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Hantavirus Interferon Regulation and Virulence Determinants

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Abstract

Hantaviruses predominantly replicate in primary human endothelial cells and cause 2 diseases characterized by altered barrier functions of vascular endothelium. Most hantaviruses restrict the early induction of interferon- β (IFN β) and interferon stimulated genes (ISGs) within human endothelial cells to permit their successful replication. PHV fails to regulate IFN induction within human endothelial cells which self-limits PHV replication and its potential as a human pathogen. These findings, and the altered regulation of endothelial cell barrier functions by pathogenic hantaviruses, suggest that virulence is determined by the ability of hantaviruses to alter key signaling pathways within human endothelial cells. Our findings indicate that the Gn protein from ANDV, but not PHV, inhibits TBK1 directed ISRE, kB and IFN^β induction through virulence determinants in the Gn cytoplasmic tail (GnT) that inhibit TBK1 directed IRF3 phosphorylation. Further studies indicate that in response to hypoxia induced VEGF, ANDV infection enhances the permeability and adherens junction internalization of microvascular and lymphatic endothelial cells. These hypoxia/VEGF directed responses are rapamycin sensitive and directed by mTOR signaling pathways. These results demonstrate the presence of at least two hantavirus virulence determinants that act on endothelial cell signaling pathways: one that regulates antiviral IFN signaling responses, and a second that enhances normal hypoxia-VEGF-mTOR signaling pathways to facilitate endothelial cell permeability. These findings suggest signaling pathways as potential targets for therapeutic regulation of vascular deficits that contribute to hantavirus diseases and viral protein targets for attenuating pathogenic hantaviruses.

Introduction

Hantaviruses predominantly infect the endothelial cell lining of vessels and nonlytically cause two diseases: hemorrhagic fever with renal syndrome (HFRS) and hantavirus pulmonary syndrome (HPS) (Duchin et al., 1994; Lahdevirta et al., 1982; Lee, 1982; Nichol et al., 1993; Schmaljohn, 2001; Yanagihara and Silverman, 1990; Zaki et al., 1995). HFRS results from infection by Eurasian hantaviruses (Hantaan virus, HTNV; Dobrava virus,

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DOBV; Puumala virus, PUUV) (Lahdevirta et al., 1982; Lee et al., 1982; Schmaljohn, 2001) while hantaviruses identified throughout the Americas (ie. Andes virus, ANDV; Sin Nombre virus, SNV; New York virus, NYV) are associated with HPS (Duchin et al., 1994; Enria et al., 1996; Lopez et al., 1996; Nichol et al., 1993; Schmaljohn, 2001). In contrast, Tula virus (TULV) and Prospect Hill virus (PHV) are hantaviruses that are not associated with any human disease (Plyusnin et al., 1994; Yanagihara et al., 1987). While TULV and PHV differ from pathogenic hantaviruses by their use of discrete integrin receptors (Gavrilovskaya et al., 1999; Gavrilovskaya et al., 1998; Raymond et al., 2005), PHV also fails to regulate early IFN induction which restricts its replication in endothelial cells and likely contributes to its inability to be a human pathogen (Alff et al., 2006; Alff et al., 2008; Geimonen et al., 2002; Spiropoulou et al., 2007). These findings suggest that hantaviruses contain virulence determinants that restrict antiviral IFN pathway signaling responses and alter normal endothelial cell signaling pathways that control vascular permeability.

Only a few viruses specifically target the endothelial cell (EC) lining of vessels and cause acute edematous or hemorrhagic disease. Mechanisms by which hantaviruses disrupt fluid barrier integrity and clearance functions of the endothelium are just beginning to be disclosed. Vascular permeability induced by hantaviruses is likely to be multifactorial in nature and result from virally altered EC responses and signaling pathways, tissue hypoxia and immune cell and platelet functions (Gavrilovskaya et al., 2012a; Gavrilovskaya et al., 2012b; Gavrilovskaya et al., 2010, 2012c, 2013; Gorbunova et al., 2010; Gorbunova et al., 2013; Gorbunova et al., 2011; Hammerbeck and Hooper, 2011; Kilpatrick et al., 2004; Koster and Mackow, 2012; Mori et al., 1999; Raymond et al., 2005; Taylor et al., 2013; Terajima et al., 1999; Vaheri et al., 2013). This is likely to occur through a collaboration of interactions which bypass redundant vascular systems that control critical fluid barrier functions. Failure of the endothelium to regulate hemorrhage or edematous fluid accumulation in tissues has severe pathologic consequences. Deficits in the regulation of vascular permeability are dramatically illustrated by findings in HPS patients which result in localized acute pulmonary edema, unprecedented pulmonary fluid accumulation rates (up to 1 liter/hour) and a ~40% mortality rate (Duchin et al., 1994; Koster and Mackow, 2012; Zaki et al., 1995). As the multifactorial nature of vascular regulation is impacted by many systems, a variety of hypotheses have been expounded, but need to be prefaced by stating that there is currently no data demonstrating that any of these theories play a causal role in vascular permeability induced by hantaviruses.

