A microtubule stability switch alters isolated vascular smooth muscle calcium flux in response to matrix rigidity.

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Abstract

During ageing, the extracellular matrix of the aortic wall becomes more rigid. In response, VSMCs generate enhanced contractile forces. Our previous findings demonstrate that VSMC volume is enhanced in response to increased matrix rigidity, but our understanding of mechanisms regulating this process remain incomplete. In this current study, we show that microtubule stability in VSMCs is reduced in response to enhanced matrix rigidity via piezo1-mediated Ca^{2+} influx. Moreover, VSMC volume and Ca^{2+} flux was regulated by microtubule dynamics; microtubule stabilising agents reduced both VSMC volume and Ca^{2+} flux on rigid hydrogels, whereas microtubule destabilising agents increased VSMC volume and Ca^{2+} flux on pliable hydrogels. Finally, we show that disruption of the microtubule deacetylase HDAC6 uncoupled these processes and increased K40 alpha tubulin acetylation, VSMC volume and Ca^{2+} flux on pliable hydrogels, but did not alter VSMC microtubule stability. These findings uncover a microtubule stability switch that controls VSMC volume by regulating Ca^{2+} flux. Together, these data demonstrate that manipulation of microtubule stability can modify VSMC matrix rigidity response.

Introduction

Maintaining aortic compliance, the ability of the aorta to change shape in response to changes in blood pressure is essential for cardiovascular (CV) health. Decreased aortic compliance is a major risk factor associated with a variety of age-related CV diseases (Glasser et al., 1997; Lacolley et al., 2020; Mitchell et al., 2010). The rigidity of the aortic wall is a major contributor to aortic compliance. In the healthy aortic wall, rigidity and compliance are determined by the balance between elastic-fibres, including elastin, that provide pliability, and non-elastic fibres, including collagen-I, that provide tensile strength, to the extracellular matrix (ECM) (Tsamis et al., 2013). However, during ageing and CV disease, elastic-fibres degrade, and collagen-I accumulates. These ECM changes increase the rigidity of the aortic wall and decrease aortic compliance (Afewerki et al., 2019; Ahmed and Warren, 2018; Johnson et al., 2021).

Vascular smooth muscle cells (VSMC) are the predominant cell type within

the aortic wall. VSMC contraction regulates vascular tone in healthy aortae, where

aortic wall rigidity and compliance are a balance between ECM rigidity and VSMC

stiffness (Johnson et al., 2021). Changes in intracellular calcium ion (Ca^{2+}) levels are

integral for regulating VSMC contractile function (Ahmed and Warren, 2018). As

such, intracellular Ca^{2+} are normally tightly regulated by mechanisms that promote

release and reuptake of Ca^{2+} from intracellular stores such as the sarcoplasmic

reticulum. During VSMC contraction, Ca^{2+} release is stimulated via receptor

signalling or stretch activated ion channels, which include the TRPC family and

piezo1, leading to the activation of sarcoplasmic IP_3R and ryanodine receptors

(Sanders, 2001). This results in a rapid Ca^{2+} spark in VSMCs that drives actomyosin

force generation. These sparks are short lived as Ca^{2+} is rapidly reabsorbed to the

sarcoplasmic reticulum via SERCA channels(Sanders, 2001).

However, the balance between ECM rigidity and VSMC stiffness is disrupted

in ageing and CV disease, resulting in VSMC dysfunction (Lacolley et al., 2020;

Tsamis et al., 2013). Age associated changes in ECM composition increase aortic

wall rigidity and triggers increased VSMC stiffness, that further decreases aortic

compliance (Lacolley et al., 2020; Sehgel et al., 2015, 2013; Tsamis et al., 2013).

This is termed VSMC stiffness syndrome, but our understanding of the mechanisms

driving this process remain poorly defined. In response to hypertension, VSMCs

undergo a process known as hypertrophy and increase their cell mass without

increasing cell number (Owens and Schwartz, 1983; Rizzoni et al., 2000; Schiffrin,

2012; Zhang et al., 2005). Increases in volume and protein synthesis are key

components in VSMC hypertrophy, which is known to increase aortic wall thickness

and rigidity and decrease aortic compliance (Owens and Schwartz, 1983; Rizzoni et

al., 2000; Schiffrin, 2012; Sehgel et al., 2013; Zhang et al., 2005). Importantly, we

have recently shown that activation of piezo1 drives a sustained Ca^{2+} influx, resulting

in an increase in VSMC volume, via the membrane translocation of the water

channel aquaporin-1, in response to matrix rigidity (Johnson et al., 2024). Increased

matrix rigidity and hypertension are known to promote VSMC dedifferentiation,

where they switch from a contractile phenotype to disease associated synthetic

phenotypes (Cao et al., 2022; Hayashi and Naiki, 2009; Xie et al., 2018). Both

aquaporin-1 and piezo1 gene expression is enhanced during VSMC dedifferentiation

in atherosclerosis, during carotid remodelling and during aortic ageing (Johnson et

al., 2024; Luu et al., 2023). Whether increased volume drives increased VSMC

stiffness in response to matrix rigidity and the VSMC phenotypes involved remain

unknown.

