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Nuclear Lamins and Oxidative Stress in Cell Proliferation and Longevity

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Abstract

In mammalian cells, the nuclear lamina is composed of a complex fibrillar network associated with the inner membrane of the nuclear envelope. The lamina provides mechanical support for the nucleus and functions as the major determinant of its size and shape. At its innermost aspect it associates with peripheral components of chromatin and thereby contributes to the organization of interphase chromosomes. The A- and B-type lamins are the major structural components of the lamina, and numerous mutations in the A-type lamin gene have been shown to cause many types of human diseases collectively known as the laminopathies. These mutations have also been shown to cause a disruption in the normal interactions between the A and B lamin networks. The impact of these mutations on nuclear functions is related to the roles of lamins in regulating various essential processes including DNA synthesis and damage repair, transcription and the regulation of genes involved in the response to oxidative stress. The major cause of oxidative stress is the production of reactive oxygen species (ROS), which is critically important for cell proliferation and longevity. Moderate increases in ROS act to initiate signaling pathways involved in cell proliferation and differentiation, whereas excessive increases in ROS cause oxidative stress, which in turn induces cell death and/or senescence. In this review, we cover current findings about the role of lamins in regulating cell proliferation and longevity through oxidative stress responses and ROS signaling pathways. We also speculate on the involvement of lamins in tumor cell proliferation through the control of ROS metabolism.

Keywords

Nuclear lamins; Lamin A; Lamin B; Oxidative stress; Reactive oxygen species (ROS)

Introduction

In mammalian cells, the nuclear lamina is a major determinant of nuclear architecture. The lamina is located at the inner membrane of the nuclear envelope (NE). The major structural components of the lamina in somatic cells are the A-type lamins (LA, LC) and the B-type lamins (LB1, LB2). LA and LC are derived from the single gene *LMNA* by alternative splicing, and LB1 and LB2 are encoded by two genes *LMNB1* and *LMNB2*, respectively [1].

In embryonic stem cells, the expression of LA and LC is low and they begin to increase at the onset of differentiation and continue to increase to relatively high levels in certain terminally differentiated cell types [2]. In contrast, LB1 and/or LB2 are expressed in all cells throughout development [2]. For example, T-cells and B-cells express only B-type lamins but not A-type lamins [1].

Lamins are type V intermediate filament proteins, which assemble into higher order filamentous structures within the peripheral lamina under the NE [3, 4]. All lamins contain a long central α -helical rod domain, flanked by globular N-terminal (head) and C-terminal (tail) domains. Many lamin subtypes are posttranslationally modified either transiently or permanently (see below). In particular, LA is transiently modified by C-terminal farnesylation and the failure to remove this farnesylation site results in nuclear defects, while B-type lamins tend to be permanently farnesylated. Electron microscopy has revealed that the lamina in *Xenopus* oocytes appears as a meshwork of ~10–15 nm filaments [5]. Lamin structures organized into meshworks have also been seen in nuclei of mouse cells by super resolution light microscopy [6]. Furthermore, it has been shown that A- and B-type lamin fibrils form separate but interacting meshworks within the lamina [7]. These lamin fibrils play important roles in assembling the lamina and contribute to the size, shape, and mechanical stability of the nucleus. Lamins are also involved in nuclear functions including chromatin organization, DNA replication, DNA repair, and transcription [7–10]. With respect to chromatin organization, the lamins provide anchorage sites for peripheral elements of heterochromatin, which are involved in the local regulation of gene expression [11–13]. Interestingly, silencing LB1 expression in HeLa cells dramatically alters the structure of the LA/C meshworks and induces LA/C-enriched NE blebs [7] that contain transcriptionally inactive gene-rich euchromatin in cancer cells [7].

The functional importance of lamins is further supported by the finding that structural changes in the lamina are among the most dramatic hallmarks of differentiation, cancer and aging and that numerous mutations in the *LMNA* gene are now known to be responsible for a wide range of genetic disorders called laminopathies. These combined studies suggest that lamins play important roles as key regulators of epigenetic events that may be critical in cellular stress responses. In particular, knowledge is accumulating to show an interdependence between oxidative stress and lamins. For example, oxidative stress modulates the expression and posttranslational modification of lamins. Conversely, mutations of lamin genes and depletion of lamins affect oxidative stress responses. Reactive oxygen species (ROS), major products of oxidative stress, are natural by-products of mitochondrial respiration which are normally eliminated in protective mechanisms such as antioxidant defenses [14–16]. Moderate increases in ROS act as a signaling mechanism to promote cell proliferation and differentiation [14–16]. However, excessive increases in ROS cause damage to DNA, proteins, and lipids, resulting in defects in proliferation and longevity that have been linked to cardiovascular and neurodegenerative diseases, as well as chronic inflammation [17]. Importantly, it is now becoming evident that lamins are involved in modulating ROS to regulate proliferation and longevity.