The primary understanding of hantavirus induced vascular deficits, remains the viruses ability to infect the endothelial cell lining of the vasculature and nonlytically cause edematous or hemorrhagic disease (Duchin et al., 1994; Lahdevirta et al., 1982; Lee, 1982; Nichol et al., 1993; Schmaljohn, 2001; Yanagihara and Silverman, 1990; Zaki et al., 1995). Hantaviruses dysregulate microvascular and lymphatic endothelial cell (MEC and LEC) functions that normally restrict fluid leakage from vessels and clear fluid from tissues (Gavrilovskaya et al., 2010, 2012c, 2013; Gavrilovskaya et al., 2008; Gorbunova et al., 2010; Gorbunova et al., 2013; Gorbunova et al., 2011; Koster and Mackow, 2012; Raymond et al., 2005; Shrivastava-Ranjan et al., 2010). The effects of hantavirus infection of endothelial cells remains enigmatic and the focus of our studies of altered endothelial cell

signaling pathways (Gavrilovskaya et al., 2013; Gorbunova et al., 2011) that are fundamental to altered vascular permeability and hantavirus virulence.

Hantavirus entry into human endothelial cells initially discriminates between pathogenic hantaviruses, whose infection is fostered by human $\alpha_v\beta_3$ integrins, and nonpathogenic TULV and PHV which are unaffected by the presence of $\alpha_{\nu}\beta_{3}$ integrins (Gavrilovskaya et al., 1999; Gavrilovskaya et al., 1998; Raymond et al., 2005). Since $\alpha_{v}\beta_{3}$ is a known regulator of vascular permeability this finding ties hantavirus receptor usage to vascular permeability (Gavrilovskaya et al., 2008; Raymond et al., 2005). Yet, in vitro, pulmonary microvascular and lymphatic endothelial cells (MECs, LECs), and human umbilical vein endothelial cells (HUVECs) are not permeablized by hantavirus infection alone suggesting that receptor usage itself is not a cause of vascular permeability (Gavrilovskaya et al., 2012c, 2013; Gavrilovskaya et al., 2008; Gorbunova et al., 2010; Gorbunova et al., 2013). Interestingly, studies indicating that cell-associated pathogenic hantaviruses bind inactive $\alpha_{v}\beta_{3}$ integrins, days after infection, tie $\alpha_{v}\beta_{3}$ integrin usage to the regulation of signaling pathways induced by a potent vascular permeability inducer, vascular endothelial growth factor (VEGF) (Gavrilovskaya et al., 1999; Gavrilovskaya et al., 2010; Raymond et al., 2005; Robinson et al., 2004). $\alpha_{v}\beta_{3}$ normally forms a complex with VEGF receptor 2 (VEGFR2), which tempers VEGFR2 directed permeability in response to localized VEGF release. Knocking out β_3 or inhibiting $\alpha_{v}\beta_3$ promotes VEGFR2 directed endothelial cell permeability (Borges et al., 2000; Byzova et al., 2000; Hodivala-Dilke et al., 1999; Reynolds et al., 2002; Robinson et al., 2004). Furthering this association during hantavirus infection, the permeability of endothelial cells infected by ANDV, SNV and NY-1V, but not nonpathogenic TULV or PHV hantaviruses, is dramatically enhanced in response to VEGF (Gavrilovskaya et al., 2010, 2012c; Gavrilovskaya et al., 2008; Gorbunova et al., 2010; Gorbunova et al., 2013; Gorbunova et al., 2011).

