

27 **Abstract**

28 The interaction between squamous cell carcinoma (SCC) cells and the tumor microenvironment
29 (TME) plays a major role in cancer progression. Therefore, understanding the TME is essential for
30 the development of cancer therapies.

31 We used four (primary and metastatic) head and neck (HN) SCC cell lines and cultured them on top
32 of or within 5 matrices (mouse sarcoma-derived_Matrigel®, rat collagen,
33 human leiomyoma-derived Myogel, human fibronectin, and human fibrin). We
34 performed several assays to study the effects of these matrices on the HNSCC behavior, such as
35 proliferation, migration, and invasion, as well as cell morphology, and molecular gene profile.
36 Carcinoma cells exhibited different growth patterns depending on the matrix. While fibrin
37 enhanced the proliferation of all the cell lines, collagen did not. The effects of the matrices on
38 cancer cell migration were cell line dependent. Carcinoma cells in Myogel-collagen invaded faster
39 in scratch wound invasion assay. On the other hand, in the spheroid invasion assay, three out of
40 four cell lines invaded faster in Myogel-fibrin. These matrices significantly affected hundreds of
41 genes and a number of pathways, but the effects were cell line dependent.

42 The matrix type played a major role in HNSCC cell phenotype. The effects of the ECMs were either
43 constant, or cell line dependent. Based on these results, we suggest to select the most suitable
44 matrix, which provides the closest condition to the *in vivo* TME, in order to get reliable results in *in*
45 *vitro* experiments.

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47 **Keywords:** Cancer, Extracellular Matrix, Invasion, Migration, Tumor Microenvironment.

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54 **Introduction**

55 Squamous cell carcinoma cells are notably affected by their microenvironment that mainly
56 includes extracellular matrix (ECM) and tumor stromal cells, such as cancer-associated fibroblasts
57 (CAF), immune and endothelial cells [\(1\)](#). ECM is a major component of the TME and it is composed
58 of a variety of proteins, proteoglycans, and polysaccharides [\(2\)](#). The structure and physical
59 properties of tumor-associated ECM differ from normal tissue stroma [\(3\)](#). Changes in the ECM
60 properties may cause variation in collagen deposition, promote the ECM stiffness, and upgrade
61 cell survival and proliferation [\(4,5\)](#). ECM could also affect tumor stroma cells, such as CAFs,
62 immune and endothelial cells [\(6\)](#). Therefore, understanding the SCC microenvironment is essential
63 for the development of cancer therapies, which targets not only the cancer cells but also their
64 environment that allows them to proliferate and spread.

65 *In vitro*, cancer cells are generally studied in 2D plastic wells. This usually leads to a loss of several
66 important elements, which could affect the cell behavior and phenotype, making the 2D system
67 not representative of the *in vivo* situation. In order to provide a more physiological environment
68 for the cells, culture systems using different ECM mimicking three-dimensional matrices were
69 introduced. Even though several matrices, which are extracted from different species such as
70 mouse, rat, bovine, or prepared from non-animal material [\(7\)](#), were proposed to be used in 3D cell
71 culture assays, selecting the most appropriate matrix for each cell type is not straightforward.

72 In spite of presence of several matrices from different origins, a human tumour-derived matrix is
73 still missing from the market. Our group has invented the first tumour-derived matrix “Myogel”
74 which is derived from leiomyoma tissue [\(8\)](#). Myogel has been used in several cancer *in vitro*
75 studies [\(1,9-12\)](#). Myogel proteome differs greatly from the commonly used mouse sarcoma-
76 derived Matrigel [\(8\)](#). We have shown recently that Myogel enhance the proliferation of freshly
77 isolated cancer cells from primary tumor compared to plastic and Matrigel [\(10\)](#). Additionally,
78 based on our recent publication, Myogel also improved the predictability of head and neck cancer
79 drug testing [\(12\)](#). This setup, applying Myogel coated wells in drug testing, could reduce the
80 number of failure clinical trials and reduce the cost of the anti-cancer drug development.

81 Here, we aimed to investigate the effects of several human- and animal-extracted ECMs, on the
82 head and neck (HN) SCC cells. We used mouse tumor-derived Matrigel[®], rat tail collagen,
83 human plasma fibronectin, human-derived fibrin, and human tumor-derived
84 Myogel; Bovine serum albumin (BSA) and uncoated wells were used as negative

85 | controls. We selected four HNSCC cell lines as a model of SCC cells: UT-SCC-24 (tongue) and UT-
86 | SCC-42 (larynx), including primary (A) and corresponding metastasis (B). We compared the effects
87 | of these matrices with the non-coated plastic analyzing cell morphology, proliferation, migration,
88 | and invasion. We also studied the effects of these matrices on the molecular profile of these cells
89 | using transcriptome profiling.

90

91 | **Materials and Methods**

92 | Cell lines

93 | UT-SCC cell line series, UT-SCC-24A (Primary tongue cancer, RRID:CVCL_7826), UT-SCC-24B
94 | (Metastatic tongue cancer, RRID:CVCL_7827) and UT-SCC-42 (larynx), including primary (A,
95 | RRID:CVCL_7847) and metastatic (B, RRID:CVCL_7848). Were kindly provided by Prof. Grenman
96 | (Department of Otolaryngology, Turku University, Turku, Finland). UT-SCC cells were grown in
97 | DMEM-F12 medium (Gibco™/Invitrogen, Tokyo, Japan) supplemented with 10% foetal bovine
98 | serum, 1% penicillin/streptomycin and 250µg/mL amphotericin B (all from Sigma-Aldrich, St. Louis,
99 | Mo, USA). All the cell lines were cultured in a humidified incubator (37°C, 5% CO₂, 95% humidity,
100 | Binder, Tuttlingen, Germany).

101 | Locally established four cell lines were isolated from two HNSCC patients, having both primary and
102 | metastatic tumors. Details of the cell lines are provided in supplementary Table 1.

103

104 | Preparation of the wells and light microscope imaging of cells morphology

105 | Ninety-six-well plates with black well walls and clear bottoms (Essen Bioscience, Ann Arbor, MI,
106 | USA) were used for coating. The plate was placed on ice and 50 µL/well of 0.5 mg/mL Matrigel and
107 | collagen (Corning, Corning, NY, USA) were dispensed using cold pipet tips. The plate was placed in
108 | the incubator for 30 minutes, then 50 µL/well of 0.01mg/mL BSA (Sigma-Aldrich), 0.01 mg/mL
109 | fibronectin (Sigma-Aldrich), 0.5 mg/mL Myogel (Lab made, see below):

119 and 0.5mg/mL fibrin
120 (For fibrin preparation, see below) was
122 added to the plate. The plate was incubated at the cell culture incubator for overnight. Cells were
123 detached from flasks using trypsin-EDTA and seeded at the density of 1000 cells/well in 100 μ L of
124 complete medium.

125 Myogel was prepared from human uterus leiomyoma tissue following the instructions in Salo et
126 al., 2015 (8). Briefly, tissue pieces were frozen using liquid nitrogen and crushed into a powder
127 with CryoMill (Retsch, Haan, Germany). A volume of 20 ml of ice cold NaCl buffer (3.4 M, pH 7.4)
128 was mixed with 10 g of tissue powder and centrifuged. 20 ml of the same NaCl buffer was used to
129 homogenize the pellet with a T18 Ultra-Turrax (IKA[®]-Werke GmbH & Co. KG, Staufen, Germany).
130 DC Protein Assay (Bio-Rad, Hercules, California, United States) was used to measure the protein
131 concentration in each preparation. The Myogel solution was stored in small (\leq 1 ml) aliquots at
132 -20°C .

133 Fibrin was prepared using 0.5 mg/mL fibrinogen (Merck, Darmstadt, Germany), 33,3 μ g/mL
134 aprotinin (Sigma-Aldrich) and 0,34 U/mL thrombin (Sigma-Aldrich).

135 To observe the effect of matrices on cell morphology, pictures of each well were taken after 3 days
136 using the reverse Nikon Digital sight DS-U3 microscope (Nikon, Tokyo, Japan) at x10 and x20
137 magnification.

138 139 Scanning electron microscope assay

140 We used two cell lines (UT-SCC-42A and UT-SCC-42B) to study the cell-matrix interaction under
141 scanning electron microscope. Glass coverslips were inserted into wells of a 24 well plate (Corning)
142 and coated using 300 μ L of coating suspension (using the same concentrations as above for each
143 matrix). Six thousand cells were seeded on each coverslip and incubated for 2 days. For fixation,

144 we performed several washing steps with 500 μ L of phosphate-buffered saline (PBS), and then we
145 added 500 μ L of 4% formaldehyde and kept it for 20 minutes at room temperature. After that, we
146 washed the wells with PBS again for 3 times, 5 minutes each. Samples were dehydrated using
147 graded ethanol series and dried using K850 critical point dryer (Quorum Technologies, UK). After
148 drying, samples were attached to aluminium stubs with double-sided carbon tape and coated with
149 5 nm of platinum using Q150T ES sputter coater (Quorum Technologies, UK). Samples were
150 imaged with Sigma HD-VP field-emission scanning electron microscope (Carl Zeiss, Oberkochen,
151 Germany).

152

153 Proliferation luminescent cell viability assay

154 For proliferation, we used the same experiment settings as in the imaging assay. After 3 days, the
155 plate was taken out from the incubator to room temperature for 15 min before starting the assay.
156 100 μ L of CellTiter-Glo was dispensed in each well. The plate was put on a plate
157 shaker (Heidolph, Schwabach, Germany) for 5 min at 450 rpm and then in plate spinner (Thermo
158 Scientific, Massachusetts, USA) for 5 min at 1000 rpm. Finally, the plate was placed in the BMG
159 PHERAstar FS (BMG Labtech, Offenburg, Germany) plate reader to detect cell viability.

160 Scratch wound cell migration assay

161 The same coating protocol was used as before, except that the gels were sucked out before
162 seeding the cells. We seeded the cells at the following density: 25000/well for UT-SCC-24A and UT-
163 SCC-42B and 30000/well for UT-SCC-24B and UT-SCC-42A. Matrigel was not used in this
164 experiment as the cells were forming clusters on top of the Matrigel leading to difficulties in
165 getting a smooth scratch. The wound maker (Essen Bioscience) was used to achieve homogeneous
166 scratch wounds.

167 Wounds were checked under the light microscope and the media was changed for all the wells.
168 The plate was placed in IncuCyte ZOOM incubator (Essen Bioscience), and wounds confluences
169 were monitored using the IncuCyte Live-Cell Imaging System (Essen Bioscience). Images were
170 taken each hour for 20 hours. Supplementary Video (Online Resource 1) shows the migration of
171 the UT-SCC-42A cells on top of Myogel.

172

173 Scratch wound cell invasion assay

174 Four gels were used in this experiment: collagen, Myogel-collagen, fibrin, and Myogel-fibrin at the
175 concentration of 1 mg/mL for all of them, as instructed by the manufacturer (Essen Bioscience).

176 Similar to the migration experiment, wells were coated, cells were seeded and a scratch wound
177 was made. After that, 50 μ L of the gels were added. Once the gels solidified, 50 μ L of media was
178 added. The plate was placed in IncuCyte ZOOM incubator, and wounds confluences were
179 monitored using the IncuCyte Live-Cell Imaging System (Essen Bioscience). Images were taken
180 each hour for 48 hours. Supplementary Video (Online Resource 2) shows the invasion of the UT-
181 SCC-42A within Myogel-collagen.

