



REVIEW

Increased Arf/p53 activity in stem cells, aging and cancer

Estefania Carrasco-Garcia,^{1*} Manuel Moreno,^{1*}
Leire Moreno-Cugnon¹ and Ander Matheu^{1,2}

¹Cellular Oncology Group, Biodonostia Institute, San Sebastian, Spain

²Ikerbasque, Basque Foundation, Bilbao, Spain

Summary

Arf/p53 pathway protects the cells against DNA damage induced by acute stress. This characteristic is the responsible for its tumor suppressor activity. Moreover, it regulates the chronic type of stress associated with aging. This is the basis of its anti-aging activity. Indeed, increased gene dosage of Arf/p53 displays elongated longevity and delayed aging. At a cellular level, it has been recently shown that increased dosage of Arf/p53 delays age-associated stem cell exhaustion and the subsequent decline in tissue homeostasis and regeneration. However, p53 can also promote aging if constitutively activated. In this context, p53 reduces tissue regeneration, which correlates with premature exhaustion of stem cells. We discuss here the current evidence linking the Arf/p53 pathway to the processes of aging and cancer through stem cell regulation.

Key words: aging; ARF; p16; p53; stem cells.

Ink4/Arf/p53 activity in cancer

Cancer is the consequence of an aberrant gain of cellular fitness linked to the accumulation of stress and cellular damage of acute intensity. This damage occasionally provides aberrant advantages to certain cells, which can eventually lead to cancer development. The *Ink4/Arf* locus and *p53* are regarded as the most relevant tumor suppressors based on their ubiquitous and frequent inactivation in human cancer. The *Ink4/Arf* locus encodes three tumor suppressor genes *p15^{Ink4b}*, *p16^{Ink4a}*, and *p14^{Arf}* (*p19^{Arf}* in mice). On one hand, *p15^{Ink4b}* and *p16^{Ink4a}* (called *Ink4* hereafter) inhibit the formation of the cyclin-dependent kinases (CDK4 and CDK6) and cyclinD complexes during the G1 phase of the cell cycle. Hence, they prevent the transcription of genes involved in the transition to S phase, importantly the Rb/E2F1 pathway, so regulating cell cycle progression (Yaswen *et al.*, 2015). On the other hand, *Arf* exerts its tumor suppressive action by inhibiting *Mdm2*, a ubiquitin ligase considered the major *p53* regulator, thereby contributing to the activation and stabilization of *p53* (Matheu *et al.*, 2008).

The *Ink4/Rb* and *Arf/p53* pathways are major sensors of stress that play a crucial role in early detection and elimination of cells that have suffered different types of stress including oncogene activation, DNA damage, oxidative stress, etc. While the activation of *Ink4/Rb* pathway

induces reversible cell cycle arrest or irreversible cellular senescence-associated changes, the activation of *p53* elicits a cellular response that might vary from restoration of cellular homeostasis by a transient blockade of the cell cycle to allow for DNA repair, senescence, or apoptosis (Fig. 1). The activation of these responses depends in a complex manner, on the intensity of the triggering stress and on the cellular context. In agreement with this damage protective role, the individual or combinatory deletion of these genes promotes cancer susceptibility in multiple tissues and contexts. On the contrary, enhanced *Ink4/Arf* and *p53* activity preserves mice from spontaneous or chemically induced cancers (Garcia-Cao *et al.*, 2002; Tyner *et al.*, 2002; Maier *et al.*, 2004; Matheu *et al.*, 2004, 2007, 2009; Mendrysa *et al.*, 2006).

Ink4/Arf/p53 activity in aging

Aging is characterized by a loss of fitness, which results from the accumulation of cellular damage induced by chronic stress of small intensity. Moreover, there is clinical evidence that tumors have a higher incidence in aged organisms, which establishes a relationship between accumulated (likely pathogenic) cell damage, aging, and cancer development. Thus, although cancer and aging may seem opposite processes, they can be regarded as two different manifestations of the same underlying process, namely the accumulation of cellular damage. Moreover, cancer and aging may share common origins (Lopez-Otin *et al.*, 2013). There are several genetic or pharmacological manipulations that simultaneously modulate cancer and aging. For example, systemic downregulation of IGF1 signaling pathway by the overexpression of *PTEN* tumor suppressor increases longevity, delays aging, and confers cancer protection in mice (Garcia-Cao *et al.*, 2012; Ortega-Molina *et al.*, 2012). Similarly, reduced expression of *c-Myc* oncogene increases lifespan and shows resistance to several age-associated pathologies, such as osteoporosis, cardiac fibrosis and immunosenescence (Hofmann *et al.*, 2015). Caloric restriction also protects from cancer and aging (Lopez-Otin *et al.*, 2013), whereas metformin and rapamycin, two pharmacological compounds, which concomitantly extend longevity and impair cancer formation and growth (Blagosklonny, 2014). These proofs demonstrate that cancer protection and longevity can be simultaneously modulated using different strategies and molecular mechanisms.

In recent years, it is deepening the knowledge of the implications that the *Ink4/Rb* and *Arf/p53* pathways have on the management of cellular damage associated with the aging process. The observation that several manipulations simultaneously modulate longevity and cancer protection establishes an interesting parallel with the expression of members of the *Ink4/Rb* and *Arf/p53* pathways, which are silent or very low during development and postnatal life, while progressively increase from adulthood to old age in a broad range of tissues and species (Zindy *et al.*, 1997; Krishnamurthy *et al.*, 2004). Moreover, human genome-wide association studies have identified genetic variants in the *INK4/ARF* locus on 9p21.3 that confer increased risk of atherosclerotic vascular diseases such as stroke, aortic aneurysm or myocardial infarction, as well as to additional age-related diseases such as type 2 diabetes or glaucoma (Liu *et al.*, 2009; Jeck *et al.*, 2012). Similarly, there is evidence of the association of single nucleotide polymorphisms located on this genomic region to exceptional longevity in centenarians although with

Correspondence

Dr. Ander Matheu, Biodonostia Institute, Paseo Dr. Beguiristain s/n, 20014 San Sebastian, Spain. Tel.: (+34) 943006073; fax: (+34) 943006250;

e-mail: ander.matheu@biodonostia.org

*These authors have contributed equally and should be considered as co-first authors.

