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Supplemental Information

Cholesterol Depletion by MetaCD Enhances Cell Membrane Tension and Its Variations-Reducing Integrity

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Table S1

Calculation of spatio-temporal parameters of membrane fluctuations					
Parameters	Description	Types of averaging	Used in figures		
Mean	Relative height of the	Averaged over 144 pixels	Figs. S1c, S3c,		
relative	membrane in a single pixel,	to give one number per	S5d		
height	calculated over 2048	FBR			
	frames	Averaged over all FBRs	Figs. S1b, S3b,		
		to give one number per	S5c		
ap.	27.6	cell	F: G1 G2		
SD_{time}	RMS value of temporal	Averaged over 144 pixels	Figs. S1c, S3c,		
	fluctuations in a single	to give one number per	S5d		
	pixel, calculated over 2048	FBR	E. 011 21 05		
	frames	Averaged over all FBRs	Figs. S1b, 3d, S5c		
		to give one number per			
SD(SD:))	Intra-FBR variation of	cell Computed from 144	Figs C1a C6a		
$SD(SD_{time}))$	single-pixel RMS of	pixels to give one number	Figs. S1c, S6a		
	temporal fluctuations,	per FBR			
	calculated over 2048	Averaged over all FBRs	Figs. 4a, S6a		
	frames	to give one number per	11gs. 4a, 50a		
	Turnes	cell			
SD SD _{time}	Intracellular heterogeneity	SD calculated after	Fig. 4d		
22 22 time	$\inf SD_{time}$	clubbing SD _{time} values of	118		
	time	all FBRs in a cell to give			
		one number per cell			
Dissimilar	Intracellular variation of	Computed from all FBRs	Figs. 4b, S6b		
pairs	single-pixel RMS of	to give one number per			
_	temporal fluctuations	cell			
SD_{space}	RMS value of spatial	Averaged in 20 frames to	Figs. 2c, S3d, S5d		
	height variation in a	give one number per FBR			
	$2.16x2.16 \mu m^2$ region	Averaged over all FBRs	Figs. 2d, S3d, S5c		
		to give one number per			
		cell			
$SD SD_{space}$	Intracellular heterogeneity	SD calculated after	Fig. 4d		
	$\inf \mathrm{SD}_{\mathrm{space}}$	clubbing SD _{space} values of			
		all FBRs in a cell to give			
2		one number per cell	E' 0 051		
λ	Correlation length obtained	Averaged over 200	Figs. 2c, S5d		
	in a 6.3x1.8 µm ² region	frames to give one			
	from fitting spatial ACFs to 3-term exponential	number per region Averaged over all regions	Figs. 2d, S5c		
	function	to give one number per	171gs. 2u, 330		
		cell			
τ	Correlation time obtained	Averaged in a 0.36x0.36	Fig. 3a		
	in a pixel by fitting	μm ² region to give one			
		number per region			

	temporal ACFs to 3-term	Averaged in a 2.16x2.16	Fig. 3a inset				
	exponential function	μm ² region to give one					
		number per region					
PSD	Power spectrum of	Averaged over 144 pixels	Figs. 2a, S3a, S5b				
	temporal fluctuations	to get one PSD per FBR					
Parameters extracted from PSD							
Parameters	Description	Types of averaging	Used in figures				
	_		-				
$\overline{\sigma(f_1,f_2)}$	Amplitude of temporal	Computed from a PSD to	Figs. S1c, S3c,				
01/2/	fluctuations in a frequency	get one number per FBR	S5d				
	regime	Averaged over all FBRs	Figs. 2a, S3b, S5c				
		to get one number per cell					
Exponent	Frequency dependent	Computed from a PSD to	Figs. S1c, S3c,				
	power law of the PSD	get one number per FBR	S5d				
		Averaged over all FBRs	Figs. 2a, S3b, S5c				
		to get one number per cell					
Parameters extracted by fitting PSD							
Parameters	Description	Types of averaging	Used in figures				
			-				
A	Active temperature	Computed from a PSD to	Figs. S1e, S3f,				
	_	get one number per FBR	S6d				
		Averaged over all FBRs	Figs. 2e, S3e, S6c				
		to get one number per cell					
$\eta_{ m eff}$	Effective cytoplasmic	Computed from a PSD to	Figs. S1e, S3f,				
	viscosity	get one number per FBR	S6d				
		Averaged over all FBRs	Figs. 2e, S3f, S6c				
		to get one number per cell					
γ	Confinement of the	Computed from a PSD to	Figs. S1e, S3f,				
	membrane	get one number per FBR	S6d				
		Averaged over all FBRs	Figs. 2e, S3e, S6c				
		to get one number per cell					
σ Membrane tension		Computed from a PSD to	Figs. S1e, 3d, S6d				
		get one number per FBR					
		Averaged over all FBRs	Figs. 2e, 3d, S6c				
1		to get one number per cell	I				

Table S1: Calculation of spatio-temporal parameters of fluctuations and mechanical parameters.