182

183 Spheroid Invasion Assay

184 Cells were seeded at the concentration of 1000 cells/well in 50 μ L using ultra-low attachment 96-
185 well round bottom plate wells (Corning). The plate was incubated for 4 days to allow spheroid
186 formation. Spheroids were embedded in 100 μ L of Matrigel (0.5 mg/mL), collagen (0.5 mg/mL),
187 Myogel-collagen (0.5 mg/mL), fibrin (0.5 mg/mL fibrinogen + 0.3 U/mL Thrombin + 3.33 mg/mL
188 Aprotinin), and Myogel-fibrin (0.5 mg/mL). Gels were allowed to solidify for 30 minutes and then
189 100 μ L of DMEM was added into each well. The plate was incubated for 4 days and pictures were
190 taken every day using Nikon Digital sight DS-U3 microscope (Nikon) at x4 magnification. The used
191 protocol is explained in detail in Naakka et al., 2019 (9)-. We analyzed the area covered by cells
192 using ilastik and ImageJ (Wayne Rasband, National Institute of Mental Health, Bethesda, MD,
193 USA). Once ilastik detected area covered by cells, we used a custom plugin, written for ImageJ, to
194 measure the area. The plugin converts the image to black and white image. All pixels outside the
195 area are set to zero, the cells area is set to one. The total area is measured as a number of pixels
196 equal to one.

197

198 Microarray

199 UT-SCC-24A and UT-SCC-24B cell lines were used to study the effects of different matrices on the
200 molecular gene profile using RNA sequencing transcriptome profiles. Wells of 24 well-plates were
201 coated with 150 μ L of gels (using the same concentrations as in the imaging assay) and seeded
202 with 150 000 cells. Cells were left on the gels for 24 hours and lysed using RLT buffer. RNA was

203 extracted using RNeasy Kit (Qiagen, Düsseldorf, Germany) according to manufacturer instructions.
204 In case some clots or fragments of gels existed in the cell lysate, sonication was used to solubilize
205 them. The quality of total RNA was assessed with a TapeStation (Agilent Technologies, Santa Clara,
206 CA, USA), and only samples of high quality (RNA integrity value >8) were included in the analyses.
207 The starting amount of total RNA was 100 ng. The labeling and hybridization were done according
208 to the manufacturer's instructions by using Applied Biosystems GeneChip™ WT PLUS Reagent Kit
209 and Manual Target Preparation for GeneChip™ Whole Transcript (WT) Expression Arrays
210 (UserGuide 23 January 2017; Thermo Fisher Scientific). Fifteen micrograms of cRNA were used for
211 single-stranded cDNA-synthesis (sscDNA) and a total of 5,5 ug of sscDNA was fragmented. A total
212 of 2.3 µg was hybridized on Clariom S Affymetrix array.

213

214 Gene set enrichment and pathway analysis

215 | Gene set enrichment analysis (GSEA, <http://software.broadinstitute.org/gsea/index.jsp>)₍₁₃₎ –was
216 carried out to connect gene expression signatures with previously known gene sets and pathways.
217 The analysis was performed for each cell line and matrix combination separately using the full
218 expression data set against C2: curated gene sets available at broad institute web page. Genes
219 were ranked using signal-to-noise ratio and gene set permutation was used for FDR estimation
220 and enrichment score adjustment. Additional analyses for Gene Ontology enrichment and KEGG
221 pathway visualization were carried out in R (v. 3.5.3) using packages gage (generally applicable
222 | gene set enrichment, v. 2.32.1)₍₁₄₎ -and pathview (v. 1.22.3)₍₁₅₎-. Both the GSEA and additional R
223 analyses were performed by the Functional Genomics Unit (FuGU) at the University of Helsinki.
224 Two samples (UT-SCC-24A/fibrin and UT-SCC-24B/fibrin) were excluded from the analysis as a
225 result of probability of mislabelling.

226

227 Analysis of cells circularity and their surface area

228 | Cells circularity and surface area were measured using ImageJ software (Wayne Rasband, National
229 Institute of Mental Health, Bethesda, MD, USA). The experiments were done four times
230 independently. Two wells were used for each condition and 3 cells were randomly selected and
231 measured from each well.

232

233 Statistical analysis

234 All experiments were repeated independently at least three times, each in duplicate or triplicate.
235 Values are given as means \pm standard deviations. All statistical analyses were performed using
236 SPSS (IBM SPSS Statistics for Windows, version 21.0; Armonk NY, IBM Corp.) To determine the
237 statistical significance, we performed one-way analysis of variance (ANOVA) followed by
238 Bonferroni correction. We set statistical significance to $p < 0.05$. P values were presented as
239 follows: * = $P \leq 0.05$, ** = $P \leq 0.01$, *** = $P \leq 0.001$, **** = $P \leq 0.0001$. Origin lab software was
240 used to create the figures.

241

242 **Results:**

243 SCC cells morphology is affected by the matrix type

244 Cancer cell morphology was affected by the type of matrix (Figure 1). While cells seeded on BSA
245 had similar morphology to the cells in the control wells, cells on Matrigel formed round clusters.
246 Cells on fibronectin had a more flattened surface than cells on the other matrices. Cells on Myogel,
247 fibrin, and collagen were more spindle in shape and there were fibers surrounding the cells. Here
248 we present the pictures of UT-SCC-24B cells only since the other cell lines behaved similarly (data
249 not shown).

250 For all the cell lines, cells on Matrigel had the ~~tendency of highest~~ highest circularity value (above 0.8 out
251 of 1) due to the formation of cell clusters, though circularity was close to 0.8 in many instances
252 with the other matrices as well (Figure 2). ~~The other matrices did not have any clear effect on cell~~
253 ~~circularity.~~

254 The majority of the cell lines, except UT-SCC-24A, seeded on top of fibronectin had higher surface
255 area than in the other conditions, but this difference did not reach statistical significance (Figure
256 3). UT-SCC-24A cultured on top of fibrin had significantly lower surface area compared to the cells
257 on plastic wells (Figure 3).

258

259 Cells-matrix interaction

260 Scanning electron microscope was used to observe the differences between structures of matrices
261 (Figure 4). While Matrigel has a fiber sheet structure, Myogel's structure was in form of thin

262 unorganized fibers together with small globular proteins. Fibrin has abundant thin fibers. Collagen
263 presented helical fibers structure. Fibronectin did not show a fibril structure.
264 The SEM pictures revealed the interaction between the cells and the matrices (Figure 5). Cells
265 cultured on BSA behaved similarly to the controls. On top of Matrigel, cells formed small clusters.
266 Cells on fibronectin tend to be flat with large surface areas. As for Myogel, cells were gathered in
267 groups and they were in contact with several fibers. For fibrin and collagen, cells were
268 embedded within the matrix fibers.

269
270 Fibrin increased and collagen reduced SCC proliferation, while matrix effect on cell migration was
271 cell line dependent.

272 The proliferation rate for all the tested cell lines was the highest on top of fibrin, and the lowest on
273 top of collagen (Figure 6). This difference was significant for the fibrin-coated wells in case of UT-
274 SCC-24A, 24B, and 42A, and for the collagen-coated wells in case of UT-SCC-24A and 42A.

275
276 The scratch wound cell migration assay showed that some matrices were able to affect cancer cell
277 migration but this effect was cell line-dependent (Figure 7). Opposite to the proliferation results,
278 collagen-induced UT-SCC-24B migration and fibrin reduced it. For UT-SCC-42A, cells cultured on
279 top of Myogel were migrating significantly slower compared to the control. Matrigel was not used
280 for migration assay since the cells formed clusters and a homogeneous wound was not possible to
281 be achieved.

282
283 Myogel induced SCC cell invasion

285 Cancer cells had different invasion speeds in the scratch wound invasion assay based on the matrix
286 used (Figure 8). Cells cultured within Myogel-collagen invaded the fastest. On the other hand, cells
287 did not invade through Myogel-fibrin and fibrin matrices. Myogel was able to induce cancer cells
288 invasion when added to the collagen, significantly in case of UT-SCC-42 A and B.

292 Figure 10). Myogel was able to significantly
293 induce cancer cell invasion when added to the fibrin in case of UT-SCC-24B and 42A (Figure 10).
304 Fibrin had the highest and BSA and fibronectin the lowest impact on SCC cell molecular profile
305 In order to understand the mechanism behind the effect of different matrices on the SCC cells
306 behavior, we studied the molecular gene profile using RNA sequencing transcriptome profiles.
307 Matrices were able to change the gene expression of hundreds of genes (Supplementary Table 2).
308 While cells seeded on fibrin had the largest difference (574 and 103 genes significantly affected,
309 $P \leq 0.05$, in UTSCC-24A and B, respectively), cells on BSA (15 and 19 genes significantly affected in
310 UTSCC-24A and B, respectively) and fibronectin (9 and 15 genes significantly affected in UTSCC-
311 24A and B, respectively) showed the least difference in their genes expression compared with the
312 cells cultured on plastic (Supplementary Table 2). The most significantly affected genes for each
313 matrix are presented in Supplementary Table 3.
314 Gene ontology enrichment analysis revealed several affected ontology groups (Supplementary
315 Table 4). These were both matrix and cell line dependent. The 10 most up- and downregulated
316 biological processes indicated by analysis are presented in Supplementary Table 5.

317 Due to the large variation between the two cell lines, we unfortunately were not able to detect
318 specific genes or ontology groups directly responsible for the changes in the SCC cells behavior.

319

320 Discussion

321 ECMs are increasingly used in cancer research to study different aspects of cancer cell behavior,
322 such as proliferation, migration, invasion and drug testing. The usage of these matrices was
323 regarded as a leap in moving ~~towards~~ closer to *in vivo* conditions than the traditional 2D cell
324 culture on plastic. This is mainly due to the ability of these matrices to provide essential elements
325 needed for the cell-cell and cell-matrix interaction. Due to the presence of several types of ECM, 326
327 such as Matrigel, Myogel, collagen, and fibrin, choosing the most suitable matrix that fits with the
328 needed assays without knowing its properties and effects could be risking the reliability of the
results ~~is difficult~~. Unfortunately, several researchers select the matrix type for their assay based
329 only on the availability, cost, and easiness of the matrix handling, without paying attention to the
330 effects of the matrix on ~~the~~ cancer cells behavior. Using a non-representative tumor matrix could
331 lead to non-reliable results. In this project, we pointed out the significant differences in SCC cells
332 behavior and their gene profile when tested with various matrices. This emphasizes the
333 importance of selecting the most suitable matrix for each assay.

334 We first studied the effect of ~~six~~ five matrices on the HNSCC cell morphology. Interestingly, all the
335 used cell lines formed cell round clusters when cultured on top of Matrigel, which is the most
336 common commercial extracellular matrix used in *in vitro* experiments. Our results are in line with
337 several other studies showing similar cell behavior on Matrigel in different cancer types_ -(16-18).
338 Forming cell clusters may be due to the presence of a large amount of basement membrane
339 proteins in Matrigel which seems to hold the cells together (19). Even though mimicking the
340 basement membrane is considered as an advantage for Matrigel, this feature is a disadvantage in
341 invasion assays due to the difficulties of cancer cells to invade through it. Opposite to the Matrigel,
342 cells cultured on top of Myogel, fibrin and collagen had a spindle shape, which represents more
343 the invasive phenotype of carcinoma cells, as reported in several publications_(16,20,21). This
344 morphology may represent an epithelial-mesenchymal transition (EMT), which is an important
345 feature for cancer cell migration and invasion_(22-24). Cells cultured on fibronectin had a unique
346 flattened shape with a large surface area. This shape could be explained by the presence of the

347 | $\alpha_5\beta_1$ integrin_(25) which is a fibronectin receptor (26), leading to an interaction that requires
348 | traction forces provided by the matrix.

