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Impaired phagocytosis of apoptotic cells causes accumulation of bone
marrow-derived macrophages in aged mice

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Running title: Accumulation of macrophage in bone marrow of aging

Abstract

Accumulation of tissue macrophages is a significant characteristic of disease-associated chronic inflammation, and facilitates the progression of disease pathology. However, the functional roles of these bone marrow-derived macrophages (BMDMs) in aging are unclear. Here, we identified age-dependent macrophage accumulation in the bone marrow, showing that aging significantly increases the number of M1 macrophages and impairs polarization of BMDMs. We found that age-related dysregulation of BMDMs is associated with abnormal overexpression of the anti-inflammatory interleukin-10. BMDM dysregulation in aging impairs the expression levels of pro-inflammatory cytokines and genes involved in B-cell maturation and activation. Phagocytosis of apoptotic Jurkat cells by BMDMs was reduced because of low expression of phagocytic receptor CD14, indicating that increased apoptotic cells may result from defective phagocytosis of apoptotic cells in the BM of aged mice. Therefore, CD14 may represent a promising target for preventing BMDM dysregulation, and macrophage accumulation may provide diagnostic and therapeutic clues.

Introduction

Macrophages protect the host from foreign pathogens, phagocytose apoptotic cells and cellular debris, and promote macrophage proliferation in response to environmental cues (1). Macrophages are highly diverse and heterogeneous cells that promptly change their function depending on their location in the body (2). Depending on the stimulus, they are categorized into two phenotypes: classically activated macrophages (M1) and alternatively activated macrophages (M2) (3). M1/M2 classification is based on the cytokines associated with macrophage phenotypes (3). Bacterial lipopolysaccharide (LPS) and/or interferon- γ stimulate production of M1 macrophages, which promote pro-inflammatory cytokines and reactive oxygen species (ROS) with antimicrobial activity. Immunosuppressive cytokines such as interleukin (IL)-4 and IL-13 yield M2 macrophages, which are mostly associated with tissue repair and suppressing inflammation (4). M2 macrophages show high phagocytic activity but produce low levels of pro-inflammatory cytokines and high levels of the anti-inflammatory cytokine IL-10 (5). Macrophage phenotype is regulated by a combination of local tissue signals and disease programming of resident macrophages, resulting in characteristic tissue-dependent macrophage phenotypic alterations (6). Therefore, macrophages are highly responsive to local tissue signals and alter their phenotypes accordingly.

Macrophages play essential roles in the rapid and efficient removal of apoptotic cells without causing damage to local tissues or initiating immune responses. Therefore, impairment of the phagocytic clearance of apoptotic cells may lead to the development of a variety of diseases, such as chronic inflammation, autoimmunity, and developmental abnormalities (7). When the number of apoptotic or dying cells increases beyond the normal rate, cellular debris is cleared by resident macrophages or those recruited from the circulation or local proliferation. Tissue macrophages are capable of self-renewal and proliferate locally under various

disease conditions, such as parasitic infection, obesity, atherosclerosis, and cancer (8-10). These macrophages exhibit the phenotype of alternatively activated M2 macrophages, a process driven by Th2 cytokines such as IL-4 and IL-10.

The contribution of macrophages to disease progression raises an important question about the nature of macrophage accumulation during aging. This study investigated the functional role of bone marrow (BM)-derived macrophages (BMDMs) in age-associated changes. We monitored the expression levels of various cytokines and surface receptors that affect macrophage phenotypes and the phagocytic function of BMDMs toward apoptotic Jurkat cells during aging. Using the macrophage-specific tracer, calcium–iron inositol hexakisphosphate (IC-IP₆) (11), we showed an age-dependent increase in BMDMs with impaired M1/M2 polarization and decreased phagocytosis of apoptotic cells in aged BM. These results might provide novel insights into the diverse physiological and pathological roles of BMDMs with potential therapeutic and diagnostic benefits in aging.

Results

BMDM accumulation and dysregulation of macrophage polarization in aged mice

Macrophages undergo significant functional changes during aging; however, age-related macrophage distribution and histopathology in BM microenvironments remain unclear. As a macrophage-specific tracer (**Fig. 1a**), intravenously injected IC-IP₆ demonstrated maximal IC-IP₆ uptake at 4 h in the liver of young mice (**Supplementary Fig. 1**). Other tissues, including the brain, heart, lung, kidney, lymph node, and thymus, showed very low levels of IC-IP₆ uptake (**Supplementary Fig. 1**), as verified by Prussian blue staining to detect the presence of iron in IC-IP₆. IC-IP₆ was injected intravenously into young and aged mice. IC-IP₆ was engulfed by tissue macrophages, especially hepatic Kupffer cells, in a manner that did not differ significantly during aging (**Fig. 1b**). In contrast, the BM of aged mice had a markedly increased number of Prussian blue-positive cells compared with young mice (**Fig. 1c, 1d**). IHC confirmed the age-dependent increase in BMDMs in the BM (**Fig. 1e**). These results strongly suggest that aging significantly increases the number of BMDMs. We investigated the effects of aging on macrophage distribution in BM by analyzing the percentage of BMDMs and the frequency of M1 and M2 macrophages in young and aged mice. We found a significant increase in the percentage of CD11b⁺MHCII⁺F4/80⁺ cells in the BM of aged mice compared with that in young mice (**Fig. 1f**). The analysis of BMDM M1 and M2 phenotypes showed that aging had a significant impact on the polarization of BMDMs (**Fig. 1g**). The percentage of M1 macrophages was approximately 2.5-fold higher in the BM of aged mice than that in young mice, whereas no significant increase in M2 macrophages was found in aged mice (**Fig. 1g**). These results strongly suggest that the age-related impairment of M1/M2 BMDM polarization may dysregulate host immune responses and the microenvironment.

