

1 **Human Airway Smooth Muscle Cell Orientation and Phenotype are altered when**
2 **cultured on Aligned Electrospun Scaffolds**

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17 **Running Head:**

18 Airway Smooth Muscle Altered Phenotype on Aligned Fibres (57 characters – max 60)

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30 **Abstract**

31

32 Human airway smooth muscle (HASM) contraction plays a central role regulating airway
33 resistance in both healthy and asthmatic bronchioles. *In vitro* studies that investigate the
34 intricate mechanisms that regulate this contractile process are predominantly conducted on
35 tissue culture plastic, a rigid, 2D geometry, unlike the 3D microenvironment smooth muscle
36 cells are exposed to *in situ*. It is increasingly apparent that cellular characteristics and
37 responses are altered between cells cultured on 2D substrates compared to 3D
38 topographies. Electrospinning is an attractive method to produce 3D topographies for cell
39 culturing as the fibres produced have dimensions within the nanometre range, similar to
40 cells' natural environment. We have developed an electrospun scaffold using the non-
41 degradable, non-toxic, polymer polyethylene terephthalate (PET) composed of uni-axially
42 orientated nanofibres, and have evaluated this topography's effect on HASM cell adhesion,
43 alignment, and morphology. The fibres orientation provided contact guidance enabling the
44 formation of fully aligned sheets of smooth muscle. Moreover, smooth muscle cells cultured
45 on the scaffold present an elongated cell phenotype with altered contractile protein levels
46 and distribution. HASM cells cultured on this scaffold responded to the bronchoconstrictor
47 bradykinin. The platform presented provides a novel *in vitro* model that promotes airway
48 smooth muscle cell development towards a more *in vivo*-like phenotype whilst providing
49 topological cues to ensure full cell alignment.

50

51 (213 words – 250 max)

52

53 **Keywords:**

54

55 Airway Smooth Muscle, Tissue Engineering, Aligned fibres, Electrospinning, *In vitro* Model

56 **Introduction**

57

58 Within the airway bronchiole, human airway smooth muscle (HASM) exists as an aligned
59 population within bundles that wrap around the bronchiole in a helical fashion (25). Smooth
60 muscle is the key effector cell regulating airway tone, with the smooth muscle contractile
61 state directly controlling the luminal capacity of the bronchiole. Dysfunction to the regulation
62 of smooth muscle contraction can alter the airway tone causing both airway hyper-
63 responsiveness (10, 44), and airway remodelling; increased smooth muscle mass is evident
64 in both mild and severe asthmatic airways (7, 18, 30). To uncover the mechanisms
65 controlling these physiological and pathophysiological roles attributed to HASM cells,
66 researchers have developed numerous experimental techniques including *in vivo*, *ex vivo*,
67 and *in vitro* platforms. The relative advantages and disadvantages in utilizing these models
68 have been discussed extensively elsewhere (3, 5, 50).

69

70 The *in vitro* culturing of primary HASM cells has provided better understanding into the direct
71 effects of bronchoconstrictors and inflammatory agonists on smooth muscle responses. One
72 such technique is the real-time imaging of intracellular signalling molecules that are
73 mobilised when muscle contraction is initiated, such as the increase in intracellular calcium
74 concentration ($[Ca^{2+}]_i$). Many important bronchoconstrictors mediate their effects through G
75 Protein-coupled receptors (GPCRs) that cause downstream activation of phospholipase C
76 (PLC), leading to the generation of the second messengers inositol 1,4,5-triphosphate (IP_3)
77 and diacylglycerol (DAG), and a subsequent increase in $[Ca^{2+}]_i$ from intracellular
78 sarcoplasmic reticulum stores and/or Ca^{2+} influx through voltage-dependent L-type Ca^{2+}
79 channels in the plasma membrane (4, 5). Bradykinin (BK) is a powerful bronchoconstrictor
80 whose effects are mediated through $G\alpha_{q/11}$ -coupled receptors present on HASM cells (21),
81 which show an increase in $[Ca^{2+}]_i$ upon stimulation (31). These, and most *in vitro* studies,
82 have been conducted on glass, or tissue culture plastic (TCP); both 2D, rigid surfaces

83 routinely used for cell culture. It is increasingly apparent that cell culture in a 3D, elastic,
84 environment can greatly alter cell phenotype and cell response to their immediate
85 surroundings (13, 14, 52).

86

87 Within a lung context, collagen gels can provide a simple 3D matrix for HASM cell culture (8,
88 32, 46), with cells residing in random planes of the gel-matrix opposed to an aligned
89 population of smooth muscle cells. Advancements in tissue engineering have provided new
90 fabrication techniques offering precise control of a substrate's nanoscale topography (29).
91 Model studies have shown multiple cell types can be aligned over extended areas using
92 different topological references such as plasma/extracellular matrix deposition (38),
93 nanoimprinting (12, 26, 54), or aligned electrospun fibres (11, 51). Electrospinning produces
94 fibrous, porous, 3D mats that closely resemble the natural extracellular matrix: The desired
95 polymer is dissolved in an appropriate solvent and this solution is passed through a syringe
96 with an electrical charge applied to the needle tip, causing the electrically charged solution to
97 become attracted to a differently charged collector plate. As the polymer jet passes through
98 the air, the solvent evaporates depositing a continuous, non-woven mesh of fibres on the
99 collector plate. Alterations to intrinsic parameters including (but not exhaustive) polymer
100 concentration, electrical field strength, or collector plate aspects, allow precise control over
101 scaffold characteristics including shape, porosity, tensile strength, and fibre diameter (16, 43,
102 47). Electrospun fibres can be manipulated to form uni-axially-aligned scaffolds through
103 alterations to fibre collection methods such as employing a rotating collector, or parallel
104 electrodes (47). This technology has been applied in culturing cells requiring directional
105 growth in oriented tissues such as neurons (9, 37, 41, 53), ligaments (42), and smooth
106 muscle (2, 19, 51), and has recently been shown to influence stem cell differentiation (40).

