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Cellular senescence is a physiological phenomenon that has both beneficial and detrimental consequences. Senescence limits tumorigenesis and tissue damage throughout the lifetime. However, at the late stages of life, senescent cells increasingly accumulate in tissues and might also contribute to the development of various age-related pathologies. Recent studies have revealed the molecular pathways that preserve the viability of senescent cells and the ones regulating their immune surveillance. These studies provide essential initial insights for the development of novel therapeutic strategies for targeting senescent cells. At the same time they stress the need to understand the limitations of the existing strategies, their efficacy and safety, and the possible deleterious consequences of senescent cell elimination. Here we discuss the existing strategies for targeting senescent cells and upcoming challenges in translating these strategies into safe and efficient therapies. Successful translation of these strategies could have implications for treating a variety of diseases at old age and could potentially reshape our view of health management during aging.
Strategies targeting cellular senescence

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Introduction

The world’s population is rapidly aging (1, 2). Living to a late age provides many opportunities but also presents a huge challenge, as it increases vulnerability to the development of chronic pathological conditions. In fact, aging is the leading risk factor for the world’s most prevalent pathologies, including cardiovascular diseases, cancer, and neurodegenerative diseases (3). Aging is heterogeneous, and some people function better than others at the same chronological age, exhibiting a longer period of good general health. Thus, a better understanding of common cellular and molecular pathways that drive the development of age-related multimorbidities is necessary. Treatment of age-related diseases based on such pathways could provide better therapies than treatment of each age-related disease individually. Recent discoveries have provided insights into the cellular and molecular events that play a role in biological aging (3, 4). One emerging factor is the accumulation of senescent cells in tissues.

Cellular senescence is an essentially irreversible cell cycle arrest that occurs in normal proliferating cells in response to various forms of cellular stress. Replicative exhaustion, oncogene activation, direct DNA damage, cell-cell fusion, and other forms of stress that elicit activation of the DNA damage response pathway can lead to senescence (5–8). Cellular senescence is a vital physiological response aimed at preventing propagation of damaged cells in the organism (9–11). It acts as a bona fide tumor suppression mechanism, limits tissue damage, and aids wound healing (12–16). Despite the protective role of cellular senescence as a cellular response to stress, studies in mouse models have shown that the long-term presence of senescent cells that form as a result of this response may be detrimental to the organism (17, 18). These cells secrete a plethora of proinflammatory factors that assist in their removal by the immune system (19, 20). Studies on diverse animal models indicate that multiple components of the immune system, including NK cells, T cells, and macrophages, are involved in controlling the presence of senescent cells in tissues (13, 21–25). The efficacy of this removal is variable among tissues and pathological conditions, and the mechanisms and rules regulating the homeostasis of senescent cells are yet to be fully understood.

At the late stages of life, senescent cells increasingly accumulate in tissues and contribute to the establishment of a chronic “sterile” inflammation that arises due to continuous secretion of proinflammatory cytokines (11, 26, 27). This condition, also known as “inflammaging,” is a pervasive feature of the majority of age-related diseases (28). Indeed, senescent cells are especially abundant at sites of age-related pathologies, and a growing body of evidence from mouse models demonstrates a causal role for senescent cells in the pathogenesis of age-related diseases including atherosclerosis, idiopathic lung fibrosis, osteoarthritis, bone loss, and hepatic steatosis (29–34). Furthermore, genetic approaches to promoting clearance of p16-expressing senescent cells in mice delay the onset of age-related deterioration of several organs and increase median survival of the mice (35, 36). Hence, elimination of senescent cells might be a promising approach for treatment and prevention of many age-related diseases, hopefully leading to healthy longevity (37–39).

Therapeutic strategies for targeting of senescent cells

There is growing interest in the possibility of targeting senescent cells therapeutically. Several promising approaches that focus on either clearance of senescent cells or prevention of their proinflammatory impact are in development (Figure 1). Current efforts are largely invested in the discovery of pharmacological agents that can induce cell death in senescent cells. These compounds are often termed “senolytic drugs” or “senolytics.” Research in this direction...
is mainly based on the biological pathways underlying senescent cell accumulation with age, and the therapies aim to utilize some of the unique molecular features that senescent cells display over other cells in the organism. One of the most prominent features of these cells is their relative resistance to apoptosis. Unlike normal cells, senescent cells are protected from both intrinsic and extrinsic proapoptotic signals, a property that allows them to persist and promote diverse biological processes under stress conditions (40–42). Targeting these apoptotic pathways preferentially in senescent cells, although different from a genetic approach in multiple respects (43), leads to selective death of these cells and prevents them from exerting their detrimental effects.

