

# Aging, Cellular Senescence, and Type 2 Diabetes: Unraveling the Links for Novel Therapeutic Approaches

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#### **ABSTRACT**

Diabetes mellitus, particularly type 2 diabetes mellitus (T2DM), is an escalating global health crisis, with projections indicating a rise from 537 million to 783 million affected individuals by 2045. Central to the progression of T2DM is the intricate interplay between insulin resistance, aging, and cellular senescence, especially in peripheral adipose tissues and pancreatic  $\beta$ -cells. This review highlights pathophysiological mechanisms underlying T2DM, emphasizing how aging exacerbates insulin resistance and impairs insulin secretion. We present evidence linking cellular senescence with increased adiposity and dysfunction in  $\beta$ -cells, illustrating how age-related changes foster a chronic inflammatory milieu that worsens metabolic dysregulation. Cellular senescence, characterized by the accumulation of dysfunctional cells producing the senescence-associated secretory phenotype (SASP), contributes significantly to the development and complications of diabetes. Notably, emerging therapies targeting senescent cells, such as senolytics and senomorphics, offer innovative strategies to mitigate diabetes-related decline. These therapeutic approaches aim to enhance insulin sensitivity and improve  $\beta$ -cell function, potentially reshaping diabetes management. This review underscores the importance of a nuanced understanding of cellular senescence in devising targeted senotherapies, advocating for a dual organ-oriented strategy to optimize treatment outcomes for diabetes and associated comorbidities.

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## 1. INTRODUCTION

Diabetes mellitus is becoming an increasingly prevalent issue worldwide. In 2021, it was estimated that 537 million people were living with diabetes mellitus, which was also linked to 6.7 million deaths. Projections suggest that the number of individuals affected by this disease will rise to 643 million by 2030 and 783 million by 2045 [1]. Additionally, the majority of those with type 2 diabetes mellitus (T2DM) are over 65 years old. Nearly 1 in 4 people suffer from T2DM, and this number continues to grow. To address this challenge, it is crucial to gain a thorough understanding of the detailed pathogenesis of diabetes and to develop new treatment strategies focused on both clinical and molecular aspects of aging [2-5].

Intensified insulin resistance and weakened insulin secretion play key roles in the causes, mechanisms and patterns of T2DM. Insulin resistance occurs due to impaired responses to insulin signaling in peripheral tissues such as muscle, liver, and fat. This resistance is further exacerbated by obesity, which leads to the accumulation of dysfunctional adipose tissues and a compensatory increase in insulin secretion [6-8]. Eventually, a decline in pancreatic  $\beta$ -cell function can lead to the development of diabetes mellitus in susceptible individuals. The quantity and function of adipose tissue are influenced by aging, as well as by factors like calorie intake, physical activity, and overall health. Consequently, aging is well known to be associated with increased insulin resistance, highlighting the significant role of cellular senescence in adipose tissue in

the onset and progression of the disease [9-11].

Insulin generation is known to decrease with age. This decline is influenced not only by the secretory capacity of individual  $\beta$  cells but also by the overall  $\beta$ -cell mass (BCM). Researches involving isolated human islets have shown an age-related reduction in insulin secretion [12]. While it is generally believed that BCM is not solely impacted by aging, significant changes in BCM can occur with the development and progression of obesity and diabetes. A reduction in BCM can have a substantial effect on diabetes management, including glycemic control and the effectiveness of anti-diabetic treatments. In response to increased insulin resistance, BCM may expand to compensate for lack of insulin generation [13,14]. However, this compensatory expansion may be restricted in part due to a decrease in  $\beta$ -cell multiplication as a result of aging. The age-related changes in  $\beta$ -cell function and multiplication highlight the role of cellular aging in the disease caution of type 2 diabetes mellitus (T2DM). That's why, a better knowledge of cellular senescence in  $\beta$  cells and adipose tissue could lead to new curative strategies for the illness [15-18].