107

108 In the present study we investigated the influence of fibre alignment and diameter on HASM
109 cell characteristics using the non-degradable, non-toxic, polymer polyethylene terephthalate

110 (PET). Whilst randomly aligned PET fibres have been employed for the culture of aortic
111 smooth muscle (34), and other polymers have been electrospun into aligned scaffolds for the
112 culture of vascular (51), and bladder (2) smooth muscle, to our knowledge this is the first
113 time HASM has been cultured on aligned electrospun fibres. The effect of this 3D aligned
114 topography on HASM cell orientation, morphology and contractile characteristics were
115 investigated and compared to HASM cells cultured on rigid 2D surfaces or within *in situ*
116 airway tissue.

117

118 **Materials and Methods**

119 **Materials**

120 All materials were purchased from Sigma-Aldrich (Dorset, UK) unless stated otherwise. The
121 Novex protein separation kit, nitrocellulose membrane, alamarBlue[®] solution, Fluo-4-AM,
122 anti-mouse and anti-rabbit fluorescein isothiocyanate-conjugated (FITC) and rhodamine-
123 conjugated secondary antibodies, and the nuclear stain Hoechst were all obtained from
124 Invitrogen Life Technologies (Paisley, UK). Rabbit polyclonal antibodies raised against
125 Calponin and SM22 α were purchased from AbCam (Cambridge, UK). The Protease Inhibitor
126 Cocktail Set III was purchased from Merk Millipore (Nottingham, UK)

127 **Scaffold Fabrication**

128 Electrospinning procedures were conducted at room temperature within in a vented chemical
129 fume hood. PET scaffolds were produced by dissolving PET (food grade drinking bottles) in
130 a 1:1 trifluoroacetic acid (TFA):di-chloromethane (DCM) (Fisher Chemicals, Loughborough,
131 UK) solution to create a 10% (*w/v*) or a 20% (*w/v*) PET polymer solution. The PET polymer
132 solution was loaded into a syringe attached to a blunt 18-gauge (G) needle (BD Falcon[™],
133 Oxford, UK) and placed in a syringe pump-driver (Harvard Apparatus Ltd., Kent, UK) set to a
134 solution feed rate of 0.5 ml/hr or 1.5 ml/hr for the 10% or 20% scaffolds respectively. A 14 kV
135 voltage charge was applied to the needle tip with fibres being collected on a stainless steel

136 grounded cylindrical mandrel positioned 15 cm from the needle tip. The mandrel was rotated
137 at 2000 revolutions per minute (equivalent to 600 metres/minute (m/min)) to establish fibre
138 alignment. Individual electrospinning parameters are summarized in Table 1.

139 **Scaffold Characterization**

140 Each scaffold-type was electrospun independently three times. Scaffolds were punctured
141 and mounted on carbon discs prior to coating with a thin layer of gold (Balzers Union SCD
142 030, Balzers Union Ltd., Liechtenstein) and imaged on a scanning electron microscope
143 (JEOL JMS-6060 LV, JEOL Ltd., Welwyn Garden City, Hertfordshire, UK). The fibre
144 diameters and fibre orientation within the scaffolds were determined through image analysis
145 from two samples taken from each scaffold. Four scanning electron microscope images
146 were taken of each sample, and twenty fibre measurements were analysed from each
147 individual image (480 fibre measurements per scaffold-type):

148 Fibre diameter measurements were calculated using the software package measureIT 5.1
149 (Olympus Soft Imaging solutions, Münster, Germany). To determine the degree of angle
150 uniformity within the scaffolds, the mean fibre angle of an individual scaffold was determined
151 using ImageJ, with the individual fibre deviation from this mean angle calculated.

152 Scaffold thickness was measured using a digital thickness gauge (accurate to 10 μm)
153 (Mitutoyo, Coventry, UK). Scaffold mass was calculated using a top-pan balance (accurate
154 to 0.1 mg) (Mettler Toledo, Leicester UK). The mass and scaffold thickness determined the
155 apparent density of the scaffold and the porosity calculated as: Porosity (%) = $1 - (ad/bd)$
156 $\times 100$ where ad is the scaffold's apparent density and bd is the density of pure amorphous
157 PET (1.38 g/cm^3). Scaffold porosity was calculated using 8 mm diameter scaffold samples.

158 Uni-axial tensile tests were performed on three samples from three independently
159 electrospun scaffolds ($n=9$). Samples (30 mm length, 3 mm width) were loaded with fibres
160 running parallel to the applied load direction on a 5969 Universal testing system (Instron,

161 High Wycombe, UK), with a 50N load cell operating with an extension rate of 5 mm/min⁻¹.
162 The Young's moduli of the samples were calculated from the resultant stress/strain curves
163 generated using an imetrum VideoGauge.

164 **Cell Culture on Aligned Scaffolds**

165 Primary HASM cells from non-asthmatic individuals were isolated from bronchial biopsies at
166 the Glenfield Hospital (Leicester, UK) as described previously (28). The research was
167 approved by the Leicestershire Ethics Committee, and patients gave their written informed
168 consent. HASM cells (passage 3–6) were grown in Dulbecco's modified Eagle medium
169 (DMEM) supplemented with 10% (v/v) foetal calf serum, 2mM L-glutamine solution, 1% (v/v)
170 antibiotic/antimycotic solution (10,000 units/mL penicillin G, 100 mg/mL streptomycin
171 sulphate and 25 µg/mL amphotericin B). Prior to cell seeding, scaffolds were sterilized by
172 UV-irradiation for 30 minutes on both scaffold surfaces (60 minutes total). Scaffolds were
173 subsequently soaked in a 20% (v/v) antibiotic/antimycotic solution (200,000 units/mL
174 penicillin G, 2000 mg/mL streptomycin sulphate and 500 µg/mL amphotericin B) overnight at
175 37°C before washing in media prior to cell seeding.

