additionally, we have reported that MDRAB clinical isolates are resistant to toxicity caused by reactive oxygen species (ROS) in macrophages (20). Taken together, *A. baumannii* is an emerging, highly pathogenic bacterium with characteristics that vary with environmental stress conditions, such as multiple antimicrobial agents and host immune responses. However, only few studies have explored the reasons for the pathogenicity of *A. baumannii* among elderly people.

Some studies on host immune responses to *A. baumannii* infection using mouse models have been performed. Neutrophils and alveolar macrophages (AM) play important roles in host resistance to respiratory infection of *A. baumannii* (21–23). In addition, Toll-like receptor (TLR) 2 and 4 play key roles in eliciting innate immune responses against *A. baumannii* infection (24, 25). TLR9 contributes to the expression of proinflammatory cytokines and chemokines (26). These results suggest that innate immune responses play important roles in host resistance to early *A. baumannii* infection. However, these studies have focused on the immune responses of young hosts against *A. baumannii* infection. In contrast, aged mice reportedly showed increased susceptibility to *A. baumannii* infection, increased inflammatory cell infiltration, increased proinflammatory cytokine levels at 24 h post-infection, as well as impaired efficacies of antibiotics and a vaccine (27). These results suggest that immune responses are altered in aged mice compared with the responses in young mice.

Klotho protein is senescence suppressor and its expression naturally reduces in the elderly and in individuals with chronic kidney disease (28). Klotho knockout (KO) mice show several aging phenotypes that are similar to human aging (29). These phenotypes include arteriosclerosis, osteoporosis, age-related skin changes, and ectopic calcifications, together with short lifespan and infertility (29). A klotho-septic mouse model that displayed hypercytokinemia with impaired bacterial clearance and increased apoptosis of lymphocytes may be related to poor survival in klotho

KO mice treated with cecal ligation and puncture (30). Additionally, a previous study reported that klotho deficiency aggravates sepsis-related multiple organ dysfunction (31). Thus, klotho KO mice are expected to be useful as an animal model for the clarification of pathogenesis of bacterial infections in aged hosts. Generally, aging affects multiple immune systems (32–34). However, although previous studies have reported acute immune responses to *A. baumannii* infection in young mice (35–37), only a few studies have investigated the immune responses of aged hosts to *A. baumannii* infection. Additionally, although immune responses are altered in aged mice compared with those in young mice at 24 h post-infection (27), the immune responses that occur for a few days after *A. baumannii* infection in aged hosts remain unknown.

In this study, we established an *A. baumannii* infection klotho KO mouse model *via* intravenous inoculation and evaluated its immune responses upon infection with *A. baumannii*. Since TLRs play an important role in eliciting immune responses to *A. baumannii* infection (24–26), we also analyzed the correlation between the expression levels of proinflammatory cytokines and those of TLRs in the lungs of the KO mice.

## MATERIALS AND METHODS

### Bacterial Strains and Growth Condition

ATCC 19606 strains of *A. baumannii* were used in this study. Bacteria were cultured on Luria-Bertani (LB) agar plates (Becton, Dickinson and Company, Sparks, MD, USA) for 16 h at 37°C. The bacteria were washed twice with PBS and resuspended at a concentration of  $5 \times 10^8$  colony forming units (CFU)/ml, with the concentration adjusted *via* optical density measurements at 595 nm. The bacterial suspensions obtained were used for animal experiments. All procedures were carried out in accordance with the relevant guidelines and regulations.

### Animal Experiments

All animal experiments were performed in accordance with the Institutional Animal Care and Use Committee of Teikyo University Animal Ethics Committee (approval #17–009). Five-week-old male klotho KO mice and littermate klotho wildtype (WT) mice were purchased from CLEA Japan, Inc. (Tokyo, Japan). Bacteria ( $10^8$  CFU/200  $\mu$ l) were inoculated intravenously into 6-week-old klotho mice. PBS were inoculated intravenously into 6-week-old klotho mice as uninfected mice. To analyze the number of bacteria in the lungs and spleen of klotho mice, 100  $\mu$ l of the homogenates in PBS was plated on LB agar plates. The bacterial count (CFU) was confirmed by the growth of serial dilutions of the bacterial suspension on LB agar after 24 h of incubation at 37°C. To analyze the number of bacteria in the blood of klotho mice, 100  $\mu$ l of blood was plated on LB agar plates. The bacterial count was confirmed using the same method.

### Flow Cytometry Analysis

Cells were harvested from the lungs, spleens, and blood of *A. baumannii*-infected klotho mice and uninfected mice. Briefly,

mice were gently killed by cervical dislocation under sedation, and the lungs and spleens were surgically harvested. The lung tissue was cut into small pieces with scissors, and processed in digestion buffer (200 µg/ml of Collagenase Type 1 CLS1 (Worthington Biochemical, Lakewood, NJ) and 2 mM of CaCl<sub>2</sub> in PBS). The homogenized lungs and spleen were prepared by gently pressing the organs through a 40-µm nylon mesh into a single-cell suspension. The cells were washed and red blood cells were lysed with PharmLyse Lysis Buffer (BD Biosciences, San Jose, CA). The harvested blood was treated with PharmLyse Lysis Buffer (BD) and the cells were washed with PBS. Isolated cells were counted and then stained with monoclonal antibodies (mAbs) for the analysis of immune cells. Briefly, isolated cells from the lungs were stained with various combinations of mAbs. The mAbs included were fluorescein isothiocyanate (FITC)-labeled anti-F4/80, phycoerythrin (PE)-labeled anti-CD11b, PE-Cy-7-labeled anti-CD45, allophycocyanin (APC)-labeled anti-CD11c, and APC-Cy-7-labeled Ly-6G mAbs (Biologend, San Diego, CA). The white blood cells from the spleen and blood were stained with various combinations of mAbs, including FITC-labeled anti-F4/80, PE-labeled anti-CD11b, PE-Cy-7-labeled anti-Ly-6C, APC-labeled anti-CD11c, and APC-Cy-7-labeled Ly-6G mAbs. To analyze the proportion of immune cells in the lungs, spleen, and blood of klothe mice, immune cells were stained with specific antibodies against surface markers for 30 min at 4°C. The cells were washed twice with stain buffer (BD), and subsequently fixed with BD Cytofix™ Fixation Buffer (BD). The stained cells were analyzed using a FACSCant II flow cytometer equipped with FACS Diva software (BD). All flow cytometry data were analyzed using FlowJo software (BD).

## Cytokine Production

Cytokine levels of tumor necrosis factor-alpha (TNF-α), IL-6, IL-10, IL-12, interferon-gamma (IFN-γ), and chemokine (C-C motif) ligand 2 (CCL2) in the sera of uninfected and *A. baumannii*-infected klothe mice were measured using the Cytometric Bead Array Mouse Inflammation Kit (BD), as per the manufacturer's protocols. Cytokine levels of mouse chemokine (C-X-C motif) ligand 1 (CXCL1) and CCL7 in the sera of uninfected and *A. baumannii*-infected klothe mice were measured using the Mouse CXCL1 ELISA Kit (Abcam) and the Mouse MCP-3 (CCL7) ELISA Kit (Abcam), as per the manufacturer's protocols. The lower limits of detection were as follows: TNF-α, 7.3 pg/ml; IL-6, 5 pg/ml; IL-10, 17.5 pg/ml; IL-12p70, 10.7 pg/ml; IFN-γ, 2.5 pg/ml; CXCL1, 5.47 pg/ml; CCL2, 52.7 pg/ml; and CCL7, 6.25 pg/ml.

## RNA Extraction and Quantitative Real-Time Polymerase Chain Reaction (qPCR)

To analyze the expression of proinflammatory cytokines, total RNA was extracted from the lungs of uninfected and *A. baumannii*-infected klothe mice using the RNeasy Plus Mini kit (QIAGEN, Tokyo, Japan). Harvested RNA samples were quantified using the NanoDrop spectrophotometer (Thermo Fisher Scientific, Waltham, MA). Total RNA was reverse-transcribed to cDNA using PrimeScript™ 1<sup>st</sup> strand cDNA

Synthesis Kit (TaKaRa Bio, Shiga, Japan). To analyze the mRNA levels of all genes, cDNA was amplified using the PowerUP SYBR Green Master Mix (Thermo Fisher Scientific) with consensus primers for detecting mouse glyceraldehyde 3-phosphate dehydrogenase (*GAPDH*), mouse *TNF-α*, mouse *IL-1β*, mouse *IL-6*, mouse *IL-10*, mouse *CXCL1*, mouse *CXCL2*, mouse *CCL2*, mouse *CCL7*, mouse *TLR2*, mouse *TLR4*, and mouse *TLR9*. The primer sequences are listed in **Table 1**. Mouse *GAPDH* was used as an internal control to quantify mouse *TNF-α*, mouse *IL-1β*, mouse *IL-6*, mouse *IL-10*, mouse *CXCL1*, mouse *CXCL2*, mouse *CCL2*, mouse *CCL7*, mouse *TLR2*, mouse *TLR4*, and mouse *TLR9*. Real-time PCR was performed using 40 cycles of denaturation at 95°C for 15 s, annealing at 62°C for 15 s, and extension at 72°C for 1 min. The amplified PCR products were quantitatively monitored using a StepOne Real-Time PCR System (Applied Biosystems, Foster City, CA). Fold changes in the expression level of each gene were calculated by the 2<sup>-ΔΔCt</sup> method using the mouse *GAPDH* gene as an internal control. The relative expression of each gene was evaluated relative to the control sample (uninfected mouse), which was assigned an arbitrary unit value of 1.