Dysregulation of cytokines and surface receptors in BMDMs in aged mice

Aging is associated with increased levels of circulating inflammatory cytokines including IL-6 and TNF- α (12); thus, we assessed the expression levels of cytokines and chemokines during aging. The expression levels of pro-inflammatory cytokines such as TNF α , IL-6, and IL1 β were reduced dramatically in BMDMs of aged mice compared with young mice (**Fig. 2a**), indicating that the dysregulation of BMDMs impairs the expression of pro-inflammatory cytokines. The expression profiles of anti-inflammatory cytokines IL-4 and IL-10 were also dysregulated in the BM of aged mice. IL-4 expression was markedly reduced in aged compared with young mice, but IL-10 expression increased significantly with age (**Fig. 2b**). As IL-10 inhibits the synthesis of pro-inflammatory cytokines (13), age-related dysregulation in BMDMs may be closely associated with the abnormal overexpression of anti-inflammatory IL-10 (**Fig. 2b**). Macrophages respond effectively to immune stimuli via surface receptors (3). We found that macrophage markers CD31 and CD36 increased significantly with aging, F4/80 and ARG1 increased slightly, and CD68 showed no significant change with aging (**Fig. 2c**).

Upregulation of genes associated with B-cell activation, survival, and surface antigen production

Surface antigens on macrophages play important roles in antigen presentation and activation of the adaptive immune response (1,2). Surface antigens CD19, CD22, IL7R, CD72, CD79A, and CD79B were overexpressed in an age-dependent manner in the BM (**Fig. 3a**). Furthermore, the expression levels of B-cell activating factor (BAFF) and BAFF receptor were increased in an age-dependent manner. In addition, the expression levels of BAFF and BAFF-R were increased significantly in the BM of aged mice (**Fig. 3a**). BAFF is responsible for B-cell survival, maturation, and

antibody production (14); therefore, excessive expression of BAFF might induce abnormally high antibody production, which could lead to autoimmune diseases especially those associated with aging (3). Furthermore, MHC class II molecules such as HLA-A-H2Q7, HLA-B-H2Q6, C1QC, and CD74 were increased significantly in the BM of aged mice (**Fig. 3b**), showing that genes involved in B-cell maturation and antigen production are highly upregulated in the BM of aged mice. CD74 plays a role in antigen presentation (15), and its loss inhibits B-cell development (16). In our study, IHC staining revealed a significant increase in CD74 expression in the BM of aged mice compared with young mice (**Fig. 3c, d**), suggesting that CD74 could be a potential therapeutic target in the BM for activation of B-cell survival and maturation during aging.

BMDM accumulation and phagocytic defects in apoptotic cells in aged mice

We hypothesized that both BMDM accumulation and increased expression of anti-inflammatory cytokines in aged mice are involved in the clearance of apoptotic cells or cellular debris. Using a fluorescently labeled suicide inhibitor with a DEVD sequence, the specific cleavage site for caspases 3 and 7 (FAM-FLICA, ImmunoChemistry Tech), we found that apoptosis was increased significantly in the BM of aged mice (**Fig. 4a**). Furthermore, immunohistochemical staining of cleaved caspase 3 showed an increased rate of apoptosis in the BM of aged compared with control mice (**Fig. 4b, c**). Consistent with previous reports suggesting that aging increases the rate of apoptosis (7,17), these results implicate that BMDM accumulation in the BM of aged mice might be due to the high incidence of apoptosis and low clearance of BMDMs. The rapid and efficient phagocytic clearance of cell debris is an essential process for maintaining tissue health and preventing autoantigen production (18). Therefore, we assessed the phagocytic function of BMDMs toward apoptotic Jurkat cells after inducing apoptosis using camptothecin.

Engulfed and non-engulfed apoptotic Jurkat cells were distinguished by labeling apoptotic Jurkat cells with a pH-sensitive dye (pHrodo) that emits red fluorescence, only when the engulfed cells are localized inside the phagosome of macrophages (pH ~5) (19). After inducing apoptosis, the BMDMs from young mice efficiently engulfed most of the apoptotic Jurkat cells (**Fig. 4d**). The BMDMs from aged mice displayed dramatically decreased engulfing activity (**Fig 4e**), indicating that this phagocytic function decreases with aging and leads to increased numbers of apoptotic cells in the BM of aged mice. To address the molecular mechanism underlying the impaired phagocytosis of apoptotic cells, we analyzed the expression levels of the genes associated with phagocytic clearance of apoptotic cells (20). The expression of TIMD4 and Stabilin-2 were dramatically upregulated on macrophages, whereas the expression levels of BAI-1, BAI-3 and TIMD1 were not detected (data now shown), suggesting these genes are not expressed in BMDMs. Most importantly, the expression of CD14, the receptor highly expressed on macrophage (21), was dramatically decreased in the BM of aged mice (**Fig. 4f**), confirming an age-related decline in phagocytosis receptors (22). Therefore, the impaired phagocytic function of BMDM toward apoptotic cells might involve low CD14 expression in aged mice.

Discussion

Tissue macrophage accumulation is a characteristic of disease-associated chronic inflammation including atherosclerosis, cancer, chronic kidney disease, obesity, and type 2 diabetes. Macrophage accumulation is considered a consequence of both monocyte migration and local proliferation, which can induce disease pathology. Therefore, monitoring tissue macrophages in aged animals might reveal the underlying mechanisms of tissue aging and age-related pathologies.

This study found that aging causes significant increases in BMDM numbers and impaired M1/M2 polarization in BM. Dysregulated M1/M2 polarization alters the

secretory pattern of pro- and anti-inflammatory cytokines. Notably, overexpression of IL-10 in aged mice may suppress the expression of pro-inflammatory cytokines by the increased number of M1 macrophages (4,13,22,23). These findings suggest that the abnormal overexpression of anti-inflammatory IL-10 in the BM of aged mice could be a cause of dysregulated polarization of BMDMs.

Macrophages play important roles in host defenses against invading pathogens, clearing cellular debris, and antigen presentation to induce the adaptive immune response (1,24). We observed age-dependent overexpression of surface antigens associated with B-cell survival, maturation, and autoantibody production in the BM of aged mice. Concomitantly, MHC class II surface antigen (HLAs, C1Q, and CD74) expression levels were increased in the BM, implying that dysregulation of BMDMs in aged mice plays an important role in B-cell activation and autoantigen presentation. Experimental evidence showed that age-related dysregulation of BMDMs might play a critical role in age-related impairment of adaptive immunity and progression of age-related diseases (13,14,23).

