Application of sequential cyclic compression on cancer cells in a flexible microdevice

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Abstract

Mechanical forces shape physiological structure and function within cell and tissue microenvironments, during which cells strive to restore their shape or develop an adaptive mechanism to maintain cell integrity depending on strength and type of the mechanical loading. While some cells are shown to experience permanent plastic deformation after a repetitive mechanical tensile loading and unloading, the impact of such repetitive compression on plastic deformation of cells is yet to be understood. As such, the ability to apply cyclic compression is crucial for any experimental setup aimed at the study of mechanical compression taking place in cell and tissue microenvironments. Here, we demonstrate such cyclic compression using a microfluidic compression platform on live cell actin in SKOV-3 ovarian cancer cells. Live imaging of the actin cytoskeleton dynamics of the compressed cells was performed for varying pressures applied sequentially in ascending order during cell compression. Additionally, recovery of the compressed cells was investigated by capturing actin cytoskeleton and nuclei profiles of the cells at zero time and 24 h-recovery after compression in end point assays. This was performed for a range of mild pressures within the physiological range. Results showed that the phenotypical response of compressed cells during recovery after compression with 20.8 kPa differed observably from that for 15.6 kPa. This demonstrated the ability of the platform to aid in the capture of minute differences in cell behaviour as a result of being compressed at various pressures in physiologically relevant manner. Differences observed between compressed cells fixed at zero time or after 24 h-recovery suggest that SKOV-3 cells exhibit deformations at the time of the compression, a proposed mechanism cells use to prevent mechanical damage. The extent of recovery of compressed cells, inferred by imaging cell membrane bulges and actin cytoskeleton and measuring the shape descriptors of cell nuclei, can give insights into the plasticity of cancer cells. Thus, biomechanical responses of SKOV-3 ovarian cancer cells to sequential cyclic compression and during recovery after compression could be revealed in a flexible microdevice. As demonstrated in this work, the microfluidic platform could control the strength and duration of cyclic compression, while enabling the observation of morphological, cytoskeletal and nuclear changes and recovery in compressed cells, thus providing a powerful new tool for the study of mechanobiological processes in cancer and cell biology.

Introduction

While some cells experience permanent plastic deformation after a repetitive mechanical tensile loading and unloading, the impact of such repetitive compression on plastic deformation of cells remains largely unknown [1,2]. As such, the ability to apply cyclic compression is crucial for any experimental setup aimed at the study of mechanical compression taking place in cell and tissue microenvironments [3–5]. For instance, in ovarian cancer, cells are exposed to chronic compressive stress from different sources such as tumour growth, displacement within stromal tissue and hydrostatic pressure out of ascitic fluid in peritoneal cavity [3,6]. Such compression is estimated to reach 18.9 kPa for human tumours and can exceed 20 kPa based on the experimental data from murine tumours, while those values can be even higher in situ with the impact of the surrounding extracellular matrix in tumour microenvironment [3,7]. Mimicking such chronic compression profiles and physiological pressure values in an *in vitro* dynamic compression system is necessary to further study impact of compressive stress in cancer.

In the presence of a mechanical stress, cells strive to restore their shape to maintain cell integrity after the removal of mechanical loading. However, cells may show a plastic response in the form of cytoskeletal bond ruptures and an incomplete cell shape recovery [1]. Meanwhile, cells produce an adaptive mechanism via these bond ruptures to reduce the mechanical cell stress and thus protect themselves against mechanical damage, while subsequently deforming under the mechanical load [1,8]. Such cell response and adaptive mechanism have been shown by Bonakdar *et al.* for cyclic mechanical tensile loading [1], but comparable investigations into compressive loading remain outstanding. In particular in relation to cancer, a knowledge gap exists in regards to cell deformation and recovery in response to cyclic mechanical compressive loading on living cells.

Asem *et al.* have recently shown that human peritoneal mesothelial cells drastically change their morphology under compression, as evidenced by formation of sub-micrometer scale intercellular projections with changes in the actin cytoskeleton and a resulting more mesenchymal phenotype [9]. Depending on the compression strength and duration, such as 3 kPa for 24-hour, mesothelial cells were observed to exhibit modified nanoscale cell surface projections and to form actin-based tunneling nanotubes (TNTs) used to physically interact with ovarian cancer cells [9].

In general, cells under compression display a change in cell height and thus cell deformation in vertical direction. The nuclei region is the most protruding section of the cell *in vivo* and contributes most to the overall cell height *in vitro* for cells adhered on a substrate. Thus, when cells are compressed, the nuclei and actin cytoskeleton components supporting the cell body are highly mechanoresponsive to the applied compression. Ho *et al.* performed volumetric scanning of the single cells being flatten and observed such a decrease in cell height as per increasing pressures applied externally through a compression control valve [2].

In itself, the nucleus is a mechanosensitive organelle that can respond to solid stress as we have recently shown by investigating nuclear deformations in ovarian cancer cells exposed to compression at increasing pressures [5]. Solid stress-induced nuclear deformations can change the activity of nuclear pore complexes and associated proteins. Such nuclear perturbations in turn modulate the nuclear import of transcription factors, which can alter gene expression and induction of DNA repair programs [10].