349 | To confirm our visual observation of cell morphology, we measured the circularity and surface
350 | area of the cells. As expected, cells cultured on top of Matrigel had the highest circularity value
351 | due to the formation of round clusters. On the other hand, cells cultured on fibronectin had the
352 | highest surface area due to the flat shape of the cells.

353 | In order to get a better understanding of the cell-matrix interaction, we visualized the cells and the
354 | matrix under scanning electron microscope. As expected, most of the used matrices, except BSA
355 | and fibronectin, have fibril structures. The fibril structure of the matrices differed from one matrix
356 | to another in the terms of the amount of the fibers (rich vs poor) and thickness of the fibers (thick
357 | vs thin). All these differences, in addition to the presence or absence of several growth factors and
358 | other proteins, explain the differences in the behavior of cancer cells from one matrix to another.
359 | The interaction between the cells and the matrix was also different from one matrix to another.
360 | For some matrices, as in Myogel, the cells were surrounded by fibers, while for others, cells were
361 | either on top of the matrix (Matrigel) or embedded in it (fibrin and collagen).

362 | As cell viability assay is one of the main assays used in *in vitro* cancer research, we studied if the
363 | matrix itself could have an effect on cancer cell proliferation. Interestingly, one pattern was found
364 | in all the tested cell lines with the highest proliferation rate detected in the fibrin-coated wells and
365 | the lowest in the collagen wells. Our results are in line with Simpson-Haidaris *et al.* who reported
366 | similar results for breast cancer cells MCF-7 cultured on fibrin_(27). On the other hand, our results
367 | are opposite to Chen *et al.*, who reported a higher proliferation rate of MCF-7 cells when cultured
368 | on a porous collagen scaffold (28), suggesting that the effect of the collagen matrix is cell line
369 | dependent. Other matrices did not have a significant effect on HNSCC cell proliferation which goes
370 | hand by hand with some studies (29,30).

371 | Next, we studied the effect of the different matrices on cancer cell migration. Our results
372 | revealed that the effects of the studied matrices on HNSCC migration were cell line dependent,
373 | and the significant effects were assured by collagen, Myogel, and fibrin for some cell lines. It
374 | was an interest to us to notice the opposite effect of collagen and fibrin matrices on the
375 | proliferation and migration behavior of the UT-SCC-24B cell line. While these cells had the
376 | highest rate of proliferation on fibrin and the lowest on collagen, the opposite happened in cell

377 migration. This may return to the fact that the low proliferative cancer cells have high migration 378
379 capacity and *vice versa* (31).

380 Our scratch wound cell invasion assay showed that cells cultured within Myogel-collagen 3D
381 matrix invaded faster than cells within other matrices. This induction of invasion was mainly due
382 to Myogel since we also cultured HNSCC cells within collagen alone and the invasion speed was
383 lower. A similar effect of Myogel was observed on other cell lines (1). Cancer cells did not invade
384 through fibrin or Myogel-fibrin, which ~~might return~~ may be due to the fibrin's compact structure.

385 Similar to scratch wound cell invasion assay, Myogel was able to induce invasion in spheroids.
386 However, in scratch wound assay, Myogel-collagen was the most invasive inductive matrix in all 4
387 cell lines, while in spheroid 3 out of 4 cell lines invaded ~~the~~ fastest in Myogel-fibrin and one in
388 Myogel-collagen. This difference is most likely due to differences in the concentration of the gels
389 in the two assays (1 mg/ml in the scratch wound invasion and 0.5 mg/ml in the spheroid invasion
390 assays). Gels concentration were chosen either following the manufacturer instruction (scratch
391 wound invasion assay) or after lab optimization (spheroid invasion assay). Based on both invasion
392 assays, adding Myogel seems to improve the speed of HNSCC cancer cells invasion. This Myogel
393 property could help in studying low invasive cancer cell lines and testing anti-invasive cancer
394 treatments.

394 Matrigel has been the mostly used matrix for *in vitro* 3D cancer research. However, it should be
395 kept in mind that it is derived from mouse sarcoma containing mostly basement membrane
396 proteins (19). Due to its nature, in our invasion assays, cells failed to invade efficiently.

397 Based on our mRNA microarray results, the matrix type was able to significantly affect hundreds of
398 genes and several pathways. Interestingly, these genes and pathways were not shared between
399 matrices or cell lines but were matrix and cell line dependent. This was the reason that
400 unfortunately we were not able to detect specific gene or pathway responsible for
401 the changes in the SCC cells behavior. These results indicate that one cell line cannot represent the
402 behavior of any studied tumor type, and always more than one cell line should be used in *in vitro*
403 experiments.

404 Our study revealed important effects of the ECMs on HNSCC cells' behavior, morphology, and
405 molecular gene profile. We showed here that the ECMs are not idle elements, but instead, they
406 have significant effects on the *in vitro* results. We believe that for each assay, selecting the

407 appropriate matrix, based on its characteristics and the studied cell line, is necessary to get
408 reliable results in *in vitro* experiments. In theory, selecting human tumor-derived matrix could
409 represent the closest condition to the *in vivo* tumor microenvironment which increases the
410 reliability of the *in vitro* cancer cells testing.

411

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416

417 **References:**

418

419 (1) Salo T, Dourado MR, Sundquist E, Apu EH, Alahuhta I, Tuomainen K, et al. Organotypic three-
420 dimensional assays based on human leiomyoma–derived matrices. *Philosophical Transactions of*
421 *the Royal Society B: Biological Sciences* 2018;373(1737):20160482.

422 (2) Whittaker CA, Bergeron K, Whittle J, Brandhorst BP, Burke RD, Hynes RO. The echinoderm
423 adhesion molecule. *Developmental Biology* 2006;300(1):252-266.

424 (3) Provenzano PP, Eliceiri KW, Campbell JM, Inman DR, White JG, Keely PJ. Collagen
425 reorganization at the tumor-stromal interface facilitates local invasion. *BMC Medicine*
426 2006;4(1):38.

427 (4) Sundquist E, Renko O, Salo S, Magga J, Cervigne NK, Nyberg P, et al. Neoplastic extracellular
428 matrix environment promotes cancer invasion in vitro. *Experimental cell research*
429 2016;344(2):229-240.

430 (5) Lu P, Weaver VM, Werb Z. The extracellular matrix: A dynamic niche in cancer progression.
431 *Journal of Cell Biology* 2012;196(4):395-406.

432 (6) Quante M, Tu SP, Tomita H, Gonda T, Wang SSW, Takashi S, et al. Bone Marrow-Derived
433 Myofibroblasts Contribute to the Mesenchymal Stem Cell Niche and Promote Tumor Growth.
434 *Cancer Cell* 2011;19(2):257-272.

- 435 (7) Lou Y, Kanninen L, Kuisma T, Niklander J, Noon LA, Burks D, et al. The Use of Nanofibrillar
436 Cellulose Hydrogel As a Flexible Three-Dimensional Model to Culture Human Pluripotent Stem
437 Cells. *Stem Cells and Development* 2014;23(4):380-392.
- 438 (8) Salo T, Sutinen M, Hoque Apu E, Sundquist E, Cervigne NK, de Oliveira CE, et al. A novel human
439 leiomyoma tissue derived matrix for cell culture studies. *BMC Cancer* 2015;15(1):981.
- 440 (9) Naakka E, Tuomainen K, Wistrand H, Palkama M, Suleymanova I, Al-Samadi A, et al. Fully
441 Human Tumor-based Matrix in Three-dimensional Spheroid Invasion Assay. *Journal of Visualized
442 Experiments* 2019(147):e59567.
- 443 (10) Al-Samadi A, Poor B, Tuomainen K, Liu V, Hyytiäinen A, Suleymanova I, et al. In vitro
444 humanized 3D microfluidic chip for testing personalized immunotherapeutics for head and neck
445 cancer patients. *Experimental Cell Research* 2019;383(2):111508.
- 446 (11) Almahmoudi R, Salem A, Murshid S, Dourado RM, Apu HE, Salo T, et al. Interleukin-17F Has
447 Anti-Tumor Effects in Oral Tongue Cancer. *Cancers* 2019;11(5).
- 448 (12) Tuomainen K, Al-Samadi A, Potdar S, Turunen L, Turunen M, Karhemo P, et al. Human Tumor–
449 Derived Matrix Improves the Predictability of Head and Neck Cancer Drug Testing. *Cancers*
450 2019;12(1).
- 451 (13) Subramanian A, Tamayo P, Mootha VK, Mukherjee S, Ebert BL, Gillette MA, et al. Gene set
452 enrichment analysis: A knowledge-based approach for interpreting genome-wide expression
453 profiles. *Proceedings of the National Academy of Sciences USA* 2005;102(43):15545.
- 454 (14) Luo W, Friedman MS, Shedden K, Hankenson KD, Woolf PJ. GAGE: generally applicable gene
455 set enrichment for pathway analysis. *BMC Bioinformatics* 2009;10(1):161.
- 456 (15) Luo W, Brouwer C. Pathview: an R/Bioconductor package for pathway-based data integration
457 and visualization. *Bioinformatics* 2013;29(14):1830-1831.
- 458 (16) Truong D, Puleo J, Llave A, Mouneimne G, Kamm RD, Nikkhah M. Breast Cancer Cell Invasion
459 into a Three Dimensional Tumor-Stroma Microenvironment. *Scientific Reports* 2016;6(1):34094.
- 460 (17) Beers J, Gulbranson DR, George N, Siniscalchi LI, Jones J, Thomson JA, et al. Passaging and
461 colony expansion of human pluripotent stem cells by enzyme-free dissociation in chemically
462 defined culture conditions. *Nature Protocols* 2012;7(11):2029-2040.