Table S2

Parameters	Notation	Values in reports	References	Values calculated in this study
Active	A	3	(1)	3.6±1.1
temperature				
Effective	$\eta_{ m eff}$	2-4*10 ⁴ Pa.s	(2–4)	4162±2010 Pa.s
cytoplasmic				
viscosity				
Bending	κ	$10-50 k_BT$	(5–7)	Kept constant at
rigidity				$15 k_B T (8)$
Confinement	γ	2.3*10 ⁸ N/m3,	(7, 9)	$8.9\pm4.2 \times 10^8 \text{N/m}^3$
		1.7*105 J/m4		
Membrane	σ	3.31*10 ⁻⁵ N/m,	(6, 10–12)	565±330 pN/μm
tension		10-100 pN/μm,		
		22-276 pN/μm		

Table S2: Description of the model which is used to extract mechanical parameters.

Figure S1

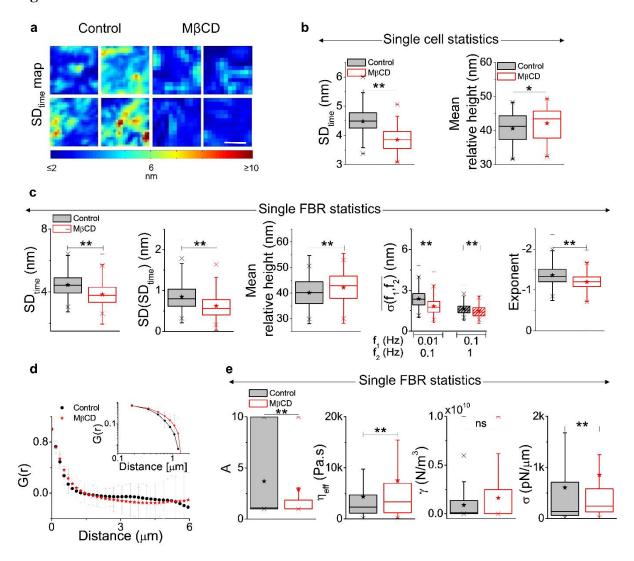


Figure S1: Effect of MβCD on detailed parameters of membrane fluctuations. (a) SD_{time} maps of representative FBRs in control and MβCD treated cells (scale bar: 1 μm). (b) Box plots of SD_{time} and mean relative height in the two conditions. N = 70 cells each. (c) Single FBR statistics of parameters of temporal fluctuations in the two conditions. $n_{control} = 1683$ FBRs, $n_{MβCD} = 1471$ FBRs, N = 70 cells each. (d) Averaged spatial ACFs (and their log-log plots, top inset) for control and cholesterol depleted cells. (e) Single FBR statistics of mechanical parameters in control vs. cholesterol depleted cells. $n_{control} = 1500$ FBRs, $n_{MβCD} = 1317$ FBRs, N = 70 cells each. * p value < 0.05, ** p value < 0.001, ns p value > 0.05, Mann-Whitney U test. See **Table S4** for statistics.