Microtubules are hollow tube-like structures, made up of alpha and beta

tubulin heterodimers, that play important roles in determining cell morphology and

resisting deformational forces applied to cells (Brangwynne et al., 2006; Stamenović,

2005). Microtubule organisation is regulated via a process called dynamic instability,

where microtubules exist in a balance between growth and shrinkage that enables

them to respond rapidly to the changing needs of a cell (Michaels et al., 2020). Post-

translational modifications of tubulin are predicted to regulate microtubule

mechanical properties (Song and Brady, 2015). Most modifications are predicted to

occur on the outer surface of polymerised microtubules. However, lysine 40 (K40) of

alpha-tubulin in the inner lumen of polymerised microtubules, can be acetylated and

deacetylated by αTAT1 and HDAC6, respectively (Osseni et al., 2020; Shida et al.,

2010). While the precise impact of K40 acetylation on microtubule mechanical

properties remains controversial and contradictory, K40 acetylation is proposed to

influence lateral associations between polymerised tubulin monomers and potentially

stabilise microtubules, making them more resistant to mechanical damage than

deacetylated microtubules (Eshun-Wilson et al., 2019; Janke and Montagnac, 2017;

Xu et al., 2017). Enhanced K40 acetylation increases cytoskeletal stiffness in striated

muscle, but the impact of K40 acetylation in VSMC matrix rigidity sensing remains

unknown(Coleman et al., 2021).

The mechanical regulation of VSMC behaviour obeys well defined rules. For

example, in response to enhanced matrix rigidity, VSMCs generate increased

actomyosin derived traction forces and increased deformational stresses are placed

upon the cell membrane (Petit et al., 2019; Sanyour et al., 2019). These

deformational stresses drive the opening of stretch-activated ion channels in VSMCs

(Johnson et al., 2024). Microtubules exist in a mechanical balance with actomyosin

activity, serving as pre-stressed compression bearing struts capable of resisting

actomyosin generated deformational stresses (Brangwynne et al., 2006; Johnson et

al., 2021; Stamenović, 2005). This relationship is known as the tensegrity model

(Stamenović, 2005). The tensegrity model predicts the interplay between actomyosin

and microtubules in VSMCs. Previous studies have shown that microtubule

stabilisation decreases VSMC actomyosin activity, whereas microtubule

destabilisation increases VSMC actomyosin activity (Ahmed et al., 2021; Zhang et

al., 2000). Other models have been proposed that describe VSMC mechano-

adaption responses, yet we lack an in-depth mechanistic understanding of pathways

linking changes in VSMC behaviour to the proposed models of mechanotransduction

(Steucke et al., 2017). For example, whether mechanical cues and stretch-activated

ion channel activity influence microtubule stability and vice versa remains unknown.

In this current study, we show that changes in microtubule stability serve as a

switch that regulates Ca^{2+} flux and VSMC volume control. Treatment with the

microtubule stabilising agent paclitaxel reduced VSMC volume on rigid hydrogels. In

contrast, treatment with the microtubule destabilising agent colchicine increased

VSMC volume on pliable hydrogels. Furthermore, we show that paclitaxel reduced

Ca^{2+} flux specifically in VSMCs on rigid hydrogels, whereas colchicine treatment

increased Ca^{2+} flux in VSMCs on pliable hydrogels. In keeping with our previous

findings, our data also implicates piezo1 in this process and depletion of piezo1

increased microtubule stability specifically on rigid hydrogels. Finally, we

demonstrate that disruption of HDAC6 increases acetylation of alpha tubulin and

induces increased volume and Ca^{2+} flux in VSMCs on pliable hydrogels. Together,

these data implicate a piezo1/Ca^{2+}/microtubule stability feedback pathway in the

regulation of VSMC matrix rigidity response.

Results

Piezo1 activity promotes microtubule destabilisation within VSMCs on rigid

hydrogels.

Previous studies have shown that matrix rigidity promotes increased VSMC

traction stress generation. To confirm that angiotensin II stimulated VSMCs seeded

on rigid hydrogels generated enhanced traction stresses, we performed traction

force microscopy. As expected, analysis confirmed that angiotensin II stimulated

VSMCs seeded on rigid hydrogels generated greater maximal and total traction

stress, compared to their counterparts on pliable hydrogels (**Supplementary Figure**
S1). Our previous study found that on pliable hydrogels, VSMCs generated
enhanced traction stresses as a result of colchicine-induced microtubule
depolymerisation. Therefore, we predicted that VSMCs would display decreased
microtubule stability on rigid hydrogels. To test this, we compared the number of
cold-stable microtubules possessed by quiescent and angiotensin II stimulated
VSMCs on pliable and rigid hydrogels. Analysis revealed that angiotensin II
stimulation resulted in a reduction of cold-stable microtubules possessed by VSMCs
on pliable and rigid hydrogels (**Figure 1 A and B**). However, this decrease was only
significantly different on rigid hydrogels (**Figure 1A and B**).

We have previously shown that piezo1 is an essential mediator of the
enhanced VSMC volume response on rigid hydrogels. We next utilised a siRNA
mediated knockdown approach to investigate whether piezo1 was required for this
change in microtubule stability. Consistent with our previous studies, western blot
analysis confirmed that piezo1 was efficiently depleted in VSMCs by piezo1 specific
siRNA compared to non-targeting scrambled siRNA (**Figure 1C**). Analysis of
microtubule stability revealed that piezo1 depleted, angiotensin II stimulated VSMCs
possessed similar numbers of cold stable microtubules on pliable hydrogels as their
scrambled control counterparts (**Figure 1 D and E**). In contrast, piezo1 depleted
angiotensin II stimulated VSMCs on rigid hydrogels possessed increased numbers of
cold-stable microtubules compared to their scrambled control counterparts (**Figure D**
and E). We have previously shown that piezo1 activation promotes Ca^{2+} influx in
VSMCs on rigid hydrogels. To confirm that Ca^{2+} influx was driving microtubule
destabilisation, we performed a cold-stable microtubule assay in the presence or
absence of extracellular Ca^{2+}. The absence of extracellular Ca^{2+} had no effect on the
number of cold-stable microtubules detected in angiotensin II stimulated VSMCs on
pliable hydrogels (**Figure 1F and G**). In contrast, the number of cold-stable
microtubules increased within angiotensin II stimulated VSMCs on rigid hydrogels
when extracellular Ca^{2+} was absent (**Figure 1F and G**). This suggests that piezo1-
mediated Ca^{2+} influx decreases microtubule stability on rigid hydrogels following
angiotensin II stimulation.