Here, we discuss current knowledge regarding the involvement of lamins in oxidative stress, cell proliferation, and longevity. Specifically, we focus our attention on the role of lamins in

mediating cell proliferation and longevity through oxidative stress responses and ROS signaling pathways. We also consider the possible involvement of this nexus in tumor proliferation.

The Expression and Stability of Lamin Proteins Is Modulated by Oxidative Stress

Several studies have indicated that the expression and stability of lamin proteins is altered in response to oxidative stress, which in turn is tightly coupled to cell proliferation, cellular senescence, apoptosis, and autophagy.

Lamin expression is regulated by the tumor suppressors p53 and retinoblastoma protein (pRb) and by telomere functions; all master regulators of the cell cycle, apoptosis, replicative senescence, and autophagy. For example, LA/C expression is significantly upregulated upon the activation of p53 [18]. The LA mutant progerin, which causes the premature aging disease Hutchinson-Gilford Progeria Syndrome (HGPS) [19] is also expressed during normal aging [20]. Progerin expression is induced by telomere dysfunctions [21]. In contrast, the expression of LB1, but not LB2, is significantly down-regulated during senescence induced by replicative exhaustion, DNA damage, and oncogenic stress [22–24] (Fig. 1). A decrease in LB1 expression has also been observed in HGPS and in atypical progeroid syndromes caused by different mutations in the *LMNA* gene [25–27]. This decrease in LB1 expression is specifically coupled to senescence, since it does not occur in quiescence induced by serum depletion [22, 23]. The activation of pRb is required for the decrease in LB1 expression in senescence [22], and this is attributable to the fact that the *LMNB1* gene is a downstream target of the pRb–E2F pathway [28]. Based on these findings, it would be predicted that LA/C expression increases and LB1 expression decreases with no change in expression of progerin and LB2 when oxidative stress activates p53 and pRb [22] (Fig. 2). However, it has been reported that fibroblasts derived from patients with the recessive autosomal genetic disorder ataxia telangiectasia show an increase in LB1 expression in response to oxidative stress mediated by the activation of p38 mitogen-activated protein (MAP) kinase [29]. It is therefore, possible that the mutation causing ataxia telangiectasia alters the pathways required for regulating LB1 expression levels.

In addition to the transcriptional regulation of lamin expression described above, other studies have indicated that lamin levels are affected by directed degradation. Rapamycin induces autophagic protein degradation by inactivating the mammalian target of rapamycin (mTOR) pathway [30]. In this fashion, the non-farnesylated form of LA, premature LA (pre-LA), and progerin are degraded by rapamycin treatment [31, 32]. Therefore, it could be that oxygen tension which is upstream of mTOR complex 1 (mTORC1) affects the protein levels of pre-LA and progerin through the mTOR pathway [33]. LB1 is degraded in transformed rat fibroblasts and a human cervical carcinoma cell line after these cells are exposed to a ROS inducer by the ubiquitin–proteasome pathway [34]. The lamina is also known to be broken down by caspases during apoptosis, and lamins are considered to be among the initial nuclear targets cleaved during the apoptotic process [35]. A- and B-type lamins are cleaved at their conserved VEID and VEVD sites by caspase-6 and 3, respectively [36–39]. Furthermore, both serine proteases, granzyme A and B, are known to cleave B-type lamins,

whereas only granzyme A but not B appear to cleave A-type lamins [40]. Since oxidative stress can induce apoptosis [41], it is also likely that these cleavages of lamin proteins are induced by oxidative stress (Fig. 2). Together, these studies strongly support the idea that oxidative stress modulates the expression and stability of lamin proteins through transcription and proteolysis.

Posttranslational Modifications of Lamins in Response to Oxidative Stress

Lamins are known to be posttranslationally processed and the resulting modifications are likely to affect their functions, their interactions with each other, and their binding partners [42]. Several studies have indicated that lamins are posttranslationally modified by oxidation and enzymes in response to oxidative stress.

