

Cell senescence and hypermitogenic arrest

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A diverse range of conditions, from mitogenic stimuli to cytotoxic stress, can induce cell senescence. Here, I propose that simultaneous stimulation of mitogen-activated pathways and downstream inhibition of cyclin-dependent kinases leads, ultimately, to cell senescence. This model distinguishes between two types of growth arrest: first, exit to G0 phase, which is caused by the withdrawal of mitogens and can lead to apoptosis; and second, hypermitogenic arrest, which is stimulated by mitogens and can lead to senescence. The concept of hypermitogenic arrest defines cell senescence as a functionally active, stable and conditionally reversible state.

EMBO reports 4, 358–362 (2003)

doi:10.1038/sj.embor.embor806

Introduction

Cell senescence has been defined as a form of replicative cell death or, in other words, a state in which proliferation is irreversibly arrested (Mathon & Lloyd, 2001). Nevertheless, senescent cells are metabolically active and can be maintained in culture for several years. Senescence may result from telomere shortening, which occurs during each cell cycle in some cell types, but senescence does not occur in others unless they are cultured under inadequate conditions (Mathon *et al.*, 2001; Tang *et al.*, 2001). These states are referred to as spontaneous and premature senescence, respectively, and some excellent recent reviews discuss their common and divergent features (for example, see Serrano & Blasco, 2001; Hanh, 2002). It has been suggested that premature senescence is triggered by the cumulative trauma of culturing cells *in vitro* (Sherr & DePinho, 2000), and also by various mitogenic, oncogenic, oxidative, cytostatic and toxic compounds, including DNA-damaging agents (McConnell *et al.*, 1998; Lundberg *et al.*, 2000; Roninson *et al.*, 2001). What do the mitogenic and anti-mitogenic stimuli listed above have in common? It seems that two simultaneous events are required and sufficient to cause senescence. First, strong stimulation of mitogen-activated pathways is required. Second, cyclin-dependent kinases (CDKs) must be blocked, either directly or through the induction of CDK inhibitors (CDKIs).

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Submitted 23 October 2002; accepted 30 January 2003

Senescence-inducing 'mitogens' inhibit CDKs

In normal cells, mitogens (growth factors) initiate and maintain the transition from G1 phase to S phase in the cell cycle. Mitogen-activated protein kinase (MAPK) signalling pathways induce cyclin D1, which results in the activation of either CDK4 or CDK6 (henceforth referred to as CDK4/6), after which the cell no longer requires mitogens to complete the cell cycle. This point in late G1 phase is known as the 'restriction point' (Pardee, 1974). CDK4/6 phosphorylates the retinoblastoma protein (Rb), causing the release of E2F. This in turn transactivates cyclin E, which then activates CDK2. The upstream, mitogen-activated pathways thus stimulate the downstream cell-cycle machinery by inducing cyclins, which are the activators of CDKs. However, MAPK pathways can also induce CDKIs, offering the cell two possibilities: proliferation or growth arrest (Fig. 1). For example, the classical MAPK signalling cascade involves the sequential activation of Ras, Raf1, MAPK kinase (MEK) and extracellular-signal-regulated kinase (ERK), which stimulates activators (such as cyclin D) and inhibitors (such as p21, p16, p15 and p57) of CDKs (Marshall, 1995; Sewing *et al.*, 1997; Woods *et al.*, 1997; Chang *et al.*, 2002). The same is true of MAPK pathways that act through the JNK (Jun kinase) and p38 kinases. Furthermore, both p21 and p27 can have opposing effects on CDK4/6 and CDK2 (Sherr & Roberts, 1999). What determines which of the two options the cell takes? It has been suggested that it is the strength or duration of the signal that is important; strong and/or sustained activation of the MAPK pathways arrests the cell cycle, whereas transient activation induces cell-cycle progression (Marshall, 1995). In support of this, low levels of Raf1 activity induce cyclin D1 and therefore proliferation, whereas high levels lead to p21 induction and growth arrest (Sewing *et al.*, 1997; Woods *et al.*, 1997). In addition, the simple explanation—that the proliferative status of the cell predetermines its response—should not be overlooked. Whereas a 'mitogen' may arrest a cycling cell or stimulate G0–G1 phase progression in a resting cell, it cannot possibly arrest a cell that is already resting. Thus, the overall cellular response may be predetermined by whether resting or cycling cells are targeted.