Figure S2

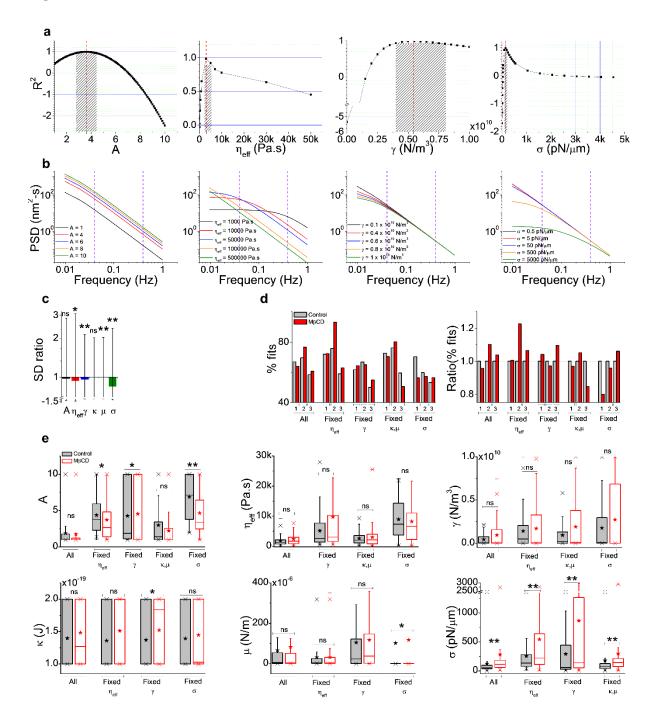


Figure S2: Alterations in membrane tension primarily rule the change in fluctuations. (a) Plots of calculated R² with different values of A, $\eta_{\rm eff}$, γ , σ to check the sensitivity of the extracted mechanical parameters. (b) Simulations of PSD $PSD(f) = \frac{4\eta_{eff}Ak_BT}{\pi} \int_{q_{min}}^{q_{max}} \frac{dq}{(4\eta_{eff}(2\pi f))^2 + \left[\kappa q^3 + \sigma q + \frac{\gamma}{q}\right]^2}$ with changing values of A, $\eta_{\rm eff}$, γ , σ from low to high

numbers to check the robustness of the parameters. The basic power spectrum was constructed with these parameters: A = 1.8083, $\eta_{eff} = 3838$ Pa.s, $\gamma = 0.16$ x 1010 N/m3, $\kappa = 0.6$ x 10-19 J, $\sigma = 74.8$ pN/ μ m. The dashed vertical lines show the regime where the exponent is calculated. (c) Average values of SD ratio simulated from using one-ON approach in M β CD treated (left,

n = 616 simulations) cells of all six parameters. Error bars represent the standard deviation values in each. PSDs $PSD(f) = \frac{4\eta_{eff}Ak_BT}{\pi} \int_{q_{min}}^{q_{max}} \frac{dq}{(4\eta_{eff}(2\pi f))^2 + \left[\kappa q^3 + \frac{9k_BT}{16\pi\kappa}\mu q + \sigma q + \frac{\gamma}{q}\right]^2}$ are simulated

from different sets of observed fitting parameters. A whole set of fitting parameters corresponding to a control set are chosen and then only one parameter is changed at a time to that of a M β CD treated set. This is done for each of the parameters – A to σ . The ratio of the SD of the original control set and the SD calculated from the simulated PSD is calculated and the log of these values are plotted to understand the different contributions. Statistics is by students' t-test. (d) Left: plot of % fits in all categories mentioned below in three sets of MβCD treated conditions. experiments control and % fits =No. of FBRs that fit to the model in the mentioned category. Right: plot of ratio (% fits) between Total no.of FBRs analyzed control and MBCD treated cells in the same criteria across three sets of experiments. Ratio (% fits) = $\frac{\% fits in M\beta CD sets}{\% fits in control treated sets}$. n = 10 cells each. (e) Mechanical parameters extracted from fitting the PSDs to the theoretical model with minor modifications of control and cholesterol depleted cells. All: none of the parameters are fixed, Fixed η_{eff} : $\eta_{eff} = 3421.27$ Pa.s, Fixed γ : $\gamma = 0.08 \times 10^{10} \text{ N/m}^3$, Fixed κ , μ : $\kappa = 1.38 \times 10^{-19} \text{ J & } \mu = 90 \times 10^{-6} \text{ N/m}$, Fixed σ : σ = 368 pN/ μ m. N = 10 cells, $n_{control}$ = 188 total FBRs, n_{MBCD} = 186 total FBRs. * p value < 0.05, ** p value < 0.001, Mann-Whitney U test. See **Table S4** for statistics.