Lamins contain some amino acid residues that could be oxidized. During senescence, increased ROS results in the oxidation of cysteine residues in the LA tail domain, which in turn appears to inhibit the formation of LA inter- and intramolecular disulfide bonds [43]. Additionally, S-thiolation of A-type lamins is induced in isolated rat kidneys subjected to ischemia and reperfusion [44].

Phosphorylation of lamins has been the most extensively studied among many lamin posttranslational modifications. Though it is well known that hyperphosphorylation of lamins drives NE disassembly in mitosis [45–48], little is known regarding interphase phosphorylation of lamins. During interphase A-type lamins are known to be phosphorylated in response to oxidative stress in human neuroblastoma cells [49]. Since extracellular signal-regulated kinase 1/2 (ERK1/2) is activated by oxidative stress [50], it is possible that A-type lamins are phosphorylated by this kinase. In support of this, A-type lamins have been identified as among the most heavily phosphorylated proteins following activation of ERK1/2 [51, 52]. LB1 is also phosphorylated by p38 MAP kinase during senescence induced by oxidative stress, which leads to an increase in LB1 expression [29].

Furthermore, it has been shown that LA posttranslational processing by farnesylation is affected by oxidative stress. This is supported by the accumulation of pre-LA in old vascular smooth muscle cells (VSMCs) but not in young healthy blood vessels [53]. This accumulation of pre-LA correlates with the downregulation of the metalloproteinase Zmpste24/FACE-1, which is required for the processing of pre-LA into mature LA. Since both the mRNA and protein level of Zmpste24 are reduced in response to oxidative stress [53], this affects pre-LA levels. Posttranslational modifications and levels of B-type lamins are also altered by oxidative stress. For example, in transformed and cancer cells, LB1 protein is oxidized by ROS, which mediates the degradation of LB1 protein [34].

It still remains unclear how these posttranslational modifications of lamins induced by oxidative stress affect their structures and functions. However, one study shows that aggregates of LA/C and LB1 are observed in the nucleus during the early response to liver injury induced by oxidative stress [54]. This suggests, but certainly does not prove, that changes in posttranslational modifications of lamins caused by oxidative stress may inhibit lamin assembly into the lamina, which may lead to dysfunctions of the nucleus.

Lamin Functions in Cell Proliferation and Longevity

As mentioned above, the lamina and lamins are known to be involved in various nuclear functions including chromatin organization, DNA replication, DNA repair, and transcription (see “ Introduction ”). Some studies have indicated that the deregulation of these lamin functions by disease causing mutations and/or alterations in the expression levels of the different types of lamins inhibit cell proliferation and induce senescence or cell death.

The functions of A-type lamins in cell proliferation have been most extensively studied in the premature aging diseases, HGPS, and atypical progeroid syndromes. Dermal fibroblasts obtained from progeria patients are commonly found to proliferate slowly and become prematurely senescent [27, 55]. In several transgenic mouse models for HGPS, the mice also show marked reduction in body size and fibroblasts derived from the progeria mice exhibit slow proliferation and premature senescence [56, 57]. Mutations in the gene encoding a pre-LA processing enzyme, *Zmpste24* and *Zmpste24* knockout mice (*Zmpste24*^{-/-}) cause a severe progeroid phenotype similar to HGPS [58, 59]. In support of this, the accumulation of pre-LA by silencing *Zmpste24* expression or the overexpression of pre-LA accelerates VSMC senescence [53]. Similarly, silencing of LA/C expression slows cell proliferation and induces premature senescence in human diploid fibroblasts (HDFs) [60]. *Lmna* knockout mice (*Lmna*^{-/-}) also have proliferation defects within 4 weeks after birth and die by 8 weeks [61].

Alterations in LA/C either caused by silencing or defects in processing by *Zmpste24* appear to increase the susceptibility of cells to DNA damage [62–65], most likely resulting in the activation of p53. In support of this, silencing LA/C expression or overexpression of LA and progerin slows proliferation and induces premature senescence in a p53-dependent but not pRb-dependent manner [60, 66] (Fig. 3). In *Zmpste24*^{-/-} mice, p53 and the p53 target genes are upregulated and induce proliferation defects and premature death [67].

