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MINIREVIEWS

Immunological aspects of age-related diseases

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Abstract

The proportion of elderly people rises in the developed countries. The increased susceptibility of the elderly to infectious diseases is caused by immune dysfunction,

especially T cell functional decline. Age-related hematopoietic stem cells deviate from lymphoid lineage to myeloid lineage. Thymus shrinks early in life, which is followed by the decline of naïve T cells. T-cell receptor repertoire diversity declines by aging, which is caused by cytomegalovirus-driven T cell clonal expansion. Functional decline of B cell induces antibody affinity declines by aging. Many effector functions including phagocytosis of myeloid cells are down regulated by aging. The studies of aging of myeloid cells have some controversial results. Although M1 macrophages have been shown to be replaced by antiinflammatory (M2) macrophages by advanced age, many human studies showed that pro-inflammatory cytokines are elevated in older human. To solve this discrepancy here we divide age-related pathological changes into two categories. One is an aging of immune cell itself. Second is involvement of immune cells to age-related pathological changes. Cellular senescence and damaged cells in aged tissue recruit pro-inflammatory M1 macrophages, which produce pro-inflammatory cytokines and proceed to agerelated diseases. Underlying biochemical and metabolic studies will open nutritional treatment.

Key words: Elderly people; Damage associated molecular patterns; Immune dysfunction; Lymphoid lineage; Myeloid lineage; Shrinkage of thymus; Cytomegalovirus; Age-related tissue damage; Cellular senescence; Pro-inflammatory

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Core tip: The authors divide immune cells, which are involved in age-related pathological changes, into two categories. First category is aging of immune cell-itself, which include age-related myeloid lineage deviation of hematopoietic stem cells, the shrinkage of thymus followed by the decline of naïve T cells, the cytomegalovirus infection-mediated decline of T-cell receptor repertoire diversity and functional decline of myeloid cells. Second category is the involvement of immune cells to agerelated pathological changes. Age-related tissue damage and cellular senescence activate pro-inflammatory (M1) macrophages, which induce many age-related diseases.



Biochemical works are needed to further elucidate agerelated immune changes.

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INTRODUCTION

The proportion of elderly people rise world wide, especial in the developed countries. Aging-related changes in the immune system contribute to the increased susceptibility of the elderly to infectious diseases, cardiovascular disease and stroke caused by atherosclerosis, autoimmune disease such as rheumatoid arthritis, cancer and late onset anemia and degenerative diseases including Alzheimer diseases^[1,2]. Further metabolic syndrome, which is caused by obesity occurs from middle age and proceeds to tissue failure such as renal failure in advanced age, is tightly related to immune system. Chronic infections such as hepatitis virus and chronic tissue damages caused by tobacco inhalation induce tissue damage, which arouses immune responses and wound repair responses. Chronic inflammation follows tissue fibrosis in advanced age proceeding to tissue failure such as chronic obstructive pulmonary disease. Most prominent cause of age-related immune dysfunction is T cell immunosenescence. There are three causes of T cell immunosenescence. One is the age-related hematopoietic stem cells (HSCs) deviation from lymphoid lineage to myeloid lineage. Second is the shrinkage of thymus. Third is expansion of T cell clones to cytomegalovirus (CMV). Changes of HSCs also affect immunosenescence. HSCs deviate to myeloid lineage by aging. Both in mouse and human myeloid-lymphoid ratio elevates by aging^[3,4], which induces the decline of lymphoid cells (T and B cells) and erythrocytes and contribute age-related anemia and decline of adaptive immunity^[5,6]. The number of aged B cells decline and affinity and diversity of antibodies are low^[7]. Ageing related myeloid deviation increases the number of myeloid cells. However, the oxidative burst and phagocytosis of both neutrophils and macrophages are decreased^[8]. The antigen presentation of aged dendritic cells and the cytolysis of Natural killer cells are low^[9]. Pro-inflammatory cytokines such as interleukin-6 (IL-6), tumor necrosis factor α (TNF- α) and IL-1 β are elevated in elderly people, which are called inflammaging and induce metabolic syndrome such as obesity, type II diabetes, atherosclerosis, cardiac diseases^[10]. However, other reports have shown that pro-inflammatory (M1) macrophages are replaced by anti-inflammatory (M2) macrophages by aging^[11]. Here to solve this discrepancy we propose to classify age-related immune changes as follows. Age-related pathological changes are classified as immune cell intrinsic changes by aging and involvement of immune cells to age-related pathological changes.

