Cell monolayers sense curvature by exploiting active mechanics and nuclear mechanoadaptation

1   **Large-scale curvature sensing by epithelial monolayers depends on active**

2   **cell mechanics and nuclear mechanoadaptation**

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22   **Abstract**

23   While many tissues fold *in vivo* in a highly reproducible and robust way, epithelial folds

24   remain difficult to reproduce *in vitro*, so that the effects and underlying mechanisms of local

25   curvature on the epithelial tissue remain unclear. Here, we photoreticulated polyacrylamide

26   hydrogels through an optical photomask to create corrugated hydrogels with isotropic wavy

27   patterns, allowing us to study how concave and convex curvatures affect cellular and nuclear

28   shape. We find that the substrate curvature leads to thicker epithelial zones in the valleys and

29   thinner ones on the crest, a feature which we show generically arises in vertex models,

30   leading us to hypothesize that curvature-sensing could arise from resulting thickness

31   modulations. We further show that concave and convex local curvatures lead to significant

32   modulations of the nuclear morphology and positioning, a feature well-explained by an

33   extension of vertex models taking into account membrane-nucleus interactions, where

34   thickness modulation generically translate into corresponding changes in nuclear aspect ratio

35   and position. Consequently, we find that the spatial distribution of Yes associated proteins

36   (YAP), the main transcriptional effector of the Hippo signaling pathway, is modulated in

37   folded epithelial tissues according to resulting nuclear density changes. Finally, we showed

38   that these deformations are also associated with variations in the relative abundance of A/C-

39   versus B1-type lamins, significant chromatin condensation and lower cell proliferation rate.

40   These findings show that active cell mechanics and nuclear mechanoadaptation are key

41   players of the mechanistic regulation of epithelial monolayers to substrate curvature, with

42   potential application for a number of *in vivo* situations.

43

Introduction

In living systems, epithelial tissues are commonly described by three-dimensional (3D) microstructures such as invaginations, folds or wavy morphologies. Curved surfaces are also observed at interfaces between tissues or at boundaries between tissues and body lumen. The geometric form and biological function of wavy epithelial tissues are inherently linked together at all scales. For instance, crypts and villi of the small intestine provide a large surface area for exchange, improving the absorbance function. Despite its wide interest, the relationship between curvature and biological function in epithelial tissues remains largely unexplored [1,2].

Despite numerous studies on the influence of cellular and subcellular-scale topography on cell fate [3–5], few studies have investigated the effect of curvature on collective cell behavior [6–9], in particular because of the technical limitations encountered to engineer soft culture substrates with curved patterns in a controlled way [10]. It remains unclear in particular whether and how curvatures at scales much larger than cell size could be sensed biologically. Early studies conducted on glass fibers have shown that cells orient themselves along the line of minimal curvature to minimize cytoskeletal deformations [11,12]. Glass wires were also used to decouple the effect of out-of-plane curvature from the lateral confinement experienced during the migration of epithelial tissues [13]. More recently, it was shown that isolate adherent cells avoid crests of ultra-smooth sinusoidal surfaces during their migration and position themselves in valleys [14] and that the persistence and speed of migration of single cells can be affected by substrate curvature [15].

Despite these efforts, our current understanding of the role of the curvature of epithelial tissues remains elusive. To answer this question, we developed well-defined soft corrugated hydrogels to investigate the response of epithelial tissues to variations of curvature. By combining *in vitro* experiments with analytical and computational vertex

69 model, we show that local changes of monolayer thickness and cell density can be interpreted

70 by energy minimization arising from the mechanics of apico-lateral tensions. We extended the

71 vertex model to consider in a minimal way changes of nuclear morphology due to active

72 tensions and cell shape, suggesting a simple mechanism via which thickness modulation

73 couple to the experimentally observed changes in nuclear shape and positioning. We then

74 showed that these changes triggered by curvature also lead to high nuclear/cytoplasmic YAP

75 ratios, demonstrating a YAP-curvature sensing of epithelial tissues, which is inhibited at high

76 cell density. Furthermore, we showed that nuclear deformations observed on corrugated

77 matrices were associated with a modulation of lamin A/C (LMAC) and B1 (LMB1), leading

78 to a lower expression of LMAC on positive curvatures and a higher expression of LMB1 on

79 negative curvatures. Finally, we demonstrate that matrix curvature can be considered as an

80 important regulatory cue of epithelial tissues, leading to high level of chromatin compaction

81 and a lower DNA synthesis rate in negative curvature zones, which correspond to high cell

82 density.

83

84 **Results**

85

86 **The overall cytoskeletal architecture of wavy epithelial monolayers is not affected by the**

87 **curvature.**

88 To study how a monolayer of epithelial cells adapt to convex and concave cell-scale

89 curvatures of their matrix (Fig. 1A), we generated corrugated hydroxypolyacrylamide

90 (hydroxyPAAm) hydrogels by photopolymerizing an hydroxyPAAm solution at 360 nm

91 through a chromium optical photomask. The polymerization was completed in transparent

92 zones and the slow diffusion of the photoinitiator towards non-illuminated zones lead to the

93 formation of a smooth and wavy profile. We used various widths of transparent and black

4

stripes to form isotropic corrugated hydrogels of 250±30 kPa with wavelengths of 20 μm ($\lambda20$), 30 μm ($\lambda30$) and 50 μm ($\lambda50$). We obtained symmetric corrugation patterns that formed wavy epithelial monolayers showing periodic wavelengths of 20 μm (Supplementary Movie S1), 30 μm (Supplementary Movie S2), 50 μm (Supplementary Movie S3) and constant amplitudes over large areas (10×10 mm^2).

To determine whether curvature changes can affect epithelial thickness and nuclear organization (Fig. 1A), we defined 3 zones of interest, as crests, valleys and interm. zones. Convex (positive) curvature zones correspond to the crests, concave (negative) curvature zones to the valleys and zero curvature zones to the interm., which corresponded to the junction between convex and concave zones (Extended Data Fig. 1A-B). We determined local maximal curvature values, amplitude (β) and wavelengths (λ) of convex and concave curvature zones using 2D (*xz*) profiles obtained from atomic force microscopy and high-resolution confocal microscopy (Extended Data Fig. 1A-C and Extended Data Table 1).

Corrugated hydrogels were functionalized with human fibronectin (FN) and Madin-Darby Canine Kidney (MDCK) cells were cultured at confluency (1×10^3 cells/mm^2) on flat hydrogels, $\lambda20$, $\lambda30$ and $\lambda50$ corrugated hydrogels for studying how convex and concave cell-scale curvatures can affect epithelial monolayers. After 48 hours in culture, wavy epithelial monolayers were immunostained with Alexafluor 488 for F-actin (Fig. 1B), 4′,6-diamidino-2-phenylindole (DAPI) for the nucleus (Fig. 1C) and imaged with a laser-scanning confocal microscope. Using Z-stack projections from confocal scanning of F-actin (Extended Data Fig. 2A-C), we found that the actin intensity was not significantly different between flat and corrugated ($\lambda20$, $\lambda30$ and $\lambda50$) hydrogels monolayers (Extended Data Fig. 2D, suggesting that corrugations of the matrix do not affect the global amount of F-actin in epithelia. We next studied the organization of cell-cell adhesive interactions in folded epithelial tissues by staining for ß-catenin, which is involved in regulation and coordination of MDCK cell-cell

119 adhesions[16]. ß-catenin staining showed that epithelial tissues remained cohesive on corrugated

120 hydrogels (Extended Data Fig. 3A-C) with well-defined cell-cell contacts between polyhedral

121 cells, independently of local concave or convex curvatures. We skeletonized ß-catenin stained

122 images (Extended Data Fig. 3D-F) to determine a mean cell area of 205.3 ± 22.8 μm^2 on flat,

123 196.7 ± 27.7 μm^2 on λ20, 183.8 ± 17.8 μm^2 on λ30 and $186.8\ 8 \pm 30.7$ μm^2 on λ50 (Extended

124 Data Fig. 3G), suggesting that corrugations did not change the mean cell area. Furthermore,

125 we found that epithelial cells were characterized by a mean side number of 5.5 ± 1.7 on flat,

126 5.8 ± 1.3 on λ20, 5.8 ± 1.3 on λ30 and 5.8 ± 1.3 on λ50 (Extended Data Fig. 3H) and that the

127 mean cell area increased with the polygon class (Extended Data Fig. 3I), regardless the

128 corrugation wavelength.

129 Altogether, these results demonstrate that wavy epithelial monolayers maintain their

130 overall architecture intact and can adapt to corrugated matrices without changing the mean

131 cell shape nor the global expression of actin.

132

133 **Epithelial thickness modulation from substrate curvature.**

134 However, when examining the shape of the cell monolayer as a function of curvature

135 (Fig. 1A), we found striking differences between convex zones (i.e. crests) and concave zones

136 (i.e. valleys). In λ20, λ30 and λ50 tissues, cells on crests displayed reduced cell height and

137 increased cell area (squamous-like), while cells in valleys were thicker and denser (columnar-

138 like). Interestingly, similar density differences were observed in human keratinocytes [4],

139 arguing this could be a general response to curvature. Given that a number of mechano-

140 sensitive cellular responses, such as YAP localization, depend on cell thickness/density [17,18] as

141 well as curvature [3], we thus hypothesized that curvature could be sensed indirectly by creating

142 thickness/density differences, and turned to a theoretical model to understand how curvature

143 could create such cell shape modulations.

144   For this, we used a vertex model [19–23], which describes the shape of epithelial

145   monolayers based on their apical, lateral and basal tensions [24,25]. We first reduced the problem

146   to 2D by neglecting the in-plane component that is translationally invariant, and assumed that

147   the basal surface could not detach from the substrate [26] (with profile $\beta\cos(qx)$ where β is the

148   amplitude of the substrate and q the wavevector), thus removing its contribution to the

149   energy. Importantly, apart from measurable geometrical quantities such as cell density or

150   volume, we found that the response of the monolayer to substrate curvature depended on a

151   single rescaled parameter, the ratio of apical to lateral tensions Γ_a/Γ_l (Fig. 1A). Indeed, when

152   apical tensions dominate, the configuration with minimum energy is the one with flat apical

153   surface (leading to maximal thickness modulation, equal to substrate height modulation).

154   However, this "flat-apical" configuration increases drastically total lateral area and is thus

155   unfavorable for large lateral tensions. In that converse case, the epithelium tends to be of

156   constant thickness, independent of curvature. In general, denoting $\alpha\cos(qx)$ the equation for

157   the apical surface, the rescaled thickness modulation $\Omega = -(\alpha - \beta)/\beta$ is thus predicted to

158   follow the law:

159
$$\Omega = \frac{\frac{1}{2}\frac{\Gamma_a}{\Gamma_l}V_c q^2 - \frac{1}{4}\Delta h^2 q^2}{1 + \frac{1}{2}\frac{\Gamma_a}{\Gamma_l}V_c q^2 - \frac{1}{8}\Delta h^2 q^2} \quad (1)$$

160   which interestingly is independent of substrate amplitude β, allowing for simpler

161   investigation of the specific effect of different substrate wavelengths (V_c is the cell volume

162   and Δh the average thickness, see Supplementary Theory Note for details). We checked this

163   analytical theory via numerical simulations of the 2D vertex model (on a fixed basal

164   substrate) by varying multiple parameters such as wavelength, amplitude, tensions, and found

165   that Eq. (1) matched well with numerical simulations at large wavelength, with small-

166   wavelength corrections (Fig. 2A and Extended Data Fig. 4A-E).

167   Turning back to the data, we quantified this effect by projecting tissues along the

168   direction of the pattern, thus building average intensity profiles in response to a curved

169   substrate in each sample, from which we could extract the average basal and apical surfaces

170   (Fig. 2B), and thus calculate Ω. Importantly, we found that thickness modulations Ω were

171   largest for $\lambda 20$ and smallest for $\lambda 50$, as predicted theoretically (Fig. 2C). More quantitatively,

172   the model could predict well the experimentally observed shape of the apical surface (Fig.