Recent studies have shown that B-type lamins are also involved in cell proliferation, differentiation, and longevity. Silencing the expression of LB1 slows cell proliferation in HDFs [22, 24] and induces premature senescence [22]. *Lmnb1* mutant mice (*Lmnb1*^Δ), *Lmnb1* knockout mice (*Lmnb1*^{-/-}), and *Lmnb1* / *Lmnb2* double knockout mice (*Lmnb1*^{-/-} *Lmnb2*^{-/-}) are born, but die immediately after birth due to developmental defects in specific differentiated cell types such as those comprising the lung and brain [68, 69]. *Lmnb2*^{-/-} mice also have severe brain abnormalities, but these developmental defects are rather minor compared to those in *Lmnb1*^{-/-} mice [69, 70]. Importantly, mouse embryonic fibroblasts (MEFs) derived from *Lmnb1*^Δ mice and the mutant mice expressing a polymorphic variant of LB1 responsible for the curly tail phenotype show proliferation defects and prematurely senesce [68, 71]. On the other hand, mice with the conditional *Lmnb1*^{-/-} *Lmnb2*^{-/-} restricted to their skin keratinocytes develop normally [72]. These findings support the idea that cell proliferation defects caused by the absence of B-type lamins are most likely to be tissue specific.

As in the case of the A-type lamins, there is evidence that p53 mediates the regulation of proliferation by LB1. Changes in proliferation rates caused by silencing LB1 expression and overexpression of LB1 can be restored by the inactivation of p53 [22, 24] (Fig. 3).

The Involvement of Lamins in Regulating Oxidative Stress

Oxidative stress activates several signaling pathways including the DNA damage response (DDR), the p38 MAP-kinase pathway, and the p53 pathway to induce slow proliferation, senescence, and cell death. Several studies have shown that lamins are involved in responses to oxidative stress.

There is significant evidence that in fibroblasts derived from HGPS patients, ROS levels are higher than those of normal fibroblasts, and these elevated levels of ROS are correlated with slow proliferation rates [73]. In addition, ROS-induced DNA double-strand breaks in HGPS fibroblasts are not repaired normally, and this appears to be related to their slow rate of proliferation [73]. This increase in ROS level in HGPS fibroblasts could be due to cysteine residues missing in progerin, which appear to be hyperoxidized during senescence and may contribute to the suppression of ROS-responsive genes [43]. Furthermore, mesenchymal stem cells (MSCs) overexpressing progerin or LA shows a decrease in expression of an antioxidant SOD2 and a SOD2-dependent increase in mitochondrial ROS, which leads to defects in chondrogenic differentiation potential [74]. Another study also shows that the viability of MSCs and VSMCs which have been differentiated from pluripotent stem cells (iPSCs) derived from HGPS fibroblasts are compromised by oxidative stress [75].

Other mutations in the *LMNA* gene have been reported to cause the accumulation of pre-LA and/or an increase in oxidative stress. Fibroblasts derived from patients with the *LMNA* mutations D47Y, L92F, L387V, R399H, L421P, and R482W, causing insulin resistance and/or lipodystrophy, accumulate pre-LA, and this is related to increased oxidative stress and the decreased expression of mitochondrial respiratory chain proteins that trigger premature cellular senescence [76]. Another study shows that cells from lipodystrophy patients with the LA mutations R439C, R482W, and H506D also accumulate pre-LA and show the expression of adipogenic proteins with brown fat-like features, an increased number of mitochondria and the overexpression of thermogenin (uncoupling protein 1, UCP1), which decreases the proton gradient generated in oxidative phosphorylation [77]. Higher levels of ROS are also induced by oxidative stress in fibroblasts from other lipodystrophy patients with a R439C *LMNA* mutation [78]. In addition, a homozygous *LMNA* mutation leads to expression of a mutated pre-LA with a deletion of 48 C-terminal amino acids, preventing its farnesylation and maturation. The resulting form of pre-LA is associated with increased oxidative stress and premature senescence [79]. *Zmpste24*^{-/-} mice also accumulate pre-LA and show an increased mitochondrial response to oxidative stress [80]. Moreover, the mitochondrial proteins related to lipid metabolism, the tricarboxylic acid cycle, and oxidative phosphorylation are all upregulated in these mice. This supports the relationship between defective pre-LA processing and mitochondrial dysfunction, in addition to highlighting the relevance of pre-LA to oxidative damage in lipodystrophy and aging [80]. These results strongly support the idea that dysfunctions of LA/C cause oxidative stress (Fig. 3).

LB1 is also known to be involved in oxidative stress responses. The proliferation defects induced by silencing LB1 expression are accompanied by a p53-dependent reduction of mitochondrial ROS in HDFs, which can be rescued by growth under hypoxic conditions [22] (Fig. 3). RT-PCR analyses show that p53-target genes are altered under the experimental conditions used for silencing LB1 expression. For example, the antioxidant genes *SOD1*, *SOD2*, *SESN1*, *SESN2*, and *GPX1* are all upregulated, resulting in lowering the levels of ROS in LB1-silenced cells [22]. Other than p53, the POU-domain transcription factor Oct-1 also appears to mediate the LB1 regulation of the genes involved in oxidative stress responses [81]. Most importantly, Oct-1 has been shown to associate with the lamina through LB1 [11, 81]. This association is disrupted in the *Lmnb1*^{-/-} fibroblasts, causing the elevation of ROS levels [81]. In autosomal dominant leukodystrophy (ADLD) fibroblasts, there is an increased expression of LB1 due to the duplication of the *LMNB1* gene, which is also coincident with an increase in the amount of Oct-1 associated with the lamina and a decrease in the nucleoplasmic fraction of Oct-1 by oxidative stress [82]. Since Oct-1 regulates the expression of genes involved in oxidative stress responses by binding to their putative octamer-binding DNA sequences [81], the sequestration of Oct-1 to the lamina by LB1 could be another mechanism by which LB1 modulates ROS levels (Fig. 3).

LB2 appears to regulate mitochondrial functions in neurons. For example, in *Lmnb2*^{-/-} mice, a neuronal layering abnormality is caused by defective neuronal migration [70]. Inhibition of *LMNB2* mRNA translation in *Xenopus* retinal ganglion cell axons in vivo does not affect guidance but causes axonal degeneration [83]. This is attributable to the finding that a form of axonal LB2 associates with mitochondria and in LB2-deficient axons mitochondria are dysfunctional causing defects in axonal transport [83]. These results suggest that axonally synthesized LB2 plays a crucial role in axon maintenance by promoting mitochondrial functions to protect against oxidative stress.

Possible Pathways Related to Lamin Functions in Response to Oxygen Metabolism and Oxidative Stress Responses

Recent studies have indicated that lamins play important roles in regulating signaling pathways mediated by SIRT1, NF- κ B, and mTORC1. These pathways are known to be involved in regulating oxygen metabolism and oxidative stress. Though in many cases, direct biochemical evidence has yet to be provided, these studies imply that lamins may regulate a cellular response to oxidative stress through these pathways.

In this regard, LA has been shown to interact with and activate an NAD-dependent deacetylase, Sirtuin 1 (SIRT1) [84]. SIRT1 deacetylates FOXO3, a member of the FOXO family of Forkhead transcription factors in response to oxidative stress, leading to cell cycle arrest, resistance to oxidative stress, and inhibition of cell death [85]. The presence of progerin or the accumulation of pre-LA in *Zmpste24*^{-/-} mice induces a decrease in deacetylase activity of SIRT1, leading to rapid depletion of adult stem cells [84]. This scenario might also explain why a mitochondrial response to oxidative stress is increased in *Zmpste24*^{-/-} mice and fibroblasts derived from HGPS patients [73, 80]. Therefore, it is possible that LA functions in oxidative stress responses are mediated by SIRT1.

It has also been shown that the accumulation of pre-LA activates an ATM- and NEMO-dependent signaling pathway, leading to the activation of NF- κ B and secretion of proinflammatory cytokines in *Zmpste24*^{-/-} and progeria mice (*Lmna*^{G609G/G609G}) [86]. The activation of NF- κ B suppresses cell death by inducing the expression of genes encoding pro-oxidants [87]. In this fashion, the accumulation of pre-LA might increase the levels of ROS through the activation of NF- κ B [80]. It is also possible that NF- κ B mediates LA functions in oxidative stress responses.

Cellular oxygen sensing is upstream of the mTOR pathway [88, 89]. Interestingly, *Lmna*^{-/-} mice show enhanced mTORC1 signaling in cardiac and skeletal muscle cells. Pharmacologic reversal of elevated mTORC1 signaling by rapamycin improves cardiac and skeletal muscle function and the longevity of the mice. In addition, this treatment also alleviates the defective autophagic-mediated degradation in *Lmna*^{-/-} mice [90]. Based on these findings, it has been suggested that there is molecular cross talk between LA and oxygen sensing mechanisms.