With the purpose of applying compression in a cyclic, dynamic and controlled manner for the investigation of cell deformation and recovery, we have recently developed a robust cyclic compression microfluidic method based on a flexible microdevice. This platform provides extensive control of the amount, duration, and mode within a pressure profile. Fabrication and characterization of the multilayer microfluidic platform, and the observation of directional growth of cells, viability and

mechanical cell lysis as end point assays under static state and pressurized states within this platform were demonstrated previously [4,5]. In this paper, we further extend the reproducibility and repeatability validation of our flexible microdevice-based compression method via comparisons of experimental and computational piston actuations for independent microdevices with different membrane thicknesses. We then use the microfluidic platform for the *in vitro* application of cyclic cell compressions that mimic biologically relevant compression profiles occurring in cellular microenvironments. In particular, we demonstrate applicability of the platform for the chronic exposure of ovarian cancer cells to repetitive compressive stress. As such, the current work extends the use of the platform to the application of cyclic compressions at and beyond physiological pressure values in a sequential fashion to study dynamic biomechanical processes by recording GFP-tagged actin dynamics of live cells under compression. In relation to our previous work which discussed initial findings on cellular deformations observed at cyclically applied ascending pressures, the current work also expands applicability by providing a more physiologically relevant setting in form of a hydrogel coating within micro-piston device. We showcase data of cellular deformations and recovery during and after cyclic compression at mild (e.g. 15.6 kPa) and upper mild (e.g. 20.8 kPa) physiological pressure values, obtained via end point assays of actin and nuclei deformations in compressed cells at zero time or at 24 h-recovery after compression, showing flexibility and novel use of our microdevice for the applications of not only cell compression but also recovery. We demonstrate that the platform can control the strength and duration of cyclic compression, thus providing a powerful new tool for the study of mechanobiological processes, by which cell deformation, cytoskeletal and nuclear changes and recovery in cells exposed to such compression can be readily captured.

Materials and methods

Device fabrication and characterization

Microdevice design, fabrication and characterization were explained in detail previously [5]. In brief, the multilayer flexible microdevice is composed of a control microchannel at the top, a polydimethylsiloxane (PDMS) membrane and monolithically attached micro-piston in the middle, suspended in a bottom microchannel enclosed by a glass substrate on which cells are cultured (Fig 1). The layers were fabricated out of PDMS via different soft-lithography methods including standard replica molding for the control layer, spin-coating for the membrane/micro-piston layer, and exclusion molding for the bottom layer. All flexible layers of the microdevice were fabricated from a 10:1 w/w mixture of PDMS (Sylgard 184, Dow) base and curing agent. Curing conditions for each layer, as well as spin-coating settings for PDMS membrane/micro-piston layer, are detailed in [5]. After the curing process was completed, individual layers were assembled via alignment marks in each layer and the aligned PDMS construct was plasma-bonded to the glass. Dimensions of the device compartments (Fig 1(a)) were measured with 3D optical profilometry (Profilm3D, Filmetrics) after each fabrication to ensure repeatability in predicting the external pressure amount via computational modelling. This ensured good control of piston contact pressures applied to cells on-chip during cell compression experiments.

Computational model and experimental validation

A previously developed finite element method (FEM) based on hyperelastic Saint Venant–Kirchhoff theory for actuation of the flexible microdevice [5] was implemented

Fig 1. Device design, operation and applied pressure profiles. (a) 3D optical profilometer image of a micro-piston device showing the compartment including total height of the channel where the micro-piston is suspended, height of the micro-piston and the gap between the piston and surface of the channel. (b) Device operation steps from hydrogel coating, to piston-retracted loading of cells and cyclic compression on cells. The device was kept in static state during hydrogel loading and put into piston-retracted state for cell loading. In this state the membrane, and thus micro-piston, are retracted from the static position towards the top control channel by the applied negative pressure. After loading and settlement of the cells, flexible components were actuated down to their initial position. Cyclic compression of cells is illustrated by the membrane deflection and micro-piston brought onto cells by pressure applied through the control channel and retracted back after compression (repeated cyclically). Representative pressure profiles for piston-retracted loading, and sequential cyclic compression process are shown in (c) and (d), respectively.

in COMSOL Multiphysics (V5.5, COMSOL) and used with different membrane 100
thicknesses obtained from experimental measurements of membranes within the 101
fabricated devices. Vertical displacement and maximum piston contact pressure as per 102
the applied external pressure were simulated and analysed. 103

Cell culture, Matrigel coating and chip loading with cells 104

SKOV-3 ovarian cancer cells (kindly supplied by the Laboratory for Cell and Protein 105
Regulation at the University of Otago, Christchurch) were cultured in Earle's salts and 106
L-glutamine-positive MEM (Gibco) supplemented with 10% fetal bovine serum (FBS, 107
Life Technologies), 1% of penicillin/streptomycin (Life Technologies), and 0.2% 108
fungizone (Life Technologies) in a humidified atmosphere of 5% CO_2 at 37°C. Prior to 109
cell loading, the cell channel of the flexible microdevice was coated with 100 µg/ml 110
Matrigel (Corning, 356237) for 60-70 minutes (Fig 1(b)). Due to the dilute Matrigel 111
solution, the height of hydrogel layer formed inside the microchannel could not readily 112
be measured, however, 100 µg/ml Matrigel was previously shown to facilitate cancer cell 113
adhesion and mimic the extracellular matrix (ECM) microenvironment in comparison to 114

other substrates [11]. Next, SKOV-3 cells at a seeding density of 1.5×10^6 cells/ml were loaded using piston-retracted loading. In this method of chip loading, the PDMS membrane with the attached micro-piston was first retracted out of the culture channel towards the top control channel using an applied negative pressure of 615 (-615) mbar to favour flow of seeding solution for homogenous distribution of cells. Seeded cells were cultured for 3-4 days on-chip until confluency (S1 Fig). One day before experiments, cell media in the channel was changed to CO_2-independent medium (Gibco) supplemented with 10% fetal bovine serum, 2% L-Glutamine (Gibco, 200 mM), 1% of penicillin/streptomycin, and 0.2% fungizone. Following this, cells were cultured overnight in the humidified atmosphere of a CO_2-free incubator at 37°C until live cell compression and imaging on a custom microscope stage (LEC060, LabIVF) pre-warmed to 37°C.