173   2D), and the evolution of the thickness modulation Ω across wavelength $(\lambda 20, \lambda 30$ and $\lambda 50)$

174   was well-predicted by Eq. (1) with the single fitting parameter $\frac{\Gamma_a}{\Gamma_l} = 0.6 \pm 0.2$ (with all other

175   parameters, such as cell seeding density, substrate deformation amplitude β, average cell

176   thickness, being constrained from independent measurements, see Supplementary Note for

177   details). Interestingly, even for substrate with strongly non-symmetric profiles (e.g. the

178   substrate with larger wavelength of 100 μm in Extended Data Fig. 5A), although the

179   analytical Eq.1 could not readily be applied, we found that vertex model simulations (taking

180   this more complex substrate shape as an input, with all other parameters kept the same as $\lambda 20$,

181     $\lambda 30$ and $\lambda 50$) still gave good agreement with the experimental shape profile (Extended Data

182   Fig. 5B). Furthermore, one important prediction of the theoretical model is that the rescaled

183   thickness modulation $\Omega = -(\alpha - \beta)/\beta$ does not change with substrate amplitudes, i.e.

184   doubling the substrate amplitude doubles the absolute thickness modulation (so that the ratio

185   of the two does not change). To further test this prediction, we varied the amplitudes of

186   corrugated hydrogels of 20 μm and 50 μm wavelength, so they both have substrate

187   amplitudes either distinct from or close to a fixed amplitude of ~2.3 μm (i.e. the amplitude of

188   30 μm wavelength substrates) (Extended Data Fig. 5C and Extended Data Table 1). We were

189   therefore able to compare the thickness modulation of varied substrate wavelengths (20, 30

190   and 50 μm) with a fixed amplitude or varied substrate amplitude with the same wavelength.

191   Importantly, we found no statistically significant differences when comparing the normalized

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192 thickness modulation (Ω) of λ20/1.2 with λ20/2.3 and λ50/2.4 with λ50/3.2 (Extended Data

193 Fig. 5D). This confirmed our model prediction, where amplitude changes have a linear effect

194 on thickness modulation (resulting in an invariance rescaled thickness modulation) and allow

195 us to compare rescaled thickness modulations for different wavelengths (as previous analysis

196 of Fig. 2C).

197   Altogether, the theory suggests that active contractile apico-lateral tensions could be a

198 very simple physical mechanism converting changes in substrate curvature into changes in

199 cellular thickness/densities.

200

201 **Substrate curvature modulates the spatial distribution of nuclei.**

202   Next, we asked whether modulation of epithelial thickness in response to substrate

203 curvature changes would impact the spatial distribution of nuclei. To answer this question, we

204 extended our vertex model to consider the mechanical interaction between the nucleus

205 (modelled as a deformable sphere under tension) and cell surfaces (Extended Data Fig. 4F for

206 sketch, see Supplementary Theory Note for details). Intuitively, apical surface in convex

207 zones (with thin monolayer) were expected to compress nuclei in-plane, whereas apical

208 surface in concave zones (with thick monolayer) were expected to compress nuclei along the

209 apico-basal axis (Extended Data Fig. 4G-I see also Supplementary Theory Note for analytical

210 expressions). Furthermore, in the presence of cell thickness gradient (which arises generically

211 from thickness modulation), nuclei are expected to move along the thickness gradient towards

212 large thicknesses to minimize nuclear deformation, and we predicted a linear relationship

213 between thickness gradient and nuclear displacement away from the cell center of mass

214 (Extended data Fig. 4I and Supplementary Theory Note).

215   Interestingly, clear differences between nuclear and cellular centers of mass in

216 corrugated epithelia were observed on confocal images (Fig. 3A), suggesting that nuclei were

217   attracted from intermediate zones to concave ones (valleys). We therefore defined a vector

218   between two centers of mass representing "nuclear offset" (Fig. 3A-B), as a function of

219   normalized position $\bar{x} = \frac{x}{\lambda}$ ($\bar{x}$=0 and 1 corresponded to the convex regions and $\bar{x}$=0.5 to the

220   concave region). To test whether the vertex model can predict these changes of nuclear

221   localization, we computed in both model and data the nuclear offset and confirmed that the

222   experimental data also followed a sinusoidal dependency with position x for all wavelengths.

223   As predicted by the model, nuclear offset was directed towards concave zones (i.e. valleys),

224   away from convex ones (i.e. crests), and maximal on the intermediate zones (Fig. 3C).

225   Importantly, the amplitude of the sinusoidal of the offset depended on one fitting parameter

226   (quantifying the relative nuclear/surface tensions, see Supplementary Theory Note for

227   details), which allowed us to jointly fit the λ30 and λ50 data (as expected, λ50 showed lower

228   amplitude than λ30).

229   One discrepancy was that the continuum model predicted that λ20 should have an

230   even larger offset than λ30, while the opposite was found in experiments (Fig. 3C). However,

231   turning to discrete vertex model simulations, we found that this was expected when

232   considering such small wavelengths. Indeed, for small wavelengths (i.e. on order of cellular

233   size), we found that numerical simulations generically predicted a "quantization" of the

234   nuclear response with substrate wavelength, i.e. when the substrate wavelength represents

235   integer values of cell lengths (Fig. 3D-G). For instance, for 2 cells per wavelength (close to

236   the λ30 scenario, Figs. 3F,G), the energy minimum of the vertex model is to have each cell

237   spanning half-a-period between one valley and one crest, maximizing the thickness gradient

238   and allowing nearly full localization of nuclei in concave zones (valleys). This cannot occur

239   for non-integer values (Fig. 3D), which can occur for λ20 substrates (Fig. 3E, see

240   Supplementary Theory Note for more detailed discussion), explaining the lower nuclear offset

241   compared to λ30 in this condition. We also tested whether other features of the model, such as

242   the curvature-dependent nuclear deformation observed in data and found that nuclear

243   projected area (Extended data Fig. 6A) followed closely the theoretical prediction of a

244   sinusoidal dependency as a function of x, with largest (resp smallest) projected nuclear area

245   (xy plane) in convex (resp. concave) regions for all wavelengths (Extended Data Fig. 6C-E),

246   i.e. the spatial dependency of projected area was similar to offset, but with a 90° phase

247   difference (see Supplementary Theory Note for details). Altogether, these quantitative

248   analyses strengthen the mechanism we propose where thinner convex zones compress nuclei

249   in-plane, whereas thicker concave zones compress nuclei along the apico-basal axis.

250   Based on this finding, we also examine the third dimension (y-direction), along the

251   corrugations, and found that the corrugation dimension modulated the orientation of nuclei.

252   Interestingly, our results showed that the nuclei densely accumulated in the concave zones

253   (i.e. valleys, Extended Data Fig. 6B) were significantly more likely to align along y-axis

254   (Extended Data Fig. 1A), with angle of orientation 15.8±6.2° on λ30 (Extended Data Fig.

255   7A), whereas nuclei formed a wider orientation distribution on convex regions (i.e. crest),

256   with angle 49.9±17.1° on λ30 (Extended Data Fig. 7B).

257   Altogether, the combined quantitative analyses and experimental measurements reveal

258   that epithelial thickness modulation due to substrate curvature can affect nuclear positioning

259   and deformation. In the following, we further explore the accompanying nuclear biochemical

260   changes arising from curvotaxis.

261

262   **YAP-curvature sensing is mediated by nuclear density modulation.**

263   Recent works have demonstrated that direct application of forces to the nucleus was

264   sufficient to regulate the activity of YAP, a central regulator of cell proliferation and fate, by

265   controlling its transport through nuclear pores [27–29], as well as a number of nuclear mechano-

266   transduction events being increasingly recognized [30,31]. Given the nuclear positioning and

267 morphological changes we observed, we take a closer look at how YAP localization is

268 affected by curvature. Interestingly, YAP is well-established to adapt its nuclear-cytoplasmic

269 localization as a function of cellular density [18], but has also been shown to be dependent on

270 substrate curvature via unknown mechanisms [4]. Given our findings that curvature generically

271 translates into cellular density changes (as well as nuclear density changes reinforced by

272 nuclear offset), we thus propose a hypothesis that the curvature sensing of YAP could be a

273 consequence of the nuclear density modulation.

274   To examine the impact of substrate corrugations on YAP translocation, epithelia were

275 grown at 1×10^3 cells/mm^2 on flat, λ20, λ30 and λ50 wavy hydrogels and immunolabeled

276 after 48h for YAP and DNA (Fig. 4A). We performed high magnification confocal imaging

277 experiments to quantify the nuclear/cytoplasmic YAP ratio in epithelial cells located on

278 convex, interm. and concave zones. Our results showed that the nuclear/cytoplasmic YAP

279 ratio increased on convex (crest) curvatures but decreased on concave (valleys) ones (Fig. 4B

280 and Extended Data Fig. 7C-E). At constant curvature, we found that the nuclear/cytoplasmic

281 YAP ratio was larger on convex (2.45 ± 0.47 for C=0.27 on λ30) than on concave (2.00 ± 0.38

282 for C=-0.27 on λ30) curvature zones (Fig. 4A), as predicted from the decreased cellular

283 density there (Extended Data Fig. 6B). Our results are thus consistent with the hypothesis that

284 YAP curvature-sensing could occur indirectly from density-sensing. To test this further, we

285 reasoned that YAP curvature-sensing should then be absent for higher overall densities, as in

286 the model density then becomes high even in crests and overall rescaled thickness

287 modulations are reduced (see Supplementary Theory Note and Extended Data Fig. 4A).

288   We thus studied very dense epithelial tissues (5×10^4 cells/mm^2) on flat, λ20, λ30 and

289 λ50 hydrogels [32,33], and found a low (<1) nuclear/cytoplasmic YAP ratio, which was

290 unaffected by the substrate curvature (Fig. 4C), confirming that high cell density can inhibit

291 YAP curvature-sensing from epithelial cells. Furthermore, superimposing the mean

292   nuclear/cytoplasmic YAP ratio on flat, λ20, λ30 and λ50 substrates versus the local nuclear

293   density (concave and convex zones) confirmed the hypothesis that YAP curvature-sensing

294   could occur indirectly from density-sensing (Fig. 4D).

295   Finally, we sought to apply our theory to previously published dataset on keratocytes,

296   and found that we could again explain multiple features of these within our theory (with

297   larger apical/lateral tension ratio $\Gamma_a/\Gamma_l=5$, see Supplementary Note and Extended Data Fig.

298   4D), including the magnitude of valley/crest density differences, and the delamination of cells

299   from the crests above a critical value of substrate curvature (see Extended Data Fig. 4D and

300   Supplementary Theory Note for details). This argues for a generality of such a mechanism of

301   density/thickness changes from curvature, with impact on YAP.

302

303   **Substrate curvature modulates the relative abundance of nuclear lamina.**

304   Given our findings that substrate curvature leads to differential nuclear deformations,

305   we sought to evaluate the lamina composition in deformed nuclei, as a potential mechano-

306   sensitive response [34]. Indeed, accumulating evidence shows that the nuclear mechanical

307   properties are viscoelastic [35] and controlled the stoichiometric ratio between B1-type (LMB1)

308   and A/C-type (LMAC) lamins [36,37,38]. By scanning confocal microscopy, we examined the

309   fluorescence intensity of A/C-type (TRITC in red) and B1-type (FITC in green) lamins on

310   convex, interm. and concave zones of λ20 corrugated hydrogels (Fig. 5A). Z-stack images

311   using a ×60 objective were collected for three channels (DAPI, TRITC and FITC) from the

312   entire volume of the nuclei using a step size of 0.15 μm. Exposure times and laser power were

313   kept constant and acquired stack of images was deconvolved to remove out of focus light.

314   Using Z-stack images of nuclei located on convex (Supplementary Movies S4 and

315   S5), interm. (Supplementary Movies S6 and S7) and concave (Supplementary Movies S8 and

316   S9) zones (Fig. 5B), we estimated the mean integrated density of DAPI, LMAC and LMB1

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(Fig. 5C) for each Z-plane and calculated the LMAC/LMB1 integrated density ratio to characterize the relative abundance of A/C- versus B1-type lamins. For each nuclear morphology, we defined three Z planes of interest: Z_0 (basal plane), Z_1 (mid plane) and Z_2 (apical plane), respectively at 15%, 50% as 85% of the total nuclear height (Fig. 5B). The mean integrated densities of DAPI, LMAC and LMB1 versus the nuclear height for the three nuclear morphologies (n=8 for each with N=3 replicates) were characterized by Gaussian-like distributions. The LMAC/LMB1 ratio for convex zone was slightly below 1 for the apical part (Z_0, Fig. 5D) and close to 1 for $Z>Z_1$, indicating that the LMA abundance was lower on convex curvatures. For nuclei in interm. zones, the LMAC/LMB1 ratio was ~1.7 at Z_0, then decreased abruptly to Z_1 to reach a value of ~0.5 for $Z_1<Z<Z_2$. The asymmetric nuclear shape on interm. zones were characterized by more LMB1 in the apical part ($Z_1<Z<Z_2$) and more LMAC in the basal one ($Z_0<Z<Z_1$). Interestingly, the relative abundance of LMAC was significantly higher for nuclei in the concave zones, with maximal values at Z_0 (basal) and Z_2 (apical) focal planes (Fig. 5D).

