

# Non-receptor tyrosine kinases and the actin cytoskeleton in contractile vascular smooth muscle

Jacqueline Ohanian, Maria Pieri and Vasken Ohanian

*Institute of Cardiovascular Sciences, Manchester Academic Health Services Centre, University of Manchester, Manchester, UK*

**Abstract** The contractility of vascular smooth muscle cells within the walls of arteries is regulated by mechanical stresses and vasoactive signals. Transduction of these diverse stimuli into a cellular response occurs through many different mechanisms, one being reorganisation of the actin cytoskeleton. In addition to a structural role in maintaining cellular architecture it is now clear that the actin cytoskeleton of contractile vascular smooth muscle cells is a dynamic structure reacting to changes in the cellular environment. Equally clear is that disrupting the cytoskeleton or interfering with its rearrangement, has profound effects on artery contractility. The actin cytoskeleton associates with dense plaques, also called focal adhesions, at the plasma membrane of smooth muscle cells. Vasoconstrictors and mechanical stress induce remodelling of the focal adhesions, concomitant with cytoskeletal reorganisation. Recent work has shown that non-receptor tyrosine kinases and tyrosine phosphorylation of focal adhesion proteins such as paxillin and Hic-5 are important for actin cytoskeleton and focal adhesion remodelling and contraction.

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**Corresponding author** J. Ohanian: Cardiovascular Research, University of Manchester, Core Technology Facility 3rd Floor, 46 Grafton Street, Manchester M13 9NT, UK. Email: johanian@manchester.ac.uk

**Abbreviations** CK, cytoskeleton; FAs, focal adhesions; FAK, focal adhesion kinase; Hic-5, hydrogen peroxide inducible clone-5; Hsp27, heat shock protein 27; LIM protein, Lin11, Isl-1 and Mec-3 domain protein; VSMCs, vascular smooth muscle cells.

## Introduction

The maintenance of normal peripheral vascular resistance and perfusion of vital organs depends on the structure and contractile tone of small arteries within the vasculature. Changes in these properties of small blood vessels lead to cardiovascular disease. Smooth muscle cells within the artery wall respond to external stimuli such as hormones and stress (increased pressure or flow) adjusting their level of tone to maintain perfusion and resistance within normal levels. However, in response to prolonged mechanical stress, for example maintained high blood pressure, vascular smooth muscle cells (VSMCs) exhibit exaggerated contractility and remodelling of the vessel wall, as occurs in hypertension or migration and proliferation as seen in atherosclerosis

(Castorena-Gonzalez *et al.* 2014). An important role for the actin cytoskeleton (CK) in the smooth muscle responses contraction, proliferation and migration, is now recognised (reviewed in Gerthoffer & Gunst, 2001; Gunst & Zhang, 2008; Yamin & Morgan, 2012). Cytosolic tyrosine kinases are emerging as important regulators of the actin CK, through tyrosine phosphorylation of focal adhesion proteins such as vinculin and paxillin (Gunst & Zhang, 2008). However, data describing actin CK remodelling and the signalling molecules involved are derived from multiple sources including smooth muscle cells in culture, freshly dispersed VSMCs settled on glass, or matrix and tissues which are composed of multiple cell types. The phenotypic properties and behaviour of the VSMCs in these different preparations may vary, and care should be taken when extrapolating between different

**Jacqueline Ohanian** is currently Senior Lecturer in the Institute of Cardiovascular Sciences at the University of Manchester, UK. Her research interests focus on understanding the signalling pathways involved in the regulation of small artery contractility. **Vasken Ohanian** is also currently Senior Lecturer in the Institute of Cardiovascular Sciences at the University of Manchester. His work focuses on regulation of the actin cytoskeleton by tyrosine kinases and mitogen-activated protein kinases in vascular smooth muscle cells.



models. In this review we will describe our current understanding of the regulation of actin CK dynamics in contractile VSMCs and tissue, focusing on the role of tyrosine kinases and adhesion site adaptor proteins.

### Actin cytoskeleton

Until recently, the CK was perceived as a relatively fixed structure; however, this view has now been completely revised and it is recognised that the CK is a highly dynamic structure constantly assembling and disassembling in response to external stimuli. Rapid reorganisation of the CK occurs through the reversible polymerisation, of monomeric globular actin (G-actin) to the polymerised filamentous form (F-actin), markedly altering cellular properties. The organisation of the CK is regulated by signalling pathways that target proteins that influence actin assembly, in particular the stress-activated protein kinases, small heat shock proteins, cytosolic tyrosine kinases and LIM (Lin11, Isl-1 and Mec-3 domain) proteins (Gerthoffer & Gunst, 2001; Gunst & Zhang, 2008).