VEGF is a potent vascular permeability factor (VPF) that locally induces vascular permeability by binding endothelial cell VEGF receptors, within 1.5 mm of its release, and directing the disassembly of inter-endothelial cell adherens junctions (Dejana et al., 2008; Dvorak et al., 1995; Gavard, 2009; Gavard and Gutkind, 2006). VEGF is induced by hypoxia to facilitate repair, and increase gas exchange within the lung, and VEGF is inactivated by circulating soluble receptors that prevent systemic vascular permeability responses. VEGF induced pulmonary edema is known to be caused by hypoxia in high altitude settings (Berger et al., 2005; Hanaoka et al., 2003; Hopkins et al., 2005; Voelkel, 2002).

HPS patients are acutely hypoxic (Bustamante et al., 1997; Zaki et al., 1995) and a recent retrospective analysis of pulmonary edema fluids in a small number of HPS patients indicated the presence of high levels of VEGF (Gavrilovskaya et al., 2012a). Hantavirus infection of MECs and LECs may disengage one or more fluid barrier regulatory mechanisms, thereby increasing vascular leakage or fluid clearance resulting in tissue edema (Dehler et al., 2006; Schraufnagel et al., 2003). These findings suggest one of many mechanisms that may participate in HPS directed pulmonary edema and vascular deficits within hantavirus patients. However, although HPS patients are hypoxic there is as yet no causal evidence for this mechanism in hantavirus disease.

Consistent with roles for $\alpha_v\beta_3$ and hypoxia directed VEGF in hantavirus pathogenesis, hypoxia and VEGF tie into complex intracellular signaling pathways and feedback regulatory mechanisms that may be altered by virulence determinants within pathogenic hantaviruses. Hypoxia and VEGFR2 are tied to mTOR (mammalian target of rapamycin) directed cell division, control of cell size and feedback regulation of hypoxic responses (Kim et al., 2009; Xue et al., 2009). Studies presented below tie virulence determinants within hantavirus proteins to altered VEGF directed mTOR activation.

In addition to regulating cell receptor signaling, hantaviruses regulate IFN signaling pathways within human endothelial cells in order to successfully replicate and be human pathogens. Hantavirus replication is highly sensitive to the early addition of IFN or IFN pretreatment and hantaviruses grow to much lower titers in IFN competent cell lines than IFN locus defective Vero E6 cells (Alff et al., 2006). Interestingly, the effects of IFN addition are nearly absent when IFN is added 1 day post-infection (Alff et al., 2006), and consistent with hantaviruses inducing high level ISG responses at late times post-infection (Geimonen et al., 2002). In contrast to pathogenic hantaviruses, PHV rapidly induces IFN β and IFN stimulated gene (ISG) responses that restrict its replication in human endothelial cells (Geimonen et al., 2002) and this response, in addition to receptor usage, are potential explanations for the absence of PHV associated human disease (Alff et al., 2006; Alff et al., 2008; Matthys et al., 2011; Matthys and Mackow, 2012). Our findings suggest that permissive hantavirus replication in human endothelial cells results from the selective restriction of early IFN induction (Alff et al., 2006; Geimonen et al., 2002; Matthys et al., 2012).

The ability of hantaviruses to regulate IFN induction and alter vascular and lymphatic endothelial cell signaling responses suggests the presence of encoded virulence determinants that permit viral replication and alter cellular responses which control fluid barrier functions of the endothelium. Here we show that hantaviruses contain virulence determinants that alter normal endothelial cell functions by regulating VEGF-mTOR signaling responses and permitting viral replication by inhibiting the early induction of Type 1 IFN. These findings suggest the presence of an IFN regulating virulence determinant in the Gn protein that is required for hantavirus replication in human endothelial cells and for subsequent vascular permeability deficits in HFRS and HPS patients. However, these clues to vascular dysfunction provide potential mechanisms by which hantaviruses induce vascular permeability and acute edema that remain to be defined *in vivo*.

