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Abstract	<p>Studying age-related changes in the physiology, biochemistry, and molecular biology of isolated skin cell populations in culture has greatly expanded the understanding of the fundamental aspects of skin aging. The three main cell types that have been studied extensively with respect to cellular aging in vitro are dermal fibroblasts, epidermal keratinocytes, and melanocytes. Serial subcultivation of normal diploid skin cells can be performed only a limited number of times, and the emerging senescent phenotype can be categorized into structural, physiological, biochemical, and molecular phenotypes, which can be used as biomarkers of cellular aging in vitro. The rate and phenotype of aging are different in different cell types. There are both common features and specific features of aging of skin fibroblasts, keratinocytes, melanocytes, and other cell types. A progressive accumulation of damage in all types of macromolecules is a universal feature of cellular aging in all cell types. A progressive failure of molecular maintenance and repair pathways is the ultimate cause of cellular aging in vitro and in vivo.</p>	
Keywords (separated by “-”)	Aging - Senescence - Hayflick limit - Longevity - Survival - Cosmetics - Cosmeceuticals	

# Aging and Senescence of Skin Cells in Culture

Suresh I. S. Rattan

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## Abstract

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 57 progressive failure of molecular maintenance 58  
 59 and repair pathways is the ultimate cause of 60  
 61 cellular aging in vitro and in vivo. 62

## Introduction

40 In modern biogerontology, the terms “cellular 41  
 42 aging,” “cell senescence,” and “replicative senes- 43  
 44 cence” most commonly imply the study of normal 45

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44 diploid cells in culture, which during serial  
45 subcultivation undergo a multitude of changes cul-  
46 minating in the permanent cessation of cell division.  
47 This process of cellular aging in vitro is generally  
48 known as the Hayflick phenomenon, and the lim-  
49 ited division potential of normal cells is called the  
50 Hayflick limit, in recognition of the observations  
51 first reported by Leonard Hayflick in 1961  
52 [1–4]. The study of age-related changes in the phys-  
53 iology, biochemistry, and molecular biology of iso-  
54 lated skin cell populations in culture has greatly  
55 expanded the understanding of the fundamental  
56 aspects of skin aging. The three main cell types  
57 that have been studied extensively with respect to  
58 cellular aging in vitro are dermal fibroblasts, epi-  
59 dermal keratinocytes, and melanocytes [5–11].

AU2 60 The aim of this article is to describe the exper-  
61 imental system of aging of skin cells in culture; to  
62 provide an overview of the age-related changes in  
63 the structural and functional aspects of cells  
64 including physiological, biochemical, and molec-  
65 ular changes; and to evaluate the use of such a  
66 system in testing and developing effective inter-  
67 ventions for maintaining and/or re-achieving a  
68 healthy skin during aging.

## 69 **Experimental Model System of Cellular** 70 **Aging in Culture**

71 Once the primary culture of normal cells is  
72 established in culture from the normal tissue  
73 (e.g., a skin biopsy), by using any of the standard  
74 methods such as the explant growth and enzymic  
75 dissociation of cells, the primary culture can then  
76 be subcultivated repeatedly each time it becomes  
77 confluent. This repeated subculturing of cells is  
78 also known as serial passaging [4]. In a descrip-  
79 tion of the Hayflick phenomenon, Phase I is the  
80 period of the establishment of the primary culture  
81 from the normal tissue, Phase II is a relatively long  
82 period of serial passaging, growth, and cell pro-  
83 liferation at a constant rate, and Phase III is the  
84 final period of slowing-down of growth, which  
85 results in the cessation of cell division and end  
86 of replicative life span of cells. The whole

87 duration of serial passaging is considered as the 87  
88 process of cellular aging, and the end-stage irre- 88  
89 versible growth arrest in  $G_1$  is termed as replica- 89  
90 tive senescence [1, 2]. 90