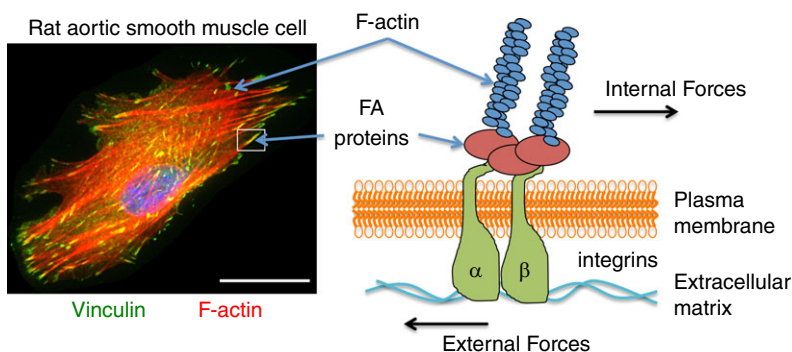
### Actin cytoskeleton in smooth muscle

Contractile VSMCs have a highly organised CK that is required for the transmission of force. In fully differentiated smooth muscle cells in tissues the CK is attached to the cell membrane at dense plaques which are thought to act as mechanosensors transducing integrin-mediated signals to the CK (Gerthoffer & Gunst, 2001; Kim *et al.* 2008; Lehman & Morgan, 2012) (Fig. 1). These membrane-localised dense plaques are multiprotein complexes that cluster many of the same proteins (vinculin,  $\alpha$ -actinin and focal adhesion kinase (FAK)) found in focal adhesions of migrating cells in culture, and are distinct from the classical dense bodies associated with the contractile machinery in the interior of the VSMCs (reviewed in Owens *et al.* 2004). Smooth muscle dense plaques are also called adhesion sites or focal adhesions, and will be referred to as focal adhesions (FAs) throughout this review. Recent studies in freshly dispersed differentiated VSMCs have led to the proposal that the

cortical actin CK and the actin contractile filaments are formed from different actin isoforms (reviewed in Yamin & Morgan, 2012). In this model  $\beta$ -actin is found around intracellular dense bodies associated with the contractile apparatus (Gallant *et al.* 2011), whereas  $\alpha$ -smooth muscle actin filaments interact with myosin forming the contractile actin fibres that run longitudinally and diagonally through the cell. Non-muscle  $\gamma$ -actin is the major component of the cortical CK at the periphery of the cell and it is the  $\beta$ - and  $\gamma$ -actin networks that alter their polymerisation state in response to stimulation (Kim *et al.* 2008). It is now clear that reorganisation of the actin cytoskeleton is an essential component of the contractile response of smooth muscle cells. Increases in filamentous actin have been reported in intact arteries during myogenic contraction (Cipolla *et al.* 2002; Moreno-Dominguez *et al.* 2013; Walsh & Cole, 2013) and following vasoconstrictor stimulation (Mehta & Gunst, 1999; Bárány *et al.* 2001; Ohanian *et al.* 2005; Srinivasan *et al.* 2008). In addition, interference with actin polymerisation blocks contraction and stabilisation of actin filaments induces relaxation in vascular tissue without affecting myosin light chain phosphorylation (Boels & Pfitzer, 1992; Gerthoffer & Gunst, 2001; Ohanian *et al.* 2005; Wang *et al.* 2014). Alterations in VSMC actin CK are also implicated in physiological (reviewed in Castorena-Gonzalez *et al.* 2014) and pathological remodelling of large and small arteries (Qiu *et al.* 2010; Saphirstein *et al.* 2013; Sehgel *et al.* 2013). However, our understanding of the mechanisms that regulate contractile VSMC actin cytoskeleton dynamics is limited.