Results

Hantavirus Regulation of Early IFN Responses Defines Virulence Determinants in the GnT

Replicating RNA viruses generate small amounts of dsRNA that are detected by cytoplasmic helicases which signal TBK1/IKK ϵ complexes (Seth et al., 2006; Yoneyama and Fujita, 2007; Yoneyama et al., 2004) to activate NF- κ B and cellular IFN response factors (IRFs3/5/7) (Hacker et al., 2011; Hiscott, 2007; Tu et al., 2013). Activated IRFs and NF- κ B translocate to the nucleus and transcriptionally induce IFN β and additional antiviral ISG responses from promoters containing IFN stimulated response elements (ISREs)

(Charoenthongtrakul et al., 2013; Daffis et al., 2009; Delhase et al., 2011; Hacker et al., 2011; Lazear et al., 2013).

We have found that GnT proteins from NY-1V, ANDV, TULV and HTNV, but not PHV, regulate polyI:C, RIG-I, MDA5 and TBK1 directed ISRE, κ B and IFN β transcriptional responses upstream of constitutively active IRF3-5D, and at the level of the TBK1-TRAF3 complex (Alff et al., 2006; Alff et al., 2008; Geimonen et al., 2002; Matthys et al., 2013). Yet IFN signaling responses are cytoplasmic and only cytoplasmic elements of Gn are likely to affect regulation. To investigate this we truncated Gn to express only the C-terminal 42 residues (C42) within its cytoplasmic tail (GnT) and evaluated the ability of C42 domains to regulate ISRE and IFN transcriptional responses. Figure 1A demonstrates that GnT C42 domains from NYV or ANDV, but not PHV, inhibit TBK1 directed ISRE transcriptional responses in a dose dependent manner. Inhibition of IFN^β transcriptional responses by GnT constructs are similar to expressing GnGc proteins (Gc level monitored by Western) from ANDV M gene segments as indicated in Figure 1B. In both Figure 1A and 1B pathogenic hantavirus GnT, C42 or GnGc expression inhibited IFN signaling pathway responses. Additional studies indicate that the ANDV GnGc inhibits RIG-I, MDA5 and TBK1, but not IRF3-5D, directed transcription from an ISRE promoter and the IFN β enhanceosome (Matthys et al., 2013; Matthys et al., 2011; Matthys and Mackow, 2012). In addition, GnGc expression also inhibited RIG-I directed IRF3 phosphorylation (Matthys et al., 2013). Collectively, these findings demonstrate that NY-1V, ANDV and TULV GnTs as well as the GnGc polyprotein inhibit RIG-I induced transcriptional responses by impacting TBK1 phosphorylation of IRF3 (Matthys et al., 2013). These findings indicate that the GnT domain contains an IFN regulating element with the potential to be a virulence determinant within hantaviruses that enhances viral replication and spread.

VEGF and ANDV Infection Enhance VE-Cadherin Internalization and EC Permeability

ANDV infects the endothelial cell lining of capillaries and results in patient hypoxia and acute pulmonary edema leading to respiratory distress (Bustamante et al., 1997; Duchin et al., 1994; Enria et al., 1996; Lopez et al., 1996; Zaki et al., 1995). Hypoxia itself induces VEGF which is a permeability factor that has the potential to cause edema during ANDV infection and contribute to HPS (Dvorak, 2006; Dvorak et al., 1995). VEGFR2 activation increases EC permeability by directing VE-cadherin internalization, but not degradation, which disassembles adherens junctions (AJs) and permits rapid reassembly of AJs (Corada et al., 1999: Corada et al., 2002: Gavard, 2009: Gavard and Gutkind, 2006: Gavard et al., 2008; Nawroth et al., 2002; Wallez et al., 2006; Zanetti et al., 2002). VE-cadherin internalization was monitored following the protocol of Gavard (Gavard, 2009; Gavard and Gutkind, 2006; Gavard et al., 2008). EC monolayers were infected with ANDV, TULV or mock infected, under hypoxic or normoxic conditions or following VEGF addition (shown). In order to assay VE-cadherin internalization, monolayers were incubated with FITC-labeled anti-VE-cadherin mAb (100 ng/ml; sc52751, Santa Cruz; 30 min, 4°C) and subsequently 1 hour at 37°C to synchronize internalization (Gavard, 2009). Cells were acid washed to remove extracellular VE-cadherin antibody and cells containing internalized VE-cadherin were quantitated by fluorescence microscopy (Gavard, 2009; Gavard and Gutkind, 2006; Gavard et al., 2008). The internalization of VE-cadherin was monitored following infection

of primary pulmonary microvascular endothelial cells (MECs) by ANDV (HPS) or TULV (nonpathogenic). Figure 2A demonstrates the hyper-responsiveness of VE-cadherin internalization in ANDV infected MECs to VEGF addition and the absence of VEGF effects on TULV or mock infected MECs. These findings demonstrate that ANDV infection combined with VEGF addition dramatically mobilizes VE-cadherin from AJs to intracellular stores.