Device operation for sequential cyclic compression

The flexible microdevices were operated with cyclic compression profiles in sequential manner. A pressure ladder with gradually increasing pressure was applied until the micro-piston reached the cells and started to apply contact pressure. The membrane, and thus micro-piston, were actuated with a pressure controller (OB1 Mk3+, Elveflow) coupled to pressure sensors (MSP4, 7 bar, Elveflow) and sensor reader (MSR, Elveflow) operated via Elveflow Smart Interface software (ESI, v3.04). This was used to set the N_2 external gas pressure values required at each step to apply the desired cyclic pressure profiles in a controlled and automated manner. Pressure values in this work are given in mbar for externally applied pressures, as per calibrated settings of the pressure pump, pressure sensor and sensor reader, while kPa is used for the piston contact pressures, which are based on conversion from values computed using mechanical modelling.

Live imaging of GFP-tagged actin cytoskeleton of cells

The actin cytoskeleton profile was quantified by comparing cancer cells with and without applied force. Both, an end point assay and live imaging of the actin cytoskeleton of cells transduced with CellLight^{TM} Actin-GFP, BacMam 2.0 (Invitrogen) were used to investigate the dynamic biomechanical processes in cancer cells under compression. The use of baculovirus for the Actin-GFP transduction was conducted under Environmental Protection Authority (EPA, New Zealand) approval number of GMD101847. Cells were seeded on-chip at a density of 1.5×10^6 cells/ml and cultured overnight, allowing sufficient time for cell adherence. Cells were then transduced on-chip for actin-GFP at 30 PPC (particle per cell) with CellLight^{TM} particles. Following this, cells were incubated for an additional 3 days and the cell media containing the particles was replenished daily until the day of the compression application.

End point assay staining method

After compression experiments, cells were fixed on-chip with 4% paraformaldehyde (Alfa Aesar) for 30 min. The nuclei and actin cytoskeleton of the cells in the cell-culture channels of the micro-piston devices were stained using Hoechst 33342 (Thermo Scientific, 1 µg/ml) and CF 488 A phalloidin (Biotium, 1:40) according to the manufacturers' instructions, with on-chip dye incubation for 20 min and 1.5 h, respectively. Stained nuclei and actin cytoskeleton of the fixed cells were captured with an inverted fluorescence microscope (Nikon Eclipse Ti) equipped with Hamamatsu camera (ORCA-Flash4.0 V2) and HCImage software.

Imaging and data analysis

Live imaging of cyclic compression was performed via time-lapse phase-contrast imaging at a rate of 1 frame per 100 ms. Fluorescence images for actin-GFP were acquired following each step of the sequential cyclic compression. Images were processed and analysed using ImageJ (Fiji) [12]. Regions of interest (ROIs) were drawn around the periphery of the micro-pistons. Corrected total cell fluorescence (CTCF) [13] was calculated as follows:

CTCF = Integrated Density - (Area of micro-piston ROI x Mean fluorescence of background readings)

For the analysis of cell nuclear deformation, ImageJ was used to extract ROIs for the cell nuclear boundaries by applying the analyse particles function to the threshold images with Hoechst epi-fluorescence signals. ROIs obtained from the Hoechst-stained cell nuclei in this way were confirmed using corresponding phase-contrast images for validation of the nuclei segmentation. Area, circularity, and aspect ratio of the cell nuclei were measured from the ROIs. Two-hundred-fold magnification images were used during the analysis of Hoechst-stained nuclei of control (non-compressed) cells in control regions (CRs) around the micro-piston and compressed cells under the micro-piston at zero time or at 24 h-recovery after compression.

Student's t-test was used to determine the statistical significance of live cell actin deformation across increasing pressure ranges, and nuclear deformations in compressed and recovered cells after compression. Statistical significance was taken as $p < 0.05$. Data were represented as mean ± standard error of mean (s.e.m.).

Results and Discussion

Comparison of the experimental and computational results in micro-piston device

To extend on what is possible with existing systems and increase robustness of cyclic cell compression results, control of the pressure application was optimized by evaluating the effect of device fabrication variations. To recall dimensions of the micro-piston device we developed in [5], the height of the cell culture channel was on average 313 µm, while micro-pistons were measured as around 206 µm in PDMS replicas. Once assembled, the gap these layers created between the bottom of the micro-piston and the glass surface was measured as on average 108 µm. In the current work, the thickness of the PDMS membrane attached to micro-pistons varied between 211 µm and 258 µm. Based on the experimental data obtained from optical profilometer measurements of actuated micro-pistons (Fig 1(a)), COMSOL simulations were performed to predict the piston contact pressure (Fig 2) and these pistons were then used to perform cyclic cell compression. Details of the sequential cyclic compression process at mild pressures, up to cell lysis at severe pressures, could be recorded and analysed, together with cell viability [14]. Computational modelling used to calculate the maximum piston contact pressure was further validated experimentally using independent microdevices with different membrane thicknesses to enhance reproducibility and repeatability of our flexible microdevice-based compression method. In our previous work we showed that simulations were in good agreement with optical profilometer measurements of vertical displacement of the PDMS membrane and thus micro-piston at given externally applied pressures [5]. Here, we extended this by providing evidence of close correlation between the computational model and cell compression applications within independent micro-piston devices fabricated with different membrane thicknesses. In particular, when small variations resulted in spin-coated PDMS membrane thicknesses, our model

