# Nuclear deformation and anchorage defect induced by DCM mutants in lamin A

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# 1 **ABSTRACT**

Dilated Cardiomyopathy (DCM) is one of the different types of laminopathies caused by the 2 mutations in A-type lamins in somatic cells. The involuntary cyclic stretching of cardiac 3 4 muscle cells, as observed in normal physiological conditions is perturbed in DCM which 5 afflict patients globally. As A-type lamins are principal components in nuclear mechanics, we 6 have investigated the effect of the DCM causing mutants- K97E, E161K and R190W on 7 nuclear stretching and deformation by static and dynamic strain inducing experiments. All 8 the mutants exhibited differential nuclear structural aberrations along with a tilt in the 9 nuclear axis compared to the direction of the cell axis and a significant decrease in the lamina 10 thickness which reflected the lower mechanical rigidity. These phenotypes could potentially 11 lead to defects in nuclear anchorage to the actin filaments thereby resulting in the misshapen 12 and misaligned nucleus.

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## 15 **INTRODUCTION**

16 Nuclear lamins which are type V intermediate filament proteins were first visualized 17 to form  $\sim$ 50 nm meshwork underlying the inner nuclear membrane (1). It was considered 18 that lamin A forms 10-nm filaments inside the cell (2-5) until recently when this notion has 19 been revised by the cryo-ET structure of vimentin null mouse embryonic fibroblast 20 suggesting that both A- and B-type lamins form tetrameric 3.5 nm thick filaments inside the 21 nucleus (6). Earlier, it was also reported by the structured illumination microscopy that both 22 lamins form distinct meshwork in the nucleoplasm (7). Lamin A is a principal regulator of 23 nuclear mechanics and the relative abundance of the lamin A is dependent on the tissue types 24 and matrix elasticity (8,9). More than 450 mutations have been discovered in the LMNA gene 25 (http://www.umd.be/LMNA/) which produces almost 16 different types of diseases like 26 Dilated Cardiomyopathy (DCM), Emery-Dreifuss Muscular Dystrophy (EDMD) etc. that are 27 collectively coined as laminopathies (10-12). These are tissue-specific in nature 28 predominantly affecting muscle and adipose tissues. The hallmark of laminopathies is the 29 formation of misshapen and even fragile nuclei (13). Two hypotheses are in vogue to explain 30 the role of mutant lamin A protein in the pathogenesis of laminopathies. The gene regulation

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31 hypothesis correlated the occurrence of aberrant gene expression due to differential 32 transcriptional regulation by lamin A while the structural hypothesis focuses on the 33 perturbation effect of mutant lamin A on higher order assembly of the nuclear lamina leading 34 to fragile nuclei (14,15). Previously, several reports based on AFM indentation, micropipette 35 aspiration and microrheological experiments showed altered nuclear elasticity for lamin A 36 null MEF or cells containing lamin A mutations (16-18). The rapidly growing area of 37 nucleoskeleton-cytoskeleton interactions has led to numerous studies involving the transduction of external mechanical cues to the nucleus. Earlier studies revealed that 38 39 cytoskeleton maintains the nucleus in a pre-stressed state by aligned actin filaments forming 40 a perinuclear actin cap (19-21). Recently, it has been shown for endothelial cells that central 41 and apical stress fibers play distinct mechanical roles in maintaining coordination between 42 the cell and nuclear shapes (20).

In this report, we studied the effect of nuclear morphology due to dilated cardiomyopathic 43 44 mutations (K97E, E161K, and R910W) under different physiological strains. These mutations were selected based on their severities of phenotypes in patients. The phenotype 45 46 of this disease is characterized by the cardiac arrhythmia with acute conduction defects and 47 myocardial infarction that can lead to sudden death (22). We took two different approaches to observe the effect of the laminopathic mutations (human origin) in the mouse myoblast 48 49 C2C12. Cells were stretched by applying cyclic strain (dynamic strain) on the PDMS membrane to mimic the physiological state of extension and relaxation of muscle and cardiac 50 51 cells. Secondly, cells were grown on the different micropatterned surfaces which exert static 52 deformation force (static strain) on the cytoskeleton via cell adhesion molecules (23,24) and 53 ultimately perturbs the lamin A network assembly. In both the cases, we observed 54 differential nuclear deformations, anchorage defect characterized by a tilt in nuclear axis 55 about actin axis and variation in the lamina thickness which suggested a reduction in nuclear rigidity and integrity. These results also suggest a probable explanation behind the 56 57 formation of elongated and misshapen nuclei in cardiomyopathy caused due to the LMNA 58 mutations.