Microtubule stability influences isolated smooth muscle cell volume following contractile agonist stimulation.

The above data shows that on rigid substrates, VSMCs generate increased traction stresses and possess decreased microtubule stability. We have previously shown that VSMCs on rigid hydrogels swell and possess increased cell area and volume. We next hypothesised that decreased microtubule stability was driving the enhanced VSMC volume response on rigid hydrogels. To test this, we utilised our screening assay described previously. Quiescent VSMCs were seeded onto pliable and rigid hydrogels and pre-treated with increasing concentrations of microtubule stabilisers prior to angiotensin II stimulation. Treatment with either paclitaxel or epothilone B had no effect on the contractile response of VSMCs seeded on pliable hydrogels (**Figure 2A, B, D and Supplementary Figure S2**). In contrast, increasing concentrations of either microtubule stabiliser was sufficient to prevent the increase in VSMC area observed following angiotensin II stimulation on rigid hydrogels (**Figure 2 A, C, D and Supplementary Figure S2**). Given that microtubule stabilisation prevented VSMC enlargement following angiotensin II stimulation on rigid hydrogels, we next hypothesised that microtubule destabilisation would trigger VSMC enlargement in angiotensin II stimulated VSMCs on pliable hydrogels. VSMCs were pre-treated with increasing concentrations of the microtubule destabilisers colchicine or nocodazole, and then stimulated with angiotensin II. On pliable hydrogels, VSMCs treated with either colchicine or nocodazole displayed increased cell area following angiotensin II stimulation (**Figure 2E, F, H and Supplementary Figure S3**). In contrast, treatment with the microtubule destabilisers had no effect on VSMCs seeded on rigid hydrogels (**Figure 2E, G, H and Supplementary Figure S3**). All microtubule targeting agents were used at concentrations that did not cause cell death, as confirmed through a viability assay for concentrations of epothilone B and nocodazole (**Supplementary Figure S4**) or previously for paclitaxel and colchicine (Ahmed et al., 2022). We next confirmed that paclitaxel and colchicine treatments were inducing microtubule stabilisation and destabilisation respectively. Quiescent VSMCs seeded on pliable and rigid hydrogels were pretreated with microtubule targeting agents prior to angiotensin II stimulation. Analysis confirmed that paclitaxel treatment (1 nM) resulted in increased numbers of cold stable microtubules in angiotensin II stimulated VSMCs on both rigid and pliable

hydrogels (**Supplementary Figure S5**). Colchicine treatment (100 nM) resulted in

reduced numbers of cold stable microtubules in angiotensin II stimulated VSMCs on

pliable hydrogels but did not significantly alter the number of cold stable microtubules

in angiotensin II treated VSMCs on rigid hydrogels (**Supplementary Figure S5**).

Having determined that microtubule stability regulated changes in VSMC

area, we then sought to confirm its regulation of VSMC volume. Quiescent VSMCs

were pre-treated with paclitaxel (1 nM) or colchicine (100 nM), prior to angiotensin II

stimulation and confocal microscopy was used to assess changes in cell volume.

Treatment with the microtubule stabiliser, paclitaxel had no effect on the volume of

VSMCs seeded on pliable hydrogels (**Figure 3A-D**). Whereas, on rigid hydrogels,

microtubule stabilisation prevented the angiotensin II induced expansion of VSMC

volume (**Figure 3A-D**). Finally, microtubule destabilisation, via colchicine pre-

treatment, increased VSMC volume on pliable hydrogels following angiotensin II

stimulation (**Figure 3D-F**) but had no additional effect on VSMC volume on rigid

hydrogels (**Figure 3D-F**).

Changes in microtubule stability alter calcium ion flux.

The above data confirm that microtubule stability influences VSMC volume

control. Previously, we have identified piezo1 mediated Ca^{2+} influx as a regulator of

VSMC volume control. Therefore, we next hypothesised that microtubule stability

was altering Ca^{2+} flux in VSMCs. To test this, quiescent VSMCs on rigid and pliable

hydrogels were loaded with the Ca^{2+} indicator Fluo-4 prior to angiotensin II

treatment. Fluorescence video time lapse microscopy was used to measure changes

in Fluo-4 fluorescence. Analysis supported our previous findings showing that Ca^{2+}

flux was heightened and prolonged in VSMCs on rigid hydrogels compared to

VSMCs on pliable hydrogels (**Figure 4A and B**). Colchicine treatment resulted in

prolonged Ca^{2+} flux in VSMCs on pliable hydrogels but had little effect on Ca^{2+} flux in

VSMCs on rigid hydrogels (**Figure 4A and B**). In contrast, paclitaxel treatment

reduced the Ca^{2+} flux in VSMCs on rigid hydrogels but had no effect on Ca^{2+} flux in

VSMCs on pliable hydrogels (**Figure 4A and B**).

HDAC6 disruption increases microtubule acetylation and induces enhanced VSMC volume response on pliable hydrogels.