Figure S3

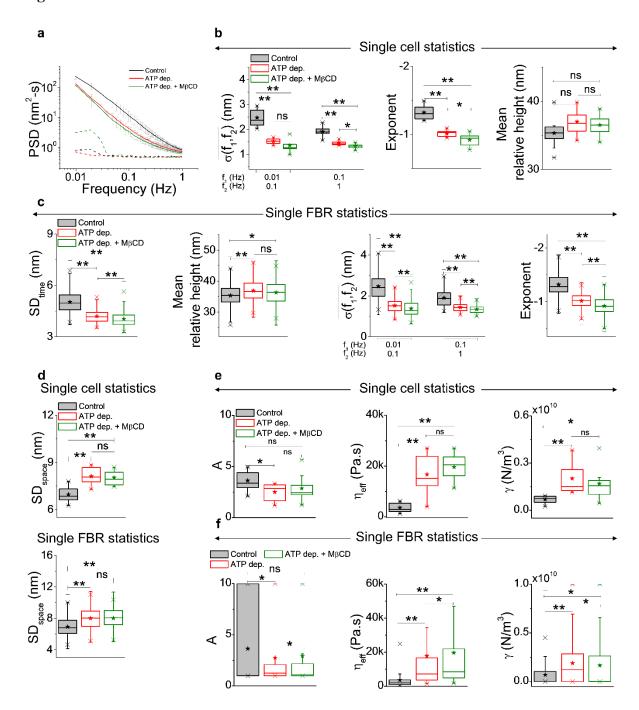


Figure S3: Effect of MβCD on ATP depleted cells. (a) Averaged PSDs of cells (solid lines) and their backgrounds (dashed lines) in control, ATP dep. and ATP dep. + MβCD cells. (b) Single cell statistics of parameters of temporal fluctuations in these conditions. N = 10 cells each. (c) Single FBR statistics of temporal fluctuations parameters in the three mentioned conditions. $n_{control} = 333$ FBRs, $n_{ATPdep.} = 235$ FBRs, $n_{ATPdep.+MβCD} = 250$ FBRs, N = 10 cells each. (d) Single cell statistics (top, N = 10 cells each) and single FBR statistics (bottom, $n_{control} = 333$ FBRs, $n_{ATPdep.} = 235$ FBRs, $n_{ATPdep.+MβCD} = 229$ FBRs, N = 10 cells each) of SD_{space} in all three conditions. (e) Box plots of A, n_{eff} and n_{eff} for single cell statistics. $n_{eff} = 305$ FBRs, Single FBR statistics of the mechanical parameters in the three conditions. $n_{control} = 305$ FBRs,

 $n_{ATPdep.} = 207 \; FBRs, \, n_{ATPdep.+M\beta CD} = 229 \; FBRs, \, N = 10 \; cells \; each. * p \; value < 0.05, ** p \; value < 0.001, \, ns \; p \; value > 0.05, \, Mann-Whitney U \; test. See {\mbox{\bf Table S4}} \; for \; statistics.$

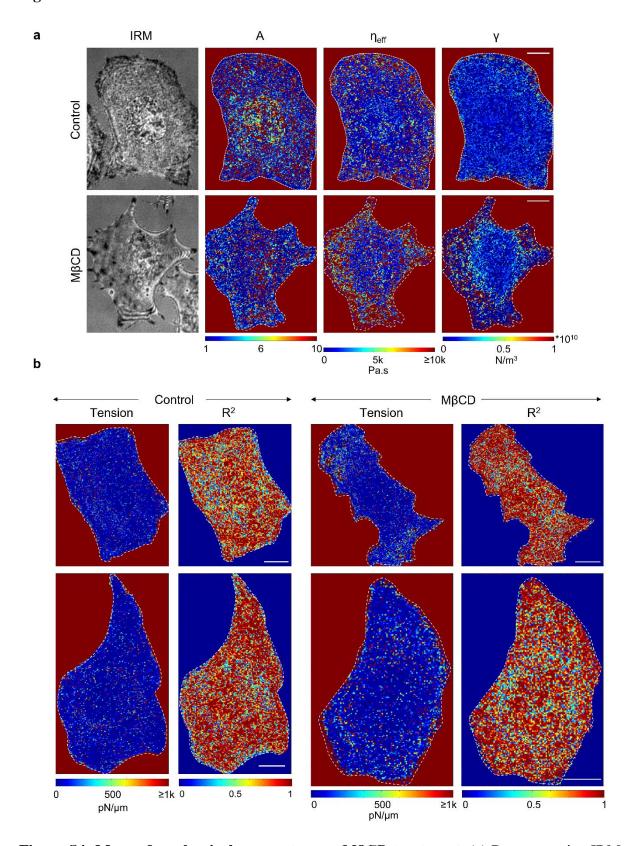


Figure S4: Maps of mechanical parameters on M β CD treatment. (a) Representative IRM images with single pixel maps of active temperature, cytoplasmic viscosity and confinement of a control and M β CD treated cell (tension map in Fig. 4e). Scale bar, 10 μ m. The white dashed

lines mark the cell boundary. Fitting was performed for pixels inside this boundary. (b) Two representative single pixel maps of tension and R^2 for control and cholesterol depleted cells. Scale bar, 10 μ m. The white dashed lines mark the cell boundary. Fitting was performed for pixels inside this boundary.