176 **Immunocytochemistry**

177 Samples were fixed in 3.8% (w/v) paraformaldehyde (PFA) before permeabilization in a
178 0.5% (v/v) Triton X-100 PBS solution. Non-specific antibody binding was reduced by
179 incubation in 3% (w/v) bovine serum albumin (BSA) solution proceeded by a 10% (v/v) goat
180 serum solution incubation. Samples were incubated with primary antibody overnight at 4°C.
181 Protein expression was visualised with species-appropriate secondary antibodies and nuclei
182 were visualised by Hoechst staining. Samples were viewed on a Leica TCS SP2 laser
183 scanning confocal inverted microscope (Leica Microsystems Ltd, Milton Keynes, UK) with
184 post-visualization image modification performed using Volocity® (Perkin Elmer, Cambridge,
185 UK).

186 Cell elongation and height were analysed using Volocity[®] to determine individual cell's
187 dimensions: XY images determined the long (*l*), and short (*s*), axes (length and width), with
188 the elongation factor (EF) determined as: $EF=(l/s)-1$. XZ images were used to determine cell
189 height (*h*) through cross sectional images. Cell alignment in relation to fibre orientation was
190 determined using HASM cell nuclei angles from 4 or more images for each time point and
191 using three HASM cell donors (n=3). Nuclei angle deviation from the fibre angle orientation
192 was calculated and expressed as a percentage of cells orientated within 10° incremental
193 steps from the mean fibre angle.

194

195 **Histological Staining**

196 Airway biopsies were fixed in phenylmethylsulphonylfluoride (PMSF)/acetone solution before
197 washing in water-free acetone. Samples were incubated in methyl benzoate followed by
198 incubating in 5% methyl benzoate/glycol methanlacrylate (GMA) processing solution in three
199 2 hour incubations at 4°C. Biopsies were embedded in GMA after polymerization for 48
200 hours at 4°C. Samples were cut into 2 µm sections, and placed onto superfrost slides
201 (Thermo Scientific, Surrey, UK). Samples were serially rehydrated through a descending
202 ethanol (in water) concentration to 0% (v/v) ethanol before staining in Harris haematoxylin.
203 Samples were serially dehydrated through an ascending ethanol concentration to 90% (v/v)
204 ethanol before staining in eosin and further dehydration in 100% ethanol and xylene.
205 Samples were air-dried, and mounted in DPX mountant prior to imaging.

206

207 **Western Blotting**

208 Airway tissue was freeze-snap dried in liquid nitrogen and stored at -80°C until use. 30 mg of
209 tissue was homogenised using a manual homogenizer (Thermo Scientific) in lysis buffer
210 (mM: 20 HEPES, 200 NaCl, 10 EDTA, 0.1% Triton X-100, and 0.5% Protease Inhibitor Set
211 Cocktail III, pH 7.4). Homogenized tissue was agitated for 2 hours on an orbital shaker at
212 4°C before centrifugation (16'000xg, 20 minutes 4°C), with samples stored at -20°C until
213 use. HASM cells cultured in 2D or 3D were cultured for 14 days prior to incubation in lysis

214 buffer. The Novex protein separation kit was used to measure protein content in HASM cells.
215 Lysates (20 µg/lane) were separated by SDS-PAGE before transfer onto nitrocellulose
216 membrane. Membranes were blocked in 5% non-fat dry milk in Tris-buffered saline
217 containing 0.1% Tween (TBST) before incubation with primary antibody overnight at 4°C.
218 Membranes were incubated with horseradish peroxidase-conjugated secondary antibody and
219 visualized by enhanced chemiluminescence (GE Healthcare, Buckinghamshire, UK). Images
220 were captured on a LAS 4000 Luminescent Image Analyser (Fujifilm, Düsseldorf, Germany).
221 Quantitative signals were derived by densitometric analysis using Advanced Image Data
222 Analyzer (AIDA) software, with total protein content corrected to GAPDH protein levels.

223 **Calcium Signalling**

224 HASM cells were cultured for 10 days prior to loading with the Ca²⁺-sensitive dye Fluo-4-AM
225 (3 µM, 60 min). Scaffolds were transferred to an imaging unit where cells were maintained at
226 37°C in Krebs-Henseleit buffer (mM: 134 NaCl, 6 KCl, 1 MgCl₂, 1.2 KH₂PO₄, 10 glucose, 10
227 HEPES, 1.3 CaCl₂, pH 7.4). Real-time images were taken using an epifluorescence Nikon
228 Eclipse TE200 microscope (Nikon, Tokyo, Japan) (×40 objective). Cells were excited at 488
229 nm and emission collected between 505 and 560 nm. Bradykinin (100 nM) was applied to
230 cells, and the fluorescence emission was measured from regions of interest within the cells
231 cytosol. [Ca²⁺]_i changes are displayed as the fluorescence emission relative to basal (F/F₀).

232 **Data and Statistical Analysis**

233 Data are presented for cells obtained from at least three individual HASM cell donors
234 cultured on aligned scaffolds electrospun independently at least three times. Data are
235 expressed as mean±standard error of mean (SEM). Data have been analysed (GraphPad
236 Prism, San Diego CA) using T-test or one-way ANOVA and appropriate *post-hoc* testing as
237 indicated.

238

239 Results

240

241 Scaffold Characteristics Manipulated through Alterations to Electrospinning 242 Parameters and Fibre Collection Protocols.