The Prospective for Future Studies of Lamin Functions in Oxidative Stress

We have discussed current evidence for roles of lamins in regulating cell proliferation and longevity through the cellular response to oxidative stress and ROS signaling pathways. Though the lamin-related diseases that we have described are not directly involved in cancer, there are significant implications that lamins could be involved in tumor cell growth and in cancer progression. Many types of cancer cells are known to produce increased reactive ROS compared to normal cells [91]. Cancer cells also develop a unique way to control their proliferation as their increased glucose metabolism is coupled to fast proliferation [22] and their mitochondrial metabolism regulates ROS production which is essential for anchorage-independent growth [14]. Cancer cells are also more susceptible to oxidative stress compared to normal cells [91]. Therefore, progress in understanding how lamins control ROS metabolism in normal and cancer cells will provide new insights for cancer treatment. The modulation of ROS metabolism by changing lamin expression might provide a biochemical basis to design therapeutic strategies, including vectors for gene therapy and small molecule compounds for chemotherapy, to selectively slow cancer development, growth, and progression.

Abbreviations

ADLD	Autosomal dominant leukodystrophy
DDR	DNA damage response
HDFs	Human diploid fibroblasts
HGPS	Hutchison-Gilford progeria syndrome
iPSCs	Inducible pluripotent stem cells
LA	Lamin A
LB1	Lamin B1

LB2	Lamin B2
LC	Lamin C
MSC	Mesenchymal stem cells
MEFs	Mouse embryonic fibroblasts
NE	Nuclear envelope
ROS	Reactive oxygen species
VSMCs	Vascular smooth muscle cells

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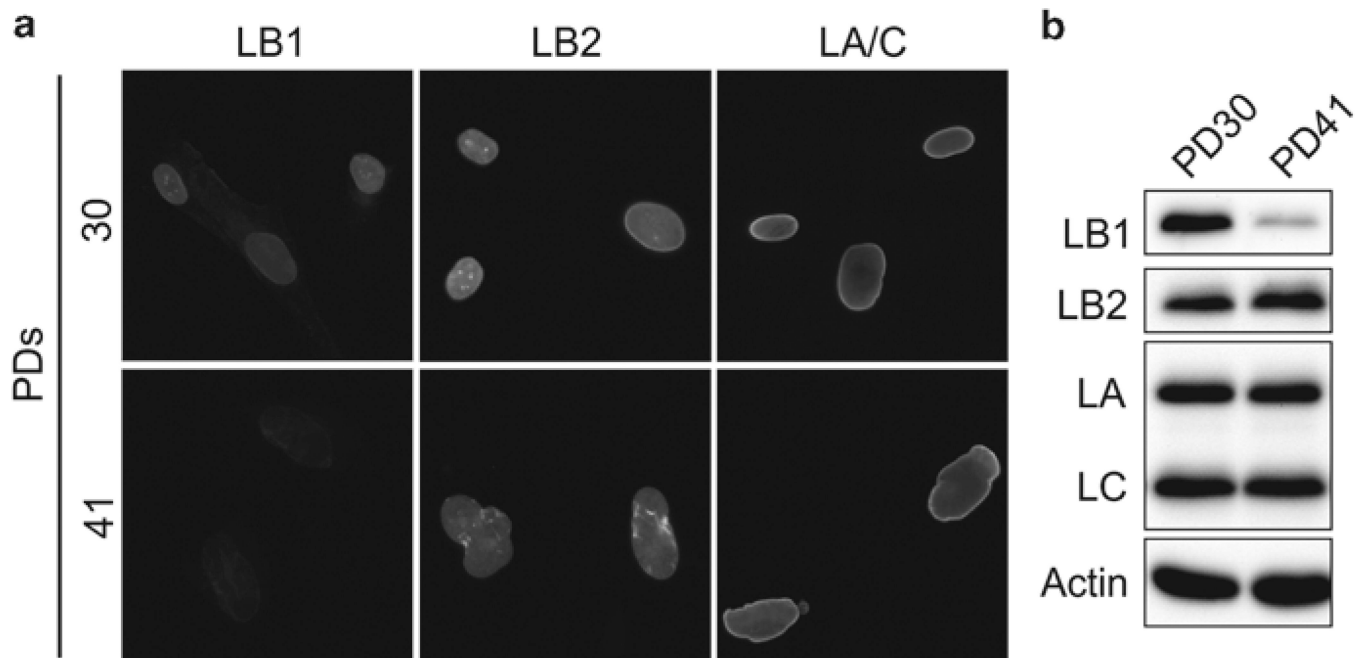