### The role of cytosolic tyrosine kinases and adhesion site remodelling in vascular smooth muscle responses

In intact small arteries we have identified two pathways activated by vasoconstrictors that are involved in regulation of the CK, p38 mitogen-activated protein kinase (p38MAPK) and cytosolic tyrosine kinase (Src and PYK2), the former through heat shock protein 27 (Hsp27) and the latter through the LIM protein paxillin and its homologue hydrogen peroxide inducible clone-5 (Hic-5) (Ohanian



**Figure 1. Vascular smooth muscle cell actin cytoskeleton and adhesion sites**

Left-hand panel: immunofluorescence image of a rat aortic smooth muscle cell cultured on collagen-coated glass. Filamentous actin (F-actin) is stained red and vinculin, a focal adhesion (FA) marker, is shown in green. Actin filaments can be seen terminating at adhesion sites. Right-hand panel: schematic diagram of an adhesion site showing sensing of external force and transmission of internal force through actin filaments and integrin focal adhesion site. The scale bar is 0 — 25  $\mu$ .

*et al.* 2001; Ward *et al.* 2002; Ohanian *et al.* 2005). Although these pathways are activated independently of each other they converge to regulate contractility through a common mechanism, promotion of F-actin formation, i.e. CK remodelling (Srinivasan *et al.* 2008). In agreement with our findings in small arteries, cytosolic tyrosine kinases and the FA proteins paxillin, zyxin and vinculin are involved in F-actin formation and contraction in response to acetylcholine in tracheal smooth muscle strips (reviewed in Gerthoffer & Gunst, 2001; Gunst & Zhang, 2008; Tang & Anfinogenova, 2008), emphasising the importance of this mechanism in smooth muscle tissue responses. In non-smooth-muscle cells in culture Src, PYK2, vinculin, zyxin, paxillin and Hic-5 are associated with FAs and are involved in assembly and disassembly of these structures (Zaidel-Bar *et al.* 2007), and activation of these proteins by vasoconstrictor hormones in smooth muscle tissues suggests that these agonists may signal to the CK through remodelling of adhesion sites. This idea is supported by two recent studies where, in A7R5 vascular smooth muscle cells, stimulation with lyso-phosphatidic acid increased cell stiffness, stress fibre formation and FA size in an Src-dependent manner (Saphirstein *et al.* 2013), and in freshly dispersed rat cremaster arteriole VSMCs where angiotensin II stimulation increased cell stiffness and cell adhesion to the extracellular matrix through actin CK and FA remodelling (Hong *et al.* 2014).

The signalling pathways and mechanisms leading to adhesion site remodelling in vascular tissues and differentiated contractile VSMCs are still unclear. Tyrosine phosphorylation is a hallmark of FA regulation and a number of cytosolic tyrosine kinases have been identified that are activated by vasoconstrictors in vascular tissue, including Src (Ohanian *et al.* 2001; Min *et al.* 2012), FAK (Min *et al.* 2012) and its homologue PYK2 (Ohanian *et al.* 2005), and Abl (Tang & Anfinogenova, 2008; Wang *et al.* 2013). Additionally, increased tyrosine phosphorylation of FA proteins paxillin (Ohanian *et al.* 2005), Hic-5 (Srinivasan *et al.* 2008) and p130 Crk-associated kinase (p130Cas) (Tang, 2009; Min *et al.* 2012) has been reported in vascular and airway smooth muscle tissue with a time course of phosphorylation consistent with the agonist-induced contractile response. Recruitment of cytosolic tyrosine kinases to FA and tyrosine phosphorylation of paxillin, Hic-5 and p130Cas regulates FA dynamics in non-smooth-muscle cells in culture (Mitra *et al.* 2005). Consistent with this, Src, PYK2, paxillin (Ohanian *et al.* 2005) and Hic-5 (Srinivasan *et al.* 2008) redistribute between cytosolic and cytoskeleton compartments following noradrenaline stimulation in intact small arteries. Inhibition of noradrenaline-induced redistribution of PYK2, paxillin and Hic-5 reduced the contractile response and actin CK remodelling without affecting myosin light chain phosphorylation (Ohanian *et al.* 2005; Srinivasan *et al.*

2008). Similarly in differentiated VSMCs, Src and p130Cas redistribute between soluble and insoluble fractions following phenylephrine stimulation and inhibition of their relocalisation reduces the rate of contraction of aortic tissue; again the reduction in contraction was independent of myosin light chain phosphorylation (Yamin & Morgan, 2012). Recently, it was shown in differentiated aortic VSMCs that following phenylephrine stimulation redistribution of FA proteins occurred through association with an endocytic recycling pathway (Poythress *et al.* 2013). Furthermore, inhibition of the endocytic pathway slowed the phenylephrine-induced rate of force development and maximum contractile response of aortic tissue (Poythress *et al.* 2013). Together these studies indicate that activation of cytosolic tyrosine kinases and phosphorylation of FA proteins occurs in response to vasoconstrictors in vascular smooth muscle tissues and cells leading to adhesion site and actin CK remodelling, and that this pathway is an essential component of the contractile response independent of myosin light chain phosphorylation (Fig. 2).