Figure S5

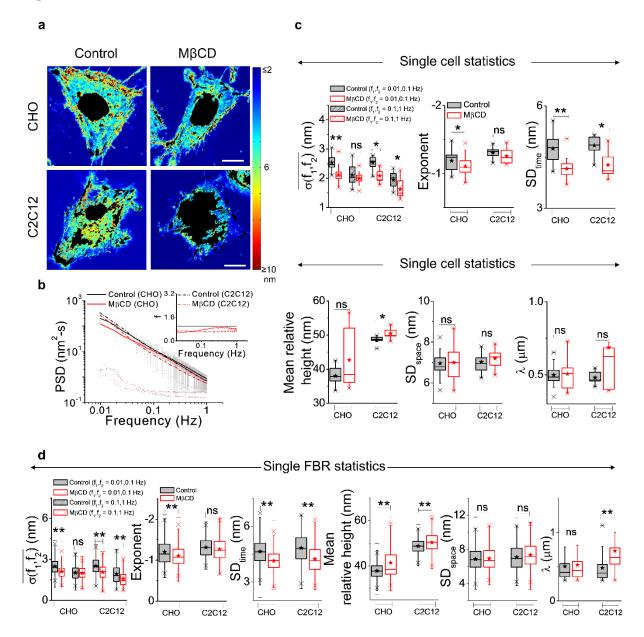


Figure S5: Effect of MβCD on detailed parameters of membrane fluctuations in different cell lines. (a) Representative whole cell SD_{time} maps of control vs. MβCD treated CHO (top, scale bar: $10 \,\mu m$) and C2C12 (bottom, scale bar: $5 \,\mu m$) cells. Non-FBRs are blackened out. (b) Averaged PSDs of CHO (solid lines) and C2C12 (dashed lines) cells in control and cholesterol depleted conditions with their respective backgrounds (dash and dotted lines); inset shows the ratio of the background subtracted PSDs of the two cell lines. (c) Box plots of single cell statistics for the parameters of temporal fluctuations and spatial undulations in both conditions for the two cell lines. N = 30 cells each in CHO, 10 cell each in C2C12. (d) Box plots of single FBR statistics for the parameters of temporal fluctuations and spatial undulations in both conditions for the two cell lines. $n_{\text{CHO control}} = 612 \,\text{FBRs}$, $n_{\text{CHO MβCD}} = 369 \,\text{FBRs}$, $N = 30 \,\text{cells}$ each; $n_{\text{C2C12 control}} = 219 \,\text{FBRs}$, $n_{\text{C2C12 MβCD}} = 179 \,\text{FBRs}$, $N = 10 \,\text{cells}$ each). * p value < 0.05, ** p value < 0.001, ns p value > 0.05, Mann-Whitney U test. See **Table S4** for statistics.

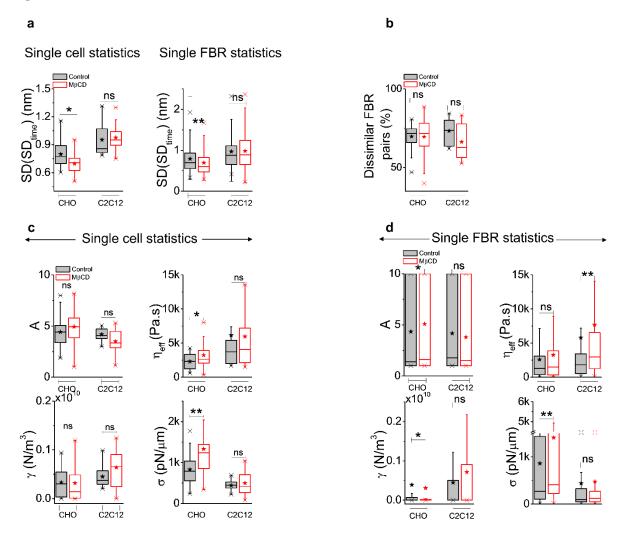


Figure S6: Effect of MβCD on fluctuations heterogeneity and mechanics in different cell lines. (a) A measure of intra-FBR fluctuations heterogeneity, SD(SD_{time}) in control and MβCD treated CHO and C2C12 cells in single cell statistics (left, N = 30 cell each for CHO, 10 cells each for C2C12) and single FBR statistics (right, $n_{CHO\ control}$ = 612 FBRs, $n_{CHO\ MβCD}$ = 369 FBRs, N = 30 cells each; $n_{C2C12\ control}$ = 219 FBRs, $n_{C2C12\ MβCD}$ = 179 FBRs, N = 10 cells each). (b) Intracellular long-range heterogeneity (dissimilar FBR pairs) in CHO (N = 30 cells each) and C2C12 cells (N = 10 cells each). (c) Membrane mechanical parameters A, η_{eff} , γ and σ obtained from fitting PSDs in CHO (N = 30 cells each) and C2C12 cells (N = 10 cells each) in control and cholesterol depletion. (d) Single FBR statistics of the mechanical parameters under the two conditions in the two cell lines. $n_{CHO\ control}$ = 495 FBRs, $n_{CHO\ MβCD}$ = 257 FBRs, N = 30 cells each; $n_{C2C12\ control}$ = 174 FBRs, $n_{C2C12\ MβCD}$ = 124 FBRs, $n_{C4C12\ control}$ = 174 FBRs, $n_{C4C12\ MβCD}$ = 124 FBRs, $n_{C4C12\ control}$ = 495 for statistics.