Although controversial, alpha tubulin K40 acetylation is proposed to alter the mechanical properties of microtubules. We next hypothesised that increased tubulin acetylation would influence the VSMC matrix rigidity response. To test this, we utilised tubastatin to inhibit HDAC6 deacetylation of the K40 position. Western blot analysis confirmed that VSMCs exposed to an increasing concentration of tubastatin possessed increased levels of acetylated alpha tubulin (**Supplementary Figure S6A and B**). Next, quiescent VSMCs were pretreated with tubastatin prior to angiotensin II stimulation and confocal microscopy was used to assess changes in the cell volume. Tubastatin treated VSMCs possessed increased volume compared to their vehicle treated counterparts on pliable hydrogels (**Figure 5A-D**). Tubastatin treatment had no effect on VSMC volume on rigid hydrogels (**Figure 5A-D**). To confirm that this altered response was driven by HDAC6 disruption, we performed siRNA mediated HDAC6 depletion experiments. WB confirmed HDAC6 was efficiently depleted in VSMCs by 2 independent siRNA oligomers (**Supplementary Figure S6C and D**). Analysis of z-stacks captured by confocal microscopy confirmed that HDAC6 depleted, angiotensin II stimulated VSMCs possessed increased volume on pliable hydrogels, compared to their scrambled control treated counterparts (**Figure 5E-H**). HDAC6 knockdown had no effect on VSMC volume on rigid hydrogels (**Figure 5E-H**). Due to the increased volume of HDAC6 disrupted VSMCs on pliable hydrogels, we next predicted that increased tubulin acetylation was altering VSMC microtubule stability and Ca^{2+} flux. To test these possibilities, we firstly performed cold-stable microtubule stability assays in tubastatin treated VSMCs on pliable and rigid hydrogels. However, the number of cold-stable microtubules remained unaffected by tubastatin treatment (**Figure 6A and B**). Secondly, quiescent VSMCs on rigid and pliable hydrogels were loaded with the Ca^{2+} indicator Fluo-4 prior to angiotensin II treatment. Fluorescence video time lapse microscopy was used to measure changes in Fluo-4 fluorescence. Tubastatin treatment resulted in prolonged Ca^{2+} flux in VSMCs on pliable hydrogels but had little effect on Ca^{2+} flux in VSMCs on rigid hydrogels (**Figure 6E and F**). Finally, we investigated whether matrix rigidity influenced acetylated K40 α-tubulin levels. However, WB analysis

291 revealed that acetylated K40 levels were similar in the lysates of VSMCs grown on

292 12 and 72 kPa hydrogels (**Supplementary Figure S7**).

Discussion

Despite much research, our understanding of the mechanisms driving VSMC

dysfunction and its contribution to decreased aortic compliance in ageing and CV

disease remains limited. Previous studies have shown that enhanced matrix stiffness

promotes the dedifferentiation of VSMCs, downregulating contractile markers whilst

increasing the expression of proliferative genes (Brown et al., 2010; Nagayama and

Nishimiya, 2020; Sazonova et al., 2015; Xie et al., 2018). Increases VSMC

migrational capacity, adhesion, proliferation and volume have also been reported

(Brown et al., 2010; Krajnik et al., 2023; Nagayama and Nishimiya, 2020; Rickel et

al., 2020; Sazonova et al., 2015; Wong et al., 2003). Furthermore, in response to

matrix stiffness, VSMC reorganise their actin cytoskeleton and generate enhanced

traction stresses, a finding we recapitulate in this study(Brown et al., 2010; Petit et

al., 2019; Sanyour et al., 2019; Sazonova et al., 2015).

Our findings show that changes in the tensegrity equilibrium defines VSMC

behaviour in response to matrix rigidity. In this model, the microtubules serve as

compression bearing struts that resist actomyosin generated strain (Brangwynne et

al., 2006; Stamenović, 2005). Healthy VSMC behaviour is therefore a balance

between microtubule stability and actomyosin activity. Actomyosin generated tension

drives VSMC contraction and microtubules maintain VSMC morphology, protecting

against strain induced cellular damage (Stamenović, 2005). The tensegrity model

predicts that disruption of the microtubule cytoskeleton alters the equilibrium of these

stresses and results in increased VSMC traction stress generation, a hypothesis

previously confirmed by both wire myography and traction force microscopy (Ahmed

et al., 2021; Zhang et al., 2000). We show that in rigid environments, the VSMC

tensegrity equilibrium becomes unbalanced. Contractile agonist stimulation

increases VSMC traction stress generation and reduces the number of cold-stable

microtubules, resulting in deregulating VSMC morphology and enlargement of VSMC

volume. Importantly, the altered tensegrity equilibrium can be induced by microtubule

destabilising agents on pliable hydrogels and restored by treatment with agents that

stabilise microtubules on rigid hydrogels. These data suggest that microtubules and

the tensegrity equilibrium are key regulators of VSMC function and dysfunction.

Given the prevalence of increased aortic wall stiffness in ageing and disease, our

findings suggest that altered tensegrity equilibrium is a key driver of VSMC

dysfunction under these conditions (Tsamis et al., 2013). We propose that changing

this equilibrium could be a mechanistic target for manipulating VSMC function. Using

microtubule agents clinically is not a viable option to modulate this equilibrium, so a

better understanding of the molecular mechanisms that regulate this equilibrium is

now required to test this idea further.