Hypoxia Induces EC Permeability via Rapamycin Sensitive mTOR Signaling

mTOR signaling is intimately tied to hypoxia directed VEGF induction and results from signaling pathways that are blocked by the mTOR inhibitor rapamycin (Gavrilovskaya et al., 2013). We recently evaluated responses of human pulmonary MECs and LECs in response to VEGF addition or hypoxic conditions (Gavrilovskaya et al., 2012c; Gorbunova et al., 2013). The permeability of MEC and lymphatic EC (LEC) monolayers was assayed by adding FITC-dextran to the upper chamber and monitoring levels in the lower chamber of confluent EC monolayers in response to VEGF addition (Gavrilovskaya et al., 2012c; Gorbunova et al., 2013). We observed little change in the permeability of ANDV infected MECs or LECs alone, but observed a dramatic increase in the permeability of VEGF or hypoxia treated ANDV infected MECs and LECs (Figure 2B). In contrast, neither VEGF nor hypoxia treatment of TULV infected cells resulted in an increase in MEC or LEC permeability. Interestingly, the hypoxia induced permeability was sensitive to the pathway specific mTOR inhibitor, rapamycin, indicating that permeability responses are mediated by mTOR directed HIF1 α activation as well as HIF-1 α , hypoxia and VEGF directed signaling responses during infection by pathogenic hantaviruses (Figure 2C)(Gavrilovskaya et al., 2013).

Consistent with this, we recently reported that ANDV infection dramatically increased HIF1a directed VEGF-A, ANG4 and EGLN3 mRNA levels within hypoxic MECs and LECs (Gavrilovskaya et al., 2013) and that human pulmonary edema fluids from HPS patients contains high levels of VEGF (Gavrilovskaya et al., 2012a). Hypoxia stabilizes the formation of HIF1a transcriptional complexes that induce VEGF, additional hypoxia responsive factors and stress regulators that impact the activation of mTOR signaling responses (Gavrilovskaya et al., 2013; Zhou et al., 2007) (Figure 2C). mTOR signaling also controls cell size and these findings support data demonstrating that VEGF and hypoxia direct the formation of giant LECs and MECs through a rapamycin sensitive mTOR dependent mechanism (Gavrilovskaya et al., 2012c, 2013). These findings suggest the presence of a second virulence determinant within pathogenic hantaviruses which, in addition to IFN pathway regulation, targets mTOR signaling pathways (Figure 2C).

Discussion

Our studies indicate that pathogenic hantaviruses contain virulence determinants that alter the normal regulation of endothelial cell signaling pathways to enhance viral replication and spread (IFN regulation) and foster the permeability of endothelial cell adherens junctions (aberrant hypoxia-VEGF-mTOR signaling) (Gavrilovskaya et al., 2012b; Matthys et al., 2013; Matthys and Mackow, 2012). Successful hantavirus replication within human

endothelial cells is at least in part due to their ability to regulate the induction of IFN β (Alff et al., 2006; Geimonen et al., 2002; Spiropoulou et al., 2007). Our findings indicate that GnTs from pathogenic ANDV, NYV, SNV as well as nonpathogenic TULV inhibit TBK1 directed ISRE, NF- κ B or IFN β transcriptional responses (Alff et al., 2006; Alff et al., 2008; Matthys et al., 2013; Matthys et al., 2011). GnTs fail to inhibit constitutively active IRF3-5D and block TBK1 directed IRF3 phosphorylation (Matthys et al., 2013). In contrast, the GnT of nonpathogenic PHV fails to regulate early IFN induction in human ECs and PHV fails to successfully replicate in human ECs or become a human pathogen (Alff et al., 2006; Alff et al., 2008; Matthys et al., 2011; Spiropoulou et al., 2007). Another virulence mechanism is suggested by failure of TULV and PHV to use $\alpha_v\beta_3$ integrins for entry or later cell association in comparison with pathogenic hantaviruses and the known role of $\alpha_v\beta_3$ in regulating vascular permeability (Coller and Shattil, 2008; Reynolds et al., 2002; Robinson et al., 2004).