was able to predict what external pressures needed to be applied through the control microchannel to obtain comparable piston contact pressures on cells, as demonstrated in Fig 2 and Table 1 for thicknesses of 246, 253 and 258 µm as measured from a PDMS membrane coated on a 4-inch photoresist master. The model could also predict the external pressure levels that needed to be applied when the same membrane thickness, such as 235 µm, was used to obtain different piston contact pressures on cells in independent microdevices (Table 1). Measurements further demonstrated that the amount of pressure required to deflect each membrane to bring the micro-piston into contact with cells was in accordance with the computationally-modelled external pressures (Fig 2(a)). A summary of the compression applied to cancer cells via different membrane thicknesses for externally applied pressures and the resulting maximum piston contact pressures is shown in Table 1. Fig 2b further shows that comparable piston contact pressures were achieved by mildly compressing cells with applied external pressures, which were experimentally matching the pressure amounts in simulations ($R^2 = 0.9967$). This indicates that the current model is indeed robust across different membrane thicknesses and independent devices, and can be used to predict the piston contact pressure resulting from externally applied pressures.

Fig 2. Computational and experimental validation of the micro-piston contact pressures across different membrane thicknesses. (a) Simulated piston displacement and maximum piston contact pressures at different membrane thicknesses as function of applied pressure. The left pattern of curves refers to the left y-axis (piston displacement (µm)), while the curve pattern on the right refers to the right y-axis (max piston contact pressure (N/m^2)), resulted at each externally applied pressure shown in x-axis. At a certain applied pressure (x-axis), micro-piston moves towards the bottom substrate and thus shows a displacement (left y-axis) depending on membrane thickness and applies a corresponding contact pressure (right y-axis). Colors of the curves refers to the membrane thicknesses indicated on the graph. (b) Correlation of independent experiments with the pressure sensor read for the externally applied pressure based on simulations run for the corresponding membrane thicknesses, resulting in a similar piston contact pressure.

While inferring these results, the cell compression process was imaged in detail in

Table 1. Compression applied to cancer cells via fabricated different membrane thicknesses, showing externally applied pressures with sensor readings (in parentheses) from experiments and the predicted resulting internal maximum piston contact pressures.

Membrane thickness (µm)	Applied pressure External (mbar)	Piston contact pressure Internal (kPa)
211	354 (350.01)	15.6
222	380 (373.74)	15.9
235	400 (397.47)	12.5
235	410 (409.34)	17.9
235	420 (415.27)	20.7
246	430 (427.94)	15.1
253	450 (450.87)	15.6
258	465 (463.61)	15.6
258	470 (468.68)	17.9

Matrigel-coated microchannels at a speed of 1 frame per 100 ms, as shown in S1 Movie. As illustrated by this time-lapse video, the platform allowed for the temporal evolution of the dynamic cell compression to be investigated in each cycle while the piston was contacting cells and applying compression. Distinct cell membrane bulges could be observed to form during such compression at piston contact pressure of 15.6 kPa and recovery of cell membranes from these bulges was visible when pistons were lifted off during rest stages. Although the platform is capable of temporal control of cell compression (e.g. to decrease, increase or remove the pressure at any certain time interval), duration of each stage in a cycle was set to 5 minutes, corresponding to a 5-min compressed stage followed by 5-min rest stage in each cycle. This was chosen to allow cells enough time during each stage to complete bulge formation or recovery, processes which were considered important for the investigation of cell deformation and recovery after compression. Sensor readings in S1 Movie showed that the micro-piston actuated according to externally applied pressures with reliable control and could be operated in short durations gradually or suddenly when needed. While compression frequencies in an *in vivo* tumor microenvironment are not yet fully known, cyclic compression is important for maintenance of cell culture during mechanical compressions in an *in vitro* experimental setup, in particular as it enhances media flow underneath the micro-pistons. To date, dynamic compression experiments in literature include frequencies of 0.1 – 30 Hz with an applied stress of 5.1, 9.3, 12.9 and 18.7 kPa on breast cancer cells over short durations (30 – 300 s), as used by Takao *et al.*, who observed a mixed mode of apoptosis and necrosis-dominant cell death in mechanically compressed monolayers in a bulk compression platform [15]. In contrast, work by Novak *et al.* used a frequency of 0.05 Hz for cyclic compression applied at 3.9 – 6.5 kPa for 24 hours [3]. They observed an increase in proliferation capacity and decrease in apoptosis for OVCAR3 and OVSAHO ovarian cancer cells cultured in 3D hydrogel components in a bioreactor at millimeter scales. In another example, Ho *et al.* applied cyclic compression by alternating between external pressures of 10 and 15 psi (68.9 and 103.4 kPa) at 0.5 Hz for 6 minutes [2]. Using a single cell microfluidic compression platform, their application resulted in full recovery and thus no permanent plastic deformation in MCF10A normal breast epithelial cells after compression [2]. In the following we add to these examples by using the capabilities of our micro-piston platform to apply cyclic compression with a low frequency in microfluidic settings. In particular, we utilize this functionality to provide a first investigation of the effects of cell compression,

deformation and recovery after compression on SKOV-3 ovarian cancer cells. As part of this we provide evidence that the amount of the pressure, duration of the application and number of the cycles clearly affects morphology and compression in this cell type.