In this study, we used Ca^{2+} flux as a readout of stretch-activated channel

function. One of these, piezo1, is emerging as an important regulator of VSMC

dysfunction (Johnson et al., 2024; Luu et al., 2023; Qian et al., 2022; Retailleau et

al., 2015; Swiatlowska et al., 2023). Our findings further implicate piezo1, as a critical

regulator of VSMC behaviour and suggest that changes in piezo1 opening drive

altered VSMC behaviour in response to enhanced matrix stiffness. In this study, we

show that VSMCs on rigid hydrogels display prolonged Ca^{2+} flux, increased traction

stresses and reduced microtubule stability. The relationship between Ca^{2+} flux and

microtubule stability were reciprocal, and changes in microtubule stability altered

Ca^{2+} flux. We have previously shown that changes in microtubule stability also

altered VSMC traction stress generation. Although untested, we predict that

microtubule stability alters the ability of VSMCs to withstand the deformational forces

placed on the plasma membrane, resulting in changes in membrane tension that

control stretch-activated ion channel opening. Therefore, stretch-activated ion

channel mediated Ca^{2+} influx serves as a mechanistic switch that triggers loss of the

microtubule bearing struts and shifts the VSMC tensegrity equilibrium. We still do not

understand the precise mechanisms that drive piezo1 mediated microtubule

destabilisation. Millimolar Ca^{2+} concentrations spontaneously induce microtubule

destabilisation in vitro, but cellular concentrations are much lower than this. We

envisage 3 possibilities: 1) increased Ca^{2+} flux induces changes in the ability of

microtubule stabilising proteins to associate with microtubules, resulting in

destabilisation; 2) increased mechanical damage induces microtubule catastrophe;

or 3) increased Ca^{2+} reduces microtubule growth/polymerisation.

The role of acetylation in microtubule stability, rigidity and resistance to
mechanical damage remains controversial with many conflicting reports. We show
that increased microtubule K40 acetylation stimulated both enhanced Ca^{2+} flux and
VSMC volume on pliable hydrogels. This suggests that changes in microtubule post
translational modification is also a factor in regulating VSMC matrix rigidity response.
Despite this finding, K40 acetylation remained unaltered in VSMCs on pliable and
rigid hydrogels in our study. This suggests that K40 acetylation does not play a role
in the altered microtubule stability observed during VSMC matrix rigidity response.
However, HDAC6 inhibition failed to alter microtubule stability and appeared to
uncouple the tensegrity equilibrium from the Ca^{2+} flux response on pliable hydrogels.
Despite this uncoupling, the increased Ca^{2+} appeared to activate downstream
signalling to enhance VSMC volume. HDAC6 has numerous targets, and we predict
that this uncoupling is being driven by increased acetylation of other targets that
remain unknown. Importantly, these findings suggest that, in addition to the
tensegrity equilibrium and our previously identified volume control pathway, other
unidentified mechanisms also contribute to the VSMC response to matrix rigidity.
Further research is now needed to identify these unknown mechanisms for a more
complete understanding of the VSMC matrix rigidity response.

In the healthy aortic wall, VSMCs adopt a quiescent, contractile differentiated
phenotype. During ageing and disease, VSMCs downregulate contractile marker
expression and modulate their phenotype towards disease relevant phenotypes
(Pedroza et al., 2020; Wirka et al., 2019; Worssam et al., 2023). An important
limitation of our study is that we have used dedifferentiated isolated VSMCs. VSMCs
are known to possess multiple stretch-activated ion channels, and these can vary
depending on phenotype. We have previously shown that piezo1 is upregulated in
disease relevant phenotypes (Johnson et al., 2024). It remains unknown whether the
VSMC tensegrity equilibrium can be altered in healthy contractile differentiated
VSMCs, but microtubule agents alter VSMC contractile function in isolated aortic
rings, suggesting that this is the case (Zhang et al., 2000). In this study, we implicate
piezo1 in the defective matrix rigidity induced tensegrity equilibrium of isolated
VSMCs. However, we cannot rule out the possibility that other stretch ion channels
can induce altered tensegrity equilibrium in other phenotypes. Despite this limitation,

our findings support the notion that the VSMC tensegrity equilibrium is modifiable
and can be targeted to restore VSMC morphology.

Materials and Methods

Polyacrylamide Hydrogel Preparation

Hydrogels were prepared as described previously (Johnson et al., 2024;
Minaisah et al., 2016). Briefly, glass coverslips were activated by treating with (3-
Aminopropyl)triethoxysilane for 2 minutes, washed 3x in dH_2O, then fixed in 0.5%
glutaraldehyde for 40 minutes. After fixation, coverslips were washed and left to air
dry overnight. Polyacrylamide hydrogel buffer was comprised as follows: 12 kPa –
7.5 % acrylamide, 0.15 % bis-acrylamide in dH_2O; 72 kPa – 10 % acrylamide, 0.5 %
bis-acrylamide in dH_2O. To prepare hydrogels for fabrication, the appropriate volume
of buffer was supplemented with 10 % APS (1:100) and TEMED (1:1,000) then
placed on a standard microscopy slide and covered by an activated coverslip. Once
set, the hydrogels were washed 3x in dH_2O, cross-linked with sulfo-SANPAH
(1:3000) under UV illumination (365 nm) for 5 minutes, then functionalised with
collagen I (0.1 mg/ml) for 10 minutes at room temperature. Hydrogel stiffness was
previously confirmed using a JPK Nanowizrd-3 atomic force microscope(Porter et al.,
2020).

Vascular Smooth Muscle Cell Culture

Human adult aortic VSMCs (passages 3-10) were purchased from Cell
Applications Inc. (354-05a). Standard VSMC culture was performed as previously
described (Ahmed et al., 2021; Johnson et al., 2024). VSMCs were seeded onto
polyacrylamide hydrogels in basal media (Cell Applications Inc, Cat# 310-500), 18
hours prior to the beginning of the experiment to induce quiescence. Briefly, VSMCs
were pre-treated for 30 minutes, prior to co-treatment with the contractile agonist,
angiotensin II (10 µM) for an additional 30 minutes. Experimental specific

concentrations are provided in the corresponding figure legends. For experiments performed in growth media, VSMCs were seeded onto hydrogels and incubated overnight. Pharmacological agents were added and VSMCs were incubated for a further 18 hours prior to fixation. Please see Table1 below for details of compounds used in this study.