Taken together, our results indicated that the nuclear deformations lead to a higher abundance of B1-type lamins on convex curvatures (i.e. crests) and a higher abundance of A/C-type lamins on concave curvatures (i.e. valleys), suggesting that the nuclear deformations induced by substrate curvature changes are accompanied by significant modulations of both A-type and B-type lamins (Fig. 5E).

Concave curvatures lead to lower cell proliferation rate and promote significant chromatin condensation.

Finally, the modulation of nuclear shape, YAP localization and the relative abundance of A/C versus B1 lamins by substrate curvature led us to question how matrix corrugations affect cell functions, such as proliferation and DNA synthesis. To answer this question, we

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342 sought to test whether our vertex 2D model can be extended to predict some functional

343 changes in deformed nuclei. We therefore extended our 2D mechanical model of the nucleus

344 to 3D to further explore how stresses arising from curvature could modulate 3D nuclear

345 deformations and volume. We did so by a combination of analytical arguments and 3D finite

346 element simulations from a minimal model of nuclear mechanics (see Supplementary Note for

347 details).   We first considered the simplest case of nuclei compressed into flat shapes (Fig.

348 6A), as observed on crests. As shown in Fig. 6B, the model predicted that increasing the

349 amount of compression from the apical surface of the monolayer also generically decreases

350 nuclear volume. Similarly, nuclei in valleys are predicted to be deformed due to the "offset"

351 forces, thus also decreasing their volume (see Supplementary Theory Note for details). To test

352 both theoretical predictions, we examined whether nuclear volume was affected by curvature,

353 using high magnification confocal Z-stacks of nuclei located on convex, interm. and concave

354 zones of λ20 (Fig. 6C) and λ30 (Fig. 6D) corrugated hydrogels. Cells on convex and concave

355 zones of λ20 and λ30 hydrogels exhibited lower nuclear volumes than those on interm. zones,

356 suggesting that both convex and concave curvatures decreased nuclear volumes. Interestingly,

357 lowest nuclear volumes were found in cells located on concave zones (i.e. valleys) of λ20

358 $(635 \pm 233 \ \mu m^3)$ and λ30 $(776 \pm 116 \ \mu m^3)$, demonstrating that prolate nuclear shapes and

359 high nuclear densities in concave zones resulted in a significant nuclear volume loss.

360   Secondly, we checked whether substrate curvature can affect chromatin organization,

361 using a quantitative procedure based on DAPI staining [39,40]. Indeed, the uptake of DAPI

362 depends on the total amount of DNA, but also on its level of condensation [41]. The average

363 spatial density corresponding to the ratio between the integrated fluorescence intensity and the

364 volume of the nucleus is therefore a reliable indicator of the *in-situ* average chromatin

365 condensation. As shown in Figs 6E-F, marked reorganization of chromatin distribution was

366 associated with nuclear deformations in concave zones of λ20 and λ30 corrugated hydrogels.

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367 Highly condensed chromatin domains showed higher fluorescence intensity with respect to

368 the less condensed ones. The quantification of the chromatin to nuclear ratio indicated that

369 nuclei deformed in concave zones of $\lambda 20$ (Fig. 6G) and $\lambda 30$ (Fig. 6H) showed the highest

370 chromatin compaction values with 0.27 ± 0.03 and 0.37 ± 0.16, respectively. Our results

371 indicated therefore that prolate nuclear shapes in concave curvature zones were associated

372 with a high level of chromatin compaction that may affect DNA synthesis.

373   To test this hypothesis, we used the incorporation of a thymidine analog, 5-ethynyl-2′-

374 deoxyuridine (EdU), as a proliferation marker. A 24-hour incubation period was chosen

375 because it allowed EdU incorporation on corrugated and flat hydrogels, without no saturation

376 of EdU incorporation in MDCK cells. The Edu-positive ratio was calculated as the ratio

377 between Edu and Hoechst-stained cells in a particular field of view (Fig. 6I). We observed

378 lower rates of positive-EdU nuclei on convex curvatures of $\lambda 20$ (0.31 ± 0.08) and $\lambda 30$

379 (0.32 ± 0.07) than on flat hydrogels (0.42 ± 0.05), regardless the positive curvature value (Fig.

380 6H). Furthermore, our findings showed that the rate of positive-EdU nuclei in concave zones

381 was roughly divided by a factor two with 0.18 ± 0.06 on $\lambda 20$ and 0.25 ± 0.11 on $\lambda 30$,

382 demonstrating that the DNA synthesis was significantly decreased in elongated cells

383 accumulated in concave curvature zones.

384   Taken together, these results indicated that concave curvature zones of corrugated

385 matrices were characterized by a high density of prolate-shaped nuclei, which were associated

386 with a high level of chromatin compaction and a lower DNA synthesis rate.

387

388 **Discussion**

389 The role of the substrate curvature has been mainly described so far at the single-cell level,

390 establishing that cells orient themselves along the line of minimal curvature to minimize

391 cytoskeletal deformations. Here, we explore the hypothesis that confluent epithelial

392   monolayers can also sense large-scale curvature via active cell mechanics and nuclear

393   mechano-sensing. Our findings reveal that substrate curvature leads to thicker epithelial zones

394   in concave zones (i.e. valleys) and thinner ones on convex zones (i.e. crests), as well as

395   corresponding cellular densities. This is fully recapitulated in a physical model of apico-

396   lateral active tension, where thickness modulation arises generically as stable states of the

397   monolayer (and consistent with findings of a pre-print released during the preparation of this

398   manuscript [42]), leading us to hypothesize that matrix curvature sensing could arise from

399   resulting density/thickness changes. Furthermore, we develop a minimal theory for how

400   thickness modulations imposed by the substrate can impact nuclear morphometrics, such as

401   nuclear deformation, aspect ratio and positioning relative to the local curvature. These

402   predicted flattened nuclei on convex curvature zones, but also a movement of nuclei towards

403   the concave zones in order to minimize their deformation in response to thickness gradients.

404   We verified these features in experiments, including non-trivial features such as enhanced

405   responses to specific wavelengths. Together, these findings show that physical processes

406   allow epithelial monolayers to respond to curvature changes, leading to the appearance of

407   different types of nuclear deformations and orientations.

408   Given accumulating evidence of nuclear mechano-transduction processes [31,43–46] we

409   then investigated how these physical changes translated into mechano-responses and

410   biochemical changes within the cells as a function of local curvature. We showed that the

411   patterns of cellular densities generated by the matrix curvature are associated with significant

412   changes in the spatial distribution of YAP [3,4], consistent with the idea of curvature-sensing

413   occurring via physically-driven thickness/density-changes. YAP in particular has been shown

414   to be is a key transcription factor that mediates the interplay between cellular mechanics and

415   signaling cascades underlying gene expression, cell proliferation, differentiation fate

416   decisions, and organ development. The spatio-temporal localization of YAP provides

417   therefore critical information about the regulatory state of the cell and in the future, it will be

418   interesting to probe the generality of these findings to other cellular lines and physiological

419   contexts.

420   In addition to YAP response, our results also indicated that lamin A/C versus B1 ratio

421   [47,48], chromatin condensation and cell proliferation rate are all modulated by substrate

422   curvature, demonstrating that substrate curvature sensing modulates multiple

423   mechanotransduction pathways in cell assemblies. The correlation of YAP and lamins with

424   stem cell differentiation and cell division has suggested recently a relationship between the

425   expression of both proteins in deformed nuclei [49], which would be a natural next step of

426   investigation. Our study therefore provides insights into the mechanistic regulation of

427   epithelial monolayers to large-scale substrate curvature, with potential impact for a large

428   number of physiological situations, given the ubiquitous presence of curvature *in vivo.*

429

430   **Materials and Methods**

431

432   **Fabrication of corrugated polyacrylamide hydrogels by UV-photocrosslinking**

433   Instead of the standard radical polymerization using catalysts such as tetramethylenediamine

434   (TEMED) and ammonium persulfate (APS), which lead to slow polymerization times, we

435   used an Irgacure 2959 photoinitiator (2-Hydroxy-4′-(2-hydroxyethoxy)-2-

436   methylpropiophenone) to polymerize hydroxypolyacrylamide (hydroxy-PAAm) hydrogels.

437   Hydroxy-polyacrylamide (hydroxy-PAAm) hydrogels were prepared by mixing acrylamide

438   (AAm), bis-acrylamide (bis-AAm), N-hydroxyethylacrylamide (HEA), 2-Hydroxy-4′-(2-

439   hydroxyethoxy)-2-methylpropiophenone (Irgacure 2959, Sigma #410896) and deionized

440   water. A solution composed of 2836 µL of acrylamide (AAm, Sigma #79-06-1) at 15% w/w

441   in deionized water, 1943 µl of N,N'- methylenebisacrylamide (BisAAm, Sigma #110-26-9) at

442   2% w/w in deionized water, and 1065 µL N-hydroxyethylacrylamide monomers at 65

443   mg/mL in deionized water (HEA, Sigma #924-42-5) were mixed together in a 15 mL

444   Eppendorf tube [50–52] and deionized water was added to reach a final volume of 6 mL. We

445   prepared a stock solution of Irgacure 2959 in sterile deionized water at 5 mg/mL. We

446   introduced 1 mL of the stock solution into the 6 mL of the hydrogel solution [53] to obtain a

447   final concentration of 0.7 mg/mL. After a gentle mixing, the solution was degassed during 30

448   min under a nitrogen flow. Glass coverslips of 22 mm^2 in diameter were cleaned with 0.1 M

449   NaOH solution during 5 min and then rinsed abundantly with deionized water during 20 min

450   under agitation. Cleaned glass coverslips were then treated during one hour with 3-

451   (trimethoxysilyl)propyl acrylate (Sigma #2530-85-0) to promote a strong adhesion between

452   the hydroxy-PAAm hydrogel and the glass coverslips and finally dried under a nitrogen

453   flow. A volume of 40 µL of the degassed mixture was squeezed between an activated glass

454   coverslip and a chromium optical photomask (Toppan photomask, France) and before

455   exposition to UV illumination at 360 nm (Dymax UV light curing lamp). Chromium optical

456   photomasks with alterning transparent stripes of 10 µm wide and black stripes of 10 µm or 20

457   µm wide were used to form corrugated hydrogels with wavelengths of 20 µm (λ20), 30 µm

458   (λ30) and 50 µm (λ50) and respectively. After UV exposition at 360 nm during 10 min at 10

459   mW/cm^2 through the optical photomask, the polymerization was completed and a corrugated

460   hydroxy-PAAm hydrogel was formed. ==The amplitude of 20 µm (λ20) and 30 µm (λ30)==

461   ==corrugated hydrogels was changed by adjusting the volume of the degassed polyacrylamide==

462   ==solution squeezed between the glass coverslip and the chromium optical photomask.== Finally,

463   hydrogels were gently removed from the photomask under water immersion, washed three

464   times in sterile deionized water under gentle agitation and stored in sterile deionized water at

465   4°C. Photocrosslinked hydroxy-PAAm hydrogels were optically transparent and did not

466   exhibit any autofluorescence background at 470±20 nm, 562±88 nm and 591±21 nm.