Accepted for publication 27 December 2016

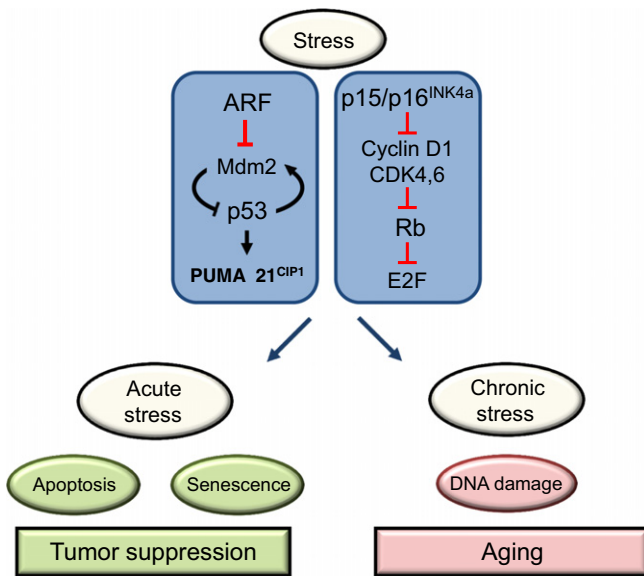


Fig. 1 *Ink4/Rb*- and *Arf/p53*-mediated responses to cellular stress. In response to stress, cells activate *Ink4/Rb* and *Arf/p53* signaling pathways in order to restore the homeostasis or eliminate themselves, preventing tumor formation in the case of acute oncogenic stress or alleviating accumulated cellular damage produced by chronic stress related to aging. p16^{INK4a} blocks the cell cycle through its inhibitory action on the CDK/Cyclin D complexes, finally avoiding the E2F transcriptional action and the progression of cells from G1 to S phase. Arf promotes the degradation of the negative p53 regulator Mdm2, inducing p53 activation and, in consequence, p21-mediated cell cycle arrest or senescence and/or PUMA-mediated apoptosis.

contradictory results (Pinos *et al.*, 2014; Congrains *et al.*, 2015). These data confirm that the response to stress mediated by the *Ink4/Arf* locus is involved in age-related pathologies. With regard to p53 in human aging, there is evidence that individuals carrying a common polymorphic variant of p53 (proline instead arginine in codon 72: p53-Pro72), that is more active inducing cell cycle arrest and senescence rather than in apoptosis, present increased lifespan (van Heemst *et al.*, 2005). Moreover, this polymorphism has implications in type 2 diabetes, wherein it is associated with a better metabolic status (Bonfigli *et al.*, 2013). There is little additional information regarding p53 and aging in human, yet there is no evidence of a pro-aging function. Indeed, it has been documented that p53 is not involved in human premature aging disorders such as Hutchinson–Gilford Progeria (O’Neill *et al.*, 2003), and it has been postulated that well-preserved p53-mediated responses are likely a key factor contributing to protection from diseases and cancer in centenarians (Salvioli *et al.*, 2009). In line with this, p53 anti-aging activity, rare alleles of two exon-derived SNPs of p21^{CIP}, well-established p53 downstream target, are significantly underrepresented among the centenarians (Gravina *et al.*, 2009). Previous studies have shown that the presence of these rare SNPs increases the susceptibility for the development of some types of cancer (Facher *et al.*, 1997; Li *et al.*, 2005). Thus, these p21^{CIP} alleles may be potentially detrimental to longevity and cancer protection.

The above raises the possibility that *Ink4/Rb* and *Arf/p53* pathways might have a role in aging. Thus, stress conditions cause an accumulation of DNA damage at the cellular level. Ultimately, it leads to the final activation of the *Ink4/Rb* and *Arf/p53* pathways in order to achieve various adaptive responses to this situation. Amidst such responses is the transient block of the cellular cycle to try to repair the damage, inducing

a state of senescence, or even apoptosis. Therefore, the empowerment of *Ink4/Rb* and *Arf/p53* pathways might play an important role not only on surveillance and suppression of tumors, but also on the accumulation of cellular damage and aging. Therefore, it is reasonable to surmise that *Ink4/Rb* and *Arf/p53* play a role also in the response to age-associated chronic stress and consequently affects aging. As activation of the *Ink4/Rb* and *Arf/p53* pathways triggers a protective mechanism against tumor-induced stresses, they could also have anti-aging activity by alleviating the load of age-associated damage (Fig. 1). However, the pleiotropic antagonism theory, which suggests that certain cellular processes that provide benefit in youth, may compromise organismal fitness in later life, postulates that tumor suppressors might have dual effects depending on the etiology of the cellular stress and the cellular and molecular context (Campisi, 2003).

The evaluation of the impact of the absence of *Ink4/Arf* locus and p53 on aging has not been studied nowadays, as null or heterozygous mice for these tumor suppressors develop several type of tumors at an early age (Donehower *et al.*, 1992; Serrano *et al.*, 1996), but it would be feasible in an inducible knockout or knockdown system. However, different mouse models with increased *Ink4/Arf* and p53 activity have been described in the last years, which allowed the understanding of the role of these tumor suppressors in aging and provided evidence supporting these two nondiscriminatory actions.

Deregulated increase in *Ink/Arf/p53* activity promotes premature aging

The pro-aging function of deregulated p53 activity has been documented in two independent mutant mouse models (Fig. 2). Tyner *et al.* (2002) developed a mutant mouse model called ‘*m*’ mice by deletion of the first six exons of the p53 gene. The mutant form consisting in a carboxy terminal p53 fragment of 24 KDa lacks the Mdm2-binding domain and, as a consequence, evades degradation by the proteasome and constitutively activates the endogenous wild-type p53 and downstream effectors. These mutant mice display higher resistance to tumor development compared to control animals but present a shorter lifespan and premature aging, including various diseases related to aging, such as osteoporosis and early and intense loss of cellularity and tissue mass (Tyner *et al.*, 2002), reflecting defects in organ homeostasis. In an additional study, Maier *et al.* (2004) developed another mouse model expressing a shorter p53 isoform lacking exons from 1 to 3 that produced a truncated and Mdm2-insensitive p53 protein of 44 KDa called p44tg. These mutant animals show increased resistance to spontaneous carcinogenesis but present a reduction in life expectancy and premature development of diseases such as osteoporosis and tissue atrophy. These effects are consequence of alterations in the signaling cascade of IGF affecting cell growth and proliferation. Moreover, they show a low rate of cellular replication resulting in small-sized individuals with a shortened breeding period, which is more pronounced in males that are sterile before reaching one year (Maier *et al.*, 2004).