243

244 To investigate whether HASM cell characteristics were affected when exposed to a 3D
245 nanofibre topography or an aligned fibre topography, we electrospun a randomly aligned
246 10% (w/v) PET solution onto a collector plate (data not shown), or a 10% (w/v) or 20% (w/v)
247 PET solution onto a rotating mandrel to produce aligned fibres with diameters in either the
248 nanometre or micrometre range. Representative SEM images of the 10% and 20% scaffolds
249 are shown in 1A, and 1B respectively. The 10% (w/v) PET solution produced fibre diameters
250 of 0.2 (± 0.002) μm , and the 20% (w/v) produced fibre diameters of 1.1 (± 0.001) μm (\pm SEM)
251 (1C). Both the 10% and 20% scaffolds had >90% fibres orientated within 10° of the mean
252 fibre angle and >50% of fibres orientated within 5° of the mean fibre angle (1D). Both
253 scaffolds had a porosity >80%, whilst the 20% aligned scaffold was both thicker than the
254 10% aligned scaffold (107 vs. 24 μm) and had a greater tensile strength (290 vs. 211 MPa).
255 All the electrospinning parameters and individual scaffold properties are summarized in
256 Table 1.

257

258 HASM cells show altered Phenotypic Characteristics when cultured on Aligned 259 Electrospun Scaffolds

260 Airway bronchiole sections displaying longitudinal HASM cell populations (2A) were used to
261 quantify *in situ* HASM cell alignment by referencing the angle of each cell's nucleus to
262 indicate cellular direction, with >60% cells orientated within 10° of the mean bundle angle
263 (2E&F). HASM cells were also cultured on 22mm glass coverslips (2B), randomly aligned
264 scaffold, 10% (2C) or 20% (2D) aligned scaffolds (cut to 22mm diameter dimensions). Cell

265 alignment was measured in three separate areas of the same sample to quantify cell
266 alignment over extended distances. Cell alignment to the underlying fibre orientation was
267 determined using nuclei angles to indicate cell direction at day 7 and day 14. With no
268 topological references, cells grown on 2D glass coverslips (2B) or 3D randomly aligned
269 scaffolds (data not shown) showed little orientation as a cell population over extended
270 distances, although cells showed local alignment upon confluency by day 14 (5A&C). Cells
271 showed consistent orientation along the 10% (2C), and 20% aligned scaffold (2D), with
272 >40% cells orientated within 10° of the mean fibre angle on both day 7 and 14 on either
273 scaffold, compared with <25% cells showing alignment when cultured on glass (2E&F).

274 The effect of a 3D aligned topography on cellular dimensions was compared to 2D and 3D-
275 randomly aligned surfaces by culturing HASM cells for 7 days prior to fixation and
276 immunostaining for SM22 α . Airway bronchiole sections displaying longitudinal and cross-
277 sectional HASM cell populations were used to quantify *in situ* HASM cellular dimensions: XY
278 images determined individual cell's long- and short-axis measurements to calculate their
279 elongation factor, whilst XZ cross-sectional images determined cell height. Cells cultured on
280 2D substrates or residing within smooth muscle bundles were significantly shorter compared
281 to cells cultured on 3D aligned scaffolds (3A, B, E&G). Cells showed an approximate
282 doubling in elongation when cultured on a 3D aligned surface compared to 2D (3C, D&H);
283 mainly due to a reduction in cell width opposed to an increase in cell length. Whilst cells
284 were significantly more elongated within smooth muscle bundles compared to 2D cultured
285 cells, there was no significant difference in elongation when compared to HASM cells
286 cultured on aligned scaffolds (3F&H). Cells cultured on 3D randomly aligned scaffolds had a
287 similar elongation factor to cells cultured in 2D, but showed a significant increase in cell
288 height compared to 2D culture (data not shown). As there was no significant variation in cell
289 characteristics between cells cultured on the 10% or 20% aligned scaffolds, the 10% scaffold
290 was used in further investigations due to the nanofibrous nature of the scaffold.

291 Where cells attach to the extracellular environment, focal adhesion complexes link the cell's
292 actin cytoskeleton to the surrounding matrix (49). To determine how the underlying
293 topological references affected focal adhesion complex formation and spatial distribution,
294 cells cultured on a 3D aligned topography were compared to those cultured upon a 2D
295 surface. Post-seeding, cells were fixed after 3, 24, or 72 hours before immunostaining for the
296 focal adhesion protein vinculin and co-stained for F-actin. On glass, HASM cell F-actin
297 organised into clear stress fibres throughout the cells with vinculin clusters at each cell's
298 distal ends (4A). This intra-cellular F-actin and vinculin expression was maintained at 24 (4B)
299 and 72 hours (4C), with cells also displaying a flattened cell phenotype. Cells cultured on the
300 aligned scaffolds showed little evidence of vinculin clusters at any of the time points (with
301 localised staining apparent around the nuclei), and F-actin was not organised into stress-
302 fibres throughout individual cells (4D). Vinculin staining became less apparent after 3 hours,
303 and cells started to exhibit an elongated shape that aligned along the fibre orientation
304 (3E&F).

305 Over a prolonged time period (3, 7, and 14 days), HASM cells cultured on 2D glass
306 coverslips or aligned fibres were fixed and immunostained for the smooth muscle specific
307 proteins SM22 α (5A&B) and calponin (5C&D). Contractile protein distribution in HASM cells
308 cultured on glass showed distinct stress-fibre organization, maintained a flat phenotype, and
309 displayed increasing local cell alignment upon cell confluency at day 14 (5A&C). Conversely,
310 HASM cells cultured on aligned scaffolds showed little evidence of contractile protein
311 localization into stress fibres, whilst maintaining an elongated phenotype and aligned cell
312 distribution at every time point (5B&D). The presence of gap-junctions were visualised by
313 immunostaining for connexin-43 with qualitatively greater localized connexin-43 expression
314 seen between adjacent HASM cells cultured on aligned scaffolds than cells cultured on glass
315 (5F&E respectively).