- 463 (18) Polo ML, Arnoni MV, Riggio M, Wargon V, Lanari C, Novaro V. Responsiveness to PI3K and
464 MEK Inhibitors in Breast Cancer. Use of a 3D Culture System to Study Pathways Related to
465 Hormone Independence in Mice. PLOS ONE 2010;5(5):e10786.
- 466 (19) Zhang Y, Lukacova V, Reindl K, Balaz S. Quantitative characterization of binding of small
467 molecules to extracellular matrix. The Journal of Biochemical and Biophysical Methods
468 2006;67(2):107-122.
- 469 (20) Chen Y, Lan H, Wu Y, Yang W, Chiou A, Yang M. Epithelial-mesenchymal transition softens
470 head and neck cancer cells to facilitate migration in 3D environments. Journal of Cellular and
471 Molecular Medicine 2018;22(8):3837-3846.
- 472 (21) Hakkinen KM, Harunaga JS, Doyle AD, Yamada KM. Direct Comparisons of the Morphology,
473 Migration, Cell Adhesions, and Actin Cytoskeleton of Fibroblasts in Four Different Three-
474 Dimensional Extracellular Matrices. Tissue Engineering Part A 2011;17(5-6):713-724.
- 475 (22) Son H, Moon A. Epithelial-mesenchymal Transition and Cell Invasion. Toxicological research
476 2010;26(4):245-252.
- 477 (23) Kalluri R, Weinberg RA. The basics of epithelial-mesenchymal transition. Journal of Clinical
478 Investigation 2009;119(6):1420-1428.
- 479 (24) Zhou P, Li B, Liu F, Zhang M, Wang Q, Liu Y, et al. The epithelial to mesenchymal transition
480 (EMT) and cancer stem cells: implication for treatment resistance in pancreatic cancer. Molecular
481 Cancer 2017;16(1):52.
- 482 (25) Ahmedah HT, Patterson LH, Shnyder SD, Sheldrake HM. RGD-Binding Integrins in Head and
483 Neck Cancers. Cancers 2017;9(6):56.
- 484 (26) Wang K, Seo BR, Fischbach C, Gourdon D. Fibronectin Mechanobiology Regulates
485 Tumorigenesis. Cellular and molecular bioengineering 2016;9:1-11.
- 486 (27) Simpson-Haidaris P, Rybarczyk B. Tumors and Fibrinogen. Annals of the New York Academy of
487 Sciences 2001;936(1):406-425.
- 488 (28) Chen L, Xiao Z, Meng Y, Zhao Y, Han J, Su G, et al. The enhancement of cancer stem cell
489 properties of MCF-7 cells in 3D collagen scaffolds for modeling of cancer and anti-cancer drugs.
490 Biomaterials 2012;33(5):1437-1444.

491 (29) Hurst RE, Kyker KD, Bonner RB, Bowditch RD, Hemstreet, George P., 3rd. Matrix-dependent
492 plasticity of the malignant phenotype of bladder cancer cells. *Anticancer Research* 2003
493 Jul;23(4):3119-3128.

494 (30) Fliedner FP, Hansen AE, Jørgensen JT, Kjær A. The use of matrigel has no influence on tumor
495 development or PET imaging in FaDu human head and neck cancer xenografts. *BMC Medical*
496 *Imaging* 2016;16(1):5.

497 (31) Mouneimne G, Brugge JS. YB-1 Translational Control of Epithelial-Mesenchyme Transition.
498 *Cancer Cell* 2009;15(5):357-359.

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500

501 **Figure legends:**

502 **Figure 1. SCC cell morphology observed under light microscope.** Different shapes of UT-SCC-24B
503 cells were observed depending on the used matrix. Cells on BSA gave similar morphology to the
504 control. Cells were clustered on Matrigel, flat on fibronectin, and spindle on Myogel, fibrin, and
505 collagen. Scale bar = 100 μ m

506 **Figure 2. SCC cell circularity.** UT-SCC cells were cultured on different matrices and plastic
507 (control) for 3 days and pictured on day 3 under light microscope. Cell circularity was measured
508 using ImageJ software. In all cell lines represented in the figure, cells cultured on top of Matrigel
509 showed the highest circularity value (above 0.8). Data are presented as means \pm standard
510 deviations.* $P \leq 0.05$. N=3.

511 **Figure 3. SCC cell surface area.** UT-SCC cells were cultured on different matrices and plastic
512 (control) for 3 days and pictured on day 3 under light microscope. Cell surface area was measured
513 using ImageJ software. UT-SCC-24B, UT-SCC-42A and UT-SCC-42B cells showed the highest cell
514 surface area when cultured on Fibronectin, but this difference did not reach statistical significance.

515 Data are presented as means \pm standard deviations.* $P \leq 0.05$. N=3.

517 **Figure 4. Matrices structure observed under scanning electron microscope.** Coverslips were
518 coated with different matrices and prepared for scanning electron microscope. Matrigel has a
519 fiber sheet structure. Myogel's structure was in form of thin unorganized fibers together with

520 small globular proteins. As for fibrin, its fibers were thin. Collagen presented helical fibers
521 structure. BSA and fibronectin did not show a fibril structure. Scale bar = 1 μm

522 **Figure 5. SCC cells and matrix interaction observed under scanning electron microscope.** UT-SCC
523 42B cells were cultured on coated coverslips with different matrices and prepared for scanning
524 electron microscope. Cells cultured on BSA coated wells did not have any interaction with the
525 matrix, similarly to the cells cultured on plastic. For Matrigel, cells formed small clusters on top of
526 the matrix. Cells on fibronectin tend to be flat, more than any studied matrix, with a large surface
527 area. As for Myogel, cells were gathered in groups and they were in contact with several fibers. In
528 fibrin and collagen, cells were embedded within the matrix fibers. Scale bar = 10 μm

529 **Figure 6. SCC cell proliferation rate on different matrices.** UT-SCC cells were cultured on different
530 matrices for three days and the cell proliferation rate was measured using luminescent cell
531 viability assay. The proliferation rate for all the cell lines was the highest on fibrin and the lowest
532 on collagen. This difference was significant for the fibrin-coated wells in case of UT-SCC-24A, 24B,
533 and 42A cell lines and also for collagen in case of UT-SCC-24A and 42A cell lines. The red line
534 represents the control value. Data are presented as means \pm standard deviations.* $P \leq 0.05$, ** \leq
535 0.01, *** ≤ 0.001 , **** ≤ 0.0001 . N=3

536 **Figure 7. SCC cell migration on different matrices.** UT-SCC cells were cultured on different
537 matrices and cell migration was evaluated using scratch wound cell migration assay. The migration
538 rate was dependent on both the matrix and the cell line. Data are presented as migration curves
539 and area under the curves as means \pm standard deviations.* $P \leq 0.05$, ** ≤ 0.01 , *** ≤ 0.001 , ****
540 ≤ 0.0001 . N=3.

541 **Figure 8. SCC cells invasion through different matrices.** UT-SCC cells were cultured on different
542 matrices and cell invasion was evaluated using scratch wound cell invasion assay. The four studied
543 cell lines showed the fastest invasion rate when cultured on Myogel-collagen and they did not
544 invade through fibrin and Myogel-fibrin. Data are presented as invasion curves and area under the
545 curves as means \pm standard deviations.* $P \leq 0.05$, ** ≤ 0.01 , *** ≤ 0.001 , **** ≤ 0.0001 . N=3.

553 **Figure 9. Spheroid invasion observed under light microscope.** UT-SCC 42A cells were cultured in
554 ultra-low attachment 96-well round bottom plate wells and embedded in different matrices.
555 Spheroids were observed under light microscope. Scale bar = 100 μm .

556 **Figure 10. SCC spheroid invasion through different matrices.** UT-SCC cells were cultured in ultra-
557 low attachment 96-well round bottom plate wells and embedded in different matrices. For UT-
558 SCC-24A, 42A, and 42B, Myogel-fibrin matrix showed the fastest spheroids invasion, followed by
559 Myogel-collagen. For UT-SCC-24B cells invaded faster in Myogel-collagen followed by Myogel-
560 fibrin. Data are presented as invasion curves and area under the curves as means \pm standard

561 deviations. * ≤ 0.05 , *** ≤ 0.001 , **** ≤ 0.0001 . N=3.

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563

564 **List of Supporting Information:**

569 **Supplementary Table 1. HNSCC cell lines details.** Clinical and pathological characteristics of the
570 HNSCC cell lines. TNM is based on

571 **Supplementary Table 2. Number of differentially expressed genes of UT-SCC-24A and B cultured**
572 **on different matrices.** Results of mRNA microarray showing the number of differentially expressed
573 genes between cells cultured on plastic and cells cultured on matrices. The genes that passed the
574 filter criteria had a $p < 0.05$ and a fold change ≤ -2 or ≥ 2 . Transcriptome analysis console software
575 was used to analyze the data

576 **Supplementary Table 3. The most affected genes of UT-SCC cells cultured on different matrices.**
577 Results of mRNA microarray showing the most significantly affected genes (up- down-regulated)

578 by each matrix we used. The genes that passed the filter criteria had a $p < 0.05$ and a fold change \leq -
579 2 or ≥ 2 . Transcriptome analysis console software was used to analyze the data.

580 **Supplementary Table 4. Number of differentially expressed pathways of UT-SCC-24A and B**
581 **cultured on different matrices.** Results of the gene set enrichment analysis (GSEA) showing the
582 number of the differentially represented pathways between cells cultured on plastic and cells
583 cultured on matrices. The pathways that passed the filter criteria had a $p < 0.05$.

584 **Supplementary Table 5. The 10 most affected pathways of UT-SCC cells cultured on different**
585 **matrices.** Results of the gene set enrichment analysis (GSEA) showing the 10 most differentially
586 expressed pathways between cells cultured on plastic and cells cultured on matrices. The
587 pathways that passed the filter criteria had a $p < 0.05$.

588

589 **Supplementary video 1:** migration of the UT-SCC-42A cells on top of Myogel.

590 **Supplementary video 2:** the invasion of the UT-SCC-42A within Myogel-collagen.

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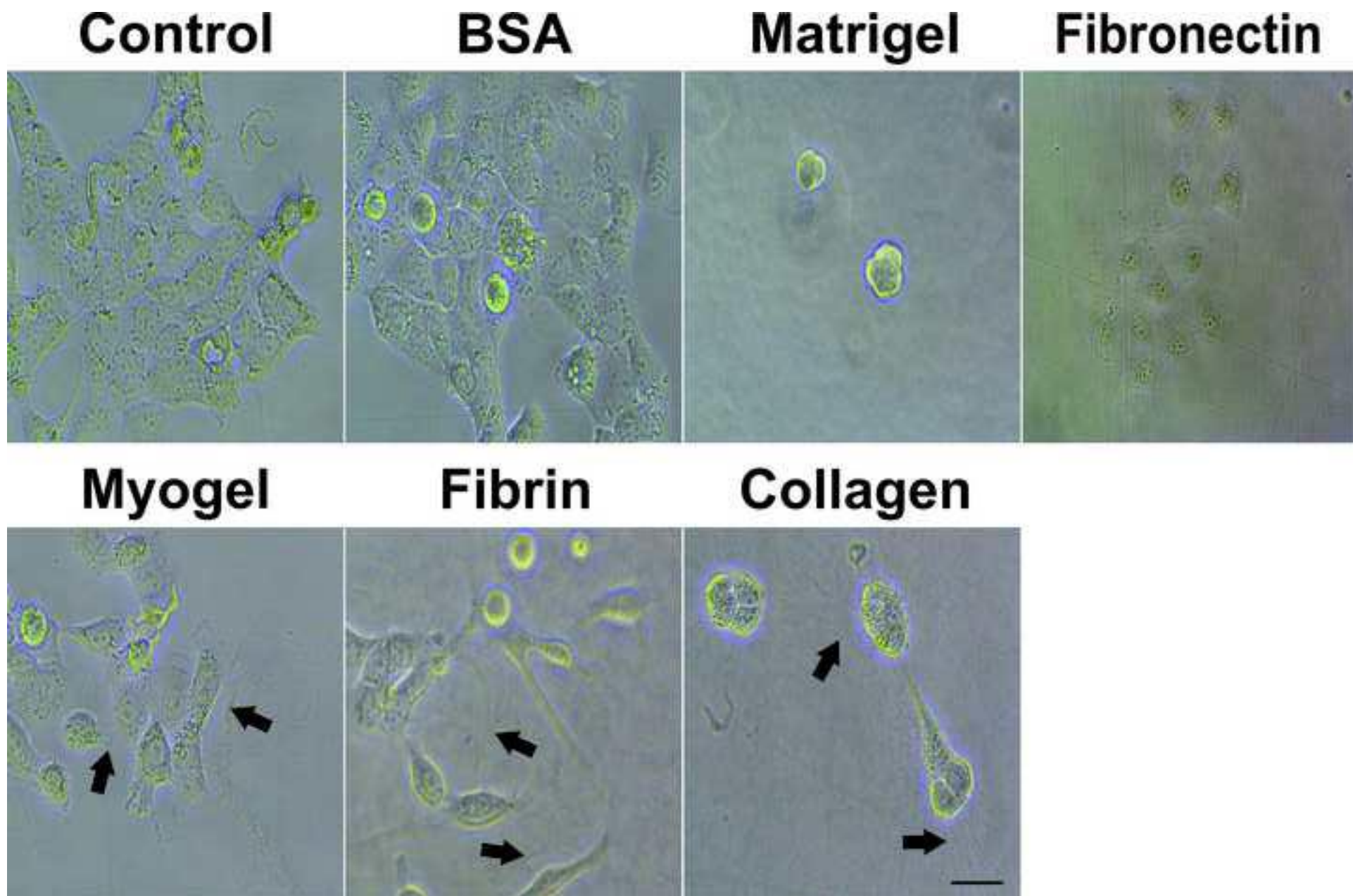