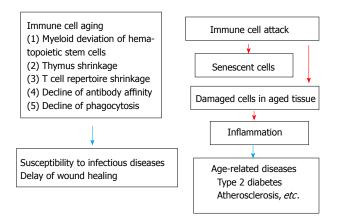


Figure 1 Two types of engagement of immune cells to aging.

Increased susceptibility of the elderly to infectious diseases is mainly caused by age-related immune cell intrinsic changes usually called immunosenescence. Metabolic syndrome and other related diseases, which occur at aged people, are mainly caused by immune cell attack to agerelated tissue changes. Age-related tissue failure is caused by repeated immune cell attack. Here we first describe the recent findings from the points of immune cell intrinsic changes by aging and then go to age-related tissue changes, which induce immune reaction (Figure 1).

IMMUNE CELL INTRINSIC CHANGES BY AGING (IMMUNOSENESCENCE)

HSCs aging

Adult immune cells develop from HSCs that originate from the bone marrow (BM) (Figure 2). From HSCs common lymphoid progenitor cells (CLPs) and myeloid precursor cells develop. T cells emigrate from BM to enter thymus, where they develop to naïve CD4 and CD8 T cells. B cells develop in bone marrow from pre-pro-B cells, pro-B cells and emigrate from bone marrow as pre-B cells to circulation and secondary lymph nodes and spleen. Finally B cells further differentiate to plasma cells to produce antibodies (Figures 2 and 3).

Although the cellularity of BM decreases by aging^[12], the number of HSCs in aged mice differ between mouse stains. The number of HSCs of aged-C57BL/6 mice increases, whereas other strain of mice decreases^[13,14]. However, in HSCs of aged mice the self-renewal activity and homing activity to BM are two-fold less efficient compared with young HSCs^[2,6,15-17]. It has been found by transplantation experiments that old HSCs tend to have a myeloid-skewed blood cell production caused by a decreased ability to produce lymphoid cells^[3,18]. These age-related changes of HSCs induce age-related abnormalities such as myeloid leukemia^[19], late onset anemia^[20]. Biochemical mechanism underlining HSC aging has been elucidated recently. The small RhoGTPase Cdc42 in aged HSCs is elevated, which correlates with a loss of polarity in aged HSCs. It has been found that

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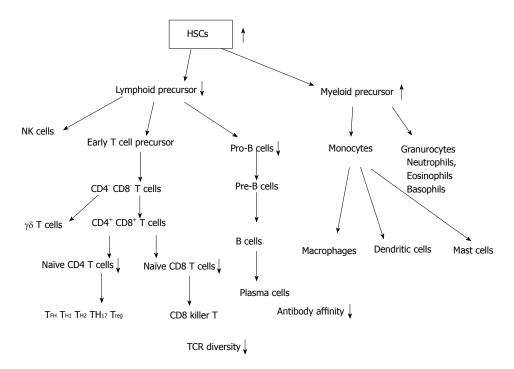


Figure 2 Hematopoietic stem cells to mature immune cells. HSCs: Hematopoietic stem cells; TCR: T-cell receptor.