316 HASM cells cultured on TCP or aligned scaffolds were lysed, and protein levels semi-
317 quantitatively compared after protein levels were normalised to levels of GAPDH protein
318 expression (6A). Cumulative data (6B) found levels of the smooth muscle-specific proteins
319 smooth-muscle α -actin (0.9 ± 0.2 vs. 1.2 ± 0.3), calponin (0.7 ± 0.1 vs. 1.3 ± 0.1), SM22 α (0.5 ± 0.2
320 vs. 1.3 ± 0.4), and desmin (0.02 ± 0.01 vs. 0.9 ± 0.3) ($n=5-6$, mean \pm SEM) all increased upon
321 aligned scaffold-culture, with levels of calponin and desmin increasing significantly. The
322 expression of smooth muscle specific proteins is often reduced/lost upon prolonged cell
323 culture (23, 39). Protein levels from freshly isolated airway tissue were determined
324 separately; with the relative levels of smooth muscle-specific proteins expressed as a ratio to
325 the level of smooth-muscle α -actin. The ratio of smooth muscle α -actin : calponin : SM22 α :
326 desmin for airway tissue was 1.0 : 0.9 ± 0.04 : 0.7 ± 0.1 : 1.2 ± 0.3 , HASM cultured on aligned
327 scaffold was 1.0 : 1.1 ± 0.2 : 1.0 ± 0.2 : 1.0 ± 0.3 , and HASM cultured on 2D TCP was 1.0:
328 0.7 ± 0.1 : 0.7 ± 0.3 : 0.03 ± 0.02 ($n=3-6$, mean \pm SEM).

329

330 To investigate HASM cell responses to bronchoconstrictor application, HASM cells cultured
331 either on 2D glass or aligned scaffolds were loaded with the Ca^{2+} sensitive dye Fluo-4-AM
332 and stimulated with the $\text{G}\alpha_{q/11}$ -agonist BK (100 nM) causing a biphasic increase in $[\text{Ca}^{2+}]_i$,
333 with an initial transient increase in Ca^{2+} concentration followed by a reduction to lower steady
334 state Ca^{2+} concentration (7A). HASM cells cultured on glass or the aligned scaffold showed
335 similar levels of peak Ca^{2+} release (7B).

336

337 **Discussion**

338 Advancements in biomaterial design allow better control to fabricate 3D matrices that are
339 specifically tailored to individual cell-types opposed to the “one rigid 2D substrate fits all”
340 approach offered by TCP. Smooth muscle exists as an aligned population of cells, and this

341 organization ensures a co-ordinated contractile response. To maintain these characteristics
342 *in vitro*, we electrospun uni-axially orientated fibres to attempt to direct the HASM cell
343 alignment. One previous study has described the alignment of PET fibres by electrospinning
344 onto a rotating mandrel (22). However the rotational velocity employed was much lower than
345 applied here (36 m/min vs. 600 m/min respectively). This large increase in velocity produced
346 thinner nanofibres (0.2 μm vs. 0.7 μm) at similar polymer concentrations (10% vs. 9% (w/v)).
347 Comparable micron diameter fibres were produced using a 20% w/v polymer solution
348 compared to the 12.5% w/v solution (1 μm vs. 1.7 μm fibres). Qualitatively, we demonstrated
349 more consistent alignment of fibres throughout the scaffolds, and greater tensile strength
350 (211 MPa vs. 90 MPa) with respect to aligned nanofibre scaffolds.

351

352 When HASM cells were cultured on either a 2D or 3D surface with no orientation reference,
353 only transitory cell alignments were seen over short distances upon contact inhibition. We
354 measured local cell alignment from separate regions of the same sample to investigate cell
355 population alignment over extended distances. The contact guidance provided by fibre
356 orientation ensured good mediation of cell alignment from 3 hours post-seeding, an effect
357 maintained over 14 days. Cells seeded on either aligned scaffold showed consistently good
358 alignment to the fibre orientations (>70% cells within 20° of the scaffold mean fibre angle
359 using either aligned scaffold), though this was still less than cell alignment within *in situ*
360 smooth muscle bundles (>90% cells within 20° of the mean fibre angle). Aligned nanofibres
361 have shown previously to enhance vascular smooth muscle alignment over an immediate
362 time course of 24 hours (51), whilst here we show this effect can be maintained over an
363 extended time period under static conditions. Other studies have achieved similar results
364 with no underlying topological reference through the application of a uni-axially mechanical
365 strain (20), with cells aligning perpendicular to the applied force.

366 HASM cells cultured in 2D showed clear vinculin localization within defined focal adhesions
367 at the distal ends of the cells, with F-actin and smooth muscle-specific contractile proteins

368 organised into stress-fibres throughout the cells, an effect most likely caused by the rigidity
369 of the substrate (TCP approximately 2-4 GPa (6)). On the aligned scaffolds, HASM cells
370 showed a reduction in vinculin localization within focal adhesion complexes, and a more
371 even distribution of intra-cellular F-actin, calponin, and SM22 α protein expression. Similar
372 reductions in focal adhesions and cytoskeletal organisation have been noted in HASM cells
373 (15, 24), and fibroblasts (1, 27) cultured on 2D surfaces coated with specific extracellular
374 matrix proteins, although HASM cells cultured on these substrates still maintained a
375 flattened cell shape (15, 24). HASM cells cultured in 2D also showed a reduction in
376 elongation compared to those cultured on aligned scaffolds where cells elongated along
377 individual fibres and displayed a more spindle-like morphology than the short, flat, and fat
378 morphology that can occur upon 2D culture. Taken together these data suggest the
379 topography encountered by the smooth muscle may play a dominant role in the cytoskeletal
380 protein organization opposed to the relative rigidity of the substrate, and the cues provided
381 by the aligned fibres ensure HASM cells orientate, and elongate along individual fibres
382 opposed to interacting with multiple fibres as seen when cultured on randomly aligned
383 scaffolds.