Blocking prosurvival pathways. Several of the prosurvival pathways that have been identified can be used for directed elimination of senescent cells (44–49). These include the BCL-2 protein family, the p53/p21 axis, PI3K/AKT, receptor tyrosine kinases, and the HIF-1α and HSP90 proteins. Currently, most identified senolytics are directed against members of the BCL-2 protein family. Antiapoptotic proteins of the BCL-2 protein family are well studied on the molecular level and are potentially attractive drug targets (50). These proteins play multiple roles in cell death regulation through their effects on apoptosis and autophagy (40, 51). Studies across different cell types have demonstrated an upregulation of the BCL-2 family members BCL-2, BCL-W, and BCL-XL during senescence (46, 48). Silencing those proteins in senescent cells leads to the activation of programmed cell death.
The BCL-2 protein family is an extensively investigated target for pharmacological intervention in cancer, offering several well-characterized candidates as senolytics (50). ABT-737, for example, is a BH3 domain mimetic that blocks the interaction of antiapoptotic family members BCL-2, BCL-W, and BCL-XL with BH3 domain–containing proapoptotic proteins, allowing senescent cells to undergo apoptosis (52, 53). ABT-737 efficiently eliminates senescent cells that were induced by DNA damage in lungs of irradiated mice, as well as senescent cells formed by p16kip1 induction in skin epidermis of transgenic mice (48). This study also suggested that elimination of senescent cells from the epidermis might positively affect hair growth, as it induced proliferation of hair follicle stem cells. ABT-263, also known as navitoclax, is a next-generation, orally available analog of ABT-737 (54). ABT-263 has demonstrated the capacity to eliminate senescent cells from sublethally irradiated mice and naturally aged mice, including senescent muscle stem cells and senescent hematopoietic stem cells (46). This action results in rejuvenation of those stem cell populations. ABT-263 also showed the capacity to eliminate senescent foam cell macrophages from atherosclerotic lesions, thereby halting progression of the disease (29).

Notably, treatment with these general inhibitors of BCL-2, BCL-W, and BCL-XL causes several mechanism-based hematological toxicities, such as neutropenia and thrombocytopenia (55). These safety issues will likely impede the development of BCL-2 family inhibitors into common clinical interventions, and emphasize the need for more specific inhibitors that might serve as superior candidates. These might include specific BCL-XL inhibitors, such as A1331852 and A1155463, which are expected to cause less toxicity to nonsenescent cells (56). A1331852 can induce apoptosis in senescent cholangiocytes and senescent fibroblasts in a genetic mouse model of biliary liver fibrosis, thereby reducing liver injury and fibrosis (57). Nonetheless, platelet survival is dependent on BCL-XL, so these compounds would likely still cause transient thrombocytopenia. They might also be less potent senolytics for senescent populations that express higher levels of BCL-2 or BCL-W (46, 48). Local administration of BCL-2 family inhibitors to the insulated sites of interest could limit those side effects. For instance, intra-articular injection of the BCL-2-targeting UBX0101 compound efficiently eliminated senescent cells in articular cartilage and synovium and reduced signs of osteoarthritis in aged mice (33). Overall, the available evidence suggests that although targeting the BCL-2 family of proteins is a molecularly validated mechanism-based approach for targeting senescent cells, its side effects will likely limit therapeutic use.

**Senolytic combinations.** An alternative route to overcoming the cytotoxic side effects of BCL-2 protein family inhibitors might be to synergistically combine low doses with compounds that target other senescence-associated prosurvival pathways. Indeed, the first published senolytic was not a single agent, but a combination of the pan–tyrosine kinase inhibitor dasatinib and a naturally occurring flavonoid, quercetin (47). When given together, these two compounds were shown to selectively kill senescent primary preadipocyte cells and senescent human umbilical vein endothelial cells (HUVECs) in tissue culture, as well as reduce senescent cell signature in a set of pathological conditions (30, 31, 47, 58). Administration of the dasatinib/quercetin combination reduced aortic calcification and osteogenic signaling in aged and hypercholesterolemic mice (34). It also improved lung function during bleomycin-induced lung fibrosis in mice (30). In both systems, however, the combination failed to alleviate the fibrotic component of these conditions, suggesting that it may be more effective as a preventative intervention during early stages of diseases. Consistent with this suggestion, dasatinib/quercetin can prevent age-related bone loss and reduce overall hepatic steatosis (a prerequisite condition of nonalcoholic fatty liver diseases) (31, 32).