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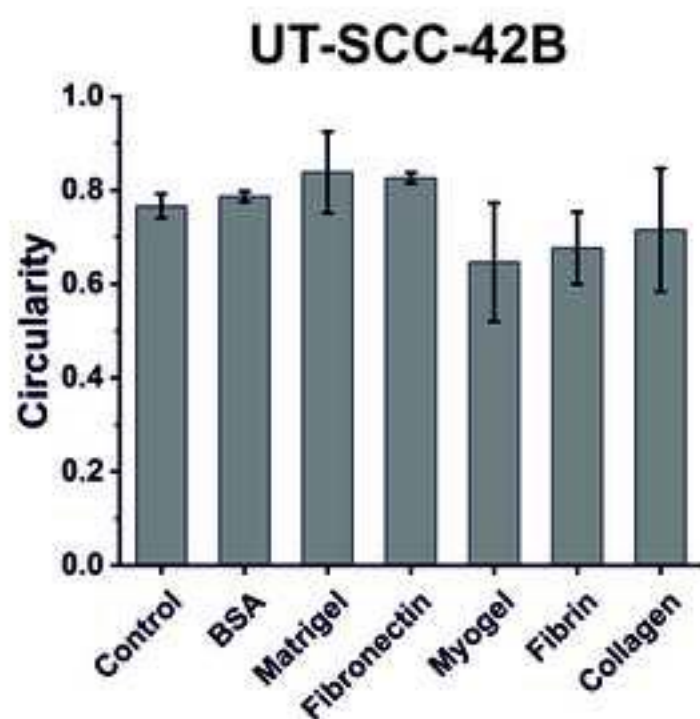
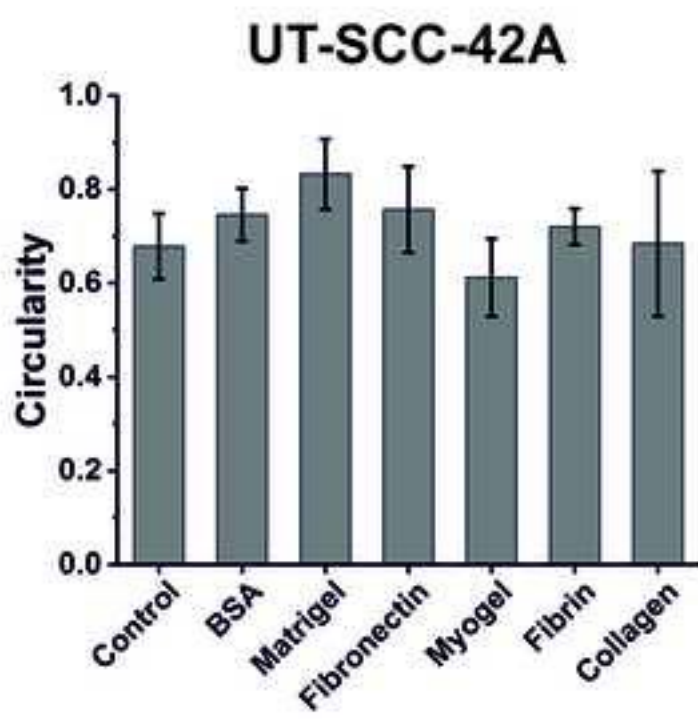
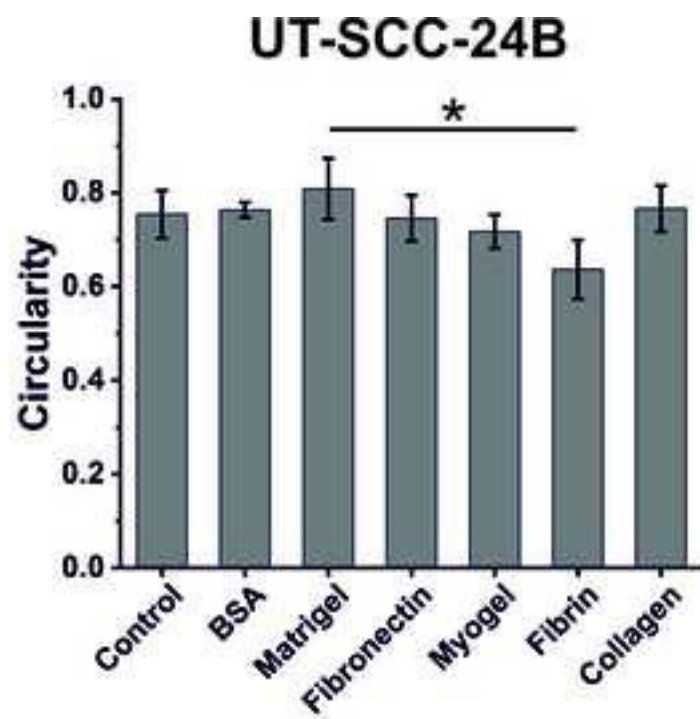
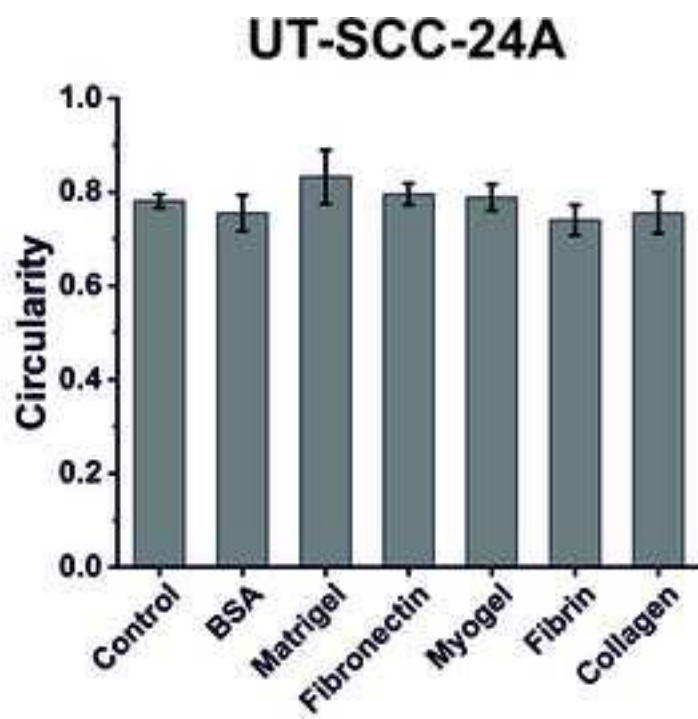
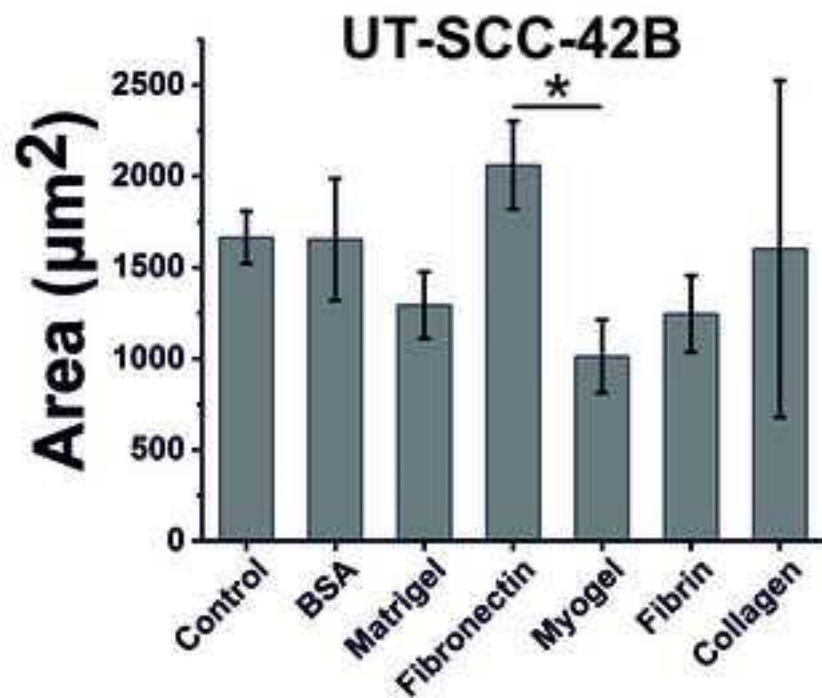
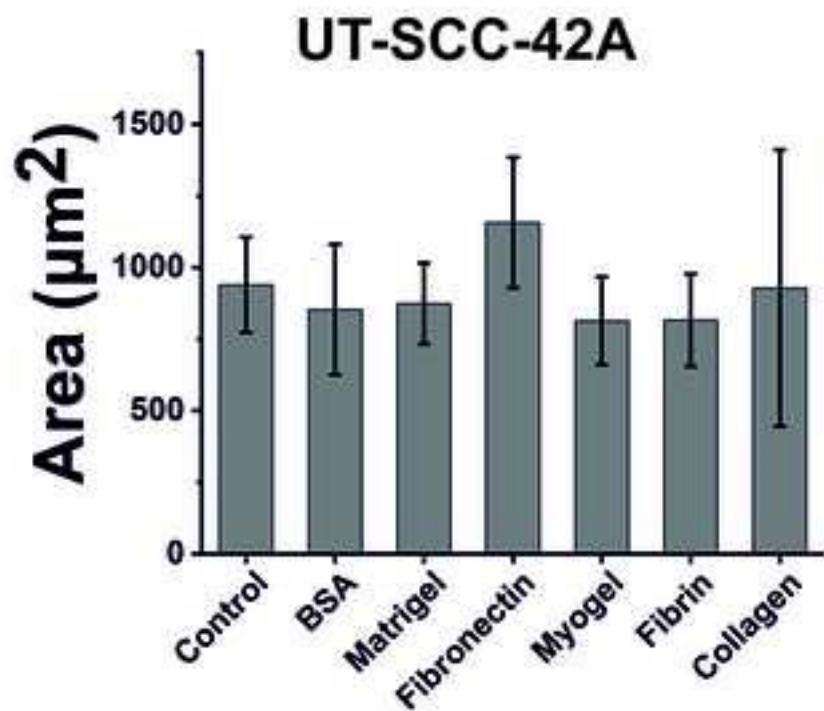
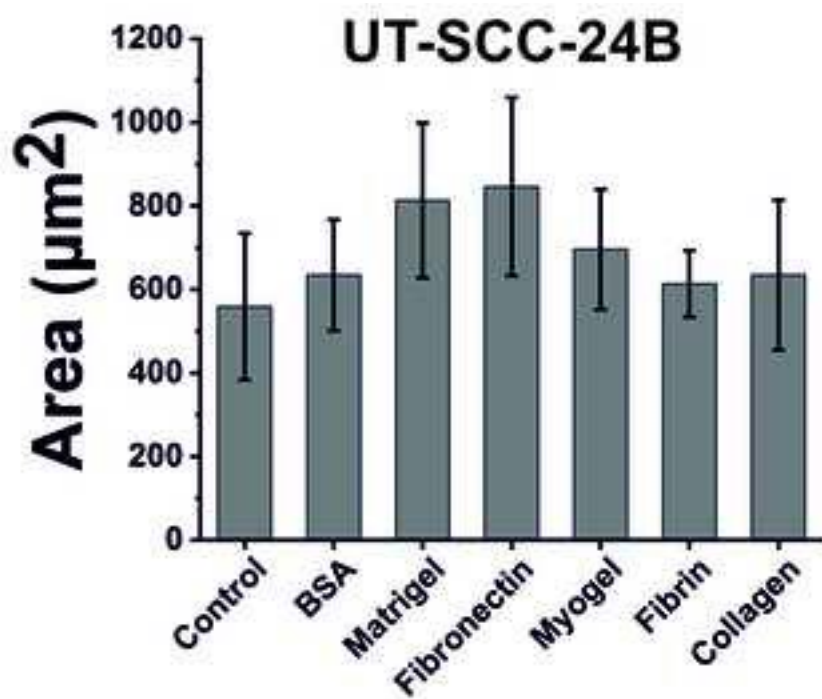
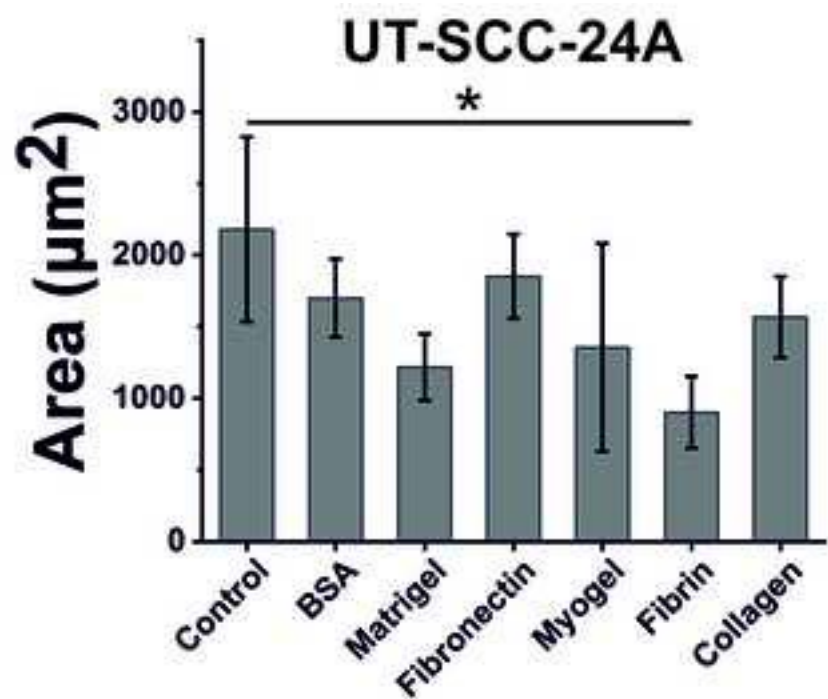