## Statistical Analyses

Survival curves with log rank test were obtained using EZR (Saitama Medical Center, Jichi Medical University, Saitama, Japan), which is a graphical user interface for R (The R Foundation for Statistical Computing, Vienna, Austria). More precisely, it is a modified version of R commander designed to add statistical functions frequently used in biostatistics (42). Quantitative results were compiled from more than two independent experiments. The results are presented as the

**TABLE 1** | Primers used for real-time PCR.

Mouse Gene	Sequence	Reference
<i>TNF-α</i>	F: ATGATCCGCGACGTGGAA	(20)
	R: CTGCCACAAGCAGGAATGAG	
<i>IL-1β</i>	F: CGCAGCAGCACATCAACAAGAGC	(20)
	R: TGTCCTCATCCTGGAAGGTCCACG	
<i>IL-6</i>	F: CCAGAGATACAAAGAAATGATGG	(20)
	R: ACTCCAGAAGACCAGAGGAAAA	
<i>IL-10</i>	F: CAGAGCCACATGCTCCTAGA	(20)
	R: GTCCAGCTGGTCCITTTGTTT	
<i>CXCL1 (KC)</i>	F: CCAGCCCACTCCAACACAGC	(38)
	R: AGGGAGCTTCAGGGTCAAGGC	
<i>CXCL2 (MIP-2)</i>	F: ACCAACCACCAGGCTACAG	(20)
	R: GCGTCACACTCAAGCTCT	
<i>CCL2 (MCP-1)</i>	F: TTA AAAACCTGGATCGGAACCAA	(39)
	R: GCATTAGCTTCAGATTTACGGGT	
<i>CCL7 (MCP-3)</i>	F: TCCACATGCTGCTATGCAAG	(40)
	R: CATGTCTAAGTATGCTATAGCC	
<i>TLR2</i>	F: TGGAATGTCACCAGGCTGC	(41)
	R: GTCCGTGGAATGGTGGC	
<i>TLR4</i>	F: AGGAAGTTTCTCTGGACTAACAAGTTTAGA	(41)
	R: AAATTGTGAGCCACATTGAGITTC	
<i>TLR9</i>	F: TTCTCAAGACGGTGGATCGC	(41)
	R: GCAGAGGTTGCTTCTCACG	
<i>GAPDH</i>	F: CTTCAACCACCATGGAGAAGGC	(20)
	R: GGCATGGACTGTGGTTCATGAG	

F, forward primer; R, reverse primer.

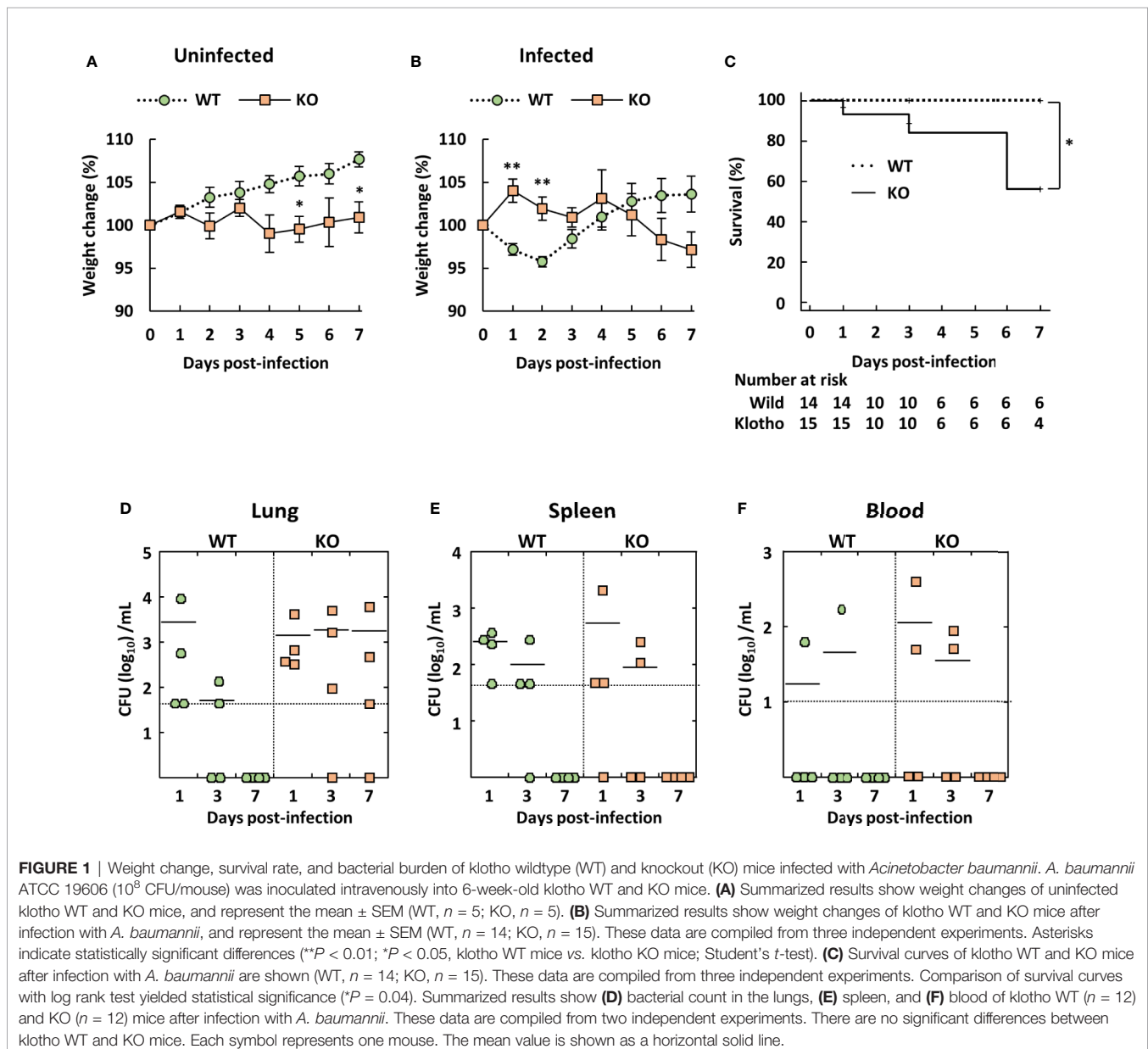
mean  $\pm$  standard error of the mean (SEM). Comparisons of numerical data were performed using the Student's *t*-test. Pearson correlation analysis was used to compare the mRNA levels of proinflammatory cytokines and *TLR2*, 4, and 9. In all analyses, a two-tailed probability of  $< 5\%$  (i.e.,  $*P < 0.05$ ) was considered statistically significant.

## RESULTS

### Klotho KO Mice Infected With *A. baumannii* Rarely Display Severe Symptoms Prior to Death

Young C57BL/6 mice reportedly showed low susceptibility to *A. baumannii* ATCC 19606 infection (35, 36). Therefore, we

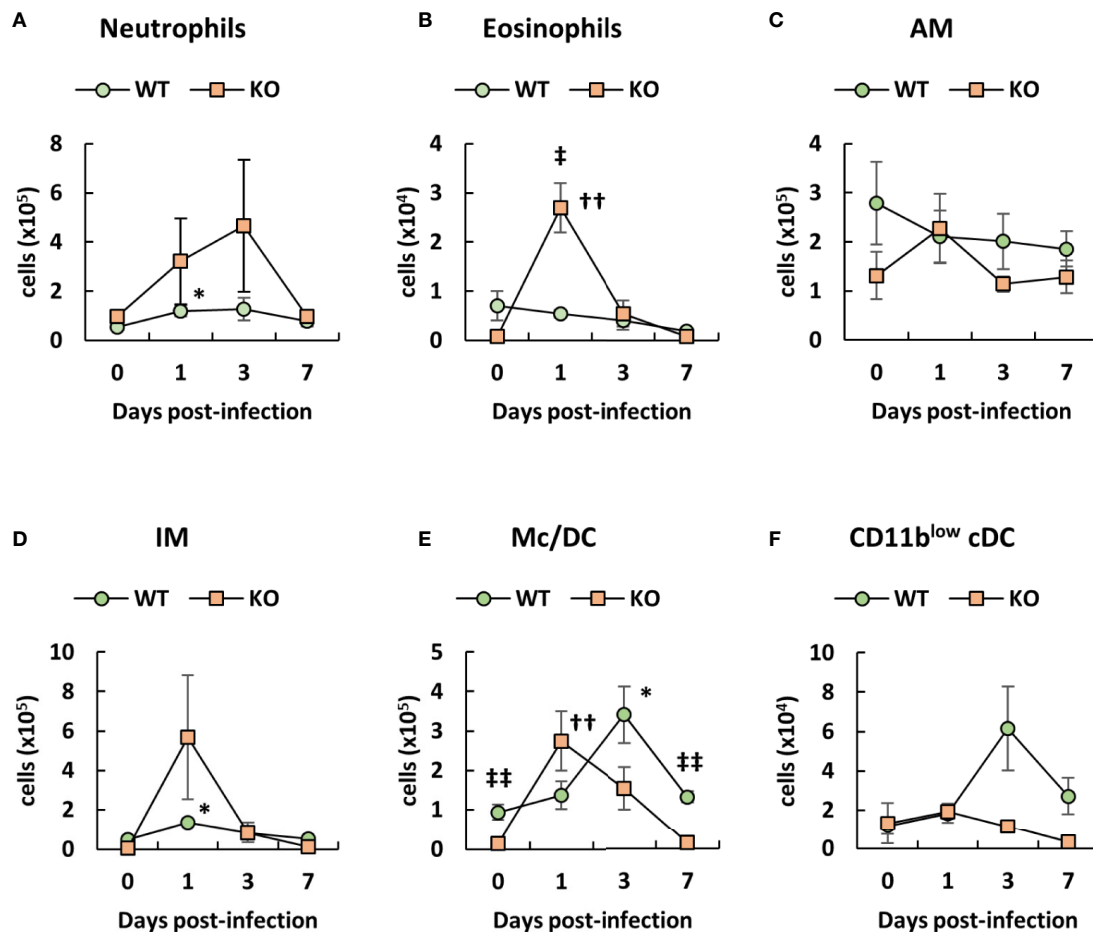
evaluated body weight, clinical score, and survival rate of klotho WT and KO mice after infection *via* intravenous inoculation of *A. baumannii* ATCC 19606. The body weight of uninfected klotho WT mice gradually increased, whereas that of uninfected klotho KO mice remained stable (**Figure 1A**). Following infection, the body weight of infected klotho WT mice significantly decreased until 2 days post-infection, but had recovered to the pre-infection level at 4 days post-infection. The body weight of klotho KO mice increased at 1 day post-infection and subsequently decreased gradually until 7 days post-infection (**Figure 1B**). A clinical score (ruffled fur) was observed for one of 15 klotho KO mice at 1 day post-infection and one of six klotho KO mice (inactive unless prodded) at 6 days post-infection. The other klotho KO mice infected with *A. baumannii* did not display any clinical



symptoms. Additionally, although klothe WT mice did not die of *A. baumannii* infection, klothe KO mice were more susceptible, as evidenced by the 56% survival rate at 7 days post-infection (Figure 1C). These results suggested that klothe KO mice rarely displayed severe symptoms of *A. baumannii* infection prior to their death. To clarify whether bacteria were eliminated from klothe KO mice, the bacteria in the lungs, spleen, and blood of klothe WT and KO mice were enumerated after infection with *A. baumannii*. Bacteria were not detected in the lungs, spleen, and blood of klothe WT mice at 7 days post-infection (Figures 1D–F; WT). In contrast, bacteria were detected in the lungs of three of four klothe KO mice at 7 days post-infection (Figure 1D; KO), but not in the spleen and blood (Figures 1E, F; KO). These results suggested that *A. baumannii* escaped from host immune responses in the lungs of klothe KO mice and subsequently colonized the lungs of the aged hosts.