In addition to transducing internal forces to the extracellular matrix during smooth muscle cell contraction, adhesion sites are important for sensing external forces such as changes in mechanical stress or strain and matrix stiffness. Signalling through adhesion sites to the actin cytoskeleton is an important mechanism for mechanosensing in VSMCs and is implicated in arterial remodelling in response to vasoconstrictors, hypertension and increased extracellular matrix stiffness (reviewed in Martinez-Lemus *et al.* 2009; Hill & Meininger, 2012; Castorena-Gonzalez *et al.* 2014; Saphirstein & Morgan, 2014). However, the signalling proteins that act as mechanosensors and mechanoresponders in vascular smooth muscle are still not fully characterised.

### LIM proteins

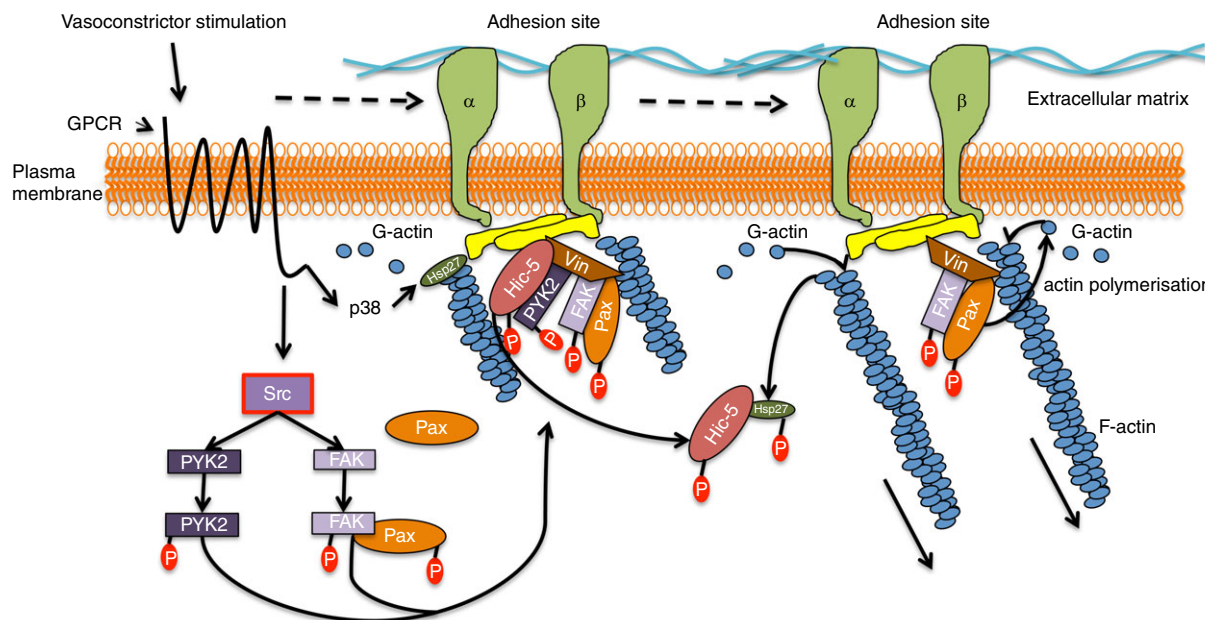
LIM domains are versatile protein binding sites found in many adaptor proteins that facilitate the assembly of multi-protein complexes involved in cytoskeletal remodelling, cell migration and gene transcription (Kadmas & Beckerle, 2004; Smith *et al.* 2014). In non-contractile cells LIM proteins are found in the nucleus where they control gene expression, and in the cytoplasm where they associate with FAs and the CK (Kadmas & Beckerle, 2004). In expression studies in non-smooth-muscle cells, in response to force a subset of LIM proteins accumulate at actin stress fibres and are implicated in the regulation of the actin CK response to mechanical stimulation (reviewed in Smith *et al.* 2014). Similarly, endogenous LIM proteins Hic-5 and cysteine rich protein (CRP2) associate with stress fibres, whereas paxillin remains at FA in response to cyclical stretch in the mouse smooth muscle cell line SVS30