467

Characterization of corrugated hydrogels by atomic force microscopy

Atomic force microscopy (AFM) measurements were performed on dried and immersed photoreticulated hydrogels to characterize their corrugations [54,55]. AFM of dried hydrogels were performed at room temperature with an ICON instrument from Bruker using the Peak Force mode in order to minimize the applied force. AFM of swollen hydrogels were performed in PBS at 7.2 pH using a NanoWizard TM 3 AFM (JPK Instruments, Berlin, Germany) operated in Quantitative Imaging (QI) mode. Once the coverslip containing the gel was placed on the stage of the microscope, the cantilever was positioned far above the glass surface and allowed to thermally equilibrate during 30 min. Cantilevers were purchased from Bruker (MLCT-BIO-DC-F: k = 0.6 N/m, f = 125 kHz; silicon nitride; front angle $35 \pm 2°$; quadratic pyramid tip shape). Cantilever spring constant and sensitivity were calibrated before each experiment using the JPK software. The force trigger was adjusted to have a high indentation without damaging the gel. Tip velocity was maximized within instrument limits and ramp size was reduced with a short baseline in order to minimize the acquisition time. Data were analyzed with the NanoWizard® Data Processing software version 6.1.96 and the stiffness was calculated according the Hertzian contact model (Young's modulus), fitting the distribution to Gaussians. We obtained a Young's modulus of 250 ± 30 kPa, which is close to the physiological situation and corresponds to the optimal window of elasticity required for vinculin assembly, actin fiber formation and, subsequently, to initiation of replication in kidney epithelial cells [56].

488

Cell culture

Epithelial cells from the Madin-Darby Canine Kidney cell line (MDCK II, Sigma #85011435) were maintained in polystyrene T75 flasks of in a cell culture incubator at 37°C and 5% CO_2.

492   MDCK cells were cultured in proliferation medium composed of Dubelcco's Modified

493   Eagle's medium (DMEM), high glucose (4.5 g/l) with L-glutamine (BE12-604F, Lonza)

494   supplemented with 10% (v/v) Fetal Bovine Serum (FBS, AE Scientific) and 1% of penicillin

495   and streptomycin antibiotics (AE Scientific) [57,58]. MDCK cells were seeded on flat (control),

496   λ20, λ30 and λ50 corrugated hydrogels at a density of 1.10^3 cells/mm^2 for 48 hours. YAP

497   immunostained experiments were also performed at a higher cell density of 5.10^4 cells/mm^2.

498   The cell density was estimated from the ratio of the mean number of nuclei to the surface

499   area, which was corrected by considering the projection effect of λ20, λ30 and λ50 corrugated

500   hydrogels.

501

502   **Proliferation assays**

503   The S-phase synthesis of the cell cycle was labeled in living epithelial tissues grown on

504   corrugated hydrogels using the Click-iT EdU (5-ethynyl-2'-deoxyuridine) (Thermofischer

505   Scientific, A20174) for 30 minutes in proliferation media. Proliferation of MDCK cells on

506   corrugated hydrogels was assessed using the Click-iT EdU Alexa (647 or 555) kit

507   (Thermofischer Scientific, C10338). Briefly, epithelial tissues were incubated with a 10 µM

508   solution of EdU in complete medium for 30 min. Then, tissues were rinsed with PBS, fixed

509   for 10 min with a 4% PFA solution and permeabilized for 20 min with a 0.5% solution of

510   Triton X-100. MDCK tissues cells were blocked with 3% BSA and incubated for 30 min in

511   the dark with a reaction cocktail composed of Click-iT reaction buffer, CuSO4, AlexaFluor

512   azide and reaction buffer additive. MDCK tissues were rinsed with PBS three times and

513   labelled with Hoechst 33342 before mounting in slow fade gold antifade.

514

515   **Immunostaining of epithelial tissues**

21

516 MDCK cells were fixed and permeabilized with 4% paraformaldehyde (Electron Microscopy

517 Sciences), 0.05% Triton X-100 (Sigma) in phosphate buffered saline (PBS 1X, Capricorn

518 scientific) for 12 min at room temperature. Fixed cells were rinsed three times in warm PBS

519 and incubated 30 min with a blocking solution containing 1% BSA (Bovine Serum Albumine,

520 GE Healthcare) and 5% FBS in PBS. MDCK cells were labeled for F-actin with Alexa Fluor

521 488 Phalloidin, 1:200, DNA with DAPI at 1:200 (ThermoFisher Scientific, #D1306) and ß-

522 catenin [59]. Yap was labeled with YAP1 monoclonal antibody produced in mouse (Abnova,

523 clone 2F12), ß-catenin with anti ß-catenin produced in mouse at 1:200 Sigma-Aldrich,

524 #C2206) for 45 min at 37°C. MDCK cells were washed three times in PBS, incubated with an

525 anti-mouse antibody produced in goat and labeled with a goat anti-mouse antibody at 1:200

526 (Molecular Probes, tetramethylrhodamine, Invitrogen, T2762) for 45 min at 37°C. Lamin B1

527 (LMB1) and Lamin A/C (LMAC) proteins were labelled with a polyclonal anti-lamin B1

528 antibody produced in rabbit (1:200, Abcam, ab16048) and a monoclonal anti-lamin A/C

529 antibody produced in mouse (1:200, Santa Cruz Biotechnology, SC-376248), respectively.

530 Epithelial tissues were incubated for 45 min at 37°C with both antibodies. Immunostained

531 cells were mounted on microscope slides with slowfade gold antifade (Thermofisher,

532 Molecular probes) for epifluorescence and confocal imaging.

533

534 **Epifluorescence and confocal imaging**

535 MDCK epithelial tissues were observed in epifluorescence and confocal mode with a Nikon

536 A1R HD25 motorized inverted microscope equipped with x20, x40, x60, x100 Plan Apo (oil,

537 NA=1.4 or silicon, NA=1.4) immersion objectives and lasers that span the violet (405 and 440

538 nanometers), blue (457, 477, and 488 nanometers), green (514 and 543 nanometers), yellow-

539 orange (568 and 594 nanometers), and red (633 and 647 nanometers) spectral regions.

540 Epifluorescence images were recorded with a photometrics Prime 95B camera (Photometrics

541   Tucson, AZ) using NIS Elements Advances Research 4.5 software (Nikon). Confocal images

542   using small Z-depth increments (0.15 μm) were processed using NIS-Elements (Nikon,

543   Advanced Research version 4.5). Photopolymerized (Irgacure 2959) hydroxy-PAAm

544   hydrogels did not exhibit any autofluorescent signal at 470±20 nm (DAPI), 562±88 nm

545   (FITC) and 591±21 nm (TRITC).

546

547   **Morphometric analysis of the nuclei**

548   The morphometric analysis of the nuclei was conducted using the trainable WeKa

549   segmentation plugin for FIJI [60] that first calculates the barycenter, extracts contour, perimeter

550   and projected area of each nuclei. Then the contour of each nuclei of 1 pixel in thickness was

551   defined as a region of interest (ROI). By using the "exclude on edges" function of FIJI, we

552   automatically exclude any ROI that have at least one pixel that overlaps the frame of the

553   image from the working population of nuclei. Then the algorithm generates an ellipse to fit

554   the nuclear contour and determine the orientation angle of each nuclei with respect to the

555   orientation of the corrugations.

556

557   **Statistical analysis**

558   Differences in means between groups were evaluated by two-tailed Student's t-tests

559   performed in Prism 9 (Graphpad Software, Inc.). For multiple comparisons the differences

560   were determined by using an analysis of variance (ANOVA) followed by Tukey post-hoc test.

561   *$p < 0.05$, **$p < 0.01$, ***$p < 0.001$, ****$p < 0.0001$ and n.s. not significant. Unless

562   otherwise stated, all data are presented as mean ± standard deviation (S.D.).

563

564   **Acknowledgments**

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724

725
726

727 **Figure 1 – Wavy epithelial monolayers on corrugated polyacrylamide hydrogels.** (A)

728 <mark>Schematic representation of an epithelial monolayer grown on a corrugated hydrogel of</mark>

729 <mark>amplitude, β, and wavelength, λ.</mark> The corrugated profile is composed of crests (convex

730 curvature), intermediate, or interm., zones (zero curvature) and valleys (concave curvature).

731 The balance between apical tension (Γ_a, in green), lateral tension (Γ_l, in red) and basal tension

732 (Γ_b, in brown) leads to the thickness modulation of the epithelial monolayer, changes of

733 nuclear morphologies and nuclear offsets. (B) Confocal volume rendering of a MDCK

734 epithelial monolayer stained for F-actin (in green) and nuclei (in blue) grown on corrugated

735 substrates of 20 µm (λ20 in grey), 30 µm (λ30 in red) and 50 µm (λ50 in blue) wavelength.

736 Scale bars are 20 µm. (C) Confocal volume rendering of the nuclei in epithelial monolayers

737 grown on corrugated substrates of 20 µm (λ20 in grey), 30 µm (λ30 in red) and 50 µm (λ50

738 in blue) wavelength. The height is color-coded. Scale bars are 20 µm.

739

740 **Figure 2 – Theoretical modelling of epithelial thickness modulations from substrate**

741 **curvature.** (A) Equilibrium configuration of a 2D vertex model representing apical and

742 lateral surfaces of an epithelial monolayer attached to a curved substrate. Increasing substrate

743 wavelength, λ, decreased epithelial thickness modulations. (B) From top to bottom:

744 representative confocal profiles (*xz*) and average thickness modulation for $\lambda 20$ (in grey), $\lambda 30$

745 (in red) and $\lambda 50$ (in blue) corrugated hydrogels (N=8 replicates for each condition) calculated

746 by subtracting apical and basal position of wavy epithelial monolayers at given position *x*. (C)

747 Thickness modulation versus a rescaled wavevector ($\Delta h/\lambda$) for experimental data (mean$\pm$

748 S.D.) $\lambda 20$ (in grey, N=6 replicates), $\lambda 30$ (in red, N=5 replicates) and $\lambda 50$ (in blue, N=6

749 replicates) with the fit from the model (dashed lines correspond to S.D.).

750

751   **Figure 3 – Substrate curvature modulates the spatial distribution of nuclei.** (A) Typical

752   confocal image of an epithelial monolayer grown on a $\lambda 50$ corrugated hydrogels stained for

753   cadherins (in red) and DNA (nuclei, in grey). The zoom panel shows the position of the

754   nuclear center (X in grey) relative to the cellular center (X in blue) as a function of thickness

755   gradients, tending to attract nuclei from intermediate zones to concave ones (valleys). The

756   vector (in red) between the center of mass of cells and nuclei represents the nuclear offset. (B)

757   Sketch of the nuclear offset (arrow in red) determined from the difference between cellular

758   and nuclear axis (lateral view). (C) Experimental and theoretical nuclear offsets versus the

759   coordinate along the x axis normalized by the wavelength for $\lambda 20$ (grey squares), $\lambda 30$ (red

760   circles) and $\lambda 50$ (blue triangles) corrugated substrates. $109 \leq n \leq 162$ per sample with N=3

761   replicates per condition. The dashed line on $\lambda 20$ corresponds to the best fit. (D-G) 2D vertex

762   models (see Supplementary Theory Note) incorporating nuclear positioning (force

763   proportional to thickness gradients, taking cells away from equilibrium center position) for

764   (D) $\lambda=1.5$ and (F) $\lambda=2$. Nuclei are predicted to be positioned towards concave zones, an effect

765   particularly accentuated for wavelengths at integer values of cell width ($\lambda=2$). Typical DIC

766   images of (E) $\lambda 20$ and (G) $\lambda 30$ wavy hydrogels with the automatic detection (in yellow) of

767   the nuclei stained with DAPI.

768

769 **Figure 4 – YAP-curvature sensing is mediated by nuclear density modulation.** (A) From

770 left to right: fluorescent images of confluent epithelial monolayers (1.10^3 cells/mm^2) on flat,

771 $\lambda20$, $\lambda30$ and $\lambda50$ hydrogels and at high cell density (5.10^4 cells/mm^2) on $\lambda20$ and flat

772 hydrogels. Tissues were stained for YAP (in red) and DNA with DAPI (in blue) after 48

773 hours in culture. Scale bars are 30 µm. Nuclear to cytoplasmic YAP ratio on convex (crest)

774 and concave (valley) zones of $\lambda20$ (grey), $\lambda30$ (red) and $\lambda50$ (blue) corrugated hydrogels for

775 (B) normal (1.10^3 cells/mm^2) and (C) high (5.10^4 cells/mm^2) cellular densities. Black bars

776 correspond to flat hydrogels (n=46, N=3 replicates). For normal cell density n=24 ($\lambda20$), n=12

777 ($\lambda30$) and n=12 ($\lambda50$) with $3\leq N\leq8$ replicates. For high cell density n=16 ($\lambda20$), n=22 ($\lambda30$)

778 and n=10 ($\lambda50$) with $3\leq N\leq8$ replicates. (D) Nuclear to cytoplasmic YAP ratio versus local

779 nuclear density on flat (circles), convex (triangles) and concave (squares) zones. All data are

780 shown as mean ± S.D. *p < 0.05, and n.s. not significant.

781

782

783 **Figure 5 – Composition of the nuclear lamina depends on substrate curvature.** (A)

784 Maximum intensity projection of the nuclei of a MDCK epithelial monolayer grown on a $\lambda 20$

785 corrugated hydrogel and stained with DAPI (in blue), lamin A/C (LMAC, in red) and lamin

786 B1 (LMB1, in green). Merge image of LMAC and LMB1. The scale bar is 20 μm. White

787 rectangles correspond to nuclei located on convex, interm. and concave zones. (B) Typical

788 confocal volume views (*xz* view) of a nucleus on convex (crest), interm. and concave (valley)

789 zones. Z_0 (basal), Z_1 (mid-height) and Z_2 (apical) zones are indicated in orange. (C) Typical

790 maximum intensity projection (*xy* view) of a nucleus on convex (crest), an interm. and a

791 concave (valley) zones and stained with DAPI in blue, LMAC in red and LMB1 in green. (D)

792 LMA/LMB intensity ratio versus nuclear height of nuclei on convex (crest), interm. and

793 concave (valley) zones. Black arrows indicate Z_0 (basal), Z_1 (mid-height) and Z_2 (apical)

794 zones. (E) Schematic representation of the balance between LMAC (in red) and LMB1 (in

795 green) for nuclei on convex, interm. and concave (valley) zones. All data are shown as mean

796 $\pm$ S.D. with n=4 nuclei per replicate (N=3).

797

798

799

800

801 **Figure 6 – Concave curvature zones lead to lower cell proliferation rate and promote**

802 **significant chromatin condensation in elongated nuclei.** (A) Schematic of the model used

803 to predict nuclear deformations and volume changes, for an axisymmetric compressed nucleus

804 (in blue) with r and z the nuclear coordinates, θ the angle between the local tangent of the

805 nuclear profile and r-axis, d_b the radius of the contact zone between nucleus and plane

806 and s the arclength of the nuclear profile (see Supplementary Note for details). The insert

807 shows a side (yz) confocal volume view of a nucleus on top of a crest zone, i.e. vertically

808 compressed. (B) Normalized nuclear volume versus normalized cell height shows that nuclear

809 deformations are generically associated to volume reductions. Black squares are Finite

810 Element Model (FEM) simulation, black line the exact solution and red dashed line the

811 analytical approximation. Nuclear volume on interm., concave and convex zones of (C) λ20

812 (in grey, n=24 with N=3 replicates) and (D) λ30 (in red, n=33 with N=3 replicates) corrugated

813 hydrogels. (E-F) Typical successive changes (from bottom to top) of the level of chromatin

814 condensation on concave curvature zones of (E) λ20 and (F) λ30 corrugated hydrogels.

815 Intensities of DNA staining were digitized in 256 bits and color coded for each Z-stack.

816 Highly condensed domains show higher fluorescence intensity with respect to the less

817 condensed ones. (G-H) Chromatin to nuclear volume ratio for cells on interm., concave and

818 convex zones on (G) λ20 (in grey, n=20 with N=3 replicates) and (H) λ30 (in red, n=29 with

819 N=3 replicates) corrugated hydrogels. (I) Typical images of DAPI (in blue) and Edu (in red)

820 staining in an epithelial monolayer grown on λ20 (top row) and λ30 (bottom row) corrugated

821 hydrogels. Scale bars are 20 μm. Rate of Edu-positive cells on (J) convex (n=22 with N=3

822 replicates) and (K) concave (n=22 with N=3 replicates) curvature zones. Dark gray bars are

823 flat hydrogels, light gray bars λ20 and red bars λ30. All data are shown as mean ± S.D. *p <

824 0.05, **p < 0.01, ***p < 0.001, ****p < 0.0001 and n.s. not significant.

825   **Extended Data Figure 1 – Curvature of corrugated hydroxy-PAAm hydrogels.** (A)

826   Representation of a 3D volume view of a corrugated substrate with a tangent plane (*xy*, in

827   blue) and two planes of principal curvatures: *xz* in green ($k_1=1/R$) and *yz* in red ($k_2=0$). The

828   Gaussian curvature is $k_1.k_2=0$. The inset shows a 3D confocal view of a $\lambda 30$ corrugated

829   epithelial monolayer immunostained for actin (in green) and DNA (in blue). The scale bar is

830   30 μm. (B) Top: confocal view (xz) of the wavy profile of a corrugated epithelial monolayer

831   ($\lambda 30$) immunostained for actin (in green) and DNA (in blue). The scale bar is 10 μm. Bottom:

832   Schematic representation of the wavy profile of a corrugated epithelial monolayer composed

833   of concave (in blue), interm. (in green) and convex (in orange) zones. Convex zones

834   correspond to the crest and concave zones to the valley. Interm. zones of zero curvature were

835   located between concave and convex zones and the tangent to the interm. zones was used to

836   determine borders with concave and convex zones. (C) The curvature, C, of convex (crest, in

837   orange) and concave (valley, in blue) zones was determined along the substrate profile (xz, in

838   black) as the reciprocal of the radius (C=1/R) of the osculating circle having its center lying

839   on the normal line. The substrate profile is characterized by a wavelength λ and an amplitude

840   β.

841

Name	Wavelength ± SD (μm)	Convex curvature (μm^{-1})	Concave curvature (μm^{-1})	Amplitude ± SD (μm)
λ20/1.2	20.6 ± 0.9	0.13	- 0.58	1.2 ± 0.1
λ20/2.3	20.1 ± 0.6	0.09	- 0.29	2.3 ± 0.6
λ30/2.3	30.1 ± 0.1	0.27	- 0.27	2.3 ± 0.1
λ50/2.4	49.9 ± 1.0	0.02	- 0.11	2.4 ± 0.8
λ50/3.2	50.2 ± 0.7	0.03	- 0.08	3.2 ± 0.5

842

843 **Extended Data Table 1 – Dimensions of the corrugated hydrogels.** Wavelength (λ),

844 convex curvature (crest), concave curvature (valley), and amplitude (β) of λ20 (n=12), λ30

845 (n=11), λ50 (n=12), λ20/4.5 (n=11), λ50/4.6 (n=11) corrugated hydrogels were determined

846 on swollen hydroxy-PAAm hydrogels by atomic force microscopy (AFM) in liquid mode at

847 37°C. Mean ± S.D.

848

849 **Extended Data Figure 2 – Corrugations do not affect the actin intensity of epithelial**

850 **monolayer.** (A) Maximum intensity projection of an epithelial monolayer grown on a flat,

851 $\lambda20$ (in grey), $\lambda30$ (in red) and $\lambda50$ (in blue) hydrogel and stained for F-actin (in green) and

852 nuclei (in blue). Scale bars are 20 μm for flat and $\lambda20$, 30 μm for $\lambda30$ and 50 μm for $\lambda50$. (B)

853 High magnification confocal images of valley and crest zones for $\lambda20$, $\lambda30$ and $\lambda50$

854 corrugated hydrogels stained for actin (in green) with AlexaFluor 488. (C) 3D volume

855 rendering of a MDCK monolayer grown on a flat hydroxy-PAAm hydrogel coated with FN.

856 Actin is labeled in green with AlexaFluor 488 and DNA in blue with DAPI. The scale bar is

857 50 μm. (D) Total actin intensity in epithelial tissues grown on flat (dark grey), $\lambda20$ (light

858 grey), $\lambda30$ (red) and $\lambda50$ (blue) hydrogels. n=8 (flat in black), n=10 ($\lambda20$ in grey), n=12 ($\lambda30$

859 in red) and n = 5 ($\lambda50$ in blue). n.s. is not significant.

860

861

862 **Extended Data Figure 3 – Mean epithelial cell area and polygon class.** Typical maximum

863 intensity projection image of a MDCK monolayer stained for ß-catenin and grown on (A)

864 λ20, (B) λ30 and (C) λ50 corrugated hydrogels and corresponding skeleton image of (D) λ20,

865 (E) λ30 and (F) λ50. Scale bars correspond to 100 µm (λ20 and λ30) and to 50 µm (λ50). (G)

866 Mean cell area and (H) distribution of polygon classes of epithelial tissues grown on flat

867 (black), λ20 (grey), λ30 (red) and λ50 (blue) hydrogels. n= 1100 cells (flat in black), 1500

868 cells (λ20 in grey), 1700 cells (λ30 in red) and 1050 cells (λ50 in blue), obtained from n=3,

869 n=6, n=5 and n=3 replicates respectively. (I) Mean cell area versus polygon class of epithelial

870 monolayers grown on flat (black), λ20 (grey) and λ30 (red) hydrogels. n.s. is not significant.

871

872

873 **Extended Data Figure 4 – Schematic of the model and sensitivity analysis.** Equilibrium

874 configuration of a 2D vertex model representing apical and lateral surfaces of an epithelial

875 monolayer attached to a curved substrate. (A) Decreasing substrate amplitude, β, or (B)

876 decreasing apical tensions, Γ_a, decreased thickness modulations. (C) Simulation of the vertex

877 model on curved substrate with increasing density (4, 6, and 8 cells per wavelength, from top

878 to bottom), showing that increasing density decreases thickness modulations. (D) Simulation

879 of human keratocyte on curved substrate (see Supplementary Note for details), modelled with

880 $\frac{\Gamma_a}{\Gamma_l} \approx 5$, showing density modulations even for large wavelengths (top), which are amplified

881 further to the point of crest/top dewetting when doubling the substrate amplitude (middle), or

882 when doubling apical tensions $\frac{\Gamma_a}{\Gamma_l} \approx 10$ (bottom). (E) Comparison for the thickness

883 modulation Ω between analytical theory (thin lines) and vertex simulations (dots), for $\frac{\Gamma_a}{\Gamma_l} = 5$

884 (purple), $\frac{\Gamma_a}{\Gamma_l} = 2$ (green) and $\frac{\Gamma_a}{\Gamma_l} = 1$ (purple), showing good agreement for large wavelengths

885 λ (normalized by average cell thickness Δh, in all panels C-E, we have taken for simplicity

886 average cell thickness $\Delta h = 1$ and average cell side length $2d = 1$), with corrections for

887 small wavelengths. (F) Schematics of contact mechanics model of a nucleus subjected to

888 apical compression, before (left) or after (right) lateral contact. (G-I) Sensitivity analysis of

889 different model parameters/observations. Both full solutions (solid line) and analytic

890 approximations (dashed line) are given (see Supplementary Note for details). We examine the

891 influence of normalized cell height $\bar{\square}$ on aspect ratio S_n (G) for the nucleus (volume ratio of

892 cell to nucleus $\bar{v} = \frac{V_c}{\pi r_0^2} = 2$ and before lateral contact). We also examine the influence of

893 local thickness gradient on aspect ratio (H) and nuclear offset (I), with $\bar{\square} = 0.5$, $\bar{d} = 1.57$,

894 and tension ratio $\frac{\Gamma_a}{\Gamma_{n0}} = 5$. We find on both metrics a sharp transition above a critical value of

895 thickness gradients, which occurs when the nucleus reaches lateral contact (and thus cannot

896   increase its offset but start adopting distorted asymmetric shape because of the asymmetric

897   contact).

898

899

Extended Data Figure 5 – Modulation of wavelength and amplitude of corrugated substrates. (A) Representative 3D confocal image of the spatial localization of the nuclei within an epithelial monolayer grown on a corrugated hydroxy-PAAm hydrogels of 100 μm in wavelength (λ100). The nuclear height is color-coded. (B) From top to bottom**:** typical confocal profile of an epithelial monolayer grown for 48 hours on a corrugated λ100 substrates and stained for actin with phalloidin. Average height profiles along the x position obtained from Z-stack confocal imaging (n=5). Vertex model showing the thickness modulation of the epithelial monolayer with thicker cells on convex zones and thinner ones on concave zones. (C) Modulation of the corrugation amplitude for λ20 (in grey) and λ50 (in blue) substrates to match the amplitude of λ30 (in red) substrates of ~2.3 μm. The amplitude of λ20 substrates was increased to ~2.3 μm (λ20/2.3, n=16 with N=3 replicates) and the amplitude of λ50 substrates was decreased to ~2.4 μm (λ20/2.4, n=18 with N=3 replicates) (B) Thickness modulation of the epithelial monolayer for λ20 (in grey) and λ50 (in blue) substrates of different amplitudes. **p < 0.01, ***p < 0.001, ****p < 0.0001 and n.s. not significant.

918   **Extended Data Figure 6 – Substrate curvature modulates the nuclear area.** (A) Local

919   nuclear projected area normalized by the mean nuclear area of each sample on convex (crest)

920   and concave (valley) zones of $\lambda 20$ (grey), $\lambda 30$ (red) and $\lambda 50$ (blue) substrates. (B) Nuclear

921   density normalized by the mean nuclear density of each sample on convex and concave zones

922   of $\lambda 20$ (grey), $\lambda 30$ (red) and $\lambda 50$ (blue) substrates. $109 \leq n \leq 162$ nuclei per sample with N=3

923   replicates for all conditions. $**p < 0.01$ and $***p < 0.001$. (C-E) Local nuclear projected area

924   (normalized by overall mean nuclear area) versus normalized position (*x* axis normalized by

925   the wavelength, so that 0 and 1 correspond to top/concave regions and 0.5 to bottom/convex

926   regions) for epithelial monolayers grown on (C) $\lambda 20$, (D) $\lambda 30$ and (E) $\lambda 50$ hydrogels, for the

927   best fit parameter of $\gamma = 0.25 \,\mu m^{-1}$ (see Supplementaty Note for details). Grey squares

928   ($\lambda 20$), red circles ($\lambda 30$) and blue triangles ($\lambda 50$) are experimental data (mean ± S.D.) and

929   plain lines the best fit model. $109 \leq n \leq 162$ nuclei per sample with 3 replicates for all

930   conditions.

931

932

933 **Extended Data Figure 7 – Nuclear orientation and YAP nuclear export are modulated**

934 **by substrate concave curvatures.** Mean orientation of the nuclei on (A) convex (crest) and

935 (B) concave (valley) curvature zones of $\lambda20$ (in grey), $\lambda30$ (in red) and $\lambda50$ (in blue) wavy

936 hydrogels. All data are shown as mean ± SD. The number of nuclei is indicated at the bottom

937 of each bar: $170 \leq n \leq 368$ for concave curvature and $109 \leq n \leq 405$ for convex curvature. Nuclear

938 to cytoplasmic YAP ratio of nuclei on interm., concave, convex zones of (C) $\lambda20$ in grey, (B)

939 $\lambda30$ in red and (C) $\lambda50$ in blue corrugated hydrogels. Black bars correspond to flat hydrogels.

940 For $\lambda20$ n=30 (interm.), n=16 (concave), 24 (convex) and n=50 (flat) obtained from 5 to 7

941 replicates, for $\lambda30$ n=18 (interm.), n=18 (concave), n=9 (convex) and n=50 (flat) obtained

942 from 5 to 9 replicates and for $\lambda50$ n=6 (concave), n=7 (convex) and n=50 (flat) obtained from

943 3 to 5 replicates. All data are shown as mean ± SD. $*p < 0.1$, $**p < 0.01$, $***p < 0.001$, $****p$

944 < 0.0001 and n.s. not significant.
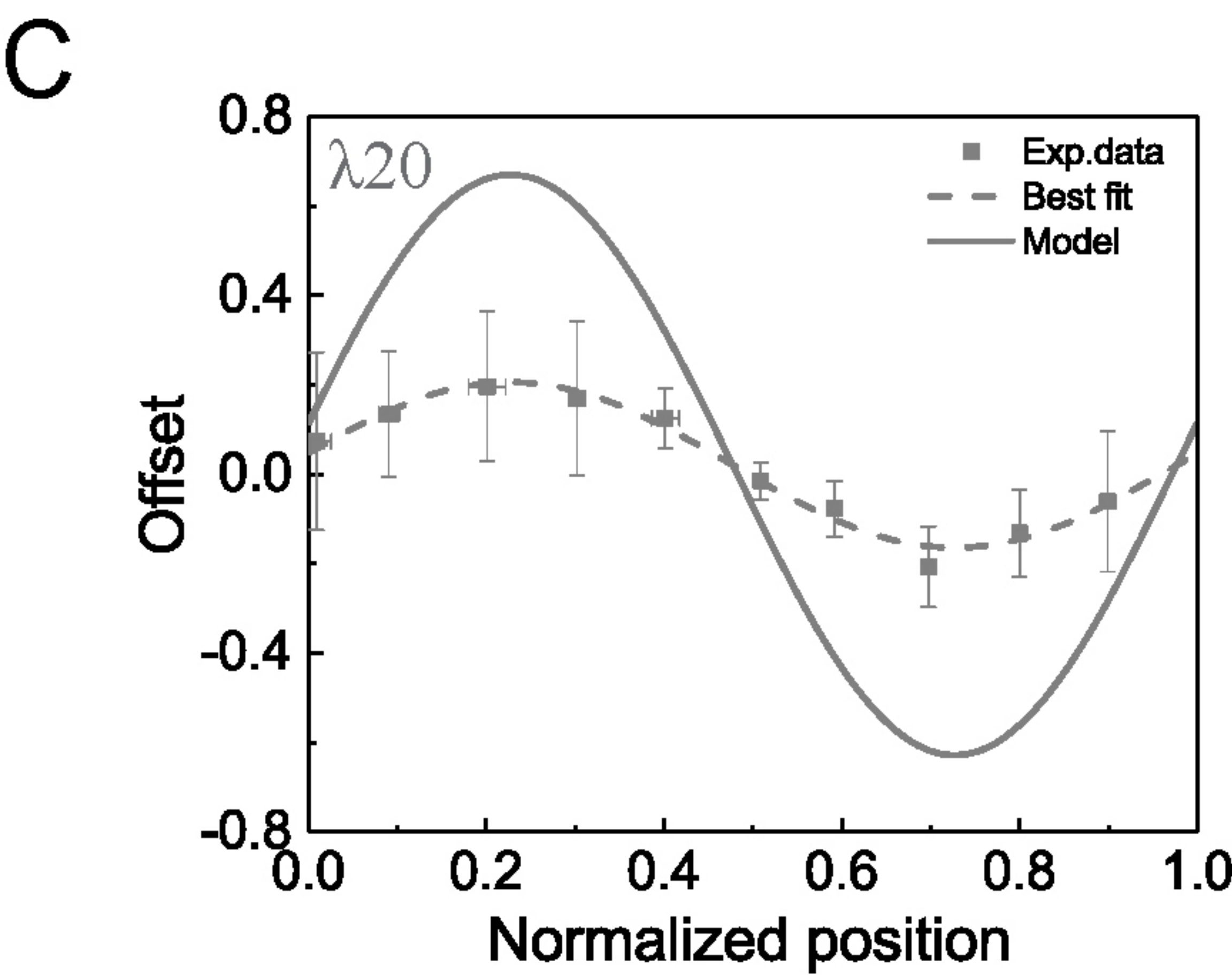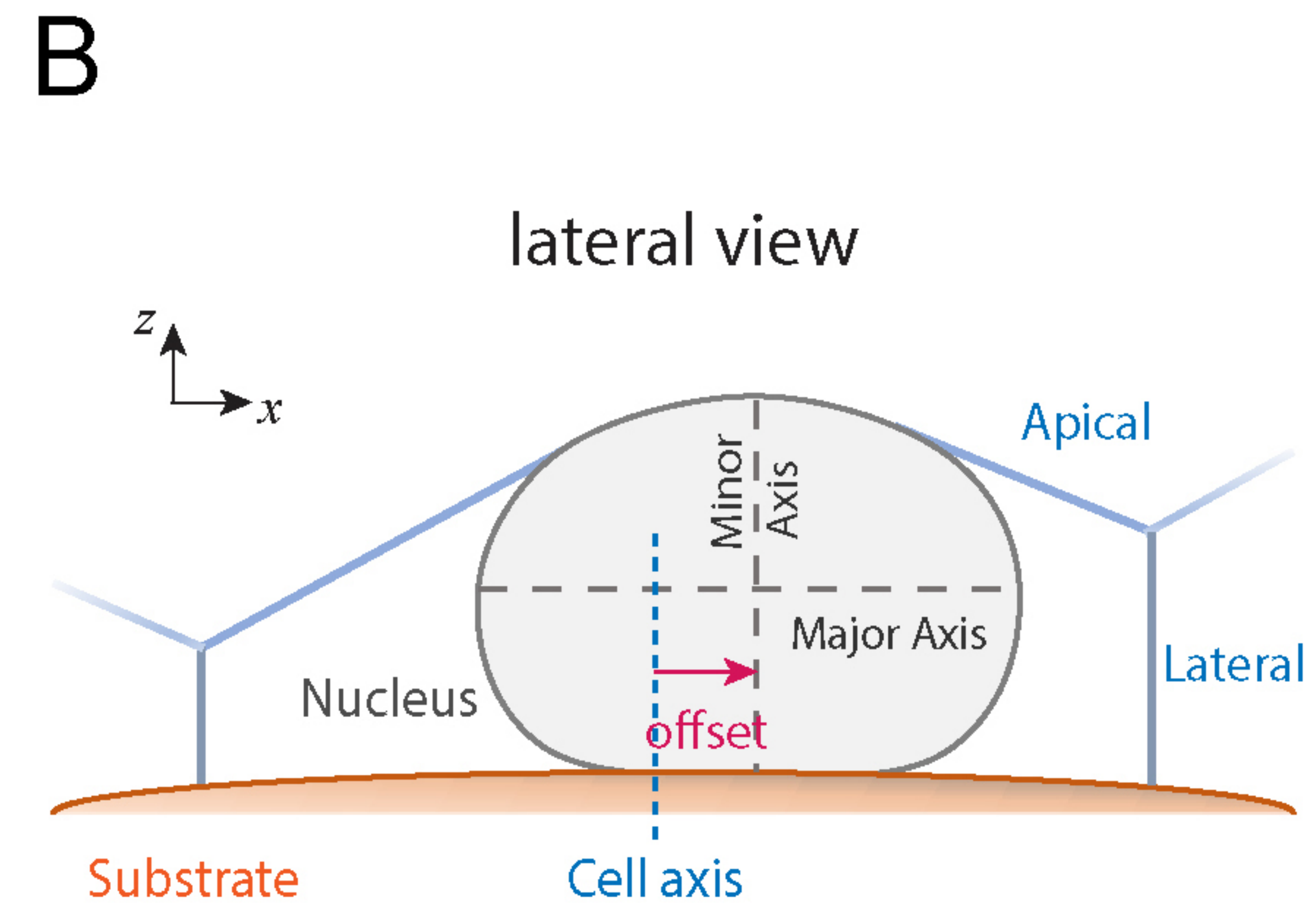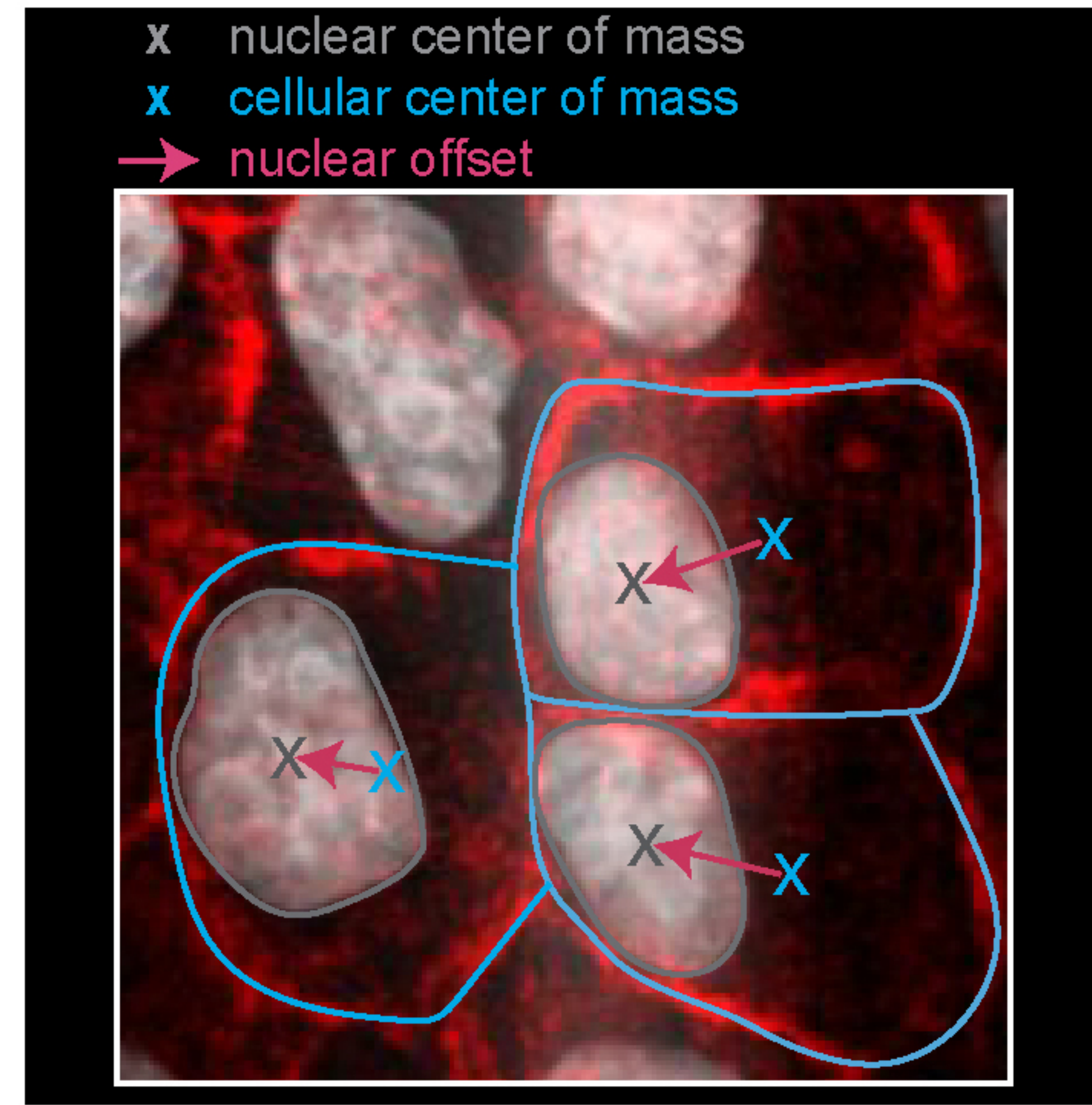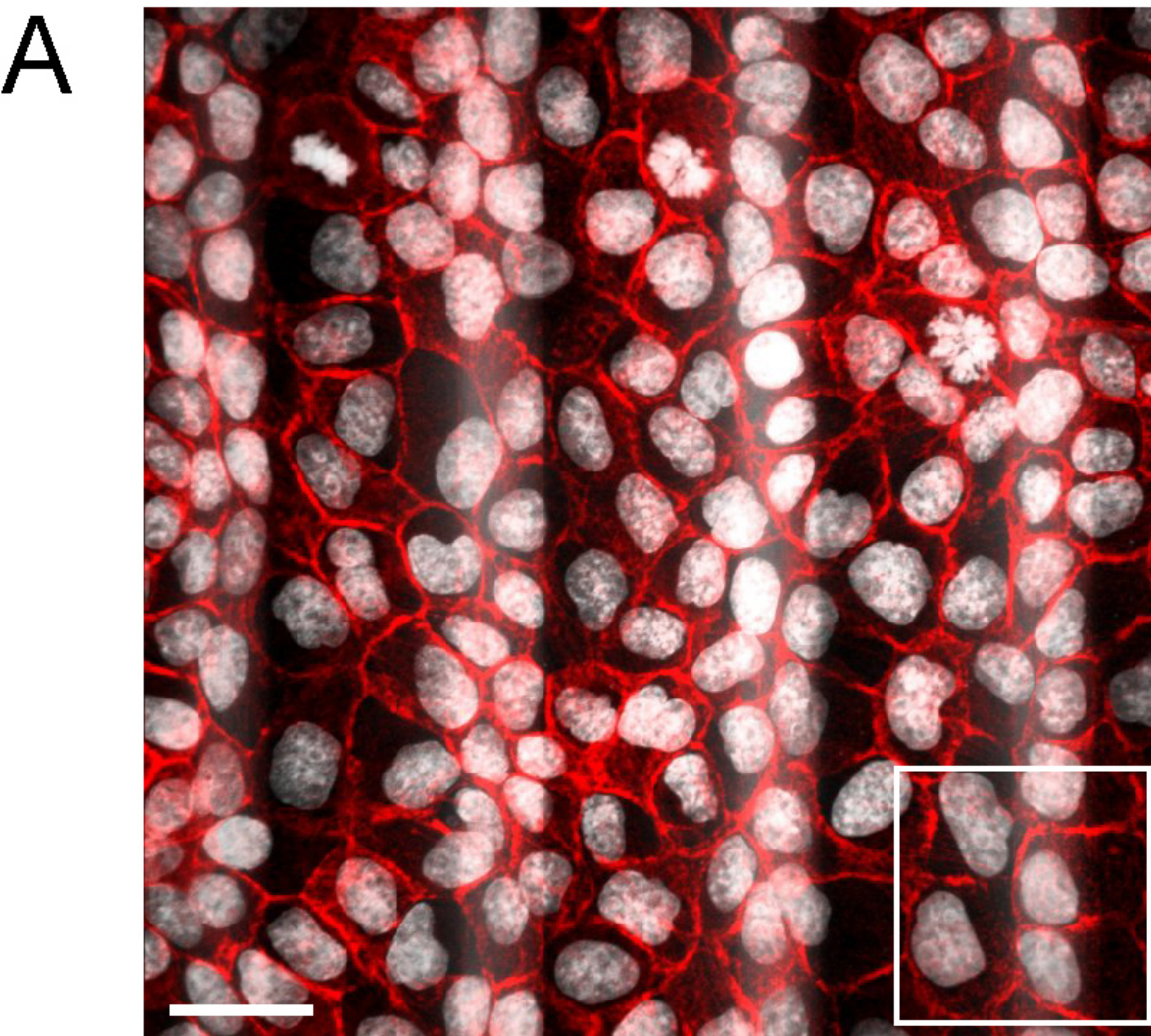
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950