The significantly decreased phagocytosis of apoptotic Jurkat cells by BMDMs from aged mice suggests that the defective clearance of apoptotic cells leads to increased apoptotic cell numbers in the BM of aged mice (23). However, a previous study reported no impairment of BMDM-mediated phagocytosis of fluorescent particles in aged mice (22). We suggest that this conflicting data could be related to the different phagocytosis assay methods employed. As anti-inflammatory IL-10 dysregulates the phagocytic function of peritoneal macrophages (22,25), high expression of IL-10 in the BM of aged mice may aggravate the defective phagocytosis of apoptotic Jurkat cells by BMDMs. Although IL-10 increases expression of the phagocytic receptor CD14 and stimulates phagocytic function toward apoptotic cells (26), there are several conflicting results regarding IL-10. Decreased subsets of CD14⁺⁺CD16⁻ monocytes were found in patients with heart

failure despite high levels of IL-10 (26), and increased expression of IL-10 was detected in CD14 knockout mice (27). Therefore, our findings suggest that dysregulation of IL-10 in aged tissues leads to weak expression of the phagocytic receptor CD14. Low levels of CD14 in aged BMDMs may underlie the age-associated impairment of apoptotic cell phagocytosis (23).

This study demonstrated that aging causes increased BMDM numbers, dysregulated BMDM polarization, increased anti-inflammatory IL-10 expression, impaired phagocytosis of apoptotic cells, and reduced expression of CD14. These findings might provide new molecular targets with potential therapeutic and diagnostic benefits for age-associated diseases.

Materials and Methods

Antibodies and reagents

The following antibodies were used for Western blotting, confocal microscopy, and flow cytometry analyses: mouse F4/80 (BM8, eBioscience, San Diego, CA, USA), β -actin (AC-15, Sigma-Aldrich), mouse CD74 (05-591, Millipore), human Annexin V (ab14196, Abcam), and cleaved caspase-3 (D175, Cell Signaling). For chemical induction of apoptosis, cells were treated with 10 μ M camptothecin at 37°C for 16 h using an apoptosis inducer kit (ab102480; Abcam).

Animal care and use.

WT C57Bl/6 mice were from Orient (Korea). Male mice were studied under protocols approved by the animal ethics committee of Gachon University, Lee Gil Ya Cancer and Diabetes Institute. We used mice aged 2- 12 months as the young control group and defined aged mice as old as 18-34 months.

Preparation of IC-IP₆

Based on our isothermal titration calorimetry results (28), a 10 mM solution of Fe³⁺-IP₆ was prepared by mixing equimolar concentrations of Fe³⁺ and phytate (pH 6.0). To minimize chelation in the blood, an equimolar concentration of Ca²⁺ was added to the Fe³⁺-IP₆ solution and adjusted to pH 6.0 by adding 1 N NaOH.

Assessment of IC-IP₆ uptake by BMDMs

IC-IP₆ (10 mg·kg⁻¹) was administered intravenously to 6-, 12-, and 18-month-old C57BL/6 mice. Femurs were harvested after 4 h, fixed in 10% neutral-buffered formalin for 48 h, decalcified in 10% EDTA (pH 7.0) for 4–5 weeks at 4°C, and embedded in paraffin for immunohistochemistry (IHC) (28). IC-IP₆ in BM sections was detected by Prussian blue staining.

Generation of apoptotic cells

Jurkat T cells were cultured in RPMI media supplemented with 10% FBS and 1% penicillin/streptomycin. Apoptosis was induced with camptothecin (10 μ M) at 37°C for 16 h.

Detection of apoptotic cells in BM tissues

For IHC, antigen retrieval was performed in Tris/EDTA (pH 9.0) with 2–3 min of boiling. Bone sections (2.5- μ m thick) from 6-, 12-, and 18-month-old mice were treated with serum-free protein blocking solution (Dako), incubated with anti-cleaved caspase 3 rabbit antibodies (9661, 1:200; Millipore) followed by horseradish peroxidase-conjugated secondary antibodies for 30 min, and detected with 3,3'-diaminobenzidine tetrahydrochloride (Dako).

***In vitro* phagocytosis assay**

Jurkat T cells were cultured overnight at a density of 5×10^6 /mL and treated with camptothecin (10 μ M) for 16 h. Growing or apoptotic Jurkat T cells were washed with PBS and incubated with 5 μ M/mL pHrodo Red AM (P353721, Life Technologies) for 30 min at 37°C. The pHrodo-labeled Jurkat T cells were washed with live cell imaging solution (A14291DJ, Life Technologies). BMDMs were cultured on 12-well plates (5×10^5 /well) for 16 h before treatment with 0.5 μ g/mL LPS for 6 h (18). For analysis of phagocytosis, pHrodo-labeled Jurkat T cells were co-incubated with BMDMs for 1.5–2 h, washed with cold PBS, fixed in 10% neutral-buffered formalin for 10 min, counterstained with DAPI to visualize the nuclei, and analyzed by confocal microscopy (Zeiss LSM 710).

Macrophage polarization and apoptosis analysis

BMDMs were isolated from 2-month-old or 22-month-old C56/BL mice. Staining was performed at 4°C for 30 min. The M1/M2 macrophage phenotype was analyzed using anti-CD11b-FITC (M1/70), anti-F4/80-PE-Cy7 (BM8), major histocompatibility complex (MHC) class II, and CD206-APC (C068C2) antibodies. All antibodies were from eBioscience except for anti-CD206 (Biolegend). For apoptotic cell analysis, BM cells were isolated from 7-, 13-, and 19-month-old C57/BL6 mice and co-stained with the caspase 3/7-specific Vybrant-FAM substrate, a fluorescein-conjugated caspase inhibitor, FAM-DEVD-fluoromethyl ketone (V35118, Molecular Probes) and Alexa 647-conjugated Annexin V. The cells were analyzed by flow cytometry using the FACS LSRII (BD Bioscience) and FlowJo software.