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# 61 **RESULTS AND DISCUSSION**

#### 62 Differential impairment of the nucleo-actin axis on the micropatterned substrate

The actin cytoskeleton is known to mechanically couple with the nuclear lamins through 63 64 Linker of Nucleoskeleton and Cytoskeleton (LINC) complexes (25). We measured the alignment of the nucleus with the cell's orientation vector in cells expressing WT and mutant 65 66 Lamins A/C. Usually, the orientation vector of the cell reflects closely the orientation of actin stress fibers (26). However, well-spread cells often had random orientations of stress fibers 67 68 as is evident from cells on non-patterned glass (Fig. 1 A, left). To be able to control this 69 variation, we used substrates in which the adhesion area was patterned by photomasks (Fig. 70 1 A, right). Two rectangular patterns denoted by RA and RB with aspect ratios 3, 2 and spread 71 areas 1200, 450 µm<sup>2</sup> respectively, were chosen. These aspect ratios were chosen based on a 72 similar ratio (1:3) (27) of hPSC-derived monolayers of cardiomyocytes. To quantify the 73 alignment of the nucleus with the cell, the cell and nucleus outlines were fitted with ellipses 74 and the absolute angle between their major axes noted as orientation angle (Fig. 1 B) for cells 75 expressing the GFP tagged versions of lamin A (Fig. 1 C). For perfect alignment of the nucleus 76 with the cell axis, the expected orientation angle is 0°, while for randomly oriented nuclei, the angles are expected to vary from 0° to 90°, therefore averaging at 45° (dashed line, Fig. 77 78 1 D). As suspected, the non-uniform shapes in non-patterned cells resulted in angles ( $\sim 21^{\circ}$ ) 79 implying low alignment (Fig. 1 D). There was also no difference in orientation angles in mutant lamins compared to wild type lamin A (denoted by WT) (Fig1 D). However, using 80 81 micropatterns, we first observed lowered angles ( $\sim 10^{\circ}$ ) for WT as expected from the 82 uniformity of the shape (Fig. 1 D). Secondly, we found that all mutants exhibited the alignment of nucleus axis to cell axis within the range 24 ° - 37 ° (Fig. 1 D, Supporting Table 1) 83 rendering it closer to the value expected from random orientation (45°, Fig. 1 D, dashed line). 84 85 However, no particular parameter of nuclear shape (eccentricity, aspect ratio, circularity, -86 Fig. 1 D) was significantly affected by the mutations. This strongly suggested the lack of 87 nuclear alignment to cell-axis presumably due to a loss of mechanical coupling between the 88 nucleus and cytoplasmic actin network (Fig. 1 D, top). Furthermore, we observed the lamin 89 A aggregation inside the nucleoplasm and quantified the size in terms of area. K97E and 90 R190W transfected nuclei showed significantly larger aggregates, ~14 and 12 µm<sup>2</sup>

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91 respectively. WT and E161K nuclei showed smaller aggregates compared.  $\sim 2.5$  and 8.3  $\mu$ m<sup>2</sup> 92 respectively on the RB micropatterned surface. But on the RA pattern, E161K and R190W 93 nuclei produced smaller aggregates compared to RB,  $\sim$ 4.6 and 5.5  $\mu$ m<sup>2</sup> respectively (shown 94 in Supplementary Fig1). The size of aggregates for WT and K97E were unchanged,  $\sim 2$  and 95  $15 \,\mu\text{m}^2$  respectively. The aggregate may constitute other subtypes of endogenous lamin also 96 because of its homo- and hetero-polymerization nature but we only monitored GFP-97 fluorescence of the exogenous lamin A. Since all these experiments were performed by 98 transient transfection, that enhance the possibility of overexpression of protein lead to 99 misfolded protein and aggregation. We ruled out this possibility by measuring the non-100 significant change in the amount of lamin upon transfection through western blotting 101 (shown in Supplementary Fig 3).