Live cell actin profiles of cancer cells during sequential cyclic compression

In addition to the observation of cell membrane bulges, we were further able to track how the application of sequential cyclic compression led to dynamic changes in GFP-tagged live cell actin as a function of applied pressure (see S2 Movie). As mentioned in the section Live imaging of GFP-tagged actin cytoskeleton of cells, for these experiments SKOV-3 ovarian cancer cells were transduced on-chip with CellLight Actin-GFP, BacMam 2.0 baculovirus to visualize the dynamic biomechanical processes during compression (Fig 3). Specifically, the actin cytoskeleton profile was quantified while cells were in compressed and rest stages (Fig 4). For this, the actin-GFP transduced cancer cells were dynamically stimulated and compressed with various intermediate actuation pressures in a sequential cyclic manner, namely following a sequence from Mild (15.6–15.9 kPa) to Intermediate 1 (23.8–26.8 kPa), Intermediate 2 (37.8–41.7 kPa) and Severe (127.8–140 kPa) piston contact pressures. The first two ranges of this sequence fell within physiological and upper physiological pressure values and cell viability was high (on average 94% for Mild and 77% for Intermediate 1), while Intermediate 2 and Severe levels constitute hyperphysiological values and cell viability decreased (on average 43% for Intermediate 2 and 20% for Severe) [5]. Similar as to the sequence shown in Fig 1(d), first, a pressure amount ladder was formed up to a pressure in the mild compression range, represented by an externally applied pressure of 350 mbar and resulting in a piston contact pressure of 15.6 kPa, during which the piston reached the cells at the bottom surface and cells started to distinctly deform. Once this was observed, cyclic compression was applied at mild pressure for a total of 1 hour, with cells compressed for 5 min at 15.6 kPa, followed alternately by a rest for 5 min at 0 kPa. To track cell response, actin-GFP signals of the compressed cells were captured before compression, as well as during the first and last cycles of the cyclic compression stage. At the end of this sequence, the externally applied pressure was sequentially increased first to 370 mbar, then to 400 mbar and finally 640 mbar, resulting in the respective piston contact pressures of 23.8 kPa (Intermediate 1), 37.8 kPa (Intermediate 2) and 140 kPa (Severe). During this stage, cells were compressed for up to 2 min at each pressure (Fig 1(d)) and actin-GFP signals of the compressed cells were captured (Fig 3).

Observations were quantified based on CTCF and results are plotted as shown in Fig 4. A clear correlation between the actin-GFP signal of the cells compressed under the piston and the applied pressure could be observed, with the signal decreasing with increasing pressure. As such, the results illustrate how the actin cytoskeleton of cancer cells changes in response to compressive stress under live-cell conditions. The change in actin signal for compressed cells under micro-piston was statistically different compared to non-compressed cells in control regions around the micro-piston at all applied compression levels ($p < 0.05$, see S1 File). Actin signal changes in the control regions on the other hand, remained statistically insignificant ($p > 0.05$). At the higher pressures of Intermediate 2 and Severe, actin of most of the cells was disrupted compared to in the initial stages of Mild compression ($p < 0.05$, see S1 File). Interestingly, a few cells usually remained resistant to the increasing pressure ranges, as evidenced by their retained fluorescent signal while under the micro-piston. CTCF during the rest stage after upper mild compression at Intermediate 2 level was statistically significantly different from the stage before compression ($p = 0.0229$) and the rest stage after first cycle of Mild compression ($p = 0.0352$), which indicates that Intermediate 2 level caused

a disruption of cell actin. This disruption was significantly higher again at Severe level compared to other levels (p <0.05, see S1 File). As such, these results expand on the work by Ho *et al.*, who used a more limited pressure and frequency range, and observed formation of new actin stress fibers and thickening of a pre-existing actin structure in a very limited number of cells [2]. In contrast, the current work observed live-cell actin in the cytoskeleton from deformation to ruptures at a wider range of applied pressures across several stages of the sequential cyclic compression.

Fig 3. Representative fluorescent microscopy images of live imaging of GFP-tagged actin cytoskeleton of cells. (a) Schematic of the on-chip BacMam transduction into cancer cells for the expression of actin-GFP fusion protein. (b) Phase contrast (right) and fluorescent (left) images of the GFP-tagged actin expressing SKOV-3 ovarian cancer cells in static state (0 kPa) or retracted back to static state (0 kPa) after compression for the given pressure range. (c) Phase contrast (left) and fluorescent (right) images showing the change in GFP-tagged actin in the cells as a result of the applied pressure in the compressed state. Pressure was applied sequentially in ascending order.

Fig 4. Corrected total cell fluorescence (CTCF) calculated from images of GFP-tagged actin cytoskeleton of cells during sequential cyclic compression at the indicated pressures. Results represent at least 3 independent experiments.

The current device also allowed the response of individual cells at the interface between compressed and non-compressed regions in the periphery of the micro-piston to be studied, thus providing a degree of control of the localization of compression (see S2 Movie). As depicted in the video, parts of cells at the interface were subject to compression, while the rest remained uncompressed. Thus, the cell actin was deformed and damaged only partially in the compressed part depending on the magnitude of the applied pressure, whereas it remained more intact in the non-compressed part. This functionality which is unique to our platform by having control and compressed groups in the same microdevice, is useful to mimic partial cellular deformation and damage that can occur *in vivo* in presence of a localized mechanical loading.