Aging in humans is associated with a range of complicated symptoms and diseases, such as diabetes and obesity, which arise from the gradual decrease in the function of different tissues and organs over time [19-21]. Strehler identified four key characteristics of aging in organisms: totality, entity, progressiveness, and harmfulness [22]. In line with the concept of totality, it is widely recognized that aging cells collect in the tissues and organs of aged individuals. As aging progresses or diseases develop, normal cells and tissues in the body are exposed to various stresses, leading to cellular damage that requires recovery, adjustment, programmed cell death, or other protection mechanisms [23-25]. Cellular aging refers to a state where cells undergo irreversible cell cycle capture and experience functional decline due to telomere shortening or stressors that provoke aging, such as DNA damage, oncogenic stress, and oxidative stress. Senescent cells differ from non-senescent ones, exhibiting unique characteristics like an enlarged, flattened shape, altered nuclear structure, formation of H2Ay foci, enhanced expression of cell cycle blockers (p16Ink4 and p21Cip1), and more [26-29].

The INK4/ARF site encrypts the neoplasm cancellers, p16Ink4 and p19Arf, both of which function as inhibitors of the cell cycle. p16Ink4 inhibits cyclin-dependent kinase (CDK)4/6, while p19Arf prevents the degradation of p53. Additionally, p21Cip1, another CDK inhibitor, is a transcriptional target of p53 [30-32]. In normal cells, the INK4/ARF site is regulated epigenetically. In young cells, it is kept quiet by polycomb repressive complexes PRC1 and PRC2, which contain chromobox protein homolog (CBX)7, B cell-specific Moloney murine leukemia virus integration site (BMI)1, and enhancer of zeste homolog 2. These complexes

maintain repressive epigenetic marks, such as trimethylation of histone H3K27 on the locus. However, in senescent cells, epigenetic activation of the INK4/ARF locus leads to unchangeable cell cycle capture [33-35]. Interestingly, the ectopic expression of CBX7 or BMI1 can beglect aging in basic cells. Furthermore, several other epigenetic factors, including Mixed lineage leukemia protein-1, Jumonji domain-containing protein-3, and Zuotin-related factor 1, also play roles in the regulation of this locus [36,37].

The idea that aging is an entirely irreversible process has been partially challenged. For instance, telomere length, a notorious marker of cellular aging in vitro, is one area of focus. Telomeres reduce as a result of replicative depletion. Recent advancements in technology have made it possible to measure telomere length in leukocytes, revealing a strong correlation between telomere length and the health status of individuals, particularly in relation to conditions such as cancer, atherosclerosis, heart failure, diabetes, depression, and chronic inflammation [38-41]. Interestingly, telomere reduction can be partially revoked through lifestyle modifications, including regular physical activity, a balanced diet without excess calories, and stress reduction. These findings indicate that certain aspects of the aging process may be correctable.

The negative aspects of aging have become a topic of debate due to the complex, dual nature of senescent cells [42-44]. Even adolescent cells with undamaged telomeres can enter a senescent state when exposed to oncogenic stress, such as the activation of oncogenic Ras, a process known as oncogene-induced senescence (OIS). OIS cells are also observed in vivo and are believed to act as a protective mechanism against the harmful development of neoplasms. However, senescent cells also exhibit increased secretion of proinflammatory cytokines and chemokines through the activation of the nuclear factor-kappa B (NF-kB) pathway, leading to what is termed a senescence-associated secretory phenotype (SASP) [45-47]. Additionally, the activation of the NF-kB pathway makes senescent cells more impervious to apoptotic incentives by amplification of anti-apoptotic factors like X-linked inhibitor of apoptosis protein and Bcl-2. The SASP contributes to chronic inflammation, which in turn fosters the development of age-related diseases. Consequently, senescent cells not only accumulate in aged tissues but also contribute to organ dysfunction associated with various lifestyle diseases, playing a key role in their illness formation [48-50].

According to these concepts, targeting chronic inflammation or removing senescent cells is becoming a promising therapeutic strategy for diseases associated with aging. Antibody therapies targeting inflammatory cytokines like interleukin (IL)-6 and tumor necrosis factor (TNF)-α are already in clinical use through molecularly aimed pharmaceuticals [51,52]. Lately, an anti-IL-1 antibody drug, canakinumab, was produced for the cure of certain autoimmune conditions. Notably, the Canakinumab Anti-inflammatory Thrombosis Outcome Study (CANTOS) demonstrated that canakinumab could impede atherosclerosis and lung cancer in approximately 10,000 elderly individuals [53-56]. This represents a groundbreaking example of the effective reduction of chronic inflammation. Additionally, last improvements in aging

research have indicated that eliminating senescent cells, a process known as senolysis, is an encouraging strategy (senotherapy) for minimizing chronic inflammation and cure age-associated illnesses [57-60].