## Transient Induction of Neutrophils, Eosinophils, Interstitial Macrophages, and Monocytes/Dendritic Cell (Subset in the Lungs of Klothe KO Mice After Infection With *A. baumannii*)

Since neutrophils and macrophages play an important role in the clearance of *A. baumannii* (21–23), we analyzed the number of immune cells in the lungs of klothe WT and KO mice after infection with *A. baumannii*, based on the classification of the lung innate immune cell population (43–45). Representative results from uninfected and *A. baumannii*-infected klothe mice at 1 day post-infection are shown in Figures S1 (WT) and S2 (KO). The results from the two groups are summarized in Figure 2. Flow cytometry revealed that the number of neutrophils in the lungs of klothe KO mice gradually increased until 3 days post-infection and decreased at 7 days post-infection, whereas that of



**FIGURE 2** | Number of innate immune cells in the lungs of klothe wildtype (WT) and knockout (KO) mice before and after infection with *Acinetobacter baumannii*. Summarized results show the number of (A) neutrophils, (B) eosinophils, (C) alveolar macrophages (AM), (D) interstitial macrophages (IM), (E) monocytes/dendritic cells (Mc/DC), and (F) CD11b<sup>low</sup> conventional DC (cDC) in the lungs of *A. baumannii*-infected klothe WT and KO mice. Line graphs were compiled from 4 independent experiments ( $n = 16$  for WT and  $n = 15$  for KO mice), and represent the mean  $\pm$  SEM. Asterisks indicate statistically significant differences (\* $P < 0.05$ , klothe WT mice before infection vs. klothe WT mice after infection, Student's  $t$ -test; <sup>‡‡</sup> $P < 0.01$ , klothe KO mice before infection vs. klothe KO mice after infection, Student's  $t$ -test; <sup>‡‡</sup> $P < 0.01$ , klothe WT mice vs. klothe KO mice, Student's  $t$ -test).

klortho WT mice significantly increased at 1 day post-infection (**Figure 2A**). The number of eosinophils in the lungs of klortho KO mice significantly increased at 1 day post-infection and subsequently decreased until 7 days post-infection, whereas klortho WT mice showed no induction of eosinophils after infection (**Figure 2B**). The number of AM in the lungs of klortho KO mice was slightly increased at 1 day post-infection and subsequently decreased to the pre-infection level at 3 days post-infection, whereas the cell number in klortho WT mice slightly decreased after infection (**Figure 2C**). Additionally, the number of AM in klortho KO mice was lower than that in klortho WT mice, except for 1 day post-infection (**Figure 2C**). The number of IMs increased in the lungs of klortho KO mice at 1 day post-infection and subsequently decreased at 3 days post-infection. (**Figure 2D**). The number of CD11b<sup>+/hi</sup>CD11c<sup>-/low</sup> monocytes plus CD11b<sup>hi</sup>CD11c<sup>low</sup> Mc/DC subset in the lungs of klortho KO mice increased at 1 day post-infection and subsequently decreased to the pre-infection level by 7 days post-infection, whereas that of klortho WT mice gradually increased until 3 days post-infection and was decreased at 7 days post-infection (**Figure 2E**). The number of CD11b<sup>low</sup> conventional DC in the lungs of klortho KO mice hardly increased after the infection, whereas that of klortho WT mice increased at 3 days post-infection and subsequently decreased at 7 days post-infection (**Figure 2F**). These results suggested that innate immune cells in the lungs of klortho KO mice were transiently induced after infection of *A. baumannii*. However, this transient change did not eliminate *A. baumannii* in the lungs (**Figure 1D**).

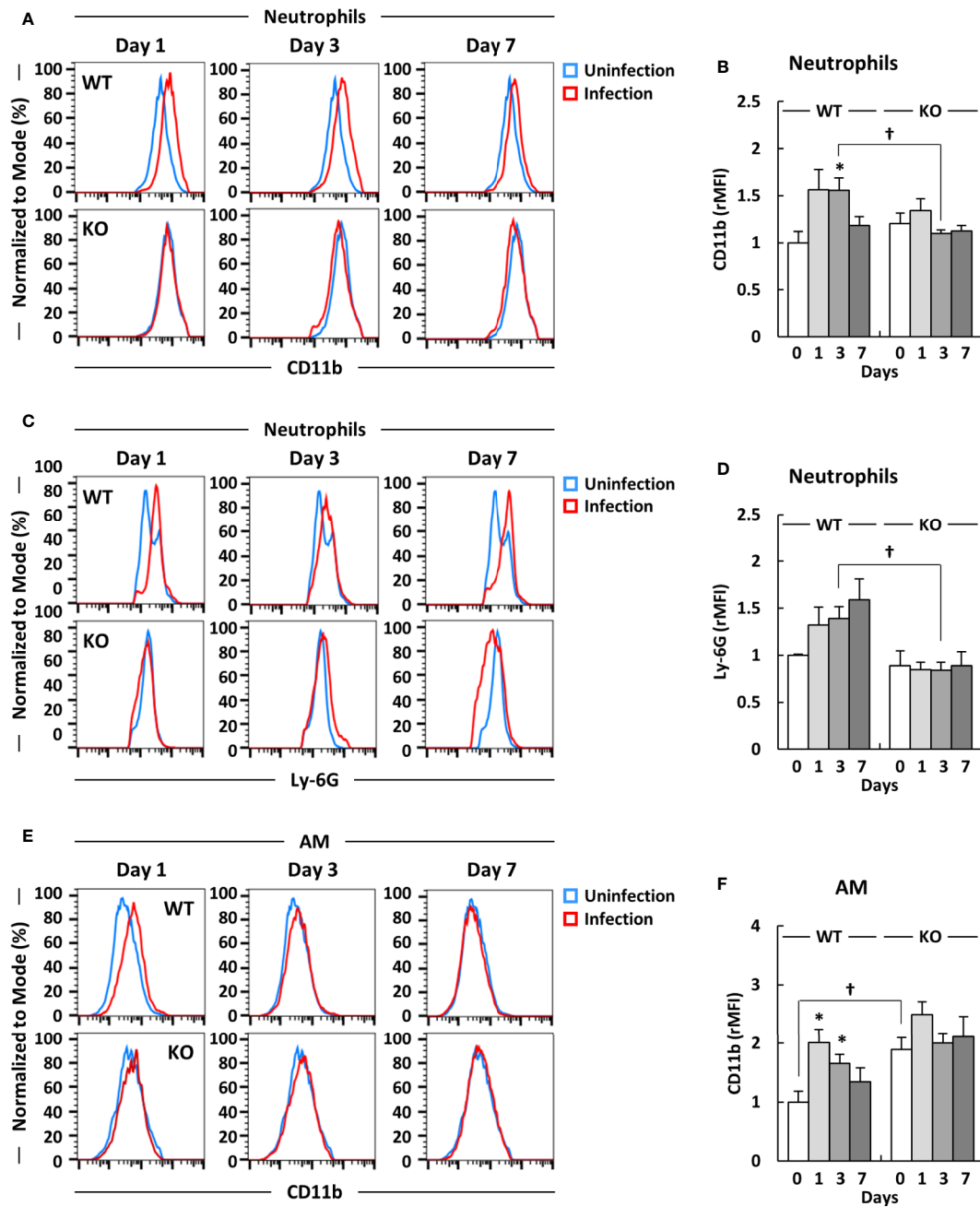
### Negligible Activation of Neutrophils in the Lungs of Klortho KO Mice Infected With *A. baumannii*

Previous studies have reported that neutrophils are activated in inflammatory conditions as well as in response to lipopolysaccharide (LPS) stimulation, with the upregulation of markers of the integrin family, such as CD11b, as well as other activation markers (46, 47). Although neutrophils were induced in klortho KO mice after infection with *A. baumannii*, the burden of the lung bacteria was not completely eliminated in the mice at 7 days post-infection. Therefore, we evaluated the expression level of CD11b on neutrophils to determine whether neutrophils were activated in the lungs of klortho KO mice after infection with *A. baumannii*. Representative results from uninfected and *A. baumannii*-infected klortho mice are shown in **Figure 3A** and the complete results are summarized in **Figure 3B**. The expression level of CD11b on neutrophils in the lungs of klortho WT mice was significantly increased after infection with *A. baumannii* (**Figures 3A, B**; WT). In contrast, the expression level of CD11b on neutrophils in the lungs of klortho KO mice was slightly increased at only 1 day post-infection (**Figures 3A, B**; KO). Moreover, there was a significant difference between the expression level of CD11b on neutrophils in the lungs of klortho KO mice and WT mice at 3 days post-infection (**Figure 3B**). In addition, the expression level of Ly-6G as a maturation marker of neutrophils was analyzed in klortho WT and KO mice after

infection with *A. baumannii*. The expression level of Ly-6G on neutrophils in the lungs of klortho WT mice gradually increased after the infection, whereas that of klortho KO mice hardly increased after infection with *A. baumannii* (**Figures 3C, D**). There was a significant difference between the expression level of Ly-6G on neutrophils in the lungs of klortho WT mice and KO mice at 3 days post-infection. These results suggest that neutrophils in the lungs of klortho KO mice impair their activation and functional maturation. We further analyzed the expression level of CD11b on AM, because the expression is upregulated by host-derived inflammatory stimuli (48, 49). Representative results from uninfected and *A. baumannii*-infected klortho mice are shown in **Figure 3E** and the collective results are summarized in **Figure 3F**. The expression level of CD11b on AM in the lungs of klortho WT mice was significantly increased at 1 day post-infection and subsequently gradually decreased (**Figures 3E, F**; WT). The expression level of CD11b on AM in the lungs of klortho KO mice before infection was significantly higher than that of klortho WT mice before infection and slightly increased at 1 day post-infection (**Figures 3E, F**; KO). These results suggested that although AM are already activated because of aging, it only slightly responded to *A. baumannii* infection.

### Proinflammatory Cytokines Are Strongly Induced in the Lungs of Klortho KO Mice Infected With *A. baumannii*

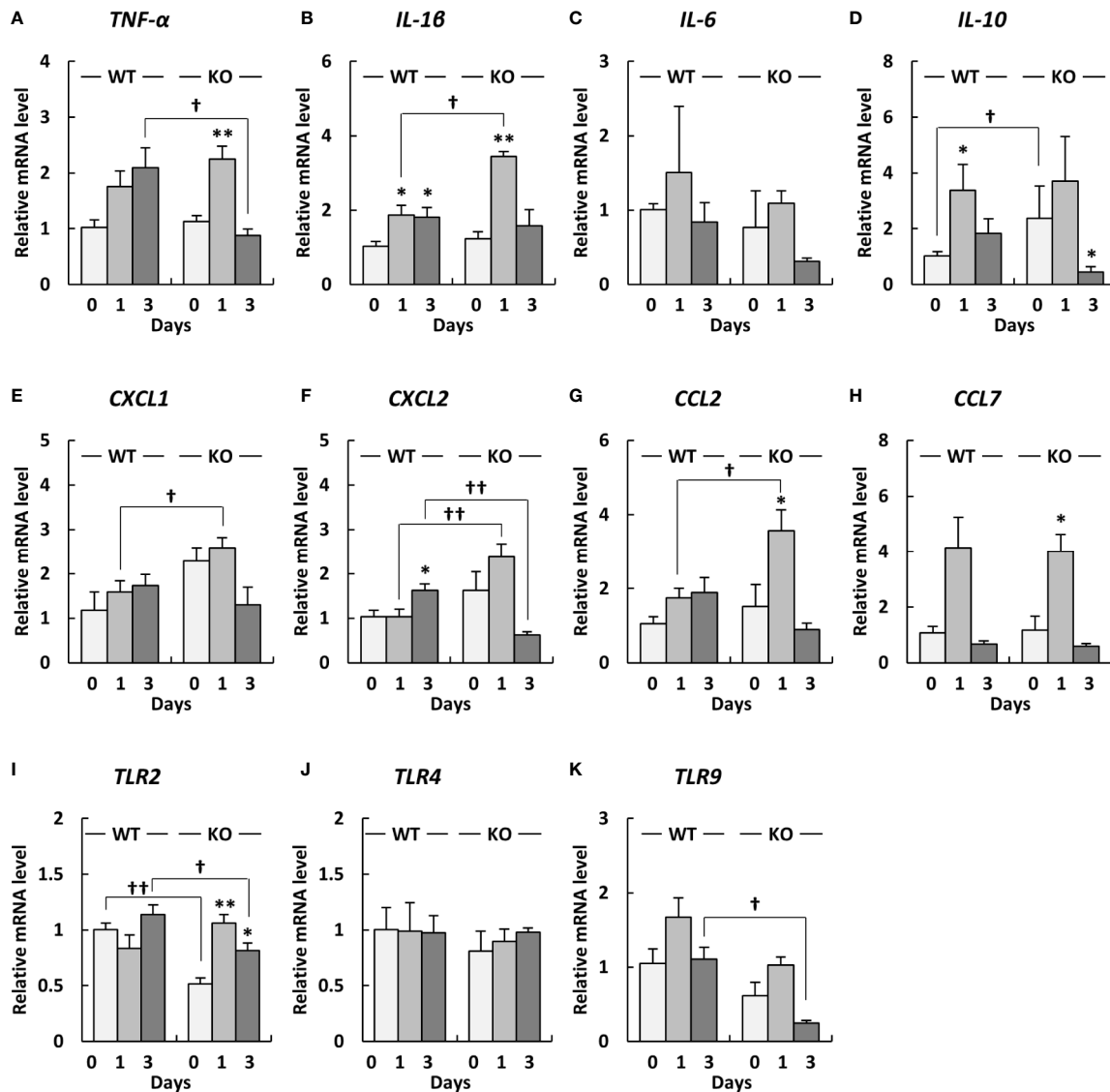
To investigate inflammatory responses in the lungs of klortho WT and KO mice after infection with *A. baumannii*, we analyzed the mRNA expression of proinflammatory cytokines in the lungs of the mice after infection with *A. baumannii*. The level of *TNF- $\alpha$*  mRNA in the lungs of klortho WT mice gradually increased after infection, whereas those of klortho KO mice significantly increased at 1 day post-infection and subsequently decreased at 3 days post-infection (**Figure 4A**). Additionally, the level of *TNF- $\alpha$*  mRNA in the lungs at 3 days post-infection was significantly lower in klortho KO mice than in klortho WT mice (**Figure 4A**). The level of *IL-1 $\beta$*  mRNA in the lungs of klortho WT mice significantly increased at 1 day post-infection and was maintained at 3 days post-infection, whereas those in the lungs of klortho KO mice strongly and significantly increased at 1 day post-infection and subsequently decreased at 3 days post-infection (**Figure 4B**). The level of *IL-1 $\beta$*  mRNA was significantly higher in the lungs of klortho KO mice than in those of klortho WT mice at 1 day post-infection (**Figure 4B**). The level of *IL-6* mRNA in the lungs of klortho WT mice slightly increased at 1 day post-infection and subsequently decreased at 3 days post-infection (**Figure 4C**). The level of *IL-6* mRNA in klortho KO mice was not increased at 1 day post-infection and was decreased at 3 days post-infection (**Figure 4C**). The level of *IL-10* mRNA before infection was significantly higher in the lungs of klortho KO mice than in those of klortho WT mice (**Figure 4D**). The level of *IL-10* mRNA in the lungs of klortho WT mice significantly increased at 1 day post-infection and decreased at 3 days post-infection (**Figure 4D**). The level of *IL-10* mRNA in the lungs of klortho KO mice increased at 1 day post-infection



**FIGURE 3** | Expression levels of CD11b and Ly-6G on neutrophils, and CD11b on alveolar macrophages (AM) in the lungs of klotho wildtype (WT) and knockout (KO) mice before and after infection with *Acinetobacter baumannii*. Representative results on (A) CD11b and (C) Ly-6G expression levels on neutrophils and (E) CD11b expression level of alveolar macrophages (AM) in the lungs of uninfected and *A. baumannii*-infected klotho WT and KO mice. Mean fluorescence intensity was measured in uninfected (blue line) and infected (red line) klotho mice using flow cytometry. Summarized results on the expression levels of (B) CD11b and (D) Ly-6G on neutrophils and (F) that of CD11b on AM in the lungs of klotho WT and KO mice before and after infection with *A. baumannii*. Bar graph data were compiled from four independent experiments ( $n = 16$  for WT and  $n = 15$  for KO mice), and represent the mean  $\pm$  SEM. Asterisks indicate statistically significant differences ( $*P < 0.05$ , before infection vs. after infection, Student's  $t$ -test;  $^\dagger P < 0.05$ , klotho WT mice vs. klotho KO mice, Student's  $t$ -test).

and subsequently decreased significantly (Figure 4D). The level of *CXCL1* mRNA in the lungs of klotho WT mice gradually increased after infection (Figure 4E). The level of *CXCL1* mRNA in the lungs before infection was higher in klotho KO mice than

in klotho WT mice (Figure 4E). Additionally, the level of *CXCL1* mRNA in the lungs at 1 day post-infection was significantly higher in klotho KO mice than in klotho WT mice (Figure 4E). The level of *CXCL2* mRNA in the lungs of klotho KO mice



**FIGURE 4** | Proinflammatory cytokine expression levels in the lungs and Toll-like receptor (*TLR*) 2/4/9 gene expression levels in the lungs of klotho wildtype (WT) and knockout (KO) mice before and after infection with *Acinetobacter baumannii*. The mRNA levels of (A) *TNF-α*, (B) *IL-1β*, (C) *IL-6*, (D) *IL-10*, (E) *CXCL1*, (F) *CXCL2*, (G) *CCL2*, (H) *CCL7*, (I) *TLR2*, (J) *TLR4*, and (K) *TLR9* in the lungs of klotho WT and KO mice before and after infection with *A. baumannii*. Bar graph data represent the mean  $\pm$  SEM and were compiled from four independent experiments ( $n = 12$  for WT and  $n = 12$  for KO mice). Asterisks indicate statistically significant differences (\*\* $P < 0.01$ ; \* $P < 0.05$ , before infection vs. after infection, Student's *t*-test; †† $P < 0.01$ ; † $P < 0.05$ , klotho WT mice vs. klotho KO mice, Student's *t*-test).

increased at 1 day post-infection and subsequently decreased at 3 days post-infection (Figure 4F). The level of *CXCL2* mRNA was significantly higher in the lungs of klotho KO mice than in those of klotho WT mice at 1 day post-infection (Figure 4F). The level of *CCL2* mRNA in the lungs of klotho WT mice gradually increased after infection, whereas those of klotho KO mice significantly increased at 1 day post-infection and subsequently decreased at 3 days post-infection (Figure 4G). The level of *CCL2* mRNA in the lungs at 1 day post-infection was significantly higher in klotho KO mice than in klotho WT mice (Figure 4G). The level of *CCL7* mRNA increased in the lungs of klotho WT and KO mice at 1 day post-infection and

subsequently decreased at 3 days post-infection (Figure 4H). The expression levels of *CCL7* mRNA were comparable between klotho WT and KO mice after infection (Figure 4H). The collective results suggested that proinflammatory responses were strongly and transiently induced in the lungs of klotho KO mice after infection with *A. baumannii*.

### Expression Levels of Proinflammatory Cytokines Correlate With Those of *TLR9* in the Lungs of Klotho KO Mice

Since TLRs play an important role in eliciting proinflammatory cytokines during infection with *A. baumannii* (24–26), we



analyzed the expression levels of *TLR2*, 4, and 9 in the lungs of klotho WT and KO mice after infection with *A. baumannii*. The level of *TLR2* mRNA was significantly lower in the lungs of uninfected KO mice than in those of uninfected klotho WT mice (Figure 4I). Following infection, the level of *TLR2* mRNA in the lungs of klotho WT mice was stable but was significantly increased in KO mice at 1 and 3 days post-infection. The level of *TLR2* mRNA was significantly lower in the lungs of klotho KO mice than in those of klotho WT mice at 3 days post-infection. In addition, the level of *TLR9* mRNA was lower in the lungs of uninfected KO mice than in those of klotho WT mice (Figure 4K). Following infection, the levels of *TLR9* mRNA in the lungs of klotho WT and KO mice increased at 1 day post-infection and subsequently decreased at 3 days post-infection. However, the level of *TLR9* mRNA in the lungs of klotho KO mice was significantly decreased compared with that of klotho WT mice at 3 days post-infection. In contrast, the levels of *TLR4* mRNA were comparable between klotho WT and KO mice, and were not altered in their lungs after the infection (Figure 4J). We further analyzed the correlation between the expression levels of proinflammatory cytokines and those of TLRs in the lungs of klotho WT and KO mice after infection with *A. baumannii*. Pearson correlation analysis revealed that the mRNA level of *CXCL2* was significantly correlated with that of *TLR2* in the lungs of klotho WT mice (Figure 5F). In contrast, mRNA levels of *TNF- $\alpha$*  and *IL-1 $\beta$*  were significantly correlated with those of *TLR2* in the lungs of klotho KO mice (Figures 5A, B). Additionally, mRNA levels of *IL-6*, *IL-10*, *CXCL1*, *CCL2*, and *CCL7* were not correlated with those of *TLR2* in the lungs of klotho WT and KO mice (Figures 5C–E, G, H). Moreover, mRNA levels of *CCL2* and *CCL7* were significantly correlated with those of *TLR9* in the lungs of klotho WT mice (Figures 6G, H). In contrast, the mRNA levels of *TNF- $\alpha$* , *IL-1 $\beta$* , *IL-10*, *CXCL1*, *CXCL2*, *CCL2*, and *CCL7* were strongly and significantly correlated with those of *TLR9* in the lungs of klotho KO mice (Figures 6A, B, D–H). Additionally, mRNA level of *IL-6* was not correlated with that of *TLR9* in the lungs of klotho WT and KO mice (Figure 6C). These results indicated the important roles of *TLR9* in eliciting proinflammatory responses in the lungs of aged hosts against *A. baumannii* infection.

### Different Immune Cell Induction Pattern in the Blood of Klotho KO Mice After Infection With *A. baumannii*

We analyzed the number of immune cells in the blood of klotho WT and KO mice after infection with *A. baumannii*. Representative results from uninfected and *A. baumannii*-infected klotho mice at 1 day post-infection are shown in Figures S3 (WT) and S4 (KO). The results from the two groups are summarized in Figures 7A–D. Flow cytometry revealed that the number of neutrophils in the blood before infection was lower in klotho KO mice than in klotho WT mice (Figure 7A). The number of neutrophils in the blood of klotho WT mice significantly decreased at 1 day-post infection and subsequently increased gradually (Figure 7A). In contrast, the number of neutrophils in the blood of klotho KO mice increased

at 3 days post-infection and subsequently decreased at 7 days post-infection (Figure 7A). The number of eosinophils significantly decreased in the blood of klotho WT and KO mice at 1 day-post infection and subsequently increased (Figure 7B). The number of monocytes in the blood of klotho KO mice significantly increased at 1 day post-infection and subsequently decreased at 3 days post-infection (Figure 7C). However, the number of monocytes in the blood before and after infection was lower in klotho KO mice than in klotho WT mice (Figure 7C). The number of macrophages in the blood of klotho KO mice significantly increased at 1 day post-infection and subsequently decreased gradually, whereas those of klotho WT mice increased at 3 day post-infection and subsequently decreased at 7 days post-infection (Figure 7D).

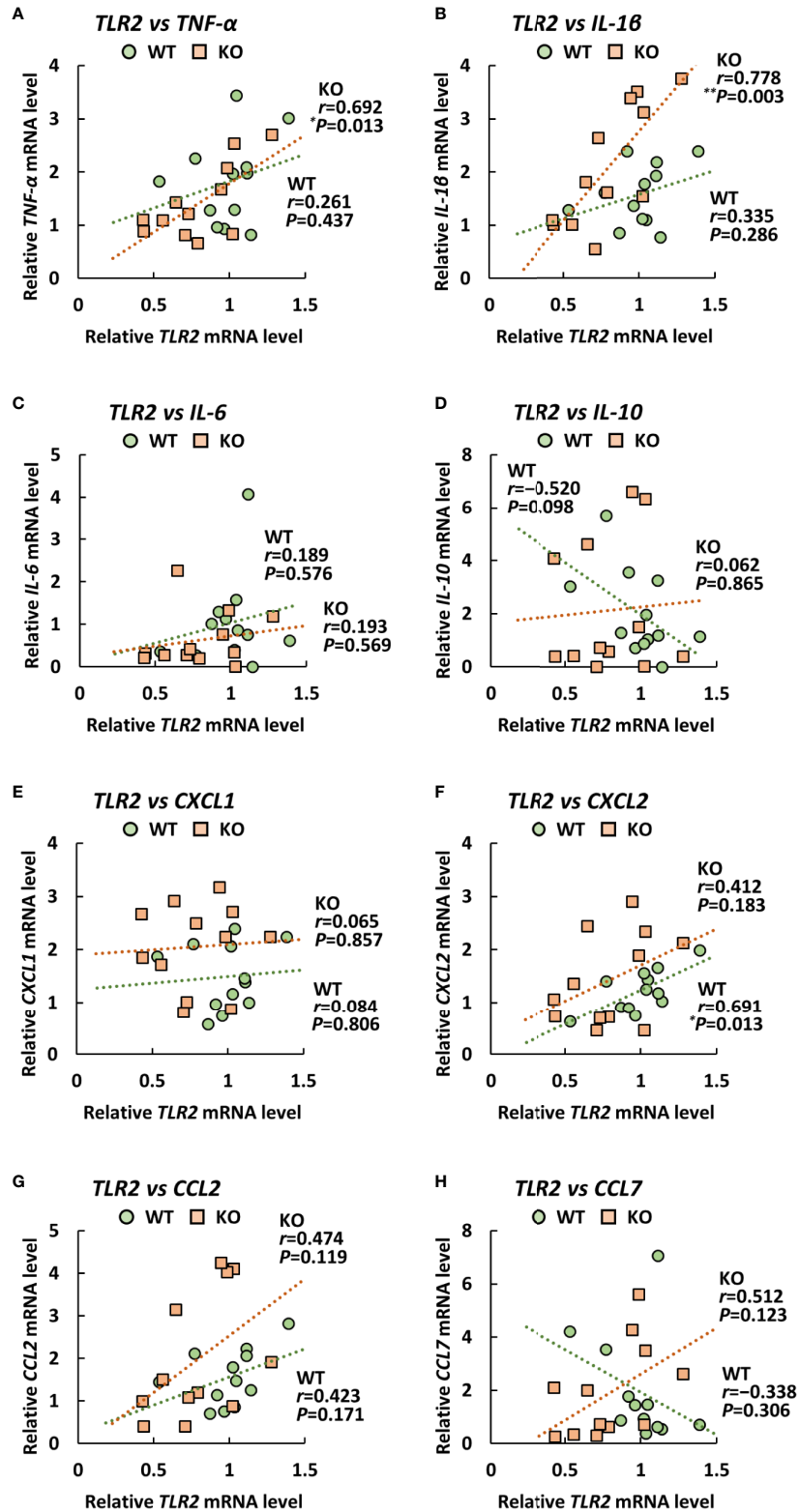
### Klotho KO Mice Infected With *A. baumannii* Display Different Cytokine Production Pattern in Their Sera

We analyzed the levels of proinflammatory cytokines in the sera of klotho WT and KO mice before and after infection with *A. baumannii*. *TNF- $\alpha$* , *IL-6*, *CXCL1*, *CCL2*, and *CCL7* levels significantly increased in the sera of klotho WT and KO mice at 1 day post-infection, while the level of *IL-12* was not (Figures 7E, F, I–K). Of note, although the serum level of *IL-6*, *CXCL1*, and *CCL7* in klotho WT mice decreased at 3 days post-infection, those in klotho KO mice was maintained until 3 days post-infection (Figures 7F, I, K). The serum level of *IL-10* increased in klotho WT and KO mice after infection and was significantly higher in klotho KO mice than in klotho WT mice at 1 day post-infection (Figure 7G). The serum level of *IFN- $\gamma$*  significantly increased in klotho WT mice at 1 day post-infection, whereas that did not increase in infected klotho KO mice (Figure 7H). These results suggested that proinflammatory responses were prolonged in the periphery of klotho KO mice infected with *A. baumannii*.