GnT is an IFN Regulating Virulence Determinant

Our studies of GnGc and GnT C42 domains point out the ability of this expressed protein to inhibit the antiviral effects of IFN induction by blocking RIG-I/MDA5 directed TBK1/IKKE signaling responses(Alff et al., 2006; Alff et al., 2008; Matthys et al., 2013; Matthys et al., 2011; Matthys and Mackow, 2012). Findings presented here demonstrate the importance of the C42 GnT domain in regulation and differences between pathogenic and nonpathogenic PHV in GnT functions that foster its role as a determinant that facilitates viral replication in human ECs (Matthys et al., 2013; Matthys et al., 2011; Matthys and Mackow, 2012). Additional studies of the NYV GnT have recently established that only 1 change, Y627 to A, S or F, prevented GnT regulation of TBK1 directed ISRE, kB or IFNB transcriptional responses (Matthys et al., 2013). Consistent with this, the Y627 residue was required for the NYV GnT to inhibit RIG-I directed IRF3 phosphorylation and cause a reduction in total IRF3 levels. Although these findings define a single tyrosine residue within the NYV GnT (Y627) required for inhibiting antiviral ISRE, κB and IFN β transcriptional responses, residues within other hantavirus GnTs required for regulation have yet to be defined (Matthys et al., 2013). Nontheless these findings identify residue specific determinants that may be used for viral attenuation and define the GnT as an IFN regulating determinant of viral replication.

Hypoxia/VEGF Enhance ANDV Directed VE-Cadherin Internalization and EC Permeability

HPS is a highly lethal disease resulting in acute rapidly progressive pulmonary edema and shock. Hypoxia, thrombocytopenia and vascular permeability are hallmark findings of hantavirus patients and contribute to acute pulmonary edema in HPS disease (Duchin et al., 1994; Koster and Mackow, 2012; Nolte et al., 1995; Zaki et al., 1995). Pathogenic mechanisms accounting for the rapid rate of pulmonary fluid accumulation have yet to be demonstrated, but appear to be a consequence of the non-cytolytic hantavirus infection of endothelial cells (Gavrilovskaya et al., 2012b; Koster and Mackow, 2012; Taylor et al., 2013; Vaheri et al., 2013). Although, MECs and LECs are not permeabilized by hantavirus infection alone, hantavirus infection of the endothelium provides a means for the virus to alter EC responses that normally regulate capillary leakage and pulmonary fluid clearance (Gavrilovskaya et al., 2012a; Gavrilovskaya et al., 2012b; Gavrilovskaya et al., 2012c;

Gavrilovskaya et al., 2008). Our findings indicate that hypoxia or VEGF addition are sufficient to induce hyper-permeability of ANDV, but not nonpathogenic TULV, infected MECs or LECs and that these responses are sensitive to the effects of the mTOR inhibitor rapamycin (Gavrilovskaya et al., 2013; Gavrilovskaya et al., 2008). These findings suggest that ANDV and other pathogenic hantaviruses encode virulence determinants which alter interrelated hypoxia-VEGF-mTOR responses (Figure 2C).

Hypoxia and VEGF Direct Permeability through Increased mTOR Signaling Responses

Constitutive mTOR activation results in the formation of giant cells, and mTOR signaling responses control HIF1a and VEGF directed permeability (Forsythe et al., 1996; Wolff et al., 2011). Genetic mutations in TSC1/TSC2 result in the formation of giant cells through the constitutive activation of mTOR and the downstream phosphorylation of p70-S6K (Laplante and Sabatini, 2012; Ruvinsky and Meyuhas, 2006). ANDV infection reportedly causes the formation of giant LECs in response to VEGF (Gavrilovskaya et al., 2012b). In fact, hypoxic conditions, 1–2% O₂, or addition of CoCl₂ (Kim et al., 2006), dramatically increased the number of ANDV infected giant MECs or LECs (80% or 70%, respectively) and their permeability (Gavrilovskaya et al., 2012a; Gavrilovskaya et al., 2012c; Gorbunova et al., 2013). Collectively these findings indicate that in the presence of hypoxia, ANDV directs the pathway specific activation of mTOR signaling responses that control lymphatic and vascular endothelial cell permeability and VE-cadherin internalization (Figure 2A–C).