91 After reaching a state of replicative senes- 91  
92 cence, some cells can still stay alive and be met- 92  
93 abolically active at a minimal level for sometime 93  
94 and generally resist undergoing apoptosis 94  
95 [3, 4]. Although the exact culturing conditions, 95  
96 such as the type of the culture medium, the source 96  
97 of growth factors, the use of antibiotics, and the 97  
98 incubation temperature, humidity, and gaseous 98  
99 composition, may vary for different cell types, 99  
100 serial subcultivation of normal diploid cells can 100  
101 be performed only a limited number of times. This 101  
102 is in contrast to the high proliferative capacity of 102  
103 transformed, cancerous, and immortalized cells, 103  
104 whose cultures can be subcultivated and 104  
105 maintained indefinitely [3, 4]. 105

106 The total number of cell divisions, measured as 106  
107 the cumulative population doublings (CPD), 107  
108 which can be achieved by a specific cell type 108  
109 in vitro, depends upon several biological factors. 109  
110 These include the maximum life span of the spe- 110  
111 cies, the developmental and adult age of the donor 111  
112 of the tissue biopsy, the site of the biopsy, and the 112  
113 health status of the donor [12]. For example, for 113  
114 human fibroblasts, the range of CPD for the cell 114  
115 strains originating from embryonic tissues is 115  
116 between 50 and 70, whereas for those originating 116  
117 from adult biopsies, it is generally less than 117  
118 50 CPD. A similar range for CPD attained by 118  
119 human keratinocytes and melanocytes has been 119  
120 reported [5–10]. Additionally, gaseous composi- 120  
121 tion, especially oxygen levels, and the quality of 121  
122 the nutritional serum and growth factors added to 122  
123 the culture medium can significantly affect the 123  
124 proliferative life span of cells in vitro. For exam- 124  
125 ple, culturing of fibroblasts in vitro in the air with 125  
126 about 20 % oxygen levels reduces their replicative 126  
127 life span which could be otherwise achieved at 127  
128 low-level (2 %) concentration akin to in vivo con- 128  
129 ditions [13, 14]. Furthermore, the site of the skin 129  
130 biopsy, for example, sun-exposed versus 130  
131 sun-protected area, has a significant effect on the 131  
132 CPD levels achieved by cells in culture [4, 15]. 132

133 **The Phenotype of Aging Skin Cells**

134 Serial passaging of normal diploid skin cells is  
 135 accompanied by a progressive and accumulative  
 136 occurrence of a wide variety of changes before the  
 137 final cessation of cell replication occurs. The  
 138 emerging senescent phenotype of serially pas-  
 139 saged normal diploid skin cells can be categorized  
 140 into the structural, physiological, biochemical,  
 141 and molecular phenotypes, which can be used as  
 142 biomarkers of cellular aging in vitro, as summa-  
 143 rized in Tables 1, 2, and 3. There are more than  
 144 200 such structural, physiological, biochemical,  
 145 and molecular characteristics that have been stud-  
 146 ied during cellular aging, and a list of major char-  
 147 acteristics that appear progressively in cell  
 148 cultures and distinguish between young and  
 149 senescent cells, generally before the end of prolif-  
 150 erative life span and their irreversible arrest in the  
 151 G<sub>1</sub> phase of the cell cycle, can be found in several  
 152 publications [3, 4]. Here a summary of such phe-  
 153 notypic changes is given below.

154 (A) *Structural phenotype of aging skin cells:*  
 155 Table 1 lists the major structural changes  
 156 observed in aging skin cells in culture. Most  
 157 commonly, a progressive increase in cell size  
 158 and the loss of homogenous morphological  
 159 pattern are the most dramatic and easily iden-  
 160 tifiable differences in early-passage young  
 161 and late-passage old or senescent skin cells  
 162 (Fig. 1).

163 Other structural changes during aging of  
 164 skin cells include cytoskeletal and membrane  
 165 rigidity, altered extracellular matrix, accumu-  
 166 lation of intracellular debris, and incomplete  
 167 cytokinesis leading to multi-nucleation.