Quantitative PCR (qPCR) analysis of cytokines and surface receptors

Total RNA was treated with RQ1 RNase-free DNase (Promega) and reverse transcribed using the PrimeScript First-Strand cDNA Synthesis kit (TaKaRa Bio). qPCR was performed using SYBR Green PCR Master Mix (TaKaRa Bio) with gene-specific primers (**Supplementary Table 1**) on the ABI PRISM 7900 Sequence detection system. All samples were analyzed in triplicate, and the results are expressed as means \pm SE.

Statistical analysis

The error bars on the graphs represent the means \pm SE of at least three independent experiments. Student's *t*-tests were used to determine statistical significance between groups, with $P < 0.05$ set as the cut-off value.

Acknowledgments

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Figure Legends

Figure 1. Increased numbers and dysregulation of bone marrow (BM)-derived macrophages (BMDMs) in aged mice. **a**, Schematic illustration of IC-IP₆. **b**, IC-InsP₆ accumulation in the liver was visualized by double-staining with Prussian blue (iron) and the macrophage-specific anti-F4/80 antibody (brown). Scale bars: top, 50 μ m; bottom, 20 μ m. **c**, Prussian blue staining for assessment of IC-IP₆ uptake by BMDMs in decalcified femur sections from age-matched mice. **d**, Quantification of Prussian blue-positive cells per mm² in the BM of mice intravenously injected with 10 μ M IC-IP₆, mean \pm SE (n=10, ***P < 10⁻³). **e**, Immunohistochemical staining of F4/80 in decalcified femur sections from age-matched mice. Scale bars: top, 200 μ m; bottom, 50 μ m. **f**, Frequency of total BMDMs assessed by flow cytometric analysis of CD11b⁺MHCII⁺F4/80⁺ cells in the BM of age-matched mice, mean \pm SE (n=6–7, ***P < 10⁻³). **g**, M1 and M2 macrophage polarization assessed by flow cytometric analysis of CD11b⁺/CD206⁺ and CD11b⁺/CD206⁻ macrophages from MHCII⁺F4/80⁺ cells in BM cells of age-matched mice, respectively, mean \pm SE (n=6–7, **P < 0.01).

Figure 2. Dysregulated mRNA expression of cytokines and surface receptors in aged mice. **a–c**, The mRNA levels of pro-inflammatory cytokines (**a**), anti-inflammatory cytokines (**b**), and surface receptors (**c**) in the BM of age-matched mice were measured by qPCR. All mRNA levels were normalized to cyclophilin A, mean \pm SE (n=4, *P<0.05).

Figure 3. mRNA expression of genes associated with B-cell activation, survival, and antigen production is upregulated in the BM of aged mice. **a**, The mRNA levels of genes involved in B-cell development and function (**a**), B-cell activating factor (BAFF), and BAFF receptor (BAFF-R) (**b**), and MHC class II molecules. **c**, Immunohistochemical staining of CD74 in decalcified femur sections from age-

matched mice. Scale bars: top, 200 μm ; bottom, 50 μm . **d**, Quantification of CD74⁺ cells per mm^2 in decalcified femur sections from age-matched mice, mean \pm SE (n=10, *P<0.05).

Figure 4. Age-dependent accumulation of BMDMs may result from phagocytic impairment of apoptotic cells through reduced levels of the phagocytic receptor CD14. **a**, Rate of apoptosis assessed using a fluorescently labeled suicide inhibitor with a DEVD sequence, the specific cleavage site for caspases 3 and 7, in BM cells of age-matched mice, mean \pm SE (n=3–4, *P<0.05). **b**, Immunohistochemical staining of cleaved caspase 3 in decalcified femur sections from age-matched mice. Scale bars: top, 200 μm ; bottom, 50 μm . **c**, Quantification of the area (%) of cleaved caspase 3 staining in decalcified femur sections from age-matched mice, mean \pm SE (n=10, *P<0.05). **d**, Representative images of pHrodo-labeled apoptotic Jurkat cell phagocytosed (red) by BMDMs in age-matched mice. Scale bar: 50 μm . **e**, Quantification of phagocytosis, mean \pm SE (n=6–8, ***P < 10⁻³). **f**, mRNA levels of genes associated with phagocytic clearance of apoptotic cells in BMs of age-matched mice measured by qPCR, mean \pm SE (n=4, *P<0.05).

Supplementary Figure 1. Assessment of IC-IP₆ biodistribution in C57BL/6 mice. **a–h**, Representative Prussian blue staining of tissues at various time points after intravenous administration of IC-InsP₆.

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Figure 1

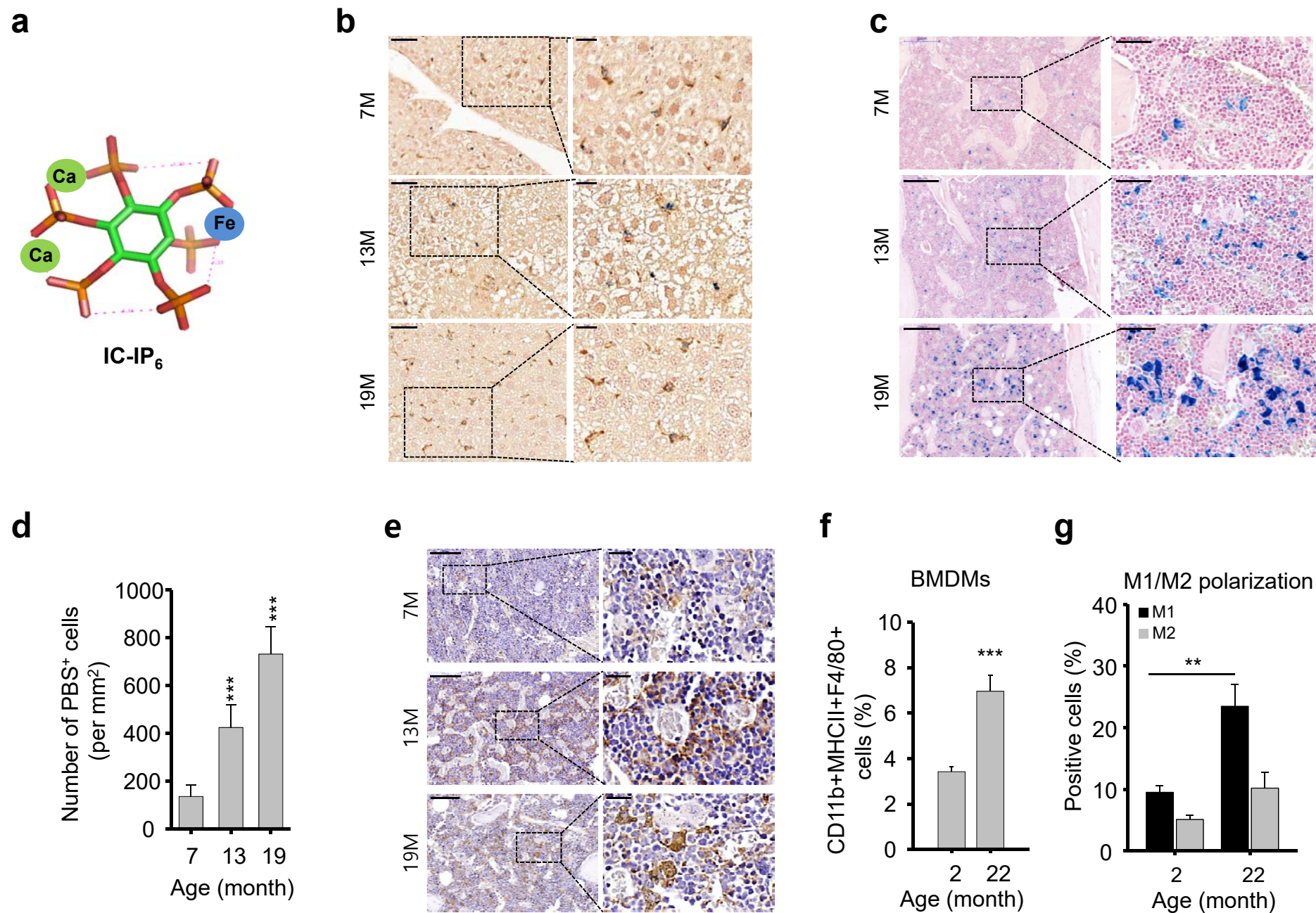