Regardless of what determines the choice between growth arrest and proliferation, cell senescence occurs only when mitogenic stimuli lead to CDK inhibition. In primary cells, Ras and the downstream MAPK pathways can induce senescence due to the induction of CDKIs (Missero *et al.*, 1996; Serrano *et al.*, 1997; Lin *et al.*, 1998; Zhu

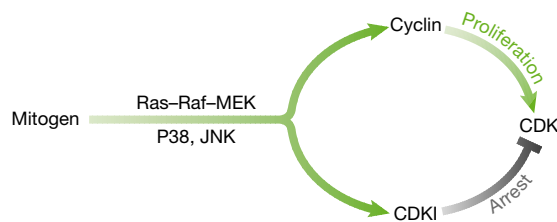


Fig. 1 | Dual mitogenic signalling. Mitogens simultaneously induce activators (such as cyclin D1) and cyclin-dependent-kinase inhibitors (CDKIs, such as p21, p16 and p15) through mitogen-activated pathways (Raf1/mitogen-activated protein kinase kinase (MEK)/extracellular-signal-related kinase (ERK), p38 and JNK (Jun kinase)), leading ultimately to cellular proliferation or arrest.

et al., 1998; Malumbres *et al.*, 2000; Wang *et al.*, 2002; Brookes *et al.*, 2002). All CDKIs induce senescence when ectopically expressed in fibroblasts (McConnell *et al.*, 1998). Either the overexpression of positive regulators acting downstream of cyclin D1 (for example, CDK4/6, E2F1 and c-Myc) or the inactivation of tumour suppressors (such as Rb, p53 and p16) can block Ras-induced senescence. Even spontaneous senescence can be delayed by the overexpression of CDK4/6 (Morris *et al.*, 2002; Holland *et al.*, 1998). We can therefore conclude that the inhibition of pathways, either at the level of CDK4/6 or downstream of CDK4/6, is essential for all forms of senescence.

Senescence-inducing cytostatic stress

Ionizing radiation, DNA-damaging drugs, the p53 tumour suppressor, microtubule-active drugs (such as Taxol), oxidative stress and hypoxia-mimicking iron chelators, inhibitors of histone acetylase, transforming growth factor- β (TGF- β) and retinoids are all able to trigger premature cell senescence (McConnell *et al.*, 1998; Chang *et al.*, 1999; Roninson *et al.*, 2001; Terao *et al.*, 2001; Itahana *et al.*, 2001). Most of these agents (especially in high doses) can also induce apoptosis, particularly in cells with 'ready-to-run' apoptotic machinery (that is, caspases). In such cells, the inhibition of apoptosis may allow cytostatic stress to induce senescence instead. For example, by inhibiting apoptosis, Bcl2 can promote p53-dependent senescence (Rincheval *et al.*, 2002). All anti-mitogens directly or indirectly inhibit CDKs, causing G1-phase arrest and/or G2-phase arrest. For example, DNA-damaging agents cause the induction of CDKIs (such as p21) and changes in the phosphorylation states of CDKs and, thereby, their inhibition. Nevertheless, MAPKs are not inhibited, and these agents activate mitogenic signalling. Radiation stimulates Ras and numerous mitogenic kinases, including Raf1 and Akt (Kasid *et al.*, 1996; Liu *et al.*, 1996; Shaulian *et al.*, 2000; Lee *et al.*, 2000; Fang *et al.*, 2001). Furthermore, p53 can induce growth factors that activate MAPK and Akt signalling, hypoxia causes autocrine secretion of mitogens and activation of mitogenic kinases, and DNA damage and hypoxia may, paradoxically, induce proliferation (Ishii *et al.*, 1995; Adelman *et al.*, 2000; Fang *et al.*, 2001; Das *et al.*, 2001; Salnikow *et al.*, 2002). This suggests that, while inhibiting CDKs, senescence-inducing cytostatic stress may actually activate mitogenic pathways.

As is the case for premature senescence, spontaneous senescence occurs in the presence of mitogens. In the latter case, senescence is caused by the stress due to telomere shortening as the cells proliferate. In mitogen-induced senescence, hyperactive MAPK pathways inhibit CDKs. The elegance of cell senescence is that one stimulus can both activate upstream mitogenic signalling and arrest the cell cycle downstream.

Upstream and downstream signalling

It is assumed that cellular senescence evolved to suppress tumorigenesis (Lin *et al.*, 1998; Campisi, 2001); that is, normal cells can respond to potentially oncogenic stimuli by entering the senescent state. This is reminiscent of the conflicting signal model, in which 'unscheduled' activation of potentially oncogenic stimuli leads to apoptosis, "ensuring that cell growth is restricted to the correct paracrine environment" (Raff, 1992; Evan & Littlewood, 1998). Does this mean that potentially oncogenic stimuli can cause both apoptosis and senescence? If so, how are the two outcomes determined? Here, we view them in the light of the upstream and downstream signalling pathways. During normal proliferation, the upstream mitogen-activated pathways stimulate downstream cell-cycle signalling (Fig. 2A). In the absence of upstream mitogenic signalling, the downstream cell-cycle pathways are inhibited (Fig. 2B), and the cells are in G0 arrest. Conversely, forced, isolated overexpression of downstream oncogenes (for example, viral E1A, c-Myc and E2F) can induce apoptosis (Fig. 2C). However, cell-cycle progression and apoptosis are not permanently connected, and can be separated; for example, apoptosis can be suppressed by mitogens (Evan & Littlewood, 1998). Finally, in the presence of mitogenic signalling, 'unscheduled' inhibition of the cell cycle, such as by CDKIs, results in senescence (Fig. 2D). Therefore, whereas oncogene-induced apoptosis is associated with 'forced' proliferation, senescence is associated with 'forced' arrest (Fig. 2C versus

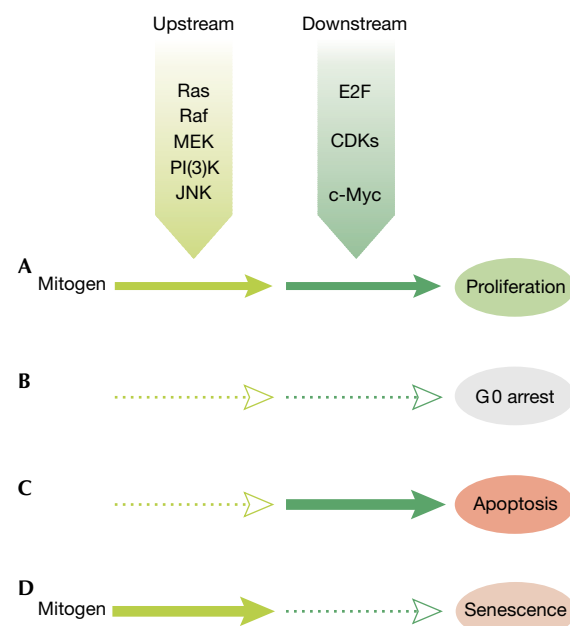


Fig. 2 | Sequential and reciprocal activation of upstream and downstream signalling pathways. (A) Normal mitogenic signalling involves the sequential activation of both upstream and downstream pathways. Note that, due to feedback loops, the distinctions between 'upstream' and 'downstream' are not always maintained, and the molecules indicated in the boxes are merely examples. (B) Classical growth arrest in G0 occurs when there is no activation of either upstream or downstream pathways. (C) The conflicting signal model of apoptosis involves the isolated overactivation of downstream pathways. (D) The conflicting signal model of senescence involves the isolated overactivation of upstream pathways.

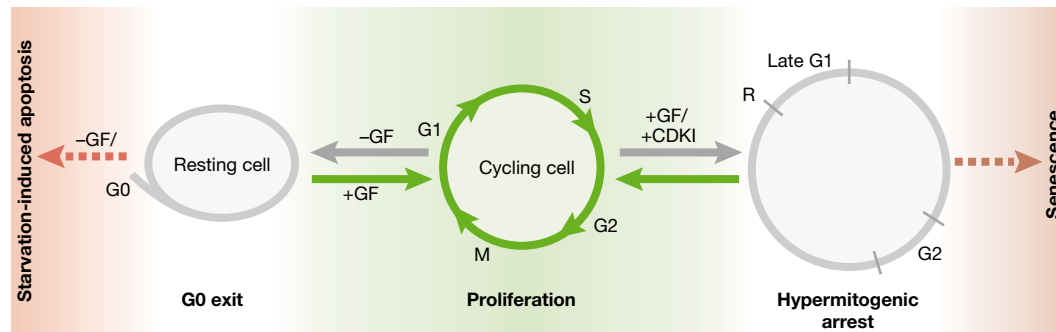


Fig. 3 | From proliferation to either G0 arrest or hypermitogenic arrest. ‘Cycling cells’ proliferate in the presence of growth factors (GFs). The withdrawal of GFs causes a reversible exit from the cell cycle, known as G0, producing a ‘resting cell’. In normal cells, a complete lack of GFs may cause apoptosis. GF stimulation in the presence of cyclin-dependent-kinase inhibitors (+CDKI) can cause hypermitogenic arrest, and continuous stimulation can lead to senescence.

2D). In this view, oncogene-induced apoptosis and senescence are mirror images, each involving the isolated activation of either upstream or downstream mitogenic pathways.

Cell-cycle arrest: hypomitogenic and hypermitogenic

The model of mitogen-induced senescence presented here leads to an unexpected prediction: there are two opposing types of growth arrest. Classical growth arrest, or G0 arrest, which is caused by growth-factor withdrawal, is essentially an exit from the cell cycle (Fig. 3), and is characterized by low levels of cyclin D1. After mitogenic stimulation, cells in the G0 state enter G1, during which they induce cyclin D1 and progress through the restriction point (Blagosklonny & Pardee, 2002). It is generally assumed that all resting cells ultimately exit the cell cycle. However, when cells arrest in response to mitogens, the consequences are different. In this state of hypermitogenic arrest, cells arrest at or beyond the restriction point, in either G1 or G2 phase (Fig. 3), and numerous anticancer drugs arrest cells when the levels of cyclins D, E, A or B are elevated (Gong *et al.*, 1995; Darzynkiewicz, 2002). The hypermitogenic arrest induced by phorbol esters is characterized by CDK inhibition in the presence of cyclin D1 (Blagosklonny, 1998), and iron chelators, contact inhibition and transforming growth factor- β induce a senescence-like late G1 arrest that is often associated with the induction of p27 (Kletsas *et al.*, 1995; Yoon *et al.*, 2002). DNA damage can induce permanent growth arrest in G2, which is dependent on Rb (Naderi *et al.*, 2002). Also, hyperactivated Raf1 can arrest cells in G2 (Kerkhoff & Rapp, 1998). In cycling cells, Ras induces cyclin D1 in G2 phase (Sherr, 2002). Therefore, forced hypermitogenic arrest is associated with elevated cyclin D1.

The existence of G0 has been questioned time after time, precisely because the idea of G0 arrest has been overgeneralized. The distinction between G0 (hypomitogenic arrest) and hypermitogenic arrest may explain much of this controversy.

Hypermitogenic arrest, cell morphology and function

Intriguingly, the model of hypermitogenic arrest predicts a ‘large-cell’ morphology, and high levels of autocrine and/or paracrine secretion by senescent cells. Hypermitogenic arrest must be accompanied by protein synthesis—as mitogens stimulate not only DNA synthesis, but also RNA and protein synthesis—and by metabolism. It is an arrest of the cell cycle, not of growth. In fact, high-intensity Raf signals are known to convert mitotic cell cycling into cellular growth (Kerkhoff & Rapp, 1998). Thus,

hypermitogenic arrest can explain the large-cell morphology that has been associated with senescence.

It is also possible that a cell may attempt to compensate for the cell-cycle block by the overinduction of mitogenic signalling. Senescent cells are highly active in the autocrine and paracrine secretion of mitogens (Roninson *et al.*, 2001). Unlike the hypomitogenic arrest at G0, hypermitogenic arrest is enhanced by stimulation by mitogens. Therefore, hypermitogenic arrest and cell senescence are stable conditions, which are often seen as irreversible. For example, consistent with the senescent phenotype, p38-induced G1 cell-cycle arrest is irreversible after four days (Haq *et al.*, 2002). As another example, when p16 expression was induced for six days, DNA synthesis remained inhibited and the cells acquired the morphological features of senescence. If p16 induction was interrupted after six days, most cells retained these morphological features (Dai & Enders, 2000).

