



OPEN

## Drug resistant pancreatic cancer cells exhibit altered biophysical interactions with stromal fibroblasts in imaging studies of 3D co-culture models

Eric Struth<sup>1</sup>, Maryam Labaf<sup>2,3</sup>, Vida Karimnia<sup>1</sup>, Yiran Liu<sup>1</sup>, Gwendolyn Cramer<sup>1,6</sup>, Joanna B. Dahl<sup>4</sup>, Frank J. Slack<sup>5</sup>, Kourosh Zarringhalam<sup>2,3</sup> & Jonathan P. Celli<sup>1,3</sup>✉

Interactions between tumor and stromal cells are well known to play prominent roles in progression of pancreatic ductal adenocarcinoma (PDAC). As knowledge of stromal crosstalk in PDAC has evolved, it has become clear that cancer associated fibroblasts can play both tumor promoting and tumor suppressive roles through a combination of paracrine crosstalk and juxtacrine interactions involving direct physical contact. Another major contributor to dismal survival statistics for PDAC is development of resistance to chemotherapy drugs, though less is known about how the acquisition of chemoresistance impacts upon tumor-stromal crosstalk. Here, we use time lapse imaging and image analysis to study how co-culture geometry impacts interactions between epithelial and stromal cells. We show that extracellular matrix (ECM) overlay cultures in which stromal cells (pancreatic stellate cells, or normal human fibroblasts) are placed adjacent to PDAC cells (PANC1) result in direct heterotypic cell adhesions accompanied by dramatic fibroblast contractility. We analyze these interactions in co-cultures using particle image velocimetry (PIV) analysis to quantify cell velocities over the course of time lapse movie sequences. We further contrast co-cultures of PANC1 with those containing a drug resistant subline (PANC1-OR) previously established in our lab and find that heterotypic cell–cell interactions are suppressed in the latter relative to the parental line. We use RNA-seq and bioinformatics analysis to identify differential gene expression in PANC1 and PANC1-OR, which shows that negative regulation of cell adhesion molecules, consistent with increased epithelial mesenchymal transition (EMT), is also correlated with reduction in the heterotypic cell–cell contact necessary for the contractile behavior observed in drug naïve cultures. Overall these findings elucidate the role of drug-resistance in inhibiting an avenue of stromal crosstalk which is associated with tumor suppression and also help to establish cell culture conditions useful for further mechanistic investigation.

Pancreatic ductal adenocarcinoma (PDAC) is among the most lethal of human malignancies, claiming the lives of more than three quarters of those who are diagnosed within the first year<sup>1</sup>. Resistance to traditional chemotherapy agents remains a major challenge in clinical management of PDAC. A significant contributor to this is the profound desmoplastic reaction which is characteristic of PDAC. The resultant dense fibrotic stroma is implicated in promoting tumor progression and survival and impeding treatment efficacy<sup>2–5</sup>. Indeed this has motivated the exploration of stromal depletion treatments, targeting the tumor stroma itself, though with mixed success<sup>6,7</sup>. Early efforts to therapeutically target cancer associated fibroblasts (CAFs) were promising, but subsequent studies

<sup>1</sup>Department of Physics, University of Massachusetts Boston, Boston, MA 02125, USA. <sup>2</sup>Department of Mathematics, University of Massachusetts Boston, Boston, MA 02125, USA. <sup>3</sup>Center for Personalized Cancer Therapy, University of Massachusetts Boston, Boston, MA 02125, USA. <sup>4</sup>Department of Engineering, University of Massachusetts Boston, Boston, MA 02125, USA. <sup>5</sup>Department of Pathology, Beth Israel Deaconess Medical Center, Harvard Medical School Initiative for RNA Medicine, Boston, MA 02115, USA. <sup>6</sup>Present address: Department of Radiation Oncology, Perelman School of Medicine, University of Pennsylvania, Philadelphia, PA, USA. ✉email: jonathan.celli@umb.edu

showed that indiscriminant depletion of stromal fibroblasts can lead to more aggressive disease progression<sup>8,9</sup>. Similarly, efforts to improve chemotherapy delivery by targeting Hedgehog pathway signaling yielded promising preclinical results<sup>10</sup>. However this approach ultimately failed in the clinic and subsequent work indicated that this approach targets stromal elements which otherwise act to restrain PDAC progression<sup>11</sup>.

Stromal involvement in PDAC is complex, with cancer associated fibroblast subpopulations that exhibit both tumor suppressive and tumor-promoting roles. Distinct phenotypic subtypes of CAFs have been identified as inflammatory CAFs (iCAFs) and myofibroblastic CAFs (MyCAFs)<sup>12–14</sup>. In the tumor microenvironment, iCAFs, which are not typically in direct physical contact with cancer cells, activate pathways that promote tumor proliferation and survival via paracrine signaling. MyCAFs interact with epithelial cells through direct cell–cell adhesions, which can play tumor suppressive roles by physically restraining invasive progression. This juxtacrine signaling is mediated by cell–cell adhesions and the genes that code for adhesion molecules, which are also associated with regulation of PDAC progression<sup>15–17</sup>.

In PDAC, as with other carcinomas, drug resistance is one of the most significant barriers to achieving successful treatment outcomes<sup>18</sup>. Through both intrinsic and acquired resistance, tumors often are, or become, non-responsive to chemotherapy agents, leading to recurrence and relapse even when initial treatment response is positive. Non-responsiveness to therapeutics is compounded by association between drug resistance and epithelial mesenchymal transition (EMT), which acts as a regulator of the tumor-initiating cancer stem cell (CSC) phenotype<sup>19</sup>. EMT, which is directly linked with drug resistance in PDAC<sup>20</sup>, also imparts increased metastatic potential<sup>21</sup>. As such, the tumor cell populations which are non-responsive to treatment are also associated with lethal metastatic progression. Less is known however about how the acquisition of chemoresistance alters heterotypic interactions between tumor and stromal cells in the PDAC microenvironment, especially in view of relatively recent knowledge of CAF subpopulations. In the present study we examine the role of drug-resistance using a chemoresistant sub-line previously established in our lab by exposure of PANC1 cells to oxaliplatin chemotherapy over multiple passages. The drug-resistant cell-line PANC1-OR, which has been stable over multiple passages and cryopreservation, exhibits a clinically-relevant oxaliplatin and gemcitabine-resistant phenotype along with increased invasive behavior, and phenotypic traits consistent with increased EMT<sup>22,23</sup>.

In this study we use in vitro 3D co-culture models to examine (A) the role of co-culture geometry in recreating interactions between adjacent and distant cells, and (B) how these interactions become altered when PDAC cells acquire chemoresistance. In both cases we leverage time lapse microscopy and particle image velocimetry analysis (PIV) of co-cultures as a means to observe growth behavior and qualitative biophysical interactions over time. To examine how phenotypic changes associated with drug resistance impact upon tumor-stroma interactions, we correlate imaging data with mRNA sequencing to examine differential gene expression between drug naïve and drug resistant PDAC cells.

In this study we initially contrast two different co-culture geometries as shown in Fig. 1. In an embedded-fibroblast-with-overlaid-cancer-cells co-culture (EOC) model, the fibroblastic cells (either Pancreatic Stellate Cells, PSC, or MRC5, normal human fibroblasts) are suspended within a layer of ECM and the epithelial PDAC cells are overlaid above the ECM, allowing biochemical crosstalk without physical contact. We contrast this with an adjacent-overlay-co-culture (AOC) model, where we incubate 3D PANC1 cells for several days to form multicellular nodules, then introduce fibroblastic cells in the same plane on the surface of the ECM bed. This AOC model, which allows for direct physical contact and juxtacrine interactions, is the main focus of the present study. Time-lapse imaging and particle-image velocimetry analysis of cell co-migration and fibroblast contractility in AOCs, measured here as the speed of nodule aggregation, allow for quantitative interrogation of biophysical interactions between tumor and stromal cells for each experimental condition.