Dasatinib and quercetin were previously tested in clinical settings, making this combination a promising senolytic candidate. Nevertheless, some issues might impede its future use. First, both compounds target a long list of biological pathways, and the combination suffers from lack of understanding of the mechanisms underlying its senolytic actions. In addition, the effects observed following the combination treatment might be mediated through alterations in multiple target pathways in multiple tissues and organs, which could result in adverse effects after long exposures. Still, short-term administration of the combination is sufficient to eliminate senescent cells and provide benefit as observed in the above studies. Since the combination might not need to be present continuously in order to be effective, off-target complications could be avoided. In addition to quercetin, other natural compounds, including fisetin and piperlongumine, have been suggested to have senolytic effects (56, 59). These molecules, too, are expected to kill senescent cells by targeting multiple molecular pathways. Owing to a lack of well-defined mechanism of action and questionable pharmacokinetics, it is unlikely that they will find their way into the clinic.

**Targeting senescence-specific pathways.** Recent studies highlight the p53/p21 axis as a promising target for development of novel senolytics (49, 60). p53, the most frequently mutated tumor suppressor gene in human cancer, is a core component of the senescence program (61). p53 is stabilized and accumulates in response to DNA damage (7, 62, 63). Its transcriptional activity controls a myriad of biological processes, positioning it on a decisional fork between transient cell cycle arrest, senescence, and apoptosis (61). Interfering in p53’s direct interaction with the transcription factor FOXO4 leads to the release of p53 from the nucleus and induction of cell-intrinsic apoptosis. Administration of a modified FOXO4/p53-interfering peptide was able to neutralize murine liver chemotoxicity in the setting of doxorubicin treatment and restore fitness, hair density, and renal function in progeroid and naturally aged mice (60). The senolytic potential of this pathway is not limited to the p53-FOXO4 interaction (64). p21 (also known as CDKN1A), a primary transcriptional target of p53, is necessary for survival and retention of senescent cells in tissues (48). Following DNA damage, it prevents NF-κB–mediated activation of JNK, caspase-3 cleavage, and subsequent cell death. In mice, p21 knockout leads to a reduction of senescent cells in fibrotic livers and alleviates liver fibrosis (49). Despite the fact that the p53/p21 axis can be indirectly modulated, for example by controlling ROS levels, there are no known drugs that can induce apoptosis in senescent cells by inhibition of either p53 or p21. Therefore, further research and development of new drug candidates is needed to fulfill the promise of the p53/p21 prosurvival pathway in targeting senescent cells. Currently, the senolytic effect of the FOXO4 peptide is a promis-
The identification of molecular pathways that preserve the viability of senescent cells provided the initial insights for the development of senolytics. In addition to pathway-based senolytic development, high-throughput approaches can also be used to identify new targets. Lately, the development of such an approach identified HSP90 proteins as a novel class of targets (44). HSP90, a family of ubiquitously expressed molecular chaperones, can promote cell survival via stabilization of AKT or ERK (65, 66), members of signaling pathways that are upregulated during senescence (67, 68). Disruption of the HSP90-AKT interaction inhibited the PI3K/AKT pathway, resulting in a selective killing of senescent cells of different origins (44). In vivo, administration of the HSP90 inhibitor 17-DMAG to progeroid mice reduced the senescence signature and extended health span. These findings mark HSP90 inhibitors as potential constituents for future combinatorial senolytics.

Directed elimination of senescent cells. Targeting of senescence-specific pathways is promising, but it is not the only way to develop senolytics. It is possible to eliminate senescent cells by exploiting other senescence-specific features. One well-described feature of senescent cells is senescence-associated β-galactosidase (SA β-gal) activity (26). A targeted delivery system using mesoporous silica nanoparticles coated with galacto-oligosaccharides was developed based on this feature (69). While the coated particles cannot be activated in nonsenescent cells, the coating is digested in senescent cells and the nanoparticle content can be released. It is conceivable that coated nanoparticles containing a cytotoxic drug could release it to the cytoplasm of senescent cells to induce apoptosis. Increased β-galactosidase activity is not a complete surrogate feature of senescent cells, as it is known that other cells in the body, e.g., macrophages, exhibit high β-galactosidase activity in some circumstances (70). In this case, it may be beneficial to increase selectivity by tethering the particles to antibodies against known surface markers for senescent cells (71). Senolitic approaches based on SA β-gal activity are yet to be tested in vivo, and more studies are needed to examine the outcomes of such treatments.