## 102 Nuclear deformations after the cell stretching

103 In cardiac muscle tissues, the cells continuously experience the cyclic and uniaxial stretching. 104 Hence, the cells also reorient themselves towards the stretching axis (28.29). Zimmermann 105 et al. stretched neonatal rat heart cells up to 10% at 2 Hz frequency to generate engineered 106 heart tissue (30) and Yu et al. showed that 10% static stretching can insert new sarcomere 107 in the neonatal rat (31). The normal resting heart rate in infant and in the case of tachycardia 108 in adult heart rate can be more than 2 Hz (32,33). We intended to test the impact of chronic 109 mechanical perturbation (cyclic stretching) on lamin A-dependent nuclear morphology by 110 mimicking physiological frequency of extension and relaxation of muscles and cardiac cells. We 111 performed the mechanical stretching experiments on the wild-type (WT) and DCM-causing 112 mutant lamin A transfected C2C12 cells. The cells were grown on the PDMS membrane and 113 the membrane was stretched cyclically up to 10% for 2.5 hr. at a frequency of 2 Hz. We 114 measured how the mutants differed from the WT in shape parameters and orientation of 115 nucleus with and without cyclic stretching. In the absence of cyclic stretch, WT and mutant 116 cells grown on PDMS displayed non-significant differences in most (8/12) parameters 117 except for the observations that R190Wwas less aligned than WT while E161K resulted in 118 nuclei with higher eccentricity and aspect ratio and lower circularity. However, after cyclic 119 stretch was imparted most (10/12) parameters were found to be different – especially the 120 ones quantifying nuclei shape changes. Eccentricity and aspect ratio of all three mutants 121 increased on stretching in contrast to E161K which showed an increase even in the absence

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122 of stretching. Correspondingly, the circularity of all three mutants decreased on cyclic 123 stretching in contrast to only E161K showing a decrease in the absence of stretching. It must 124 be emphasized that, K97E showed a tendency to deviate from the WT in mechanically 125 unperturbed condition (Fig. 2). On cyclic stretching nuclear deformation of K97Ewas seen to 126 largest among all mutants. Thus, we quantified alignment of the nucleus with the cell body 127 (orientation angle difference) by employing micropatterning to reduce the initial 128 heterogeneity of cell shape in the population (reduced standard deviation of  $\theta$ , Supporting 129 Table 1). Here, we could elucidate that lamin A mutations may lead to weaker coupling of the 130 nucleus shape to the cell's shape (Supporting Table 2). Next, we demonstrated that when 131 these mutants undergo mechanical perturbation, they are less resistant to shape changes 132 and therefore underwent deformations, unlike WT nuclei that did not show any significant alteration in nucleus-to-cell alignment or shape on cyclic stretching (Supporting Table 3). 133 134 We have calculated the meshwork size of lamin A for WT and R190W (Supplementary Fig 2). 135 R190W nuclei dilated lamin A meshwork compared to the WT-nuclei which reflected the 136 lower mechanical rigidity of the nuclei. K97E and E161K nuclei produced lamin A aggregates 137 only.

### 138 **Reduction in the lamina thickness due to laminopathic mutation**

139 Nuclear stiffness is primarily determined by nuclear lamina meshwork. We measured lamina 140 thickness which could be the direct readout of stiffness as reported earlier as well (34). We 141 assessed the impact of laminopathic mutations on nuclear lamina thickening behavior. We 142 also included the effect of spreading as reported earlier that lowering in the spread area 143 reduces the stress fiber driven compression of the nucleus and nuclear volume which in turn 144 reduces physical strain on the nucleus (35). We chose two spread area (RA,1200  $\mu$ m<sup>2</sup> and 145 RB, 450  $\mu$ m<sup>2</sup>) which has ~ 3-fold spread area difference. We measured the lamina thickness 146 by structured illumination microscopy (SIM) whose lateral resolution was calculated to be 147  $\sim$ 120 nm (shown in Supplementary Figure 3). We measured average thickness 0.35 ± 0.03 148  $\mu$ m of the WT lamina on the RA patterned that reduces to 0.30 ± 0.03  $\mu$ m on RB surface 149 (shown in Fig. 3 A-D, left) suggesting thickening of the lamina on highly spread cells. We 150 believed that lamina in the low spread area cells is less stiff thereby resulting in the less 151 aligned network and showed lamina softening behavior. On comparing lamina thickness of

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152 WT with R190W and E161K transfected nuclei, we observed that the lamina thickness did 153 not alter for wild-type ( $\sim 0.3 \pm 0.05 \text{ }$  µm) and R190W ( $\sim 0.29 \pm 0.02 \text{ }$  µm), however, E161K 154 mutant lamina ( $\sim 0.23 \pm 0.03 \mu m$ ) showed significant softening behavior compared to WT on 155 the RB surface (Figure 3D). But on RA, the lamina for both E161K and K97E were not clearly 156 visible and both the nuclei showed a significantly large number of aggregates. All the values 157 are tabulated in the supporting table 3. The effect of the geometrical constraint on the K97E-158 lamin A transfected nuclei were similar. In both cases (RA & RB) for K97E nuclei, the lamina 159 was not clearly visible and nuclei showed larger aggregates. On the spread cell area (RA), the 160 mechanical strain perturbed the lamin A assembly for the mutant nuclei, hence, lamina 161 thickness was significantly decreased or abolished. This lamina softening behavior might 162 explain the lower mechanical rigidity of the nuclei among the cardiomyopathic patients.

## 163 **CONCLUSION**

164 In this article, we studied the nuclear deformation and misalignments of the nuclear axis 165 compared to the cell axis under different physiological strain. We chose different DCM 166 causing *LMNA* mutants (K97E, E161K, and R190W) and applied external mechanical strain 167 to mimic the physiological condition. In addition, we measured the variation in lamina 168 thickness which plausibly explained the origin of DCM disease and its severity. We observed 169 all the mutations produced significant nuclear deformations and showed significant 170 misalignments of nuclear axes when cyclic stretching was applied. On different 171 micropatterned surfaces, K97E and E161K nuclei produced significantly large lamin A 172 aggregates. Previously, it was reported that no apparent changes in the nuclear lamina 173 occurred for E161K mutation in heart tissue (36). As similarly overexpressed WT nuclei 174 showed less or no visible aggregates, hence the aggregation in the mutant nuclei was due to 175 the overexpression of the improperly assembled mutants lamin A. Previously, it has also 176 been reported that the perinuclear actin and TAN lines anchor the nucleus in pre-stressed 177 condition and help to maintain its orientation (37) and nesprin-1 (an important LINC 178 partner) also helps in tethering the nucleus to actin in C2C12 cells (38). A plausible 179 explanation for impairment of nucleo-actin axis might be the progressive weakening of 180 lamina at points such that the contact nodes with the actin filaments are abrogated thereby 181 resulting in impairment about the axis of the TAN lines. It is important to note that we had

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182 previously observed bundling behavior in E161K & R190W which lead to misshapen nuclei 183 with reduced viscoelastic properties. But in the case of the K97E, the lamin A network 184 formation was significantly altered which was reflected on the nuclear elasticity (39,40). 185 Therefore, the cell might not respond to the external mechanical cues effectively which in 186 turn can lead to altered cell response. This might be the reason behind significant nuclear 187 deformations under cyclic stretching. Furthermore, we studied the variation in lamina 188 thickness by SIM. The thickness of the lamina is in the range of  $\sim 20-50$  nm and electron 189 microscopic study revealed that lamina thickness can vary from ~18 nm to 100 nm in human 190 cartilage due to injury(41). Earlier, Schermmelleh et al. showed by 3D-SIM that thickness of 191 the lamin B1 thread is ~100 nm laterally and ~300 nm axially in C2C12 cells (because of its 192 resolution limit) (42). But in our experiments, lamina thickness was detected around  $\sim 0.3$ 193 um which could be due to both overexpression of lamin A and response to mechanical stress. 194 But within the resolution limits ( $\sim$ 120 nm) of our experiments, we confirmed the thickening 195 behavior of lamina. We observed that WT lamina thickness slightly increased on the 196 geometrically constrained surface area. But the mutant nuclei showed drastic effect. We 197 could not detect any rim staining of lamin A on the RA surface for the E161K mutant, instead, 198 they formed larger aggregates in the nucleoplasm. The lamina thickness for the R190W 199 mutant nuclei also decreased when cells were grown on the RB surface. We also observed 200 that the total lamin A expression did not change because of the mutation (Supplementary Fig 201 3). These results suggested that soluble lamin A might get recruited to the lamina in response 202 to an increase in the mechanical strain, but mutant lamin A showed the defect in the lamina 203 assembly. Therefore, R190W nuclei showed less thick lamina and E161K nuclei showed 204 larger lamin A aggregates in the nucleoplasm. Due to mechanical stress, changes in the lamin 205 A structure and the phosphorylation status may lead to a change in the lamina organization 206 (43,44). It is already established that at higher mechanical stress, lamin A accumulates at the 207 nuclear periphery that leads to the increment in nuclear stiffness and at low strain, changes 208 in the phosphorylation status produce more soluble lamin A. Hence, lamin A follows a 209 'mechanostat equilibrium' in the nucleoplasm (45). The change in lamina thickness could be 210 due to the accumulation of the lamin A in the aggregates and the change in the 211 phosphorylation pattern. The increase in WT lamina thickness due to the increase in