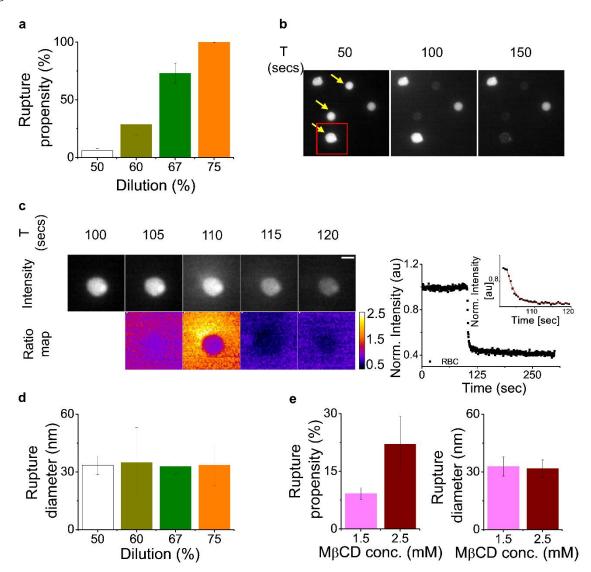


Figure S7: Rupture characteristics of RBCs. (a) Rupture propensity of RBCs with increase in osmotic stress. (b) Time lapse images of Calcein AM loaded RBCs undergoing rupture (arrows in yellow). (c) Intensity (top) and ratio map (bottom) of a rupturing RBC marked in (b) followed in time shows single point rupture. Right: A time profile of normalized mean intensity of a ruptured RBC; inset shows the double exponential fit to the profile. (d) Rupture diameter in RBCs with change in osmotic stress. (e) Rupture propensity (left) and rupture diameter (right) of RBCs treated with increasing concentrations of MβCD without hypoosmotic shock administration. Mean \pm SD of at least two experiments is plotted in each set. See **Table S4** for statistics.

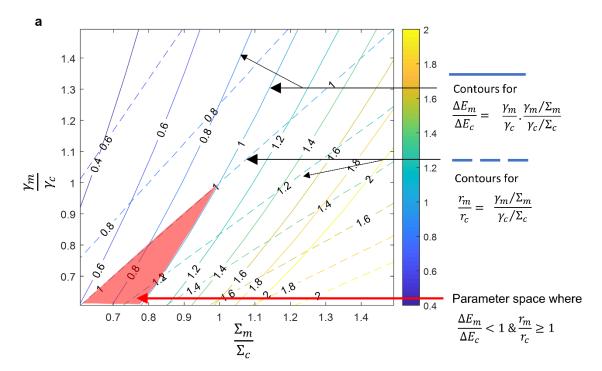


Figure S8: Lowered lysis surface and line tension on MβCD treatment. Contour plots of dependence of ratio of rupture radius of MβCD with control $(\frac{r_m}{r_c})$ & ratio of ΔE of MβCD with control $(\frac{\Delta E_m}{\Delta E_c})$ on the ratio of lysis surface tension between MβCD and control conditions $(\frac{\Sigma_m}{\Sigma_c})$ and on ratio of line tension between MβCD and control conditions $(\frac{\gamma_m}{\gamma_c})$. Contour lines closest to 1 are used to roughly map out the region in the parameter space where the observation of increased rupture propensity of ΔE ratio <1 and unchanged or increased rupture size ratio $r \ge 1$ is true. Note that in this region both lysis surface tension ratio and line tension ratio (MβCD to control) is < 1, indicating lowered lysis surface and line tension on MβCD.