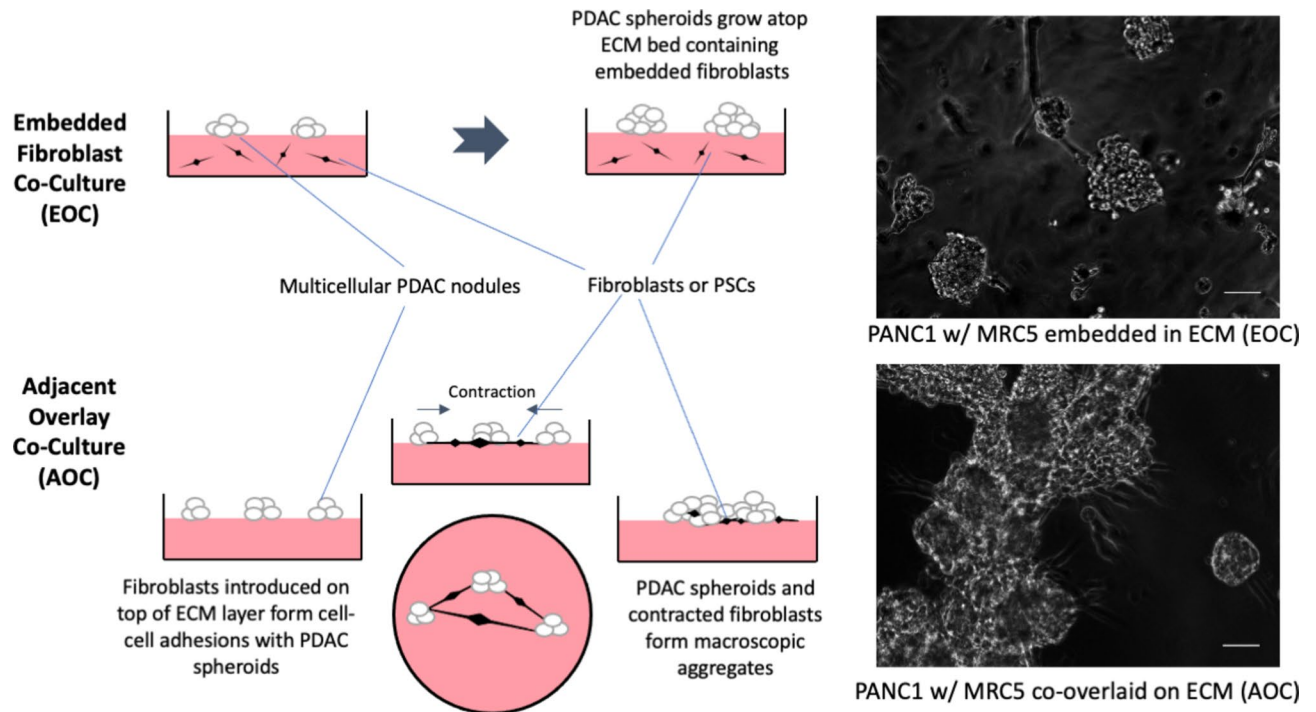
## Materials and methods

### Cell culture

PANC1 and MRC5 cell lines were acquired from ATCC and maintained in T75 cell culture flasks according to ATCC guidelines. DMEM and MEM (HyClone) were supplemented with 10% FBS (HyClone), 100 IU/mL penicillin, 1% streptomycin (HyClone), and 0.5 mg/mL amphotericin B (Corning). PSC cells were obtained from ScienCell. SteCM (Stellate cell media, ScienCell) was supplemented with 2% FBS, 1% SteCGS (Stellate cell growth supplement, ScienCell), and 1% penicillin. PSC and MRC5 cells were used for experiments between the 3rd and 10th passages. PANC1-OR is a stable oxaliplatin resistant sub-line of PANC1 developed and characterized previously<sup>17</sup>. PANC1-OR cells were maintained in T75 flasks according to ATCC guidelines for the PANC1 parent line.

### Analysis of mRNA-seq data

RNA was extracted from triplicate cell cultures of PANC1 and PANC1OR cells using the TRIzol reagent (Life Technologies Corporation, Carlsbad, CA). RNAseq was performed by the Center for Personalized Therapy Genomics Core at University of Massachusetts Boston. The quality of the raw fastq files were assessed using FastQC (v.0.11.5)<sup>24</sup>. Adaptor sequences, “AGATCGGAAGAGCACACGTCTGAACTCCAGTCA”, and “AGATCGGAAGAGCGTCGTGTAGGGAAAGAGTGT” were trimmed from the 3’ end of the reads using Cutadapt commandline tool from the Trime Galore package (v.0.4.2). The trimmed reads were mapped against the human reference genome (Ensemble, GRCh38) using STAR/2.5 using default parameters<sup>25</sup>. The average alignment rate was 96%. The sorted BAM files generated by STAR were used to estimate the transcript abundance per sample using featureCount from the Subread package (v.1.6.2)<sup>26</sup>. Gene expression analysis was performed using the edgeR Bioconductor R package (v.3.24.3)<sup>27</sup>. The edgeR TMM method (trimmed mean of M values) was applied to the filtered genes utilizing the DGElist(), calcNormFactors(), estimateGLMCommonDisp(), estimateGLMTrendedDisp(), estimateGLMTagwiseDisp() functions. The glmFit and glmLRT functions from edgeR were used to fit



**Fig. 1.** Overlay and adjacent tumor-fibroblast co-culture geometries. Left: Schematic diagrams of overlay and embedded co-culturing methods. In the embedded fibroblast with overlaid spheroid co-culture (EOC) fibroblasts are embedded in the underlying ECM layer and cannot make physical contact with PDAC spheroids overlaid above. In the adjacent fibroblast-spheroid overlay co-culture (AOC) PDAC spheroids are grown on a Matrigel bed and fibroblasts are then introduced triggering contractile behaviour. Right:  $10\times$  phase contrast image data of each co-culture platform. The EOC image is taken after seven days of spheroid formation and the AOC image is taken 48 h after fibroblasts have been introduced. Scale bars =  $200\ \mu\text{m}$ .

a negative binomial generalized log-linear model to the read counts. The expression of the genes was ranked by logFoldChange (logFC) and false discovery rate (FDR). Differentially expressed genes (DEGs) were determined using  $\text{abs}(\log\text{FC}) > 2$  and  $\text{FDR} < 0.01$  cutoffs, resulting in 1342 protein coding DEGs.

### Enrichment analysis

Gene ontology (GO) analysis was performed on differentially expressed genes using the g: Profiler with g:SCS (<https://biit.cs.ut.ee/gprofiler>)<sup>28</sup>. GO terms with significant overlap with up and down regulated genes were determined ( $\text{FDR} < 0.05$ ) and visualized with Heatmaps using the ComplexHeatmap R package (v.1.20.0)<sup>29</sup>. To run the gene set enrichment analysis (GSEA), we used the fgsea Bioconductor R package (v.1.9.5)<sup>30</sup>.

### Spheroid preparation and co-culturing

Prior to plating cells, GFR Matrigel™ was thawed overnight at  $4\ ^\circ\text{C}$  and kept on ice until use. For the adjacent overlay co-cultures, 225  $\mu\text{l}$  of growth factor reduced (GFR) Matrigel™ was added to the center of each well of a pre-chilled black-walled 24-well plate (Ibidi USA inc.). After the addition of Matrigel™ the plate was agitated to ensure an even coat on the bottom of the well and then the plate was incubated at  $37\ ^\circ\text{C}$  for 20 min allowing the Matrigel™ to solidify. PANC-1 cells or PANC-1-OR cells were collected, and cell density was determined using an automated cell counter. Preparations of each line were made at a concentration of 7500 cells/mL of media. 1 mL of the PANC1 cell preparation was plated in each of six Matrigel™ coated wells of the prepared 24 well plate. This was repeated with the PANC-1-OR subline. The cells were allowed to incubate at  $37\ ^\circ\text{C}$  and 5%  $\text{CO}_2$  for 7 days. After 7 days of spheroid growth, MRC5 or PSC cell lines were collected and counted. The cells were pelleted and resuspended in DMEM at a concentration of  $1 \times 10^5$  cells/mL of media. Media was removed from each well containing tumor spheroids. A 1 mL volume of the prepared MRC5-media solution was added to each of three wells of PANC1 spheroids and three wells of PANC1-OR spheroids. One mL of the prepared PSC-media solution was added to each of three wells of PANC1 spheroids.