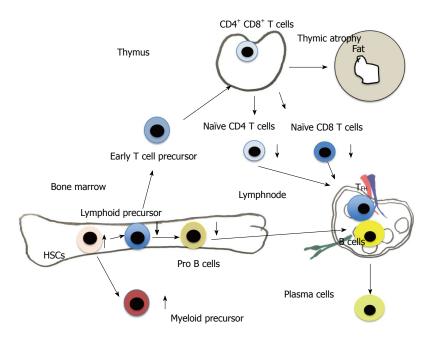


Figure 3 Differentiation of hematopoietic stem cells to mature immune cells.

in young HSCs Cdc42 and tubulin are asymmetrically distributed at the same location, whereas in aged HSCs Cdc42 and tubulin were distributed throughout the cell body in an unpolarized fashion^[21]. HSCs, which produce all mature blood cells, must counterbalance the deleterious effects of chronic stress by generating new cells to replace those that become damaged or destroyed. Cells adapt chronic stress by autophagy and apoptosis^[22]. Autophagy adapts starvation by using degraded own cell organelles for nutrient. One report has shown that autophagy is suggested to be decreased by aging^[23]. However, another report has shown that

by electron microscopic analysis autophagy is increased by aged mice, and old HSCs retain an intact FoxO3adriven pro-autophagy gene program^[24]. *FOXO* genes are key regulators of longevity downstream of insulin and insulin like growth factor signaling^[25]. MicroRNAs, a class of small-noncoding RNAs, regulate normal function of HSCs, including cell cycling and engraftment potential^[26]. The microRNA-212/132 cluster (Mirc19) is enriched in HSCs and is up regulated during aging and plays a role in maintaining balanced hematopoietic output by buffering FOXO3 expression^[27]. There exist many other molecules, which relate HSCs aging. For instance recent work done by Chang *et al*^[28] showed that ABT263 (a specific inhibitor of the anti-apoptotic proteins BCL-2 and BCL-xL) rejuvenated aged HSCs.</sup>

T cell aging

One of the most early and dramatic changes in aging of immune system is the decrease of naïve T cells. Naïve T cells are derived from thymus and continuously recirculate through the peripheral lymphoid tissues. Naïve T cells differentiate into effector/memory T cells after primed with cognate antigen presented by dendritic cells. Naïve T cells originate from thymus. Naïve T cell production from thymus declines very early only after first year of life. These naïve T cell decline comes from the early thymic involution^[29,30] (Figures 2 and 3). This dramatic decline of human naïve T cell output is compensated by homeostatic proliferation^[31,32]. CD4⁺CD45RA⁺CCR7⁺ human naive T cells are recently found to be divided by CD31. CD31⁺ naïve T cells are recent thymic emigrants (RTEs). By aging CD31⁺ CD4⁺CD45RA⁺ decrease, while CD31⁻ CD4⁺CD45RA⁺ increase during healthy aging by homeostatic proliferation^[33-35]. Recent work using neonatal thymectomy has shown that total number of CD3⁺, CD4⁺ and CD8⁺ cell numbers decrease in the first year after neonatal thymectomy. CD4⁺CD45RA⁺CCR7⁺ and CD31⁺ CD4⁺CD45RA⁺ naive T cells decrease, while effector memory CD4⁺ T cells (CD45RA⁻CCR7⁻) and central memory CD4⁺ T cells (CD45RA⁻CCR7⁺) increase^[36]. A human memory T cell subset with stem cell-like properties (CD4⁺CD45RA⁺CCR7⁺ CD28⁺CD27⁺FAS⁺), which has been shown recently^[37] increase relatively. Wijk's group has shown recently that after more than 10 years of thymectomy most children have thymic regeneration and show almost the same number of CD31⁺ naïve T cells compared to healthy control. Only small numbers of children have no thymic regeneration with lower T cell count and low percentage of naïve T cells compared to thymectonized CD31⁺ children^[38]. Another report, which uses in vivo ²H₂O labeling, has shown that compensatory increase of peripheral T- cell division rates are not required to compensate the age-related decrease of thymic output^[34]. From these findings, Goronzy *et* al^[39] summarized that even in early adulthood, thymic output contributes little to the maintenance of the size of the naïve T cell compartment. In order to build T cell repertoire, thymic output is necessary only in early life. To maintain T cell repertoire, thymic output is not necessary. Homeostatic proliferation of the existing T cell pool is enough to maintain CD4 T cell repertoire. Both IL-2 and IL-7 drive homeostatic proliferation and IL-2 receptor (CD25) expression is increased by aging. In contrast to CD4 naïve T cells, the CD8 naïve T cell compartment shrinks with age, which induces higher homeostatic proliferation in aged-CD8 naïve T cells than that in aged-CD4 naïve T cells^[39]. Another cause of T cell dysfunction of aged people instead of decreased production of naïve T cells is the decline in TCR repertoire diversity, caused by cytomegalovirus-driven T cell clonal

expansions^[40]. The naïve T cell repertoire are in the range of approximately 2.5×10^7 in humans. These repertoires are hijacked by CMV infection^[41]. Virus-specific clonally expanded CD8 T cells occupy more than 50% of memory pool and reached to 90% in aged mice^[42]. Naïve T cells are thought to be the precursor of memory T cells. By infection memory T cells respond quickly by producing cytokines, granzyme and perforin. Recently, new type of CD8 T cells have been discovered called "memory T cells with a naive phenotype" (TMNP cells), increased with age. TMNP cells have Naïve T cell phenotypes, but quickly respond to infection by producing cytokines, granzyme and perforin. TMNP cells have unique V 14 T cell receptor. TMNP cells bound to multimers of CMV or EBV peptide and MHC class I^[43]. These results mean that some population of CD8 T cells, which are expanded by CMV, work as effective protection of other virus infection. Further works are needed to elucidate the effects on age-related T cell dysfunction caused by CMV infection. Here we do not describe B cell dysfunction of aged people. B cell function decreases directly by B cell aging itself and by the result of age-related T cell functional decline, because helper T cell affect B cells by direct binding and secreting cytokines.

Myeloid cell aging

Innate immune cells include granulocytes (neutrophils, eosinophils, basophils), macrophages, dendritic cells, natural killer cells and innate lymphoid cells and others^[44]. Among them "myeloid cell" here include granulocytes, macrophages and dendritic cells. We review recent findings, which relate to myeloid cell intrinsic aging. Phagocytosis is the most important effector function of myeloid cells especially neutrophils. Neutrophils play a crucial role in the immune defense against bacterial and fungal pathogens^[45]. Neutrophils survive for only 8-12 h in the circulation and up to 1-2 d in tissues^[46]. Neutrophils from aged human have impaired phagocytosis of bacteria^[46,47] DHEA (Dehydroepiandrosterone sulfate), which increases the phosphorylation of p47phox via PKC-β and generate neutrophil superoxide to kill bacteria, decreases in old age^[48]. Neutrophils also work as an important player in wound healing. Neutrophils normally begin arriving at the wound site within minutes of injury, continuing for several days. Molecules called damage associated molecular patterns (DAMPs) are produced from wounded tissue cells, which activate neutrophils. Neutrophils engulf damaged cells and produce inflammatory cytokines and chemokines, which facilitate wound repair. We have shown that in aged C57BL/6 mice, wound healing is relatively inefficient. Anti-Gri-1 treatment delayed wound healing in aged mice, whereas G-CSF injection increased wound repair^[49]. In bacterial infection and wound healing, macrophages come to work after neutrophil and engulf dead cells including dead neutrophils.

Every tissue contains tissue macrophages, which are sometimes called tissue specific names such as microglia in brain, Langerhans cells in skin and kupper cells in liver.



Important functions of tissue macrophages include innate defense against infection and wound healing. Origin and turnover of tissue macrophages depend on tissue types. Adult's microglia derive from primitive myeloid progenitors of yolk sac that arise before embryonic day 8^[50]. Further non-parenchymal macrophages such as perivascular, meningeal and choroid plexus macrophages, which mediate immune responses at brain boundaries, also arise from yolk sac precursors during embryonic development and remain a stable population^[51]. Langerhans cells predominantly derived from embryonic fetal liver with a minor contribution of yolk sac^[52,53]. Tissue-resident macrophages with Low turn over are self-maintain locally throughout adult life with minimal contribution from circulating monocytes^[54] and self renew after birth^[55]. In heart cardiac macrophages are primarily embryonic and are self-renewed, which reside throughout myocardium^[56]. Recent data have shown that by advanced age these embryo-derived tissue macrophages are replaced by blood monocytes derived from HSCs. In the heart embryo-derived cardiac macrophages decline self-renewal by age and are progressively substituted by monocytederived macrophages^[57]. Intestinal macrophages have a high turnover rate and are constantly replaced from bone marrow derived Ly6C⁺blood monocytes from early lifetime^[58]. In peritoneal cavity, embryo-derived macrophages retain self-renewal at least until 4 mo in mice. By aging embryo-derived macrophages in peritoneal cavity are replaced by bone marrow-derived monocytes^[59].

There exist two types of macrophages namely M1 macrophages and M2 macrophages. M1 (inflammatory) macrophages produce NO radical form arginine with IFN γ or LPS stimulation, whereas M2 macrophages produce TGF β 1 and inhibit NO synthase and stimulate arginase. Arginase converts arginine to ornithine, which induces cell division^[60]. M2 macrophages are further subdivided into M2a and M2c. M2a is activated by IL-4/IL13, and relates to wound-healing process^[61]. M2c is activated by IL10, which has anti-inflammatory activity^[62]. It has been shown that macrophages of aged mice polarize to M2 macrophages^[11,63]. M2 macrophages promote angiogenesis in age-related eye disease^[11]. M2a macrophages have been shown to induce muscle aging by inducing fibrosis. Transplantation of young BM cells to aged mice shows fewer M2a macrophages and less accumulation of collagen^[64]. These findings of age-related M2 predominance contradicts so called "inflammaging", which predominates pro-inflammatory state.

AGE-RELATED TISSUE CHANGES, WHICH INDUCE IMMUNE CELL ATTACK

Although aged HSCs have reduced regenerative potential, aged HSCs are biased toward myeloid differentiation at the expense of lymphoid differentiation^[65]. Chronic, low-grade, systemic inflammation is the primary risk factor for major human chronic diseases, including

cardiovascular disorders, cancer, type 2 diabetes, and neurodegenerative disease. Increased production of inflammatory mediators that accompany this process, referred to as "inflammaging"^[66]. This inflammatory tendency in aged human has wide variability. The mechanism of inflammaging has still not reach complete understanding.

Damaged macromolecules and self-debris released as a consequence of cell/organelle injury work as DAMPs, which may activate pro-inflammatory macrophages. These DAMPs include free radicals from oxidative stress and advanced glycation end products, ATP, fatty acids and amyloid. Another sources, which are up regulated by aging, are leaked microbial products from gut working as pathogen associated molecular patterns (PAMPs)^[65]. These DAMPs and PAMPs may work as stimulator to M1 macrophages. These inflammaging and age-related M2 macrophage activation described previously seem to contradictive. How to consider this discrepancy? By acute infection or injury, neutrophils and M1 macrophages first predominate to the infected or injured site, then M2 macrophages resolve inflammation and induce tissue repair. In chronic inflammation, M1 macrophages continue to exist in inflamed tissue with late coming M2 macrophages. We speculate that both tissue fibrosis caused by M2 macrophages and pro-inflammatory status caused by M1 macrophages continue to exist in chronic inflammation. In acute infection or acute tissue damage, M2 macrophages compensate inflammation caused by M1 macrophages and induce tissue repair. However, in persistent infection such as hepatitis C or repeated tissue damage such as smoking, M1 macrophages, which work as inflammation and M2 macrophages, which resolve tissue damages by producing matrix materials, exist at the same time. Predominance of either M1 or M2 macrophages may dependent on disease process.