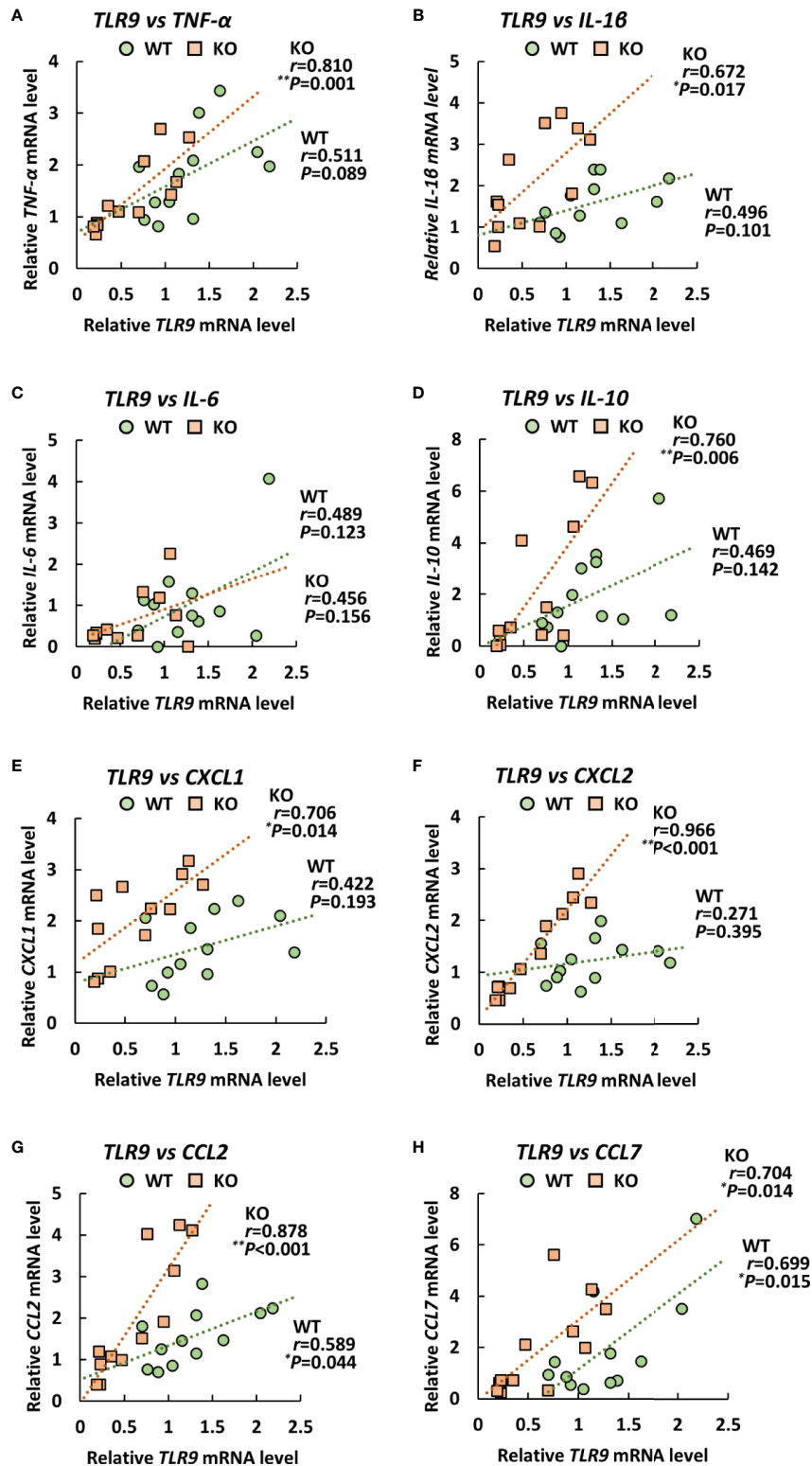
## DISCUSSION

*A. baumannii* has recently emerged as a major nosocomial pathogen (1, 2). Although we have previously reported the virulence characteristics of *A. baumannii* clinical isolates in human epithelial cells and macrophages *in vitro* (13, 14, 20), the pathogenicity of *A. baumannii* infection in elderly people has remained elusive. In the current study, we established an aged mouse model of *A. baumannii* infection using klotho KO mice and evaluated the immune responses in the mice before and after infection.

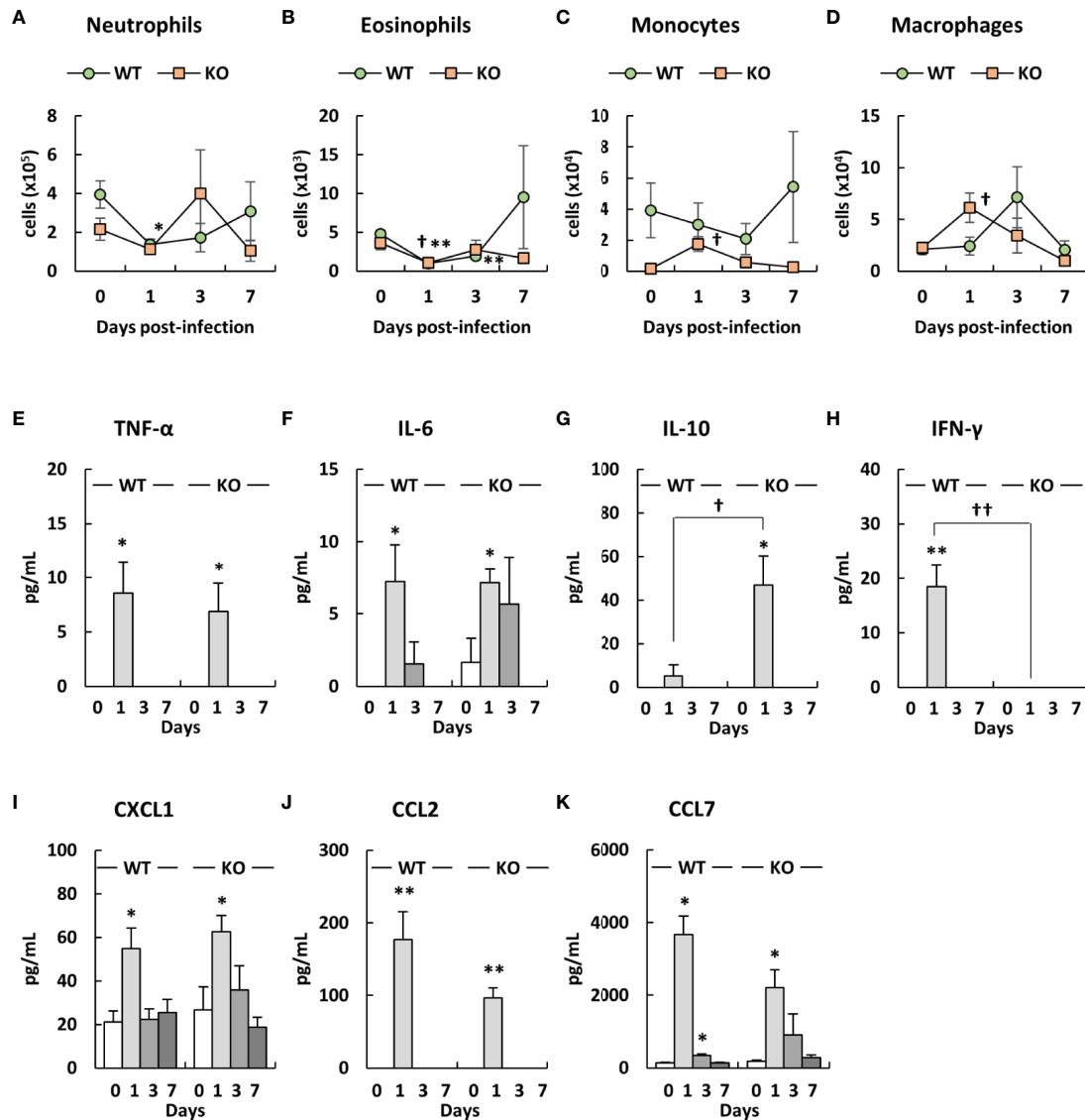
Previous study reported an aged pneumonia mouse model by non-invasive intratracheal inoculation with *A. baumannii* LAC-4 strain (27). Higher mortality was observed in aged mice along with increased bacterial burdens and more severe lung injury. In addition, increased inflammatory cell infiltration and enhanced pro-inflammatory cytokines at 24 h post-infection were detected in aged mice. *A. baumannii* LAC-4 strain is much more virulent than *A. baumannii* ATCC 19606 strain (36). As *A. baumannii*



**FIGURE 5** | Relationship between the expression of proinflammatory cytokines and Toll-like receptor 2 (*TLR2*) in klothe wildtype (WT) and knockout (KO) mice. Pearson correlation coefficient was calculated between the mRNA level of *TLR2* and that of (A) *TNF- $\alpha$* , (B) *IL-1 $\beta$* , (C) *IL-6*, (D) *IL-10*, (E) *CXCL1*, (F) *CXCL2*, (G) *CCL2*, (H) *CCL7* in klothe WT and KO mice. Each symbol represents one mouse. Asterisks indicate statistically significant correlations (\*\* $P < 0.01$ ; \* $P < 0.05$ ).



**FIGURE 6** | Relationship between the expression of proinflammatory cytokines and Toll-like receptor 9 (*TLR9*) in klotho wildtype (WT) and knockout (KO) mice. Pearson correlation coefficient was calculated between the mRNA level of *TLR9* and that of (A) *TNF- $\alpha$* , (B) *IL-1 $\beta$* , (C) *IL-6*, (D) *IL-10*, (E) *CXCL1*, (F) *CXCL2*, (G) *CCL2*, (H) *CCL7* in klotho WT and KO mice. Each symbol represents one mouse. Asterisks indicate statistically significant correlations (\*\* $P < 0.01$ ; \* $P < 0.05$ ).



**FIGURE 7** | Number of innate immune cells in the blood and proinflammatory cytokine expression levels in the sera of klotheo wildtype (WT) and knockout (KO) mice before and after infection with *Acinetobacter baumannii*. Summarized results show the number of (A) neutrophils, (B) eosinophils, (C) monocytes, and (D) macrophages in the blood of *A. baumannii*-infected klotheo WT and KO mice. Line graphs were compiled from four independent experiments ( $n = 16$  for WT and  $n = 16$  for KO mice), and represent the mean  $\pm$  SEM. Asterisks indicate statistically significant differences ( $*P < 0.05$ , klotheo WT mice before infection vs. klotheo WT mice after infection, Student's  $t$ -test;  $^{\dagger}P < 0.01$ , klotheo KO mice before infection vs. klotheo KO mice after infection, Student's  $t$ -test;  $^{**}P < 0.01$ , klotheo WT mice vs. klotheo KO mice, Student's  $t$ -test). The production of (E) TNF- $\alpha$ , (F) IL-6, (G) IL-10, (H) IFN- $\gamma$ , (I) CXCL1, (J) CCL2, and (K) CCL7 in the sera of klotheo WT and KO mice before and after infection with *A. baumannii*. Bar graph data represent the mean  $\pm$  SEM and were compiled from four independent experiments ( $n = 16$  for WT and  $n = 16$  for KO mice). Asterisks indicate statistically significant differences ( $^{**}P < 0.01$ ;  $*P < 0.05$ , before infection vs. after infection, Student's  $t$ -test;  $^{\dagger}P < 0.01$ ;  $^{\ddagger}P < 0.05$ , klotheo WT mice vs. klotheo KO mice, Student's  $t$ -test).