Table 1 – Concentrations and details of compounds used in this study.

Compound	Product Code	Supplier	Vehicle	Concentration Range Tested	Working Concentration
Angiotensin II	A9525	Merck	dH$_2$0		10 µM
Colchicine	C9754	Merck	DMSO	0.1 - 1000nM	100 nM
Paclitaxel	T7402	Merck	DMSO	0.001 - 10 nM	1 nM
Nocodazole	ab120630	Abcam	DMSO	0.001 - 10 nM	-
Epothilone B	E2656	Merck	DMSO	0.001 - 10 µM	-
Tubastatin A	#6270	Tocris	DMSO	0.01 – 1 µM	1 µM
A23187		Tocris	DMSO		20 µM

siRNA Knockdown

VSMC siRNA transfection was performed using HiPerFect (Qiagen) as per manufacturer's instructions, the day before cells were seeded onto hydrogels. VSMCs were transfected with either scrambled siRNA control, piezo1, or HDAC6 targeting siRNA (listed below) oligonucleotides. The next afternoon, cells were seeded onto hydrogels as above, serum was withdrawn overnight to induce quiescence and the next morning VSMCs were stimulated with angiotensin II (10 µM) for 30 minutes prior to fixation and downstream immunofluorescent analysis.

Piezo1 siRNA:

siRNA #5 – CCGCGTCTTCCTTAGCCATTA

siRNA #7 – CGGCCGCCTCGTGGTCTACAA

HDAC6 siRNA:

siRNA #5 - CACCGTCAACGTGGCATGGAA

siRNA #10 - CCGGAGGGTCCTTATCGTAGA

Western Blotting

Western blotting was performed as previously described (Ragnauth et al.,

2010). When looking for piezo1 expression specifically, lysates were run on a

TruPAGE precast 4-20 % gradient gel (Sigma-Aldrich) at 120 V for 2 hours. Protein

was transferred onto PVDF membrane at 30 V for 3 hours prior to the membrane

being blocked in 5 % milk/TBST. The following antibodies were used: anti-piezo1

(1:500) (Novus Cat# NBP1-78537, RRID:AB_11003149), anti-GAPDH (1:4000) (Cell

Signaling Technology Cat# 2118, RRID:AB_561053), anti-acetyl K40 alpha tubulin

(Cell signaling technology Cat #3971, RRID:AB_2210204), anti-total alpha tubulin

(Cell Signaling Technology Cat#3873, RRID:AB_1904178), anti-HDAC6 (D2E5) (Cell

Signaling Technology Cat# 7558, RRID:AB_10891804), anti-Rabbit-HRP (1:2000)

(Sigma-Aldrich Cat# GENA934, RRID:AB_2722659) and anti-Mouse-HRP (1:2000)

(Sigma-Aldrich Cat# AP160P, RRID:AB_92531). 12 v 72 kPa lysates were repeated

in triplicate for each individual experimental repeat.

Immunofluorescence and VSMC Area/Volume Analysis

Cells were fixed in 4% paraformaldehyde for 10 minutes, permeabilised with

0.5 % v/v NP40 for 5 minutes, then blocked with 3 % w/v BSA/PBS for 1 hour.

Primary staining against lamin A/C (1:200) (Sigma-Aldrich Cat# SAB4200236,

RRID:AB_10743057) was performed overnight at 4 $^\square$C in 3% BSA/PBS. Secondary

antibody staining was performed using the appropriate Alexa Fluor™ 488 antibody

(1:400) (Thermo Fisher Scientific Cat# A-11001, RRID:AB_2534069) in the dark for

2 hours. F-actin was visualised using Rhodamine Phalloidin (1:400) (Thermo Fisher

Scientific Cat# R145). Images were captured at 20x magnification using a Zeiss

LSM980-Airyscan confocal microscope. Cell area and volume was measured using

FIJI, open-source software as described previously (Johnson et al., 2024; Schindelin
et al., 2012).

Traction Force Microscopy

VSMCs were seeded onto polyacrylamide hydrogels containing 0.5 µm red
fluorescent (580/605) FluoSpheres (1:1000) (Invitrogen). Following angiotensin II
stimulation (30 minutes), cell lysis was achieved by the addition of 0.5 % v/v Triton
X-100. Images were captured at 20x magnification before and after lysis at 2-minute
intervals using a Zeiss Axio Observer live cell imaging system. Drift was corrected
using the ImageJ StackReg plugin and traction force was calculated using an
ImageJ plugin that measured FluoSphere displacement (Tseng et al., 2012). Briefly,
bead displacement was measured using the first and last image of the movie
sequence. The cell region was determined by overlaying the traction map with the
phase image, selecting the cell traction region with an ROI and extracting the traction
forces in each pixel using the XY coordinate function in FIJI (Ahmed et al., 2021;
Porter et al., 2020).

Cold-Stable Microtubule Stability Assay

Cold-stable microtubules were identified as per previous studies (Atkinson et
al., 2018). Following treatment, cells were placed on ice for 15 minutes before being
washed once with PBS and twice with PEM buffer (80 µM PIPES pH 6.8, 1 mM,
EGTA, 1 mM, $MgCl_2$, 0.5 %, Triton X-100, and 25 % (w/v) glycerol) for 3 minutes.
Cells were fixed in ice-cold methanol for 20 minutes then blocked with 3 % BSA/PBS
for 1 hour. Microtubules were visualised by staining for α-tubulin (1:200) (Cell
Signalling Technology Cat# 3873, RRID:AB_1904178) whilst cell nuclei were
visualised using a mounting medium containing DAPI. Images were captured at 40x
magnification using a Zeiss AxioPlan 2ie microscope.