Additional studies reinforce the pro-aging function of deregulated p53 activity. Cao *et al.* show that the deficient activity of *BRCA1* and the consequent DNA damage stress cause increased senescence and premature aging in mice. Interestingly, these consequences are directly dependent on a strong activation of p53, as inferred from the fact that blocking its main effector, p21^{CIP}, is enough to reverse the process (Cao *et al.*, 2003). In the same line, Varela *et al.* noted that another model, in which permanent cell stress is induced by abnormalities in the nuclear structure due to the deficiency of the metalloproteinase Zmpste24, exhibits cellular senescence and premature aging. Notably, these

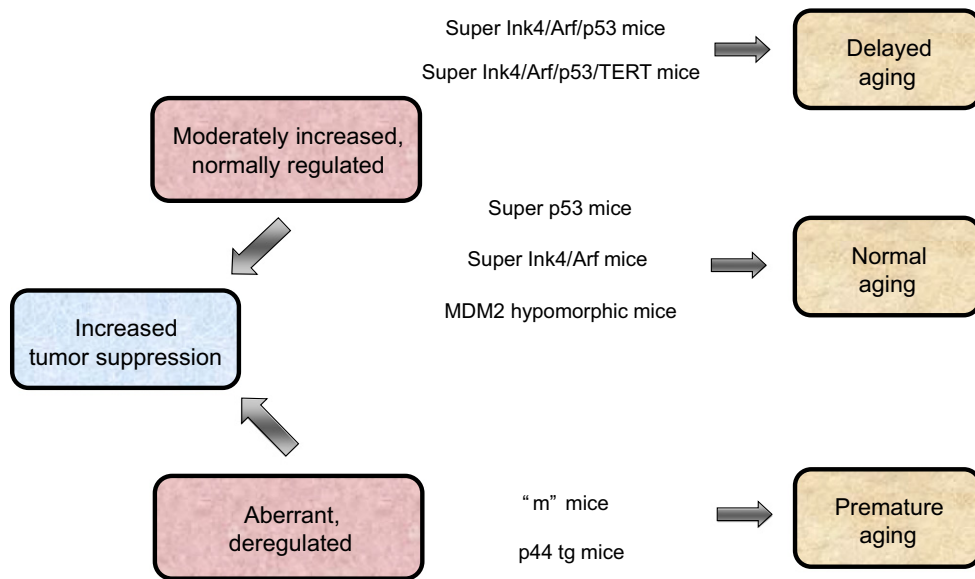


Fig. 2 Mouse models with regulated or deregulated *Ink4/Rb* and/or *Arf/p53* pathways and their phenotypes with regard to tumor prevention and aging. Mutant mice with deregulated p53 hyperactivity display reduced tumor development but present premature aging and aging-related diseases. Mice presenting a modest increase in normally regulated *Ink4/Arf* and p53 exhibit an enhanced tumor suppression capacity and normal aging. Mice presenting modest and regulated increases in both p53 and *Ink4/Arf* in the absence or presence of a constitutive telomerase reverse transcriptase (TERT) display resistance to tumor formation and present delayed aging. ‘m’ mice: Mice expressing a constitutively active truncated form of p53 lacking exons from 1 to 6 (Tyner *et al.*, 2002); p44 tg mice: Mice presenting a constitutively active truncated form of p53 lacking exons from 1 to 3 (Maier *et al.*, 2004), super p53 mice: Mice harboring an extra copy of the p53 intact gene (Garcia-Cao *et al.*, 2002); super *Ink4a/Arf* mice: mice with an extra intact *Ink4/Arf* locus (Matheu *et al.*, 2004); MDM2 hypomorphic mice: mice with reduced Mdm2 levels (Mendrysa *et al.*, 2006); ‘super Arf/p53’ mice: mice harboring an extra copy of p53 and *Ink4/Arf* (Matheu *et al.*, 2007); ‘super Arf/p53/TERT’ mice: super Arf/p53 mice additionally presenting constitutive telomerase reverse transcriptase (TERT) (Tomas-Loba *et al.*, 2008).

processes are driven by p53 hyperactivation and the phenotype was partially reversed in a p53-null genetic background (Varela *et al.*, 2005).

Normally regulated increase in *Ink4/Arf/p53* activity in aging

Different researchers have generated genetically manipulated mice with a modest increase in normally regulated *Ink4/Arf* and p53 (Fig. 2). These include mice carrying transgenic alleles encoding the intact p53 gene (Garcia-Cao *et al.*, 2002), the intact *Ink4/Arf* locus (Matheu *et al.*, 2004), or mutant mice with reduced *Mdm2* activity (Mendrysa *et al.*, 2006). In all these three models, the tumor suppressors retain their natural regulation, yet their levels are moderately increased. As example, those mice with additional regulated copies of p53 or *Mdm2* inactivation present enhanced apoptosis induction in the thymus, revealing the expected p53 augmented activity (Garcia-Cao *et al.*, 2002; Mendrysa *et al.*, 2006). Interestingly, all these models exhibit elevated tumor suppression potency, as are less prone to develop chemically or genetically induced sarcomas, adenomas or carcinomas, and present increased spontaneous cancer protection (Garcia-Cao *et al.*, 2002; Matheu *et al.*, 2004; Mendrysa *et al.*, 2006). Noteworthy, the proper regulation and moderate overexpression of the suppressors do not accelerate organismal aging and/or decrease longevity in any of the three models, providing a more potent and optimal response to chronic stress, along with cancer prevention, than p53 deregulated models. In agreement with the idea of enhanced and regulated p53 not affecting longevity and aging, the presence of the transgenic p53 or *Ink4/Arf* allele in mice deficient in telomerase and apolipoprotein E-null mice does not aggravate the accelerated aging or the atherosclerosis-prone characteristic phenotypes of these mice (Garcia-Cao *et al.*, 2006; Sanz-Gonzalez

et al., 2007; Fuster *et al.*, 2012). Similarly, iPS cells derived from *Ink4/Arf* or p53 transgenic mice do not present any alteration in their pluripotency, but they had limited their tumorigenicity (Menendez *et al.*, 2012). Notably, *Ink4/Arf* transgenic mice show a modest increase in their lifespan albeit not statistically significant (Matheu *et al.*, 2009). In line with this result, the presence of an extra copy of the *Ink4/Arf* protects against insulin resistance and glucose intolerance, two syndromes associated with aging (Sanz-Gonzalez *et al.*, 2007). These results indicate that normally regulated activation of the *Ink4/Rb* and *Arf/p53* pathways trigger a protective mechanism against tumor-induced stresses and show that these genes might also have anti-aging activity in some contexts by alleviating the load of age-associated damage.

Interestingly, the combined effects of modest and regulated increases in p53 and *Ink4/Arf* (*s-Ink4/Arf/p53*) or two extra copies of *Ink4/Arf* (*s-Ink4/Arf tg/tg*) result in a significantly elongated lifespan and delayed organismal aging (Matheu *et al.*, 2007, 2009). The longevity of the transgenic mice is still longer than wild-type mice when the survival analysis excludes the cases of cancer from both groups. Thereby, the *Ink4/Rb* and *Arf/p53* pathways not only prevent cancer, but also contribute to attenuate the deleterious effects of aging (Matheu *et al.*, 2007). Furthermore, these long-lived phenotypes of *s-Ink4/Arf/p53* mice are further potentiated in the presence of a constitutive telomerase reverse transcriptase (*s-TERT*) (Tomas-Loba *et al.*, 2008). In support of the anti-aging activity of p53, the elimination of p53 exacerbates the degenerative phenotype of different progeroid mouse models presenting alterations in telomere maintenance or DNA repair (Begus-Nahrman *et al.*, 2009; Murga *et al.*, 2009; Ruzankina *et al.*, 2009). Moreover, the inhibition of the p53-mediated apoptosis entails the formation of neoplasia and accelerates aging-associated signs and reduces lifespan in *ATM* mutant mice (Armata *et al.*, 2007).