Recent studies suggest that bradykinin may contribute to hantavirus directed vascular permeability (Taylor et al., 2013; Vaheri et al., 2013). Interestingly, hypoxia is linked to ANDV dysregulation of normal endothelial cell functions through effects on bradykinin, VEGF and thrombocytopenia, all of which regulate vascular permeability (Dehler et al., 2006; Gavard and Gutkind, 2006; Hanaoka et al., 2003; Liesmaa et al., 2009). Although not evaluated in their reports (Taylor et al., 2013; Vaheri et al., 2013), bradykinin and VEGF synergistically increase VEGFR2 phosphorylation (Thuringer et al., 2002) and secreted bradykinin induces VEGF (Knox et al., 2001). In fact, hypoxia itself induces bradykinin receptors on endothelial cells (Liesmaa et al., 2009), fostering the potential interrelationship of VEGF, bradykinin and hypoxia induced responses in hantavirus directed permeability (Gavrilovskaya et al., 2012b; Gavrilovskaya et al., 2013; Gorbunova et al., 2013; Liesmaa et al., 2009; Taylor et al., 2013; Thuringer et al., 2002; Vaheri et al., 2013).

Hypoxia induced VEGF causes high altitude-induced pulmonary edema (HAPE) (Berger et al., 2005; Dehler et al., 2006; Hanaoka et al., 2003; Scherrer et al., 2010), and the ability of hypoxia alone to induce edema and thrombocytopenia suggests that hypoxia may play a critical role in the HPS disease process (Berger et al., 2005; Christou et al., 1998; Dehler et al., 2006; Dvorak, 2006; Gavrilovskaya et al., 2010, 2013; Gorbunova et al., 2013; Koster and Mackow, 2012). HPS patients are clearly hypoxic and HPS patient pulmonary edema fluid contains elevated VEGF-A levels (Gavrilovskaya et al., 2012a). However, hypoxia directs a number of additional cellular responses which act on endothelial cell and platelet functions and which may participate in vascular leakage during ANDV infection (Irigoyen et al., 2007; Kulshreshtha et al., 2008; Liesmaa et al., 2009). Hypoxia increases endothelial NO synthase (eNOS) responsible for lymphatic vessel contraction and fluid clearance

functions (Hagendoorn et al., 2004; Miao et al., 2008). Hypoxia also causes thrombocytopenia in mice and the production of the platelet inhibitor prostacyclin which renders platelets quiescent (Birks et al., 1975; Farmer et al., 2001).

Hypoxia-VEGF-mTOR Signaling Activation as a Determinant of Hantavirus Virulence

Hantavirus infected lymphatic EC are also hyper-responsive to the permeabilizing effects of VEGF (Gavrilovskaya et al., 2012b; Gavrilovskaya et al., 2012c, 2013; Gavrilovskaya et al., 2008). Although the role of LECs and lymphatic vessels in HPS have not been defined in HPS patients or animal models, there is also a compelling rationale for hantavirus infected LECs to impede fluid clearance functions of pulmonary lymphatic vessels that exacerbate pulmonary fluid accumulation (Alitalo, 2011). Hypoxic conditions and hantavirus infection of MECs or LECs induced permeability and giant cell formation via mTOR signaling responses that result in the phosphorylation of S6K (Laplante and Sabatini, 2012). Interestingly, we found that both hypoxia directed permeability and giant cell responses of ANDV infected MECs and LECs were inhibited by rapamycin, an mTOR inhibitor (Laplante and Sabatini, 2012). In fact, rapamycin is a known negative effector of hypoxia/ VEGFA induced permeabilizing responses (Land and Tee, 2007; Wolff et al., 2011). Our findings indicate that hypoxia activates mTOR signaling responses within ANDV infected ECs (Gavrilovskaya et al., 2013; Gorbunova et al., 2013; Gorbunova et al., 2011; Robinson et al., 2004). This suggests that ANDV encodes a virulence determinant which activates mTOR pathways, although it is not clear how ANDV induces mTOR signaling responses.

