

Aging and Its Molecular Mechanism

Subjects: **Cell Biology**

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Aging is a continuous process over time that is mainly related to natural alterations in mechanical–biological processes. This phenomenon is due to several factors, including the time and energy of biological processes. Aging can be attributed to biological factors such as oxidative stress, cell longevity, and stem cell senescence.

aging

senescence

1. Introduction

Higher organisms are organized into organs and tissues with complex interconnections. During the life of a higher organism, the loss of functional and structural cells is directly compensated for by stem cells. However, over time, the ability of these stem cells to generate newly differentiated cells weakens for two reasons: the depletion of stem cells and the reduction in the energy necessary for the survival of these cells (ΔG), which is lost in the form of ΔS [1][2][3][4]. This mechanical–biological process results in aging, which is essentially associated with the senescence of the cells constituting the organism (their inability to divide and the inability of cells to be renewed). Thus, this process is becoming more important with the increase in life expectancy leading to the appearance of a certain number of so-called age-related diseases. Several recent works have deciphered the molecular nature of certain pathologies. They have demonstrated a close relationship between the aging process and several human pathologies, particularly cancer, Alzheimer's, and Parkinson's diseases [5][6][7].

2. Aging and Its Molecular Mechanism

Human aging, unlike sickness, is a progressive time-related process. It varies from person to person, and corresponds biologically to a loss of homeostasis, an increase in the organism's sensitivity and susceptibility to disease and death, and the progressive degeneration of cells, tissues, and organs associated with advancing age [8]. These aspects of deterioration are called senescence, and are responsible for the weakening of an individual's health. They also cause physiological changes in “regular” aging, such as menopause and decreased kidney function, and age-related disorders, such as coronary heart diseases, in “ordinary” aging. Several variables contribute to the loss of homeostasis, which is ultimately the consequence of a genetic program; some models propose that genes function to increase or decrease the relative risk of death by increasing the likelihood of disease [8]. Changes in the crystal structure or macromolecular aggregation at the molecular level, the loss and shortening of telomeres at the chromosomal level, changes in mitochondria, and the accumulation of lipofuscins inducing cellular aging are all monitored as part of the aging process [8], along with the appearance of cross-linking

lesions of collagen and elastic fibers, and amyloid deposition. This usually causes a change in an organism's appearance, function, and behavior [8][9].

The human average life duration has risen substantially over time; the most remarkable possible lifespan has remained stable, ranging from 90 to 100 years, and differs from person to person. The average human life expectancy has increased in recent years due to changes and developments in disease management and the socio-economic status and nutritional status of individuals, as well as fewer accidents; the improvement in all these factors has contributed to an increase in the average human life expectancy [10]. At the very least, increased mortality after maturation [11][12][13], changes in the biochemical composition of tissues [14], progressive declines in physiological capacity [15][16][17], and reduced abilities to respond adaptively to environmental stimuli are among the numerous patterns associated with aging in mammals. However, these are as yet unclear and unverified, because the mechanisms of aging could be highly different between animals, tissues, and cells.

Aging mechanisms in humans differ significantly from those of one organism, tissue, or cell, making it impossible to establish a specific mechanism. On the other hand, other academics have focused on understanding the evolutionary foundation of senescence in the aging mechanism so as to understand the aging process and study the various genes involved in senescence, which can be divided into three groups; namely, genes regulating somatic maintenance and repair, genes promoting early survival, and genes causing late deleterious mutations [18][19]. These essential genes have been shown to impact the evolution and survival of a species, and may play a role in reducing longevity due to higher growth and repeatability. The published research in molecular genetics has revealed that cellular senescence could have an opposite pleiotropic effect, preventing cancer while simultaneously contributing to organism aging. **Figure 1** describes the molecular mechanisms involved in the aging process.