## Fundamental ageing mechanisms contribute to diabetes pathogenesis

The mechanisms underlying aging are numerous and interconnected. A few models have been proposed to categorize agerelated processes, superposing with abnormalities observed in obesity and diabetes. This overlap has been particularly well-documented in the situation with adipose tissue. Generally, the fundamental mechanisms of aging can be grouped into the next categories: (1) macromolecular dysfunction, which includes the loss of proteostasis, impaired DNA damage repair, and abnormal mRNA processing; (2) sterile inflammation, characterized by the infiltration of immune cells and the emission of proinflammatory cytokines without the presence of a specific pathogen, along with fibrosis; (3) progenitor cell dysfunction, which encompasses the depletion of progenitor cell pools, reduced differentiation capacity, or abnormal lineage distribution; and (4) cellular aging [61-64]. These groups are explored more thoroughly below, particularly in relation to their roles in the pathogenesis and development of diabetes. Special attention is given to adipose tissue due to its significant role in the progression of inflammation, insulin impedance, and related maladies, and because several mechanistic studies on senescent cells within adipose tissue have been conducted [65-67].

#### Senescent cell burden is increased in diabetes

The primary risk factors for progression type 2 diabetes are age and adiposity, both of which are linked to a higher accumulation of aging cells. While cellular senescence is thought to play a role in the onset of diabetes, as mentioned earlier, the diabetic microhabitat also appears to contribute to the increased charge of senescent cells [68,69]. For instance, heightened glucose and lipid levels, similar to inflammation, can trigger cellular aging. Either type 1 or type 2 diabetes are connected to a higher risk of glucose-related microvascular intricacies affecting the eyes, nerves, and kidneys. However, the role of cellular aging in the evolution of these intricacies remains poorly understood [70].

While most of the current researches on the consequences of cellular senescence has been performed using animal models, several studies have also examined senescence in human cells from both diabetic and non-diabetic individuals. The following is a summary of the latest data on cellular senescence in key tissues linked to the development and clinical manifestations of type 2 diabetes [71-74].

## Beta cells

The onset of insulin resistance requires a compensatory growth in insulin secretion to maintain normal glucose levels. Type 2 diabetes develops when this compensatory insulin secretion becomes insufficient to counteract the level of insulin resistance. A few investigations have demonstrated that the gene expression profile in beta cells changes with age, with an upregulation of genes associated with cellular senescence, such as **Cdkn2a** and **Cdkn2b** [75-77]. While this typically results in a reduced capacity for cell proliferation, an unexpected discovery was that insulin secretion was actually enhanced in **p16Ink4a**-induced senescent cells, rather than diminished. The extent to which this finding explains the age-related increase in basal insulin secretion, generally attributed to the concurrent rise in obesity and insulin resistance, remains unclear but presents an intriguing hypothesis [78-81]. More recently, research has shown that eliminating senescent beta cells in a mouse model of type 1 diabetes improved insulin secretion and preserved the cells' insulin-producing ability, suggesting a novel connection between cellular senescence and serious insulin lack [82-86].

## Abdominal/visceral obesity

Inflammation in adipose tissue increases as a result of adipocyte hypertrophy, which in turn can lead to the aggregation of aging cells. Fatness, particularly when linked to hypertrophic expansion, is associated with elevated indicators of cellular senescence. These markers include increased β-galactosidase activity in adipose tissue, which indicates high lysosomal activity and content, as well as elevated levels of plasminogen activator inhibitor 1 (PAI-1), p53, and cyclin-dependent kinase inhibitors like p16Ink4a [87-91]. Senescent ancestry cells impede adipogenesis, leading to ectopic lipid collection, increased visceral fat, and abdominal fatness. Correspondingly, ageing is connected with the buildup of non-dividing senescent cells in adipose tissue. Additionally, age-related increases in visceral fat have been observed independently of BMI [92-96].

### Fatty liver disease

Type 2 diabetes is linked to a higher risk of developing non-alcoholic fatty liver disease (NAFLD). Recent studies have demonstrated that the burden of senescent cells in the liver is elevated in individuals with NAFLD, with the degree of steatosis corresponding to indicators of senescence. In mouse models, inducing senescence specifically in hepatocytes led to increased fat accumulation, suggesting that senescent hepatocytes play a direct role in the development of NAFLD. Furthermore, treatment with senolytics (D+Q) was found to reduce steatosis [97-100].