Conclusions

At present there is little understanding of how hantaviruses alter redundant vascular barrier regulating systems to coordinately dysregulate pulmonary responses and cause acute pulmonary edema. Pathogenic hantaviruses appear to contain virulence determinants that facilitate viral replication and spread as well as alter normal MEC and LEC responses resulting in vascular hyper-permeability. Virulence determinants that selectively alter human endothelial cell functions are likely to be targets for therapeutics that resolve altered viral-cell responses at late stages of infection and for attenuating pathogenic hantaviruses.

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Hantaviruses regulate early IFN responses to successfully negotiate human endothelial cells and cause disease. The Andes (ANDV) hantavirus causes hantavirus pulmonary syndrome, a highly lethal disease culminating in hypoxia, acute pulmonary edema and respiratory distress. This paper summarizes data on virulence determinants within hantaviruses that permit their replication in human microvascualr and lymphatic endothelial cells and their ability to alter normal signaling pathways that control vascular permeability. Our findings indicate that elements within the hantavirus GnTs regulate IRF3 phosphorylation by restricting total IRF3 levels. Further our findings indicate that hypoxic responses observed in HPS patients are sufficient to elicit permeabilizing endothelial cell responses that likely contribute to acute pulmonary edema and which are inhibited by rapamycin regulation of mTOR signaling pathways.

- Hantaviruses regulate EC signaling pathways to successfully infect human ECs, alter normal EC functions and cause highly lethal diseases.
- Virulence determinants permit hantavirus replication in ECs by altering signaling pathways that control vascular permeability.
- Virulence elements within GnTs inhibit IRF3 phosphorylation required for the induction of antiviral IFNβ responses.
- ANDV virulence determinants enhance mTOR signaling responses to hypoxic conditions that are blocked by rapamycin.
- These findings suggest the potential for therapeutically targeting pathways altered by hantavirus infection.

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Figure 1. Gn and GnT Protein Expression Regulates TBK1 Directed ISRE Promoter Transcription

A) HEK293T cells were transfected with ISRE promoter directed luciferase reporter and Renilla luciferase plasmids (Matthys et al., 2013). Cells were co-transfected with a TBK1 expressing plasmid and increasing amounts of ANDV, NYV or PHV GnT-C42 expression plasmids. Cells were harvested 1 day post-transfection and assayed for firefly luciferase activity. Results are presented as the percent induction compared to pcDNA3 induction control (100%) after standardization to Renilla luciferase levels as previously described (Matthys et al., 2013). **B**) HEK293T cells were transfected with IFN β promoter directed luciferase reporter and Renilla luciferase plasmids and co-transfected with TBK1 plasmid and either ANDV or PHV Gn-T or the ANDV M segment expression plasmid. Cells were harvested and luciferase reporter responses analyzed as in A. ANDV Gc and β -actin (loading control) were detected by Western Blot analysis as described (Matthys et al., 2013).

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Figure 2. VEGF and Hypoxia Induced Permeability Responses are Rapamycin Sensitive MECs were mock, ANDV (CHI-7913), or TULV (Tula/Moravia/MA 5302V/94) infected in BSL3 at an MOI of 0.5. **A**) Analysis of ANDV induced VE-cadherin internalization was performed 3 days post-infection as previously described on pulmonary MECs (Gavrilovskaya et al., 2008; Gorbunova et al., 2010). **B**) Two days post infection, MECs were grown in normoxic or hypoxic conditions for 18 hours and treated with rapamycin (20 ng/ml) for 1 hour prior to evaluating permeability as previously described (Gavrilovskaya et al., 2012c; Gavrilovskaya et al., 2008; Gorbunova et al., 2010; Gorbunova et al., 2013; Gorbunova et al., 2011). Monolayers were treated as indicated with VEGF (100 ng/ml) prior to assessing monolayer permeability to FITC-dextran (40,000; 0.5 mg/ml) as previously described (Gavrilovskaya et al., 2008; Gorbunova et al., 2010). **C**) Interrelationship of Hypoxia, HIF1a, VEGFA/VEGFR2 responses and the TSC1/2 regulated mTOR signaling pathway.