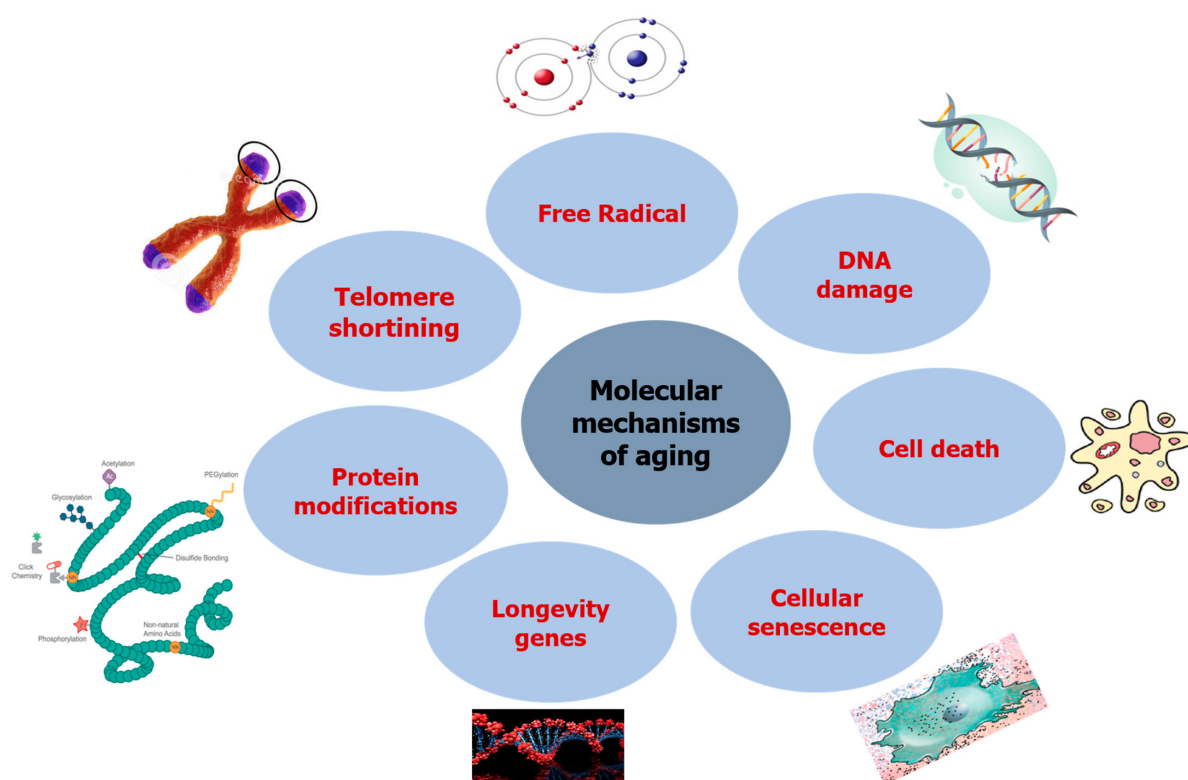


Figure 1. Molecular mechanisms inducing aging.

2.1. DNA Damage

The molecular mechanisms of aging are still unknown. According to researchers, aging is simply a physiological decline caused by the accumulation of random damage to vital molecules in aging populations. It is an example of background radiation causing genetic damage that leads to mutations, resulting in functional impairment and death [20][21]. The primary example of this theory is the ability to repair DNA damage caused by intrinsic sources such as replication defects and chemical modifications of DNA [22][23], or by external sources, including UV and genotoxic drugs [22][23] in species with different lifespans [24]. Mutations, transcription, replication halts, and the DNA damage response are triggered when DNA is damaged (DDR). These DDRs block cell cycle progression and activate signaling pathways affecting the cell via repair, apoptosis, or cellular senescence [25]. In addition, the presence of defects in genes involved in the DNA repair system causes the accumulation of unrepaired DNA and chromosomal damage [26].

On the other hand, exonuclease 1 (EXO1) and postmeiotic segregation increased 2 (PMS2) are two components that play a vital role in the DNA repair system. Their function contributes to human longevity [27][28]. Indeed, various investigations in human patients and mouse models have highlighted the relevance of DNA repair in the aging process [29]. Thus, it is not the DNA damage that causes aging; instead, it is about how the cell reacts to DNA damage and how this response will affect the organism's life. In this respect, any abnormality or mutation in DNA repair pathways—such as Werner's syndrome caused by a mutation in Werner's syndrome protein (WRN), a gene that encodes a RecQ DNA helicase essential for replication stress management and telomere stability [25][30]—causes rapid aging and shorter lifespan [31]. Furthermore, researchers discovered several clinical disorders caused by problems with the genome maintenance system and aging-related syndromes. For example, Cockayne syndrome is affected by transcription-coupled nucleotide excision repair (TC-NER) [9][32]—these DNA damage repairs are generally induced after the exposure of genetic material to UV rays; ataxia-telangiectasia (AT) affected by DNA damage response—an immune deficiency affecting the humoral pathway and manifested by progressive cerebellar ataxia, linked to genetic instability of the genes that code for a protein kinase (coded by the MRE11 gene) controlling the repair of DNA double-strand breaks, especially in cerebellar and endothelial cells; Werner syndrome impacted by telomere maintenance and replication stress [8][33]—an autosomal recessive genetic disease inducing premature aging due to genetic instability, which affects the DNA repair system; Rothmund–Thomson syndrome, affected by DNA replication starting codes [34][35]—a hereditary genetic dermatosis, manifesting in dermatological signs linked to premature aging, which predisposes one to skin cancer.