168 In addition to the gross structural alter-  
 169 ations listed in Table 1, there are several  
 170 ultrastructural changes reported by using  
 171 electron microscopic methods. These include  
 172 the presence of distorted mitochondria,  
 173 increased level of chromosomal aberrations,  
 174 overcondensation of chromatin, increased  
 175 nucleolar fragmentation, and the accumula-  
 176 tion of lipid-protein conjugate lipofuscin in  
 177 lysosomes [16–18].

**Table 1** Structural phenotype of skin cells undergoing aging in vitro

Increased cell size
Change of shape from thin, long, and spindle-like to flattened and irregular
Loss of whorl-like arrangement in parallel arrays on the cell culture substrate
Rodlike polymerization of the cytoskeletal actin filaments and disorganized microtubules
Increased membrane rigidity
Increased multi-nucleation
Increased number of vacuoles and dense lysosomal autophagous bodies

**Table 2** Physiological phenotype of skin cells undergo- ing aging in vitro

Altered calcium flux, pH, viscosity, and membrane potential
Reduced activity of ionic pumps
Reduced mobility
Reduced respiration and energy production
Reduced response to growth factors and other mitogens
Increased sensitivity to toxins, drugs, irradiation, and other stresses
Increased basal levels of autophagy
Increased basal levels of stress proteins

**Table 3** Biochemical and molecular phenotype of skin cells undergoing aging in vitro

Permanent growth arrest in late G <sub>1</sub> phase of the cell cycle near the S phase boundary
Increased mRNA and protein levels of cell cycle inhibitors
Increased mRNA and protein levels of inhibitors of proteases
Decreased expression, levels, and activities of numerous housekeeping enzymes
Decreased expression, levels, and activities of macromolecular turnover pathways
Reduced levels of methylated cytosines in the DNA
Reduced length of telomeres
Increased levels of damage in nuclear and mitochondrial DNAs
Altered profiles of micro-RNAs
Increased levels of damaged and abnormal proteins
Increased levels of macromolecular cross-linking
Increased levels of reactive oxygen species
Altered profile of secreted proteins

**Fig. 1** Figure shows Giemsa-stained light microscopic phase contrast pictures of serially passaged human skin fibroblasts at various points in their in vitro life span. Sparse and confluent cultures at three stages during replicative life span are compared: (1) early-passage young

adult skin fibroblasts with less than 30 % life span completed, (2) middle-aged cells with 60–80 % replicative life span completed, and (3) late-passage senescent cells with more than 95 % life span completed

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178 (B) *Physiological phenotype of aging skin cells:*

179 Numerous studies have been performed elucidating changes in various functional and  
180 physiological parameters of skin cells undergoing aging. Table 2 lists some of the main  
181 such changes, which clearly indicate that almost all aspects of cellular function and  
182 physiology become impaired during aging. Collectively, these data show that aging skin  
183 cells progressively become less active, have reduced the ability to maintain various phys-  
184 iological functions, and become more prone to the negative effects of harmful substances.

185 Altered responsiveness of cells during aging is one of the most significant age-related changes, which can be a rate-  
186 limiting factor for the use of any potential modulators of aging. Several studies have  
187 been performed in order to understand the mechanisms for age-related alteration of  
188 responsiveness, and the pathways include unaltered receptor numbers and affinities,  
189 ineffective signal transduction, and interrupted networks [12]. Furthermore, increased levels of intracellular stress as  
190 manifested in increased basal levels of stress proteins and autophagy in serially passaged  
191 senescent skin cells are a hallmark of cellular aging [19].

192 (C) *Biochemical and molecular phenotype of aging skin cells:* At the biochemical and  
193 molecular levels, a large body of data is available which indicates that skin cells undergo a  
194 plethora of changes, which form the mechanistic bases of structural and physiological  
195 alterations. Table 3 gives a list of main categories of biochemical and molecular changes  
196 that have been reported in aging skin cells in culture.

