Nuclear Lamins and Neurobiology

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Much of the work on nuclear lamins during the past 15 years has focused on mutations in *LMNA* (the gene for prelamin A and lamin C) that cause particular muscular dystrophy, cardiomyopathy, partial lipodystrophy, and progeroid syndromes. These disorders, often called "laminopathies," mainly affect mesenchymal tissues (e.g., striated muscle, bone, and fibrous tissue). Recently, however, a series of papers have identified important roles for nuclear lamins in the central nervous system. Studies of knockout mice uncovered a key role for B-type lamins (lamins B1 and B2) in neuronal migration in the developing brain. Also, duplications of *LMNB1* (the gene for lamin B1) have been shown to cause autosome-dominant leukodystrophy. Finally, recent studies have uncovered a peculiar pattern of nuclear lamin expression in the brain. Lamin C transcripts are present at high levels in the brain, but prelamin A expression levels are very low—due to regulation of prelamin A transcripts by microRNA 9. This form of prelamin A regulation likely explains why "prelamin A diseases" such as Hutchinson-Gilford progeria syndrome spare the central nervous system. In this review, we summarize recent progress in elucidating links between nuclear lamins and neurobiology.

The nuclear lamina has attracted considerable scrutiny from biochemists, cell biologists, and geneticists. Much of this attention has been stimulated by the discovery that mutations in *LMNA* (the gene for the A-type lamins, prelamin A and lamin C) cause multiple human diseases, including muscular dystrophy, cardiomyopathy with conduction system disease, partial lipodystrophy, and progeroid syndromes (1–3). These diseases, often called "laminopathies," are largely confined to mesenchymal tissues. For the past decade, many laboratories have worked to define disease mechanisms and to devise therapeutic strategies. These efforts have been summarized in many reviews (2–11).

While "*LMNA* diseases" have attracted most of the research efforts, a new topic has slowly emerged in the field—nuclear lamin biology in the brain. Three lines of investigation have contributed to the emergence of this topic. The first was work by developmental biologists that uncovered a role for the B-type lamins (lamins B1 and B2) in neuronal migration in the developing brain (12–14). The second was work showing that a demyelinating disorder, autosomal dominant leukodystrophy, is caused by *LMNB1* gene duplications (15, 16). The third was work by clinical investigators to understand the spectrum of disease phenotypes in children with Hutchinson-Gilford progeria syndrome (17, 18), a precocious aging syndrome caused by a toxic form of prelamin A (19). For years, the field marveled that children with progeria have multisystem aging-like phenotypes but are spared common features of physiologic aging in the central nervous system, notably, senile dementia. Recent studies have identified a likely mechanism—regulation of prelamin A by a brain-specific microRNA (17, 18).

PROTEINS OF THE NUCLEAR LAMINA

The nuclear lamina in somatic cells is mainly composed of four nuclear lamins (lamins A, C, B1, and B2), which form a filamentous meshwork lining the inner nuclear membrane. Prelamin A (the precursor to mature lamin A) and lamin C are products of the *LMNA* gene and are produced by alternative splicing (20). Prelamin A and lamin C are identical through the first 566 amino acids but have distinct carboxyl-terminal sequences (20). Lamin C contains sequences from exons 1 to 10 and terminates with six unique amino acids not found in prelamin A. Prelamin A contains

sequences from two additional exons (exons 11 and 12) and terminates with 98 unique residues (20). Lamins B1 and B2 are produced from separate genes, *LMNB1* and *LMNB2* (21, 22). The nuclear lamina interacts with chromatin and nuclear membrane proteins and is important for providing structural support for the cell nucleus and tethering the nucleus to the cytoskeleton (7, 23, 24).

Prelamin A terminates with a "CaaX" motif (CSIM), which triggers three posttranslational modifications: farnesylation of the carboxyl-terminal cysteine, endoproteolytic release of the last three amino acids of the protein, and carboxymethylation of the newly exposed farnesylcysteine. Subsequently, the last 15 residues of prelamin A, including the farnesylcysteine methyl ester, are removed by a zinc metalloprotease, ZMPSTE24, releasing mature lamin A (25–27). The generation of mature lamin A is utterly dependent on protein farnesylation; when protein farnesylation is inhibited, the subsequent processing steps do not occur, and nonfarnesylated prelamin A accumulates in cells (28).

Prelamin A's farnesylcysteine methyl ester is generally assumed to associate with the inner nuclear membrane and facilitate the delivery of lamin A to the nuclear lamina (29). However, studies of *Lmna* knock-in mice have suggested that farnesylation and methylation might have little impact on the targeting of lamin A to the nuclear rim (10). Coffinier et al. (30) created *Lmna* knock-in mice that produce mature lamin A directly, bypassing prelamin A synthesis and all of the subsequent posttranslational processing steps. These mice were quite healthy, even after 2 years of observation, and the targeting of mature lamin A to the nuclear rim in mouse tissues appeared normal. Also, in *Lmna* knock-in mice that produce a nonfarnesylated version of prelamin A, there was no obvi-

ous defect in the delivery of the nonfarnesylated prelamin A to the nuclear rim (31).

Lamin C does not have a CaaX motif and therefore is not farnesylated or methylated. Lamins B1 and B2 terminate with a CaaX motif and undergo farnesylation and methylation. Unlike the situation with prelamin A, however, the farnesylcysteine methyl ester in lamins B1 and B2 is not removed by a subsequent endoproteolytic cleavage step.

Lamins B1 and B2 are expressed at high levels in virtually all cells from the earliest stages of development (32). Lamins A and C are expressed at low levels early in development (33), but expression levels increase late in development (14, 34). The low levels of *Lmna* expression early in development (14, 34), along with the fact that *Lmna* knockout mice survive development (35), have led to the notion that lamins A and C are mainly important for differentiated cells (4).