Actin cytoskeleton and nuclei profiles of cancer cells after compression and recovery

Applicability of the cyclic compression on the platform was further demonstrated by recording actin and nuclei deformation as an end point assay in cancer cells fixed at zero time or after 24 h-recovery following 1 hour-long cyclic compressions. To this end, we show progression of the compressed cells when live culture was maintained for 24 h-recovery after compression (S2 Fig). Nuclear shape analysis was performed on fluorescence images of control (non-compressed) and compressed cells at zero time or 24 h-recovery after compression (S3 Fig, Fig 5(a) and S4 Fig). No significant change was observed between the control and compressed cells at 15.6 kPa for areal and axial cell nuclei deformation, while compressed cells had larger nuclei than 24 h-recovered compressed cells (Fig 5(b,d)). Conversely, circularity of the cell nuclei was significantly reduced in the compressed cells compared to the control (non-compressed) at zero time after compression, while not significant in the 24 h-recovered compressed group compared to control (Fig 5(c)). Differences between the compressed cells fixed at zero

time or after 24 h-recovery suggest that cells exhibit deformations at the time of the ₃₄₂
compression, a proposed mechanism cells use to prevent mechanical damage [1,16]. ₃₄₃
These deformations were not permanent morphologically however, and could be ₃₄₄
recovered at mild pressures, such as 15.6 kPa. The absence of statistical difference in ₃₄₅
nuclei circularity between the 24 h-recovered compressed cells under micro-pistons and ₃₄₆
that of the non-compressed cells in the control region within those recovery samples ₃₄₇
(Fig 5(c)) shows that 24 h-recovered compressed cells started to behave like ₃₄₈
non-compressed cells during the recovery after compression at 15.6 kPa. Nonetheless, ₃₄₉
the absence of statistical difference for nuclei circularity between the zero-time and ₃₅₀
24 h-recovered cell groups after compression suggests that cells exists that show no full ₃₅₁
recovery. This divergence supports the existence of heterogeneity in the response of ₃₅₂
cancer cells to mild compression [17]. Although the developed platform is not a ₃₅₃
single-cell compression platform, cells could be assessed as individual cells for their ₃₅₄
nuclei despite overlapping cellular parts due to dense culture formation on Matrigel ₃₅₅
coatings. At the same time, the platform could be considered superior compared to ₃₅₆
single-cell systems in that it allows for cell-to-cell communication within cell monolayers ₃₅₇
and cultures on Matrigel coatings on-chip, with conditions in the presence of mechanical ₃₅₈
compression. Overall, the above experiments constitute a first determination of ovarian ₃₅₉
cancer cell response to lower physiological compression values, such as 15.6 kPa, applied ₃₆₀
in a cyclic manner and over 1 hour. To further understand whether cells exhibit ₃₆₁
different cell recovery profiles after compression, upper physiological values were tested ₃₆₂
as well, results of which are discussed in the following. ₃₆₃

Utilising the full pressure range capabilities of the platform, we also investigated cell ₃₆₄
recovery after compression at upper physiological values, such as 20.8 kPa (S2 Fig). For ₃₆₅
this, cells that were compressed at upper mild pressures were assessed for zero time and ₃₆₆
24 h-recovery after the compression and compared to non-compressed control cells in ₃₆₇
each group (Fig 6 and S5 Fig). Cells treated this way developed distinct actin ₃₆₈
deformation, as shown by the increased fluorescence signals at the edges of the ₃₆₉
compressed cells under the micro-piston at zero time after cyclic compression at ₃₇₀
20.8 kPa for 1 hour. Some cells had their actin visibly disrupted over the nuclei and ₃₇₁
cytoplasm, while highly increased stress fibers presented at the cell membrane ₃₇₂
(Fig 6(a)). This result is consistent with changes in actin cytoskeleton and formation of ₃₇₃
cell surface projections observed in compressed peritoneal mesothelial cells in the study ₃₇₄
by Asem *et al.* [9]. Interestingly, in the current work cell actin signal was localized ₃₇₅
mostly in cells under micro-piston at 24 h-recovery after cyclic compression ₃₇₆
(Fig 6(a)), compared to control cells (S5 Fig). For compressed cells this outcome ₃₇₇
indicates no full recovery from the impact of the compression at 20.8 kPa, and thus an ₃₇₈
altered actin cytoskeleton profile. Shape descriptors of the cell nuclei were measured for ₃₇₉
zero-time and 24 h-recovery after the cyclic compression. There was no statistical ₃₈₀
difference in nuclei area between compressed or 24 h-recovered compressed groups and ₃₈₁
with their control groups (Fig 6(b)). On the other hand, circularity of the compressed ₃₈₂
and 24 h-recovered compressed cells was statistically different both between each and ₃₈₃
respective control groups (Fig 6(c)). Furthermore, the aspect ratio of the 24 h-recovered ₃₈₄
compressed cell nuclei was statistically significantly higher than its 24 h-recovered ₃₈₅
control group and compressed group at zero time after compression (Fig 6(d)). These ₃₈₆
results suggest that cells did not fully recover after cyclic compression at upper mild ₃₈₇
physiological pressure values such as 20.8 kPa. The phenotypical response of ₃₈₈
compressed cells during recovery after compression at 20.8 kPa differed observably from ₃₈₉
that for 15.6 kPa, demonstrating the capability of the flexible microdevice to capture ₃₉₀
minute differences in cell behaviour after being compressed at various pressures in a ₃₉₁
physiologically relevant manner. In fact, such changes in cell actin and nuclei shape ₃₉₂
during recovery after upper mild compression as observed here may indicate formation ₃₉₃

Fig 5. End point assay staining and analysis for actin and nuclei of cancer cells fixed at zero time or after 24 h-recovery, following 1 hour-long cyclic compression between 0 and 15.6 kPa. (a) Representative images of control and compressed cell groups stained for actin (green) and nuclei (blue) for their shape at zero time and 24 h-recovery after cyclic compression. Dashed areas are under piston, while the surrounding is control region (CR). Merged: merged form of the phalloidin (stain for actin) and Hoechst (stain for nuclei) epi-fluorescence images; merged with PH: merged form of the epi-fluorescence images with the corresponding phase-contrast (PH) image; magnified: two-hundred-fold magnification images of the region under micro-piston. Representative arrows (white) show the distinct actin deformation by the increased fluorescence signals at the edges of the cells in the compressed groups under the micro-piston at zero time after compression. Nuclei deformation in control cells at control regions (CR) versus in compressed cells under piston at zero time or 24 h-recovery after cyclic compression, was measured for area (b), circularity (c) and aspect ratio (d) of the nuclei (mean ± s.e.m. n = 588, 230, 358, 174 cell nuclei). Dashed horizontal bars show significant differences between the control region (CR) group and compressed group under micro-piston. Continuous horizontal bars show the comparison pairs between the cell groups under the micro-piston at zero time and 24-recovery after compression.

of more aggressive and metastatic cells [18].