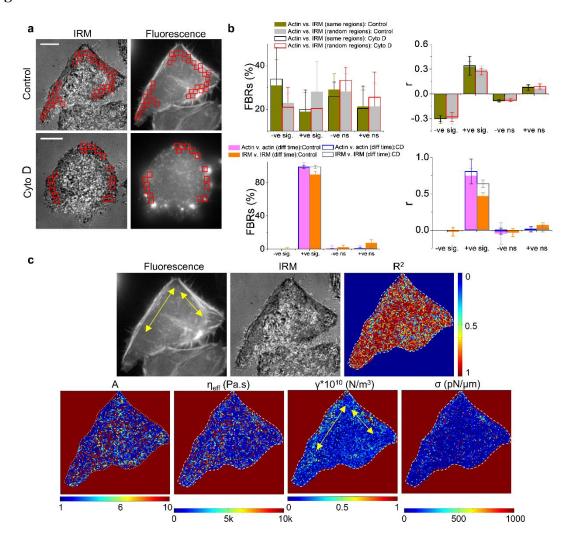


Figure S9: Actin density does not strongly affect IRM intensity at FBRs. (a) IRM and epifluorescence images of mEmerald-Lifeact-7 transfected control (top) and Cyto D treated (bottom) HeLa cells. Scale bar, 5 μ m. (b) Top: plots of no of FBRs that show positive/negative correlation in actin and IRM intensities of same and random FBRs in control and Cyto D treated cells with their corresponding average Pearson correlation coefficients (r). Bottom: plots of no of FBRs that show positive/negative correlation in actin vs. actin and IRM vs. IRM intensities of same FBRs in and Cyto D treated cells with their corresponding average Pearson correlation coefficients (r). (c) Clockwise from top left: Fluorescence and IRM image of a control transfected cell, with its corresponding single pixel R^2 , σ , γ , η_{eff} and A maps. The white dashed line marks the boundary of the cell. The yellow arrows mark the areas in confinement maps that faintly show the presence of stress fibres. Scale bar, 10 μ m.

Figure S10

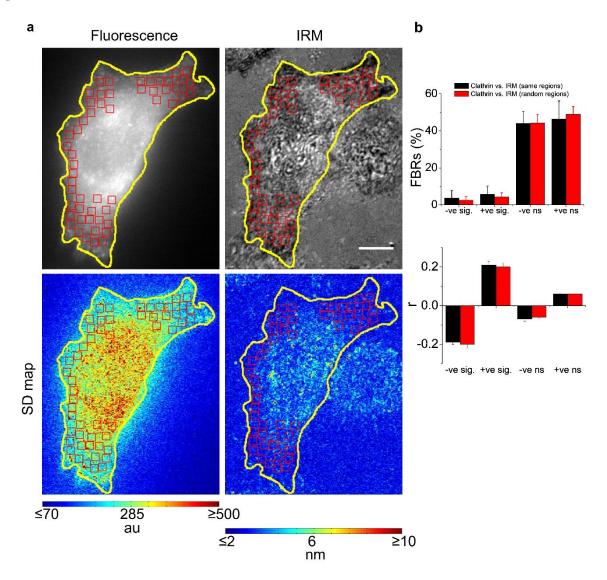


Figure S10: Mobile clathrin pits do not strongly affect IRM intensity fluctuations at **FBRs.** (a) Epi-fluorescence and IRM (top) images of mCherry-Clathrin LC-15 transfected control HeLa cells, with their corresponding SD maps (bottom). The boundary of the cell is marked in yellow. Scale bar, 10 μm. (b) Plots of no of FBRs that show positive/negative correlation in clathrin and IRM intensity fluctuations of same and random FBRs in cells (top) with their corresponding average Pearson correlation coefficients (r, bottom).

Supplementary Discussion

As elaborated in the reference (13), the energy needed to form a circular pore in the lipid bilayer can be written in terms of the line tension (γ), surface tension (Σ) and the radius of the pore (r):

$$E(r) = 2\pi r \gamma - \pi r^2 \Sigma$$

The surface tension at the time of rupture can be termed as lysis tension.

Minimizing energy
$$(\frac{dE}{dr} = 0)$$
, yields $r = \frac{\gamma}{\Sigma}$ (1)

The energy required to cross this critical radius r can be written as:
$$\Delta E = \frac{\pi \gamma^2}{\Sigma}$$
 (2)

In this study, we have two conditions where control is denoted as 'c' and M β CD treatment is denoted 'm'. From experiments, we see that M β CD treatment increases propensity (Fig. S6e). Since the propensity, or probability to rupture in isotonic conditions is expected to be related $_{-\Delta E}$

thus:
$$P \propto e^{\frac{-\Delta E}{k_B T}}$$
, this implies

$$\frac{\Delta E_m}{\Delta E_c} < 1$$
 or, $\frac{\gamma_m^2/\Sigma_m}{\gamma_c^2/\Sigma_c} < 1$ or, $\frac{\gamma_m}{\gamma_c} \cdot \frac{\gamma_m/\Sigma_m}{\gamma_c/\Sigma_c} < 1$ (3)

But, we also know that rupture diameter is unaltered on MβCD treatment (Fig. S6d)

$$\frac{r_m}{r_c} = 1$$
 or, $\frac{\gamma_m/\Sigma_m}{\gamma_c/\Sigma_c} = 1$ (4)

From Eq. 3 and 4, it is evident that:

$$\frac{\gamma_m}{\gamma_c} < 1 \tag{5}$$

This information, together with Eq. 4 implies that $\frac{\Sigma_m}{\Sigma_c} < 1$

or, that the lysis tension of M β CD treated cells needs to be lower than that of control cells.