217 Depending on the available technologies and  
218 the prevailing trends, changes in the amounts and

219 activities of thousands of proteins and in the levels of thousands of mRNAs have been reported for  
220 aging skin cells. Recently, data are beginning to be collected for age-related changes in the so-called  
221 epigenome, micro-RNAs, metabolome, secretome, and proteome, including posttranslational  
222 modifications [15, 20]. All such data will further strengthen the descriptive understanding  
223 of the phenomenon of aging of skin cells.

224 Although every single piece of descriptive data for aging skin cells is yet to be collected, a gener-  
225 alized picture of the aging phenomenon has emerged, Therefore, based on the large amount  
226 of data collected so far, important inferences and generalizations can already be made, which have  
227 implications with respect to developing effective interventions for a healthy skin. These are as  
228 follows:

- 229 1. The rate and phenotype of aging are different in different cell types. There are both common  
230 features and specific features of aging of skin fibroblasts, keratinocytes, melanocytes, and  
231 other cell types.
- 232 2. A progressive accumulation of damage in all types of macromolecules is a universal feature  
233 of cellular aging in all cell types.
- 234 3. Replicative senescence of cells in culture is not due to the activation of any aging-specific  
235 genes but is an indirect consequence of occurrence and accumulation of molecular damage  
236 and molecular heterogeneity.
- 237 4. A progressive failure of molecular maintenance and repair pathways is the ultimate  
238 cause of cellular aging.

253 **From Cellular Aging In Vitro**  
 254 **to Understanding Aging In Vivo**

255 The Hayflick system of aging of skin cells in  
 256 culture has proved to be very useful in developing  
 257 the cellular and molecular understanding of the  
 258 overall process of aging. A loss of proliferative  
 259 capacity of any of the cell types has a deteriorative  
 260 impact on the functioning and survival of the  
 261 entire organism. A loss or slowing-down of pro-  
 262 liferation of osteoblasts, glial cells, myoblasts,  
 263 epithelial cells, lymphocytes, and fibroblasts can  
 264 lead to the onset of many age-related diseases and  
 265 impairments including osteoporosis, arthritis,  
 266 immune deficiency, altered drug clearance,  
 267 delayed wound healing, and altered functioning  
 268 of the brain. Furthermore, the occurrence of fully  
 269 senescent or near-senescent heterogenous cells  
 270 in vivo can promote dysfunctioning of the other  
 271 tissues by producing harmful signals and can also  
 272 promote and stimulate the growth of other precancerous  
 273 and cancerous cells [21, 22].

274 However, the existence of the Hayflick-type  
 275 senescent cells in vivo is not very well established  
 276 so far. A commonly used biomarker of senescent  
 277 cells is the so-called senescence-associated beta-  
 278 galactosidase (SABG), which has been used to  
 279 demonstrate the presence of senescent cells in  
 280 the human skin and some other tissues [23]. How-  
 281 ever, there are several limitations regarding the  
 282 use of SABG as a marker of cellular aging  
 283 in vitro, since SABG can also be detected in  
 284 immortal cells under various conditions  
 285 [24]. More and multiple independent markers of  
 286 senescent cells are needed for this purpose.

287 The correlation between cellular aging in vitro  
 288 and in vivo is often based on the evidence gath-  
 289 ered from studies on the effects of donor age,  
 290 species life span, and premature aging syndromes  
 291 on cellular proliferative capacity in culture. These  
 292 studies indicate that the genetic and intrinsic  
 293 Hayflick limit of diploid cell strains in culture is  
 294 a true reflection of what is going on during aging  
 295 of an organism. However, there are some recent  
 296 critiques of this based on the replicative potential  
 297 of stem cells, which in the case of the skin appear  
 298 to be maintained throughout the life span [25,

299 26]. Similarly, some changes observed in the  
 300 Hayflick system in vitro, such as increased basal  
 301 levels of autophagy, may not be present in the  
 302 aging skin tissues [19]. In contrast to this, there  
 303 is evidence showing that the stem cell population  
 304 in the skin also undergoes aging and the number  
 305 of stem cells declines as a function of donor age  
 306 and during aging of the skin equivalents  
 307 in vitro [27].