In the study by Ho *et al.*, the height of breast epithelial cells recovered fully after 6 minutes of cyclic compression applied between the external pressures of 10 and 15 psi at 0.5 Hz [2]. No statistical difference in cell height before and after compression was observed, indicating that the breast epithelial cells did not exhibit permanent plastic deformation. Measurement of the viscoelastic relaxation of the cells is difficult however, especially when vertical and temporal resolution in imaging are limited, and long image acquisition times required for 3D visualization might prevent capturing the changes in cell height at the time of compression. In the current study, we thus deliberately limited

Fig 6. End point assay staining and analysis for actin and nuclei of cancer cells fixed at zero time or after 24 h-recovery, following 1 hour-long cyclic compression between 0 and 20.8 kPa. (a) Control and compressed cell groups stained for actin (green) and nuclei (blue) for their form at zero time and 24 h-recovery after cyclic compression. Dashed areas are under piston, while the surrounding is control region (CR). Merged: merged form of the phalloidin (stain for actin) and Hoechst (stain for nuclei) epi-fluorescence images; merged with PH: merged form of the epi-fluorescence images with the corresponding phase-contrast (PH) image; magnified: two-hundred-fold magnification images of the region under micro-piston. Representative arrows (white) show distinct actin deformation in form of increased fluorescence signals at the edges of the cells in the compressed groups under the micro-piston at zero time after compression. Several cells can be observed to have their actin disrupted over the nuclei and cytoplasm, with highly increased stress fibers present at the membrane. Nuclei deformation in control cells at control regions (CR) versus in compressed cells under piston at zero time or 24 h-recovery after cyclic compression, measured for area (b), circularity (c) and aspect ratio (d) of the nuclei (mean ± s.e.m. n = 406, 162, 342, 143 cell nuclei). Dashed horizontal bars show significant differences between the control region (CR) group and compressed group under micro-piston. Continuous horizontal bars show the comparison pairs between the cell groups under the micro-piston at zero time and 24-recovery after compression.

cell deformation assessment to planar dimensions including area, circularity, and aspect ratio [12], as wide-field imaging is considered fast enough (e.g. milliseconds long) to capture cell states during the steps of the compression process compared to real-time volumetric imaging (e.g. minutes long) using confocal microscopy [2,5,19–22].

Cyclic compression in the study of Ho *et al.* was applied only for 6 minutes in total, significantly shorter than the entire duration of a cyclic compression in this work, and results presented here suggest that cell recovery after compression depends on the temporal length of the compression on cells. The overall extent of recovery of

compressed cells represented by cell membrane bulges, the actin cytoskeleton and measurements of shape descriptors of cell nuclei can yield further insight into the plasticity of the cancer cells. Particularly observable in S1 Movie, cell plasticity depends on the amount of pressure applied, as well as the timing. Cell bulge formation slowly developed during the compression stage while cells were responding to contact pressure at the interface with the PDMS piston. This was despite the fact that the piston was actuated onto the cells rapidly as directed by the external pressure controller. As shown by the pressure sensor readings in the same movie, PDMS membrane and piston were consistently responding to amount, duration and mode of the applied pressure. Thus, the gradual deformation of cells and bulge formation during compression stages must originate in the response type of the cells to mechanical load when in solid contact with the PDMS piston. During rest stages, where the compression was lifted, cell bulges did not fully recover right after compression was removed, suggesting that the cells require a certain amount of time to recover compressed cellular parts (S1 Movie). Based on these observations of temporal evolution of the cell compression application, and comparison of application durations in this study with the study by Ho *et al.* [2], we hypothesize that this recovery also depends on the duration and number of the cycles. Based on this we conclude that the presented sequential cyclic compression and cell recovery after compression method should be employed in the future to study the effects of compression and decompression in intratumor vessels or endothelial cells to help understand the structure and functionality of decompressed vessels, importance of which was shown in a study by Padera *et al.* [23].

In summary, our platform was shown to be able to control the strength and duration of cyclic compression, all while enabling the observation of morphological, cytoskeletal and nuclear changes in cells. It thus provides a powerful new tool for the study of mechanobiological processes in cancer and cell biology. Furthermore, the data obtained in this work suggest that ovarian and likely other cancer cells can exhibit a plastic cell response to applied sequential cyclic compression. Cells in the current work were able to do so by deformations and perturbations in actin cytoskeleton and nuclei. It is proposed that cells exhibit such deformations at the time of compression to compensate for the applied compressive stress and protect themselves against mechanical damage. Those deformations were recovered fully or partially depending on the amount of applied pressure. At upper mild pressures, such as 20.8 kPa, cells appear to develop and use an adaptive mechanism through incomplete shape recovery. Whereas recovery after mechanical loading is important to maintain cell integrity, an adaptive mechanism in the form of incomplete shape recovery might indicate the emergence of more aggressive and metastatic cancer cells.