Cell Viability Assay

Cell viability was determined using a RealTime-Glo™ MT Cell Viability Assay,

as per manufactures instructions. Briefly, 5,000 cells per well seeded in a 96-well

plate and exposed to a range of drug concentrations for 1 hour. Luminescence was

subsequently measured using a Wallac EnVision 2103 Multilabel Reader

(PerkinElmer).

Fluo-4 Calcium imaging

Cells seeded on 12 and 72kPa hydrogels and incubated in basal medium for

48 hours. Cells were loaded with 3 µM Fluo-4 AM (ThermoFisher Scientific cat#

F14201) diluted in basal media for 30 minutes. Paclitaxel, colchicine and tubastatin

were used at working concentrations listed in Supplementary Table 1. Cells were

co-treated with vehicle control or compounds of interest during the Fluo-4 loading

step. Cells were washed in PBS and returned to basal media prior to imaging. Cells

were placed onto a Zeiss Axiovert 200M inverted microscope stage and images

were captured every 500 msec. Cells were imaged for 5 minutes prior to angiotensin

II (10 µM) stimulation. Cells were subsequently imaged for a further 20 minutes after

addition of angiotensin II. Finally, the ionophore A23187 was added for a further 5

minutes to confirm the calcium imaging had worked. Images were analysed in

ImageJ by selecting a ROI and extracting the fluorescence intensities for each time

point using the plot Z-axis profile option. Background was subtracted and the ΔF and

$\Delta F/F_0$ values were calculated and analysed using GraphPad prism.

Statistical Analysis

Statistical analysis was performed using GraphPad Prism 9. Results are

presented as mean ± SEM, with individual data points shown. The number of

independent repeats performed, and total number of cells analysed per experiment

are detailed in the corresponding figure legend. To compare more than two

conditions a one-way ANOVA was performed, with a Tukey's multiple comparison

post-hoc test performed. Concentration response curves are presented as mean ±

SEM plotted on a semi-logarithmic scale. Comparisons between concentration

ranges on different hydrogel stiffness were performed using a two-way ANOVA

followed by Sidak's post-hoc test. Differences between conditions were considered
statistically significant when P < 0.05.

Author Contributions: RTJ, FW, RS, TM, CT, AB, OS, YB and DTW were
responsible for performing experiments and analysing data. DTW was responsible
for the conceptualisation; RTJ, SB, YB and DTW were responsible for the
methodology of the study. SB, CJM and YB reviewed and edited this manuscript.
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Figure legends

Figure 1. Extracellular calcium influx via piezo1 ion channels triggers

microtubule destabilisation. A) Representative images of isolated VSMCs cultured

on 12 or 72 kPa polyacrylamide hydrogels with/without angiotensin II treatment.

Cold-stable microtubules, (α-tubulin, green) and nuclei (DAPI, blue). Scale bar = 50

μm. B) Graph shows number of cold-stable microtubules per cell on pliable and rigid

hydrogels with/without angiotensin II treatment. Graph represents combined data

from 3-independent experiments, analysing >56 cells per condition, and black dots

mark mean data for each independent repeat. C) Representative WB confirming

efficient siRNA-mediated piezo1 depletion in VSMCs (siCon = non-targeting

scrambled siRNA, siP#5 and siP#7 = independent piezo1 targeting siRNA #5 and

#7). D) Representative images of angiotensin II stimulated siCon, siP#5 and siP#7

treated VSMCs cultured on 12 or 72 kPa polyacrylamide hydrogels. Cold-stable

microtubules, (α-tubulin, green) and nuclei (DAPI, blue). Scale bar = 50 μm. E)

Graph shows number of cold-stable microtubules per cell on 12 kPa and 72 kPa

hydrogels in angiotensin II stimulated siCon, siP#5 and siP#7 VSMCs. Graph

represents combined data from 3-independent experiments, analysing >65 cells per

condition, and black dots mark mean data for each independent repeat. F)

Representative images of angiotensin II stimulated VSMCs cultured on 12 or 72 kPa

polyacrylamide hydrogels in calcium containing or calcium free media. Cold-stable

microtubules, (α-tubulin, green) and nuclei (DAPI, blue). Scale bar = 50 μm. G)

Graph shows number of cold-stable microtubules per cell on 12 kPa and 72 kPa

hydrogels in angiotensin II stimulated VSMCs in the presence/absence of

extracellular calcium. Graph represents combined data from 3-independent

experiments, analysing >71 cells per condition, and black dots mark mean data for

each independent experimental repeat. Significance determined using a two-way

ANOVA followed by Tukey's post-hoc test. (* = p < 0.05, ** = p < 0.01 and *** = p <

0.001. Error bars represent ± SEM)

Figure 2. Microtubule stability influences VSMC spreading following

angiotensin II stimulation. A) Representative images of isolated VSMCs cultured

on 12 or 72 kPa polyacrylamide hydrogels pretreated with a concentration range of

paclitaxel prior to angiotensin II stimulation. Actin cytoskeleton (purple) and lamin

A/C labelled nuclei (green). Scale bar = 50 µm. Graphs show VSMC area on B) 12

kPa hydrogels, C) 72 kPa hydrogels, and D) 12 kPa and 72 kPa hydrogels. Graphs

represent the combined data from 3-independent experiments, analysing <80 cells

per condition. Black dots represent mean data from each individual experimental

repeat. E) Representative images of isolated VSMCs cultured on 12 or 72 kPa

polyacrylamide hydrogels pretreated with a concentration range of colchicine prior to

angiotensin II stimulation. Actin cytoskeleton (purple) and lamin A/C labelled nuclei

(green). Scale bar = 50 µm. Graphs show VSMC area on F) 12 kPa hydrogels, G) 72

kPa hydrogels, and H) 12 kPa and 72 kPa hydrogels. Graphs represent the

combined data from 3-independent experiments, analysing <80 cells per condition.