384

385 Levels of contractile proteins are known to rapidly decrease when HASM cells are passaged
386 upon isolation from fresh tissue (23, 39), with levels of desmin virtually abolished within 7
387 days post-isolation (23). The culturing of HASM cells on a 3D aligned scaffold was sufficient
388 to significantly increase calponin and desmin protein levels compared to 2D culture, with
389 desmin expression upregulated from negligible levels (2D culture) to consistent expression
390 (3D aligned culture). These expression levels were compared to freshly isolated airway
391 tissue. Relative protein levels, when normalised to smooth muscle α -actin levels, displayed
392 similar expression patterns between airway tissue and 3D-aligned cultured cells; smooth
393 muscle α -actin, calponin and desmin all showing a 1:1 ratio, whilst 3D-aligned cultured cells
394 showed a relative enhancement in calponin expression (1.0 : 0.6). 2D cultured cells showed

395 a relative reduction in calponin expression and a significant reduction in desmin expression
396 as reported elsewhere (23). Whilst expression patterns are comparable between airway
397 tissue and aligned scaffold cultured cells, the magnitude in protein expression is likely to be
398 underestimated within this study due to the heterogeneous airway cell population used to
399 determine levels of specific smooth muscle proteins within tissue opposed to a homogenous
400 smooth muscle population. Alterations to smooth muscle-specific contractile proteins have
401 also been observed in 3D cultures of HASM cells when cocultured with fibroblasts (48), or
402 fibroblasts and epithelial cells under pulsatile conditions, albeit over an extended time course
403 of 28 days (33). Mechanical strain causes similar alterations to HASM cell contractile protein
404 mRNA and protein levels, including desmin and calponin (2, 20, 45, 55), suggesting similar
405 SMC phenotypic modulations (SMC alignment and contractile protein levels) can be induced
406 through mechanical stimulation, or exposure to a 3D aligned topography. These
407 enhancements in the underlying contractile machinery of the HASM cells could help more
408 accurately model smooth muscle functionality within an *in vitro* setting.

409 Smooth muscle contraction can be studied using a variety of techniques. Measuring
410 changes in key second-messenger signalling molecules such as $[Ca^{2+}]_i$ have previously
411 been employed in airway- (17) and vascular-smooth muscle (35, 36) contractile profiling.
412 When the bronchoconstrictor BK was applied to HASM cells cultured on 2D or aligned
413 scaffolds, peak Ca^{2+} levels, and Ca^{2+} release profiles were comparable. Given the
414 pronounced alterations in the HASM cellular characteristics and protein levels when cultured
415 on aligned scaffolds compared to 2D culture one may have expected an increase in Ca^{2+}
416 responses. Other studies employing alternative outputs to measure smooth muscle
417 contraction have noted an increase in the comparative shortening of asthmatic smooth
418 muscle compared to healthy smooth muscle (32, 46). A novel multi-cell micro-tissue culture
419 model co-culturing HASM cells with 3T3 fibroblast cells showed alterations in contractile
420 profiles in response to some agonists (such as histamine), but not others (such as KCl) (48).

421 These suggest underlying differences in Ca^{2+} responses may be uncovered if diseased cells
422 were employed in the present model.

423 In conclusion, we have developed a fully-aligned nanofibrous scaffold, whose orientation
424 guidance is sufficient to produce fully aligned HASM cell sheets that can be cultured for
425 prolonged periods whilst maintaining a more *in vivo*-like phenotype.

426

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431

432 **Figure 1: PET electrospun scaffolds can be manipulated to form aligned nano- and**
433 **microfibre scaffolds**

434 Polyethylene terephthalate (either 10% or 20% (w/v)) was electrospun onto a rotating
435 mandrel (600 metres/min) to produce aligned fibres with diameters in the nano- or
436 micrometre range. Representative SEM images of the 10% (nano) and 20% (micro)
437 scaffolds are shown in 1A and 1B respectively. Intra-scaffold fibre diameter distributions are
438 shown in 1C (n=480 from 3 separate scaffolds), and intra-scaffold fibre alignments are
439 shown in 1D (deviation of individual fibre angle from scaffold's mean fibre angle, n=480 from
440 3 separate scaffolds).

441 **Table 1: Characteristics of aligned PET scaffolds**

442 The electrospinning parameters used to create PET scaffolds consisting of uni-axially
443 orientated fibres of different diameters are listed in Table 1. The table also shows the
444 average intra-scaffold fibre diameters, the percentage of fibres orientated within +/- 10° mean
445 fibre angle, tensile strength, scaffold thickness, and scaffold porosity as described in material
446 and methods. Data are mean (±SEM) from 3 independently electrospun scaffolds.

447 **Figure 2: HASM cells orientate along scaffold fibres**

448 Haematoxylin and eosin stained immunohistological sections from ASM bundles were used
449 to quantify cell directionality *in situ* (2A). HASM cells (1.5×10^5) were cultured on glass
450 coverslips, the 10%-, or 20%-aligned scaffolds for 7 or 14 days prior to fixation. Nuclei were
451 visualised by Hoechst staining (blue) and nuclei angles used as reference to cell
452 directionality. Deviation of individual cell nuclei from fibre orientation was determined, and %
453 cell population range plotted. Representative images of HASM nuclei cultured on glass,
454 10%-, or 20%-aligned scaffolds at day 7 are shown in 2B, 2C, and 2D respectively. 2E
455 shows distribution of cell alignment on all 3 *in vitro* topographies and in ASM bundles. 2F
456 shows % cell population within +/-10° fibre orientation at day 7 and day 14 (mean±SEM,
457 HASM cells cultured on 3 independently electrospun scaffolds). Statistical significance is
458 indicated as $p=^* < 0.05$, $^{**} < 0.01$, and $p=^{****} < 0.0001$ (glass versus), or $p=^{\neq} < 0.05$, and
459 $p=^{\neq} 0.01$ (bundle versus), one-way ANOVA, Tukeys post-test. Scale bar indicates 40 µm.
460 Arrow indicates orientation of scaffold fibres.