We are beginning to understand the impact that the individual members of the Ink4/Rb and Arf/p53 pathways have on aging. Thus, the ablation of $p16^{\text{Ink4a}}$, but not $p19^{\text{Arf}}$, $p53$, or $p21^{\text{CIP}}$, alleviates the premature aging of mice with constitutively high levels of endogenous chromosome damage due to hypomorphic mutant alleles of *BubR1* (Baker *et al.*, 2008, 2013). These results reveal an anti-aging activity for the p53 pathway, likely attributable to the protection against aging-accumulated oxidative and DNA damage (Matheu *et al.*, 2007, 2009). Moreover, these works suggest that $p16^{\text{Ink4a}}$ might have pro-aging activity. Additional evidences indicate that $p16^{\text{Ink4a}}$ might contribute to promote aging through its function as senescence inductor. Indeed, it has been shown that the clearance of $p16^{\text{Ink4a}}$ positive cells from the organism ameliorates aging and shortens healthy lifespan in mice (Baker *et al.*, 2011, 2016). Furthermore, transgenic mice in which $p16^{\text{Ink4a}}$ is conditionally overexpressed display strong inhibition of proliferation of normal intestinal cells in young mice and premature induction of several features of aging such as reduced hair density, variable lightening of hair color, lower weight, and kyphosis (Boquoi *et al.*, 2015). However, this activity does not seem related to senescence as the induction of $p16^{\text{Ink4a}}$ did not detectably increase senescence-associated β -galactosidase staining in intestine and the de-induction of $p16^{\text{Ink4a}}$ revealed that the premature aging features were reversible (Boquoi *et al.*, 2015). Additional evidence supports a more complicated picture of the role of $p16^{\text{Ink4a}}$ on aging. Thus, the effects of a transgenic allele with constitutive and systemic overexpression of $p16^{\text{Ink4a}}$ are minimal in the pancreatic islets and in the brain (Krishnamurthy *et al.*, 2006; Molofsky *et al.*, 2006), while a luciferase knockin mouse model ($p16^{\text{LUCL}}$), which faithfully reports the expression of $p16^{\text{Ink4a}}$, revealed an exponential increase in luminescence with aging, but could not predict overall mortality and development of spontaneous malignancy (Burd *et al.*, 2013). Alternatively, it has been also postulated the possibility that $p16^{\text{Ink4a}}$ is indeed an anti-aging gene, through its capacity to slow down proliferation (Matheu *et al.*, 2009). Moreover, there are some integration and cross talk between Ink4/Rb and Arf/p53 pathways that might be important when studying their function in aging. Thus, Arf/p53 pathway collaborates with $p16^{\text{Ink4a}}$ to activate Rb through the induction of $p21^{\text{CIP}}$ expression (McConnell *et al.*, 1999; Mitra *et al.*, 1999). Moreover, inactivation of $p16^{\text{Ink4a}}$ /Rb pathway induces upregulation of $p14^{\text{Arf}}$ expression through the activation of E2F1 (Bates *et al.*, 1998), whereas inactivation of p53 causes compensatory upregulation of $p16^{\text{Ink4a}}$ expression (Leong *et al.*, 2009; Yamakoshi *et al.*, 2009).

Increased Arf / p53 activity in stem cells

The fundamental manifestation of aging is an overall decline in the functional capacity of organs to maintain tissue homeostasis and respond adequately to physiological needs under conditions of chronic stress. The regenerative potential of tissues relies on the activity of stem cells, becoming the maintenance of stem cell quiescence and self-renewal critical processes involved on tissue homeostasis. Thereby, stem cell exhaustion has been postulated as one of the 'Hallmarks of aging' (Lopez-Otin *et al.*, 2013). In line with this idea, stem cell rejuvenation is presented as a promising strategy to mitigate organismal aging. In the particular case of the central nervous system, the impairment of the brain is a common feature of normal aging, which coincides with a reduction in the number of neural stem/progenitor cells (NSCs), resulting in functional decline of the brain activity, cognitive impairment, and neurodegenerative diseases (Fuentetaja *et al.*, 2012).

The above raises the possibility that Ink4/Rb and Arf/p53 might play their role in aging through the modulation of the stem cell activity.

Indeed, p53 deregulated premature aging mouse models present decreased cell replacement in tissues, which has been linked to a premature depletion of stem cell niches. Thereby, the 'm' mouse model exhibits exhaustion of hematopoietic stem cells (HSCs) along with reduced hematopoiesis (Dumble *et al.*, 2007), and also disruption in mammary gland morphogenesis attributable to the depletion of stem cells (Gatza *et al.*, 2008). Similarly, adult $p44tg$ mice present a reduced number of NSCs and impoverished olfactory capacities (Medrano *et al.*, 2009). These mice also present an impaired replacement of beta cells in the pancreas and the consequent alteration of glucose homeostasis (Hinault *et al.*, 2011). Accordingly to these findings, mice with p53 overactivation in the epidermis by Mdm2 ablation show diminished epidermal stem cell activity and skin prematurely aged (Gannon *et al.*, 2011). These results indicate that deregulated p53 impacts on lifespan by a decline in tissue stem cell regenerative function likely as a consequence of premature stem cell exhaustion (Fig. 3).