Conclusion

We have demonstrated coating of cell culture chambers in micro-piston devices with hydrogels and successful culture of SKOV-3 ovarian cancer cells therein, mimicking the *in vivo* cellular microenvironment. Furthermore, micro-piston devices fabricated with different membrane thicknesses were used to cyclically compress cells with controlled micro-piston contact pressures, verified using FEM simulations to account for possible membrane thickness variations. Computational results showed good agreement with experimentally applied cyclic compressions. Live cells were imaged in the micro-piston devices and mild compression was applied to mechanically stimulate and record changes in actin profile dynamics. This was extended to dynamic cyclic compressions and recording of the long-term cellular changes in mechanosensitive subcellular structures, such as actin cytoskeleton and nuclei, while allowing cells to recover for 24 hours after compression. Details of morphological cell deformations during sequential cyclic

compression were captured via high-speed imaging. Membrane bulges in cells fixed ₄₆₁ directly after compression appeared as increased actin signal at cell edges. We also ₄₆₂ conclude that recovery of the compressed live cells was dependent on magnitude of the ₄₆₃ applied compression. As demonstrated here with controlled micro-scale mechanical cell ₄₆₄ compression and recovery in a flexible microdevice, our comprehensive method will ₄₆₅ provide a more accurate replication of cell-physiological mechanisms to study both ₄₆₆ short- and long-term effects of compression in cellular microenvironments. ₄₆₇

Supporting information

S1 Fig. On-chip growth of SKOV-3 cells on Matrigel-coated glass surfaces in micro-piston devices. Representative images of two samples during culture from zero time to Day 4 after piston-retracted loading of cells at -615 mbar.

S2 Fig. Examples of 24 h-recovery of mildly and upper-mildly compressed SKOV-3 cells on Matrigel-coated glass surfaces in micro-piston device.

S3 Fig. Image analysis of nuclear shape after cell compression on Matrigel-coated surfaces. Regions of interest (ROIs) extracted for the cell nuclear boundaries in Hoechst (stain for nuclei) epi-fluorescence channel (blue in merged/composite frames) were used to measure area, circularity, and aspect ratio of the cell nuclei, as independent of phalloidin (stain for actin) epi-fluorescence channel (green in merged/composite frames). Merged - 200X: merged form of the phalloidin and Hoechst epi-fluorescence images obtained at two-hundred-fold magnification; merged with PH - 200X: merged form of two-hundred-fold magnification epi-fluorescence images with the corresponding phase-contrast (PH) image for the region under micro-piston; merged with PH - 100X: merged form of one-hundred-fold magnification epi-fluorescence images with the corresponding phase-contrast (PH) image for the region under and around micro-piston in channel.

S4 Fig. End point assay staining and analysis for actin and nuclei of cancer cells in control region (CR) fixed at zero time or after 24 h-recovery, following 1 hour-long cyclic compression between 0 and 15.6 kPa. Control cell groups stained for actin (green) and nuclei (blue) for their form at zero time and 24 h-recovery. Merged: merged form of the phalloidin (stain for actin) and Hoechst (stain for nuclei) epi-fluorescence images; merged with PH: merged form of the epi-fluorescence images with the corresponding phase-contrast (PH) image; Random region 200X: two-hundred-fold magnification images of control cells in random region as part of the control region.

S5 Fig. End point assay staining and analysis for actin and nuclei of cancer cells in control region (CR) fixed at zero time or after 24 h-recovery, following 1 hour-long cyclic compression between 0 and 20.8 kPa. Control cell groups stained for actin (green) and nuclei (blue) for their form at zero time and 24 h-recovery. Merged: merged form of the phalloidin (stain for actin) and Hoechst (stain for nuclei) epi-fluorescence images; merged with PH: merged form of the epi-fluorescence images with the corresponding phase-contrast (PH) image; Random region 200X: two-hundred-fold magnification images of control cells in random region as part of the control region.

S1 Movie. Temporal evolution of the dynamic cell compression within Matrigel-coated micro-piston devices. Time-lapse live cell imaging with 100 ms per frame was recorded for ladder pressure increase from 0 kPa up to 15.6 kPa and a short cycle at 15.6 kPa with gradual increase (from 0 kPa to 15.6 kPa) and decrease (from 15.6 kPa to 0 kPa) in 30 s for each. These steps were sequentially followed by 1 h-long cyclic compression between 0 kPa and 15.6 kPa (piston contact pressure from simulation). Cells were compressed in the temporal evolution of the dynamic pressure control within Matrigel-coated micro-piston devices and dynamic changes in cells during the cyclic compression were captured in detail at a rate of 100 ms per frame.

S2 Movie. Representative data recording of the dynamic changes of GFP-tagged live cell actin during application of sequential cyclic compression at various applied pressures. SKOV-3 ovarian cancer cells were transduced on-chip with CellLight Actin-GFP, BacMam 2.0 baculovirus (Invitrogen) to study the dynamic biomechanical processes under compression. The actin cytoskeleton profile was quantified for the cancer cell response under applied forces at compressed and rest stages during the compression process.

S1 File. Corrected total cell fluorescence (CTCF) data for actin integrated density and statistical test results.

Acknowledgments

The authors thank Helen Devereux, Gary Turner, and Nicole Lauren-Manuera for assistance. We also thank Mathieu Sellier for help with COMSOL, and Campbell Sheen for assistance with the EPA protocol to use the Actin-GFP BacMam construct. Financial support was provided by the MacDiarmid Institute for Advanced Materials and Nanotechnology and the Biomolecular Interaction Centre. Additional funding for V.N. was provided by Rutherford Discovery Fellowship RDF-19-UOC-019. We also thank University of Canterbury for Faculty of Engineering PhD Publishing Scholarship to S.O.

Competing interests

The authors have declared that no competing interests exist.

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