Black dots represent mean data from each individual experimental repeat.

Significance determined using one-way ANOVA (graphs B, C, F and G) or two-way

ANOVA (Graphs D and H) followed by Tukey's post-hoc test. (* = p < 0.05. Error

bars represent ± SEM)

Figure 3. VSMC volume is regulated by microtubule stability. A) Representative

images of isolated VSMCs cultured on 12 or 72 kPa polyacrylamide hydrogels

pretreated with vehicle control (dH$_2$0) or paclitaxel prior to angiotensin II stimulation.

Actin cytoskeleton (purple) and DAPI labelled nuclei (green). Top panel shows

representative XY images of VSMC area, scale bar = 50 µm. Bottom panel shows

representative XZ images of VSMC height, scale bar = 20 µm. Graphs show VSMC

B) area, C) volume and D) height and represent the combined data from 3

independent experiments with ≥45 cells analysed per condition. Mean data from

each individual repeat of the 3 independent experiments is shown by a black dot.

Significance was determined using two-way ANOVA followed by Tukey's post-hoc

test. E) Representative images of isolated VSMCs cultured on 12 or 72 kPa

polyacrylamide hydrogels pretreated with vehicle control (dH$_2$0) or colchicine prior to

angiotensin II stimulation. Actin cytoskeleton (purple) and DAPI labelled nuclei

(green). Top panel shows representative XY images of VSMC area, scale bar = 50

µm. Bottom panel shows representative XZ images of VSMC height, scale bar = 20

µm. Graphs show VSMC F) area, G) volume and H) height and represent the

combined data from 3 independent experiments with ≥42 cells analysed per

condition. Mean data from each individual repeat of the 3 independent experiments

is shown by a black dot. Significance was determined using two-way ANOVA

followed by Tukey's test. (* = $p < 0.05$, ** = $p < 0.01$ and *** = $p < 0.001$. Error bars

represent ± SEM)

Figure 4. Microtubule stability influences Ca^{2+} flux in VSMCs. A) Graphs show

representative ΔF/F$_0$ values over time for vehicle control, paclitaxel and colchicine

pre-treated angiotensin II stimulated Fluo-4 loaded VMSCs on 12 and 72 kPa

hydrogels. Each graph represents the mean combined data of 9 individual fields of

view combined from 3 individual experiments. B) Graph shows area under the curve

values (ΔF/F$_0$ x T). Mean data for individual repeats of the 3 independent

experiments are shown as black dots. Significance was determined using two-way

ANOVA followed by Tukey's multiple comparison test. (** = $p < 0.01$ and *** = $p <$

0.001. Error bars represent ± SEM)

Figure 5. HDAC6 disruption increases microtubule acetylation and enhances

VSMC volume. A) Representative images of isolated VSMCs cultured on 12 or 72

kPa polyacrylamide hydrogels pretreated with vehicle control (dH$_2$0) or tubastatin

prior to angiotensin II stimulation. Actin cytoskeleton (purple) and DAPI labelled

nuclei (green). Top panel shows representative XY images of VSMC area, scale bar

= 50 µm. Bottom panel shows representative XZ images of VSMC height, scale bar

= 20 µm. Graphs show VSMC B) area, C) volume and D) height and represent the

combined data from 3 independent experiments with ≥41 cells analysed per

condition. Mean data from each individual repeat of the 3 independent experiments

is shown by a black dot. Significance was determined using two-way ANOVA

followed by Tukey's test. E) Representative images of isolated VSMCs cultured on

12 or 72 kPa polyacrylamide hydrogels treated with scrambled (siCon) or HDAC6

targeting (HDAC6 #5 and #10) siRNA oligos prior to angiotensin II stimulation. Actin

cytoskeleton (purple) and DAPI labelled nuclei (green). Top panel shows

representative XY images of VSMC area, scale bar = 50 μm. Bottom panel shows

representative XZ images of VSMC height, scale bar = 20 μm. Graphs show VSMC

F) area, G) volume and H) height and represent the combined data from 3

independent experiments with ≥30 cells analysed per condition. Mean data from

each individual repeat of the 3 independent experiments is shown by a black dot.

Significance was determined using two-way ANOVA followed by Tukey's test. (* = p

< 0.05 and ** = p < 0.01. Error bars represent ± SEM)

Figure 6. Tubastatin treatment alters Ca^{2+} flux in VSMCs. A) Representative

images of isolated angiotensin II stimulated VSMCs pre-treated with vehicle control

or tubastatin on 12 or 72 kPa polyacrylamide. Cold-stable microtubules, (α-tubulin,

green) and nuclei (DAPI, blue). Scale bar = 50 μm. B) Graph shows number of cold-

stable microtubules per cell on pliable and rigid hydrogels. Graph represents

combined data from 6-independent experiments, analysing >60 cells per condition,

and black dots mark mean data for each independent repeat. C) Graphs show

representative $\Delta F/F_0$ values over time for vehicle control and tubastatin pre-treated

angiotensin II stimulated Fluo-4 loaded VMSCs on 12 and 72 kPa hydrogels. Each

graph represents the mean combined data of 9 individual fields of view combined

from 3 individual experiments. D) Graph shows area under the curve values ($\Delta F/F_0$ x

T). Mean data for individual repeats of the 3 independent experiments are shown as

black dots. Significance was determined using two-way ANOVA followed by Tukey's

multiple comparison test. (* = p < 0.05. Error bars represent ± SEM)

Figure 1

Figure 2

Figure 3

A.

B.

Figure 4

Figure 5

Figure 6