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Innate immunity in peripheral tissues is differentially impaired under normal and endotoxic conditions in aging

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Chronic low-grade inflammation is a hallmark of aging, aka "inflammaging", which is linked to a wide range of age-associated diseases. Immune dysfunction increases disease susceptibility, and increases morbidity and mortality of aging. Innate immune cells, including monocytes, macrophages and neutrophils, are the first responders of host defense and the key mediators of various metabolic and inflammatory insults. Currently, the understanding of innate immune programming in aging is largely fragmented. Here we investigated the phenotypic and functional properties of innate immune cells in various peripheral tissues of young and aged mice under normal and endotoxic conditions. Under the steady state, aged mice showed elevated proinflammatory monocytes/macrophages in peripheral blood, adipose tissue, liver, and colon. Under lipopolysaccharide (LPS)-induced inflammatory state, the innate immune cells of aged mice showed a different response to LPS stimulus than that of young mice. LPS-induced immune responses displayed differential profiles in different tissues and cell types. In the peripheral blood, when responding to LPS, the aged mice showed higher neutrophils, but lower pro-inflammatory monocytes than that in young mice. In the peritoneal fluid, while young mice exhibited significantly elevated pro-inflammatory neutrophils and macrophages in response to LPS, aged mice exhibited decreased proinflammatory neutrophils and variable cytokine responses in macrophages. In the adipose tissue, LPS induced less infiltrated neutrophils but more infiltrated macrophages in old mice than young mice. In the liver, aged mice showed a more robust increase of pro-inflammatory macrophages compared to that in young mice under LPS stimulation. In colon, macrophages showed relatively mild response to LPS in both young and old mice. We have further tested bonemarrow derived macrophages (BMDM) from young and aged mice, we found that BMDM from aged mice have impaired polarization, displaying higher expression of pro-inflammatory markers than those from young mice. These data collectively suggest that innate immunity in peripheral tissues is impaired in aging, and the dysregulation of immunity is tissue- and cell-dependent. Our findings in the rodent model underscore the complexity of aging immunity. Further investigation is needed to determine whether the immune profile observed in aged mice is applicable in age-associated diseases in humans.

KEYWORDS

aging, inflammaging, immunometabolism, innate immunity, macrophages, monocytes, neutrophils

1 Introduction

The aging population is on the rise worldwide. Increasing health span is in the forefront of aging research and is a challenging task for the global health care system (1). Aging is a major risk factor for chronic illnesses (2). Inflammaging is the condition of age-associated chronic low-grade inflammation (3), which is linked to the development and progression of various age-related chronic diseases such as obesity, type 2 diabetes, cardiovascular disease, and inflammatory bowel disease (4). There is clear evidence that elevated inflammatory cytokines, such as TNF α , IL- β , and IL-6, are correlated with high mortality rates in old individuals (5–8).

The immune system is critical for survival, as it defends the host from various inflammatory stimuli such as pathogens and mutated endogenous substances. Immune dysfunction in aging increases susceptibility to disease; it diminishes vaccination responses and contributes to high morbidity and mortality in the elderly (9). To date, the T and B cells in aging have been well-documented (10, 11), whereas the innate immunity in aging remains understudied. Given that innate immunity is a first-line defense and plays a critical role in initiation and resolution of inflammation (12), it is of paramount importance to understand innate immunity in aging, which will have important implications in age-associated diseases and healthspan.

Innate immune cells are highly plastic, showing dynamic response to inflammatory stimuli in order to rapidly adapt to the microenvironment and perform specific functions; this is referred to polarization. The polarization of innate immune cells leads to different functional outcomes, which is traditionally described as a dichotomous concept; e.g., the M1-M2 paradigm of macrophage polarization (13–16). This concept is widely applied *in vitro*, but evidences now suggest that tissue macrophages are heterogeneous and present across a wide spectrum that is not restricted to the dichotomous M1-M2 characterization (17–19).

It has been shown that aging impacts the programming of innate immune cells and contributes to inflammaging and ageassociated diseases (20–22). However, it is largely unknown if ageassociated innate immune programming is tissue- and/or cellspecific. Here we uncovered the phenotypic and functional properties of innate immune cells, particularly neutrophils and macrophages, in a wide range of peripheral tissues in young and aged mice under both steady state and LPS-induced inflammatory state. We assessed properties of neutrophils and macrophages, as well as inflammatory cytokines produced by these immune cells, to assess functional outcomes using flow cytometry. The observed tissue-specific differential changes in these immune cells underscore the dysregulated immune responses in aging, thus increasing the susceptibility to the development of age-associated chronic diseases.

2 Materials and methods

2.1 Animal model

Animal experiments were conducted with C57BL/6J background adult male young (4-to-7-month-old) and aged (17to-27-month-old) mice. All experimental animal protocols were approved by the Texas A&M University Institutional Animal Care and Use Committee (protocol code: IACUC 2022-0180, approval date: October 4, 2022). The mice were bred and housed under 12 h light-dark cycles, at 23 \pm 1°C, with *ad libitum* access to water and semi-purified diet (Harlan-Teklad, Madison, WI, USA) which provided 58% kcal from carbohydrates, 24% kcal from protein, and 18% kcal from fat. To determine the inflammatory response, young (4-to-7-month-old) and aged (17-to-27-month-old) male mice received a single *i.p.* injection of lipopolysaccharide (LPS) 0.75 mg/kg and tissues were collected 6 hours after LPS exposure. At termination, the mice were euthanized using isoflurane anesthesia. The data from with and without LPS treatment groups were collected together within the same age group.

2.2 Tissue collection, digestion, and single cell preparation

2.2.1 Peripheral blood

Mice were anesthetized using isoflurane and then whole blood was collected from retro-orbital into EDTA-coated MiniCollect tubes (Greiner, USA). For flow cytometry, 50 μ L of whole blood was incubated with BD Fc blockTM (CD16/32 clone, BD Bioscience, 553141) for 15 min on ice followed by surface marker staining with

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CD45-BV510 (Biolegend, 103138), CD11b-PE (Biolegend, 101208), Ly6C-eFluor450 (Thermofisher, 48-5932-82), Ly6G-BV785 (Biolegend, 127645), CD115-APC (Biolegend, 135509), CCR2-FITC (Biolegend, 150607) for 30 min. Red blood cells were then lysed with FACS Lysing Solution (BD Bioscience, 349202) in 1:20 ratio for 20 min at room temperature followed by 2% PFA for 30 min on ice. AccuCheck counting beads (Thermofisher, PCB100) were added to samples before data acquisition to enable the computation of absolute cell counts in PBMC, as per the manufacturer's instructions. Fluorescence was detected using BD LSR FortessaTM X-20 (BD Bioscience, San Jose, CA, USA) and/or Cytek® Aurora (Cytek®, Fremont, CA, USA). The gating strategy is noted in Supplementary Figure 1A. Briefly, neutrophils were gated as CD45⁺CD11b⁺Ly6G⁺ and monocytes were gated as CD45⁺CD11b⁺CD115⁺. Then, monocyte subpopulations were further distinguished by differential expression of Ly6C. We confirmed that the Ly6C^{hi} subset expressed high levels of CCR2, whereas Ly6C^{inter}and Ly6C^{low} subsets had low expression of CCR2.