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212 mechanical stress suggested that changes in the phosphorylation profiles led to the 213 accumulation of the insoluble lamin A to the envelope. But the mutant nuclei lack this ability 214 leading to improper self-assembly and aggregate formation. Ectopic expression of lamin A in 215 the background of the endogenous lamins can significantly contribute to the nuclear stiffness 216 (46). But the background lamin expression was not sufficient to resist the lamina 217 deformation due to different mechanical strain. The aspect ratio of the micropattern was like 218 the hPSC-derived monolaver of the immature cardiomyocytes. The change in the nuclear 219 organization due to the geometrical constraint might be irreversible in the case of 220 cardiomyocytes which might lead to laminopathies. Based on our experimental findings 221 which we have summarized all the results in Supporting Table 1, 2 and 3. We hereby propose 222 that external mechanical cue can alter lamin A meshwork density significantly in presence 223 of laminopathic mutations; lamina shows stress induced thinning behavior and soluble lamin 224 A forms large insoluble aggregates inside the nucleoplasm. These factors cumulatively affect 225 the nuclear rigidity which alters the nuclear shape and mechanics. A plausible model for 226 occurring of laminopathy diseases due to single-point mutation is shown in Fig 4. In all 227 cases, we noticed differential alterations in the nuclear shape and the lamin A meshwork for 228 the DCM mutations. The mutant nuclei produced a severe deformation when exposed to the 229 external mechanical strain. In the physiological state also, we can predict that mutation of 230 lamin A in the tissues which are continuously exposed to the external strain can produce 231 significant damage to the nuclear shape and lamin meshwork. Eventually, that can lead to 232 differential gene expression programme and thereby produce the disease.

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## 241 **METHODS**

#### 242 Site-directed mutagenesis

- All the mutations were generated using site-directed mutagenesis methods in EGFP-LA plasmid and
- the details of the primer are reported in Bhattacharjee et al. (40). The mutations were confirmed by
- the Sanger sequencing method(47).

#### 246 Micropatterning glass coverslip

247 22 x 22 mm glass coverslips were etched in ethanol: acetic acid (19:1) mixture for 30 min, following 248 by ethanol washing and air drying. Dried coverslips were treated with UV Ozone cleaner (Jelight 249 Company, USA) for 5 min and incubated with 0.2 mg/ml PLL-g-PEG (SuSos, Switzerland) solution 250 (prepared in 10 mM HEPES buffer, pH 8.3) for 1 h. Photo-masks (JD, Photo Data, UK) is cleaned with 251 acetone and isopropanol followed by 5 min UV Ozone cleaning. PEG-coated coverslips are attached 252 on the chrome side of already cleaned photo-masks with the help of a drop of water. Excess water 253 was soaked with tissue paper for firm adherence of coverslip to mask. Coverslips attached with 254 photo-mask were illuminated with deep UV for 5 min by placing non-chrome side facing UV lamp. 255 Patterned coverslips were detached by floating into water followed by 45 min incubation in 20 ul 256 fibronectin solution (Sigma) (25 µg/ml solution in NaHCO<sub>3</sub>, pH 8.6). Finally, patterned coverslips 257 were used for cells seeding.

#### 258 Transfection and cells seeding

70 % of confluence C2C12 cells were transfected with 3 μg GFP lamin mutants by lipofection (Lipofectamine 3000, Life Technologies) following manufacturer instructions. Post 16 h of transfection, cells were detached from culture dish using 0.02 % EDTA solution (Calbiochem, USA) prepared in cell culture grade PBS (Sigma). 2 X 10<sup>5</sup> cells were seeded on each patterned coverslips and unattached cells washed off after 30 using warm media. For cell stretching, 10<sup>5</sup> cells per 500 μl culture media on each PDMS sheet were seeded. After transfection, Lamin A expression was checked mouse monoclonal anti-lamin A/C antibody at 1:1000 dilution.

#### 266 Cell Stretching

Custom made cells stretching device was employed for cell stretching as used earlier (ref). Briefly,
Silicon elastomer (SYLGRAD 184, Dow Corning) and the curing agent was mixed in 10:1 proportion,
an air bubble was removed by centrifuging 5 min at 3000 rpm. 3 ml mixture was spread over each
90 mm dish. Placed vertically to get excess mixture flow away. The dish at this configuration was

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271 baked at 60° C to get 100 um thick PDMS sheet. PDMS sheet was treated 0.5 mg/mlSulfo-SANAPAH 272 (Pierce chemicals) under deep UV for 5 min (UVO cleaner, Jelight, Inc., USA), functionalized with 273 fibronectin (25 µg/ml solution in NaHCO<sub>3</sub>, pH 8.6 Sigma). Fibronectin functionalized coverslips were 274 mounted and custom builds stretchers, GFP Lamin transfected cells were seeded on 25 x 25 mm area 275 encircled with silicon grease. Cells were grown for 24 h on PDMS loaded stretcher inside the 276 incubator. For cyclic cell stretching, PDMS sheet containing C2C12 cells grown in a well was attached 277 to two motors (Physik Instrumente (PI), GmbH and Co KG) using custom-designed adapters. Cyclic 278 stretching at 2 Hz frequency and 10% amplitude was performed with MATLAB for 2.5 h. Stretching 279 was performed inside the cell-culture incubator.