For the EOC model, MRC5 or PSC cells collected and suspended in Matrigel™ and 225  $\mu\text{l}$  of the fibroblast-Matrigel™ suspension were added to the center of each well. After the addition of Matrigel™ the plate was agitated to ensure an even coat on the bottom of the well and then placed in the incubator until the Matrigel™ set. PANC1 cells were collected, and cell density was determined using an automated cell counter. A preparation of PANC1 cells in media was made at a concentration of 7500 cells/mL of media. A 1 mL volume of the PANC1 cell-media preparation was plated in three MRC5-Matrigel™ suspension coated wells and three PSC-Matrigel™ suspension coated wells. The cells were allowed to incubate at  $37\ ^\circ\text{C}$  and 5%  $\text{CO}_2$  for 7 days.

## Imaging and image analysis

Time-lapse  $10\times$  phase contrast images and  $1.25\times$  bright field images were taken using an inverted time lapse microscope (EVOS, Thermo Fisher Scientific) at a rate of 10 min per frame for 4 days. Image data was collected from three independent cell platings containing AOC, EOC and homotypic culture conditions and processed with custom Matlab image processing routines to obtain spheroid size distribution histograms from sets of segmented images. In area-weighted histograms the distribution represents the fraction of total growth area occupied by nodules of each size bin. Calculation of  $p$  values for comparison of object sizes across culture conditions used the two-tailed student's  $t$ -test.

Velocimetric data was obtained using PIVlab v1.43, an open source PIV toolbox for Matlab<sup>31,32</sup>. For  $10\times$  image data pre-processing was done in PIVlab. CLAHE, high pass, intensity capping, and denoise filters were all enabled in order to optimize in-app segmentation for analysis. For  $1.25\times$  bright field image data, images were first segmented, and preprocessing was disabled in PIVlab. In both cases an  $800\times 800$  pixel region of interest was chosen in the middle of the well to eliminate anomalies at the edge of wells. Multi-pass PIV analysis of each experimental condition was completed in triplicate utilizing the fast Fourier transform (FFT) window deformation algorithm generating velocimetric data. Here pairs of images are analyzed in multiple passes. Interrogation windows within the region of interest were cross-correlated in the frequency domain to generate a the velocity vector map of object speed across the region of interest. Object speed for each of the three iterations of each experimental condition was then averaged across each time-matched frame providing a basis for comparing the relative changes in average speed under each experimental condition. Curve Fits were performed using Origin-Pro software. For statistical analysis of variation in velocimetry profiles between conditions, the mean velocity and standard error across replicates was computed for each timepoint and used to calculate chi-squared values, and ultimately overall  $p$ -value for each comparison between conditions based on chi-squared and the number of degrees of freedom.

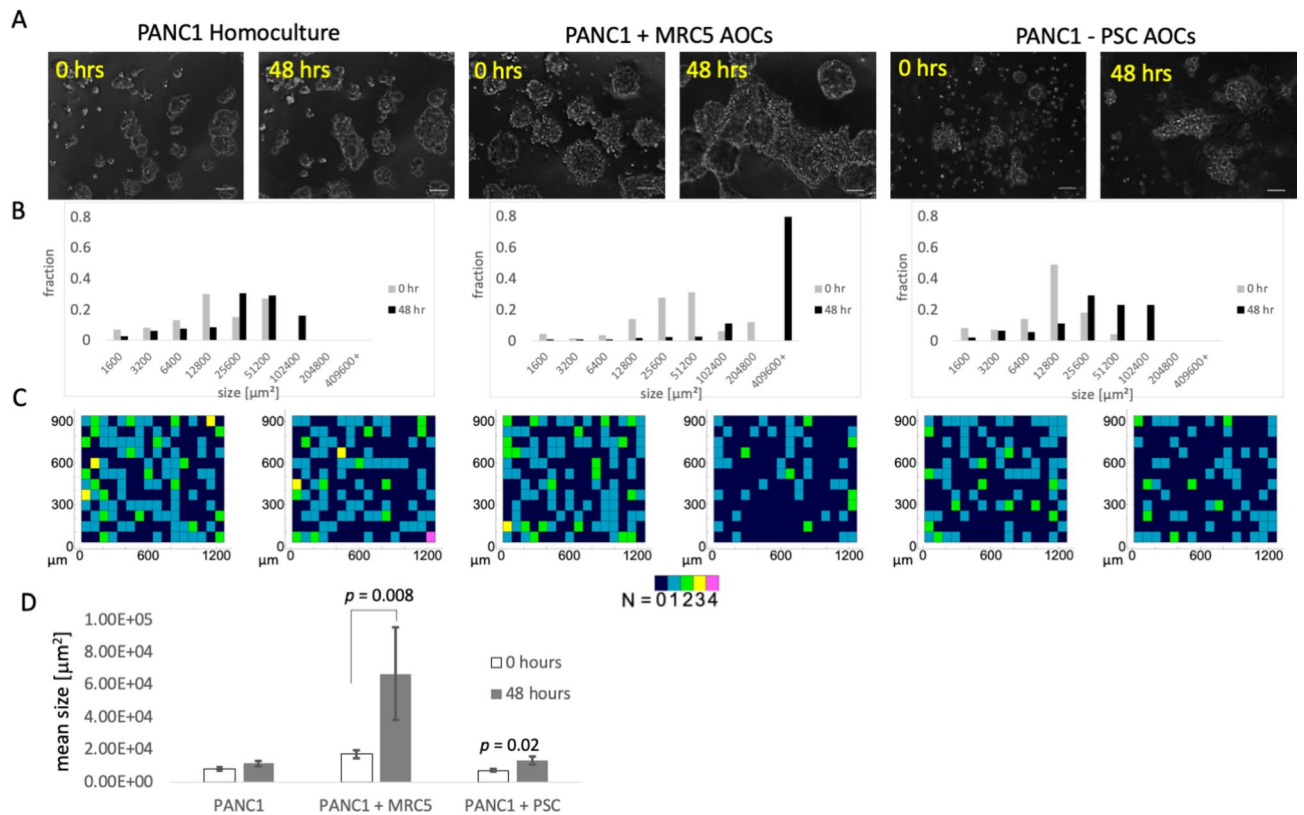
## Results and discussion

### Imaging-based analysis of PDAC-fibroblast co-culture growth, aggregation and contractile behavior

Co-cultures of adherent PDAC (PANC1) 3D nodules and fibroblastic cells (MRC5 or PSC) were initially carried out in two different geometries (Fig. 1): (1) an embedded fibroblast/PDAC overlay co-culture (EOC) in which the fibroblastic cells are embedded in ECM but cancer cells are overlaid on the ECM surface, and (2) an adjacent overlay co-culture (AOC) in which both 3D PDAC nodules are formed on the spheroid surface and fibroblast are also added in the same plane. It has been shown that intercellular interactions between tumor cells and stroma are mediated by a combination of paracrine signaling and biophysical juxtacrine interactions which involve direct physical contact between tumor and stromal cells. When fibroblasts are suspended within the ECM bed and PANC-1 spheroids are grown on top (EOC), only paracrine signaling is able to occur by secretion of growth factors and cytokines that can diffuse through ECM. In EOC cultures the PANC1 nodule growth behavior is qualitatively similar to homotypic 3D cultures which have been extensively characterized in previous reports<sup>33–35</sup>. Size distributions for PANC1 homotypic ECM overlay, PSC EOCs and MRC5 EOCs are shown in Supplemental Fig. 1.