#### Cardiovascular disease

Cells in the aortic media and atherosclerotic plaques of hypercholesterolaemic (ApoE-/-) and ageing mice exhibit elevated pointers of senescence. Research has shown that the removal of senescent cells improves vascular smooth muscle sensitivity to nitric oxide donors and reduces plaque calcium deposits, suggesting that senescent cells contribute to endothelial dysfunction in atherosclerosis [101-103]. Additionally, clearing senescent cells in obese mice improved cardiac diastolic function, a finding with potential relevance for diabetic patients, who often experience heart failure with preserved ejection fraction. During aging, senescent cardiac ancestry cells develop a hypertrophic, pro-fibrotic phenotype and lose their ability to replicate. Removing these senescent cells in mice reduced age-related dysfunction in cardiac ancestry cells and reduced fibrotic area formation following myocardial infarction [104-107].

## **Renal dysfunction**

Cellular senescence is more prevalent in kidney cells from individuals with type 2 diabetes and also enlargers with age in non-diabetic individuals. The clinical significance of this observation is underscored by recent findings that senolytic treatment (D+Q) reduced proteinuria in obese, insulin-resistant mice [108-110].

Cognitive dysfunction and Alzheimer's disease

Cellular senescence has been implicated in mental disability in both obese, insulin-resistant mice and aged mice. Senolytic therapy (D+Q) decreased the abundance of senescent cells in the brain, restored neurogenesis, and alleviated neuropsychiatric defect in obese animals. In aged mice with an Alzheimer's-like condition induced by Tau protein overexpression, these agents also reduced neuroinflammation, restored neurogenesis, and partially reversed brain atrophy [111-113].

Collectively, these findings from both human and animal studies strongly support the role of cellular senescence in the development of diabetes and its associated complications. Moreover, they suggest that reducing the burden of senescent cells could be a promising new therapeutic approach for managing diabetes and its related difficulties [114-116].

# Senotherapy: Senolytics and Senomorphics

Table 1. Senotherapeutic Strategies for Managing Type 2 Diabetes Mellitus (T2DM)

Therapy Type	Examples	Mechanism of Action	Clinical Benefits	Research Status
Senolytics	Dasatinib, Quercetin (D+Q)	Induces apoptosis in senescent cells	Reduces senescent cell burden, enhances tissue function	Clinical trials ongoing
	ABT263, ABT199	Targets BCL-2 to promote β-cell apoptosis	1	Preclinical and clinical data available
Senomorphics	Metformin		Improves insulin sensitivity, decreases inflammation	Widely used in clinical practice
	Curcumin	Activates autophagic pathways, counters oxidative stress	Protects against β-cell dysfunction	Emerging evidence, clinical studies ongoing
Antibody Therapies	Anti-IL-6, Anti-TNF-α	Neutralizes inflammatory cytokines	Reduces chronic inflammation, improves insulin sensitivity	Under investigation in clinical trials

Therapy Type	Examples	Mechanism of Action	Clinical Benefits	Research Status
Lifestyle Interventions	Diet, Exercise	Promotes metabolic health and reduces adiposity		Well-established; recommended for all T2DM patients
Combination Therapies	Metformin + Senolytics		Potentially greater improvements in glycemic control	Research ongoing; promising preliminary results
Nutraceuticals	Omega-3 Fatty Acids	Modulates inflammatory pathways	Reduces inflammation, supports cardiovascular health	Some evidence supports efficacy
Gene Therapy		Targets senescence- related genes	Potential to delay or reverse age-related dysfunction	-

Senotherapies, which include senolytics and senomorphics, have the potential to reduce the onset of age-related pathologies. Senolytics work by selectively eliminating senescent cells through the targeting of SCAP (Senescence-Associated Anti-apoptotic Pathways). RNA interference techniques have identified drugs that target critical nodes within SCAP pathways, such as BCL-2/BCL-XL, PI3K/AKT, and p53/p21/serpins [117-119]. This approach has led to the discovery of several senolytics that induce apoptosis in senescent cells. On the other hand, senomorphics modulate SASP (Senescence-Associated Secretory Phenotype) pathways without killing the senescent cells [120,121].