On the contrary, the extended lifespan and delayed aging of mice with an extra copy of *Ink4a/Arf/p53* have been linked to an increased maintenance of stem cells pools. For example, *s-Ink4/Arf/p53* aged transgenic animals present greater preservation of the capacity of proliferation and self-renewal of NSCs compared to control littermates demonstrated by the relative increase in the number of neurospheres in culture and the higher expression of several markers of NSCs *in vivo* (Carrasco-Garcia *et al.*, 2015). This positive effect of extra *Ink4/Arf/p53* on the maintenance of NSCs directly impacts on mice behavior and neuronal activity, as revealed by enhanced functional activity in various behavioral tests such as neuromuscular coordination or cognitive performance (Carrasco-Garcia *et al.*, 2015). According with this evidence, *Ink4/Arf/p53* aged transgenic mice also display delayed epithelial stem cell exhaustion in the skin and improved hair regeneration compared to control mice (Matheu *et al.*, 2007; Tomas-Loba *et al.*, 2008). These results confirm that modest and regulated increases in Ink4/Arf/p53 tumor suppressors result in systemic organismal benefits ameliorating stem cell aging and maintaining tissue homeostasis (Fig. 3). In this case, the anti-aging activity of Arf/p53 is dependent on the conventional p53 pathway affecting its major role as regulator of cellular proliferation. Thus, moderate upregulation of the p53 pathway during aging likely results in slower proliferation rates, probably contributing to their quiescence and long-term maintenance and delaying stem cell exhaustion. In support of this idea, $p53$ or $p21^{\text{CIP}}$ inactivation results in loss of quiescence and exhaustion of long-term stem cells at advanced ages (Cheng *et al.*, 2000; Kippin *et al.*, 2005; Meletis *et al.*, 2006). Moreover, p53 also maintains the self-renewal of the nephron progenitor pool in the mouse kidney (Li *et al.*, 2015). With regard to the role of the $p16^{\text{Ink4a}}$, there is no clear conclusion. In one hand, it negatively affects the maintenance of stem cell pools in the brain, hematopoietic system, or pancreas (Janzen *et al.*, 2006; Krishnamurthy *et al.*, 2006; Molofsky *et al.*, 2006), and its inducible overexpression inhibits intestinal $Lgr5^+$ stem cell proliferation (Boquoi *et al.*, 2015), supporting an *Ink4/Arf* pro-aging activity. On the other hand, aged transgenic mice overexpressing specifically $p16^{\text{Ink4a}}$ do not present any significant deleterious impact in NSC or pancreatic progenitors (Krishnamurthy *et al.*, 2006; Molofsky *et al.*, 2006).

Ink4/Arf/p53 aged transgenic mice also present an increase in markers of neuroblasts (DCX and PSA-NCAM) and in markers of new neurons (NeuN) in the olfactory bulb and dentate gyrus (Carrasco-Garcia *et al.*, 2015). Similarly, deletion of $p53$ does not influence the HSC pool size in *Wip1* mutant mice but rescues their multilineage repopulation defect (Chen *et al.*, 2015), together indicating that p53 might also regulate stem cell aging modulating differentiation in a manner independent of

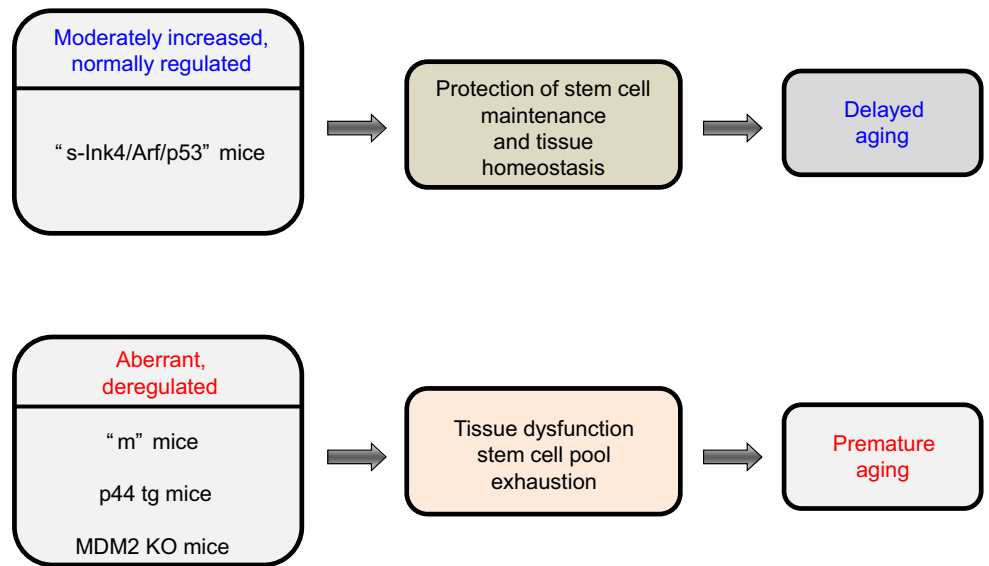


Fig. 3 Ink4/Rb and Arf/p53 pathways action on stem cell niches and aging modulation. Mice models with deregulated hyperactivation of p53 exhibit stem cell exhaustion along with tissue homeostasis disruption that promotes premature aging. However, mice with an extra copy of Ink4a/Arf/p53 display an improved maintenance of stem cells that provides tissue homeostasis delaying aging.

conventional p53 pathways. An additional mechanism to explain the function of p53 in stem cell aging derives from its activity to maintain the genome stability necessary to prevent the aging process and/or to eliminate the damaged cells. Thus, p53 deletion impairs clearance of chromosomal-unstable stem cells in aging telomere-dysfunctional mice (Begus-Nahrman *et al.*, 2009). This process is likely to be mediated by p21^{CIP} and Puma, p53 targets, which represent cooperating checkpoints limiting self-renewal and chromosomal instability of stem cells in response to telomere dysfunction (Sperka *et al.*, 2011). In support of this idea, the constitutive expression of *TERT*, concomitant with the extra *Ink4/Arf* and *p53* transgenes in mice, extends longevity, not only median but also maximum lifespan, together with a delayed exhaustion of epidermal stem cells (Tomas-Loba *et al.*, 2008). These results demonstrate that constitutive and regulated expression of p53 exerts anti-aging activity in the context of mammalian organisms maintaining stem cell function through different mechanisms.

Conclusion

Increasing evidence in humans and mice indicates that cancer and aging might be considered as two different manifestations of the accumulation of cellular damage, and both processes share common origins and molecular mechanisms. Among them, significant efforts have been made to determine the impact of Ink4/Rb and Arf/p53 tumor suppressor pathways on them. While their protective function against cancer is firmly established, their role in aging remains controversial.

In mice, it has been demonstrated that modest increases of regulated Arf/p53 activity are anti-aging while deregulated activation of p53 promotes aging. These observations are not in conflict per se and indicate that the activity of Arf/p53 could be beneficial or detrimental for aging depending on their intensity and regulation. Significant amount of data has recently demonstrated that these effects are mediated through the activity of stem cells concept of a reciprocal trade between tumor suppression, aging, and stem cell biology. Based on these data, we postulate a model by which high or deregulated Arf/p53 impacts on lifespan by a decline in tissue stem cell regenerative function, but modest and regulated increases in Arf/p53 result in systemic organismal benefits ameliorating stem cell aging and maintaining tissue homeostasis.

Additional work is necessary to establish the detail role and mechanism of action of p16^{Ink4a} in aging and stem cell biology.

Acknowledgments

L.M.-C. is recipient of a predoctoral fellowship from the Department of Education, University and Research of the Basque Government.