When fibroblasts are introduced on top of the ECM bed upon which adherent PANC-1 spheroids have been grown, both paracrine and juxtacrine interactions may occur. In this scenario, PDAC and fibroblastic cells rapidly co-migrate and form heterotypic adhesions. Once attached, the fibroblasts contract and pull the largely microscopic tumor spheroids into a macroscopic mass of mingled stroma and cancer cells near the center of the culture vessel. The 2D spatial distributions of nodules atop ECM bed after 48 h in PANC-1 spheroid-fibroblast co-cultures show fewer and larger objects and with large swaths of empty ECM surface compared to PDAC spheroid homo-cultures. This is seen clearly in the PANC1-MRC5 AOC co-cultures where after 48 h wells are dominated by single very large co-mingled nodule (Supplemental Movie 1). This data suggests that, at least in this in vitro co-culture geometry, interactions between PANC1 and fibroblasts are mediated by juxtacrine communication. This qualitative change in growth behavior prompted further image analysis to quantify changes in spheroid/fibroblast co-migration and contractile motion using the AOC model, which is the major focus of this study.

Analysis of time lapse image data from AOC cultures reveals contrasting fibroblast contractility observed in co-cultures with MRC5 versus those with PSCs (Fig. 2). Here we observed that over the course of 48 h, PANC1 cells grown in homoculture remain distributed across the ECM layer with small changes in position, while nodule-area weighted histograms confirmed an expected shift in size due to normal cell proliferation (Fig. 2B). Distributions of nodule centroid positions similarly show there are fewer nodules but they remain distributed relatively uniformly across the well (Fig. 2B). On the other hand, in MRC5 AOCs aggregation occurred rapidly, resulting in a large mass of PANC1 3D nodules with enmeshed fibroblasts. Area-weighted histograms quantify aggregation with a dramatic shift towards few very large nodules over time accounting for most of the tissue coverage on the plate. PSC AOCs exhibit aggregation also, though less pronounced than the MRC5 co-cultures. Centroid distributions (Fig. 2C) present a similar case with MRC5 AOCs showing rapid convergence into a small number of aggregates leaving large areas of the ECM surface vacated. This contrast in contractility is consistent with previous characterization of co-cultures of these two cell types by us, suggesting more myofibroblastic CAF (MyCAF)-like phenotypic traits in MRC5 cells compared to the PSCs<sup>35</sup>. Mean size of aggregates (Fig. 2D) also increases over time, where again the change is dramatic in the PANC1 + MRC5 AOCs (increase in mean size of about 4 times in 24 h,  $p < 0.01$ ) and less so in the PANC1 + PSC AOCs (factor of 1.8 increase with modest significance of  $p = 0.02$  over the same time period). In PANC1 overlay homocultures the mean aggregate size increase over this 48 h period is not significant relative to the wide distribution of nodule sizes. Over longer

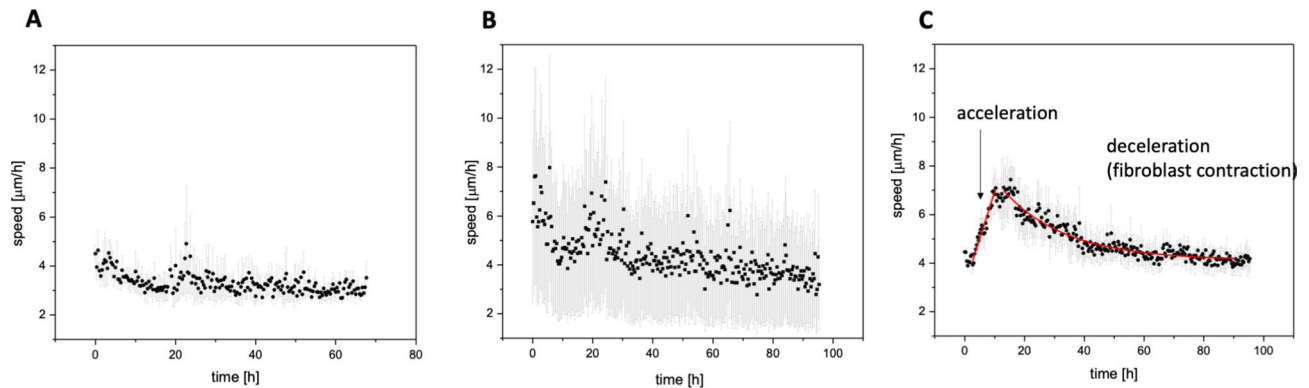


**Fig. 2.** Contrasting growth behavior of homocultures and co-cultures of PDAC cells with fibroblasts or pancreatic stellate cells. **(A)** Representative 10× phase contrast images from time lapse sequences of AOCs immediately (within ~30 min) after introduction of fibroblasts and 48 h later. Scale bars = 200 μm. **(B)** Total area weighted histograms of nodule size distribution before and after initiation of fibroblast contractility. MyCAF-like MRC5 AOCs show a dramatic shift towards few very large objects, while PANC1 grown in homoculture and iCAF-like PSC data show only a subtle shift towards larger objects. **(C)** Centroid distributions of nodules in each AOC at 0 and 48 h. After 48 h the MRC5 AOCs are spatially well separated on the ECM surface. **(D)** Mean size of PANC1 homocultures and co-cultures with MRC5 and PSCs at indicated time points. Error bars indicate standard error of the mean. In homocultures the aggregate size increases but the increase in mean is not significant relative to the breadth of the distribution ( $p > 0.05$ ). In the PANC1 + MRC5 AOCs co-cultures aggregation is significant with an increase in mean size of approximately a factor of 4 and  $p < 0.01$ . In PSC co-cultures there is a modest increase in size with mean increasing by a factor of 1.8 and  $p = 0.02$ .

times the MRC5 AOCs would eventually collapse into a single condensed aggregate visible in roughly the center of each well.

### PIV analysis of PDAC homocultures and fibroblast co-cultures

PIV analysis was conducted to further analyze speed and patterns of migration and contractile motion occurring in PANC1 AOCs and PANC1 homocultures (Fig. 3). This method involves quantification of shifts in object positions in segmented timelapse image sequences to generate a 2D map of velocity vectors at each time point (Supplemental Fig. 2). From these velocity maps we calculate average velocity overall space in each field as shown in Fig. 3, where the error bars on each time point represent standard error in time resolved velocity measurements over all replicate movie sequences. The velocity profile for PANC-1 overlay homocultures is essentially flat (Fig. 3A). In PSC AOCs (Fig. 3B) there is some evidence of reproducible acceleration and deceleration, while in the MRC5 AOCs (Fig. 3C) there is a well-defined and reproducible velocity profile that correlates with epithelial fibroblast co-migration, and fibroblast contraction events (Supplemental Fig. 3). In the first 12 h, a linear fit to the speed data shows acceleration of  $0.365 \pm 0.019 \mu\text{m}/\text{h}^2$  with adjusted R-squared of 0.94. The deceleration phase after 12 h, corresponding to ongoing fibroblast contraction fits to an exponential decay in speed of the form  $v = v_0 + Ae^{-t/\tau}$  with  $v_0 = 4.10 \pm 0.034 \mu\text{m}/\text{h}^2$ ,  $A = 5.05 \pm 0.27 \mu\text{m}/\text{h}^2$  and  $\tau = 22.3 \pm 1.3 \text{ h}$ , and adjusted R-squared of 0.85. The distinct velocity profile of AOC cultures contrasts with EOC experiments in which growth behavior of tumor nodules on the ECM surface that are not in direct contact with fibroblastic cells is qualitatively the same as in homocultures. Overall these experiments indicate that direct cell–cell juxtacrine interactions between fibroblasts and cancer cells are required for contractile behavior, and that these interactions are most pronounced in the fibroblastic cells shown previously to exhibit a more myofibroblast-like phenotype.