2.2.2 Peritoneal fluid cells

Peritoneal fluid was collected as described previously (23). Briefly, 3 ml of ice-cold phosphate-buffered saline (PBS) containing 2% fetal bovine serum (FBS) was injected into the peritoneum. After gentle shaking for 1 min, peritoneal fluid was collected with an 18G syringe needle. The samples were then centrifuged at 500 g for 5 mins at 4°C. To lyse red blood cells, ACK lysing buffer (Fisher scientific, 50-101-9080) was added to the cell pellets and incubated for 5 min. Then, cells were collected by centrifugation at 500 g for 10 min at 4°C followed by washing twice with PBS. Golgi plug (BD Bioscience) was added to the cells for 4 hours followed by antibody staining as described in the Flow cytometry staining and cell analysis section below. The surface staining markers were, CD45-eFluor450 (eBioscience, 50-112-9409), Ly6G-BV785 (Biolegend, 127645), CD11b-APC-Cy7 (Biolegend, 101226), F4/80-PE-Cy7 (eBioscience, 25-4801-82), CD38-PE (BD Bioscience, 553764), and CD206-Alexa Fluor 488 (Biolegend, 141710). Additionally, to assess functional outcome more directly, intracellular antibodies were used to test cytokine expression in specific cells. TNFα-PE-Dazzle594 (Biolegend, 506345), IL-1β-PE (Thermofisher, 12-7114-80), iNOS-Alexa Fluor 488 (Thermofisher, 53-5920-80) and Arg1-Alexa Fluor700 (Thermofisher, 56-3697-82) were used to stain cells for 30 min on ice. Cells were then washed with the perm wash buffer. Data were acquired using a BD LSR FortessaTM X-20 (BD Bioscience, San Jose, CA, USA) and/or Cytek® Aurora (Cytek®, Fremont, CA, USA). The flow cytometry gating strategy is detailed in Supplementary Figure 2. Briefly, peritoneal neutrophils were defined as CD45⁺CD11b⁺Ly6G⁺. The intracellular markers (TNF α , IL-1 β , and iNOS) were evaluated from neutrophil population (CD45⁺CD11b⁺Ly6G⁺). Peritoneal macrophages were defined as CD45⁺CD11b⁺Ly6G⁻F4/80⁺. The surface markers (CD38 and CD206) and intracellular markers (TNFa, IL-1B, iNOS, and Arg1) were evaluated from macrophage population (CD45⁺CD11b⁺Ly6G⁻F4/80⁺).

2.2.3 Stromal vascular fraction from epididymal adipose tissue

Stromal vascular fraction was collected from epididymal adipose tissue as described previously (24). In brief, approximately 1 g of epididymal fat was minced using scissors and incubated in digestion buffer (1 mg/ml collagenase type I in RPMI medium) for 30 min at 37°C in a shaking water bath. The samples were filtered with 70 µm cell strainers to remove undigested tissue. After centrifuging at 1,700 g for 3 min, the cell pellets were collected and incubated with 4 ml of ACK lysing buffer (Fisher scientific, 50-101-9080) on ice for 10 min. The samples were then centrifuged at 1,700 g for 3 mins and the pellet of stromal vascular fraction (SVF) was re-suspended in 1 ml of PBS with 2% FBS. The antibody staining was described in the Flow cytometry staining and cell analysis section below. Surface markers were stained using CD45-BV510 (Biolegend, 557235), Ly6G-AlexaFluor 700 (Biolegend, 127622), CD11b-APC-cy7 (Biolegend, 101226), F4/80-PE-cv7 (eBioscience, 25-4801-82), and CX3CR1-BV421 (Biolegend, 149023) for 30 min on ice. The cells were washed with FACS buffer (1xPBS and 2% FBS) and then stained with 7AAD (BD Bioscience) before data collection. The data was acquired using MoFlo Astrios EQ (Beckman Coulter Life Sciences, Indianapolis, IN, USA) and/or Cytek® Aurora (Cytek®, Fremont, CA, USA). The flow gating strategy and representative plots are shown in Supplementary Figure 3. Neutrophils were gated as CD45⁺CD11b⁺Ly6G⁺. Infiltrating macrophages were defined as CD45⁺CD11b⁺Ly6G⁻F4/80^{hi} and resident macrophages were defined as CD45⁺CD11b⁺Ly6G⁻F4/80^{low}. The surface markers (CD38 and CD206) were evaluated from the macrophage population (CD45⁺CD11b⁺Ly6G⁻F4/80⁺).

2.2.4 Liver

The gallbladder was carefully removed first, and then the liver was dissected. A portion of liver tissue was digested into single cell suspension using the Liver Dissociation Kit for mouse (Miltenyi Biotec, Auburn, CA, USA, 130-105-807) following the manufacturer's instructions. Briefly, liver was rinsed with DMEM then transferred into MACS C tubes containing a dissociation enzyme mix, which was pre-heated for 30 min at 37°C. The samples were then digested using GentleMACSTM Octo Dissociator with heaters (Miltenyi Biotec, Auburn, CA, USA). The cell suspensions were filtered with 100 μ m strainers, then centrifuged at 300 g for 10 min. Resuspended cells were stained with antibodies as described in the Flow cytometry staining and cell analysis section below. The antibodies used for cell surface staining were CD45-PerCP (BD Bioscience, 557235), Ly6G-BV785 (Biolegend, 127645), F4/80-PE-Cy7 (eBioscience, 25-4801-82), CD11b-APC-Cy7(Biolegend, 101226), CD38-PE (BD Bioscience, 553764), and CD206-BV650 (Biolegend, 141723). The intracellular staining antibodies were TNFa-Alexa Fluor 700 (BD Bioscience, 558000), IL-1β-PE (Thermofisher, 12-7114-80), and iNOS-Alexa Fluor 488 (Thermofisher, 53-5920-80). The data was acquired using Cytek® Aurora (Cytek®, Fremont, CA, USA). The flow gating strategy and representative plots are shown in Supplementary

Figure 4. Neutrophils were selected as CD45⁺CD11b⁺Ly6G⁺. The intracellular markers (TNF α , IL-1 β , and iNOS) were evaluated from neutrophil population (CD45⁺CD11b⁺Ly6G⁺). Infiltrating macrophages were defined as CD45⁺CD11b^{hi}Ly6G⁻F4/80^{low} and resident macrophages were defined as CD45⁺CD11b^{hi}Ly6G⁻F4/80^{hi}. The surface markers (CD38 and CD206) and intracellular markers (TNF α , IL-1 β , and iNOS) were evaluated from macrophage population (CD45⁺CD11b⁺Ly6G⁻F4/80⁺).