2.2. Free Radicals

It has been discovered that the majority of age-related alterations are caused by molecular damage induced by free radical [36][37][38] atoms or molecules possessing an unpaired and reactive electron, constituting another possible cause of aging. These oxygen-derived species can react with macromolecules to produce free radicals from the attacked molecules [38][39], and act as secondary messengers in signaling pathways implicated in the

control of various mechanisms, such as changes in gene expression, cell replication, differentiation, and apoptotic cell death [40][41][42]. The generation of these free radicals in human organs such as the heart, kidney, and liver affects maximal lifespan [43][44]. In this context, nutritional antioxidants have been found to reduce the risk of vascular dementia, heart disease, and cancer in humans [45][46]. Reactive oxygen species (ROS), in turn, play a role in the somatic accumulation of mutations in mitochondrial DNA, which is one of the developmental–genetic aspects of aging. These mutations result in a gradual loss of bioenergetic capacity, as well as aging and cell death [47][48][49]—“Mitochondrial aging’s redox mechanism” [50]. As people age, oxygenated free radicals play a role in mitochondrial DNA (mtDNA) damage [51][52][53]. This damage causes inefficient mitochondrial respiration, which increases with age, as well as the onset of age-related illnesses, including Parkinson’s disease [54][55], Alzheimer’s disease [56][57], Huntington’s chorea [58][59], and others. mtDNA is passed down through the generations and replicates. Moreover, it is much more susceptible to mutations than nuclear DNA, and the rate of mutations rises with age [60][61].

2.3. Telomere Shortening

In addition to DNA damage, telomere length is associated with age-related disorders, whereby telomere shortening can impair somatic stem cell function. Findings indicate that telomerase-deficient mice have short telomeres and age prematurely, whereas cancer-resistant mice with high telomerase expression have long telomeres and age more slowly [62][63]. The phenomenon of telomere shortening also remains a feature and a counting mechanism of senescent cells [64][65]. Telomeres consist of long stretches of TTAGGG repeats located at the ends of chromosomes, and act as protectors that prevent them from degrading or fusing with other chromosome ends [51]. Telomerase expression is limited in human somatic cells, leading to telomere reduction and replicative senescence [52]. In fibroblasts and peripheral blood lymphocytes, the average length of the terminal restriction fragment of chromosomes decreases with age [66][67][68]. These findings show that telomere length, rather than telomerase activity, is the most critical determinant in cellular aging. Furthermore, the shortest telomere affects cell viability and chromosome stability, rather than the average-length telomeres. In this respect, several studies have demonstrated that in aged animals and humans, telomeres shorten over time [62][69]. Shorter telomeres are associated with an increased risk of death [70] and replicative senescence; they prevent cancer cells from dividing indefinitely [71]. In contrast, non-enzymatic telomere elongation extends cell lifetime in the laboratory [72].

2.4. Protein Modifications

Proteins are the building blocks of living creatures’ cellular and physiological functions, and their physical and chemical qualities determine their activities and functions. Protein folding and final conformation, and biochemical activity, stability, and half-life are all affected by the primary sequence [73]. Researchers have shown that protein repair and modification might play a role in longevity in certain situations [74]. The oxidation of amino acid residues, metal-catalyzed oxidation, and change caused by lipid oxidation products reduce the specific activity of numerous enzymes, affect thermal stability, and increase the carbonyl content of proteins [75][76]. Moreover, protein acetylation has also been proposed to play a significant role in the aging process by improving the function of specific genes,

most notably the AMP-activated protein kinase (AMPK) regulatory subunit, which has been linked to increased longevity [77].

2.5. Longevity Genes

Scientists propose that the aging process is primarily caused by a genetically programmed continuum of growth and maturation. The maximum lifespan is very species-specific, as humans have a maximum lifespan 30 times longer than mice. In this context, it was discovered that specific genes found in many animals play a role in determining the full lifetime potential [78][79][80]. Indeed, the existence of these genes results in the synthesis of products that are engaged in the control of the species' life via several mechanisms, including the modulation of stress and resistance, the increase in metabolic capacity, and the silencing of genes that promote aging [81][82][83]. In this respect, numerous investigations have suggested that the overexpression of the SIR2 gene and its homolog increases the lifespan of yeasts and nematodes [84].

Sir2 has been linked to reduced histone acetylation at the amino group of N-terminal lysine residues and global hypoacetylation in yeast [85]. Sirtuins are thought to play an essential role in cell response to several stimuli, including oxidative and genotoxic stress, and are necessary for cell metabolism [86]. Many studies have questioned the direct involvement of sirtuins in extending human lifespan, and their intervention in many human body systems such as the liver and cardiovascular system. It has been suggested that the main activity of sirtuins is the deacetylation of lysine residues [87][88]. Sirtuins cleave nicotinamide adenine dinucleotide (NAD) to nicotinamide. Then, an acetyl/acyl group is transferred from the substrate to the ADP-ribose moiety of NAD, resulting in 2'-O-acetyl-ADP-ribose and a deacetylated substrate [89].