#### 280 Cell fixation and microscopy

281 C2C12 cells either grown on glass or PDMS sheet were fixed using 4% PFA on their respective 282 surfaces for 12 min, following PBS wash. Cells were permeabilization by 0.2 % Triton X-100 solution 283 (Sigma) in PBS, stained with DAPI (2.8  $\mu$ M, Sigma) and Phalloidin Alexa Fluor 568 (0.35  $\mu$ M, 284 Molecular probe, Life Technologies) for 2 h at room temperature. Finally, coverslips were mounted 285 on with slides (Sigma) for imaging. In case of cells grown on a PDMS sheet, a number one glass 286 coverslips were first attached on top of the PDS sheet with mounting media (Mowiol, Sigma). The 287 entire stretching device was flipped upside down and captured images using oil objective through 288 the glass. Z-stack (0.5 µm step size) images were captured by Olympus epi-fluorescence microscope 289 (Olympus Corporation Japan) with 100X, 1.49NA objective and sCMOS camera (Orca-Flash 4.0, 290 Hamamatsu Photonics Japan) with pixel size 65 nm.

# 291 Nucleus to cell orientation angle difference and nucleus shape parameter extraction

Image analysis was done using software ImageJ/Fiji ((https://imagej.net/Fiji)). Nucleus (GFP lamin) to cell (actin) orientation angle difference was measured in two-parts, first drawing ROI manually on maximum intensity projection image to outline object, second fitting ROI to an ellipse for measuring orientation angle. Major (D1) and minor (D1) axis of the fitted ellipse on the nucleus was used for eccentricity ( $\sqrt{(1 - \frac{D2}{D1})^2}$ ) and aspect ratio (D1/D2) measurement. Perimeter and area of object (nucleus) outline on maximum intensity projection image were used for circularity ( $4\pi \times \frac{Area}{Perimeter^2}$ ) measurement.

#### 299 Mesh size and width of the peripheral lamin measurements

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300 Cells were washed three times with ice-cold 1X PBS (pH 7.4) and fixed with 4% paraformaldehyde. 301 Actin was stained with Alexa-561 conjugated phalloidin (Thermo Fisher Scientific) with 1:1000 302 dilutions. After proper washing with PBS, the coverslips were stained with Vectashield that contained 303 DAPI (Vector Laboratories). The coverslips were sealed with watercolor nail polish. The slides were 304 visualized with NIKON Inverted Research Microscope ECLIPSE TiE with Plan Apo VC 100X oil DIC N2 305 objective/1.40 NA/1.515 RI with a digital 4X zoom. Images were analyzed using Ni Elements AR Ver 306 4.13 and Image J (Fiji). The X-Y resolution of N-SIM was calibrated to 120-150 nm using 20 nm 307 FluoSphere beads (Thermo Scientific, F887) in 580 nm laser. We measured the mesh sizes using the 308 area selection tool of Ni-Elements software, but the lines were drawn considering the FWHM method 309 where lower values were considered as the limit of the sides. To measure the lamina thickness, first 310 nuclear rim staining of lamin A were straightened using Image I and then lines were drawn across 311 the linearized lamina and intensity profiles were plotted. From the intensity profile, FHWM was 312 considered to be the thickness or width of the lamina. The thickness of each nucleus was calculated 313 by averaging 10 randomly chosen line profiles and more than 15 separate nuclei were considered for 314 each condition. Sizes of aggregates were measured using an area selection tool of Ni-Elements. The 315 statistics were generated using Origin Pro 8.5 software.

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# 322 AUTHORS CONTRIBUTION

- 323 KSG, BS, MB & RK designed all the experiments. RK & MB performed all the experiments. KSG, MB,
- RK & BS analyzed the data and wrote the paper. KSG conceived the entire project and responsible for
- 325 finances related to the project.