951   **Extended data Figure 8. Finite element simulations of nuclei on curved substrates.** (A)

952   Snapshots of nuclear 3D morphologies in different regions, with the normalized average

953   monolayer thickness $\Delta\bar{h}=0.3$. Dependence of (B) the normalized nuclear volume $\bar{V}_n$ and (C)

954   the nuclear aspect ratio in x-y plane on $\Delta\bar{h}$. A nucleus in the concave region is either in

955   contact with the neighboring nucleus (or cell membrane) on the right side, or confined on both

956   sides, with cell side length (along x-axis) proportional to monolayer thickness.
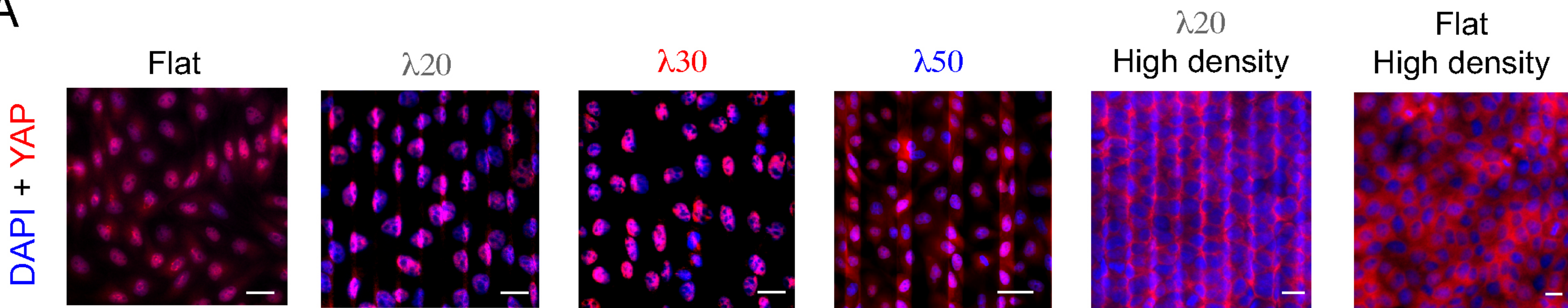
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958

959   **Extended data Figure 9. Large-scale curvature sensing by epithelial monolayers depends**

960   **on active cell mechanics and nuclear mechanoadaptation.** Schematic representation of the

961   epithelial thickness modulation and the three main nuclear morphologies observed on crest

962   (convex), interm. zones and valleys (concave). Composition of the nuclear lamina depends on

963   substrate   curvature,   whereas   YAP-curvature   sensing   is   mediated   by   nuclear

964   density modulation. Concave curvature zones lead to lower cell proliferation rate and promote