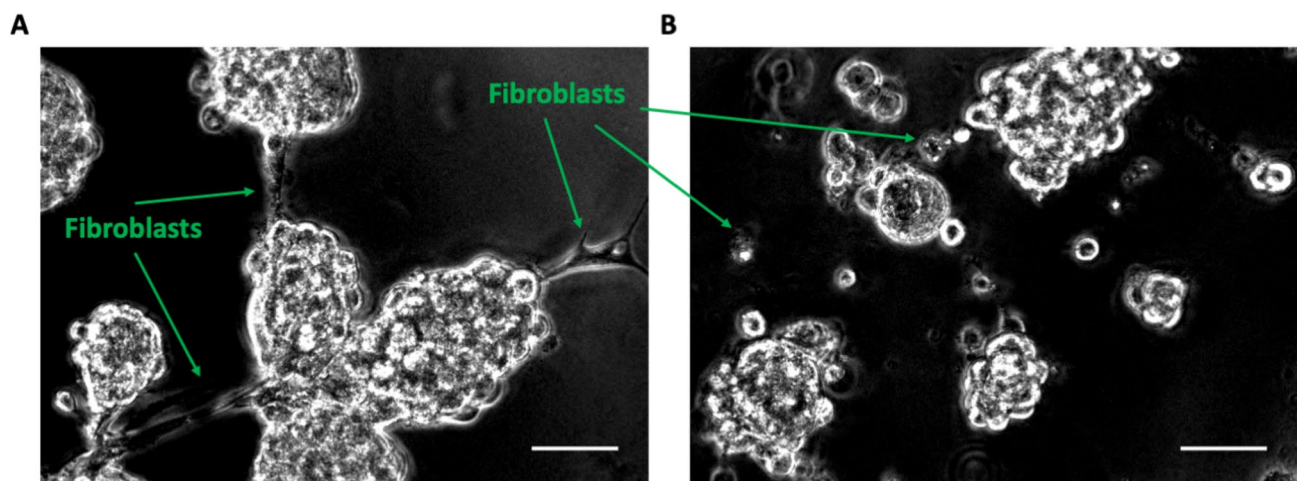
**Fig. 3.** Particle image velocimetry analysis. Time-dependent velocimetry plots for PANC1 homocultures (A), PANC1 + PSC AOC (B) and PANC1 + MRC 5 AOC (C). Each data point shows mean velocity at each time point from 3 experiments and error bars indicate standard error. In homocultures the velocity profile is relatively flat while time dependent acceleration is evident in the two cultures. In the MRC5 AOCs velocity changes map to co-migration and fibroblast contraction events with linear increase in speed in the first 12 h where in the first 12 h and exponential decay after 12 h (curve fits shown in red). The linear fit to the speed data shows initial acceleration of  $0.365 \pm 0.019 \mu\text{m}/\text{h}^2$  with adjusted R-squared of 0.94. The deceleration phase after 12 h, corresponds to ongoing fibroblast contraction fits to an exponential decay in speed of the form  $v = v_0 + Ae^{-t/\tau}$  with  $v_0 = 4.10 \pm 0.034 \mu\text{m}/\text{h}^2$ ,  $A = 5.05 \pm 0.27 \mu\text{m}/\text{h}^2$  and  $\tau = 22.3 \pm 1.3$  h, and adjusted R-squared of 0.85.

### Altered behavior of adjacent overlay cultures of chemoresistant PDAC cells

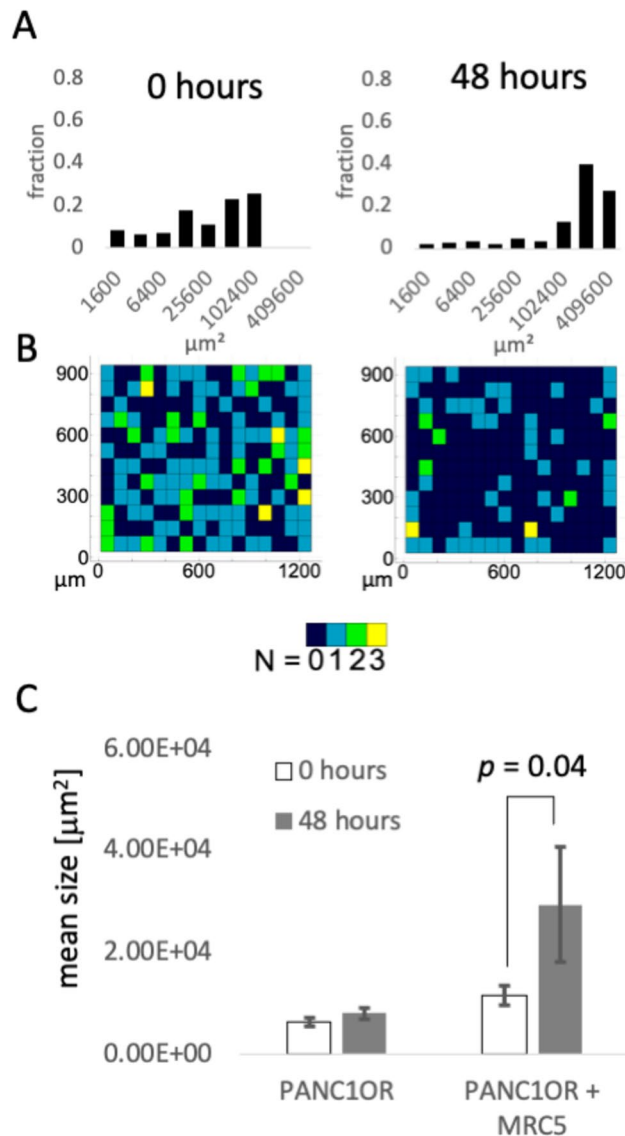
To evaluate how interactions between PDAC and stromal cells is altered due to acquisition of chemoresistance we evaluated co-cultures of a chemoresistant subline of PANC1, PANC1-OR with previously reported resistance to oxaliplatin and gemcitabine chemotherapy, and which exhibit increased mesenchymal phenotypic traits and invasive behavior<sup>22</sup>. Here, following the same imaging and PIV analysis protocols described above, we examine contrasting growth behavior in PANC1 and PANC1OR cells co-cultured in AOC geometry with MRC5 following from the conditions which produce the most notable heterotypic interactions of those characterized above.

When grown on ECM beds, drug resistant PANC1-OR cells do form multicellular 3D nodules, but when MRC5 cells are introduced, there is a marked difference in the interaction with stromal cells when compared with drug-naïve PANC1 cells (Fig. 4). Twelve hours after fibroblasts were introduced in PANC1 AOCs we observe fibroblasts stretched out and making physical contact with PANC1 nodules as they begin to merge into compact aggregates. (Fig. 4A). In contrast, in PANC1-OR AOCs there are few extended fibroblasts and very few direct adhesions between PDAC nodules and fibroblasts (Fig. 4B).

Visualizing quantitative aggregation data, area-weighted nodule size histograms shows a decrease in aggregation in PANC1-OR AOCs (Fig. 5) relative to PANC1 (non drug resistant) MRC5 AOCs shown in Fig. 2. The mean size of PANC1OR + MRC5 aggregates does increase over the 48 h time window documented in Fig. 5 but the extent of increase and its significance (mean size increases by a factor of 2.5, with  $p = 0.04$ ) is in stark contrast



**Fig. 4.** Representative images showing contrasting behavior of PANC1 versus PANC1-OR and MRC5 AOCs. (A)  $10\times$  phase contrast image of a PANC1 with MRC5 AOC 12 h after introduction of fibroblasts. Here fibroblasts make physical contact with spheroids leading to subsequent contraction. (B)  $10\times$  phase contrast image of a PANC1-OR with MRC5 AOC 12 h after addition of fibroblasts in overlay cultures.