Observations of increased rupture diameter $(\frac{r_m}{r_c} > 1)$ on M β CD treatment (Fig. 4 e) in hypotonic condition are in line with this inference since enhanced propensity would still need the line tension and hence lysis tension to be lowered by M β CD treatment.

Table S3: Statistical parameters for data presented in main figures.

This is provided as a separate Excel sheet

Table S4: Statistical parameters for data presented in supplementary figures.

This is provided as a separate Excel sheet

Supplementary Movies

Movie S1: Time-lapse imaging of single HeLa cells under control and MβCD treated

conditions at 37 °C under IRM mode.

The movie shows the time evolution of the interference pattern of the basal membrane of single

HeLa cells in control (left) and MβCD treated (right) condition. Scale bar: 10 μm. Stacks of

2048 images are captured at 19.91 frames/sec.

Movie S2: Time-lapse imaging of control and MBCD treated HeLa cells after

administration of 95% hypo-osmotic shock at 37 °C.

The movie shows the time evolution of the fluorescence of Calcein AM loaded HeLa cells

under control (left) and with MβCD (right), after the application of a 95% hypo-osmotic shock.

Scale bar: 100 µm. Images are captured every 2 secs for 5 mins.

Movie S3: Time-lapse imaging of RBCs before and after administration of 67% hypo-

osmotic shock at 37 °C.

The movie shows the time evolution of the fluorescence of Calcein AM loaded RBCs before

(left) and after (right) the application of a 67% hypo-osmotic shock. Scale bar: 100 µm. Images

are captured every 0.5 secs for 5 mins.

Movie S4: Time-lapse imaging of MBCD treated RBCs without the administration of 67%

hypo-osmotic shock at 37 °C.

The movie shows the time evolution of the fluorescence of Calcein AM loaded RBCs that are

treated with 1.5 mM (left) and 2.5 mM (right) MBCD without the application of a 67% hypo-

osmotic shock. Scale bar: 100 µm. Images are captured every 0.5 secs for 5 mins.

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Supplementary References

- 1. Gov, N., A.G. Zilman, and S. Safran. 2003. Cytoskeleton confinement and tension of red blood cell membranes. Phys. Rev. Lett. 90: 228101.
- 2. Thoumine, O., O. Cardoso, and J.-J. Meister. 1999. Changes in the mechanical properties of fibroblasts during spreading: a micromanipulation study. Eur Biophys J. 28: 222–234.
- 3. Kim, T., M.L. Gardel, and E. Munro. 2014. Determinants of Fluidlike Behavior and Effective Viscosity in Cross-Linked Actin Networks. Biophys. J. 106: 526–534.
- 4. Joanny, J., and J. Prost. 2009. Active gels as a description of the actin-myosin cytoskeleton. HFSR J. 3: 94–104.
- 5. Betz, T., M. Lenz, J.-F. Joanny, and C.C. Sykes. 2009. ATP-dependent mechanics of red blood cells. PNAS. 106: 15320–15325.
- 6. Peukes, J., and T. Betz. 2014. Direct Measurement of the Cortical Tension during the Growth of Membrane Blebs. Biophys. J. 107: 1810–1820.
- 7. Alert, R., J. Casademunt, J. Brugués, and P. Sens. 2015. Model for Probing Membrane-Cortex Adhesion by Micropipette Aspiration and Fluctuation Spectroscopy. Biophys. J. 108: 1878–1886.
- 8. Simunovic, M., and G.A. Voth. 2015. Membrane tension controls the assembly of curvature-generating proteins. Nat. Commun. 6: 1–8.
- 9. Curie, M., S. Bernard, and P. Cedex. 2008. 3D Processing and Analysis with ImageJ e. Microscopy. Vi: 1–6.
- 10. Sens, P., and J. Plastino. 2015. Membrane tension and cytoskeleton organization in cell motility. J. Phys. Condens. Matter. 27: 273103.
- 11. Lieber, A.D., S. Yehudai-Resheff, E.L. Barnhart, J.A. Theriot, and K. Keren. 2013. Membrane Tension in Rapidly Moving Cells Is Determined by Cytoskeletal Forces. Curr. Biol. 23: 1409–1417.
- 12. Betz, T., and C. Sykes. 2012. Time resolved membrane fluctuation spectroscopy. Soft Matter. 8: 5317.
- 13. Moroz, J.D., and P. Nelson. 1997. Dynamically Stabilized Pores in Bilayer Membranes. Biophys. J. 72: 2211–2216.