# 326 **COMPETING FINANCIAL INTERESTS**

327 The authors declare no competing financial interests.

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# **FIGURE LEGEND**

Figure 1. Lamin A mutations lead to the altered alignment of the nucleus with the cell axis. (A) Schematic representations depicting variations possible in the nucleus and cell shape along with nucleus alignment with cell axis in non-patterned (grown on glass) (left) and patterned cells (right). Ellipses are fits to cell-outline and nucleus-outline; the corresponding major axes represented as dashed lines. Wide-field transmission images of patterns on photomasks used to create micropatterns – RA (60  $\mu$ m x 20  $\mu$ m) and RB (30  $\mu$ m x 15  $\mu$ m) are shown here. Scale bar, 50  $\mu$ m. (B) Schematic representation of cell to nuclear orientation angle ( $\theta$ ) and major (D1) and minor (D2) diameters of nucleus used for calculating nuclear eccentricity and aspect ratio. (C) Representative images of cells expressing GFP-Lamin A/C (green) mutants stained with DAPI (blue, DNA) and Phalloidin Alexa Fluor 568 (red, F-actin). Images show basal plane for actin and the maximum intensity projection for DAPI and GFP-Lamin A/C. Scale, 10  $\mu$ m. (D) Orientation angle difference ( $\theta$ ), eccentricity, aspect ratio, and circularity of nucleus measured from three different conditions Glass (left column), RA pattern (middle column), RB pattern (right column). The asterisk marks significance (p < 0.001) difference when compared with WT cells grown on the glass. n = 27, 32, 52, 21 cells for glass from one experiment, n = 70, 41, 48, 55 cells for RA in three independent experiment, n = 41, 37, 52, 36 cells for RB in two independent experiment. \* p < 0.05, \*\* p < 0.01, \*\*\* p < 0.001, Wilcoxon based Mann-Whitney U test were performed for statistical testing.

**Figure 2.** Nuclei of cells with lamin mutations are less resistant to cyclic stretch. Blue, green and red channels in images denote DAPI, GFP-Lamin A transfection and Phalloidin staining in C2C12 cells with wild type (WT) and mutant Lamin A/C (WT, K97E, E161K, R190W). Cells were either (A) grown on PDMS and fixed – without imparting cyclic stretch or (B) grown, imparted cyclic stretch (10% cyclic stretched with 2 Hz frequency for 2.5 h.) and subsequently fixed. Scale, 10  $\mu$ m. The arrow shows the direction of stretching. (C) Orientation angle difference ( $\theta$ ), eccentricity, aspect ratio, and circularity of nucleus measured from unstretched and stretched conditions. n = 26, 80, 72, 119 cells for unstretched and n = 77, 55, 77, 123 cells for stretched condition in three independent experiment. \* p < 0.05 \* p < 0.01, \*\* p < 0.001, Wilcoxon based Mann-Whitney U test were performed for statistical testing.

**Figure 3. Reduction in the lamina thickness due to laminopathic mutation.** eGFP-lamin A (containing wild-type and R190W, E161K & K97E mutations) transfected cells were grown on RA (A) and RB (C) patterned surface. The lower panels of A and C are representing linearization of the nuclear lamina for wild-type and mutant nuclei. The representative FWHMs of the intensity profiles

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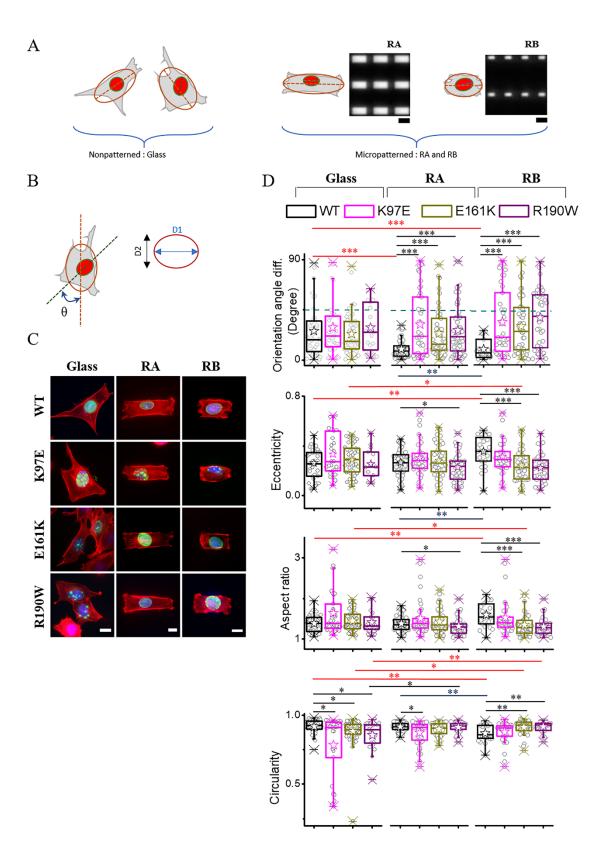
of white the line across the linearized lamina (after Gaussian Fittings) are shown in panel B for RA and in panel D for RB surface. The box plots in the panel B and D represent the FWHM (thickness) of the lamina for RA and RB micropatterned surface respectively. Each dot in the box plot denotes the raw data obtained from each nucleus. The scale bar is 5  $\mu$ m. \*\*\*p<0.001, \*\*p<0.05, ns= not significant.