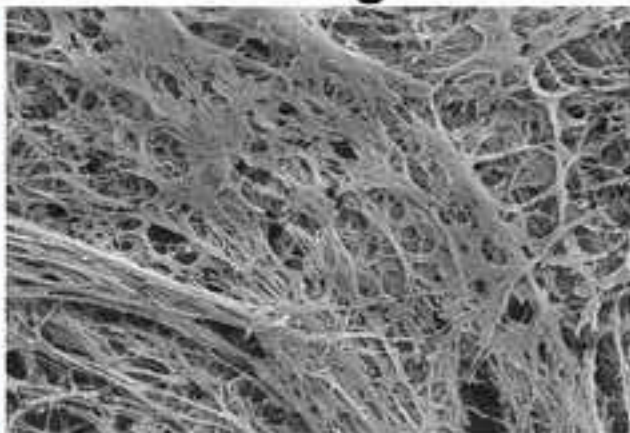
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BSA



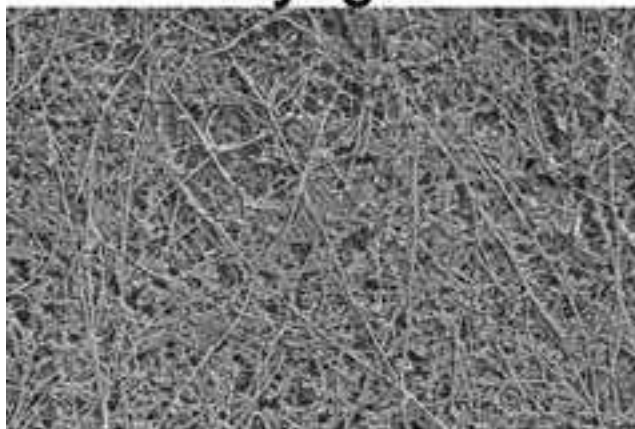
Matrigel



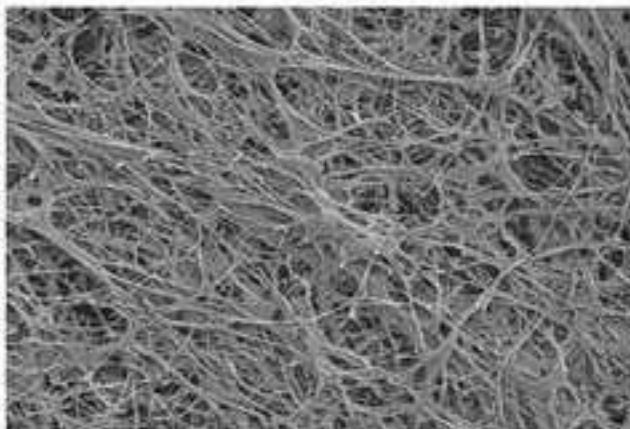
Fibronectin



Myogel



Fibrin



Collagen



Figure 5
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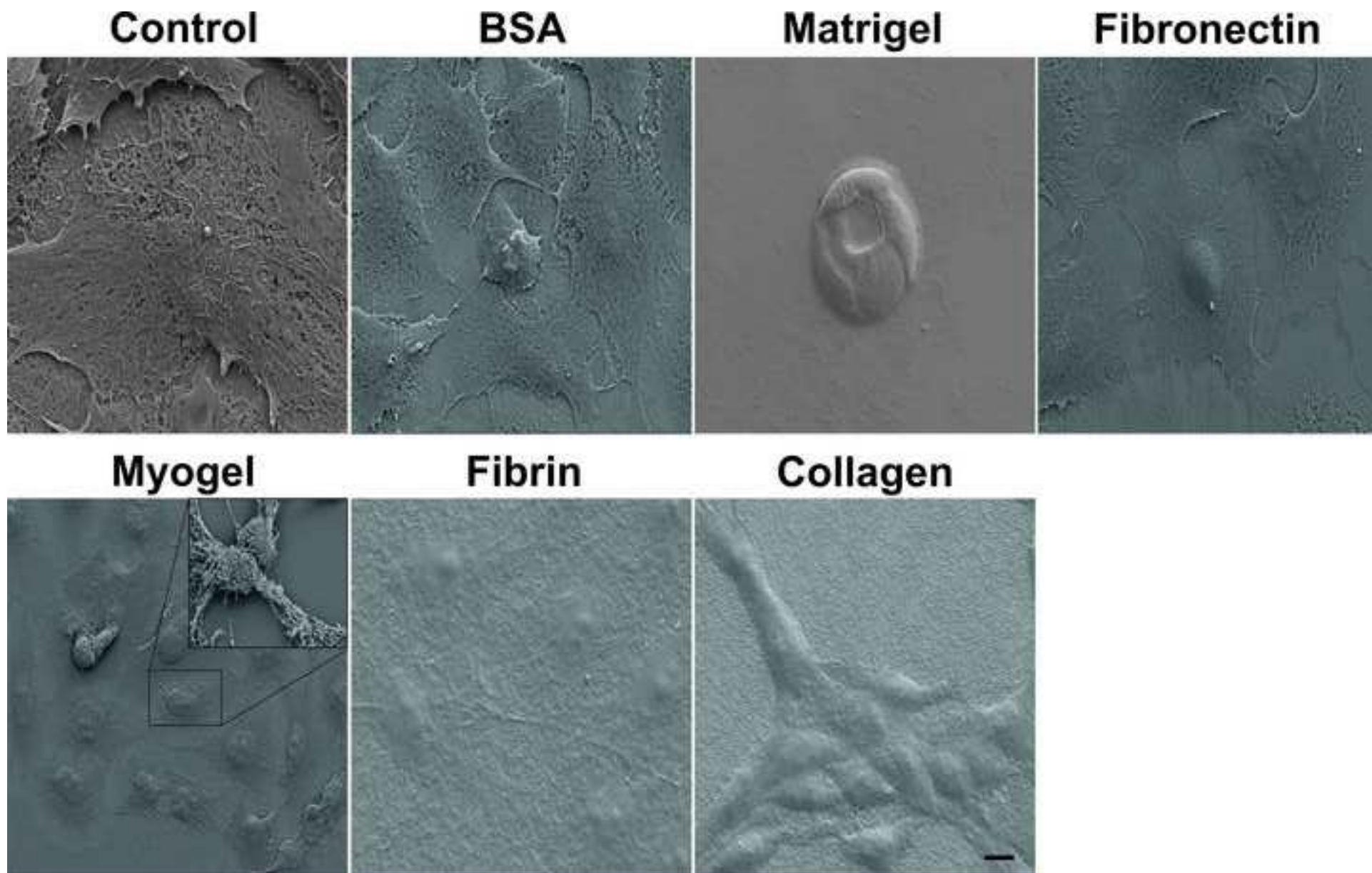