Immune-mediated interventions. Apart from directed elimination of senescent cells based on their intrinsic properties, therapeutic interventions could be also mediated by the immune system. Senescent cells are immunogenic in nature and are subject to immune surveillance mechanisms (13, 20, 21, 23, 72, 73). In a model of hepatocellular carcinoma, induction of senescence by p53 restoration triggers an innate immune response consisting of infiltrating leukocytes, neutrophils, macrophages, and NK cells (21). In a model of liver fibrosis, senescent hepatic stellate cells promote their own NK cell-mediated elimination (13). Senescent cells upregulate the immune recognition molecule NKG2D, which is usually not present on the surface of normal cells (72). Thus, NKG2D expression is a potential asset in immunotherapy approaches. The mechanisms of senescence surveillance are context-dependent. Induction of senescence by expressing of NRASG12V in hepatocytes results in a mixed phenotype of both innate and adoptive immune response, as CD4+ T cells cooperate with monocytes/macrophages to execute the clearance of senescent hepatocytes (23). Accumulation of senescent cells in aged tissues could be attributed, at least in part, to an age-related decline in immune function. Therefore, repotentiating the immune system may result in successful clearance of senescent cells from aged tissues. In support of this notion, administration of the immune stimulator polyI:C facilitates NK cell-mediated clearance of senescent cells in fibrotic livers (13). However, in vivo treatment with such a strong immune stimulator would probably overwhelm the aging immune system, which is often chronically activated. Therefore, more precise immune modulators should be considered.

Adapting common immunotherapy practices from the field of cancer might contribute to this effort. For example, blockade of the immune checkpoint programmed death 1 (PD-1) is a powerful treatment for cancer that can also improve pathological symptoms in a murine model of Alzheimer’s disease (74). In addition to immune-stimulating strategies, a growing arsenal of tools such as chimeric antigen receptor (CAR) T cells provide the option to redirect immune responses against senescent cells (75). The bottleneck in this and other immune-targeting approaches is a shortage of highly selective senescence markers. Very recently, however, novel senescence cell-surface antigens have been suggested. In one study, senescent fibroblasts were shown to express an oxidized form of membrane-bound vimentin on their cell surface (76). In another study, mass spectrometry analysis identified dipeptidyl peptidase 4 (DPP4) as a cell surface marker for senescent fibroblasts (77). The selective expression of DPP4 on the surface of senescent cells enables their preferential killing by NK cells in an antibody-dependent cell-mediated cytotoxicity (ADCC) assay (77). Senescent cells are partly protected from NK cell-mediated cytotoxicity because of their upregulation of decoy receptor 2 (DcR2) (41). DcR2 inhibits the activation of death receptors 4 and 5 (DR4/5) by TNF-related apoptosis-inducing ligand (TRAIL), confining the actual killing to perforin- and granzyme-mediated pathways. Finding ways to boost this process, e.g., by blocking DcR2, could render senescent cells more vulnerable to natural immune surveillance and immunotherapy-based targeting. Overall, while the findings regarding the immune targeting of senescent cells are quite preliminary and require further investigation, they highlight the potential in harnessing the immune system to eliminate senescent cells in the context of age-related pathologies. Identification of novel and specific cell surface markers of senescent cells will further facilitate the development of immune system-based senotherapies.