ATCC 19606 strain was used in this study, klotheo KO mice seldom displayed severe symptoms of *A. baumannii* infection but 44% died. Moreover, bacteria were not completely eliminated in the lungs of these mice. These results suggest that *A. baumannii* increases its bacterial burdens and inflammatory cell infiltration in the lungs of aged hosts and can eventually lead to death. Neutrophils and AM play important roles in the clearance of *A. baumannii* (21–23). Although the number of

neutrophils increased in the lungs of klotheo KO mice after infection with *A. baumannii*, the expression level of CD11b on neutrophils was hardly upregulated. Previous studies have reported that the expression level of CD11b on neutrophils is upregulated by LPS stimulation via TLR4 (46, 47). However, signal transduction and functional changes were observed in neutrophils with aging (50). Considering the above findings, our results suggest that TLR4-mediated signaling was weak in the

neutrophils of klotho KO mice. The age-related disruption in the function and/or differentiation of these cells in klotho KO mice resulted in a bacterial burden in the lungs of the mice. Additionally, there were fewer AM in klotho KO mice than in klotho WT mice, except for 1 day post-infection. The level of expression of CD11b on AM in the lungs of uninfected klotho KO mice was significantly higher than that of uninfected klotho WT mice. Previous studies demonstrated the upregulated expression of CD11b on AM because of initial inflammation or LPS stimulation (48, 49). However, another study reported that the expression of CD11b is upregulated in AM in old mice and the cells are significantly better in phagocytosing *Mycobacterium tuberculosis* than the CD11b<sup>-</sup> AM population, which enhanced bacterial growth (51). These results suggest *M. tuberculosis* phagocytosis and survival are enhanced in aged hosts. Likewise, *A. baumannii* resist toxicity caused by ROS and survive in macrophages (20). Considering the above findings, the phagocytosis of *A. baumannii* by AM protects the bacteria from the lethal action of neutrophils and allows the bacteria to establish an intractable infection in aged hosts. Conversely, there were more IMs in the lungs of klotho KO mice infected with *A. baumannii*. A previous study reported that IMs are induced by acute lung injury caused by, for instance, high doses of LPS, and perform important immune functions, including the maintenance of lung homeostasis (52). These results suggest that acute inflammation is elicited in the lungs of aged hosts early during *A. baumannii* infection.

Inflammatory monocytes are recruited into tissues and kill pathogens via phagocytosis, ROS activity, production of nitric oxide (NO), and the action of inflammatory cytokines (53, 54). Although we analyzed ROS production in monocytes as well as neutrophils in klotho KO mice infected with *A. baumannii*, it was not clear whether those cells induced ROS in response to the infection. Further studies are required to clarify the production of ROS in those cells. On the other hands, inflammatory monocytes upregulate levels of CD11c and MHC class II, which can promote T cell proliferation (53). In the present study, inflammatory monocytes in the lungs, spleen (**Figures S5, S6, and S7**), and blood (**Figures 7, S3, and S4**) of klotho KO mice infected with *A. baumannii* were transiently induced. However, the number of inflammatory monocytes in those tissues were lower in klotho KO mice than in klotho WT mice. These results suggest that *A. baumannii* infection is complicated by the low induction of inflammatory monocytes in aged hosts. Additionally, klotho KO mice infected with *A. baumannii* showed lower induction of DC subsets than infected klotho WT mice (**Figures 2 and S7**). These results suggest that age-associated changes in the proportion of monocyte and DC subsets affect adaptive immune function, thereby reducing the efficacies of antibiotics and vaccines in aged hosts (27).

Although the present study suggests the disruption in the function and/or differentiation of immune cells in klotho KO mice, it remains unclear how klotho KO mice died from infection with *A. baumannii*. A previous study reported that klotho deficiency aggravates sepsis-related multiple organ dysfunction (31). Thus, one possible reason for eventually death might be

incomplete bacteria clearance and worsen multiple organ dysfunction. Hearps *et al.* reported that monocytes from older individuals exhibit impaired phagocytosis but contain shortened telomeres, suggesting a dysregulation of monocytes function in the aged hosts (55). Moreover, it is suggested that telomere shortening in peripheral monocytes reflects telomere shortening in haematopoietic bone marrow precursor cells (55). Considering the above findings, impaired activation of immune cells in klotho KO mice may be ascribed to telomere shortening in haematopoietic bone marrow precursor cells and causes incomplete bacteria clearance in the lungs, and then led to death with sepsis-related multiple organ dysfunction.

IL-1 $\beta$  is a potent proinflammatory cytokine. Its overexpression can lead to tissue damage (56). For instance, decreased IL-1 $\beta$  in inflammatory responses leads to a better outcome in acute lung infections with *Pseudomonas aeruginosa* (57). Pulmonary inflammation in mice after infection with *A. baumannii* has been associated with IL-1 $\beta$  production and the severity of lung pathology (58). In addition, CC and CXC chemokines attract neutrophils and mononuclear cells to sites of inflammation (59–62). Mediators of inflammation, including IL-1 and TNF- $\alpha$ , or bacterial products such as LPS elicit the production of those chemokines (59). Presently, the induction level of IL-1 $\beta$ , CXCL1 (KC), CXCL2 (MIP-2), and CCL2 (MCP-1) mRNA in the lungs after infection was significantly higher in klotho KO mice than in klotho WT mice. This result suggests that klotho KO mice infected with *A. baumannii* showed severe lung inflammation compared with klotho WT mice, which sometimes led to death. A previous study reported that TLR2 KO mice displayed increased lung concentrations of MCP-1 and MIP-2 after infection with *A. baumannii* (24). The authors discussed the possibility that TLR2 mediates anti-inflammatory pathways that downregulate MCP-1 production. We also demonstrated that the expression level of TLR2 mRNA was lower in the lungs of uninfected klotho KO mice than in those of uninfected klotho WT mice. However, the expression level in the lungs of klotho KO mice was upregulated at 1 day post infection of *A. baumannii*. These results suggest the possibility that initial infection of *A. baumannii* in aged hosts causes severe inflammation. Moreover, the expression level of TLR9 mRNA in the lungs of klotho KO mice was lower than that of klotho WT mice. TLR9 KO mice challenged intranasally with *A. baumannii* reportedly displayed a significantly increased bacterial burden and neutrophil recruitment in the lungs, with reduced levels of proinflammatory cytokines, such as TNF- $\alpha$ , IFN- $\gamma$ , and MCP-1, in their sera during systemic *A. baumannii* infection (26). These phenotypes are similar to those of klotho KO mice infected with *A. baumannii* in the present study. Additionally, we demonstrated that the expression of proinflammatory cytokines was strongly and significantly correlated with that of TLR9 mRNA in the lungs of klotho KO mice. The above findings suggest that TLR9-mediated immune responses may play an important role in the lung inflammation of elderly people with *A. baumannii* infection.

In this study, lung bacterial burden remained in klotho KO mice at 7 days post-infection in spite of the infection established by the intravenous inoculation of *A. baumannii*. Previous studies have demonstrated that the virulence factors of *A. baumannii*,

such as OmpA and Bap, mediate bacterial adherence to bronchial and lung epithelial cells (11, 12). Additionally, we have reported that the levels of *ompA* expression in MDRAB clinical isolates correlate with the adherence of the bacteria to lung epithelial cells (13). We also reported a novel bacterial transport mechanism, where *A. baumannii* exploits human neutrophils by adhering to the cells (63). These results support the novel idea concerning *A. baumannii* infection in elderly people, in which the bacteria infect aged hosts *via* the peripheral blood, respiratory system, and other routes, followed by the movement with neutrophils to the lung tissue and subsequent colonization of the lungs. Additionally, the production of proinflammatory cytokines in the sera after infection with *A. baumannii* were prolonged in *klotho* KO mice compared with *klotho* WT mice. However, although *klotho* WT mice showed decreased weight after infection with *A. baumannii*, *klotho* KO mice showed only a slight weight change after the infection. Additionally, most of *klotho* KO mice infected with *A. baumannii* did not display any clinical symptoms. These results suggest that weak inflammation lasting for prolonged periods are observed in aged hosts.

A previous study reported that *A. baumannii* infection inhibits airway eosinophilia and lung pathology in a mouse model of allergic asthma (64). Our results indicate that eosinophils are decreased in the lungs of *klotho* WT mice after infection with *A. baumannii*, implying the inhibition of airway eosinophilia in young hosts with allergic asthma. However, the significant increase in eosinophils in the lungs of *klotho* KO mice after infection with *A. baumannii* indicates further induction of airway eosinophilia in elderly people with allergic asthma. Thus, the pulmonary inflammation in *A. baumannii* infection in elderly people may be a different clinical condition compared with that in young people.

In summary, our results suggest that *A. baumannii* infection in aged hosts involves accumulation of the bacteria in the lungs, which elicits pulmonary inflammation during early infection and can eventually lead to death. The results also suggest that pulmonary inflammation correlates with altered TLR9 expression in *klotho* KO mice. This study provides insights into the pathogenicity of *A. baumannii* infection, particularly in elderly people. Further studies are required to understand the mechanism of *A. baumannii* infection in elderly people.

## DATA AVAILABILITY STATEMENT

The original contributions presented in the study are included in the article/supplementary material. Further inquiries can be directed to the corresponding author.

## ETHICS STATEMENT

The animal study was reviewed and approved by Institutional Animal Care and Use Committee of Teikyo University Animal Ethics Committee (no. 1705236A1b).

## AUTHOR CONTRIBUTIONS

Conceptualization, data curation, formal analysis, investigation, visualization, writing—original draft preparation: YS. Funding acquisition, project administration, supervision: YS, YO. Methodology: YS, ST-N, TU. Resources, validation, writing—review and editing: YS, ST-N, TU, YO. All authors contributed to the article and approved the submitted version.

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## SUPPLEMENTARY MATERIAL

The Supplementary Material for this article can be found online at: <https://www.frontiersin.org/articles/10.3389/fimmu.2020.601614/full#supplementary-material>

**SUPPLEMENTARY FIGURE 1** | Phenotypic analysis of innate immune cells in the lungs of *Acinetobacter baumannii*-infected and uninfected *klotho* wildtype (WT) mice. Innate immune cells in the lungs of *A. baumannii*-infected and uninfected *klotho* WT mice were analyzed for their expression of the following cell-surface markers: CD45, CD11b, CD11c, Ly-6G, and F4/80. The results represent 4-color flow cytometric analysis of the innate immune cell population in lung homogenates from (A) an uninfected *klotho* WT mouse and (B) an *A. baumannii*-infected mouse at 1 day post-infection. Np, Neutrophils (CD45<sup>+</sup>CD11b<sup>+</sup>Ly-6G<sup>+</sup>); AM, Alveolar macrophages (CD45<sup>+</sup>CD11b<sup>low</sup>CD11c<sup>high</sup>Ly-6G<sup>-</sup>F4/80<sup>+</sup>); IM, Interstitial macrophages (CD45<sup>+</sup>CD11b<sup>high</sup>CD11c<sup>low</sup>Ly-6G<sup>-</sup>F4/80<sup>+</sup>FSC<sup>mid/high</sup>SSC<sup>mid</sup>); Mc, Monocytes (CD45<sup>+</sup>CD11b<sup>+</sup>CD11c<sup>-</sup>Ly-6G<sup>-</sup>F4/80<sup>low</sup>); Mc/DC, Monocytes/dendritic cells (CD45<sup>+</sup>CD11b<sup>high</sup>CD11c<sup>+</sup>Ly-6G<sup>-</sup>F4/80<sup>low</sup>); Ep, Eosinophils (CD45<sup>+</sup>CD11b<sup>high</sup>CD11c<sup>low</sup>Ly-6G<sup>-</sup>F4/80<sup>+</sup>FSC<sup>mid</sup>SSC<sup>high</sup>); CD11b<sup>low</sup> cDC, CD11b<sup>low</sup> conventional dendritic cells (CD45<sup>+</sup>CD11b<sup>low</sup>CD11c<sup>+</sup>Ly-6G<sup>-</sup>F4/80<sup>low</sup>).