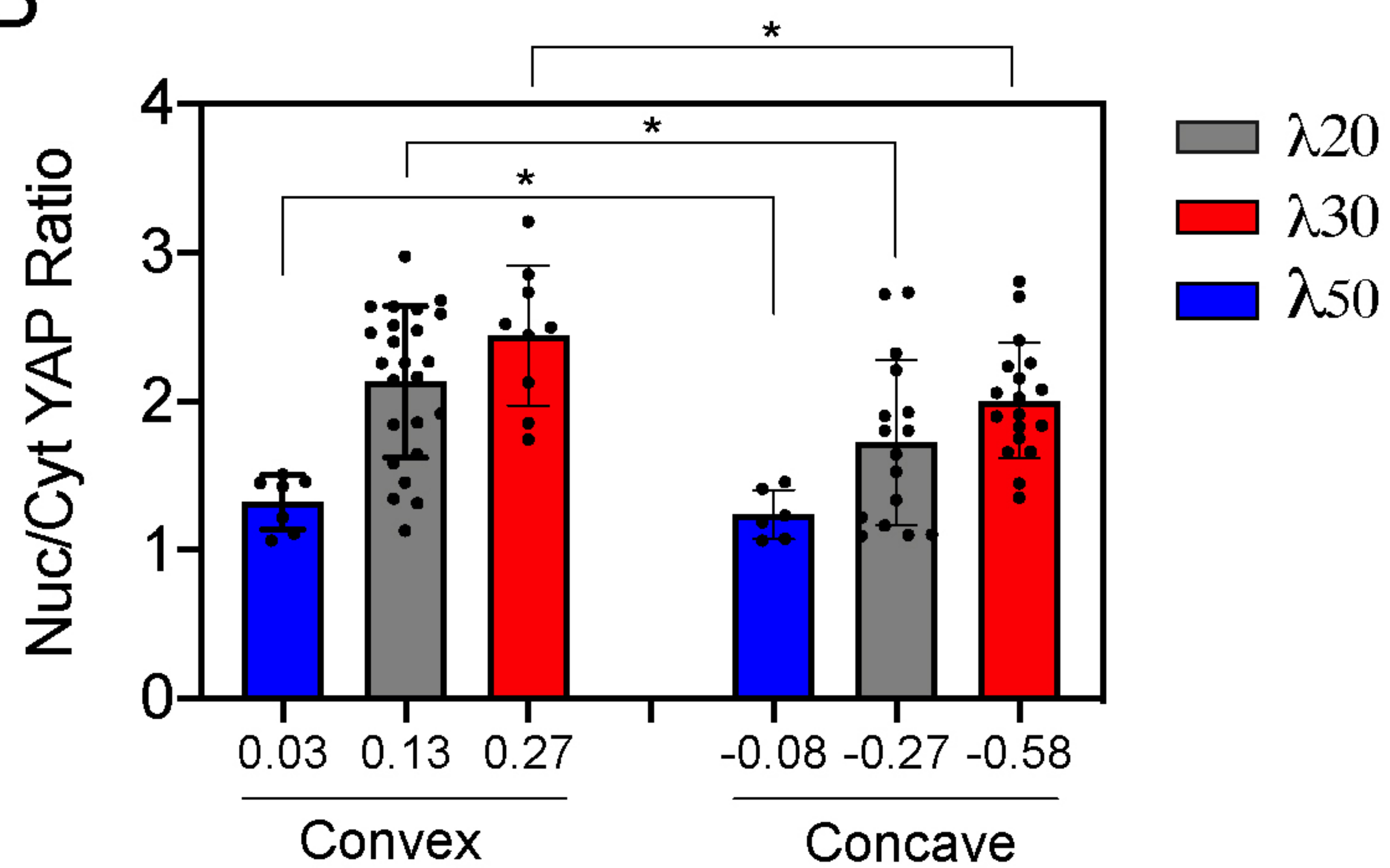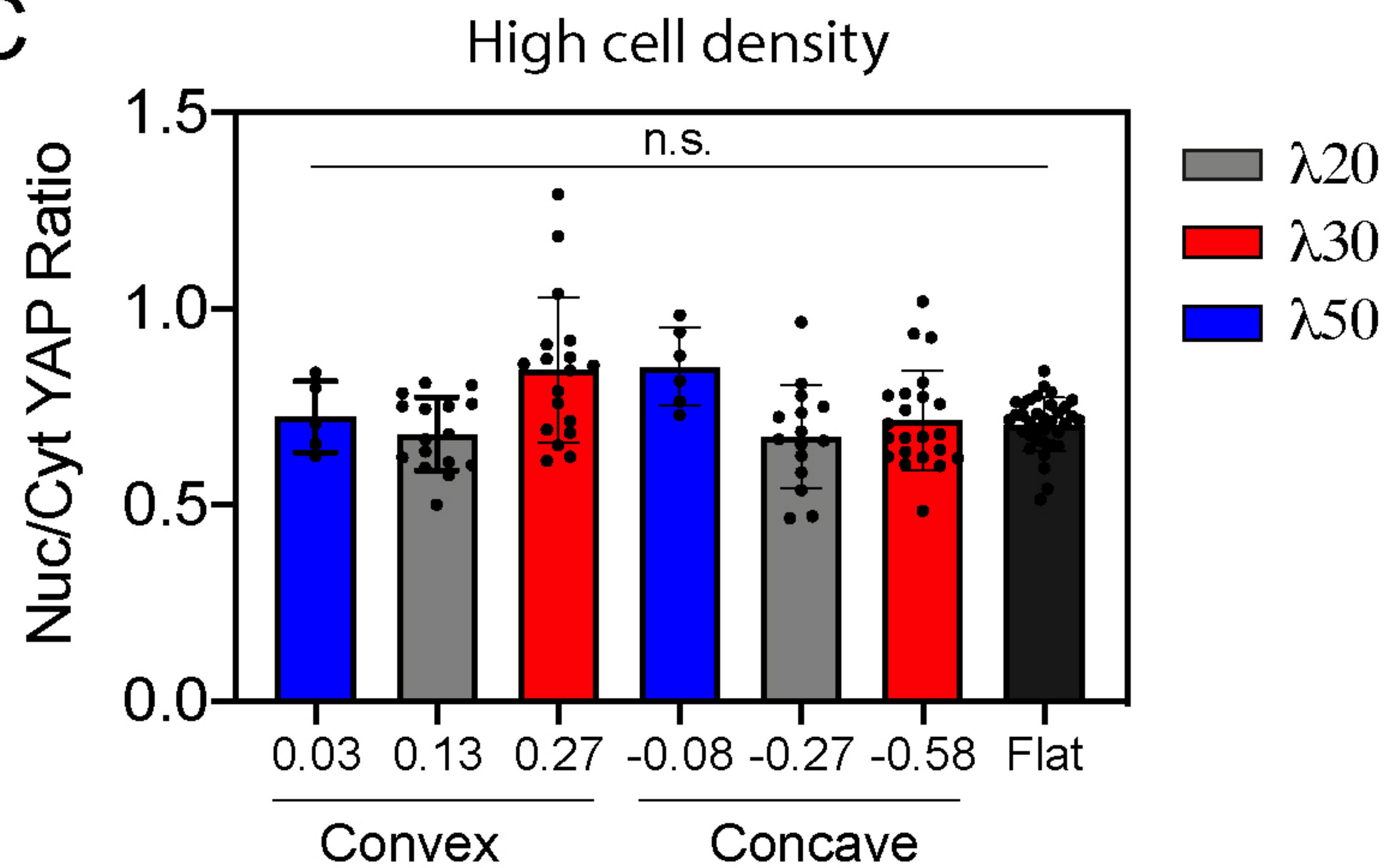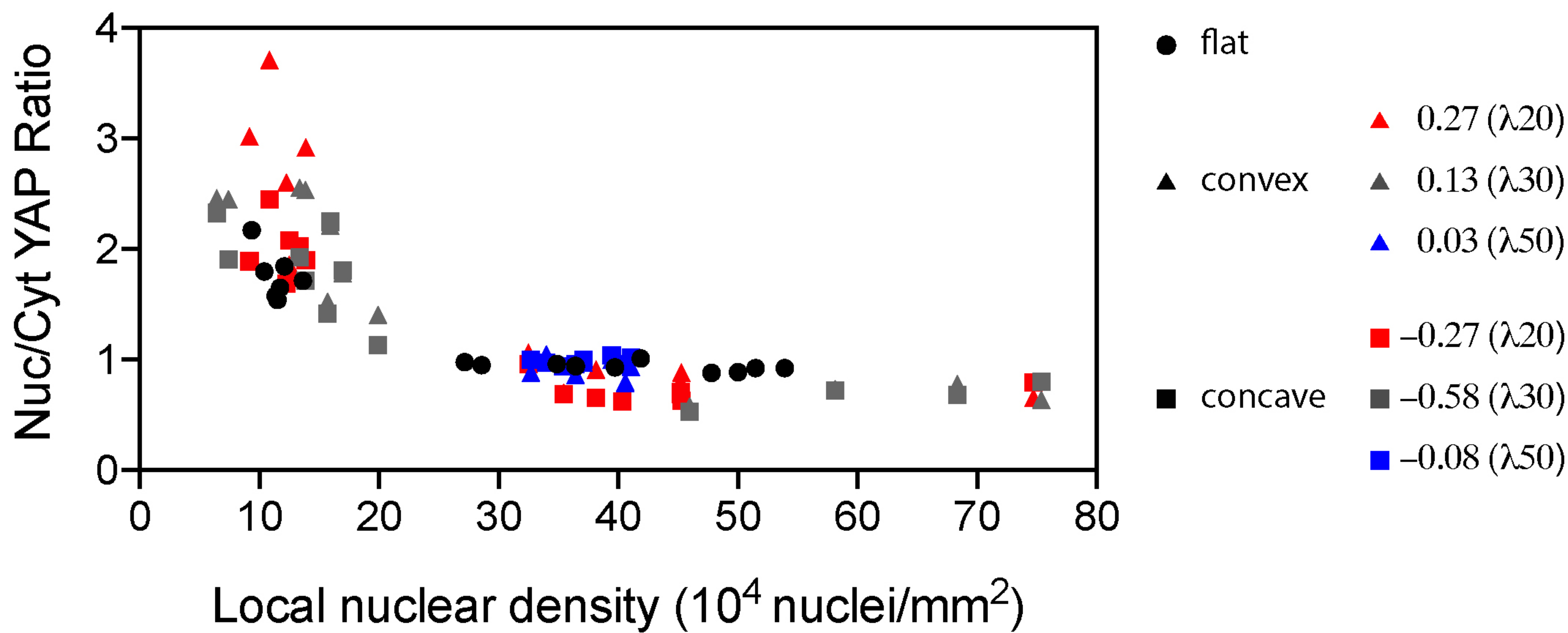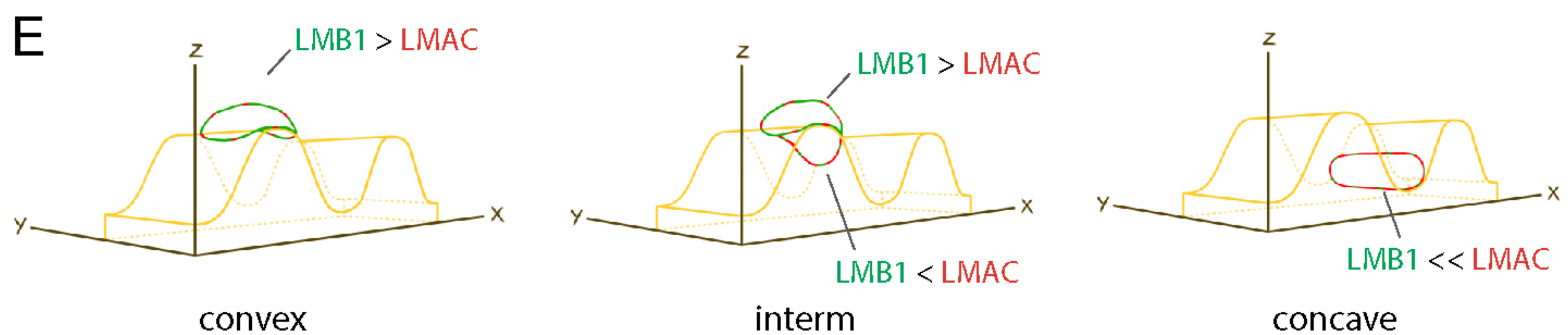965   significant chromatin condensation in elongated nuclei.

A

λ

CREST (convex)

β

INTERM (zero curvature)

VALLEY (concave)

Epithelial thickness

Convex

Concave

⟷ Apical surface Ta
⟷ Lateral surface Tℓ
⟷ Basal surface Tb

● Nuclear morphology and offset

B

λ20

λ30

λ50

C

λ20

8 μm
6 μm
4 μm
2 μm
0 μm

λ30

12 μm
10 μm
8 μm
6 μm
4 μm
2 μm
0 μm

λ50

12 μm
10 μm
8 μm
6 μm
4 μm
2 μm
0 μm

A

B lateral view

C

D Wavelength=1.5

E

F Wavelength=2

G

A

Flat   λ20   λ30   λ50   λ20 High density   Flat High density

DAPI + YAP

B

Nuc/Cyt YAP Ratio

Legend: λ20 (gray), λ30 (red), λ50 (blue)

	Convex			Concave		
	0.03	0.13	0.27	-0.08	-0.27	-0.58

C High cell density

Nuc/Cyt YAP Ratio

n.s.

Legend: λ20 (gray), λ30 (red), λ50 (blue)

	Convex			Concave			
	0.03	0.13	0.27	-0.08	-0.27	-0.58	Flat

D

Nuc/Cyt YAP Ratio

- ● flat
- ▲ convex
- ■ concave

- ▲ 0.27 (λ20)
- ▲ 0.13 (λ30)
- ▲ 0.03 (λ50)

- ■ −0.27 (λ20)
- ■ −0.58 (λ30)
- ■ −0.08 (λ50)

Local nuclear density (10^4 nuclei/mm^2)

A DAPI | LMAC | LMB1 | LMAC + LMB1

(LMAC panel labels: convex, interm, concave)

B convex | interm | concave

(z_2, z_1, z_0 dashed lines indicated in each 3D rendering; axes z, x)

C
convex: DAPI | LMAC | LMB1 (axes y, x)
interm: DAPI | LMAC | LMB1 (axes y, x)
concave: DAPI | LMAC | LMB1 (axes y, x)

D

convex — LMAC/LMB1 ratio (a.u.) vs Height (µm); markers z_0, z_1, z_2; legend LMAC/LMB1

interm — LMAC/LMB1 ratio (a.u.) vs Height (µm); markers z_0, z_1, z_2; legend LMAC/LMB1

concave — LMAC/LMB1 ratio (a.u.) vs Height (µm); markers z_0, z_1, z_2; legend LMAC/LMB1

E

convex: LMB1 > LMAC

interm: LMB1 > LMAC ; LMB1 < LMAC

concave: LMB1 << LMAC

A

B

FEM simulation
Exact solution
Analytic approximation

Normalized nuclear volume

Normalized cell height

height

Nucleus

C λ20

Nuclear Volume (μm³)

Interm   Concave   Convex

D λ30

Nuclear Volume (μm³)

Interm   Concave   Convex

E Concave curvature - λ20

F Concave curvature - λ30

G λ20

Chromatin / Nuclear volume ratio

n.s

Interm   Concave   Convex

H λ30

Chromatin / Nuclear volume ratio

n.s

Interm   Concave   Convex

I

DAPI   EdU   Merged

λ20

λ30

J

DNA synthesis (% of labeled nuclei)

n.s

Flat   0.13   0.27

Convex curvature (μm⁻¹)

K

DNA synthesis (% of labeled nuclei)

Flat   - 0.27   - 0.58

Concave curvature (μm⁻¹)