2.2.5 Colon

The colon was first washed with ice-cold phosphate-buffered saline, then cut longitudinally in the middle. Half colon was digested to obtain a single-cell suspension using the Lamina Propria Dissociation Kit for mouse (Miltenyi Biotec, Auburn, CA, USA, 130-097-410) according to the manufacturer's instructions. In brief, the colon samples were washed with 1x HBSS without Ca2+ and Mg2+ containing 10 mM HEPES, 5mM EDTA and 5% FBS, and then the samples were digested by the enzyme mix from the Lamina Propria Dissociation kit (Miltenyi Biotec, Auburn, CA, USA) using GentleMACSTM Octo Dissociator with heaters (Miltenyi Biotec, Auburn, CA, USA). The cell suspensions were filtered with 100 µm strainers. The single-cell suspensions were then incubated with Golgi plug (BD Bioscience) for 4 hours then stained with fluorescenceconjugated antibodies of targeted markers for flow cytometry as described in the Flow cytometry staining and cell analysis section below. Surface marker antibodies used were CD45-FITC (eBioscience, 11-0451-82), CD11b-APC-cy7 (Biolegend, 101226), Ly6G-BV785 (Biolegend, 127645), Ly6C-eFluor450 (eBioscience,48-5932-82), MHCII-Alexa Fluor700 (Biolegend, 107622), CD64-Alexa Fluor647 (Biolegend, 139322), and CD206-PerCP-cy5.5 (Biolegend, 141716). Intracellular markers used were TNFq-PE-Dazzle594 (Biolegend, 506345) and IL-1β-PE (Thermofisher, 12-7114-80). The data were acquired using MoFlo Astrios EQ (Beckman Coulter Life Sciences, Indianapolis, IN, USA) and/ or Cytek® Aurora (Cytek®, Fremont, CA, USA). The flow gating strategy and representative plots are shown in Supplementary Figure 5. Briefly, neutrophils were identified as CD45⁺CD11b⁺Ly6G⁺. The intracellular markers (TNF α and IL-1 β) were evaluated from neutrophil population (CD45⁺CD11b⁺Ly6G⁺). The monocyte/macrophage population was selected as CD45⁺CD11b⁺Ly6G⁻CD64⁺. The surface markers (Ly6C and CD206) and intracellular markers (TNF α and IL-1 β) were evaluated from monocytes/macrophage population (CD45⁺CD11b⁺Ly6G⁻CD64⁺).

2.2.6 Bone marrow-derived macrophage differentiation and polarization

Bone marrow cells were collected from mouse femurs and tibias as previously described (25). 4×10^6 cells were seeded in a 10 cm plate with differentiation medium (RPMI 1640 medium supplemented with 10 ng/mL macrophage colony-stimulating factor (M-CSF) (Peprotech, 315-02-250UG), 10% fetal bovine serum, 100U/ml penicillin, and 100 µg/ml streptomycin). The cells were incubated in a humidified incubator at 37°C with 5% CO₂ for 7 days. On day 7, 2×10^5 cells/well were seeded to 96 well plates and treated with LPS 100 ng/mL for either 4 or 24 hours, and then incubated with Golgi PlugTM (BD Bioscience) for 4 hours to enhance the detectability of cytokine-producing cells in subsequent immunofluorescent staining. The cells were subsequently stained as described in the Flow cytometry staining and cell analysis section below. The surface staining antibodies were CD11b-eFluor450 (eBioscience, 48-0112-82), F4/80-PE-Cy7 (eBioscience, 25-4801-82), and CD38-PE (BD Bioscience, 553764) for 30 min on ice. The intracellular staining antibodies used were TNF α -Alexa Fluor700 (BD Bioscience, 558000), IL-1 β -PE (Thermofisher, 12-7114-80), and iNOS-Alexa Fluor 488 (Thermofisher, 53-5920-80). The data were acquired by BD LSR FortessaTM X-20 (BD Bioscience, San Jose, CA, USA). The flow gating strategy and representative plots are shown in Supplementary Figure 6. BMDM were defined as CD11b⁺F4/80⁺. The intracellular markers (TNF α , IL-1 β and iNOS) were evaluated from the macrophage population (CD11b⁺F4/80⁺).

2.3 Flow cytometry staining and cell analysis

The prepared single-cell suspensions from various tissues were stained with fluorescence-conjugated antibodies. Dead cells were stained using LIVE/DEADTM Fixable Aqua Dead Cell Stain Kit (Thermofisher, USA) for 30 mins, followed by BD Fc blockTM (CD16/32 clone, BD Bioscience, 553141) for 10 mins. The cell surface staining antibodies were resuspended in FACS buffer (1xPBS and 2% FBS) and incubated for 30 mins at 4°C followed by 2% fixation buffer in PBS (Thermofisher, J19943.K2) for 20 mins at 4°C. The cells were permeabilized by BD Perm/Wash Buffer (BD Bioscience, 554723) for 20 mins at 4°C followed by incubation with intracellular staining antibodies which were resuspended in Perm/Wash Buffer (BD, USA). The cells were then washed twice with the Perm/Wash Buffer prior to analysis.

The fluorescence intensity of the cells was detected by flow cytometers, MoFlo Astrios EQ (Beckman Coulter Life Sciences, Indianapolis, IN, USA), BD LSR FortessaTM X-20 (BD Bioscience, San Jose, CA, USA) and/or Cytek® Aurora (Cytek®, Fremont, CA, USA). Unstained cells and fluorescence minus one (FMO) controls were used for compensation in the gating strategies. Data were analyzed using FlowJo software v10 (Tree Star Inc., Ashland, OR, USA).

2.4 Statistics

Statistical analysis was performed using GraphPad Prism 8.0.1 (GraphPad Software, La Jolla, CA, USA). Unpaired Student's t-test was used. The data are presented as mean \pm SEM. The statistical significance was considered as * p < 0.05; ** p < 0.01; *** p < 0.001.