Funding info

This work is supported by grants from Instituto de Salud Carlos III and FEDER Funds (CP10/00539, PI13/02277), Diputacion Guipuzcoa (DFG16/002), and Marie Curie Career Integration Grant 2012/712404 to Ander Matheu.

Conflict of interest

None declared.

References

- Armata HL, Garlick DS, Sluss HK (2007) The ataxia telangiectasia-mutated target site Ser18 is required for p53-mediated tumor suppression. *Cancer Res.* **67**, 11696–11703.
- Baker DJ, Perez-Terzic C, Jin F, Pitel K, Niederlander NJ, Jeganathan K, Yamada S, Reyes S, Rowe L, Hiddinga HJ, Eberhardt NL, Terzic A, van Deursen JM (2008) Opposing roles for p16Ink4a and p19Arf in senescence and ageing caused by BubR1 insufficiency. *Nat. Cell Biol.* **10**, 825–836.
- Baker DJ, Wijshake T, Tchkonja T, LeBrasseur NK, Childs BG, van de Sluis B, Kirkland JL, van Deursen JM (2011) Clearance of p16Ink4a-positive senescent cells delays ageing-associated disorders. *Nature* **479**, 232–236.
- Baker DJ, Weaver RL, van Deursen JM (2013) p21 both attenuates and drives senescence and aging in BubR1 progeroid mice. *Cell Rep.* **3**, 1164–1174.
- Baker DJ, Childs BG, Durik M, Wijers ME, Sieben CJ, Zhong J, Saltness RA, Jeganathan KB, Verzosa GC, Pezeshki A, Khazaie K, Miller JD, van Deursen JM (2016) Naturally occurring p16(Ink4a)-positive cells shorten healthy lifespan. *Nature* **530**, 184–189.
- Bates S, Phillips AC, Clark PA, Stott F, Peters G, Ludwig RL, Vousden KH (1998) p14ARF links the tumour suppressors RB and p53. *Nature* **395**, 124–125.
- Begus-Nahrman Y, Lechel A, Obenauf AC, Nalapareddy K, Peit E, Hoffmann E, Schlaudraff F, Liss B, Schirmacher P, Kestler H, Danenberg E, Barker N, Clevers H,