This part will concentrate on the senolytics and senomorphics that have been used in clinical tests and animal studies, particularly in the context of type 2 diabetes (T2DM) and its intricacies. Senolytics such as D, Q, fisetin, and ABT263 have been investigated, while metformin has been explored as a senomorphic.

## 2. DASATINIB AND QUERCETIN

D is a tyrosine kinase inhibitor used in the treatment of chronic myeloid leukemia and Philadelphia chromosome-positive acute lymphoblastic leukemia that is firm or prejudiced to other treatments. Q is an innate flavonoid that hinders PI3K [122,123]. The first clinical use of a senolytic involved the combination of D+Q, which was reported in 2015. In studies involving insulin resistance using S961, an insulin receptor antagonist, D+Q was shown to reduce the number of  $\beta$ -galpositive dispersed islet cells both in vitro and in vivo, and to lower blood glucose levels in a mouse model of insulin resistance [124-126].

The D+Q combination also diminished the number of naturally occurring senescent cells and their SASP emission in human adipose obese extracts. This research utilized omental adipose tissue obtained from eight obese subjects undergoing gastric bypass surgery. Notably, D+Q has potential adverse reactions including hematologic dysfunction, fluid retention, skin rash, and QT prolongation. Further clinical studies are required to assess its potential efficacy in treating type 2 diabetes (T2DM) [127-129].

BCL-2 Inhibitors: ABT263 (NAVITOCLAX), ABT199 (VENETOCLAX), and ABT737

ABT263 (Navitoclax) aims the BCL-2 course. The studies found that ABT263 effectively eliminated a significant portion of  $\beta$ -gal-positive dispersed islet cells in vitro. Additionally, ABT263 reduced blood glucose grades and p16Ink4a emission in a medication-associated insulin-repellent mouse model [130-132]. It also reduced p16Ink4a levels in peripheral tissues in a high-fat diet (HFD)-induced insulin resistance example. When used in combination with D+Q, Navitoclax selectively aimed and eliminated senescent cells, and improved outcomes in SARS-CoV-2-infected hamsters and mice with reduced lung disease. Despite its potential, common side effects such as diarrhea, nausea, and thrombocytopenia restrict its clinical

applicability [133-135].

ABT199 (Venetoclax) optionally hinders BCL-2, an apoptosis-suppressing protein. Venetoclax is an oral BCL-2 hinder endorsed for use in relapsed/refractory chronic lymphocytic leukemia and acute myeloid leukemia. In studies, administration of ABT199 to normoglycemic non-obese diabetic (NOD) mice suppressed the development of diabetes in contradiction of controls in a type 1 diabetes mellitus (T1DM) example [136,137].

ABT737, another BCL-2 inhibitor, has demonstrated productive antitumor activity in cancerous lymphoma and small-cell lung cancer. Treatment with ABT737 in NOD mice resulted in a 30% reduction in  $\beta$ -cells emitting cyclin-dependent kinase inhibitor 2A (Cdkn2a) and a diminished incidence of T1DM [138-141].

## **Senomorphic: Metformin**

Metformin has been used for decennaries to cure type 2 diabetes (T2DM) by reducing circulating glucose levels through the restraint of hepatic gluconeogenesis. Beyond its glucose-lowering effects, metformin may also possess anti-aging characteristics by avoiding DNA damage and inflammation and has been shown to exhibit senomorphic effects. In a mouse model of kidney disease, a brief course of metformin inhibited acute senescence in bone marrow mesenchymal stem cells (MSCs) [142,143]. Other studies have demonstrated that metformin reduces ROS levels and delays the onset of senescence in mouse adipose-derived MSCs. Additionally, treating human adipose stem cells with metformin at therapeutic concentrations for 6 weeks led to a decline in β-galactosidase activity. Measuring circulating SASP levels before and after metformin control could help to define patients who might profit most from its senomorphic consequences [144-146].

#### **Others: Curcumin**

Curcumin, a phytocompound found in turmeric, has been investigated in numerous clinical tests for type 2 diabetes (T2DM). Research has shown that a combination of curcumin and hesperetin, two antioxidant polyphenolic compounds, can enhance cellular senescence outcomes. This combination has been reported to reduce  $\beta$ -galactosidase staining, p16Ink4a, and p21Cip1 in neurons and rats induced with D-galactose. Additionally, curcumin has been shown to mitigate D-galactose-induced senescence in cardiomyocytes by promoting autophagy through the SIRT1/AMPK/mTOR pathway [147-149].

While numerous clinical studies have explored the effects of curcumin on type 2 diabetes (T2DM), there is currently no evidence directly demonstrating its impact on senescent pancreatic β-cells. Many trials have shown that curcumin significantly reduces T2DM incidence among individuals with prediabetes. In diabetic mouse models, curcumin treatment has improved β-cell function, as indicated by higher homeostasis model assessment of β-cell function (HOMA-β) and lower homeostasis model assessment of insulin resistance (HOMA-IR) in contradiction of placebo [150,151]. Additionally, curcumin has been reported to protect against diabetes-induced pathological changes in the aorta, primarily by inhibiting JNK2 and upregulating nuclear factor erythroid 2-related factor 2 (Nrf2) expression and function. Furthermore, a combination of metformin and curcumin provided greater myocardial protection in diabetic rats than metformin alone, seems to indicate that inhibition of the JAK/STAT pathway and activation of the Nrf2/heme oxygenase 1 (HO-1) pathway may negotiate these effects. Notably, the JAK/STAT pathway is also a known regulator of SASP secretion, highlighting a possible healing mechanism. Although there are no clinical studies yet examining curcumin's role in elimination of senescent cells in diabetic patients, further research is warranted [152-154].

## 3. FUTURE PERSPECTIVES

Over the past twenty years, research has significantly advanced our knowledge of cellular senescence in adipose tissue and pancreatic  $\beta$  cells related to diabetes, involving its morbid effect and impact on the growth and progression of the disease. Recent studies on senolysis have introduced the concept of senotherapy as a potential new treatment for diabetes. Since both aging and metabolic stress—such as insulin resistance—affect the onset, progression, and potential reversal of cellular senescence, individuals may experience varying degrees of senescence even at the same age [155-158]. Additionally, distinct indicators and molecular mechanisms of senescence have been identified in adipose tissue and pancreatic  $\beta$  cells, suggesting that the extent and impact of senescence on diabetes can vary between these two critical components of diabetes pathology. In adipose tissue, p21Cip1 serves as a marker of early-stage senescence and a primary target for senolysis to reduce insulin resistance. Conversely, targeting p16INK4a can enhance  $\beta$  cell function in pancreatic  $\beta$  cells, while targeting p21high cells has shown limited effects [159,160]. Therefore, a targeted approach that addresses both organs with specific molecular targets and therapeutic strategies could be a promising senotherapeutic method for diabetes. Further research into the molecular mechanisms of senescence in diabetes and the clinical efficacy and safety of senolytic agents is needed to make this concept a viable clinical treatment.

## 4. CONCLUSION

In conclusion, the intricate relationship between aging, cellular senescence, and type 2 diabetes mellitus (T2DM) reveals

the critical roles that these biological processes play in the pathogenesis of this widespread condition. As the prevalence of T2DM continues to rise globally, a deeper understanding of how aging influences insulin resistance and  $\beta$ -cell dysfunction is essential for developing effective therapeutic strategies. The accumulation of senescent cells in adipose tissue and pancreatic  $\beta$ -cells not only exacerbates metabolic dysregulation but also highlights the potential for senotherapy as a groundbreaking approach in diabetes management.

Emerging evidence supports the efficacy of senolytics and senomorphics in reducing the burden of senescent cells and mitigating the inflammatory responses they invoke. By targeting specific molecular pathways, these therapies have the potential to restore insulin sensitivity and enhance  $\beta$ -cell function, offering hope for improved outcomes in individuals with T2DM. This review emphasizes the need for continued research to elucidate the underlying mechanisms of cellular senescence and to evaluate the clinical efficacy and safety of these novel therapies. A targeted approach that addresses the unique senescence markers and mechanisms in both adipose tissue and pancreatic  $\beta$ -cells could significantly advance our ability to manage diabetes and its associated complications.

Ultimately, integrating insights from aging research with diabetes treatment strategies holds the promise of not only alleviating the immediate challenges posed by the disease but also improving long-term health outcomes for millions worldwide. As we move forward, a concerted effort to understand and leverage the biology of aging will be essential in shaping the future landscape of diabetes management.

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