---

**Modulators of Aging Skin Cells**

308  
 309 The Hayflick system of cellular aging in culture is  
 310 primarily a model for the study of slow and pro-  
 311 gressive accumulation of damage resulting in the  
 312 arrest of cells in a non-proliferative state [4]. This  
 313 system has been proved to be very useful for  
 314 testing various physical, chemical, and biological  
 315 conditions for their harmful or beneficial effects  
 316 and for understanding other aspects of cellular  
 317 aging with implications in the origin of  
 318 age-related diseases. For example, irradiation;  
 319 severe oxidative stress by UV, hydrogen peroxide,  
 320 or dicarbonyls; and gene transfection have been  
 321 used to induce a sudden and rapid increase in  
 322 molecular damage, resulting in premature appear-  
 323 ance of the senescent phenotype [28]. On the other  
 324 hand, insertion of catalytically active component  
 325 of the telomerase gene can completely bypass the  
 326 Hayflick limit in many cell types including skin  
 327 cells, and such cells can proliferate indefinitely  
 328 with or without becoming transformed [29]. Sim-  
 329 ilarly, normal diploid cells can be transformed and  
 330 immortalized by chemical carcinogens, irradiation,  
 331 and viral genes. Such approaches are helpful  
 332 for unraveling the molecular details of cell cycle  
 333 regulation in normal cells and its dysregulation in  
 334 cancer cells [29].

335 The Hayflick system of cellular aging in cul-  
 336 ture has also been very useful for testing various  
 337 natural and synthetic molecules as potential  
 338 antiaging and health-promoting compounds for  
 339 the skin. Some of the well-tested examples are  
 340 cytokinins kinetin and zeatin [30], a dipeptide  
 341 carnosine [31], curcumin which is a component  
 342 of the spice turmeric [32, 33], and extracts from  
 343 medicinal plants and sea algae [34]. Several of

344 these tests have resulted in the successful devel- 384  
 345 opment, production, and marketing of various 385  
 346 products with pharmaceutical, cosmeceutical, 386  
 347 and nutritional applications [34].

348 Another use of the model system of cellular 387  
 349 aging in culture has been to test the principle of 388  
 350 mild stress-induced beneficial and antiaging 389  
 351 effects, which is the phenomenon of hormesis 390  
 352 [35]. For example, human skin fibroblasts and 391  
 353 keratinocytes exposed to repeated mild heat stress 392  
 354 (41 °C, 1 h, twice a week) show several hormetic 393  
 355 effects, such as improved protein degradation 394  
 356 pathways, higher levels of chaperones, increased 395  
 357 resistance to other stresses, improved differentia- 396  
 358 tion, and increased proliferative life span 397  
 359 [10]. Such studies can form the basis of testing 398  
 360 novel hormetic agents, including potential 399  
 361 hormetins of natural or synthetic origin, for 400  
 362 improved skin care during aging [10, 35, 36].

## 363 Conclusion

364 It could be reemphasized that the present under- 401  
 365 standing of the cellular and molecular basis of 402  
 366 aging of the skin owes a lot to the use of the 403  
 367 Hayflick system of aging of skin cells in culture. 404  
 368 Most importantly, studies performed by using this 405  
 369 model system have demonstrated that aging of 406  
 370 cells is characterized by the accumulation of dam- 407  
 371 age in various molecules that results in the failure 408  
 372 of maintenance and repair systems. Detailed 409  
 373 genomic, proteomic, and metabolomic studies 410  
 374 using this system can further identify the 411  
 375 interacting networks of regulatory pathways, 412  
 376 which will then be accessible to modulation for 413  
 377 the maintenance of the structural and functional 414  
 378 integrity of the skin. 415

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