Metabolic syndrome and macrophages

Metabolic syndrome, which has the risk to proceed to Type 2 diabetes and atherosclerosis, is caused by obesity. Now it is established that visceral and subcutaneous white adipose tissues (WAT) are different concerning metabolic activity. Lipolytic effect of catecholamines is lower in visceral WAT than that in subcutaneous WAT^[67]. Two main causes of insulin resistance (which causes type 2 diabetes) are saturated fatty acids and pro-inflammatory cytokines. Saturated fatty acids can bind to the Toll-like receptors (TLR) 2 and 4, especially in adipocytes and macrophages, which induce c-Jun N-terminal kinase (JNK), phosphorylate IRS1 and decrease insulin receptor signaling. Saturated fatty acids also produce pro-inflammatory cytokines^[68,69]. The expansion of WAT leads to necrosis of hypertrophic adipocytes, which produce DAMPs. DAMPs recruit macrophages and produce pro-inflammatory cytokines, which aggravate insulin resistance^[70]. Atherosclerosis is mediated by the recruitment of monocyte-derived macrophages into the subendothelial space, where they ingest the lipoproteins^[71]. Type 2 diabetes proceeds to

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many other age-related diseases such as renal failure. Atherosclerosis also proceeds to stroke, coronary heart disease. Thus macrophages especially M1 macrophages are engaged in early stage of age-related diseases.

Clearance of senescent cells by immune cells

Not only M1 macrophages but also aged-other tissue cells produce pro-inflammatory cytokines. Vascular aging enhanced secretion of inflammatory mediators from aging endothelial cells and vascular smooth muscle cells from disease free animals^[72]. One explanation of this age-related secretion of inflammatory mediators is senescence associated secretory phenotype (SASP). SASP is produced by senescent cells. From the studies using mice and aged human, aged innate immune system show elevation of basal inflammation^[73]. Especially older human have elevated level of pro-inflammatory cytokines^[74,75]. However, we should keep in mind that cellular senescence is an independent phenomenon to in vivo aging. Cellular senescence triggers tissue remodeling in embryonic tissue development and in tissue damage. Cellular senescence also engages in age associated loss of tissue functions^[76,77]. Cellular senescence has been first described by Heyflick at 1961 as limited replicative potential of human fibroblasts in vitro^[78]. Cellular senescence is cell cycle arrest with or without telomere shortening after cell division. There exist p53/ p21^{Waf1} pathways and p16^{INK4a}/pRB pathways in cellular senescence^[79]. Whether senescent cells contribute to age-related diseases remains unclear. Some reports have shown that senescent cells have a protective role in atherosclerosis. Mice lacking senescent markers such as p53, p21 or p19Arf show accelerated atherosclerosis^[76]. Senescent cells produce SASPs, which attract immune cells and eliminate senescent cells. Senescent cells attract not only granulocytes and macrophages but also NK cells, CD8 T cells and CD4⁺ T helper 1 (Th1 CD4 T) cells^[80,81]</sup>. However recent report showed that senescent foamy macrophages produced inflammatory cytokines (TNFa), monocyte recruiting chemokines (MCP1) and matrix proteases, which engaged in atherosclerosis^[82]. In this case, macrophages themselves became senescent.

CONCLUSION

From the demand to solve worldwide increase of agerelated diseases, the future biochemical studies are needed, which elucidate the phenomena of immunosenescence and immune cell attack to eliminate senescent cells or damaged cells by age-related stresses. In advanced age, diseases are caused by M1 or M2 macrophages as mentioned above. Which type of macrophages are involved depend on the situation. In order to prevent or treat age-related diseases we have to carefully evaluate the immune status. Because metabolic reprogramming of oxidative phosphorylation to glycolysis occurs in M1 pro-inflammatory macrophages and urea cycle intermediates such as arginine, ornithine, citrulline regulate M1/M2 polarization^[60], the measurement of metabolic changes will elucidate the predominant type of macrophages. Studies of metabolic pathway of immune cells in aging will open the nutrient treatment of agerelated diseases.

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