Figure 2

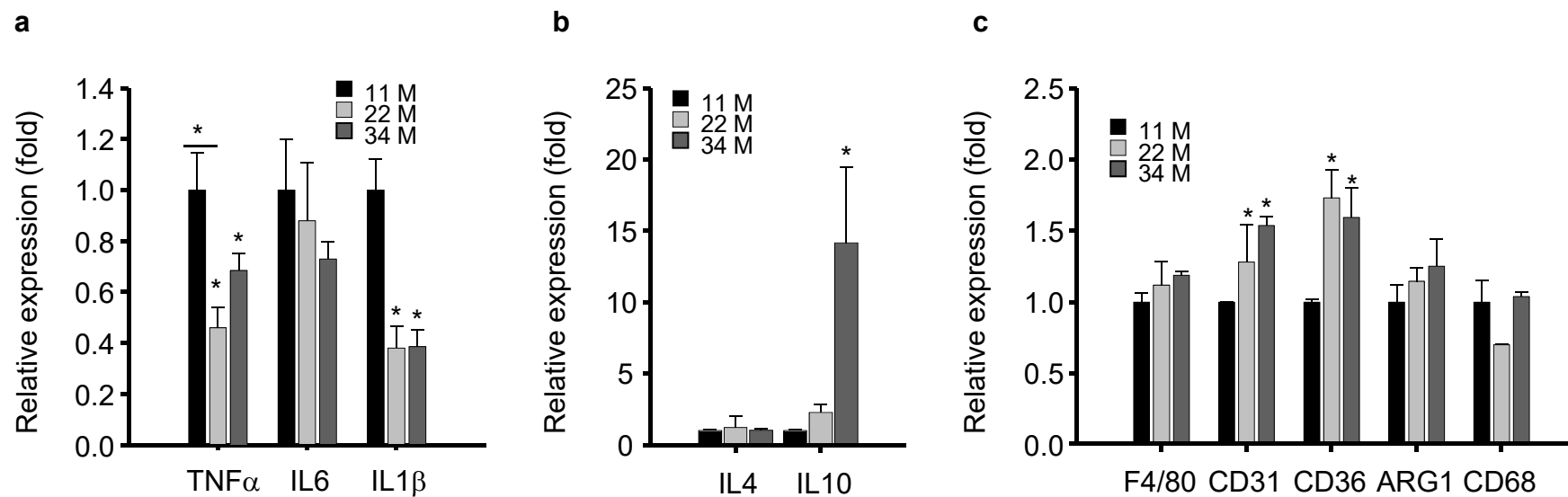


Figure 3

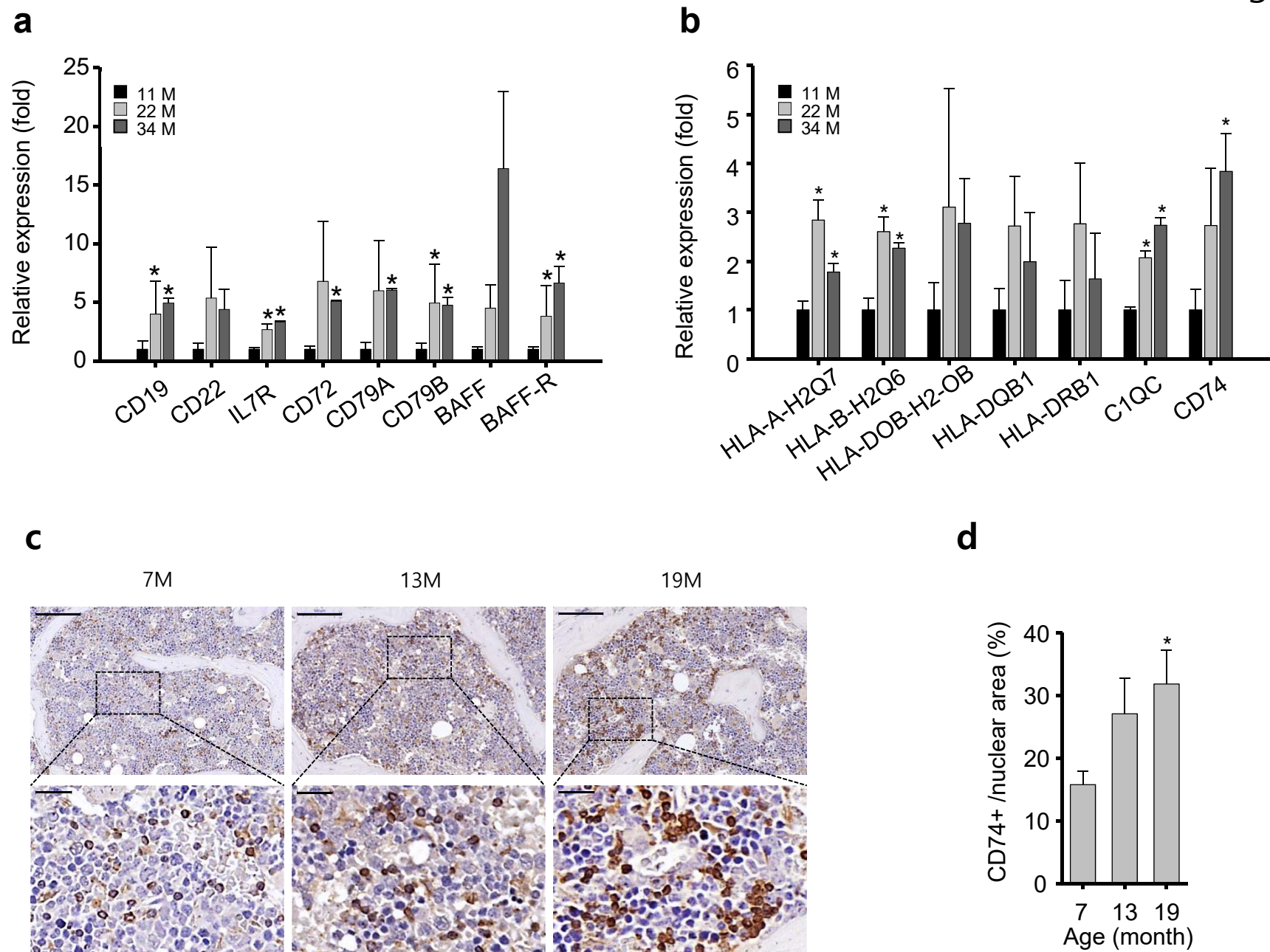