**SUPPLEMENTARY FIGURE 2** | Phenotypic analysis of innate immune cells in the lungs of *Acinetobacter baumannii*-infected and uninfected *klotho* knockout (KO) mice. Innate immune cells in the lungs of *A. baumannii*-infected and uninfected *klotho* KO mice were analyzed for their expression of the following cell-surface markers: CD45, CD11b, CD11c, Ly-6G, and F4/80. A representative result of 4-color flow cytometric analysis of the innate immune cell population in lung homogenates from (A) an uninfected *klotho* KO mouse and (B) an *A. baumannii*-infected mouse at 1 day post-infection. Np, Neutrophils (CD45<sup>+</sup>CD11b<sup>+</sup>Ly-6G<sup>+</sup>); AM, Alveolar macrophages (CD45<sup>+</sup>CD11b<sup>low</sup>CD11c<sup>high</sup>Ly-6G<sup>-</sup>F4/80<sup>+</sup>); IM, Interstitial macrophages (CD45<sup>+</sup>CD11b<sup>high</sup>CD11c<sup>low</sup>Ly-6G<sup>-</sup>F4/80<sup>+</sup>FSC<sup>high</sup>SSC<sup>mid</sup>); Mc, Monocytes (CD45<sup>+</sup>CD11b<sup>+</sup>CD11c<sup>-</sup>Ly-6G<sup>-</sup>F4/80<sup>low</sup>); Mc/DC, Monocytes/dendritic cells (CD45<sup>+</sup>CD11b<sup>high</sup>CD11c<sup>+</sup>Ly-6G<sup>-</sup>F4/80<sup>low</sup>); Ep, Eosinophils (CD45<sup>+</sup>CD11b<sup>high</sup>CD11c<sup>low</sup>Ly-6G<sup>-</sup>F4/80<sup>+</sup>FSC<sup>mid</sup>SSC<sup>high</sup>); CD11b<sup>lo</sup> cDC, CD11b<sup>lo</sup> conventional dendritic cells (CD45<sup>+</sup>CD11b<sup>low</sup>CD11c<sup>+</sup>Ly-6G<sup>-</sup>F4/80<sup>low</sup>).

**SUPPLEMENTARY FIGURE 3** | Phenotypic analysis of innate immune cells in the blood of *Acinetobacter baumannii*-infected and uninfected klotho wildtype (WT) mice. Innate immune cells in the blood of *A. baumannii*-infected and uninfected klotho WT mice were analyzed for their expression of the following cell-surface markers: CD11b, CD11c, Ly-6C, Ly-6G, and F4/80. A representative result of 4-color flow cytometric analysis of the innate immune cell population in blood cells from (A) an uninfected klotho WT mouse and (B) an *A. baumannii*-infected mouse at 1 day post-infection. Np, Neutrophils (CD11b<sup>+</sup>Ly-6G<sup>+</sup>); MΦ, Macrophages (CD11b<sup>high/+</sup>CD11c<sup>-</sup>Ly-6C<sup>+/-</sup>Ly-6G<sup>-</sup>F4/80<sup>+</sup> FSC<sup>high/mid</sup>SSC<sup>mid/lo</sup>); Mc, Monocytes (CD11b<sup>high/+</sup>CD11c<sup>lo/-</sup>Ly-6C<sup>high</sup>Ly-6G<sup>-</sup>F4/80<sup>+</sup>FSC<sup>high/mid</sup>SSC<sup>mid/lo</sup>); Ep, Eosinophils (CD11b<sup>high/+</sup>CD11c<sup>-</sup>Ly-6G<sup>-</sup>FSC<sup>mid</sup>SSC<sup>high</sup>); DC, Dendritic cells (CD11b<sup>low/-</sup>CD11c<sup>high</sup>Ly-6G<sup>-</sup>).

**SUPPLEMENTARY FIGURE 4** | Phenotypic analysis of innate immune cells in the blood of *Acinetobacter baumannii*-infected and uninfected klotho knockout (KO) mice. Innate immune cells in the blood of *A. baumannii*-infected and uninfected klotho KO mice were analyzed for their expression of the following cell-surface markers: CD11b, CD11c, Ly-6C, Ly-6G, and F4/80. A representative result of 4-color flow cytometric analysis of the innate immune cell population in blood cells from (A) an uninfected klotho WT mouse and (B) an *A. baumannii*-infected mouse at 1 day post-infection. Np, Neutrophils (CD11b<sup>+</sup>Ly-6G<sup>+</sup>); MΦ, Macrophages (CD11b<sup>high/+</sup>CD11c<sup>-</sup>Ly-6C<sup>+/-</sup>Ly-6G<sup>-</sup>F4/80<sup>+</sup> FSC<sup>high/mid</sup>SSC<sup>mid/lo</sup>); Mc, Monocytes (CD11b<sup>high/+</sup>CD11c<sup>lo/-</sup>Ly-6C<sup>high</sup>Ly-6G<sup>-</sup>F4/80<sup>+</sup>FSC<sup>high/mid</sup>SSC<sup>mid/lo</sup>); Ep, Eosinophils (CD11b<sup>high/+</sup>CD11c<sup>-</sup>Ly-6G<sup>-</sup>FSC<sup>mid</sup>SSC<sup>high</sup>); DC, Dendritic cells (CD11b<sup>low/-</sup>CD11c<sup>high</sup>Ly-6G<sup>-</sup>).

**SUPPLEMENTARY FIGURE 5** | Phenotypic analysis of innate immune cells in the spleen of *Acinetobacter baumannii*-infected and uninfected klotho wildtype (WT) mice. Innate immune cells in the spleen of *A. baumannii*-infected and uninfected klotho WT mice were analyzed for their expression of the following cell-surface markers: CD11b, CD11c, Ly-6C, Ly-6G, and F4/80. A representative result of 4-color flow cytometric analysis of the innate immune cell population in splenocytes

from (A) an uninfected klotho WT mouse and (B) an *A. baumannii*-infected mouse at 1 day post-infection. Np, Neutrophils (CD11b<sup>+</sup>Ly-6G<sup>+</sup>); MΦ, Macrophages (CD11b<sup>high/+</sup>CD11c<sup>-</sup>Ly-6C<sup>+/-</sup>Ly-6G<sup>-</sup>F4/80<sup>+</sup> FSC<sup>high/mid</sup>SSC<sup>mid/lo</sup>); Mc, Monocytes (CD11b<sup>high/+</sup>CD11c<sup>lo/-</sup>Ly-6C<sup>high</sup>Ly-6G<sup>-</sup>F4/80<sup>+</sup>FSC<sup>high/mid</sup>SSC<sup>mid/lo</sup>); Ep, Eosinophils (CD11b<sup>high/+</sup>CD11c<sup>-</sup>Ly-6G<sup>-</sup>FSC<sup>mid</sup>SSC<sup>high</sup>); DC, Dendritic cells (CD11b<sup>low/-</sup>CD11c<sup>high</sup>Ly-6G<sup>-</sup>).

**SUPPLEMENTARY FIGURE 6** | Phenotypic analysis of innate immune cells in the spleen of *Acinetobacter baumannii*-infected and uninfected klotho knockout (KO) mice. Innate immune cells in the spleen of *A. baumannii*-infected and uninfected klotho KO mice were analyzed for their expression of the following cell-surface markers: CD11b, CD11c, Ly-6C, Ly-6G, and F4/80. A representative result of 4-color flow cytometric analysis of the innate immune cell population in splenocytes from (A) an uninfected klotho WT mouse and (B) an *A. baumannii*-infected mouse at 1 day post-infection. Np, Neutrophils (CD11b<sup>+</sup>Ly-6G<sup>+</sup>); MΦ, Macrophages (CD11b<sup>high/+</sup>CD11c<sup>-</sup>Ly-6C<sup>+/-</sup>Ly-6G<sup>-</sup>F4/80<sup>+</sup> FSC<sup>high/mid</sup>SSC<sup>mid/lo</sup>); Mc, Monocytes (CD11b<sup>high/+</sup>CD11c<sup>lo/-</sup>Ly-6C<sup>high</sup>Ly-6G<sup>-</sup>F4/80<sup>+</sup>FSC<sup>high/mid</sup>SSC<sup>mid/lo</sup>); Ep, Eosinophils (CD11b<sup>high/+</sup>CD11c<sup>-</sup>Ly-6G<sup>-</sup>FSC<sup>mid</sup>SSC<sup>high</sup>); DC, Dendritic cells (CD11b<sup>low/-</sup>CD11c<sup>high</sup>Ly-6G<sup>-</sup>).

**SUPPLEMENTARY FIGURE 7** | Number of innate immune cells in the spleen of klotho wildtype (WT) and knockout (KO) mice before and after infection with *Acinetobacter baumannii*. Summarized results show the number of (A) neutrophils, (B) eosinophils, (C) monocytes, (D) macrophages, and (E) dendritic cells (DC) in the spleen of *A. baumannii*-infected klotho WT and KO mice. Line graphs were compiled from 4 independent experiments ( $n = 16$  for WT and  $n = 16$  for KO mice), and represent the mean  $\pm$  SEM. Asterisks indicate statistically significant differences as follows:  $^{**}P < 0.01$ ,  $^{*}P < 0.05$ , klotho WT mice before infection vs. klotho WT mice after infection, Student's  $t$ -test;  $^{++}P < 0.01$ ,  $^{+}P < 0.05$ , klotho KO mice before infection vs. klotho KO mice after infection, Student's  $t$ -test; and  $^{++}P < 0.01$ ,  $^{+}P < 0.05$ , klotho WT mice vs. klotho KO mice, Student's  $t$ -test).

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**Conflict of Interest:** The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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