(Kim-Kaneyama *et al.* 2005). Additionally, in cultured rat VSMCs stably expressing the LIM protein zyxin–green fluorescent protein, zyxin accumulated at sites of force transmission (Sun *et al.* 2012). Given that zyxin, paxillin and Hic-5 are all expressed in contractile VSMCs and vascular tissues (Ward *et al.* 2002; Kim-Kaneyama *et al.* 2005; Ohanian *et al.* 2005; Srinivasan *et al.* 2008; Sun *et al.* 2012; Poythress *et al.* 2013), a role for these LIM proteins in VSMC mechanotransduction and contractility seems likely.

### Hic-5

Hic-5 is a FA scaffold protein, first identified as a TGF $\beta$  and H<sub>2</sub>O<sub>2</sub> inducible gene (Shibanuma *et al.* 1994). Through its scaffolding activity Hic-5 is implicated in apoptosis (Kim-Kaneyama *et al.* 2011; Hornigold *et al.* 2013; Desai *et al.* 2014), protein degradation (Ryan *et al.* 2012; Desai *et al.* 2014; Lei *et al.* 2014), fibrosis (Kim-Kaneyama *et al.* 2012; Hornigold *et al.* 2013; Desai *et al.* 2014), force sensing (Fujita *et al.* 1998; Thomas *et al.* 1999; Hetey *et al.* 2005; Kim-Kaneyama *et al.* 2005; Guignandon *et al.* 2006; Deakin *et al.* 2012; Smith *et al.* 2014), contractility (Kim-Kaneyama *et al.* 2005; Srinivasan *et al.* 2008), migration (Kim-Kaneyama *et al.* 2008; Deakin *et al.* 2012) and proliferation (Wang *et al.* 2011; Kim-Kaneyama

*et al.* 2012). However, the effects of Hic-5 are cell-type specific, for instance Hic-5 is anti-apoptotic in cultured VSMCs (Kim-Kaneyama *et al.* 2011) and pro-apoptotic in mesangial myofibroblasts and lung fibroblasts (Hornigold *et al.* 2013; Desai *et al.* 2014), and stimulates migration in epithelial and endothelial cells, but inhibits migration in fibroblasts and smooth muscle cells (Wu *et al.* 2005; Avraamides *et al.* 2007; Dabiri *et al.* 2008; Kim-Kaneyama *et al.* 2008). Hic-5 is a homologue of paxillin (Thomas *et al.* 1999) and in addition to sharing a similar domain structure and localising to the same sites (FAs) Hic-5 shares many binding partners with paxillin (Fujita *et al.* 1998; Thomas *et al.* 1999). However, they must not entirely duplicate each other's functions as paxillin knockout is embryonic lethal (Hagel *et al.* 2002). Furthermore, Hic-5 has unique binding partners such as Hsp27 (Jia *et al.* 2001; Srinivasan *et al.* 2008), CRP2 (Kim-Kaneyama *et al.* 2005) and the ubiquitin ligase Cblc (Ryan *et al.* 2012) and lacks the binding site for adaptor protein Crk that is present in paxillin (Thomas *et al.* 1999). These differences indicate that Hic-5 and paxillin will regulate different downstream pathways and so regulate different cellular functions. Indeed, forced expression of Hic-5 in fibroblasts decreases tyrosine phosphorylation of paxillin, possibly through sequestration of FAK, leading to the suggestion that Hic-5 acts as a counterbalance to



**Figure 2. The possible role of paxillin and Hic-5 in vasoconstrictor-induced focal adhesion and actin cytoskeleton remodelling in vascular smooth muscle**

Vasoconstrictor agonist acting through G protein-coupled receptor (GPCR) activates Src tyrosine kinase, leading to activation of PYK2 and FAK, and tyrosine phosphorylation of paxillin (Pax) and Hic-5. Remodelling of adhesion sites occurs such that paxillin, FAK and PYK2 relocate to, and Hic-5 moves away from, the adhesion sites. Simultaneously vasoconstrictor activation induces p38MAPK-dependent phosphorylation of the actin-capping protein Hsp27. Phosphorylated Hsp27 and Hic-5 associate, freeing the barbed end of actin filaments promoting actin polymerisation. Phosphorylation of paxillin also promotes actin polymerisation inducing actin cytoskeleton remodelling. Abbreviation: Vin, vinculin.