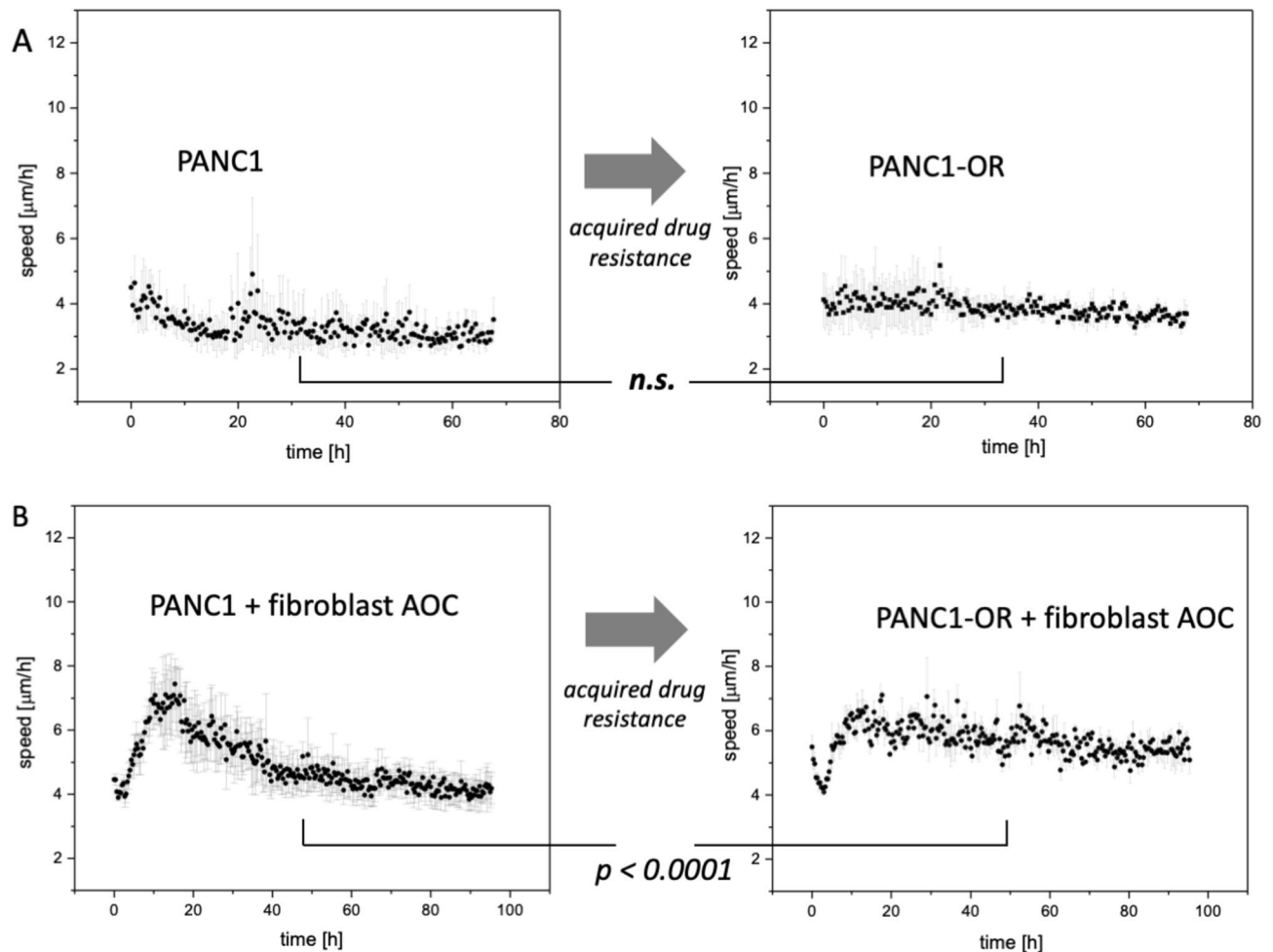
**Fig. 5.** Analysis of size distributions over time for PANC1-OR AOCs. **(A)** Area weighted AOC nodule size histograms at 0 and 48 h. PANC1-OR + MRC5 AOCs shift towards fewer and larger objects, though to a lesser extent than corresponding the PANC1 + MRC5 AOCs presented in Fig. 2. **(B)** Spatial distributions at 0 and 48 h after stromal cell overlay also show lower clustering relative to PANC1 + MRC5 AOCs shown in Fig. 2. **(C)** Mean size of PANC1OR and PANC1OR + MRC5 aggregates at indicated time points with error bars indicating standard error of the mean. In the PANC1 + OR co-cultures aggregation is evident though in contrast to the parental line the significance is borderline ( $p = 0.04$ ) as the distribution includes numerous individual PANC1OR nodules not adhered to stromal cells remaining after 48 h.

to the parental PANC1 cells in identical co-culture conditions shown in Fig. 2. As seen in Supplemental Movie 2, the formation of adhesions between epithelial and stromal cells is less frequent than in co-cultures of the parental line and unaggregated cell clusters that persist through the duration of the experiment.

Comparison of velocimetry profiles obtained from sequences of PANC1-OR and PANC1 co-cultures reveals marked change in behavior concomitant with acquisition of drug resistance (Fig. 6). While chi-squared analysis comparing PANC1 and PANC1OR homoculture velocimetry plots (Fig. 6A) shows no significant change (overall  $p$ -value of 0.41), the drug resistant co-culture velocity curve (Fig. 6B) changes significantly ( $p < 0.0001$ ). The linear acceleration and exponential deceleration phases characterized above for PANC1 + MRC5 AOCs are less well defined the co-culture velocity profiles, which are noisier, and with higher mean speed over the duration of the experiment. This analysis is consistent with the qualitative observations that drug resistant cells are more motile and less likely to form cell–cell adhesions.

#### Analysis of differential gene expression in chemoresistant versus chemosensitive PDAC cells

To examine changes in gene expression concomitant with acquisition of drug resistance that could account for suppression of adhesions to stromal cells observed above, we conducted mRNA sequencing and bioinformatics



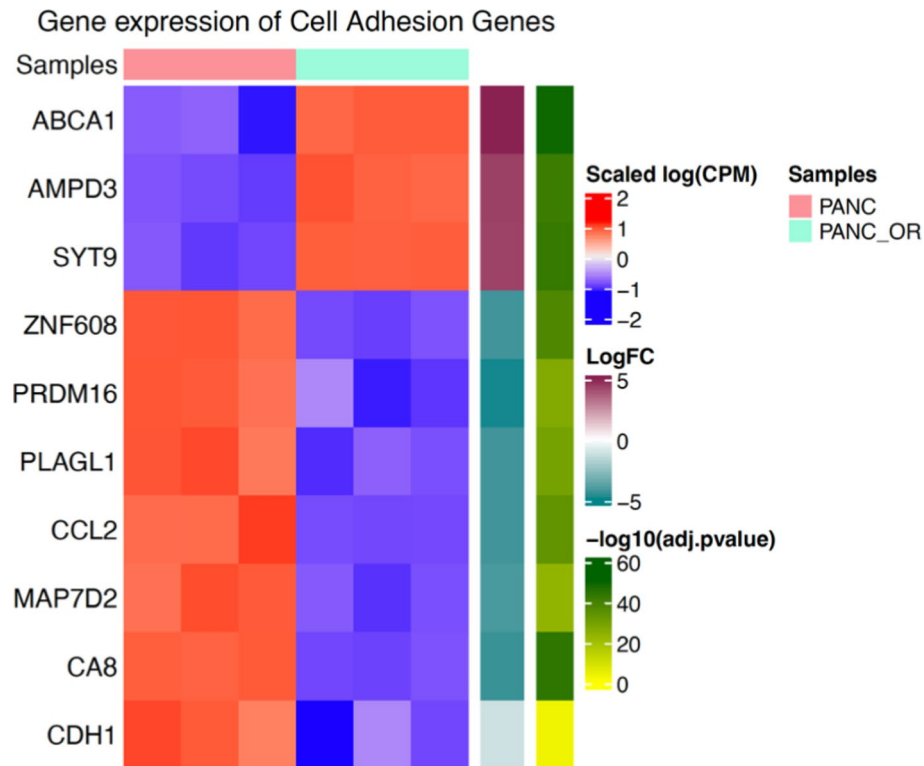
**Fig. 6.** Comparative velocimetry analysis for PANC1 versus PANC1-OR AOCs. **(A)** Comparison of velocimetry profiles for homocultures of PANC1 versus PANC1-OR. In both cases the overall time-dependent trend is flat with no significant change ( $p=0.41$  from analysis of sum of chi-squared weighted by standard error at each time point) with the acquisition of drug resistance. **(B)** Comparison of mean speed in PANC1 vs PANC1-OR AOCs shows a highly significant change. The well-defined acceleration phases of the PANC1 + MRC5 AOCs is no longer present when PANC1 are substituted with PANC1-OR. Chi-squared analysis shows the two curves differ significantly with  $p < 0.0001$ .