Strategies aimed to limit detrimental roles of senescent cells. While elimination of senescent cells appears to be a favorable strategy, it might be unsuitable for some conditions in which senescent cells occur. In those cases, specific targeting of detrimental components of senescence could potentially serve as an alternative strategy. Senescent cells exhibit a profound proinflammatory secretory profile that is largely conserved between different senescent states and cell origins (19, 78–80). The senescence-associated secretory phenotype (SASP) is mainly composed of inflammatory chemokines and cytokines, growth factors, and matrix-remodeling proteases. The SASP promotes non–cell-autonomous tumor suppression by enforcing cell cycle arrest in damaged and cancer cells, and by recruiting immune cells to eliminate them. Nonetheless, it has also been strongly implicated in disruption of tissue homeostasis and development of age-related pathologies (81). Several signaling pathways that converge into activation of the NF-κB pathway and the C/EBPβ pathway orchestrate the complex secretome (78, 80,
82–84). These signaling pathways are well studied and can be modulated by a broad spectrum of drugs, some of which are approved by the FDA. For instance, IL-1α/IL-1 receptor signal transduction is upstream of NF-κB, and the use of neutralizing antibodies against either IL-1α or its receptor is sufficient to reduce NF-κB transcriptional activity (85). mTOR inhibitors, such as rapamycin and its analogs, can abolish SASP by reducing the expression of membrane-bound IL-1α (86). This action has been shown to ameliorate the ability of senescent fibroblasts to stimulate prostate tumor growth in mice. Metformin, a commercially approved drug for type 2 diabetes, blocks NF-κB translocation to the nucleus, thus restricting its transcriptional activity (87). Treating mice with metformin limits inflammatory processes, alleviates different age-related pathologies, and extends lifespan (88). Reports of similar effects in human diabetic patients have initiated investigations into metformin’s effects on aging in the general population (89, 90). C/EBPβ transcriptional activity can be induced by JAK/STAT signaling (91). JAK1/2 inhibitors, such as the FDA-approved drug ruxolitinib, can reduce systemic inflammation and improve fitness in aged mice (92). Therefore, there is evidence that existing inhibitors can dampen the effects of SASP in vivo and potentially reduce the deleterious effects of senescent cells.

While restricting inflammation is a potentially promising way to limit the non–cell-autonomous effects of senescent cells, treatment with strong antiinflammatory drugs can potentially cause a wide spectrum of side effects (93). These circumstances might not permit a prolonged therapeutic regimen, which is prerequisite to preserving their beneficial effects. The solution to this problem might come from new studies on epigenetic regulators, including mixed-lineage leukemia protein 1 (MLL1), bromodomain-containing protein 4 (BRD4), and high-mobility group box 2 (HMGB2), which are responsible for reshaping the epigenetic landscape of senescent cells in a manner that would dictate expression of SASP components (24, 94, 95). Experimental knockdown of these chromatin modifiers results in sustainable antiinflammatory effects without compromising cell cycle arrest. Relevant small-molecule modulators, such as the bromodomain and extra-terminal motif (BET) inhibitors, are available and should be further tested for this exciting possibility (96). Targeting specific components of SASP could also provide a safer way to mitigate the deleterious effects of SASP. Cytokines, well-defined components of SASP, such as IL-6, IL-8, and matrix-remodeling proteases such as ADAM17, could serve as possible targets. An attractive method to block these molecules is the application of neutralizing antibodies, which can be developed based on the plethora of commercially available monoclonal antibodies. For example, targeting of IL-6 or its receptor could be achieved by the approved drugs siltuximab or tocilizumab, blocking the cytokine or the receptor, respectively (97–99). However, these agents were not developed for use in the context of cellular senescence, and their effect on SASP and aging must be studied in detail before their effect as senescence modifiers can be exploited therapeutically.

### Moving toward clinical trials

Successful translation of strategies aimed at targeting senescent cells into the clinic could have a vast impact on the treatment of a variety of diseases at old age and potentially revolutionize our view of aging. Several technologies aiming at senescent cells are now taking their first steps in the challenging, time-consuming, and expensive journey of drug development (Table 1). In order to maximize their chances of becoming common treatments, critical issues of efficacy, safety, and tolerability should be considered during the early stages of development.

**Upcoming challenges of strategies to target cellular senescence**

The main question that still remains in the field is the efficiency of different senotherapies in different tissues of human patients. Many of the present data on these treatments were obtained from experiments in cell culture, which might only partially represent the situation in vivo and are far from the context of a disease. Efficacy of compounds in vivo was often tested in mouse models, some of which are only partly relevant to aging and age-related pathologies. Therefore, the relevance to specific human conditions and pathologies must be properly investigated before senotherapies can be enrolled for clinical trials. Moreover, current strategies target a general population of senescent cells, whereas senescent cells are a highly heterogeneous population. Diversity in cell of origin, trigger of senescence induction, or pathological context can create disparate vulnerability of cell subpopulations to a certain treatment. For this reason, it would be rational to expect that some treatments would be more suitable than others for a certain tissue, organ, or age-related disease. Therefore, a current concept that considers senotherapies as pan-antisenescence treatments is somewhat concerning and needs to be carefully reevaluated. It will be necessary to investigate the influence of every treatment on diverse cell populations while using the most relevant model for each condition to which the treatment may be applied.

