Cellular Senescence and Stem Cell Aging

Cellular senescence is the cessation of the process of division of the cells. Aging is an insidious definite decline in a cell’s functionality, vigor, and vitality. Recent advances in biomedical research, cytogenetic and molecular studies have brought forth fascinating theories to expatriate aging and, more importantly, to tame aging to modulation by genetic and pharmacological manipulations. This article focuses on the effect of cellular senescence and stem cell exhaustion on aging and the subsequent clinical implications.

Introduction to Cellular Senescence and Stem Cell Aging

Aging is defined as a diminished response to stress, escalation of homeostatic imbalance and the enhanced threat from aging-related pathologies. With the advances in biomedical research, a number of theories proposed to expound aging have emerged and many are on their way. The most likely cause of aging is a multi-targeted attack on the normal functioning of the cell leading to disruption of the milieu interior of the cell and consequent detrimental effects which finally culminate in a diminution of a cell’s lifespan.

There is a multitude of diverse physiological traits described which characterize aging. The same can be tabulated as follows for easy memorization:

- Trait
- Stem cell exhaustion
Mitochondrial dysfunction
Cellular senescence
Epigenetic alterations
Loss of proteostasis
Altered intercellular communication
Telomere attrition
Deregulated nutrient sensing
Genomic instability

We now focus on cellular senescence and stem cell exhaustion and their role in aging.

Cellular Senescence

Overview

Cellular senescence implies irrevocable arrest of dividing cells. These cells possess grossly dysregulated phenotype, altered genotype, morphology and behavior distinct from their growth competent counterparts.

Note: The stable arrest of the cell cycle and associated phenotypic changes are intended to prevent the propagation of damaged cells and trigger their clearance by the immune system. Insidious aggregation of these cells in normal tissues induces aging and age-related diseases. They are responsible for the development of incapacitated abnormal tissue function. Cellular senescence is an antagonistic hallmark of aging, a response to damage.

Different pathways

Telomere attrition, Oxidative stress, and Oncogene activation are the different pathways which can induce cellular senescence.

Repetitive senescence

Another important terminology closely related to cellular senescence is “Repetitive senescence”. Telomere shortening occurs every time a cell divides. Once the telomere length reaches a critical threshold, it can no longer be protected by telomeric proteins.

This culminates in the exposure of DNA ends, subsequent DNA damage and ultimately
cellular senescence. Thus, cellular senescence secondary to telomere attrition and DNA damage as a result of multiple cell divisions is known as “Replicative senescence”.

**The key mechanisms** by which senescent cells contribute to the diminution of the lifespan of a cell are enlisted as follows:

- Alteration of phenotypic behavior of surrounding cells
- Cellular dysfunction
- Reduction of growth-competent mitotic cells conglomerations
- Dysregulation of extracellular matrix and similar significant structural elements of the cell
- A propensity towards cancer formation

**The biological impact** of senescent cells at the molecular level is multi-pronged and can be summarized as follows:

- Up-regulation of growth factors
- Escalated levels of extracellular matrix (ECM)-degrading proteins
- Increased pro-inflammatory cytokines
- Up-regulation of matrix metalloproteinases (MMPs)
- Increased IL6, TNF α, Interleukin-8
- Abnormal secretion of growth factors
- Altered transcriptional profile

Cellular senescence has been incriminated in a number of disorders. The principal diseases associated with it are intervertebral disc degeneration, chronic hepatitis C, hepatocellular carcinoma, benign prostatic hyperplasia, type-II diabetes mellitus, a glomerular disease of the kidney and emphysema.

**Clinical implications of Cellular Senescence and Stem Cell Aging**

The concept of cellular senescence has deep far-reaching clinical consequences. Active, vigorous research concentrates on **3 key areas of cellular senescence, namely; prevention, removal, and replacement of senescent cells** in tissues. Various therapies are taking shape in experimental setups already.

**Telomerase therapy**

The notion is to develop strategies to **transiently turn on the telomerase activity in cells**. This will limit telomere attrition and thus tame replicative senescence. However; not all cells are subject to this ideology. There exists a specific sub-population of cells which age independent of telomere shortening. Some are directly afflicted by DNA damage. These cells are unaffected by telomerase therapy.

**Apoptosis-inducing pharmacotherapy**

Cellular senescence is modulated by a fine balance maintained between the production of young cells from stem cells and removal of local damaged cells by scavenging and the immune system. The inspiration comes from the development of cell-surface marker specific anti-cancer drugs.

**Note:** Scientists working tirelessly in this direction hope to generate drugs which would
recognize senescent specific cell-surface markers and induce apoptosis in these cells.

**Immune system modulation**

Rejuvenation therapy believes that **accumulation of damaged senescent cells in tissues is predominantly a culmination of dysregulated and the aged immune system** itself. There are various junctures in the immune system which, if reinstated, could curb the processes which accelerate aging.

The various important steps in the immune system affected by aging are:

1. Regulation of dendritic cell function
2. Recognition of proper target cell
3. Display of accurate target cell-specific marker
4. Responsive secondary activation of lymphocytes

**Stem cell therapy**

The deleterious effects of senescent cells are compounded by defective clearance of these cells associated with the diminished production of young healthy cells by a local niche of stem cells. Aging might itself decrease the vigor of the local stem cells. In these circumstances, stem cell rejuvenation therapy by the addition of stem cell populations into the tissues after removal of senescent cells is an interesting notion.

**Stem Cell Exhaustion**

Adult stem cells preservation is crucial for the maintenance of tissue homeostasis; especially as the cell grows older. Aging is associated with the attrition of stem cells. This is known as stem cell exhaustion. Exhaustion of the stem cell pool is **brought in by a multitude of diverse processes such as dysregulated metabolic signaling, premature aging secondary to oncogenic insult to the somatic genome** and other pathways which contribute to the decline in the functionality of the cell.

Adult stem cells regulate the key signaling pathways responsible for the emergence of age-related pathologies. Stem cell exhaustion leads to anemia, myelodysplasia, osteoporosis and decreased fracture repair, decreased repair of muscle fibers, decreased intestinal function and cancer. Highly advanced therapeutic manipulations of a stem cells niche to affect healthspan and lifespan of an individual is not a distant dream now.

Stem cell exhaustion is an integrative hallmark of aging, the culprit of the phenotype. It is an integrated consequence of so many others of these hallmarks. Age-related genomic instability and DNA damage are proposed to result in altered differentiation of stem cells which fail to provide optimum replacement of healthy cells and hence ultimately lead to decreased longevity of life.

**Stem cells and the immune system are in the positive feedback loop when it comes to accelerating the aging phenotype.** The genomic and transcriptomic profile of stem cells also undergoes changes as stem cell exhaustion occurs.

**Stem cell and immune system**

Stem cell exhaustion and dysregulated aged immune system are in a vicious cycle with positive feedback mechanism linking the two. Thus, while the altered function of stem cells leads to the further dysfunctional immune system, the defunct immune system
creates a pro-inflammatory aging milieu which further hampers the functionality of the stem cells. This vicious cycle is thought to culminate in accelerated senescent phenotype.

**Altered gene expression profile in stem cell exhaustion**

Network analysis studies have revealed differential regulation of the different set of genes in adult stem cells as they are subjected to exhaustion and subsequently cellular senescence. Studies on aging from the cytogenetic perspective reveal a fine balance between proto-oncogenes and tumor suppressor genes, each pushing the growth curve in opposite directions. Aging can be looked upon as a disruption of this fine balance regulating alternative cellular states.

The key genes involved are as follows:

<table>
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<tr>
<th>Proto-oncogenes:</th>
<th>BMI 1</th>
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<tr>
<td>Wnt/β-catenin</td>
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</table>

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<tr>
<th>Tumor suppressor genes:</th>
<th>INK4</th>
<th>ARF</th>
<th>AIMP3/p18</th>
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</table>

The principal changes which occur at the molecular level to bring about this altered genomic and transcriptomic profiling can be summarized as follows:

- Up-regulation of glycan metabolism
- Slower DNA unwinding
- Dysfunctional regulators of senescence
- Perturbation of germline genetic heterogeneity
- Increased collagen cross-linking
- Capillary basement membrane thickening
- Overexpression of pro-inflammatory genes and inflammatory adipocytokines; especially in females
- Aberrant insulin-IGF1, FoxO and mTOR signaling, glucose and fatty acid metabolism
- Overexpression of pro-inflammatory genes; especially in females
- Decreased Na+K+ ATPase activity

**Stem cells and Metabolic stress**

Adult stem cells are victimized by a number of metabolic stressful insults. These stressors challenge the reparative and regenerative potential of the stem cells. Stem cell exhaustion secondary to these stressful circumstances results in aging. The key metabolic stressors can be summarized as follows:

- Oxidative stress
- Nutrient deprivation
- Abnormal proteome; especially involving proteins in cytoskeletal organization and anti-oxidant defense armamentarium
- Altered protein infrastructure dealing with the redox status of the cell

The critical cellular pathways implicated in stem cell exhaustion and aging can be summarized as follows:

**Explanation of Wnt signaling pathway**

The canonical Wnt signaling pathway is crucial in the fate determination of stem cells.
Stem cell exhaustion is regulated in a disciplined manner by Wnt pathway through **extensive crosstalk with multiple other key regulatory pathways of the cell** such as the SMAD signaling pathway; BMP, NOTCH, and prostaglandin E2 pathways.