461

462 **Figure 3: HASM cell morphology is altered when cultured on aligned fibre topography**

463 HASM cells (1.5×10^5) were cultured on glass coverslips, 10%- or 20%-aligned scaffolds for
464 7 days. Cells were fixed and immunostained for the contractile protein SM22 α (red) with
465 nuclei stained with Hoechst (blue). Representative XZ images of HASM cells grown on glass
466 or 10%-aligned scaffolds are shown in 3A&B respectively. Representative 3D opacity
467 images of HASM cells grown on glass or 10%-aligned scaffolds are shown in 3C&D
468 respectively. Representative hematoxylin and eosin stained immunohistological sections
469 from ASM bundles sectioned cross-sectionally and longitudinally are shown in 3E&F
470 respectively. Dashed arrows indicate electrospun-fibre orientation (3B&D). Scale bars
471 represent 20 μ m. The heights (h) of individual cells were calculated from XZ images through
472 cells. Individual cell's long (l) and short (s) axis were calculated from XY images, and cell
473 elongation factors determined. Cumulative cell height data is shown in 3G ($n=20-82$
474 individual cells grown on 3 independently electrospun scaffolds and 3 separate airway tissue
475 donors). Cumulative cell elongation data is shown in 3H ($n=42-73$ individual cells grown on 3
476 independently electrospun scaffolds and 3 separate airway tissue donors). Statistical
477 significance is indicated as $p=**** < 0.0001$ (glass vs. scaffold), $p=\# < 0.05$ (ASM bundle vs
478 glass), $p=#### < 0.001$ (ASM bundle vs scaffold) one-way ANOVA, Tukeys post-test.

479 **Figure 4: HASM cells show a reduction in intracellular-stress fibres when cultured on**
480 **aligned fibre topographies**

481 HASM cells (1.5×10^5) were cultured on glass coverslips or 10%-aligned scaffolds for 3, 24,
482 or 72 hours prior to fixation. Cells were immunostained for the focal adhesion protein vinculin
483 (red), with F-actin stained with phalloidin (green) and nuclei with Hoechst (blue).
484 Representative images of HASM cultured on glass coverslips for 3-, 24- and 72-hours, and
485 10%-aligned scaffolds for 3-, 24- or 72-hours are shown in 4A, 4B, 4C, and 4D, 4E, 4F
486 respectively. Scale bar indicates 20 μ m. Arrow indicates orientation of scaffold fibres.

487 **Figure 5: Smooth muscle specific protein expression in HASM cells cultured over 14**
488 **days on 2D or aligned fibre topography**

489 HASM cells (1.0×10^5) were cultured on glass coverslips (5A,C,&E) or 10% aligned scaffolds
490 (5B,D,&F) for 3, 7, or 14 days prior to fixation. Cells were immunostained for the smooth
491 muscle-specific contractile proteins SM22 α (5A&B, red) or calponin (5C&D, green), and the
492 gap-junction protein connexin-43 (5E&F, yellow), with nuclei stained with Hoechst (blue).
493 Scale bar indicates 70 μ m. Arrow indicates orientation of scaffold fibres.

494 **Figure 6: Relative HASM cells protein levels are altered when cultured on either 2D or**
495 **3D aligned topography**

496 HASM cells (1.0×10^5) were cultured on tissue culture plastic (TCP) or 10%-aligned scaffolds
497 for 14 days. Cells were lysed and 20 μ g protein loaded for SDS-PAGE separation and
498 immunoblotting. Representative immunoblots for the smooth muscle-specific proteins
499 calponin, SM22 α , smooth muscle specific α -actin, desmin, and GAPDH controls are shown
500 in 6A. Cumulative densitometric data showing protein levels normalised to GAPDH protein
501 expression are shown in 6B (data shown are mean(\pm SEM) n=5-6). Statistical significance is
502 indicated as $p=^* < 0.05$ and $p=^{**} 0.01$ (TCP vs. scaffold) unpaired T-Test.

503 **Figure 7: Primary HASM cells respond to BK when cultured on aligned scaffolds**

504 HASM cells (1.5×10^5) were cultured on glass coverslips or aligned scaffolds for 10 days.
505 Cells were loaded with the calcium-sensitive dye Fluo-4 AM and stimulated with BK (100
506 nM). Representative images showing Ca^{2+} changes, and representative Ca^{2+} traces from
507 HASM cells cultured on glass (black trace) or aligned scaffold (grey trace) and stimulated
508 with BK (100 nM) are shown in 7A. 7B shows cumulative peak Ca^{2+} release from HASM cells
509 stimulated with BK (100 nM) when cultured on glass (black bar) or aligned scaffold (grey
510 bar). Data are mean \pm SEM (30-60 cells, n=3 separate HASM donors).