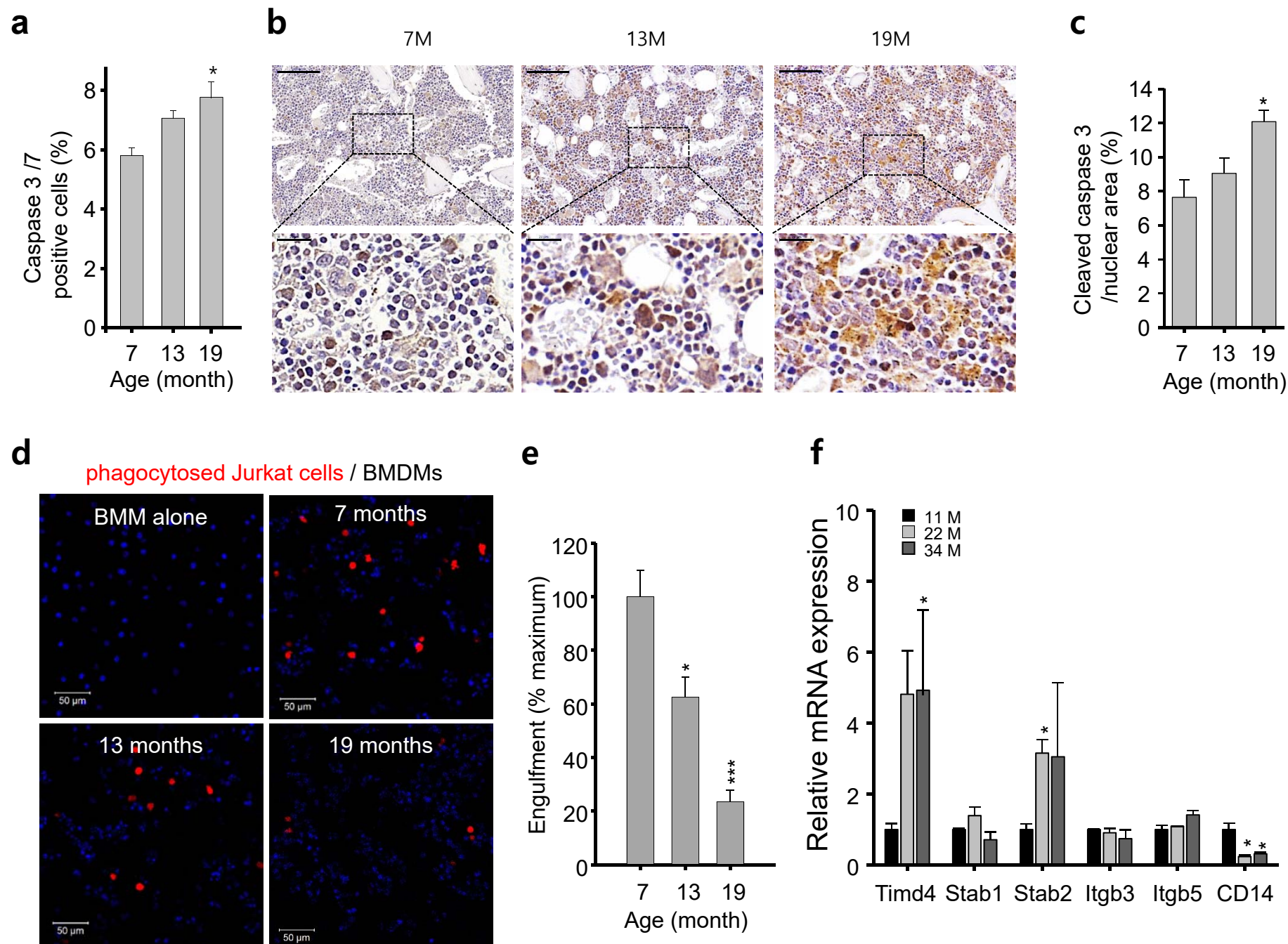
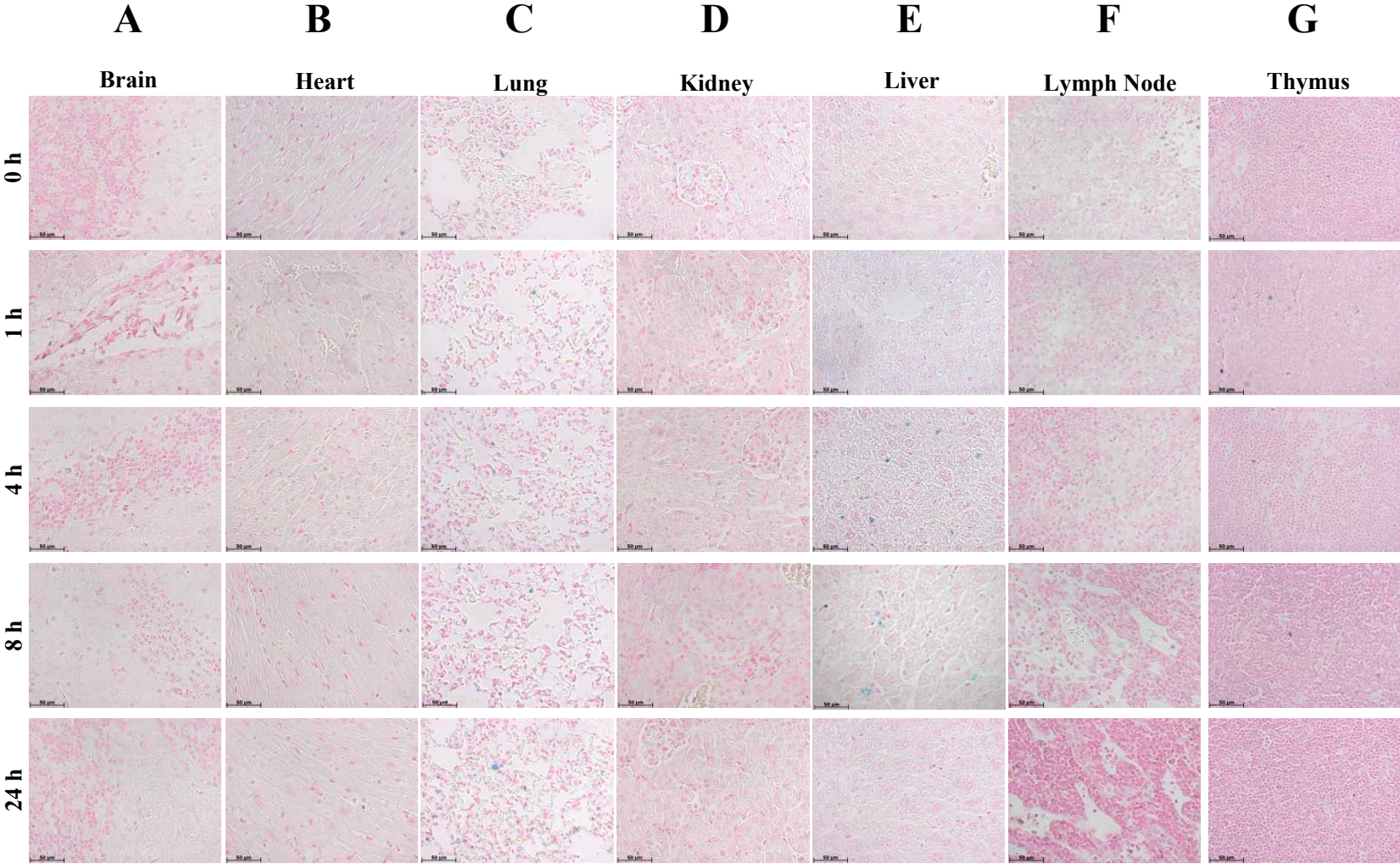