3 Results

3.1 Innate immune cells reprogram in tissue-specific manner in aging

To study immune reprogramming in specific tissue microenvironments in aging, we studied innate immune cells in peripheral blood, peritoneal fluid, epididymal fat, liver, and colon in young and aged mice.

3.1.1 Circulating monocytes are increased in peripheral blood of aged mice

Monocytes in circulation migrate to sites of inflammation and then differentiate into macrophages and dendritic cells to refill the pool of tissue immune cells (26). Circulating monocytes in mice are categorized into three subsets based on the surface expression of Ly6C; Ly6C^{hi}, Ly6C^{inter} and Ly6C^{low} (27). Ly6C^{hi} cells are proinflammatory monocytes which rapidly migrate into inflamed tissue driven by the expression of CC-chemokine receptor 2 (CCR2). Ly6C^{inter} cells are pro-inflammatory monocytes which are differentiated from Ly6C^{hi}. Ly6C^{low} subsets express low levels of CCR2, patrolling the vascular endothelium and engaging in tissue repair (27). To investigate neutrophils and monocytes in the circulation, peripheral blood of young and aged mice was investigated using flow cytometry. Aging did not change total neutrophil count and percentage, but significantly increased monocyte percentage in peripheral blood (Figure 1A). As expected, aging increased all three subsets of monocytes, proinflammatory Ly6C^{hi}, Ly6C^{inter}, and patrolling Ly6C^{low} compared to peripheral blood from young mice (Figure 1A). In summary, both pro-inflammatory and patrolling monocytes were increased in aging peripheral blood.

3.1.2 Peritoneal fluid of aged mice exhibited mild trend of change in neutrophils and macrophages

The peritoneal fluid has abundant immune cells, predominantly with mature macrophages and neutrophils, and plays an important role in modulating immune response of the host (28–30). The total neutrophil percentage was not changed with aging, but the percentage of TNF α^+ neutrophils showed a trend of increase in the aged mice (Figure 1B). While IL-1 β^+ neutrophils remained unchanged, iNOS⁺ neutrophils significantly decreased in aged mice (Figure 1B). TNF α^+ macrophages showed a trend of increase, whereas macrophages with other pro-inflammatory markers of CD38, IL-1 β , and iNOS did not differ between young and aged mice (Figure 1B). The anti-inflammatory CD206⁺ macrophages also did not change with age (Figure 1B). Collectively, the aging effect was quite mild in neutrophils and macrophages of peritoneal fluid.

3.1.3 Pro-inflammatory myeloid cells are increased in epididymal fat of aged mice

We previously showed that aging increases adiposity and inflammation in adipose tissues, which further exacerbates insulin resistance in aging (31, 32). To further elucidate whether aging reprograms innate immune cells of neutrophils, monocytes, and macrophages in epididymal fat, we studied stromal vascular fraction (SVF) of epididymal fat in young and aged mice. While aging did not alter the percentage of total neutrophils and macrophages, the percentage of infiltrating macrophages showed a trend of increase in aging (Figure 2A). In addition, CD38⁺ proinflammatory macrophages were significantly elevated in aging. While the percentage of CD206⁺ anti-inflammatory macrophages also showed a trend of increase in aging, that trend was highly varied in aged mice (Figure 2A). Collectively, aging increased proinflammatory myeloid cells, mainly by increasing infiltrating macrophages and CD38⁺ pro-inflammatory macrophages in aging epididymal fat.



Characterization of neutrophils, monocytes, and macrophages in peripheral blood and peritoneal fluid of young and aged mice. Peripheral blood and peritoneal fluid from young (5-to-7-month-old, n=5-7) and aged (17-to-27-month-old, n=5-8) mice were subjected to flow cytometry for cell surface markers and cytokine expression analysis. (A) Peripheral blood neutrophils, monocytes, and monocyte subsets. The count data is measured in 50 uL blood. (B) Peritoneal fluid: Total neutrophils and macrophages (left), neutrophils (middle), and macrophages (right). Unpaired Student's t-test was used. * p<0.05, *** p<0.001 aged vs. young.



FIGURE 2

Characterization of myeloid population in epididymal fat, liver, and colon of young and aged mice. Young (5-to-7-month-old, n=5-7) and aged (17-to-27-month-old, n=5-8) mice were subjected to flow cytometry assessment of innate immune cells in epididymal fat, liver and colon. (A) Epididymal fat: Total neutrophils and macrophages (left), characteristics of macrophages (middle) and infiltrating/resident macrophages (right). (B) Liver: Total neutrophils and macrophages (far left), neutrophils (left), macrophages (right), and infiltrating/resident macrophages (far right). (C) Colonic total neutrophils and macrophages (left), and characteristics of colonic neutrophils (middle) and macrophages (right). Unpaired Student's t-test was used. * p<0.05, ** p<0.01 aged vs. young.

3.1.4 The infiltration of pro-inflammatory macrophages is enhanced in the liver of aged mice

Fatty liver diseases, including non-alcoholic fatty liver disease (NAFLD) and nonalcoholic steatohepatitis (NASH), are more prevalent in aging (33–35). Here we examined whether innate immune cells in liver are altered in aging. Aging did not change the total percentage of neutrophils and pro-inflammatory $\text{TNF}\alpha^+$, IL-1 β^+ , and iNOS⁺ neutrophils (Figure 2B). However, aging significantly elevated the total percentage of macrophages, and the pro-inflammatory CD38⁺, TNF α^+ , and IL-1 β^+ macrophages were significantly higher in the aging liver. Notably, although the infiltrating macrophages were significantly increased in aging liver, the resident macrophages (Kupffer cells) did not change (Figure 2B). Collectively, aging primarily affects macrophages, exhibiting the infiltration of pro-inflammatory macrophages in the liver.

3.1.5 Myeloid cells in colon undergo proinflammatory programming in aging

The gastrointestinal tract has the largest pool of immune cells in the body due to the greatest amount of antigen exposure from diet and microbiome (36). To investigate if innate immune populations are altered in the aging colon, we used flow cytometry to assess innate immune cells of young and aged mice. The percentage of the total colonic neutrophils was similar between young and aged mice. However, aging significantly increased IL-1 β^+ pro-inflammatory neutrophils in colon compared to young mice colon, whereas aging did not significantly alter TNF α^+ neutrophils (Figure 2C). Macrophage total percentage did not change in the aging colon (Figure 2C). However, a robust change of macrophage subpopulations was detected. Ly6C^{hi} infiltrating monocytes and macrophages were significantly increased in aging (Figure 2C). Also, IL-1 β^+ pro-inflammatory macrophages were significantly elevated in aged mice colon, whereas TNF α^+ macrophages remained similar in aging (Figure 2C). Furthermore, aging colon showed significantly decreased CD206⁺ anti-inflammatory macrophages (Figure 2C). Collectively, aging significantly enhances pro-inflammatory neutrophils and macrophages, and reduces anti-inflammatory macrophages in the colon.