511

512

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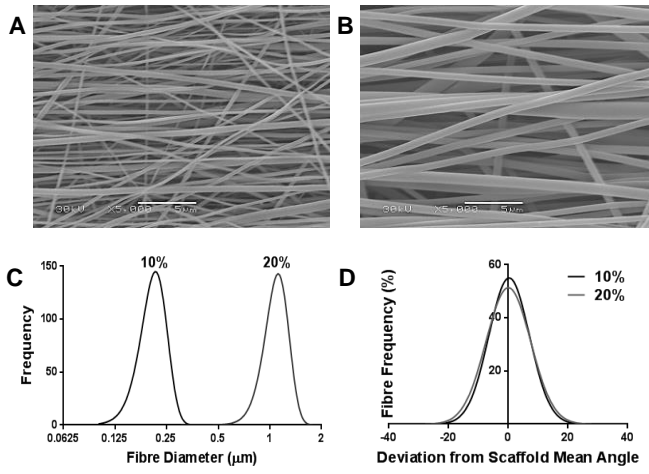
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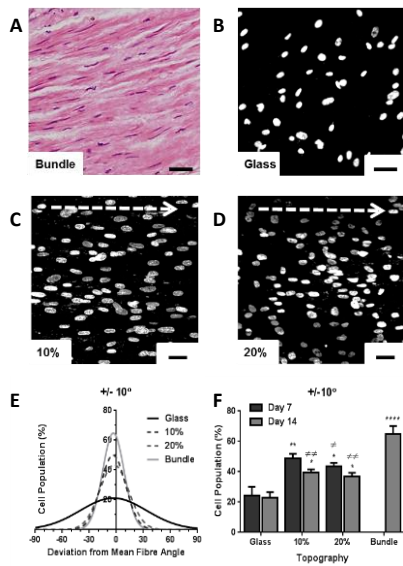
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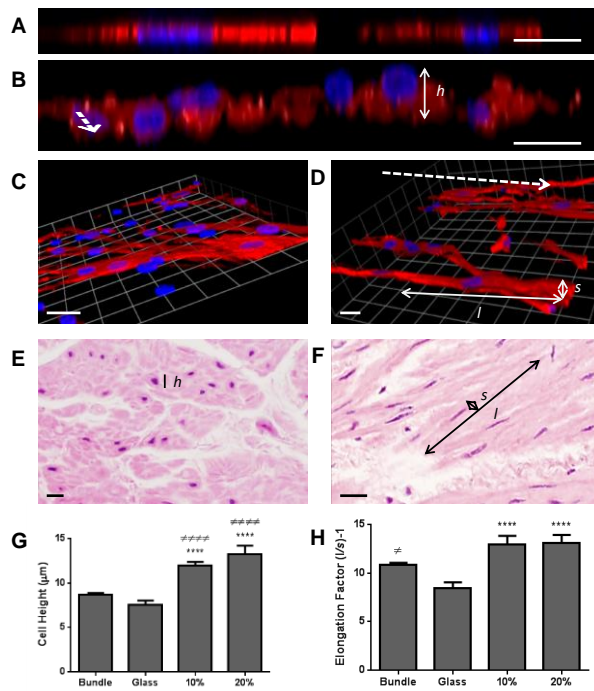
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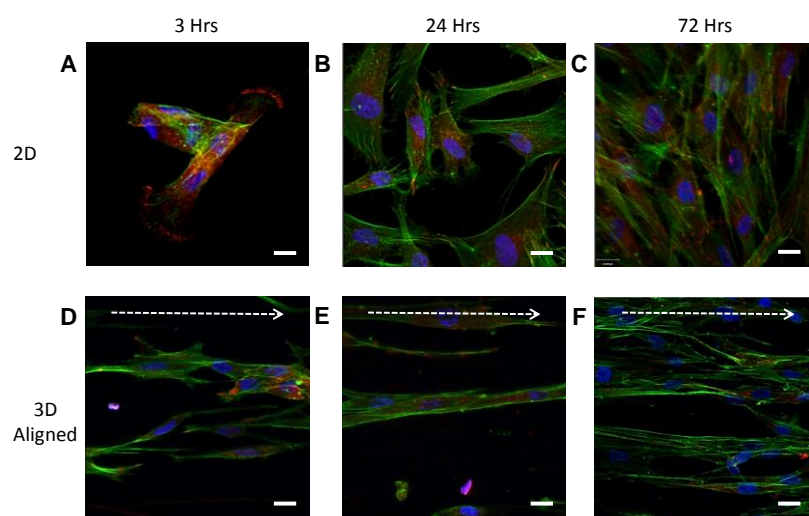
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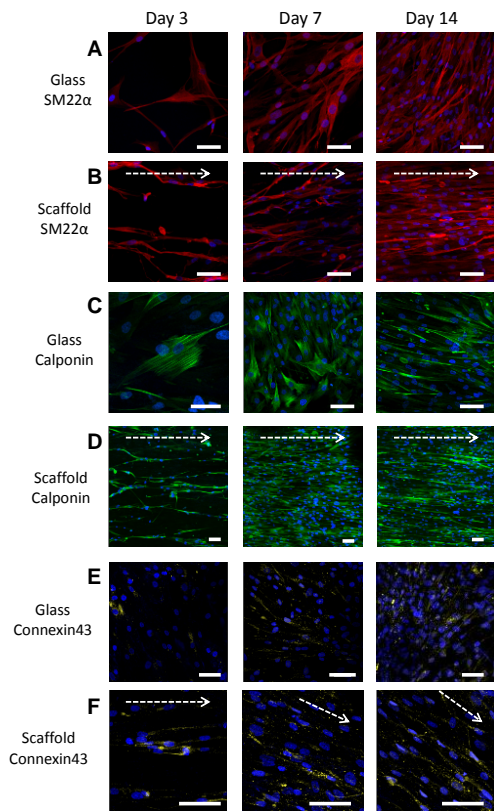
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659 Figure 4



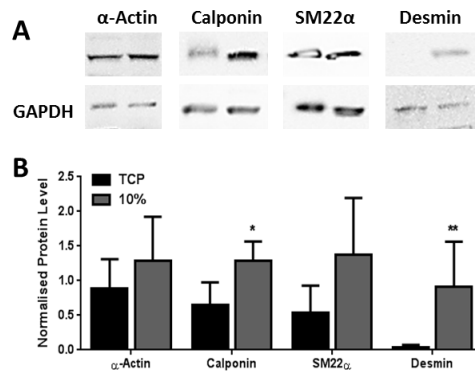
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661 Figure 5



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663 Figure 6



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