Sullivan et al. (35) created mice lacking both lamin A and lamin C (*Lmna*$^{-/-}$). Although the mice survived embryonic development, they succumbed to muscular dystrophy and cardiomyopathy by 4 to 5 weeks of age. Embryonic fibroblasts exhibited striking abnormalities in nuclear shape. Subsequent studies revealed that the knockout mice expressed trace amounts of an internally truncated prelamin A (36). However, that product appeared to be relatively unimportant because the phenotypes of a newer line of *Lmna*$^{-/-}$ mice were similar (37). To gain insights into unique roles of prelamin A and lamin C, Fong et al. (38) and Davies et al. (31) created knock-in mice that produced exclusively lamin C or exclusively prelamin A. However, both mouse models were healthy and free of pathology, limiting decisive insights into the unique functions for the two *Lmna* isoforms.

Until recently, dogma held that the B-type lamins were essential proteins with many crucial functions in the cell nucleus (4), ranging from roles in DNA replication and gene transcription (39–41) to the formation of the mitotic spindle (42) and the organization of heterochromatin (43). A paper holding that cultured cells undergo growth arrest and apoptosis after small interfering RNA (siRNA) inhibition of *LMNB1* and *LMNB2* expression lent credence to this view (44). However, recent studies with genetically modified mice have undercut this view—at least in the case of peripheral tissues. Mice lacking both lamin B1 and lamin B2 in keratinocytes had no skin pathology, had normal keratinocyte proliferation rates, and had no abnormalities in the nuclear envelope or in heterochromatin distribution (45). Also, an absence of both lamin B1 and lamin B2 in hepatocytes did not lead to liver pathology (46). One obvious explanation for these findings is that expression of lamins A and C is sufficient for the vitality of keratinocytes and hepatocytes. However, Kim et al. (47) recently reported that mouse embryonic stem cells lacking all nuclear lamins were able to proliferate and differentiate—and even form large teratomas in mice. Those studies were surprising and at face value further undermine the view that lamins play important roles in mitotic spindle assembly, DNA replication, and other vital functions in the cell nucleus. Additional efforts to define the consequence of a complete deficiency of nuclear lamins in specific cell types are needed.

A ROLE FOR B-TYPE LAMINS IN BRAIN DEVELOPMENT

The first clue about the *in vivo* importance of B-type lamins in mammals came from lamin B2 knockout mice (12). *Lmnb2*-deficient embryos (*Lmnb2*$^{-/-}$) were virtually normal in size during
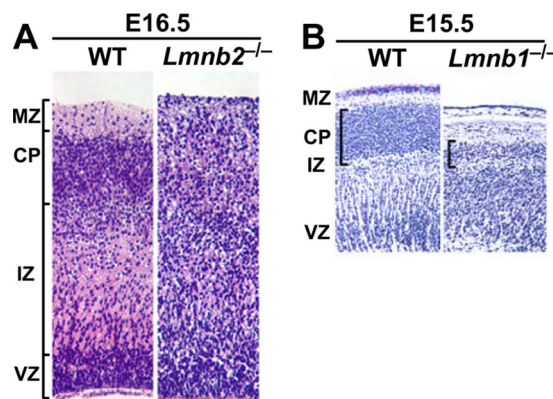

FIG 1 Defects in neuronal migration and neuronal survival in *Lmnb1* and *Lmnb2* knockout mice. (A and B) Hematoxylin and eosin staining of cerebral cortex sections from E16.5 wild-type (WT) and *Lmnb2* knockout (*Lmnb2*$^{-/-}$) embryos (A) and sections from E15.5 WT and *Lmnb1* knockout (*Lmnb1*$^{-/-}$) embryos (B). (Panel A was reprinted from reference 12 and panel B was reprinted from reference 14 with permission of the publishers.) MZ, marginal zone; CP, cortical plate; IZ, intermediate zone; VZ, ventricular zone.

development, but they died shortly after birth (12) with obvious neuropathology. The cerebellum was small and devoid of sulci; the cerebral cortex was also small, and the layering of cortical neurons was abnormal. Further studies revealed that the neuronal layering abnormality was present as early as embryonic day 16.5 (E16.5) (Fig. 1A). Because this pathology was similar to pathology in mice with well-characterized defects in neuronal migration (48–50), Coffinier et al. (12) performed immunochemistry studies with layer-specific neuronal markers and bromodeoxyuridine (BrdU) birth-dating studies. In E19.5 *Lmnb2*$^{-/-}$ embryos, many NeuN-positive neurons accumulated in lower levels of the cortical plate. In wild-type littermates, most NeuN-positive neurons migrated past Ctip2-positive neurons into the superficial layers of the cortex. Also, FoxP1-positive neurons in *Lmnb2*$^{-/-}$ embryos accumulated in lower levels of the cortical plate, and few neurons reached the more superficial layers. The neuronal migration defect was confirmed by BrdU birth-dating studies. In wild-type mice, most neurons that originated at E13.5 (i.e., when the BrdU was injected) were found deep in the cortical plate at E18.5. Neurons that originated at later time points (and which had little BrdU) migrated past the BrdU-positive neurons into more-superficial layers of the cortical plate. In contrast, most BrdU-positive neurons in *Lmnb2*$^{-/-}$ embryos were located in superficial layers of the cortical plate, suggesting that newer, BrdU-negative neurons lacked the capacity to migrate into more-superficial layers (12). Neuronal migration defects were also documented in an independently created line of *Lmnb2*$^{-/-}$ mice (51).

In hindsight, a role for the nuclear lamina in neuronal migration was not surprising. Neuronal migration depends on cytoplasmic motors pulling the cell nucleus forward toward the centrosome in the leading edge of the cell (nucleokinesis) (52). The trailing portion of the migrating neuron is then remodeled, and the net effect is forward movement of the cell. Repeated cycles of this process make it possible for neurons to traverse from the ventricular zone to the cortical plate. When *Lmnb2*$^{-/-}$ mice were initially created (12), the cytoplasmic proteins required for "tugging" on the cell nucleus had already been investigated in considerable detail (49, 53, 54), but the identity of nuclear proteins re-

quired for this process was unclear. Finding defective neuronal migration in $Lmnb2^{-/-}$ embryos implied that neuronal migration depends on B-type lamins.

The cytoplasmic motors required for nuclear translocation cannot interact directly with the nuclear lamina, simply because the nuclear lamina is located within the nucleus. Instead, the forces on the cell nucleus are almost certainly exerted through the LINC (Linker of nucleoskeleton and cytoskeleton complex) (13, 55). The LINC complex involves KASH proteins (e.g., nesprins) in the outer nuclear membrane and SUN proteins in the inner nuclear membrane. On the cytoplasmic side, the nesprins interact with elements of the cytoskeleton, while on the nucleoplasmic side, the SUN proteins interact with the nuclear lamina (13). Nuclear translocation and neuronal migration depend on the LINC complex; the absence of SUN1 and SUN2, or the removal of the KASH domain from nesprin 1 and nesprin 2, results in neuronal migration defects similar to those in $Lmnb2^{-/-}$ mice (56).

Coffinier et al. (12) suggested that the neuronal migration defects in $Lmnb2^{-/-}$ mice were a consequence of reduced integrity of the nuclear lamina—leading to impaired nuclear translocation. Studies on the brain pathology in $Lmnb2^{-/-}$ mice supported this model (14). Many of the cell nuclei in the cortical plate neurons of $Lmnb2^{-/-}$ embryos were markedly "stretched out" (comet shaped), and they suggested that this abnormality was due to the deformational stresses during nuclear translocation. Presumably, the cytoplasmic motors in $Lmnb2^{-/-}$ neurons simply "stretched out" a weakened nuclear envelope rather than moving the nucleus into the leading edge of the cell.

In the report on $Lmnb2^{-/-}$ mice, Coffinier et al. (12) speculated that lamin B1 might also play a role in brain development. At that time, $Lmnb1$ knockout mice from a gene-trap ES cell clone had already been described (57). The $Lmnb1^{-/-}$ embryos, which expressed a lamin B1–β-galactosidase fusion protein, were small and died shortly after birth with an abnormally shaped cranium. More-detailed studies by Coffinier et al. (14) revealed that $Lmnb1$ knockout embryos had a small brain with abnormal layering of neurons in the cerebral cortex (Fig. 1B). Immunohistochemical studies with antibodies against Otx1 and TBR1 indicated a neuronal migration defect, and that was confirmed with BrdU birth-dating studies (14). Another group, using an independent line of $Lmnb1^{-/-}$ mice, reached similar conclusions (51).

The brain pathology in the $Lmnb1^{-/-}$ embryos was more severe than in the $Lmnb2^{-/-}$ embryos; there were fewer neurons in the cortical plate, and numbers of neuronal progenitors were reduced (14). Also, nuclear shape abnormalities were different. There were almost no comet-shaped nuclei in $Lmnb1^{-/-}$ embryos, but a large fraction of $Lmnb1^{-/-}$ neurons contained a single large nuclear bleb. Also, lamin B2 was asymmetrically distributed and much of it was located in the nuclear bleb. The authors suggested that reduced integrity of the nuclear envelope during nucleokinesis may have contributed to the formation of the nuclear bleb (14).

As discussed previously by Coffinier and coworkers and Young and coworkers (12, 13, 58), the discovery that lamins B1 and B2 are essential for neuronal migration in the mammalian brain was foreshadowed by a publication by Patterson et al. on eye development in *Drosophila* (59). They found that a *Drosophila* lamin and Klarsicht, a KASH domain-containing protein, were essential for photoreceptor nuclear migration in the *Drosophila* eye. They spec-

ulated that the lamin-Klarsicht interaction was relevant to human laminopathies, such as muscular dystrophy and cardiomyopathy.

A ROLE FOR THE B-TYPE LAMINS IN NEURONAL SURVIVAL

To explore the importance of lamin B1 and lamin B2 in neurons postnatally, Coffinier et al. (14) used recombination techniques involving *Cre* and *loxP* to generate forebrain-specific *Lmnb1* knockout mice and forebrain-specific *Lmnb2* knockout mice. Mice from both models survived, but they had a small forebrain with neuronal layering abnormalities. In both models, the number of forebrain neurons in adult mice was markedly reduced (more so than in the knockout embryos), suggesting that the loss of either B-type lamin impairs neuronal survival. Coffinier et al. (14) went on to breed forebrain-specific *Lmnb1* and *Lmnb2* double-knockout mice. Those mice had complete atrophy of the forebrain and no detectable neurons in the forebrain, indicating that the loss of both B-type lamins is not compatible with neuronal survival. This situation is obviously different than with keratinocytes and hepatocytes, where the loss of B-type lamins had no significant consequences (45, 46). Of note, *Lmna* is expressed in the adult mouse brain, but it is not expressed in neurons during development (14). We suspect that the complete absence of nuclear lamins in the forebrain neurons of forebrain-specific *Lmnb1-Lmnb2* knockout embryos "seals their fate" and places them on a pathway to cell death. The general idea that nuclear lamins are relevant to cell survival in response to stress is supported by other studies. For example, lamin A expression promotes survival of cells forced to migrate through 3-μm micropores (60). Other studies have shown that nuclear lamin expression is a key determinant of cell stiffness and the ability of cells to migrate through micropores (61–65).

The mouse studies leave little doubt that the two B-type lamins play important roles in brain development and neuronal survival in mice, but what is the situation in humans? Surprisingly, no one has yet identified *LMNB1* or *LMNB2* loss-of-function mutations in humans with neurodevelopmental abnormalities. There have been suggestions that *LMNB1* polymorphisms may be associated with neural tube defects (66, 67), but more data are needed to be confident in that association. In the NHLBI exome sequencing database (http://evs.gs.washington.edu/EVS/), there are many rare *LMNB1* and *LMNB2* missense mutations, and there were several *LMNB1* frameshift or splicing mutations. However, homozygous *LMNB1* or *LMNB2* mutations have not yet been encountered in humans, likely because the relevant patient populations have not yet been screened. Sooner or later, we suspect that *LMNB1* and *LMNB2* loss-of-function mutations will be encountered in human fetuses with neurodevelopmental defects, and we would not be surprised to find missense mutations in outpatient populations in neurology clinics.

INVESTIGATING THE IMPORTANCE OF THE FARNESYL LIPID ANCHOR IN LAMIN B1 AND LAMIN B2

Lamins B1 and B2 are farnesylated proteins, and it is likely that the farnesyl lipid is embedded in the inner nuclear membrane, thereby attaching the nuclear lamina to the inner nuclear membrane. Jung et al. (68) speculated that a tight association between the nuclear lamina and nuclear membranes might be crucial for migrating neurons. To test that idea, they created knock-in mice expressing nonfarnesylated versions of lamin B1 and lamin B2 (by replacing the cysteine of the CaaX motif with a serine). The tar-
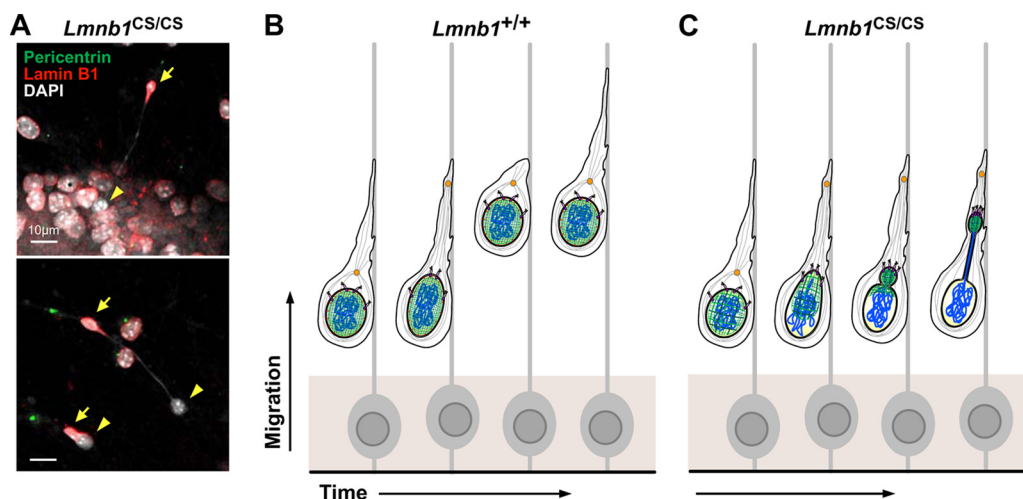
FIG 2 Nuclear abnormalities in *Lmnb1*$^{CS/CS}$ mouse embryonic fibroblasts (MEFs) and *Lmnb1*$^{CS/CS}$ neurons. (A) Immunofluorescence microscopy of migrating *Lmnb1*$^{CS/CS}$ neurons in cell culture, stained with antibodies against pericentrin (green) and lamin B1 (red). In the dumbbell-shaped *Lmnb1*$^{CS/CS}$ nuclei, lamin B1 was always in the leading edge of the cells (arrows), leaving "naked chromatin" behind in the trailing edge (arrowheads). (B and C) A model for the neuronal migration defect in *Lmnb1*$^{CS/CS}$ embryos. Unlike wild-type neurons (*Lmnb1*$^{+/+}$) (B), the nuclear lamina (green) in *Lmnb1*$^{CS/CS}$ neurons (C) is not tightly affixed to the inner nuclear membrane because of the absence of the farnesyl lipid anchor on lamin B1 (red) (creating a potential space between the nuclear lamina and the inner nuclear membrane). Also, the nuclear lamina meshwork in *Lmnb1*$^{CS/CS}$ cells is not as tightly woven, exhibiting a honeycomb distribution. During neuronal migration, the nuclear lamina is pulled forward by the LINC complex (SUN1/2, purple; nesprin-1/2, pink) by microtubule (gray strands)-associated dynein motors (yellow) in the direction of the centrosome (orange) in the leading edge of the cell. However, the chromatin (blue) escapes from the bounds of the nuclear lamina into the space between the nuclear lamina and inner nuclear membrane. (Reprinted from reference 68 with permission of the publisher.)

geted missense mutations worked as planned, abolishing lamin B1 and lamin B2 farnesylation. Absent farnesylation led to lower steady-state levels of both B-type lamins (68).

The "nonfarnesylated lamin B2" knock-in mice survived and were free of neuropathology or behavioral abnormalities, but the situation was very different for mice expressing nonfarnesylated lamin B1. Like *Lmnb1*$^{-/-}$ mice, the "nonfarnesylated lamin B1" mice died soon after birth with neuronal layering abnormalities in the cerebral cortex. However, the nonfarnesylated lamin B1 appeared to retain partial function. The size of newborn nonfarnesylated lamin B1 mice was nearly normal, whereas *Lmnb1*$^{-/-}$ embryos were extremely small (14, 57). Also, the brain pathology in the nonfarnesylated lamin B1 mice was milder than in *Lmnb1*$^{-/-}$ mice.

The neurons expressing nonfarnesylated lamin B1 exhibited a remarkable nuclear shape abnormality—dumbbell-shaped nuclei in which the nuclear lamina was separated from the bulk of the chromosomal DNA (Fig. 2A) (68). The nuclear lamina was "bunched up" in one end of the dumbbell within the leading edge of the cell, while most of the chromatin was found at the other end of the dumbbell and was "naked"—in that it was entirely free of a nuclear lamina. Both ends of the dumbbell were surrounded by nuclear membranes, as judged by staining for the inner nuclear membrane protein LAP2β. Dumbbell-shaped nuclei were detected in the midbrain and cortex and were also found in cultured neurons as they migrated away from neurospheres. Jung et al. (68) proposed that the dumbbell-shaped nuclei and "naked chromatin" were the consequence of weakened interactions between the nuclear lamina and the inner nuclear membrane (Fig. 2B). They suggested that the absence of lamin B1 farnesylation abolishes the hydrophobic interactions that normally affix the nuclear lamina to the inner nuclear membrane. They further suggested that migrating neurons from the "nonfarnesylated lamin B1 mice" would

retain the ability to pull the nuclear lamina forward into the leading edge of the cell but that the chromatin would not "come along" and would instead "leak" into the potential space between the inner nuclear membrane and the nuclear lamina. Neurons expressing nonfarnesylated lamin B1 also had a honeycombed nuclear lamina, which likely facilitated the escape of genomic DNA from the nuclear lamina. Of note, the nonfarnesylated version of lamin B1 appeared (by immunofluorescence microscopy) to be located mainly at the nuclear rim—even in cells with a honeycombed nuclear lamina. In wild-type neurons, DNA cannot escape the bounds of the nuclear lamina because the farnesyl lipids anchor the nuclear lamina to the inner nuclear membrane and because of the more tightly woven pattern of the nuclear lamina.

INVESTIGATING THE FUNCTIONAL REDUNDANCY OF LAMINS B1 AND B2 IN BRAIN DEVELOPMENT

Lamins B1 and B2 are ~60% identical at the amino acid level (11), and both are expressed ubiquitously beginning at the earliest stages of development (32). Also, both proteins play crucial roles in neuronal migration and neuronal survival (12). The similarities in lamin B1 and lamin B2 sequences and expression patterns, along with the similar neurodevelopmental abnormalities in the knockout mice, naturally raise the issue of whether the two proteins have redundant functions. To address that issue, Lee and coworkers (69) generated "reciprocal knock-in mice," which made it possible to determine if increased expression of one of the B-type lamins would compensate for the loss of the other. In the first knock-in mouse (*Lmnb1*$^{B2/B2}$ mice), they inserted a lamin B2 cDNA into exon 1 of *Lmnb1*, resulting in lamin B2 expression from the *Lmnb1* locus while eliminating *Lmnb1* transcripts. In the other mouse (*Lmnb2*$^{B1/B1}$ mice), they inserted a lamin B1 cDNA into exon 1 of *Lmnb2*, resulting in lamin B1 expression from the *Lmnb2* locus while eliminating *Lmnb2* transcripts.
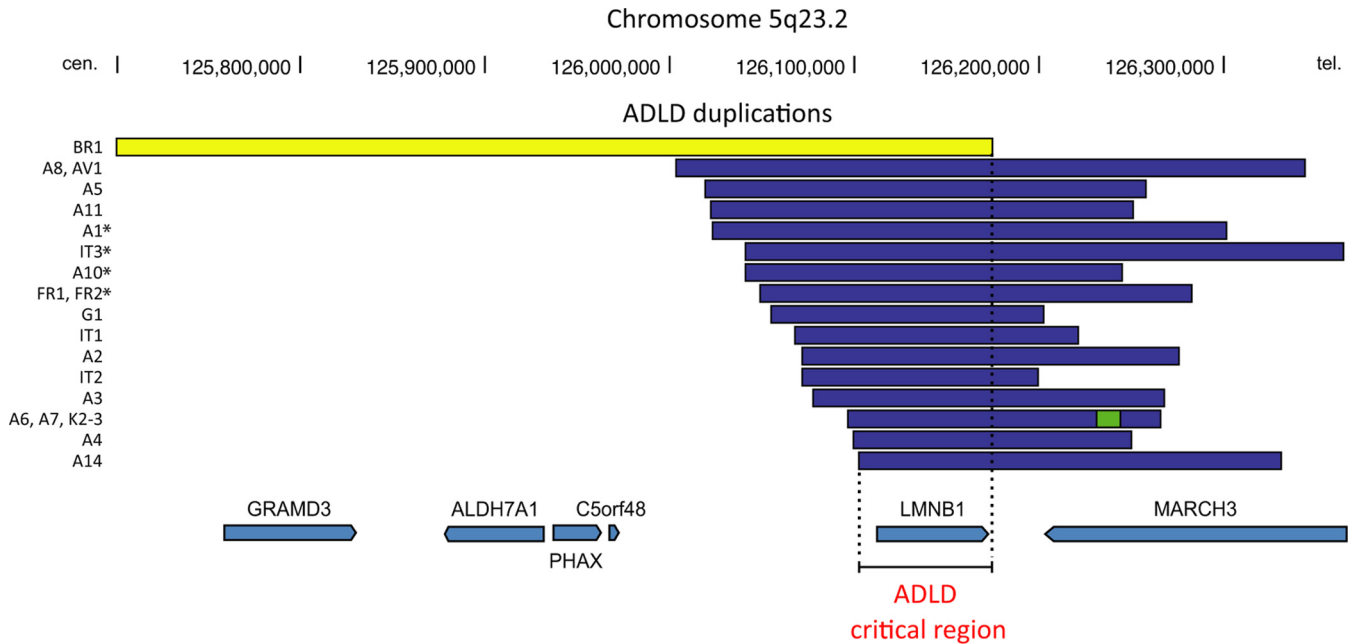
FIG 3 Genomic rearrangements in autosomal dominant leukodystrophy (ADLD) families. The modified output from the University of California, Santa Cruz (UCSC), genome browser shows *LMNB1* gene duplications in 20 ADLD families (16 unique *LMNB1* duplications). The duplications are marked in blue, with the exception of the BR1 duplication/inversion, which is in yellow; a "triplicated" segment is marked in green. Duplications marked with asterisks have sequence insertions at their duplication junctions and show a clustering of their centromeric (cen.) breakpoints within a 25-kb segment. The minimal critical region duplicated in ADLD of ~75 kb is also shown. (Reprinted from reference 75 with permission of the publisher.)

The *Lmnb1*$^{B2/B2}$ mice worked as planned; *Lmnb2* transcript levels in the cerebral cortex were >2.5-fold higher than in wild-type mice (reflecting the production of *Lmnb2* transcripts from both *Lmnb2* and *Lmnb1*) (69). Lamin B2 levels in the cortex of *Lmnb1*$^{B2/B2}$ embryos were ~3-fold higher than in wild-type embryos. Despite increased levels of lamin B2 expression, the *Lmnb1*$^{B2/B2}$ mice were small and died shortly after birth with neuronal layering abnormalities in the cerebral cortex, indicating that increased lamin B2 production cannot prevent the developmental abnormalities associated with lamin B1 deficiency (69). However, the body and brain weights of *Lmnb1*$^{B2/B2}$ embryos were higher than in *Lmnb1*$^{-/-}$ embryos, and the density of cortical neurons in *Lmnb1*$^{B2/B2}$ embryos was greater—implying that increased lamin B2 production partially compensates for the loss of lamin B1 (69).

The story with *Lmnb2*$^{B1/B1}$ mice was similar. Those mice, which expressed increased amounts of lamin B1 and no lamin B2, manifested neurodevelopmental abnormalities (69) that were virtually identical to those in *Lmnb2*$^{-/-}$ mice (12). Thus, surplus lamin B1 could not fully compensate for the loss of lamin B2. Again, however, there was evidence for partial compensation. The pathology in the cerebral cortex of *Lmnb1*$^{B2/B2}$ *Lmnb2*$^{B1/B1}$ embryos was milder than in *Lmnb1*$^{B2/B2}$ embryos, implying that lamin B1 synthesis from the *Lmnb2*^{B1} allele was capable of lessening the severity of the developmental defects found in *Lmnb1*$^{B2/B2}$ embryos.

LAMIN B1 GENE DUPLICATIONS AND AUTOSOMAL DOMINANT LEUKODYSTROPHY

A major advance in the human genetics of B-type lamins was the discovery that an adult-onset demyelinating disorder, autosomal dominant leukodystrophy (ADLD), is caused by *LMNB1* duplica-

tions (15, 16). ADLD was first identified in a large American-Irish kindred (70) but was soon identified in multiple families of various ethnicities (15). ADLD typically begins between the fourth and sixth decades with autonomic dysfunction (manifested by orthostatic hypotension, impotence, and bladder abnormalities), followed by progressive signs of cerebellar and pyramidal disease. Cerebellar disease is generally manifested by ataxia and tremors, while the pyramidal signs include weakness in the extremities and spasticity. ADLD is diagnosed by magnetic resonance imaging (MRI) scanning, which reveals widespread and symmetrical loss of white matter, most prominently in the frontal and parietal lobes and cerebellum (71). At autopsy, patients with ADLD have vacuolated white matter with a loss of myelin but minimal or no loss of neurons or oligodendrocytes (the cells that produce myelin) (15, 16, 71). As revealed by Western blotting, however, brain tissue from ADLD patients has reduced amounts of oligodendrocyte and myelin proteins (72). There is little astrogliosis in ADLD brains, and there are no inflammatory infiltrates (distinguishing ADLD from multiple sclerosis) (15, 71).

The ADLD gene was first localized to chromosome 5q31 (73) and later narrowed to a 1.5-Mb segment within that region (74). In 2006, Padiath et al. (16) showed that ADLD is caused by head-to-tail duplication events involving *LMNB1*. The *LMNB1* duplication led to increased expression of lamin B1 at both the RNA and protein levels. Recently, Giorgio et al. (75) characterized 16 *LMNB1* gene duplication events in 20 ADLD families (Fig. 3). The minimum duplication required for ADLD is ~72 kb—spanning the entire *LMNB1* gene and including 9.9 kb of sequences upstream from *LMNB1* and 1.8 kb downstream. The expression of the duplicated copy of the gene is equivalent to that of the other copies of the gene. In one instance, the *LMNB1* duplication event

involved recombination between *Alu* sequences, but most of the duplication events appeared to involve short (<6-bp) regions of homology upstream and downstream of *LMNB1* (75). Interestingly, one family with clinical features of ADLD manifested lamin B1 overexpression without a gene duplication event, presumably from a *LMNB1* regulatory abnormality (76).

IMPACT OF LAMIN B1 OVEREXPRESSION IN CULTURED CELLS AND IN *DROSOPHILA* AND MOUSE MODELS

Cell culture studies have shown that overexpression of B-type lamins leads to increased nuclear membrane formation and nuclear shape abnormalities (blebbing) (16, 77, 78). Padiath et al. (16) showed that lamin B1 overexpression in neurons or glia of *Drosophila* results in lethality, while overexpression in the eye leads to a degenerative abnormality. Subsequent studies by Lin and Fu (72) showed that overexpression of lamin B1 in oligodendrocyte cell lines results in an arrest of oligodendrocyte differentiation along with reduced expression of oligodendrocyte markers.

Recently, Heng et al. (79) examined the effects of lamin B1 overexpression by creating transgenic mice with a 177-kb bacterial artificial chromosome (BAC) clone spanning the *Lmnb1* gene. The transgenic mice manifested cognitive abnormalities, as judged by the Morris water maze test, and progressive motor impairment, as judged by rotarod and balance beam tests. Electroencephalography uncovered frequent spontaneous seizures (79). Myelination defects were not apparent in 12-month-old BAC transgenic mice but were evident by 24 months of age, as judged by electron microscopy. The ultrastructural defects consisted mainly of "outfoldings, extensions, and invaginations" in the myelin sheath; axon disintegration was also observed. Heng et al. (79) also created transgenic mice that overexpressed lamin B1 in oligodendrocytes. Those mice developed a rapidly progressive motor abnormality leading to death by 12 months of age. The mice also had seizure activity. These abnormalities were accompanied by evidence of demyelination and axonal degeneration (79). Lamin B1 overexpression in oligodendrocytes also led to reduced expression of the major myelin protein, proteolipid protein (PLP), at both the protein and RNA levels (79). Reduced expression of PLP was attributed to reduced binding of a transcriptional activator, Yin Yang 1, to the PLP promoter.

THE IMPORTANCE OF miR-23 FOR LAMIN B1 REGULATION AND MYELIN FORMATION

LMNB1 is predicted to contain multiple binding sites for microRNA 23 (miR-23), which is among the most highly expressed microRNAs in oligodendrocytes. miR-23 overexpression led to reduced lamin B1 expression in reporter assays as well as reduced amounts of lamin B1 protein in cells, as judged by Western blotting. Lentiviral transduction of miR-23 into primary glial cultures led to increased expression of oligodendrocyte markers, whereas lamin B1 expression had the opposite effects (72). Of note, miR-23 expression appeared to mitigate the oligodendrocyte maturation defects elicited by lamin B1 overexpression. The impact of miR-23 on oligodendrocyte differentiation and myelin formation was recently tested in transgenic mice that overexpress miR-23a driven by an oligodendrocyte-specific promoter (80, 81). Interestingly, the miR-23 transgenic mice manifested a variety of neurologic abnormalities accompanied by increased myelination in the corpus callosum and increased expression of myelin-specific pro-

teins (80). Electron microscopy revealed hypermyelination of axons and other abnormalities in myelin formation (80).

HGPS AND miR-9 REGULATION OF PRELAMIN A EXPRESSION IN THE BRAIN

Hutchinson-Gilford progeria syndrome (HGPS) is a pediatric progeroid syndrome characterized by multiple disease phenotypes resembling premature aging (e.g., thin skin, osteoporosis, alopecia, and atherosclerotic coronary heart disease) (82). However, some common phenotypes with normal aging, for example, senile dementia, are absent. For this reason, HGPS and other progeroid syndromes are often referred to as "segmental aging syndromes" (83–85).

HGPS is caused by a *de novo* point mutation in exon 11 of *LMNA* that alters mRNA splicing and leads to the synthesis of a mutant prelamin A (progerin) containing an internal deletion of 50 amino acids (19, 86). Lamin C synthesis is not affected. The internal deletion does not affect prelamin A's CaaX motif; hence, progerin is farnesylated and methylated (25, 87–89). However, the site for the subsequent ZMPSTE24 cleavage step (the step that would ordinarily release mature lamin A) is eliminated by the internal deletion; hence, progerin retains its farnesyl lipid anchor. Progerin causes misshapen nuclei in cultured cells and is solely responsible for the disease phenotypes of HGPS. *Lmna* knock-in mice that synthesize progerin have many of the same phenotypes as those found in children with HGPS (8, 10, 90).

For many years, the absence of primary neurological disease in HGPS was a mystery. Jung et al. (18) hypothesized that the explanation might be straightforward—that the level of prelamin A/lamin A expression in the brain might simply be lower than in other tissues. Indeed, surveys of nuclear lamin expression in different tissues supported this explanation. In most tissues, the amounts of lamin A and lamin C are roughly equivalent, but the brain produces mainly lamin C and little lamin A (18). By immunohistochemistry, lamin C is expressed at high levels in neurons and glia of the mouse brain, while lamin A expression in the brain is virtually absent—except in capillary endothelial cells and the meningeal cells (18). The findings were similar at the RNA level—high levels of lamin C transcripts and low levels of prelamin A transcripts. Most scientists would have wagered that the distinct pattern of lamin A/lamin C expression in the brain was caused by alternative splicing, but studies with "lamin A-only" knock-in mice (30) showed that this was not the case. The targeted mutation in the lamin A-only mice eliminates lamin C splicing; all of the output from *Lmna* in these mice is channeled into prelamin A transcripts. The lamin A-only mice produced large amounts of lamin A in peripheral tissues, but lamin A expression was negligible in the brain (18). Similarly, progerin-only knock-in mice (where all of the output of *Lmna* is channeled into progerin transcripts) produced large amounts of progerin in peripheral tissues but only trace amounts in the brain (18). These findings eliminated alternative splicing as a potential explanation for low levels of prelamin A/lamin A expression in the brain. Subsequent studies showed that prelamin A expression in the brain is regulated by miR-9, which is expressed at high levels in the brain (18). miR-9 binds to a single site in prelamin A's 3′ untranslated region (UTR) and reduces prelamin A expression. When the miR-9 binding site is mutated, miR-9 has no effect on prelamin A expression. Overexpression of miR-9 in fibroblasts or HeLa cells reduces levels of prelamin A transcripts and lamin A protein but has no effect on
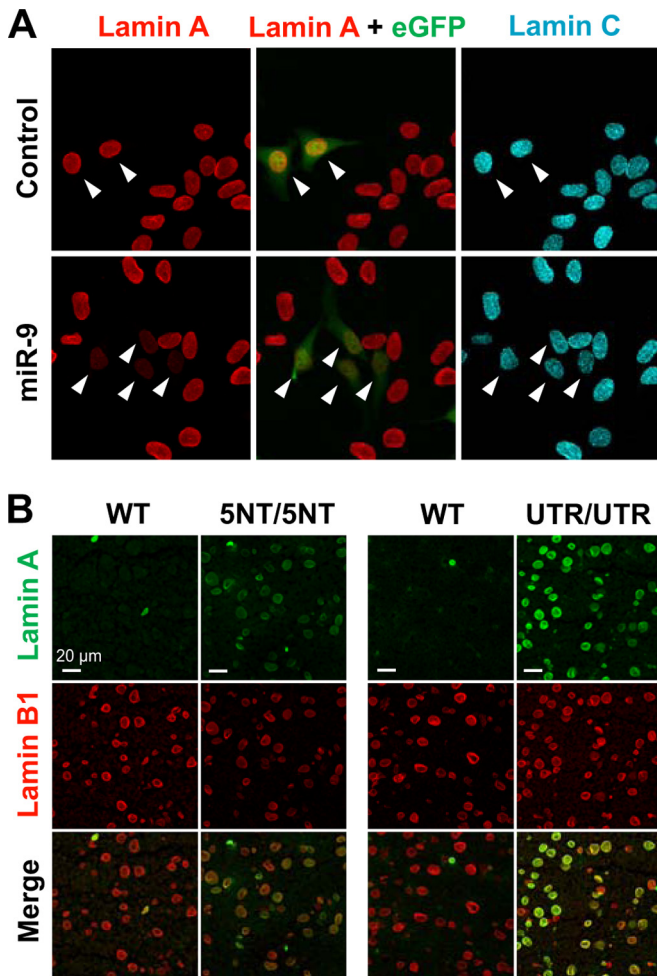
FIG 4 Downregulation of lamin A expression by miR-9. (A) Immunofluorescence microscopy of HeLa cells transfected with an empty vector (control) or a miR-9 expression vector. Transfected cells could be identified by enhanced green fluorescent protein (eGFP) expression (arrowheads). (Reprinted from reference 18 with permission of the publisher.) (B) Immunofluorescence microscopy of the cerebral cortex of $Lmna^{\mathrm{PLAO\text{-}5NT/PLAO\text{-}5NT}}$ (5NT/5NT), $Lmna^{\mathrm{PLAO\text{-}UTR/PLAO\text{-}UTR}}$ (UTR/UTR), and wild-type (WT) littermates, stained with antibodies against lamin A (green) and lamin B1 (red). Scale bar, 20 μm. (Reprinted from reference 17 with permission of the publisher.)

lamin C (Fig. 4A) (18). Nissan et al. (91) reported similar findings and also showed that miR-9 regulation of prelamin A occurs in neurons generated from induced pluripotent stem cells.

The regulation of prelamin A expression by a brain-specific microRNA suggested a possible explanation for the absence of neurodegenerative disease in HGPS. However, there were legitimate questions about the *in vivo* significance of the cell culture studies, mainly because microRNA regulation of target transcripts can be context dependent (92–95). Another issue was whether other prelamin A sequences, aside from the miR-9 binding site, might be relevant to the regulation of prelamin A. To assess the *in vivo* relevance of miR-9-mediated prelamin A regulation in the brain, Jung et al. (17) created two *Lmna* knock-in mouse lines with mutations in prelamin A's 3′ UTR. One knock-in line, $Lmna^{\mathrm{5NT/5NT}}$, contained a 5-nucleotide (5NT) mutation in the miR-9 binding site in prelamin A's 3′ UTR. In the other knock-in line ($Lmna^{\mathrm{UTR/UTR}}$), prelamin A's 3′ UTR was replaced with lamin

C's 3′ UTR. The levels of prelamin A transcripts and lamin A protein in the brain of $Lmna^{\mathrm{5NT/5NT}}$ and $Lmna^{\mathrm{UTR/UTR}}$ mice were far higher than in control mice (17). By immunohistochemistry, cortical neurons in $Lmna^{\mathrm{5NT/5NT}}$ and $Lmna^{\mathrm{UTR/UTR}}$ mice contained large amounts of lamin A (Fig. 4B) (17). In contrast, lamin A expression in the brain of control mice was mainly confined to capillary endothelial cells. Interestingly, the expression of lamin A in the brain was higher in $Lmna^{\mathrm{UTR/UTR}}$ mice than in $Lmna^{\mathrm{5NT/5NT}}$ mice (17), implying that additional 3′ UTR sequences (aside from the miR-9 site) play a role in regulating prelamin A expression. The existence of additional regulatory sequences is not particularly surprising because large segments of prelamin A's 3′ UTR—and not just the miR-9 binding site—have been conserved in mammalian evolution.

The $Lmna^{\mathrm{5NT/5NT}}$ and $Lmna^{\mathrm{UTR/UTR}}$ models provided proof that miR-9 has a crucial role in regulating prelamin A in the mammalian brain. As yet, however, the "physiologic rationale" for the preferential synthesis of lamin C in the brain is not clear. The $Lmna^{\mathrm{5NT/5NT}}$ and $Lmna^{\mathrm{UTR/UTR}}$ mice produced large amounts of prelamin A/lamin A in the brain, and yet there was no obvious neuropathology or behavioral abnormality in those mice. Thus, the explanation for the preferential synthesis of lamin C in neurons remains elusive. One possibility is that lamin C is better suited for interacting with the spectrum of nuclear envelope proteins that are expressed in neurons and glia (17). Another possibility is that lamin A expression in the brain, while innocuous in laboratory mice, is poorly suited to extenuating situations, for example, in the setting of injury or metabolic stress (17). It is also possible that prelamin A has subtle adverse effects on the function of B-type lamins.

In our opinion, understanding the "rationale" for the preferential expression of lamin C in the brain will require a far better understanding of the biochemical properties of lamins A and C and their binding partners within the nucleus. Also, while miR-9 plays a key role in lamin A expression and helps to explain the absence of neurodegenerative disease in HGPS, this argument is not entirely complete, simply because we still do not know whether progerin—if it were to be expressed in neurons—would be toxic to neurons or glia. Other cell types, for example, hepatocytes, produce abundant amounts of progerin and yet manifest little or no toxicity. In the future, it would be worthwhile determining whether progerin expression would be toxic in the brain, and if so, which cell types would be most affected.

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