3.2 Bone marrow-derived macrophages from aged mice show impaired polarization in response to proinflammatory LPS stimulation

Chronic exposure of pro-inflammatory cytokines in aging results in defective ability of BMDM to clear pathogens, suggesting aging driven functional alteration of BMDM (37, 38). To determine whether aging itself affects the inflammatory response of BMDM, BMDM from young (4-month-old) and aged (17-to-27-month-old) mice were treated with LPS 100 ng/mL for either 4 or 24 hours, and then pro-inflammatory markers were tested using flow cytometry. In BMDM from both young and aged mice, the expression of pro-inflammatory marker CD38 was expressed the most at 24 hours after LPS treatment (Figure 3). The BMDM from aged mice showed significantly increased CD38⁺ expression in 24 hours of LPS treatment. The expressions of the pro-inflammatory cytokines of TNF α and IL-1 β peaked at 4 hours after LPS treatment,

whereas iNOS expression was increased at 24 hours after LPS treatment (Figure 3). Interestingly, BMDM from aged mice had significantly elevated baseline expression of IL-1 β compared to BMDM from young mice (Figure 3). Also, BMDM from aged mice showed significantly higher expression of iNOS at baseline and at 4 hours post-LPS treatment (Figure 3). Collectively, BMDM from aged mice exhibited higher expression of pro-inflammatory markers (CD38, IL-1 β , and iNOS) compared to those in BMDM from young mice.

3.3 Aging innate immune cells show cell- and tissue-specific response to LPS-induced inflammation

Overwhelming evidences support that the elderly are more susceptible to infectious disease due to immune decline (39, 40). Although many vaccines are developed to combat life-threatening infections, the vaccine efficacy is lower in the elderly (41, 42) due to impaired immunity in the elderly. Our BMDM data above revealed that macrophages, which are key innate immune cells, have altered polarization in aging. To further test the differential response of aging innate immunity and its characteristic changes across the peripheral tissues in LPS-induced inflammation, low-dose LPS (0.75 mg/kg) was *i.p.* injected into young and aged mice, and tissues were collected after 6 hours. The dosage and the time of LPS treatment were chosen based on our preliminary titration study (data not shown). Considering that the elderly are more susceptible to infection (9), low-dose LPS injection was used in mice to induce acute inflammation to mimic the condition of the elderly when



FIGURE 3

Polarization potential of bone-marrow derived macrophages (BMDM) from young and aged mice. Bone-marrow from young (5-month-old, n=3) and aged (17-to-18-month-old, n=6) mice were differentiated to BMDM by 10 ng/mL M-CSF for 7 days and then treated with 100 ng/mL LPS for 4 or 24 hours to promote pro-inflammatory polarization. Flow cytometry analysis of macrophages: (A) % of CD38,TNF α , IL-1 β and iNOS; (B) MFI of CD38, TNF α , IL-1 β and iNOS. Unpaired Student's t-test was used. * p<0.05, ** p<0.01 aged vs. young.

exposed to infection. Below, we have analyzed innate responses of the peripheral tissues in young and aged mice in response to the low dose LPS exposure.

3.3.1 Myeloid cells in peripheral blood of aged mice exhibit a robust response to LPS

In peripheral blood, young mice showed significantly increased neutrophils upon LPS exposure (Figure 4A). Aged mice also exhibited increased neutrophils upon LPS treatment, whereas the response was more robust than in young mice (Figure 4A). Interestingly, total monocyte percentage was significantly reduced in both young and old mice in response to LPS (Figure 4B). Furthermore, the monocyte subsets of Ly6Chi, Ly6Cinter and Ly6C^{low} were all decreased by LPS in both young and aged mice (Figure 4C). Of note, old mice showed a more pronounced reduction of Ly6C^{hi}, Ly6C^{inter} and Ly6C^{low} monocytes as compared to that of young mice (Figure 4C). Collectively, peripheral blood neutrophils were increased, but total monocytes and monocyte subsets were decreased in both young and aged mice. On the other hand, aged mice showed more robust change of neutrophils and monocyte subsets in response to LPS as compared to young mice.

3.3.2 The responses of peritoneal neutrophils and macrophages to LPS are impaired in aging

In the peritoneal fluid, young mice showed significantly elevated neutrophil percentage in response to LPS. The aged mice also showed a trend of increase in neutrophil percentage by LPS treatment (Figure 5A). Interestingly, the secretion of proinflammatory cytokines in neutrophils was the opposite between young and aged mice in response to LPS (Figure 5A): In young mice, $TNF\alpha^+$ and $iNOS^+$ neutrophils were increased in peritoneal fluid; counterintuitively, aged mice exhibited decreased $TNF\alpha^+$ and iNOS⁺ neutrophils upon LPS exposure. Next, we found that the young mice displayed a trend of decrease in total macrophage percentage in response to LPS, whereas aged mice exhibited a significantly lower macrophage percentage under LPS treatment (Figure 5B). On the other hand, $TNF\alpha$ and iNOS expressing proinflammatory macrophages were significantly increased in both young and aged mice (Figure 5B). Of note, aged mice showed a less robust increase of $TNF\alpha^+$ macrophages than did young mice (Figure 5B). On the contrary, aged mice had a more significant increase of iNOS⁺ macrophages compared to young mice under LPS (Figure 5B). The response of pro-inflammatory CD38⁺ macrophages in LPS was also different in young and aged mice. Young mice showed increased CD38⁺ pro-inflammatory macrophages in response to LPS, while aged mice showed significantly decreased CD38⁺ macrophages (Figure 5B). In addition to pro-inflammatory markers, anti-inflammatory markers Arg1 and CD206 were also tested in macrophages. Under LPS exposure, young mice showed merely a trend of decreased Arg1⁺ macrophages, whereas aged mice showed significantly decreased Arg1⁺ macrophages (Figure 5C). Similarly, young mice did not show an altered CD206⁺ macrophage percentage, whereas aged mice showed a significantly reduced CD206⁺ macrophage percentage (Figure 5C). Collectively, aged mice showed an impaired response of neutrophils and macrophages to LPS. The macrophage subsets in aged mice



The response of peripheral blood immune cells to LPS in young and aged mice. Young (5-to-7-month-old, n=5-4) and aged (15-month-old, n=5-7) mice were subjected to either vehicle or LPS (*i.p.* 0.75 mg/kg) for 6 hours and the neutrophils and monocytes in peripheral blood were tested by flow cytometry. (A) Neutrophils (B) Monocytes (C) Monocyte subsets. Unpaired Student's t-test was used. * p<0.05, ** p<0.01, *** p<0.001 LPS vs. control.



showed differential responses to LPS depending on cytokine production.