analysis to identify differentially expressed genes in PANC1-OR relative to PANC1. Extensive RNAseq analysis is shown in Supplemental Figs. 4 through 8 and Supplemental Tables 1 to 3. The gene ontology analysis for differential expression of gene families related to biological processes, molecular functions, and cellular components in Supplemental Figs. 3 through 5, respectively shows that PANC1-OR exhibit upregulation of genes related to cellular motility/migration, developmental and morphogenic processes, and decreased cell–cell adhesions. The overall pattern is generally consistent with increased EMT, though with some exceptions. The downregulation of CDH1 ( $\log_2(\text{FC}) = -1.2$ ,  $p = 0.001$ ), which codes for E-cadherin, is consistent with loss of epithelial characteristics. Other less obvious changes fit with this trend. For example, expression of KLF8, which has been shown to promote EMT by upregulating vimentin expression and downregulating E-cadherin<sup>36</sup>, was increased in PANC1-OR ( $\log_2(\text{FC}) = +5.15$ ,  $p = 8.09 \times 10^{-18}$ ). On the other hand SNAI3 expression was decreased in PANC1-OR, and SNAI1 expression was not significantly changed. Focusing on differential expression of genes potentially relevant to the observations here we found broadly that there was significantly decreased expression of genes which code for cell adhesion molecules in PANC1-OR (Fig. 7). Immunofluorescence imaging of fixed cultures (Supplemental Fig. 9) also shows loss of E-cadherin in PANC1-OR. Also, while the vimentin gene was not differentially expressed, the immunofluorescence imaging shows a clear pattern of cell elongation with increased cytoskeletal vimentin in PANC1-OR. Further phenotypic analysis of these drug resistant cells has been reported previously<sup>22</sup>.

## Conclusion

In this study we use a set of image analysis tools to contrast growth behavior in adjacent overlay PDAC 3D co-cultures. The analysis performed in this study points to the important role of direct biophysical interaction requiring cell–cell contacts between epithelial and stromal cells in PDAC. Only when fibroblasts are free to physically attach to PDAC nodules (as in the AOC model), do we see strong evidence of direct cell–cell contacts





**Fig. 7.** Heat map showing differential expression of genes relevant to cell–cell adhesion as determined from mRNA sequencing of triplicate samples of PANC1 and PANC1-OR. All genes included were differentially expressed with  $p < 0.001$ . More comprehensive analysis of differential gene expression is shown in Supplemental Figs. 3 through 7 and Supplemental Tables 1 to 3.

which mediate fibroblast contractility. This growth behavior drives toward densely compressed aggregates of intermingled tumor and stromal cells. This system could be further leveraged to model desmoplastic PDAC tissues and study how compressive stress in fibrotic stroma impacts upon disease progression in a controlled laboratory model.

A major goal of this study was to examine how these biophysical interactions between epithelial and adjacent stromal cells become altered in drug resistant PDAC. We observe through imaging and image analysis that the physical contacts required for contractility and aggregation are suppressed in drug resistant as compared to drug naïve PDAC cells co-cultured with fibroblasts. This finding is supported by RNA sequencing and immunofluorescence imaging showing downregulation of genes required for cell–cell adhesions between tumor and stromal cells, which is consistent with increased EMT and invasive behavior in these cells. This is significant in view of evidence that these contractile interactions between epithelial PDAC and myofibroblastic cancer associated fibroblast subpopulations act to constrain invasive progression of disease<sup>37</sup>. On the other hand it is also important to note that heterophilic E-cadherin/N-cadherin adhesions between cancer cells and stromal fibroblasts have been shown to promote an invasion mechanism in which CAFs guide collective migration of epithelial cells with intact adherens junction<sup>38</sup>. However, the drug resistant cells studied here, which have almost no expression of E-cadherin at the gene or protein level, would not be able to participate in this collective migration. Taken together, this suggests that single cell EMT-mediated invasion could be enabled by loss of adhesion to myofibroblastic CAFs which could otherwise contribute to mechanical confinement through exertion of contractile compressive stresses as seen in co-cultures with the parent PDAC line. In other words, the acquisition of drug resistance, which itself contributes to poor outcomes in PDAC<sup>20</sup>, is found here to also contribute to inhibition of stromal interactions that could otherwise help constrain invasive PDAC progression.

In these imaging-based biophysical studies we found that MRC5 co-cultures were more characteristically myofibroblastic, producing profound contractile behavior and much greater shift in object size and clustering than PSC aggregates. MyCAF interactions are known to be mediated by juxtacrine signaling, and the differential response between MRC5 and PSC AOCs suggests fibroblast contractility, and that the intercellular forces induced by stromal crosstalk, are mediated by juxtacrine interactions within the PDAC tumor microenvironment.

The macroscopic AOC aggregates cultured in this study may also be a useful tool to recapitulate desmoplasia characteristic of human PDAC in a versatile *in vitro* experimental platform. From the time evolution of contractile behavior we can glean insight into the mechanics of these tumor fibroblast interactions. Also, as a 3D culture methodology, the development of these highly fibrotic intermingled tumor and stroma tissue models could be used to study drug delivery in desmoplastic tumors<sup>39</sup>, or detached from ECM beds for xenograft implantation or further downstream analysis.

## Data availability

The datasets used and/or analysed during the current study available from the corresponding author on reasonable request.