The following are modulated by the Wnt signaling pathway:

- Regulation of stem cell proliferation
- Self-renewal
- Differentiation
- Embryonic patterning
- Stem cell maintenance
- Stem cell senescence

The Klotho mouse model of aging points to aberrant Wnt signaling as the culprit pathway in stem cell exhaustion and premature senescence of stem cells with decreased self-renewal and early senescent phenotype. There is evidence to correlate gradual accumulation of Progerin and diminishing Klotho levels with human aging.

**Explanation of FoxO family**

The FoxO transcription factor signaling pathway **regulates the cellular response to oxidative stress**. It coordinates the composite response of a cell to tumor suppression, metabolism, and longevity of lifespan of a cell. It regulates multiple intersecting cellular pathways and affects a multitude of key fundamental cell functions, such as the following:

- Stress resistance
- Gluconeogenesis
- Cell cycle arrest
- Autophagy
- Apoptosis
- Neuropeptide secretion
- Atrophy

Bestowed with ubiquitous expression, FoxO functions are still **highly specialized and customized uniquely to distinct cell types**. Redirection of β-catenin function as a result of oxidative stress-mediated through FoxO family of transcription factors culminates in the altered transcriptomic profile of stem cells and affects cellular homeostasis and cellular senescence. FoxO transcription factors also play a role in stem cell exhaustion by altering the structural and functional integrity of the aging immune system.

**Clinical implications**

Extensive understanding of the role of stem cell exhaustion in aging has triggered a lot of fascination and vigor in interventional regenerative medicine. Engineering cells and tissues for replacement and regeneration of diseased and injured cellular entities, such as in neuromuscular systems, can confer resistance to aging.

**Ex vivo aging of stem cells in rodent cells is an upcoming area of research. Partial reprogramming involves moving adult stem cells from limited multi-potency towards pluripotency** by altering the transcriptional phenotype of stem cells. This approach can be used to preserve the proliferative potential and regenerative capacity of the stem cells; thus evading exhaustion.
Summary of Cellular Senescence and Stem Cell Aging

Aging is defined as the diminished response to stress, escalation of homeostatic imbalance and the enhanced threat from aging-related pathologies. There is a multitude of diverse physiological traits described which characterize aging.

Cellular senescence implies irrevocable arrest of dividing cells. These cells possess grossly dysregulated phenotype, altered genotype, morphology and behavior distinct from their growth competent counterparts. Cellular senescence is an antagonistic hallmark of aging; a response to damage.

There are different pathways which can induce cellular senescence. Cellular senescence secondary to telomere attrition and DNA damage as a result of multiple cell divisions is known as “Replicative senescence”. Cellular senescence has been incriminated in a number of disorders.

The concept of cellular senescence has deep far-reaching clinical consequences. It opens avenues for therapeutic manipulations such as telomerase therapy, apoptosis-inducing pharmacotherapy, stem cell therapy, and immune system modulation. Adult stem cells preservation is crucial for the maintenance of tissue homeostasis; especially as the cell grows older. Aging is associated with the attrition of stem cells. This is known as stem cell exhaustion.

Stem cell exhaustion leads to anemia, myelodysplasia, osteoporosis and decreased fracture repair, decreased repair of muscle fibers, decreased intestinal function and cancer.

Stem cell exhaustion is an integrative hallmark of aging. It is influenced by multiple factors, such as oxidative stress, metabolic stress, dysregulated immune system, altered genomic and transcriptomic profiling, disrupted balance between proto-oncogenes and tumor suppressor genes. Wnt signaling pathway and FoxO transcription factors are key regulators of stem cell exhaustion.

Ex vivo aging of stem cells, partial reprogramming, and tissue engineering are potential areas of research in relation to stem cell exhaustion.

A summary of the hallmarks of cellular aging

<table>
<thead>
<tr>
<th>Primary hallmarks (Causes of damage)</th>
<th>Antagonistic hallmarks (Response to damage)</th>
<th>Integrative hallmarks (Culprits of the phenotype)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Genomic instability</td>
<td>Deregulated nutrient sensing</td>
<td>Stem cell exhausting</td>
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<tr>
<td>Telomere attrition</td>
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Review Questions

The correct answers can be found below the references.

1. Which of the following entities is best described by the following statement: “Cellular senescence secondary to telomere attrition and DNA damage as a result of multiple cell divisions.”?
A. Replicative senescence  
B. Reproductive senescence  
C. Regenerative senescence  
D. Reparative senescence

2. Which type of hallmark of aging is cellular senescence?
   
   A. Primary hallmark  
   B. Integrative hallmark  
   C. Antagonistic hallmark  
   D. Inflammatory hallmark

3. Which of the following pathway crucially influences stem cell exhaustion?
   
   A. AIMPK pathway  
   B. Wnt signaling pathway  
   C. Hedgehog signaling pathway  
   D. RAS pathway

References


Correct answers: 1A, 2 C, 3B

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