Other researchers suggest that mutations in specific genes, such as the daf-16 gene implicated in various signal transduction pathways, including insulin signaling [84][90][91], may cause greater longevity in mutants [92]. In contrast, genetic research on mammalian longevity has revealed the presence of immunological loci in mice and humans, with these loci having implications for longevity [71]. The role of genetics in longevity was highlighted after scientists discovered that siblings and parents of long-lived people also live longer, and researchers found the presence of multiple genes on chromosome 4 linked to exceptional longevity [93][94][95], most of which are pleiotropy genes. However, since age causes differences in gene expression in muscles and the brain, several studies have shown that caloric restriction prevents age-related gene expression changes in mice. Furthermore, several investigations have focused on the cellular pathway by suggesting that aging is a cellular model, and that an individual capacity is relatively proportional to the functional capacity of the cells [96][97].

Intracellular enzymes such as collagenases, elastases, and tyrosinase are increased by intrinsic aging and photoaging factors, resulting in skin aging [98]. Extrinsic aging is generated by external stimuli, such as chronic exposure to pollutants or UV rays. At the same time, it is believed that internal aging is controlled and established by several hereditary genes [99][100]. Collagen and elastase in the dermis denature as a result of persistent UV exposure and other external factors, leading to wrinkles and the photoaging of the skin; by stimulating intracellular signal transcription pathways such as p38 mitogen-activated protein kinase and c-Jun-N-terminal kinase, this

mechanism will induce the creation of MMPs, which can arise from extracellular matrix (ECM) degradation [101][102][103]. Numerous proteins present inside the ECM have been discovered. Elastase is a member of the chymotrypsin family of proteases that is primarily responsible for the degradation of elastin and collagen, essential biomolecules protecting the skin against damages [104]. Several studies have emphasized the critical role of disposing foreign proteins within the ECM during neutrophil phagocytosis and tissue healing in standard settings [98]. Many research works have focused on the direct impact of inhibiting those enzymes. It has been found that inhibiting elastase and MMP-1 enzymes may positively affect skin aging due to their usefulness in avoiding skin sagging and losses of elasticity [98][105]. Furthermore, scientists have investigated the impact of inhibiting tyrosinase, a copper-containing monooxygenase that catalyzes the O-hydroxylation of tyrosine to 3,4-dihydroxyphenylalanine and then to dopaquinone, which is synthesized by epithelial, mucosal, retinal, and ciliary body melanocytes, and which is deeply involved in the protection of the skin from melanogenesis [106][107].

2.6. Cellular Senescence

Cellular senescence and the number of divisions are required to determine proliferative lifespan [1][108]. Several studies have shown that a decrease in the rate of cell proliferation corresponds to aging in animals [19][109]. Multiple intrinsic and extrinsic factors, including oxidative stress (OS), DNA damage, oncogene activation, epigenetic stress, and mitotic spindle stress, can cause cellular senescence. When stem cells or progenitor cells are damaged, cellular senescence compromises tissue function and reduces tissue regeneration capacity [110]. In this context, several studies on the senescence of many cells and their relationship with aging, such as glial cells [111], keratinocytes [112], vascular smooth muscle cells [113], lens cells [114], endothelial cells [115], and lymphocytes [116], have been conducted. The results indicate that eliminating senescent cells in adult wild-type mice delays tumor formation; thus, eradicating senescence could help people live longer [117]. In addition, normal cells in life forms have a limited proliferative ability; cells divide less frequently in humans, and their latency before proliferation increases. Despite these investigations, there is no conclusive evidence that senescent cells grow in vivo as people age.

2.7. Cell Death

Traditionally, cell death methods have been divided into energy-dependent programmed cell death apoptosis mechanisms and necrotic cell death mechanisms. The body can preserve equilibrium in the apoptosis mechanism by committing active "suicide". This is a preprogrammed death caused by the progressive activation of a group of cysteine proteases known as caspases. Most cells have caspases in their cytoplasm, but they are generally inactive (procaspase) [118]. Because there is no release beyond the cell, there is no inflammation. This occurs in response to environmental stimuli and is controlled by genes [119][120]. In addition to its essential role in the immune system, where up to 95% of T cells undergo cell death (presumably because they recognize self-antigens) [121], it involves the compaction and segregation of chromatin adjacent to the nuclear membrane and the condensation of the cytoplasm, eventually evolving into nuclear/cellular fragmentation. Meanwhile, apoptosis may play a role in aging and age-related disorders. Apoptosis, followed by the replacement by division of another cell, may occur if cells are unable to repair DNA damage; it is also essential for wound healing, which is often reduced as people

age, often in conjunction with local inflammation ^[122]. The central nervous system shows the most transparent relationship between apoptosis and aging, whereby neuronal apoptosis increases with aging. Similarly, cancer rates rise with age due to decreased apoptotic defenses ^[123].

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