Received: 3 October 2023; Accepted: 27 August 2024

Published online: 05 September 2024

## References

- Kamisawa, T., Wood, L. D., Itoi, T. & Takaori, K. Pancreatic cancer. *Lancet* **388**, 73–85. [https://doi.org/10.1016/S0140-6736\(16\)00141-0](https://doi.org/10.1016/S0140-6736(16)00141-0) (2016).
- Hwang, R. F. *et al.* Cancer-associated stromal fibroblasts promote pancreatic tumor progression. *Cancer Res.* **68**, 918–926 (2008).
- Korc, M. Pancreatic cancer-associated stroma production. *Am. J. Surg.* **194**, S84–S86 (2007).
- Rucki, A. A. & Zheng, L. Pancreatic cancer stroma: Understanding biology leads to new therapeutic strategies. *World J. Gastroenterol.* **20**, 2237 (2014).
- Landry, B. D. *et al.* Tumor-stroma interactions differentially alter drug sensitivity based on the origin of stromal cells. *Mol. Syst. Biol.* **14**, 1–15 (2018).
- Celli, J. P. Stromal interactions as regulators of tumor growth and therapeutic response: A potential target for photodynamic therapy?. *Isr. J. Chem.* **52**, 757–766 (2012).
- Piersma, B., Hayward, M. K. & Weaver, V. M. Fibrosis and cancer: A strained relationship. *Biochim. Biophys. Acta Rev. Cancer* **1873**, 188356 (2020).
- Feig, C. *et al.* Targeting CXCL12 from FAP-expressing carcinoma-associated fibroblasts synergizes with anti-PD-L1 immunotherapy in pancreatic cancer. *Proc. Natl. Acad. Sci. U S A* **110**, 20212–20217 (2013).
- Özdemir, B. C. *et al.* Depletion of carcinoma-associated fibroblasts and fibrosis induces immunosuppression and accelerates pancreas cancer with reduced survival. *Cancer Cell* **25**, 719–734 (2014).
- Olive, K. P. *et al.* Inhibition of hedgehog signaling enhances delivery of chemotherapy in a mouse model of pancreatic cancer. *Science* **1979**(324), 1457–1461 (2009).
- Rhim, A. D. *et al.* Stromal elements act to restrain, rather than support, pancreatic ductal adenocarcinoma. *Cancer Cell* **25**, 735–747 (2014).
- Öhlund, D. *et al.* Distinct populations of inflammatory fibroblasts and myofibroblasts in pancreatic cancer. *J. Exp. Med.* **214**, 579–596 (2017).
- Biffi, G. & Tuveson, D. A. Diversity and biology of cancer-associated fibroblasts. *Physiol. Rev.* **101**, 147–176 (2021).
- Elyada, E. *et al.* Cross-species single-cell analysis of pancreatic ductal adenocarcinoma reveals antigen-presenting cancer-associated fibroblasts. *Cancer Discov.* **9**, 1102–1123. <https://doi.org/10.1158/2159-8290.CD-19-0094> (2019).
- Zeisberg, M. & Neilson, E. G. Biomarkers for epithelial-mesenchymal transitions. *J. Clin. Investig.* <https://doi.org/10.1172/JCI36183> (2009).
- Horioka, K. *et al.* Suppression of CD51 in pancreatic stellate cells inhibits tumor growth by reducing stroma and altering tumor-stromal interaction in pancreatic cancer. *Int. J. Oncol.* **48**, 1499–1508 (2016).
- Sommariva, M. & Gagliano, N. E-Cadherin in pancreatic ductal adenocarcinoma: A multifaceted actor during EMT. *Cells* <https://doi.org/10.3390/cells9041040> (2020).
- Vasan, N., Baselga, J. & Hyman, D. M. A view on drug resistance in cancer. *Nature* **575**, 299–309 (2019).
- Shibue, T. & Weinberg, R. A. EMT, CSCs, and drug resistance: The mechanistic link and clinical implications. *Nat. Rev. Clin. Oncol.* **14**, 611–629 (2017).
- Arumugam, T. *et al.* Epithelial to mesenchymal transition contributes to drug resistance in pancreatic cancer. *Cancer Res.* **69**, 5820–5828 (2009).
- Dongre, A. & Weinberg, R. A. New insights into the mechanisms of epithelial-mesenchymal transition and implications for cancer. *Nat. Rev. Mol. Cell Biol.* **20**, 69–84 (2019).
- Cramer, G. M., Jones, D. P., El-Hamidi, H. & Celli, J. P. ECM composition and rheology regulate growth, motility, and response to photodynamic therapy in 3D models of pancreatic ductal adenocarcinoma. *Mol. Cancer Res.* **15**, 15–25 (2017).
- Kumar, N. *et al.* Stochastic modeling of phenotypic switching and chemoresistance in cancer cell populations. *Sci. Rep.* **9**, 10845 (2019).
- Andrews, S. FastQC: A Quality Control Tool for High Throughput Sequence Data. (2010).
- Dobin, A. *et al.* STAR: ultrafast universal RNA-seq aligner. *Bioinformatics* **29**, 15–21 (2013).
- Liao, Y., Smyth, G. & Shi, W. featureCounts: An efficient general purpose program for assigning sequence reads to genomic features. *Bioinformatics* **30**, 923–930 (2014).
- Robinson, M., McCarthy, D. & Smyth, G. edgeR: A Bioconductor package for differential expression analysis of digital gene expression data. *Bioinformatics* **26**, 139–140 (2010).
- Raudvere, U. *et al.* g:Profiler: A web server for functional enrichment analysis and conversions of gene lists (2019 update). *Nucleic Acids Res.* <https://doi.org/10.1093/nar/gkz369> (2019).
- Gu, Z. Complex heatmaps reveal patterns and correlations in multidimensional genomic data. *Bioinformatics* <https://doi.org/10.1093/bioinformatics/btw313> (2016).
- Korotkevich, G., Sukhov, V. & Sergushichev, A. Fast gene set enrichment analysis. *bioRxiv* <https://doi.org/10.1101/060012> (2019).
- Thielicke, W. & Stamhuis, E. J. PIVlab—Towards user-friendly, affordable and accurate digital particle image velocimetry in MATLAB. *J. Open Res. Softw.* **2**, 30 (2014).
- Thielicke, W. & Sonntag, R. Particle image velocimetry for MATLAB: Accuracy and enhanced algorithms in PIVlab. *J. Open Res. Softw.* **9**, 1–14 (2021).
- Glidden, M. D. *et al.* Image-based quantification of benzoporphyrin derivative uptake, localization, and photobleaching in 3D tumor models, for optimization of PDT parameters. *Theranostics* **2**, 827 (2012).
- Celli, J. P. *et al.* An imaging-based platform for high-content, quantitative evaluation of therapeutic response in 3D tumour models. *Sci. Rep.* **4**, 3751 (2014).
- Karimnia, V., Rizvi, I., Slack, F. J. & Celli, J. P. Photodestruction of stromal fibroblasts enhances tumor response to PDT in 3D pancreatic cancer coculture models. *Photochem Photobiol* **97**, 416–426 (2021).
- Yan, Q. *et al.* KLF8 promotes tumorigenesis, invasion and metastasis of colorectal cancer cells by transcriptional activation of FHL2. *Oncotarget* **6**, 25402–25417 (2015).
- Hingorani, S. R. Epithelial and stromal co-evolution and complicity in pancreatic cancer. *Nat. Rev. Cancer* **23**, 57–77 (2023).
- Labernadie, A. *et al.* A mechanically active heterotypic E-cadherin/N-cadherin adhesion enables fibroblasts to drive cancer cell invasion. *Nat. Cell Biol.* **19**, 224–237 (2017).
- Karimnia, V. *et al.* Photodynamic stromal depletion enhances therapeutic nanoparticle delivery in 3D pancreatic ductal adenocarcinoma tumor models. *Photochem. Photobiol.* **99**, 120–131 (2023).

## Acknowledgements

We gratefully acknowledge funding from NIH/NCI grants, R35CA232105 (FJS) and U54CA156734 (JPC and FJS), which supports the UMass Boston-Dana Farber/Harvard Cancer Center Partnership, and the University of Massachusetts Boston Healey Research Grant Program (JBD and JPC). We would also like to thank Fathi Mohamed for helping prepare samples for RNA sequencing and assistance from the Center for Personalized Cancer Therapy Genomics Core at UMass Boston.

## Author contributions

E.S. performed experiments and prepared the initial manuscript draft. J.P.C. guided experimental design and revised manuscript text. F.J.S. and J.B.D. guided experimental design and revised manuscript text. G.M.C. generated and characterized drug resistant cells. V.K. contributed to design of cell culture methods and data for Fig. 1. M.L. carried out bioinformatics analysis under supervision of K.Z. Y.L. assisted with bioinformatics analysis and writing. M.L. and K.Z. also revised the manuscript text.

## Competing interests

The authors declare no competing interests.

## Additional information

**Supplementary Information** The online version contains supplementary material available at <https://doi.org/10.1038/s41598-024-71372-9>.

**Correspondence** and requests for materials should be addressed to J.P.C.

**Reprints and permissions information** is available at [www.nature.com/reprints](http://www.nature.com/reprints).

**Publisher's note** Springer Nature remains neutral with regard to jurisdictional claims in published maps and institutional affiliations.

**Open Access** This article is licensed under a Creative Commons Attribution-NonCommercial-NoDerivatives 4.0 International License, which permits any non-commercial use, sharing, distribution and reproduction in any medium or format, as long as you give appropriate credit to the original author(s) and the source, provide a link to the Creative Commons licence, and indicate if you modified the licensed material. You do not have permission under this licence to share adapted material derived from this article or parts of it. The images or other third party material in this article are included in the article's Creative Commons licence, unless indicated otherwise in a credit line to the material. If material is not included in the article's Creative Commons licence and your intended use is not permitted by statutory regulation or exceeds the permitted use, you will need to obtain permission directly from the copyright holder. To view a copy of this licence, visit <http://creativecommons.org/licenses/by-nc-nd/4.0/>.

© The Author(s) 2024