### Table 1. Moving into the clinic: current translational activities in the field of cellular senescence

<table>
<thead>
<tr>
<th>Strategy</th>
<th>Company</th>
<th>Class of drug</th>
<th>Target</th>
<th>Stage</th>
</tr>
</thead>
<tbody>
<tr>
<td>Senolytics</td>
<td>Unity Biotechnology</td>
<td>Small molecules</td>
<td>BCL-2 family</td>
<td>Preclinical</td>
</tr>
<tr>
<td></td>
<td>Oisin Biotechnologies</td>
<td>Nanoparticle-based gene therapy</td>
<td>p16INK4A</td>
<td>Preclinical</td>
</tr>
<tr>
<td></td>
<td>Senolytic Therapeutics</td>
<td>Nanoparticle-based drug delivery</td>
<td>ji-Galactosidase</td>
<td>Preclinical</td>
</tr>
<tr>
<td></td>
<td>Antoxerene</td>
<td>Small molecules</td>
<td>p53-FOXO4 interaction</td>
<td>Discovery</td>
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<td></td>
<td>CellAge</td>
<td>Gene therapy</td>
<td>Multigene senescence signature</td>
<td>Discovery</td>
</tr>
<tr>
<td>Senescence immunotherapy</td>
<td>Everon Biosciences</td>
<td>Small molecules and immunotherapeutics</td>
<td>Senescence-associated macrophages</td>
<td>Preclinical</td>
</tr>
<tr>
<td></td>
<td>Siwa Therapeutics</td>
<td>Antibodies</td>
<td>AGE-modified proteins</td>
<td>Preclinical</td>
</tr>
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AGE, advanced glycation end-products.
In using senescence-targeted treatments, safety issues might become a major challenge, owing to the physiological role of senescent cells and the side effects of the existing senolytics. Some senolytics, especially among BCL-2 family inhibitors, pose a cytotoxic threat to nesencessent cells when administered systemically (55). In the case of immunotherapy, certain senescence epitopes might be also expressed to some extent in nesencessent cells, such as stressed cells or secretory cells. There are several possible ways to overcome this problem. For instance, drugs could be administered directly to the area of interest. More importantly, combinatorial approaches based on nesencessent cell–specific delivery of senolytics could have a synergistic impact, thus allowing the use of lower, less toxic doses with higher efficacy and better safety. Another possibility to increase targeting specificity is to modify the agent to be active only after processing by an enzyme that is preferentially expressed by nesencessent cells. In addition to the side effects of the targeting moiety itself, compromised safety might also stem from possible interference with the beneficial roles of senescence. The senescence program contributes to embryogenesis, tissue repair, regeneration, and tumor suppression, but the effect of targeting senescence on these processes is still unexplored in the context of senolytic treatments (8, 100–105). Fortunately, existing senolytics induce apoptosis in senescent cells after short exposures, and their administration can be limited to short periods of time. Future treatments will have to take this notion into account and will have to be conducted in a spatiotemporally controlled manner.

Careful design of clinical studies would have a strong impact on their fate. Indications for first-wave senotherapies should be carefully chosen and could involve an uncommon age-related pathology with only a symptomatic rather than therapeutic treatment as standard of care. A causal role of cellular senescence in the pathogenesis of each disease or condition must be well ensured, and the mechanism of action should be clear. Availability of adequate preclinical models will help to evaluate the effect of each treatment’s risks and benefits. For successful clinical studies there is also a cardinal need for senescence-specific and disease-oriented biomarkers for prognostic and predictive use. Basic research is still constrained by the lack of satisfactory senescent markers, and quantitative methods are limited to ex vivo analysis of cells (106, 107). There is an urgent need to develop novel senescence biomarkers in order to reliably assess senescence burden in tissues in a noninvasive manner. Candidate patients must be considered according to their senescence burden. Careful consideration of the scientific evidence, experimental settings, and patient management will increase the chances of successful trial completion.

Overall, cellular senescence is a comprehensive phenomenon with both beneficial and detrimental consequences. Senotherapies hold promise to block detrimental effects and treat a large set of age-related pathologies, but they will have to be implemented in an accurate manner in order to assure both efficient and safe use. Understanding the mechanisms of possible side effects while designing the dosage and the regimen of senolytic treatments will help facilitate their way to the clinic and provide new hope for improvement of human health span and longevity.

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