3.3.3 Aging increases the responsiveness of adipose macrophages to LPS stimulation

In epididymal fat of young mice, LPS treatment did not change total myeloid cells whereas aged mice showed significantly increased myeloid cells (Figure 6A). The percentage of neutrophils was significantly increased following LPS stimulation in both groups (Figure 6B); however, the response was less pronounced in the aged mice compared to young mice (Figure 6B). On the other hand, macrophages in aged epididymal fat were more responsive to LPS than macrophages in epididymal fat from young mice. Total macrophages in young mice were not altered under LPS exposure, whereas the total macrophages in aged mice exhibited a significantly increased in epididymal fat (Figure 6C). Similarly, the macrophage subsets showed significant changes in aged mice but not in young mice (Figure 6C): While young mice did not show an altered macrophage subpopulation in response to LPS, aged mice revealed significantly increased CD38⁺ pro-inflammatory macrophages. The CD206⁺ anti-inflammatory macrophage population was not changed in either young or aged mice in response to LPS treatment (Figure 6C). Furthermore, in aged mice, there was significantly elevated macrophage infiltration and reduced resident macrophages upon LPS treatment (Figure 6D). Collectively, aging decreased neutrophil response but increased macrophage responses to inflammatory stimuli in epididymal fat.

3.3.4 Liver macrophages are more responsive to LPS in aged mice

Liver neutrophils were significantly elevated in both young and old mice in response to LPS (Figure 7A). Young mice showed significantly increased TNF α^+ and IL-1 β^+ neutrophils in response to LPS, whereas these cells in livers of aged mice did not respond to LPS (Figure 7A). Old mice presented a robust change of pro-



n=4-7) mice were subjected to either vehicle or LPS (*i.p.* 0.75 mg/kg) for 6 hours, and the innate immune cells from SVF of epididymal fat were used for flow cytometry analysis. (A) Myeloid cells (B) Neutrophils (C) Macrophages, as well as CD38⁺ or CD206⁺ macrophage subset, and (D) Infiltrating and resident macrophages. Unpaired Student's t-test was used. * p<0.05, ** p<0.01 LPS vs. control.

inflammatory macrophages compared to young mice in response to LPS (Figure 7B). CD38⁺ macrophages in young mice did not change, whereas old mice showed significantly elevated CD38⁺ macrophages upon LPS challenge (Figure 7B). In young mice, TNF α^+ macrophages did not respond to LPS; however, this cell type was significantly reduced in old mice (Figure 7B). The IL- $1\beta^+$ and iNOS⁺ macrophages were significantly increased in both young and old mice, but aged mice showed a more robust increase in those macrophages (Figure 7B). Interestingly, young mice showed increased infiltrating macrophages in response to LPS exposure, whereas aged mice did not show a change in macrophage infiltration (Figure 7C). The resident macrophages were not significantly altered in both young and aged mice following LPS treatment (Figure 7C). Collectively, liver neutrophils responded similarly between young and old mice, whereas pro-inflammatory macrophages from the livers of aged mice showed a more potent response to LPS.

3.3.5 Colonic macrophages have dampened response to LPS

Our data above support a significant aging effect of increased pro-inflammatory innate immune cells in the colon. Interestingly, colonic innate immune cells from both young and aged mice displayed relatively mild responsiveness to LPS challenge. Total neutrophil percentage was significantly higher in both young and aged mice colons, but aged mice colons showed a smaller increase of neutrophils in response to LPS as compared to young mice (Figure 8A). TNF α^+ neutrophils were steady in both young and aged mice upon LPS (Figure 8A). While in young mice, IL-1 β^+ neutrophils remained non-responsive to LPS, aged mice showed significantly escalated IL-1 β^+ neutrophils in LPS exposure (Figure 8A). Total macrophage population from young mouse colons was increased by LPS, whereas the percentage of the total macrophages in aged mouse colons did not change (Figure 8B). Also, young mice showed a trend of increased Ly6C^{hi} macrophages,



The response of liver innate immune cells to LPS in young and aged mice. Young (5-to-7-month-old, n=4-5) and aged (20-month-old, n=5) mice were subjected to either vehicle or LPS (*i*, *p*. 0.75 mg/kg) for 6 hours, and the neutrophils and macrophages were assessed by flow cytometry. (A) Total neutrophils, $TNF\alpha^+$ or IL-1 β^+ neutrophils; (B) Total macrophages, $CD38^+$, $TNF\alpha^+$, IL-1 β^+ or iNOS⁺ macrophages; (C) Infiltrating and resident macrophages. Unpaired Student's t-test was used. * p<0.05, ** p<0.01, *** p<0.001 LPS vs. control.



FIGURE 8

The response of colonic neutrophils and macrophages to LPS in young and aged mice. Young (5-to-7-month-old, n=4-5) and aged (18-to-24-month-old, n=5-7) mice were subjected to either vehicle or LPS (*i.p.* 0.75 mg/kg) for 6 hours, and the neutrophils and macrophages were assessed by flow cytometry. (A) Total neutrophils, $TNF\alpha^+$ or IL-1 β^+ neutrophils; (B) Total macrophages, Ly6C^{hi} infiltrating macrophages, $TNF\alpha^+$ or IL-1 β^+ macrophages. Unpaired Student's t-test was used. * p<0.05 LPS vs. control.

whereas aged mice showed a trend of decreased Ly6C^{hi} macrophages; however, neither reached statistical significance (Figure 8B). LPS did not alter TNF α^+ and IL-1 β^+ macrophages in both young and aged mice (Figure 8B). Overall, total neutrophils and macrophages increased in response to LPS. While pro-inflammatory neutrophils were increased significantly with LPS stimulation in aged mice, pro-inflammatory macrophages barely responded to LPS in either young or aged colons.

The collective data of the age-associated alterations of immune cells in each tissue are summarized in Figure 9. Also, responses of the immune cells to LPS in each tissue are summarized in Table 1.

4 Discussion

Emerging evidence highlights the importance of innate immunity in aging and age-associated diseases (2, 4, 12, 43). We studied innate immune programming of a wide range of peripheral tissues under both normal aging and an LPS-induced inflammatory state. Our current study reveals that aging impairs innate immunity in peripheral tissues in both normal and endotoxic conditions. Interestingly we found that the activation/response of the innate immune cells upon inflammatory stimulus is distinctively tissue/ cell-dependent in aging.

Our data from peripheral blood showed that pro-inflammatory Ly6C^{hi} monocytes were significantly increased in aging (Figure 1A). The data are in agreement with previous studies showing higher monocytes in older adults compared to young adults (44, 45), and in further support of previous findings that aging is associated with chronic systemic inflammation (2, 3, 37). Under steady state in

aging, we observed that colons from the aged mice exhibited enhanced infiltration of macrophages, and pro-inflammatory activation of neutrophils and macrophages (Figure 2C), which could contribute to systemic inflammation during aging. It is known that colon is chronically exposed to inflammatory stimuli due to microbial dysbiosis and increased gut permeability during aging (37, 46–50). The chronic exposure to inflammatory stimuli reduces phagocytic activity of macrophages in removing pathogens (37), which would perpetuate colonic inflammation. The ageassociated colonic inflammation likely further contributes to increased vulnerability to inflammatory bowel disease in aging as we and others have reported (51–53). Collectively, our data revealed pro-inflammatory activation of colonic neutrophils and macrophages, contributing to systemic inflammation in aging.

Aging is associated with obesity, a condition influenced by dietary and lifestyle choices which significantly contribute to the prevalence of insulin resistance and metabolic diseases in the elderly (54, 55). Epididymal fat and liver are major metabolic tissues extensively studied in chronic inflammatory diseases in the context of inflammation and insulin resistance (56-59). We demonstrated that macrophages in epididymal fat and liver are responsive to age-associated inflammation, exhibiting increased infiltration of macrophages (Figure 2A, B). This is in line with previous reports by us and others showing accumulation of monocytes and macrophages in adipose tissue and liver in aging (31, 60). It has been reported that accumulation of senescence celldriven mediators (SASP) in aging impairs leukocyte trafficking, thus exhibiting dysregulated adhesion/chemokine mediator expressions in aged endothelium, immune cells in the circulation, and the tissue microenvironment (61-63). We observed increased CD38⁺ pro-



Schematic diagram of the age-associated alterations of immune cells in peripheral tissues. Overall, pro-inflammatory neutrophils, monocytes and macrophages were increased in aging in peripheral blood, peritoneal fluid, epididymal fat, liver, colon, and bone marrow-derived macrophages. \uparrow or \downarrow aged vs. young.

	Neutrophils			Monocytes/Macrophages		
	Population	Young	Aged	Population	Young	Aged
Peripheral blood	Total	Ť	<u>t</u> tt	Total Ly6C ^{hi} Ly6C ^{inter} Ly6C ^{low}	+ - +++	111 1 1 1
Peritoneal fluid	Total TNFα iNOS	1 1 -	11 -	Total CD38 TNFα iNOS Arg1 CD206	- - 1 - -	↓↓ ↑ ↓ ↓
Epididymal fat	Total	<u>↑</u>	Ť	Total CD38 CD206 Infiltrating Resident		↑ ↑ ↑ ↓↓
Liver	Total TNFα IL-1β	11 11 1	++++ -	Total CD38 TNFα IL-1β iNOS Infiltrating Resident	1 - - 11 1 -	- 11 111 111 - -
Colon	Total TNFα IL-1β	1 - -	1 - 1	Total Ly6C ^{hi} TNFα IL-1β	1 - -	

TABLE 1 Summary of aging under LPS.

↑ or ↓ p<0.05, ↑↑ or ↓↓ p<0.01, ↑↑↑ or ↓↓↓ p<0.001 LPS vs. control.

inflammatory macrophages in epididymal fat and liver in aging (Figure 2A, B), which likely is due to increased SASP-driven CD38 expression in macrophages (64). Though it is not clear if macrophage senescence itself could directly increase level of CD38, the senescence of macrophages is known to induce defective insulin signaling in human adipocytes (65). While ample evidence supports increased infiltration of macrophages into adipose tissue in aging, the change of resident macrophages in adipose tissue has been debatable. In our study, resident macrophages (CD45⁺CD11b⁺F4/80^{low}) were not changed in aging (Figure 2A). As another study reported increased resident macrophages in adipose tissue (F4/80⁺CD11b⁺CD11c⁻) (64), these discrepancies suggest a heterogeneity of resident macrophages. Assessment of functional differences between subpopulations within resident macrophages using single-cell RNA sequencing would be beneficial. Interestingly, we found that CD206⁺ antiinflammatory macrophages were also increased in aging liver (Figure 2B); This is in line with the report showing counteract response of the macrophages to adapt to age-associated proinflammatory state (66). Recent study unraveled a novel mechanism of the gut-liver axis, suggesting that commensal microbe-derived components lead to suppression of liver-derived molecules which in turn regulate proliferation of intestinal stem cells (67). Investigating this mechanism in the context of aging would be interesting in order to determine whether there is a possible link between tissue crosstalk, immune cell activation, and age-associated inflammation. Collectively, our data suggest that

alteration of the tissue microenvironment in aging underlies the pro-inflammatory activation of macrophages in epididymal fat and liver in aging, thus contributing to systemic inflammation.

Given that impaired immune responses to vaccines and infection leads to high mortality in the elderly, we further studied how the innate immune cells respond to systemic inflammation in peripheral tissues in young and aged mice. The myeloid cell populations in tissue consist of resident immune cells and the recruited infiltrating immune cells from the circulation. In our study, we observed that the aged mice showed a greater increase of blood neutrophils in LPS exposure as compared to young mice (Figure 4A). On the other hand, we found that the neutrophil expansion in peripheral tissues (peritoneal fluid, epididymal fat, and colon) was less robust in aged mice than young mice with LPS treatment (Figures 5A, 6B, 8A). Our data are in line with the reports of low chemotaxis of neutrophils in the elderly (68, 69). Together, our data suggest that aging delays neutrophil recruitment into tissues, in response to an inflammatory insult.

Compared to neutrophils, infiltrating monocytes/macrophages showed distinctive profiles in response to LPS. We found that total circulating monocytes and their subsets were all significantly reduced in response to LPS in both young and aged mice, but aged mice showed a more significant reduction compared to young mice (Figure 4C). This result is likely attributed to the higher baseline number of circulating monocytes in aged mice in a steady state (Figure 1A). Our data is consistent with other reports that endotoxin-induced inflammation decreases the number of monocytes in humans in the early stages of inflammation (70–73). In contrast, another study reported that immature monocytes are promptly released to circulation in response to bacterial and viral infection (72, 74, 75). While emergency monopoiesis can enhance production of monocytes in the bone marrow under inflammatory stress (76), it replenishes the monocyte population within 4-8 hours after endotoxin exposure (72). We used low-dose LPS for 6 hours in our experiment, which is in line with the report above. We observed the differential expansion of monocytes/macrophages within tissues in aging under LPS (Figures 5B, 6D, 7C, 8B). Collectively, under LPS, monocytes in circulation exhibited a significant reduction whereas the tissue monocytes/macrophage were differentially regulated in young and aged mice.

Our data demonstrated that the inflammatory response of innate immune cells is cell- and/or tissue-dependent. We observed that young mice showed enhanced pro-inflammatory macrophages in peritoneal fluid and liver upon LPS exposure (Figures 5B, 7B). Our results are in line with the report showing that the peritoneal macrophages can rapidly infiltrate liver through non-vascular routes within 2 hours of post-injury (77). On the contrary, we observed the immune cells in epididymal fat from young mice were barely responsive to LPS (Figure 6C), suggesting an immune tolerance specific to the tissue niche under an inflammatory state. Interestingly, depending on tissue/ cell type, we found differential responses to LPS challenge in immune cells in aged mice, showing either hyper-reactivity, hypo-reactivity, or desensitization/tolerance (Figures 4B, 6C, 7A, 7B). Accumulating evidences indicate a correlation between inflammaging and trained immunity which is the response of immune cells becoming higher and quicker upon subsequent inflammatory exposure (78, 79). On the other hand, others have reported reduced responsiveness or tolerance of macrophages, exhibiting downregulated toll-like receptor 4 in consequential low-grade inflammatory challenge; this is a protective mechanism to limit cell damage and prevent septic shock (80-82). The macrophage response resulting in trained immunity or immunetolerance could be determined by multiple factors such as dose/time of LPS, and/or tissue microenvironmental factors (77, 83, 84). Better understanding of cell-specific responses and factors determining immune responses in aging will be beneficial for future development of therapeutic approaches for inflammaging-related chronic disease in the elderly (85). Taken together, our study revealed that the response of immunity to LPS challenges in aged mice is impaired tissue- and cell-dependent.

Unlike other tissues, colonic macrophages with proinflammatory cytokine expressions were not altered in response to LPS (0.75 mg/kg) in young mice (Figure 8B). However, others have shown that a higher dose of LPS (2-5 mg/kg) induces colon inflammation (86, 87), indicating the response of the colon to exogenous LPS is dose-dependent. Of note, colonic immune cells responded to intrinsic sterile inflammation induced by aging, thus exhibiting significantly elevated pro-inflammatory profiles (Figure 2C). This suggests that colonic immune cells are likely more responsive to intrinsic inflammatory factors rather than extrinsic factors. Notably, colonic macrophages from aged mice also did not show response to LPS (Figure 8B). The tolerance of aged colonic cells to LPS could be due to macrophage dysfunction caused by chronic exposure to sterile inflammation in aging. Indeed, it has been reported that prolonged exposure to pro-inflammatory mediators suppresses the macrophage functions (37).

Sexual dimorphism is an important biological variable in immune response. Age-associated changes of immune cells show sexdependent variability, reflected in cell types as well as the degree of immune responses (88); this potentially explains why males are more susceptible to infections while females are more vulnerable to autoimmune diseases (89-92). In chronic low-grade inflammation, peripheral blood monocytes in women exhibited higher immune activation transcriptomes compared to men (93). Leukocyte trafficking, especially of macrophages into peritoneal fluid, was higher in young female mice than young male mice, but it was similar in both old male and old female mice (61, 62). Even though the numbers of infiltrating macrophages are similar between sexes, a greater pro-inflammatory shift of macrophages was reported in male mice versus an anti-inflammatory shift in female mice was reported under infection and aging (94, 95). In the current long-term aging study, we focused on male mice. Future studies in females are needed to elucidate phenotype and functional change of innate immunity across various tissues during aging and under disease conditions.

Ample evidence shows that neutrophils, monocytes, and macrophages have essential roles in wound repair (96–98). These immune cells also shift their phenotype and functional properties during the repair stage. The previous report showed that the mature macrophages infiltrated from the peritoneum to the local site of injury can be alternatively activated 12 hours after local injury to secure the tissue integrity (99). Aging reduces the tissue regeneration ability after injury, which possibly can be attributed to age-associated impaired functions of immune cells (100, 101). The information is limited on what mechanisms of innate immune cells are involved in defective tissue repair in aging. While our data have shed light on aging innate immunity, it is important to note that in our study, the phenotypical and functional characteristics of myeloid cells were studied at 6 hours after LPS injection, a snapshot of the inflammatory state. Long-term longitudinal studies throughout aging are critically needed.

Age-associated chronic diseases, including obesity, type 2 diabetes, and atherosclerosis, are tightly associated with immune dysfunction (102). Current available therapeutic approaches such as anti-integrin/chemokine therapies, peptide inhibitors, and antiaging drugs like metformin and rapamycin, are known to regulate leukocyte trafficking and block pro-inflammatory mediators in the treatment of inflammation-driven diseases (62). However, these treatments are often accompanied with various side effects (62, 103, 104) due to lack of cell specificity and/or unsatisfactory efficacy in older adults. The current knowledge in tissue-specific immune programming in aging is very limited, and the mediators responsive to the local tissue environment in aging are largely unexplored. Our study underscores the cell-specific and tissuespecific alteration of innate immunity in aging. We investigated the dynamics of neutrophils and macrophages in various peripheral tissues under normal aging as well as innate immune responses to inflammatory stimulus in aging. The insights gained from our study offer a crucial foundation for developing tissue-specific prevention and therapeutic strategies for age-related chronic diseases. Future studies to further elucidate cell-autonomous effects and the interaction among local and systemic mediators using single-cell

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RNA sequencing, spatial transcriptomics, and cell-specific gene targeting would help to obtain better understanding of immunoregulation in aging and advance the development of celland tissue-specific therapeutic opportunities.

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 approved by Texas A&M University Institutional Animal Care and Use Committee (protocol code: IACUC 2022-0180, approval date: October 4, 2022). The study was conducted in accordance with the local legislation and institutional requirements.

Author contributions

JN: Conceptualization, Data curation, Formal analysis, Investigation, Methodology, Project administration, Software, Validation, Visualization, Writing – original draft, Writing – review & editing. HH: Data curation, Formal analysis, Methodology, Software, Writing – review & editing. DK: Data curation, Formal analysis, Methodology, Writing – review & editing. EG: Writing – review & editing. YF: Writing – review & editing. GW: Writing – review & editing. YS: Conceptualization, Funding acquisition, Investigation, Methodology, Project administration, Resources, Supervision, Writing – review & editing.

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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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Supplementary material

The Supplementary Material for this article can be found online at: https://www.frontiersin.org/articles/10.3389/fimmu.2024. 1357444/full#supplementary-material

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