paxillin opposing some of the effects of paxillin such as growth promotion and cell spreading (Fujita *et al.* 1998; Nishiya *et al.* 2001). Additionally, in the SVS30 smooth muscle cell line in response to cyclical stretch endogenous Hic-5 moved out of FAs and bound to CRP2 at actin stress fibres whereas paxillin remained within FAs, further demonstrating differences between them (Kim-Kaneyama *et al.* 2005). An expression study in fibroblasts has shown that the pathways regulating Hic-5 and paxillin association with vinculin at FAs also differ, with Rac1 activation regulating paxillin association with vinculin in immature FAs and RhoA activation important for Hic-5 association with vinculin in mature FAs (Deakin *et al.* 2012). Furthermore, when cells were placed under mechanical strain by growing in 3-D matrices Hic-5 but not paxillin interacted with vinculin at adhesion sites (Deakin *et al.* 2012), suggesting that in tissues that co-express Hic-5 and paxillin, Hic-5 would be the main mechanosensor through preferential interaction with vinculin at adhesion sites. Whilst paxillin is ubiquitously expressed (Brown & Turner, 2004), Hic-5 is restricted to smooth muscle, myofibroblast and epithelial cells in adults (Yuminamochi *et al.* 2003; Kim-Kaneyama *et al.* 2005) and Hic-5 is recognised as a phenotypic marker of differentiated smooth muscle cells (Wang *et al.* 2011). Consequently, Hic-5 would be expected to have an important role in mechanosensing and counterbalancing the effects of paxillin in smooth muscle.

### Hic-5 in vascular cells and tissues

Hic-5 is abundantly expressed in differentiated contractile vascular smooth muscle cells in both large and small arteries (Yuminamochi *et al.* 2003; Kim-Kaneyama *et al.* 2005; Srinivasan *et al.* 2008). In femoral artery, electron microscopy studies have shown Hic-5 is localised at the cell periphery of smooth muscle cells indicative of association with adhesion sites (Kim-Kaneyama *et al.* 2012). Expression of Hic-5 in endothelial cells in arteries appears to be low, as it is not detectable by immunofluorescence. However, Hic-5 has been detected in endothelial cells by electron microscopy following immunogold labelling of mouse pulmonary arterioles (Kim-Kaneyama *et al.* 2012). The Hic-5 was present at the abluminal plasma membrane adjacent to the extracellular matrix indicating localisation to adhesion sites. *In vitro* studies have shown that Hic-5 regulation of FA dynamics plays a key role in endothelial cell migration (Wu *et al.* 2005; Avraamides *et al.* 2007; Komorowsky *et al.* 2010) stimulating interest in Hic-5 as a regulator of angiogenesis. However, the function of Hic-5 in endothelial cell responses in arteries remains unclear.

### Hic-5 in vascular contractility

Two studies have implicated Hic-5 in vascular smooth muscle contractility. In a smooth muscle cell line endogenous Hic-5 translocated from FA to stress fibres in response to cyclical stretch; in mouse aorta Hic-5 and CRP2 localised to filamentous structures shown by electron microscopy. Additionally, mouse embryonic fibroblasts with forced expression of Hic-5 had a slower contraction of the 3-D gel matrix; in contrast, cells over-expressing paxillin had greater contraction. This led to the proposal that in response to cyclic stretch Hic-5 localises to stress fibres and negatively regulates contractility (Kim-Kaneyama *et al.* 2005). We have shown in intact small arteries that noradrenaline promoted Hic-5 association with PYK2 and increased Hic-5 tyrosine phosphorylation, which induced Hic-5 translocation from the actin cytoskeleton to the cytosol and interaction with Hsp27. This was accompanied by actin cytoskeleton remodelling and contraction, and inhibition of Hic-5 tyrosine phosphorylation-attenuated contraction suggesting Hic-5 positively regulates contractility in differentiated smooth muscle cells in the artery wall (Srinivasan *et al.* 2008) (Fig. 2). However, a possible explanation for the apparently contradictory effects on contractility in these two studies is an effect on paxillin activity. It has been shown that over-expression of Hic-5 in fibroblasts reduces tyrosine phosphorylation of paxillin (Fujita *et al.* 1998; Nishiya *et al.* 2001) and that paxillin activation is important for tension development in fibroblasts (Deakin *et al.* 2012). Additionally, in airway smooth muscle tissue tyrosine phosphorylation of paxillin is important for acetylcholine-induced contraction (Gerthoffer & Gunst, 2001), and in intact small arteries noradrenaline induces paxillin tyrosine phosphorylation and association with the actin cytoskeleton (Ohanian *et al.* 2005). In our study in intact arteries (Srinivasan *et al.* 2008) we used an Src inhibitor to block Hic-5 tyrosine phosphorylation and to study contractility. Src inhibition also blocks paxillin tyrosine phosphorylation (Ohanian *et al.* 2005) raising the possibility that the effect on contractility was mediated through inactivation of paxillin. Obviously, further work is required to elucidate the interplay between Hic-5 and paxillin in smooth muscle contractility.