Fig. 1. The distribution and expression of lamin proteins were determined at population doublings (PD) 30 and 41 in WI-38 human embryonic lung fibroblasts. These cells are proliferating at PD 30 and senescent by PD 41. **(a)** LA/C, LB1, and LB2 are localized by immunofluorescence. **(b)** The expression of LA/C, LB1, and LB2 was determined by immunoblotting. The expression of LB1 but not LA/C or LB2 was significantly decreased during replicative senescence (permission to reproduce these data from Cold Spring Harbor Lab Press [22])

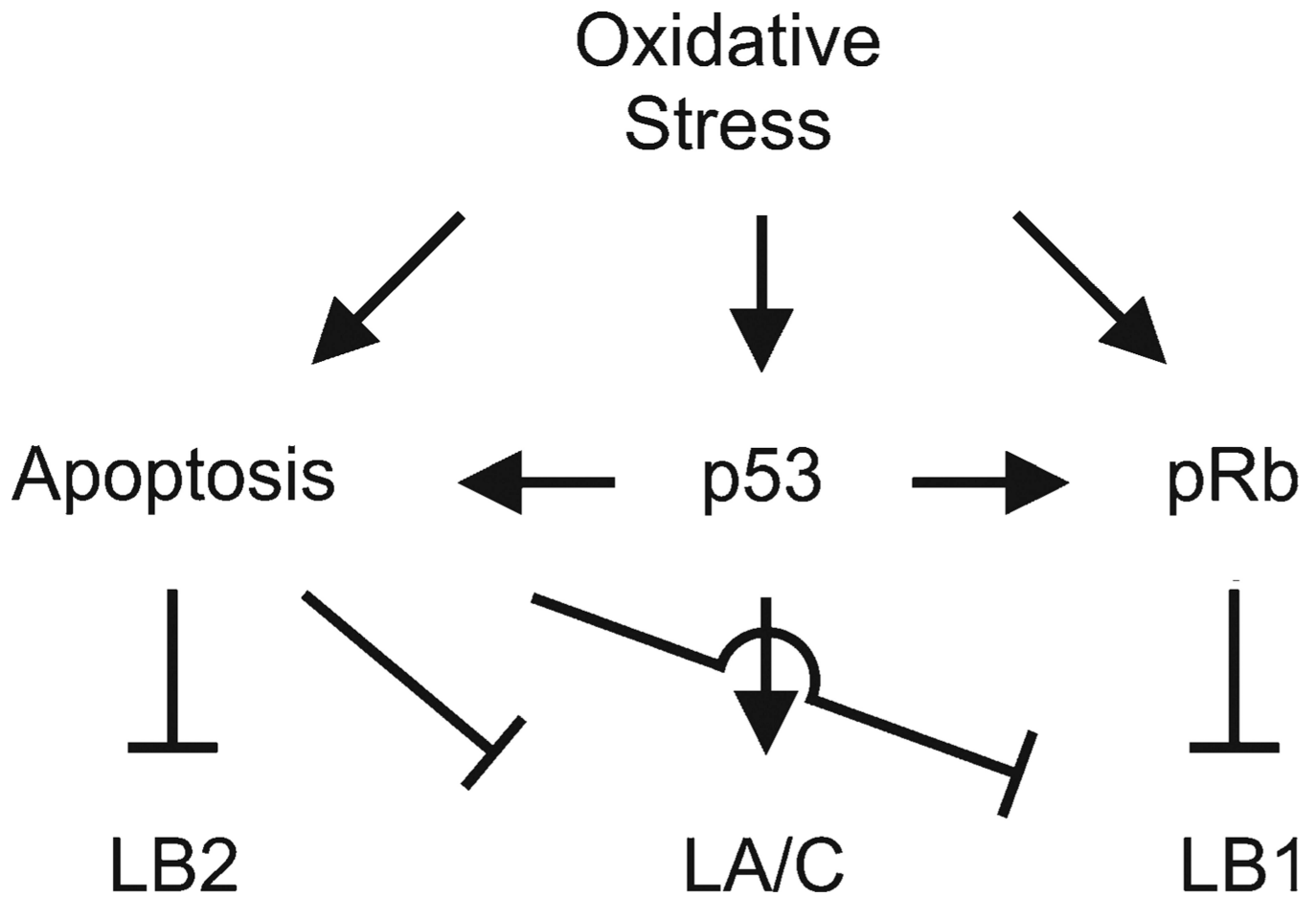
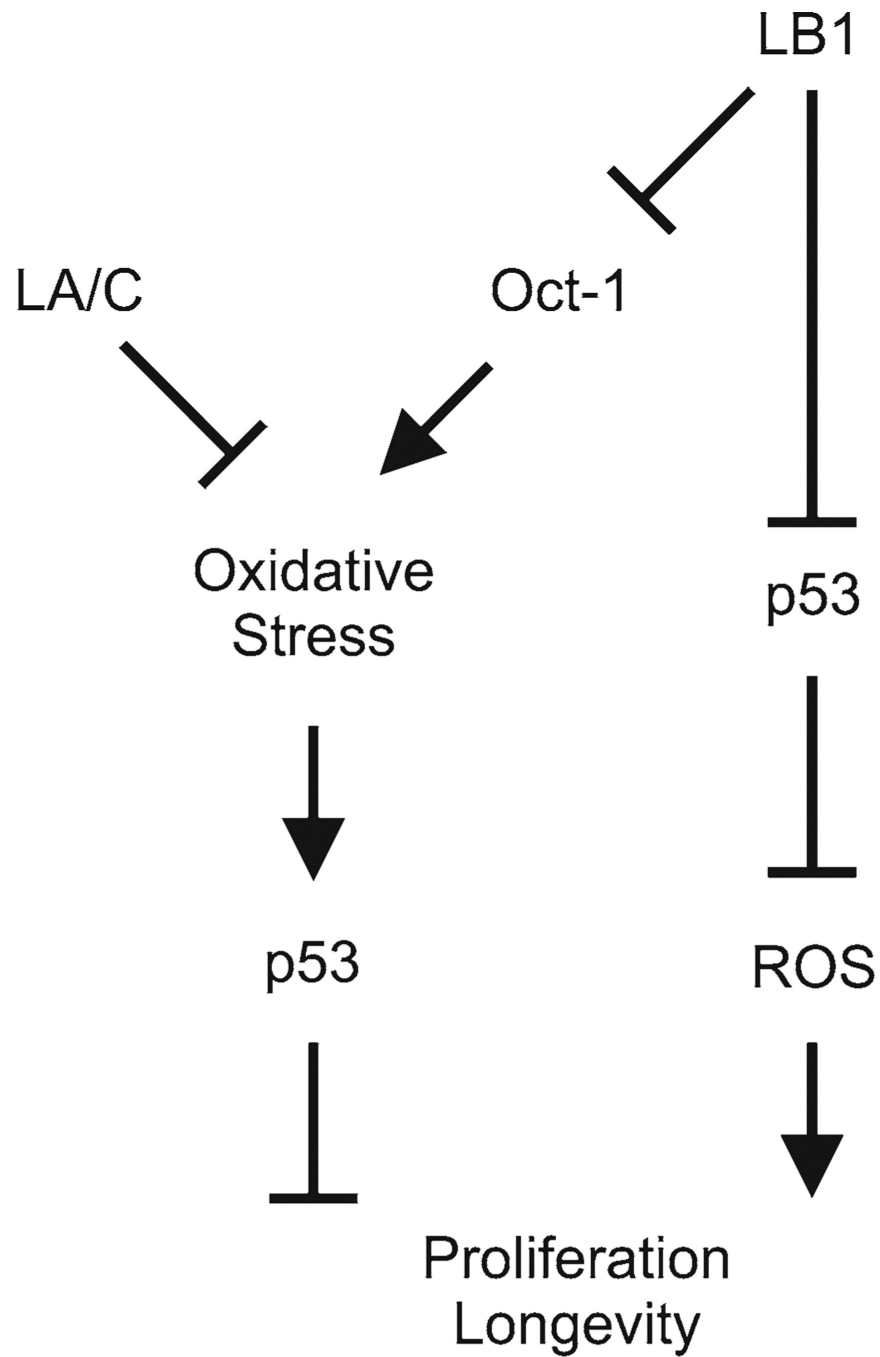


Fig. 2.

A summary of the p53, pRb, and apoptotic pathways known to modulate the levels of lamin expression in response to oxidative stress. Oxidative stress induces an increase in LA/C expression, a decrease in LB1 expression, but no change in the expression of LB2 through the p53 and pRb pathways. Lamin proteins can be cleaved by oxidative-stress induced apoptosis

**Fig. 3.**

A summary of p53- and Oct-1 dependent pathways linking lamin functions to oxidative stress responses, cell proliferation, and longevity. LA/C and LB1 suppress oxidative stress to promote proliferation and longevity through the p53 pathway. The suppression of oxidative stress by LB1 is mediated by Oct-1. LB1 also maintains ROS metabolism through the p53 pathway