- Speicher MR, Rudolph KL (2009) p53 deletion impairs clearance of chromosomal-instable stem cells in aging telomere-dysfunctional mice. *Nat. Genet.* **41**, 1138–1143.
- Blagosklonny MV (2014) Koschei the immortal and anti-aging drugs. *Cell Death Dis.* **5**, e1552.
- Bonfigli AR, Sirolla C, Testa R, Cucchi M, Spazzafumo L, Salvioli S, Cieriello A, Olivieri F, Festa R, Procopio AD, Brandoni G, Boemi M, Marra M, Franceschi C (2013) The p53 codon 72 (Arg72Pro) polymorphism is associated with the degree of insulin resistance in type 2 diabetic subjects: a cross-sectional study. *Acta Diabetol.* **50**, 429–436.
- Boquoi A, Arora S, Chen T, Litwin S, Koh J, Enders GH (2015) Reversible cell cycle inhibition and premature aging features imposed by conditional expression of p16Ink4a. *Aging Cell* **14**, 139–147.
- Burd CE, Sorrentino JA, Clark KS, Darr DB, Krishnamurthy J, Deal AM, Bardeesy N, Castrillon DH, Beach DH, Sharpless NE (2013) Monitoring tumorigenesis and senescence in vivo with a p16(INK4a)-luciferase model. *Cell* **152**, 340–351.
- Campisi J (2003) Cellular senescence and apoptosis: how cellular responses might influence aging phenotypes. *Exp. Gerontol.* **38**, 5–11.
- Cao L, Li W, Kim S, Brodie SG, Deng CX (2003) Senescence, aging, and malignant transformation mediated by p53 in mice lacking the Brca1 full-length isoform. *Genes Dev.* **17**, 201–213.
- Carrasco-Garcia E, Arrizabalaga O, Serrano M, Lovell-Badge R, Matheu A (2015) Increased gene dosage of Ink4/Arf and p53 delays age-associated central nervous system functional decline. *Aging Cell* **14**, 710–714.
- Chen Z, Yi W, Morita Y, Wang H, Cong Y, Liu JP, Xiao Z, Rudolph KL, Cheng T, Ju Z (2015) Wip1 deficiency impairs haematopoietic stem cell function via p53 and mTORC1 pathways. *Nat. Commun.* **6**, 6808.
- Cheng T, Rodrigues N, Shen H, Yang Y, Dombkowski D, Sykes M, Scadden DT (2000) Hematopoietic stem cell quiescence maintained by p21cip1/waf1. *Science* **287**, 1804–1808.
- Congrains A, Kamide K, Hirose N, Arai Y, Oguro R, Nakama C, Imaizumi Y, Kawai T, Kusunoki H, Yamamoto H, Onishi-Takeya M, Takeya Y, Yamamoto K, Sugimoto K, Akasaka H, Saitoh S, Miura T, Awata N, Kato N, Katsuya T, Ikebe K, Gondo Y, Rakugi H (2015) Disease-associated polymorphisms in 9p21 are not associated with extreme longevity. *Geriatr. Gerontol. Int.* **15**, 797–803.
- Donehower LA, Harvey M, Slagle BL, McArthur MJ, Montgomery CA Jr, Butel JS, Bradley A (1992) Mice deficient for p53 are developmentally normal but susceptible to spontaneous tumours. *Nature* **356**, 215–221.
- Dumble M, Moore L, Chambers SM, Geiger H, Van Zant G, Goodell MA, Donehower LA (2007) The impact of altered p53 dosage on hematopoietic stem cell dynamics during aging. *Blood* **109**, 1736–1742.
- Facher EA, Becich MJ, Deka A, Law JC (1997) Association between human cancer and two polymorphisms occurring together in the p21Waf1/Cip1 cyclin-dependent kinase inhibitor gene. *Cancer* **79**, 2424–2429.
- Fuentealba LC, Obner K, Alvarez-Buylla A (2012) Adult neural stem cells bridge their niche. *Cell Stem Cell* **10**, 698–708.
- Fuster JJ, Molina-Sanchez P, Jovani D, Vinue A, Serrano M, Andres V (2012) Increased gene dosage of the Ink4/Arf locus does not attenuate atherosclerosis development in hypercholesterolaemic mice. *Atherosclerosis* **221**, 98–105.
- Gannon HS, Donehower LA, Lyle S, Jones SN (2011) Mdm2-p53 signaling regulates epidermal stem cell senescence and premature aging phenotypes in mouse skin. *Dev. Biol.* **353**, 1–9.
- Garcia-Cao I, Garcia-Cao M, Martin-Caballero J, Criado LM, Klatt P, Flores JM, Weill JC, Blasco MA, Serrano M (2002) “Super p53” mice exhibit enhanced DNA damage response, are tumor resistant and age normally. *EMBO J.* **21**, 6225–6235.
- Garcia-Cao I, Garcia-Cao M, Tomas-Loba A, Martin-Caballero J, Flores JM, Klatt P, Blasco MA, Serrano M (2006) Increased p53 activity does not accelerate telomere-driven ageing. *EMBO Rep.* **7**, 546–552.
- Garcia-Cao I, Song MS, Hobbs RM, Laurent G, Giorgi C, de Boer VC, Anastasiou D, Ito K, Sasaki AT, Rameh L, Carracedo A, Vander Heiden MG, Cantley LC, Pinton P, Haigis MC, Pandolfi PP (2012) Systemic elevation of PTEN induces a tumor-suppressive metabolic state. *Cell* **149**, 49–62.
- Gatza CE, Dumble M, Kittrell F, Edwards DG, Dearth RK, Lee AV, Xu J, Medina D, Donehower LA (2008) Altered mammary gland development in the p53+/m mouse, a model of accelerated aging. *Dev. Biol.* **313**, 130–141.
- Gravina S, Lescai F, Hurteau G, Brock GJ, Saramaki A, Salvioli S, Franceschi C, Roninson IB (2009) Identification of single nucleotide polymorphisms in the p21 (CDKN1A) gene and correlations with longevity in the Italian population. *Aging (Albany NY)* **1**, 470–480.
- van Heemst D, Mooijaart SP, Beekman M, Schreuder J, de Craen AJ, Brandt BW, Slagboom PE, Westendorp RG (2005) Variation in the human TP53 gene affects old age survival and cancer mortality. *Exp. Gerontol.* **40**, 11–15.
- Hinault C, Kawamori D, Liew CW, Maier B, Hu J, Keller SR, Mirmira RG, Scrabble H, Kulkarni RN (2011) Delta40 Isoform of p53 controls beta-cell proliferation and glucose homeostasis in mice. *Diabetes* **60**, 1210–1222.
- Hofmann JW, Zhao X, De Cecco M, Peterson AL, Pagliaroli L, Manivannan J, Hubbard GB, Ikeno Y, Zhang Y, Feng B, Li X, Serre T, Qi W, Van Remmen H, Miller RA, Bath KG, de Cabo R, Xu H, Neretti N, Sedivy JM (2015) Reduced expression of MYC increases longevity and enhances healthspan. *Cell* **160**, 477–488.
- Janzen V, Forkert R, Fleming HE, Saito Y, Waring MT, Dombkowski DM, Cheng T, DePinho RA, Sharpless NE, Scadden DT (2006) Stem-cell ageing modified by the cyclin-dependent kinase inhibitor p16INK4a. *Nature* **443**, 421–426.
- Jeck WR, Siebold AP, Sharpless NE (2012) Review: a meta-analysis of GWAS and age-associated diseases. *Aging Cell* **11**, 727–731.
- Kippin TE, Martens DJ, van der Kooy D (2005) p21 loss compromises the relative quiescence of forebrain stem cell proliferation leading to exhaustion of their proliferation capacity. *Genes Dev.* **19**, 756–767.
- Krishnamurthy J, Torrice C, Ramsey MR, Kovalev GI, Al-Regaiey K, Su L, Sharpless NE (2004) Ink4a/Arf expression is a biomarker of aging. *J. Clin. Invest.* **114**, 1299–1307.
- Krishnamurthy J, Ramsey MR, Ligon KL, Torrice C, Koh A, Bonner-Weir S, Sharpless NE (2006) p16INK4a induces an age-dependent decline in islet regenerative potential. *Nature* **443**, 453–457.
- Leong WF, Chau JF, Li B (2009) p53 Deficiency leads to compensatory up-regulation of p16INK4a. *Mol. Cancer Res.* **7**, 354–360.
- Li G, Liu Z, Sturgis EM, Shi Q, Chamberlain RM, Spitz MR, Wei Q (2005) Genetic polymorphisms of p21 are associated with risk of squamous cell carcinoma of the head and neck. *Carcinogenesis* **26**, 1596–1602.
- Li Y, Liu J, Li W, Brown A, Baddoo M, Li M, Carroll T, Oxburgh L, Feng Y, Saifudeen Z (2015) p53 Enables metabolic fitness and self-renewal of nephron progenitor cells. *Development* **142**, 1228–1241.
- Liu Y, Sanoff HK, Cho H, Burd CE, Torrice C, Mohlke KL, Ibrahim JG, Thomas NE, Sharpless NE (2009) INK4/ARF transcript expression is associated with chromosome 9p21 variants linked to atherosclerosis. *PLoS ONE* **4**, e5027.
- Lopez-Otin C, Blasco MA, Partridge L, Serrano M, Kroemer G (2013) The hallmarks of aging. *Cell* **153**, 1194–1217.
- Maier B, Gluba W, Bernier B, Turner T, Mohammad K, Guise T, Sutherland A, Thorner M, Scrabble H (2004) Modulation of mammalian life span by the short isoform of p53. *Genes Dev.* **18**, 306–319.
- Matheu A, Pantoja C, Efeyan A, Criado LM, Martin-Caballero J, Flores JM, Klatt P, Serrano M (2004) Increased gene dosage of Ink4a/Arf results in cancer resistance and normal aging. *Genes Dev.* **18**, 2736–2746.
- Matheu A, Maraver A, Klatt P, Flores I, Garcia-Cao I, Borrás C, Flores JM, Vina J, Blasco MA, Serrano M (2007) Delayed ageing through damage protection by the Arf/p53 pathway. *Nature* **448**, 375–379.
- Matheu A, Maraver A, Serrano M (2008) The Arf/p53 pathway in cancer and aging. *Cancer Res.* **68**, 6031–6034.
- Matheu A, Maraver A, Collado M, Garcia-Cao I, Canamero M, Borrás C, Flores JM, Klatt P, Vina J, Serrano M (2009) Anti-aging activity of the Ink4/Arf locus. *Aging Cell* **8**, 152–161.
- McConnell BB, Gregory FJ, Stott FJ, Hara E, Peters G (1999) Induced expression of p16(INK4a) inhibits both CDK4- and CDK2-associated kinase activity by reassembly of cyclin-CDK-inhibitor complexes. *Mol. Cell. Biol.* **19**, 1981–1989.
- Medrano S, Burns-Cusato M, Atienza MB, Rahimi D, Scrabble H (2009) Regenerative capacity of neural precursors in the adult mammalian brain is under the control of p53. *Neurobiol. Aging* **30**, 483–497.
- Meletis K, Wirta V, Hede SM, Nister M, Lundeberg J, Frisen J (2006) p53 suppresses the self-renewal of adult neural stem cells. *Development* **133**, 363–369.
- Mendrysa SM, O’Leary KA, McElwee MK, Michalowski J, Eisenman RN, Powell DA, Perry ME (2006) Tumor suppression and normal aging in mice with constitutively high p53 activity. *Genes Dev.* **20**, 16–21.
- Menendez S, Camus S, Herreria A, Paramonov I, Morera LB, Collado M, Pekarik V, Maceda I, Edel M, Consiglio A, Sanchez A, Li H, Serrano M, Belmonte JC (2012) Increased dosage of tumor suppressors limits the tumorigenicity of iPSC cells without affecting their pluripotency. *Aging Cell* **11**, 41–50.
- Mitra J, Dai CY, Somasundaram K, El-Deiry WS, Satyamoorthy K, Herlyn M, Enders GH (1999) Induction of p21(WAF1/CIP1) and inhibition of Cdk2 mediated by the tumor suppressor p16(INK4a). *Mol. Cell. Biol.* **19**, 3916–3928.
- Molofsky AV, Slutsky SG, Joseph NM, He S, Pardoll R, Krishnamurthy J, Sharpless NE, Morrison SJ (2006) Increasing p16INK4a expression decreases forebrain progenitors and neurogenesis during ageing. *Nature* **443**, 448–452.
- Murga M, Bunting S, Montana MF, Soria R, Mulero F, Canamero M, Lee Y, McKinnon PJ, Nussenzweig A, Fernandez-Capetillo O (2009) A mouse model of

- ATR-Seckel shows embryonic replicative stress and accelerated aging. *Nat. Genet.* **41**, 891–898.
- O'Neill M, Nunez F, Melton DW (2003) p53 and a human premature ageing disorder. *Mech. Ageing Dev.* **124**, 599–603.
- Ortega-Molina A, Efeyan A, Lopez-Guadamillas E, Munoz-Martin M, Gomez-Lopez G, Canamero M, Mulero F, Pastor J, Martinez S, Romanos E, Mar Gonzalez-Barroso M, Rial E, Valverde AM, Bischoff JR, Serrano M (2012) Pten positively regulates brown adipose function, energy expenditure, and longevity. *Cell Metab.* **15**, 382–394.
- Pinos T, Fuku N, Camara Y, Arai Y, Abe Y, Rodriguez-Romo G, Garatachea N, Santos-Lozano A, Miro-Casas E, Ruiz-Meana M, Otaegui I, Murakami H, Miyachi M, Garcia-Dorado D, Hinohara K, Andreu AL, Kimura A, Hirose N, Lucia A (2014) The rs1333049 polymorphism on locus 9p21.3 and extreme longevity in Spanish and Japanese cohorts. *Age (Dordr)* **36**, 933–943.
- Ruzankina Y, Schoppy DW, Asare A, Clark CE, Vonderheide RH, Brown EJ (2009) Tissue regenerative delays and synthetic lethality in adult mice after combined deletion of Atr and Trp53. *Nat. Genet.* **41**, 1144–1149.
- Salvioli S, Capri M, Bucci L, Lanni C, Racchi M, Uberti D, Memo M, Mari D, Govoni S, Franceschi C (2009) Why do centenarians escape or postpone cancer? The role of IGF-1, inflammation and p53. *Cancer Immunol. Immunother.* **58**, 1909–1917.
- Sanz-Gonzalez SM, Barquin L, Garcia-Cao I, Roque M, Gonzalez JM, Fuster JJ, Castells MT, Flores JM, Serrano M, Andres V (2007) Increased p53 gene dosage reduces neointimal thickening induced by mechanical injury but has no effect on native atherosclerosis. *Cardiovasc. Res.* **75**, 803–812.
- Serrano M, Lee H, Chin L, Cordon-Cardo C, Beach D, DePinho RA (1996) Role of the INK4a locus in tumor suppression and cell mortality. *Cell* **85**, 27–37.
- Sperka T, Song Z, Morita Y, Nalapareddy K, Guachalla LM, Lechel A, Begus-Nahrman Y, Burkhalter MD, Mach M, Schlaudraff F, Liss B, Ju Z, Speicher MR, Rudolph KL (2011) Puma and p21 represent cooperating checkpoints limiting self-renewal and chromosomal instability of somatic stem cells in response to telomere dysfunction. *Nat. Cell Biol.* **14**, 73–79.
- Tomas-Loba A, Flores I, Fernandez-Marcos PJ, Cayuela ML, Maraver A, Tejera A, Borrás C, Matheu A, Klatt P, Flores JM, Vina J, Serrano M, Blasco MA (2008) Telomerase reverse transcriptase delays aging in cancer-resistant mice. *Cell* **135**, 609–622.
- Tyner SD, Venkatachalam S, Choi J, Jones S, Ghebranious N, Igelmann H, Lu X, Soron G, Cooper B, Brayton C, Hee Park S, Thompson T, Karsenty G, Bradley A, Donehower LA (2002) p53 mutant mice that display early ageing-associated phenotypes. *Nature* **415**, 45–53.
- Varela I, Cadinanos J, Pendas AM, Gutierrez-Fernandez A, Folgueras AR, Sanchez LM, Zhou Z, Rodriguez FJ, Stewart CL, Vega JA, Tryggvason K, Freije JM, Lopez-Otin C (2005) Accelerated ageing in mice deficient in Zmpste24 protease is linked to p53 signalling activation. *Nature* **437**, 564–568.
- Yamakoshi K, Takahashi A, Hirota F, Nakayama R, Ishimaru N, Kubo Y, Mann DJ, Ohmura M, Hirao A, Saya H, Arase S, Hayashi Y, Nakao K, Matsumoto M, Ohtani N, Hara E (2009) Real-time in vivo imaging of p16Ink4a reveals cross talk with p53. *J. Cell Biol.* **186**, 393–407.
- Yaswen P, MacKenzie KL, Keith WN, Hentosh P, Rodier F, Zhu J, Firestone GL, Matheu A, Carnero A, Bilsland A, Sundin T, Honoki K, Fujii H, Georgakilas AG, Amedei A, Amin A, Helferich B, Boosani CS, Guha G, Ciriolo MR, Chen S, Mohammed SI, Azmi AS, Bhakta D, Halicka D, Niccolai E, Aquilano K, Ashraf SS, Newsheem S, Yang X (2015) Therapeutic targeting of replicative immortality. *Semin. Cancer Biol.* **35**(Suppl), S104–S128.
- Zindy F, Quelle DE, Roussel MF, Sherr CJ (1997) Expression of the p16INK4a tumor suppressor versus other INK4 family members during mouse development and aging. *Oncogene* **15**, 203–211.