### Hic-5 in vascular remodelling

Hic-5 is also implicated in vascular remodelling. Wire injury of the mouse carotid artery led to decreased Hic-5 expression in the medial smooth muscle cells and delivery of Hic-5 to the injured artery decreased neointima formation (Kim-Kaneyama *et al.* 2008). *In*

*in vitro*, over-expression of Hic-5 in a smooth muscle cell line repressed uPA expression, an activator of matrix metalloproteinase (MMP), an effect that was not duplicated by paxillin expression (Kim-Kaneyama *et al.* 2008). This suggests that the protective effect of Hic-5 against vascular injury is mediated through inhibition of MMP activation. However, in a mouse model of abdominal aortic aneurysm (AAA) loss of Hic-5 in aortic vascular smooth muscle cells suppressed AAA. Mechanistically, Hic-5 increased c-Jun N-terminal kinase activity by acting as a scaffold for c-Jun N-terminal kinase and its activator MKK4, leading to increased MMP activity, elastin degradation and increased susceptibility to AAA (Lei *et al.* 2014). These studies again highlight how the effects of Hic-5 are context specific. However, studies in non-smooth-muscle cells showing that Hic-5 regulates fibrosis (Kim-Kaneyama *et al.* 2012; Hornigold *et al.* 2013; Desai *et al.* 2014) support a role for Hic-5 in vascular remodelling.

### Studies of Hic-5<sup>-/-</sup> mice

Given the above evidence that Hic-5 is involved in smooth muscle contractility and artery remodelling, it was surprising that genetic deletion of Hic-5 in mice produced no obvious vascular or other phenotype (Kim-Kaneyama *et al.* 2011). However, following wire injury of the femoral artery there was increased medial smooth muscle cell apoptosis and increased neointima formation in the Hic-5<sup>-/-</sup> mice compared to wild-type (Kim-Kaneyama *et al.* 2011) supporting a protective role of Hic-5 in VSMCs following vascular injury. In support of Hic-5 as a mechanosensor and regulator of the actin cytoskeleton and adhesion sites, aortic VSMCs from the Hic-5<sup>-/-</sup> mice had fewer stress fibres and in response to mechanical stimulation the FA protein vinculin relocated to the cytoplasm. Additionally, Hic-5<sup>-/-</sup> aortic VSMCs were more susceptible to stretch-induced apoptosis, suggesting that Hic-5 decreases VSMC sensitivity to stretch-induced apoptosis by stabilising stress fibres and vinculin at FAs (Kim-Kaneyama *et al.* 2011). Hic-5 did not protect VSMCs from cytokine- or H<sub>2</sub>O<sub>2</sub>-induced apoptosis (Kim-Kaneyama *et al.* 2011), further emphasising its role as a mechanoresponder in VSMCs.

### Summary

It is only relatively recently that the role of the actin CK in VSMC responses has been appreciated. It is now clear that dynamic restructuring of the CK is required for contraction – independent of myosin light chain phosphorylation – and for maintenance of vascular tissue integrity. Actin CK remodelling occurs in response to both mechanical stresses and vasoactive

agonist stimuli, and is implicated in increased large artery stiffness, neointima formation and small artery remodelling in cardiovascular disease. In addition to actin CK remodelling evidence is now accumulating for concomitant adhesion site remodelling in response to vasoconstrictor and mechanical stimuli. However, although it is clear that non-receptor tyrosine kinases and tyrosine phosphorylation of FA proteins is an important component of the response, the major pathways involved and their regulation and cross talk remain unclear. The goal of future research must be to unravel the complexities of VSMC actin CK and adhesion site remodelling in order to offer new areas for development of novel therapeutic intervention in cardiovascular disease.

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## Additional information

### Competing interests

None declared.

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