Overtuning hypermitogenic arrest and senescence

The concept presented here predicts that, unlike G0 arrest, hypermitogenic arrest cannot be blocked by mitogenic stimuli and, indeed, senescent cells have been shown to be resistant to mitogens (Mathon & Lloyd, 2001). However, as illustrated by the following examples, it can be overcome by induction of downstream ‘oncogenes’. Both c-Myc and E1A override the hypermitogenic arrest caused by PMA (phorbol-12-myristate-13-acetate) (Blagosklonny, 1998). Reactivation of cyclin E1, which acts downstream of p16, is sufficient to trigger escape from Ras-induced senescence (Peeper *et al.*, 2002). In contrast to stimulation by mitogens, stimulation by viral proteins can reactivate terminally differentiated cells of the muscle system: for example, the large-T viral oncoprotein induces the cycling of terminally differentiated skeletal muscle myotubes (Ohkubo *et al.*, 1994), E2F1 can override ‘irreversible’ arrest in post-mitotic ventricular myocytes (Agah *et al.*, 1997), and the E1A oncogene can reactivate the cell cycle in terminally differentiated skeletal muscle cells (Tiainen *et al.*, 1996). Also, the inactivation of either p53 or simian virus 40 (SV40) induces senescent cells to re-enter S phase and to revert to their ‘young’ morphology (Gire & Wynford-Thomas, 1998). Finally, oncogenes that act downstream of the restriction point (such as c-Myc and CDK4), and thus may overcome senescence, cooperate with hypermitogenic oncogenes (such as Ras), leading to malignant transformation (Land *et al.*, 1983).

The physiological role of cell senescence

The activation of apoptosis, by 'unscheduled' oncogenic signalling, is thought to contribute to the prevention of cancer. Similarly, cell senescence has been proposed as a mechanism to block immortalization and tumorigenesis. For example, it may limit the transforming potential of excessive Ras mitogenic signalling. However, this view may be too focused on cancer. Similar to apoptosis, which has many physiological roles in development and tissue turnover, senescence may have other physiological functions. By analogy to apoptosis, it is generally perceived as a pre-death stage that results in the elimination of cells. The fact that senescence is a stable and metabolically active state, however, argues that the functions of senescence and apoptosis oppose one another: senescence does not eliminate a cell, but instead preserves it in a maximally stable and metabolically active state. For example, senescent fibroblasts have a useful purpose: they express proteins that are normally induced on wounding, including those that remodel the extracellular matrix (Benanti *et al.*, 2002). Furthermore, senescence in fibroblasts is associated with resistance to apoptosis caused by radiation (Yeo *et al.*, 2000). By contrast, mitogen deprivation results in an unstable condition that can lead to apoptosis because mitogens are also survival factors.

Many terminally differentiated cells may be arrested in a hypermitogenic (senescence-like) state, rather than in the unstable and apoptosis-prone G0 arrest, owing to phenomena such as contact inhibition. One can speculate that muscle cells, dermal fibroblasts and certain neuronal cells may be preserved in this way. For example, contact inhibition of growth may be accompanied by p27 accumulation and cellular senescence (Munro *et al.*, 2001). Unlike arrest due to growth-factor withdrawal, arrest due to contact inhibition occurs in the presence of mitogens. Interestingly, cell-cell interactions between biliary cells and hepatocytes result in the mid-G1-phase arrest of hepatocytes, which are insensitive to mitogens (Loyer *et al.*, 1996). If quiescence in 'contact-inhibited' organs, such as the liver, is controlled by CDKs (such as p27), then the downregulation of these inhibitors might be necessary for liver regeneration, for example after partial hepatectomy.

It is possible that certain mitogens secreted by contact-inhibited tissue might maintain hypermitogenic arrest, essentially acting like growth inhibitors. More than 30 years ago, it was postulated that tissues produce inhibitors, known as chalones, to control cell homeostasis (Bullough & Laurence, 1968). Despite thousands of publications describing putative chalones, these mysterious inhibitors have never been purified. Perhaps these elusive inhibitors are simply ordinary mitogens, acting as mitogens in G0 cells and as chalones in cycling cells. Finally, if cells in our body are arrested in a hypermitogenic state, this explains why mammalian viral oncoproteins (for example, E1A, E7, E6 and T antigen) act downstream of the restriction point (Fig. 2, downstream pathways). In fact, oncoproteins that act upstream would simply strengthen the senescent state. The sequence of oncogenic events that lead to cancer may therefore vary depending on the prevalence of hypermitogenic versus G0 arrest in different tissues.

ACKNOWLEDGEMENTS

I apologize to those authors whose work was not cited due to page limitations.

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