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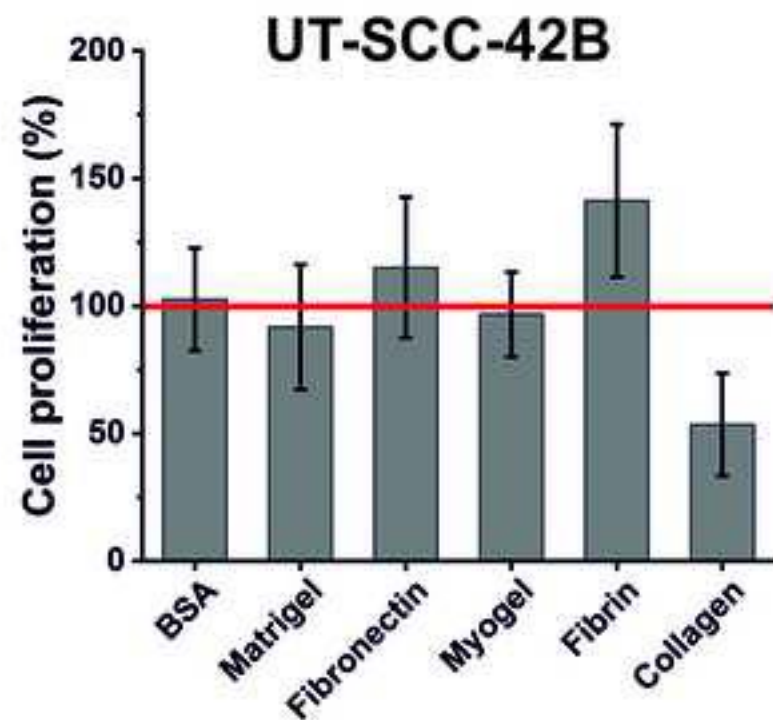
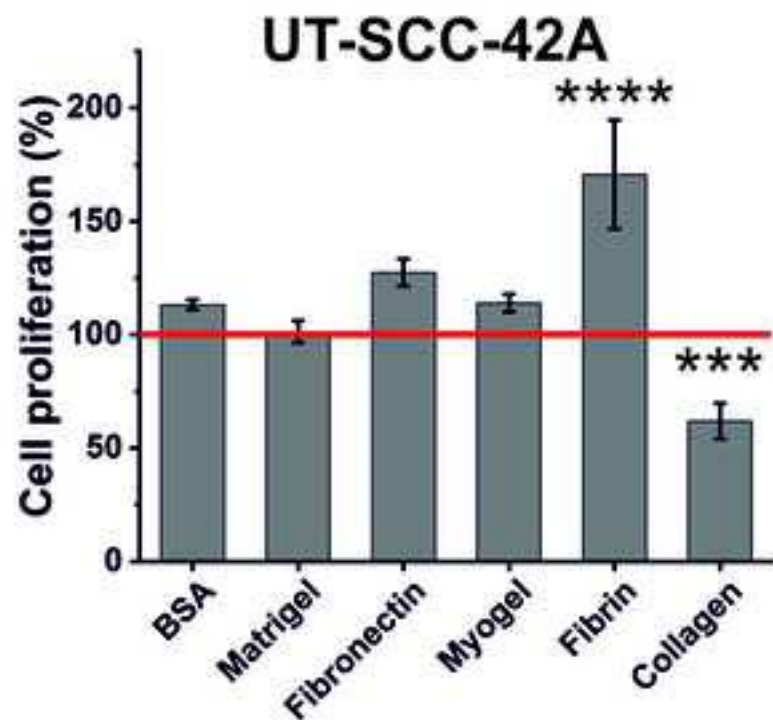
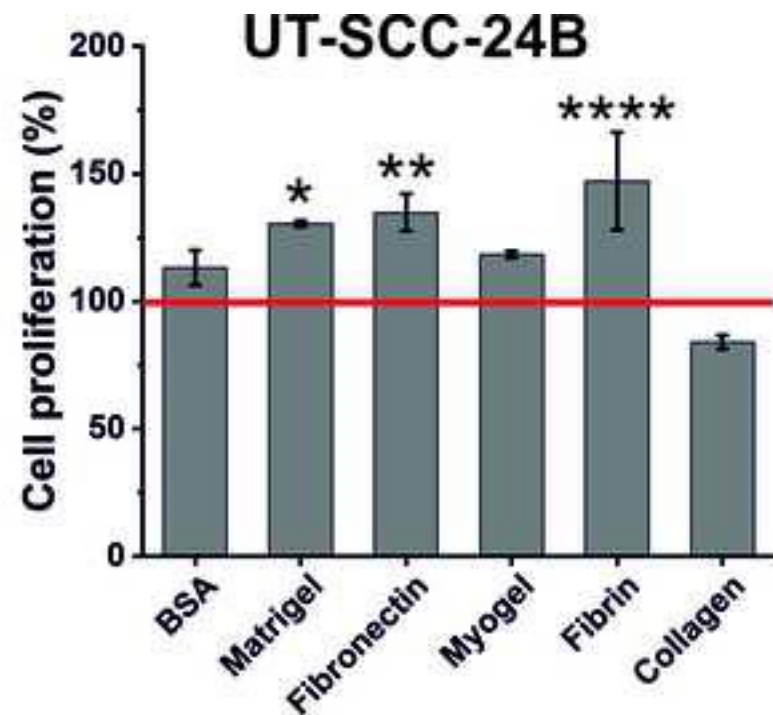
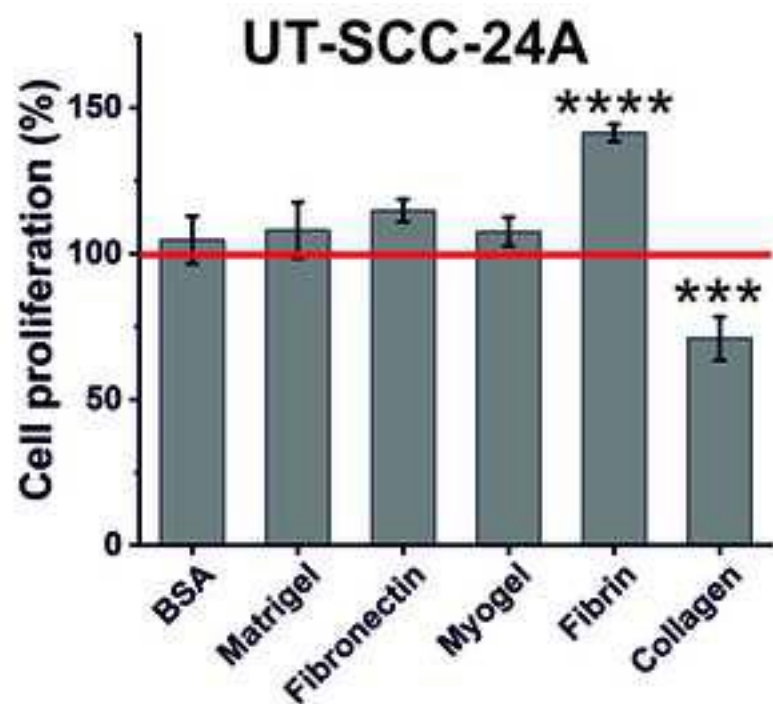


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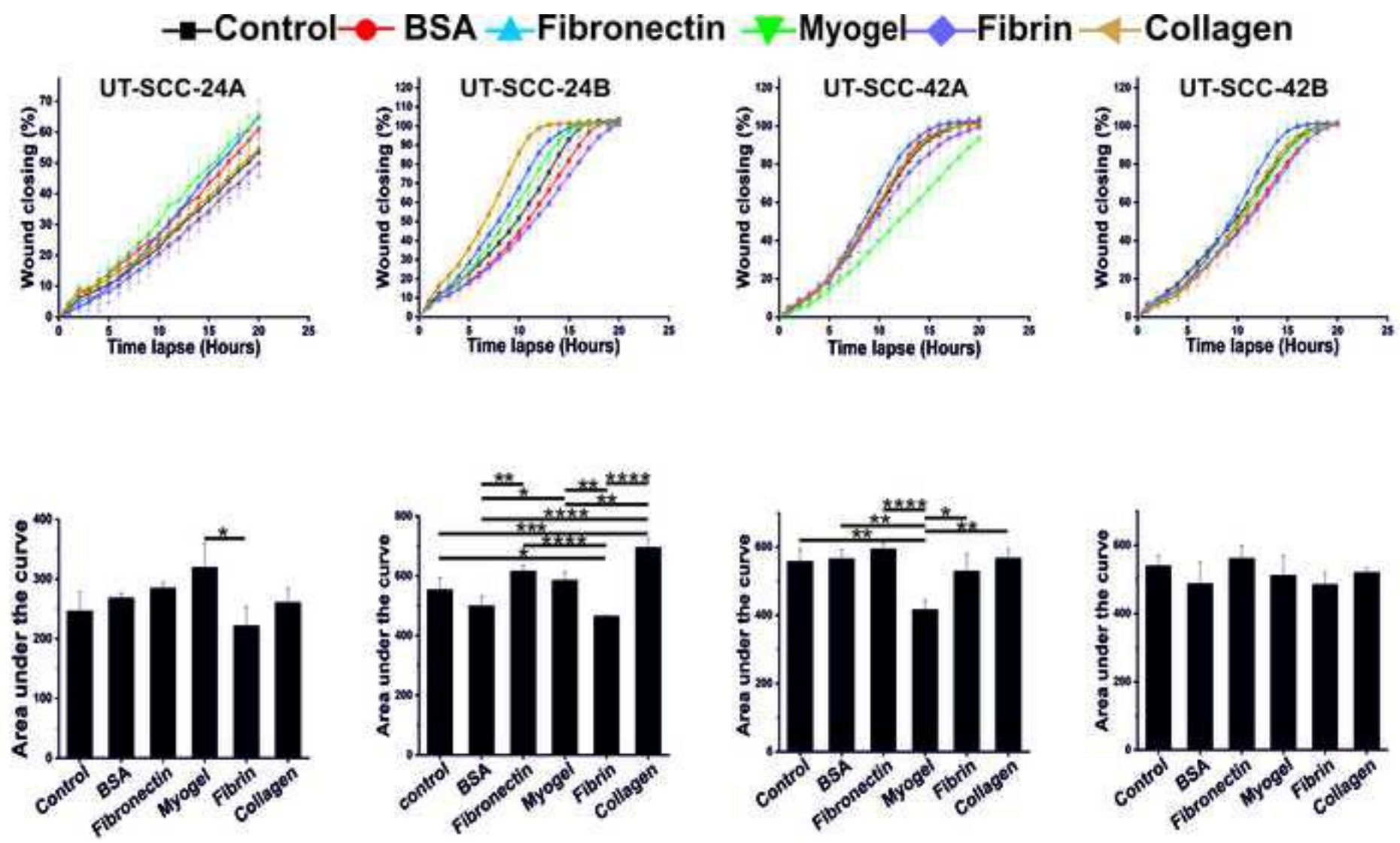


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■ Collagen ● Myogel-Collagen ▲ Fibrin ▼ Myogel-Fibrin