#### Figure 4. Model for lamin meshwork in the presence of mechanical cue in laminopathic cells.

In presence of the external mechanical cue, wild-type lamin A form dense meshwork inside the nucleoplasm and width of the lamina also increases. In the case of laminopathic nuclei, lamin A meshwork density decreases and the lamina also shows thinning behavior in the presence of the same external mechanical force. The nuclear axis also shows large impairment along the acting stress fiber compared to the wild-type nucleus.

Nuclear deformations in cardiomyopathic nuclei

# Figure 1

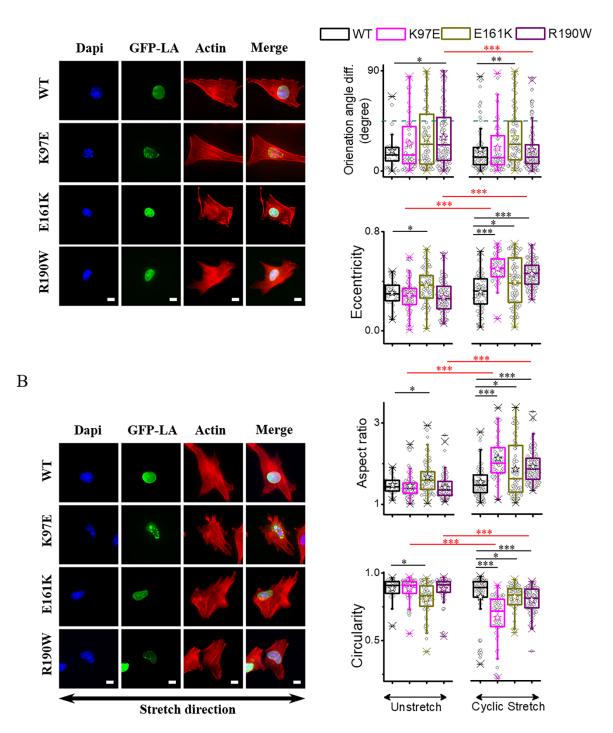


Nuclear deformations in cardiomyopathic nuclei

# Figure 2

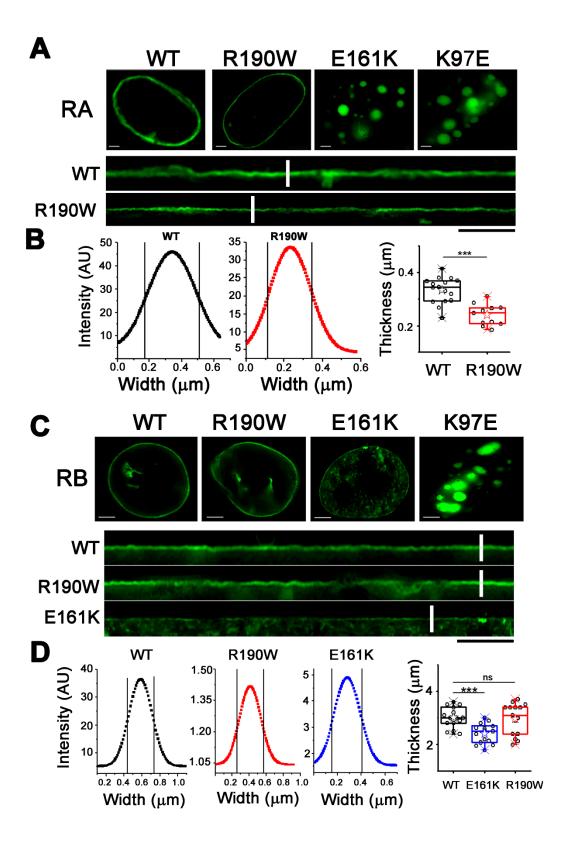


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Nuclear deformations in cardiomyopathic nuclei

# Figure 3



Nuclear deformations in cardiomyopathic nuclei

# Figure 4