Figure 4



Supplementary Figure 1



Supplementary Table 1. Primers used for qRT-PCR

Gene	GenBank accession no.	Primer	Primer sequences	Gene	GenBank accession no.	Primer	Primer sequences
CD19	NM_009844	Forward	5'- GACCCTTTGCTTTCCCAAA	CD74	NM_001042605	Forward	5'- ACATGACCAGGACCATGTGA
		Reverse	5'- CTTCCTCTCATCCACACAA			Reverse	5'- CTGGGAAGTCCCTTCAG
CD72	NM_007654	Forward	5'- CTCTGTGAACCTGCTGCTCTCA	CD36	NM_007643	Forward	ATACAGAGTTCGTTATCTAGCCAAGGA
		Reverse	5'- ACGAGAAGGCCAAGCAAGAA			Reverse	CCATTGGGCTGTACAAAAGACA
CD74	NM_001042605	Forward	5'- ACATGACCAGGACCATGTGA	CD31		Forward	5'-TACTGCGGATCGGCAAA
		Reverse	5'- CTGGGAAGTCCCTTCAG			Reverse	5'-ATACTGGGCTTCGAGGACATT
CD79A	NM_007655	Forward	5'- GCCTCTCCTCTCTTCTGTGAT	ARG-1	NM_007482.2	Forward	5'-GAACCTCAACGGGAGGGTAACC
		Reverse	5'- AAGTTCACCGTCAGGGATGGT			Reverse	5'-TTTTCATGTGGCGATTAC
CD79B	NM_008339	Forward	5'- CTGGGATGGAGGAAGATCACA	F4/80	NM_153100.1	Forward	5'-GCCTATTATCTATACCTCCAGCACATC
		Reverse	5'- TGTCCGAAGAGTCACTATGCTTCA			Reverse	5'-TCCATCTCCCATCCTCCACATCAG
HLA-DRB1	NM_010382	Forward	5'- AGCCCTGGAACACCACAA	CD68	NM_009853	Forward	5'- ATGCGGCTCCTGTGTGT
		Reverse	5'- TCCGGAACCATCTGACTTCAA			Reverse	5'- GGCTTTTTGTGAGGACAGTCT
HLA-DQB1	NM_207105	Forward	5'- CATGGGCGAGTGCTACTTCA	CD64	NM_010186	Forward	5'-TTGAGATGACATGTGGCTCTAA
		Reverse	5'- ACGTACTCTCCCGTTGTAGA			Reverse	5'-CAGCCTTGGTGGCATTAAACC
CIITA	NM_001243760	Forward	5'- GGTGTCCTGCGTGTGATG	CD14	NM_009841.3	Forward	5'-GCTCAACTTTCTCGGAACA
		Reverse	5'- GTTTCACCTGAGGGCACTTC			Reverse	5'-TCGCCCAATTCAGGATTGTC
C1QC	NM_007574	Forward	5'- GCGATGAGGTGTGGCTATCA	C1QA	NM_007572	Forward	5'- TGGCAGTGTCTGCTGATCA
		Reverse	5'- GGAAACCAGAGAAGACGCTGTT			Reverse	5'- GGTCCGATCAAATGGATGCT
CHST1	NM_023850	Forward	5'- GCTCAGCTGGGTACAAGATG	C1QB	NM_009777	Forward	5'- TCGCTTCGAAAGGTGATCA
		Reverse	5'- GGAAGTCACGCTCCTCCACTA			Reverse	5'- AGGCCAGGCACCTTGCA
IL7R	NM_008372	Forward	5'- CGATCCATCCCCATAACGA	HLA-DOB-H2-OB	NM_010389	Forward	5'- ACAGGCAACCGCAGTGAAGT
		Reverse	5'- GGGTTCTGGAGTTCAAGGT			Reverse	5'- TTGTGTTTGACCTGAGCAAGTG
HLA-A	NM_010391	Forward	5'- CTGATCTGCTGTGCCATCA	C1QC	NM_007574	Forward	5'- GCGATGAGGTGTGGCTATCA
		Reverse	5'- TCCCTCTTTTCTACCTGTGTTT			Reverse	5'- GGAAACCAGAGAAGACGCTGTT
HLA-B	NM_008199	Forward	5'- GGTTTCATCTGTGCGTACGT	HLA-B-H2-Q6	NM_207648	Forward	5'- AACTCGCTGCGGTATTTC
		Reverse	5'- GCGGCTCAGATCTCGGATT			Reverse	5'- TCCACGTAGCCGACGATAATG
HLA-DOB-H2-OB	NM_010389	Forward	5'- ACAGGCAACCGCAGTGAAGT	HLA-DQB1	NM_207105	Forward	5'- CATGGGCGAGTGCTACTTCA
		Reverse	5'- TTGTGTTTGACCTGAGCAAGTG			Reverse	5'- ACGTACTCTCCCGTTGTAGA
HLA-B-H2-Q6	NM_207648	Forward	5'- AACTCGCTGCGGTATTTC	HLA-A-H2-Q7	NM_001198560	Forward	5'- AGCCCTCAGTCTCTTAGACAACA
		Reverse	5'- TCCACGTAGCCGACGATAATG			Reverse	5'- GCTGGGCTCCACAGTTCT
HLA-A-H2-Q7	NM_001198560	Forward	5'- AGCCCTCAGTCTCTTAGACAACA	HLA-DRB1	NM_010382	Forward	5'- AGCCCTGGAACACCACAA
		Reverse	5'- GCTGGGCTCCACAGTTCT			Reverse	5'- TCCGGAACCATCTGACTTCAA
POU2AF1	NM_011136	Forward	5'- GAAAGCACGCCAGTCACAT	HLA-B	NM_008199	Forward	5'- GGTTTCATCTGTGCGTACGT
		Reverse	5'- GAGCATGGCTTGACAGGAT			Reverse	5'- GCGGCTCAGATCTCGGATT
SPIB	NM_019866	Forward	5'- CGCAAGCGCATGACGTATC	HLA-A	NM_010391	Forward	5'- CTGATCTGCTGTGCCATCA
		Reverse	5'- GCGTTTGACCTTGCGGATT			Reverse	5'- TCCCTCTTTTCTACCTGTGTTT
CD22	NM_001043317	Forward	5'- CGCCACAAGACTGAGAAGGA	BLNK	NM_008528	Forward	5'- TTCCACAGCTTCCCATTAATTC
		Reverse	5'- CTATTCTGTTCCCGAAGATGTACTC			Reverse	5'- ACAGGCGCCAGCATACCA
Tnfrsf13c	NM_028075	Forward	5'- TTCAGAAGGAGTCCAGCAAGAGT	MAPK13	NM_011950	Forward	5'- CCCTTCGGAGCTTCATGAT
		Reverse	5'- CAGGCCCCCAGGAGATG			Reverse	5'- TGAATTCATCCCCATGATCTT
Tnfrsf13b	NM_021349	Forward	5'- GAGGCAGCGCACCTGTACA	CEBPD	NM_007679	Forward	5'- CGACTTCAGCGCTACATTG
		Reverse	5'- CAGGCCCCCAGGAGATG			Reverse	5'- TGAAGAGTCCGCGAAGAGT