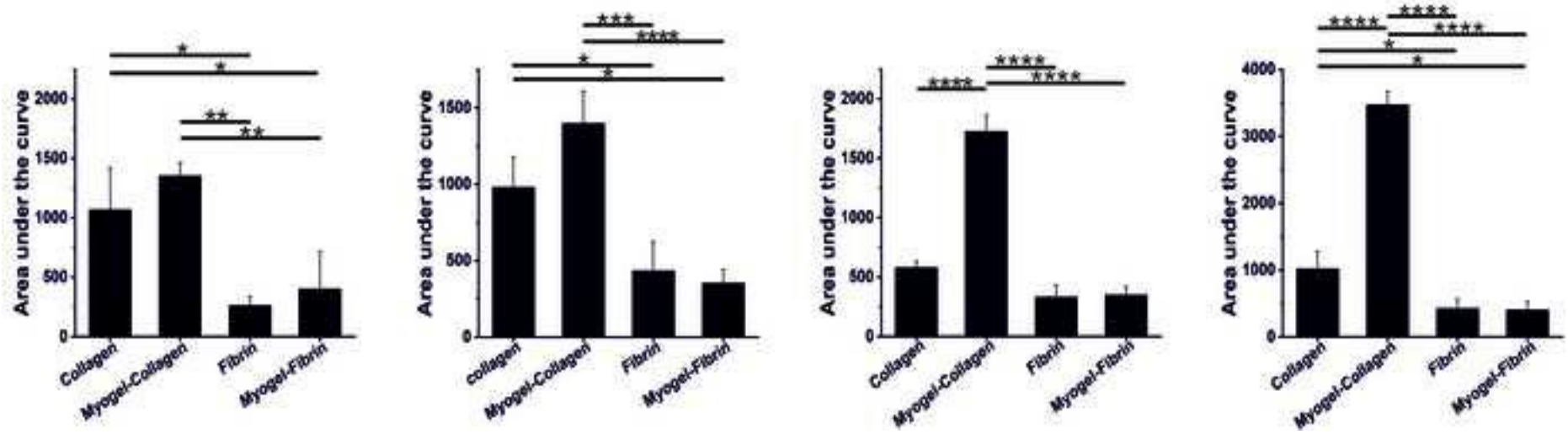
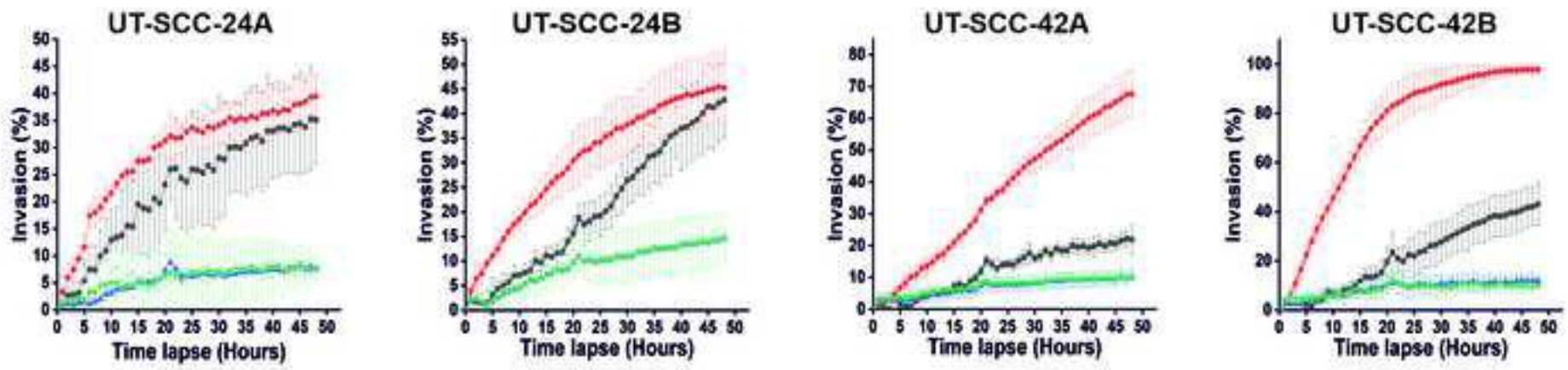


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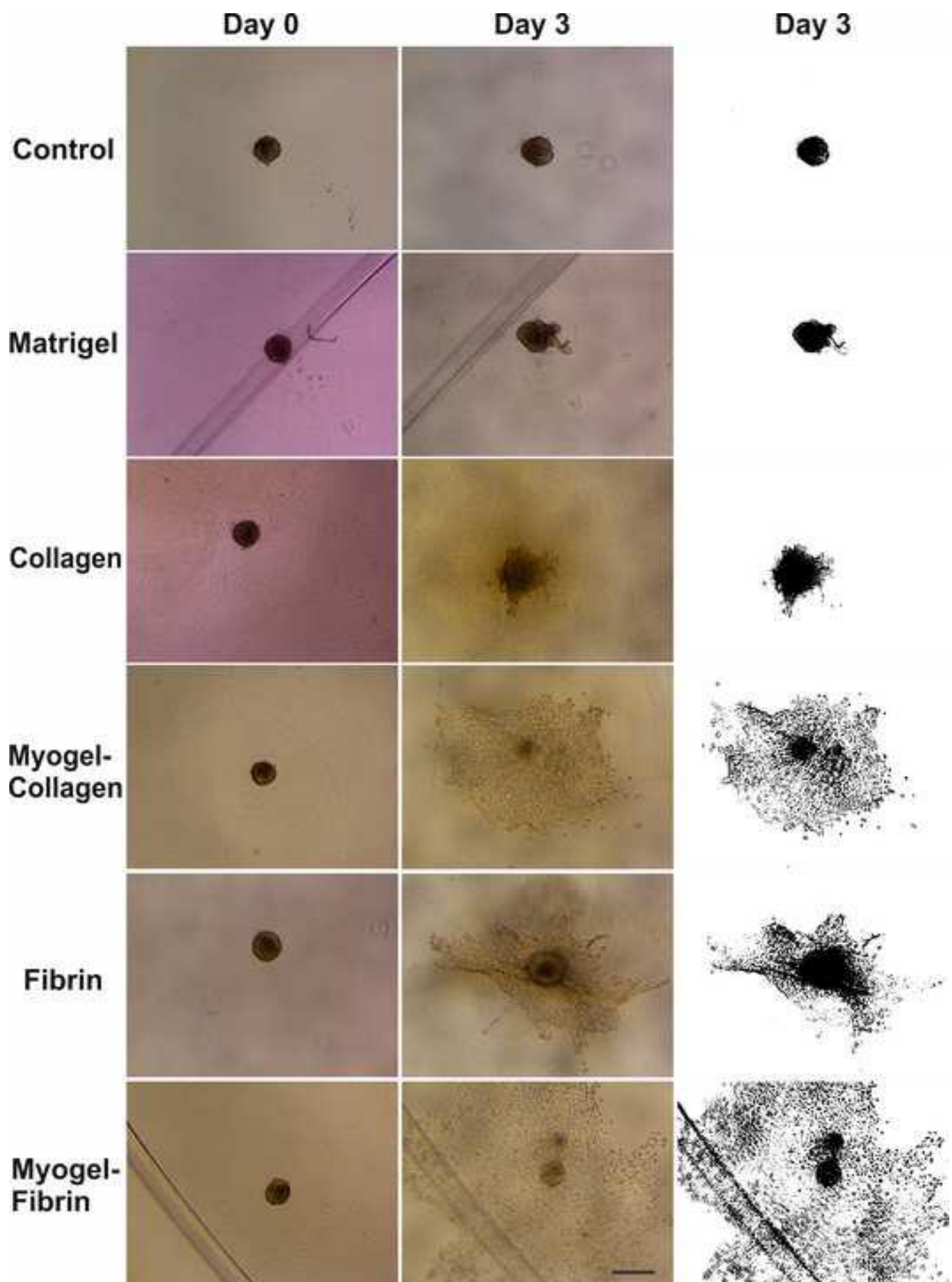
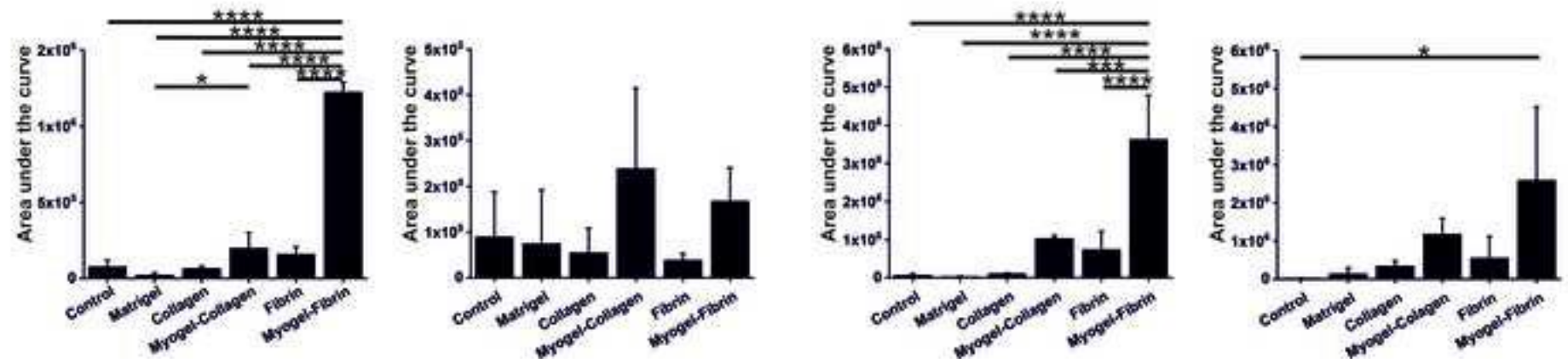
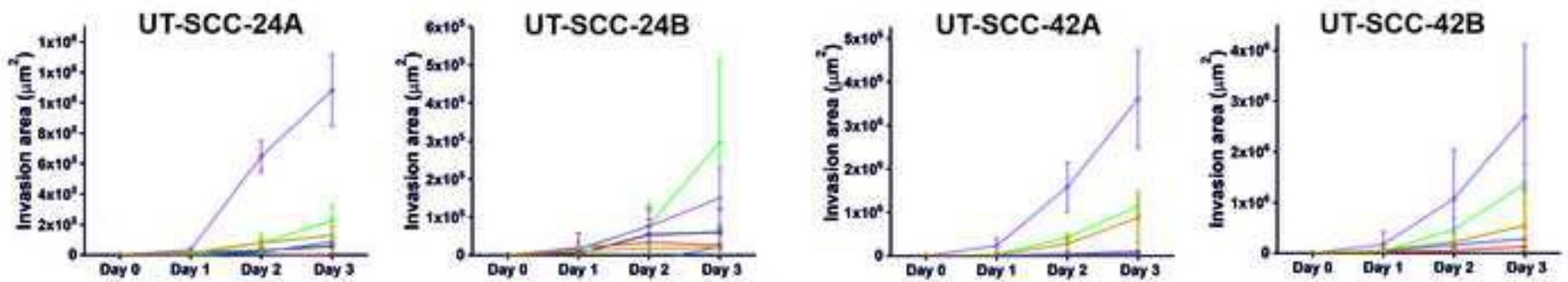


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■ Control
 ● Matrigel
 ▲ Collagen
 ▼ Myogel-Collagen
 ◀ Fibrin
 ◆ Myogel-Fibrin



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Ahmed Al-Samadi: Conceptualization, Methodology, Validation, Formal analysis, Software, Investigation, Data Curation, Writing - Review & Editing, Visualization, Project administration, Supervision.

Tuula Salo: Conceptualization, Methodology, Supervision, Writing - Review & Editing.

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Declaration of interests

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

The authors declare the following financial interests/personal relationships which may be considered as potential competing interests: