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Hepatocyte Growth Factor preferentially activates the anti-inflammatory arm of NF- κ B signaling to induce A20 and protect renal proximal tubular epithelial cells from inflammation

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Abstract

Inflammation induces the NF- κ B dependent protein A20 in human renal proximal tubular epithelial cells (RPTEC), which secondarily contains inflammation by shutting down NF- κ B activation. We surmised that inducing A20 without engaging the pro-inflammatory arm of NF- κ B could improve outcomes in kidney disease. We showed that hepatocyte growth factor (HGF) increases A20 mRNA and protein levels in RPTEC without causing inflammation. Upregulation of A20 by HGF was NF- κ B/RelA dependent as it was abolished by overexpressing I κ B α or silencing p65/RelA. Unlike TNF α , HGF caused minimal I κ B α and p65/RelA phosphorylation, with moderate I κ B α degradation. Upstream, HGF led to robust and sustained AKT activation, which was required for p65 phosphorylation and A20 upregulation. While HGF treatment of RPTEC significantly increased A20 mRNA, it failed to induce NF- κ B dependent, pro-inflammatory MCP-1, VCAM-1, and ICAM-1 mRNA. This indicates that HGF preferentially upregulates protective (A20) over pro-inflammatory NF- κ B dependent genes. Upregulation of A20 supported the anti-inflammatory effects of HGF in RPTEC. HGF pretreatment significantly attenuated TNF α -mediated increase of ICAM-1, a finding partially reversed by silencing A20. In conclusion, this is the first demonstration that HGF activates an AKT-p65/RelA pathway to preferentially induce A20 but not inflammatory molecules. This could be highly desirable in acute and chronic renal injury where A20-based anti-inflammatory therapies are beneficial.

Keywords

Inflammation; Cell signaling; Nuclear Factor kappa B; Hepatocyte Growth Factor; Acute Kidney Injury

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Introduction

Renal inflammation is an invariable finding in acute and chronic kidney injury (Eddy, 2005; Liu, 2002; Remuzzi and Bertani, 1998). It is a critical initiating and aggravating factor in kidney damage (Akcaý et al., 2009; Remuzzi et al., 1997). Specifically, RPTEC are central players in tubulointerstitial inflammation, being both producers of inflammatory cytokines and chemokines and early casualties of inflammation and apoptosis (Remuzzi et al., 1997; van Kooten et al., 1999). Clinical studies suggest that inflammation is a major aggravating factor in decline of renal function in patients with chronic kidney disease, or acute renal failure due to sepsis, acute graft rejection, or other causes (Akcaý et al., 2009; Robertson and Kirby, 2003; Segerer et al., 2000). Accordingly, control of the inflammatory response has been beneficial in attenuating kidney injury in several animal models of chronic and acute renal failure (Tamada et al., 2003; Tashiro et al., 2003; Vielhauer et al., 2004). Clinically, more needs to be done to define optimal and safe anti-inflammatory therapies to prevent or treat acute kidney injury and forestall progression of chronic kidney disease.

Our laboratory has long focused on defining the functions and molecular targets of A20/TNFAIP3, a potent inhibitor of NF- κ B activation and inflammation in many cells, including human RPTEC (Arvelo et al., 2002; Cooper et al., 1996; Grey et al., 1999; Kunter et al., 2005; Patel et al., 2006). A20 was first identified as a TNF α -inducible gene in endothelial cells (Opipari et al., 1990). Although originally characterized as an inhibitor of TNF α -induced apoptosis (Daniel et al., 2004; Opipari et al., 1992), its anti-apoptotic effect does not apply to RPTEC (Kunter et al., 2005). A20 is now recognized primarily as a central regulator of inflammation due to its ubiquitous inhibitory effect on NF- κ B activation in response to a broad spectrum of pro-inflammatory mediators (Beyaert et al., 2000; Cooper et al., 1996; Longo et al., 2003). In previously published work, we demonstrated that overexpression of A20 in RPTEC strongly inhibits TNF α mediated up-regulation of the pro-inflammatory adhesion molecule ICAM-1 and the chemokine MCP-1, through inhibition of NF- κ B activation (Kunter et al., 2005). A20 knockout mice die prematurely due to cachexia and massive multi-organ inflammation (Lee et al., 2000), demonstrating its fundamental role in the hierarchy of inflammatory responses. As a NF κ B-dependent gene, A20 is part of a negative regulatory loop limiting NF- κ B activation and inflammation (Bach et al., 1997; Krikos et al., 1992). Our laboratory is searching for means of inducing A20 to control inflammation without initiating an inflammatory response.

HGF is a multifunctional, pleiotropic protein with mitogenic, motogenic, and morphogenic effects in a wide variety of cells (Matsumoto and Nakamura, 1996). It is an important regulator of kidney function and a potent renoprotective agent (Matsumoto and Nakamura, 2001), preserving normal structure and function and accelerating recovery in experimental models of acute renal failure and ischemic injury (Kawaida et al., 1994; Miller et al., 1994). HGF also ameliorates chronic renal injury in a variety of models, including remnant kidney (Gong et al., 2004), unilateral ureteral obstruction (Mizuno et al., 2001) and diabetic nephropathy (Mizuno and Nakamura, 2004). These benefits are ascribed to its anti-apoptotic, anti-fibrotic (Liu, 2002), and anti-inflammatory effects in myriad pathologies including inflammatory bowel disease (Ohda et al., 2005), airway inflammation (Ito et al., 2005) and acute and chronic kidney disease (Gong et al., 2006a; Gong et al., 2006b; Kawaida et al., 1994). Some of the advantageous effects of HGF have been linked to its ability to induce (Muller et al., 2002) or inhibit NF- κ B activation both *in vitro* (Gong et al., 2008; Min et al., 2005) and *in vivo*, including in models of inflammatory kidney diseases (Giannopoulou et al., 2008; Gong et al., 2006a; Gong et al., 2006b).

In this work, we questioned whether HGF treatment of RPTEC affects expression of the anti-inflammatory protein A20, and if so, how this relates to the anti-inflammatory effects of HGF and its ability to activate or inhibit NF- κ B in these cells.

Material and Methods

Reagents

We purchased human recombinant tumor necrosis factor-alpha (TNF α) and hepatocyte growth factor (HGF) from R&D Systems (Minneapolis, MN, USA), and 2-(4-morpholinyl)-8-phenyl-1(4H)-benzopyran-4-one hydrochloride (LY294002), gelatin and fetal bovine serum (FBS) from Sigma-Aldrich Co (St. Louis, MO, USA).

Cell culture

Primary human RPTEC were purchased from Lonza (Allendale, NJ, USA) and cultured in suggested medium: REBM/REGM bullet kit (Biowhittaker Inc, Walkersville, MD, USA) or RECGM2 (Promocell, Heidelberg, Germany), as described (Kunter et al., 2005). Confluent cells between passages 6–9, derived from 3 different single donor preparations were used in experiments.

Western blot analysis

RPTEC were incubated with HGF (50 ng/mL) or TNF α (100 U/mL) for varying time periods. In some experiments cells were pre-incubated with the PI3K/AKT inhibitor LY294002 for 30 min. Total protein cell lysates were recovered, and analyzed (Laemmli, 1970) by Western blot (WB), as described (Kunter et al., 2005) using: rabbit anti-I κ B- α , mouse anti- β actin and rabbit anti-p65 antibodies (Santa Cruz Biotechnology, Inc., Santa Cruz, CA, USA), rabbit anti-phospho-p65 (Ser536; P-p65), rabbit anti-phospho-I κ B- α (Ser32; P-I κ B α), rabbit anti-phospho RelB (Ser552; P-RelB), and rabbit anti-phospho Glycogen Synthase 3 beta (Ser9; P-GSK3 β) (Cell Signaling Technology, Danvers, MA, USA), chicken anti-TNFAIP3 (A20), and rabbit anti-A1 (Abcam Inc., Cambridge, MA, USA), and mouse anti-glyceraldehyde 3-phosphate dehydrogenase (GAPDH) (Calbiochem, San Diego, CA, USA). Membranes were incubated with appropriate secondary antibodies (Thermo Scientific, Rockford, IL, USA). Immunoblots were scanned and the band intensity was quantified by densitometry using NIH ImageJ 1.41 software.

Quantitative real time polymerase chain reaction (qPCR)

Messenger RNA was isolated from RPTEC using RNeasy mini kits (Qiagen, Valencia, CA, USA) and cDNA was synthesized using iScript cDNA synthesis kit (Bio-Rad, Hercules, CA, USA). Quantitative PCR was performed to measure human p65, MCP-1, ICAM-1, VCAM-1, A1 and A20 mRNA levels (primer sequences available upon request) using iTaq Fast SYBR Green Supermix with ROX (Bio-Rad), and ABI 7500 Fast Real-time PCR system (Applied Biosystems, Inc., Foster City, CA, USA). Expression of target genes was normalized to that of the housekeeping gene or β actin.

Adenoviral-mediated gene transfer

Recombinant adenovirus (rAd.) vector expressing the porcine I κ B α gene (ECI-6) (rAd.I κ B α) was generated by C.J. Wrighton (Wrighton et al., 1996). Control rAd. β -galactosidase (rAd. β gal) was a kind gift from Dr. R. Gerald (University of Texas Southwestern Medical Center). rAd. were generated and tittered, as described (Ferran et al., 1998). RPTEC were transduced at a multiplicity of infection (MOI) of 50 plaque-forming units per cell (pfu/cell), as previously described (Kunter et al., 2005). Transgene expression

(I κ B α and β gal) was analyzed by Western blot and X-gal (5-bromo-4-chloro-3-indolyl- β -D-galactoside) staining (Sanes et al., 1986).

SiRNA transfection

Predesigned p65/RelA siRNA, TNFAIP3 siRNA and AllStars negative control siRNA, as well as HiPerfect transfection reagent, were purchased from Qiagen. Transfections were carried out according to the manufacturer's fast-forward transfection protocol. Experiments were performed 24–48 h following transfection. Knockdown of target genes was confirmed by qPCR and Western Blot.

Statistical analysis

Results are presented as means \pm standard error of mean (SEM). Statistical analysis was performed by one-way analysis of variance (ANOVA) followed by post-hoc Tukey multiple comparison test when F was significant, using Prism 5 for Mac software (GraphPad, Inc., La Jolla, CA, USA). Concentration-dependent effects were tested by Pearson correlation and linear regression analysis. Differences between groups were rated significant at a probability error (P) < 0.05 .

RESULTS

HGF upregulates A20 mRNA and protein expression in RPTEC

To determine whether HGF regulates A20 expression in RPTEC, we treated human RPTEC cultures with HGF at concentrations ranging from 10 to 50 ng/mL for 2 to 24 h, and determined A20 protein levels by Western blot analysis. Our results indicated that HGF significantly upregulated A20 protein expression in a time and concentration dependent manner (Pearson (r)=0.9703, r^2 = 0.9415, p <0.05), with the maximum (up to 3-fold) observed 6 h after the addition of 50 ng/mL of HGF (Fig. 1A&B). Correspondingly, A20 mRNA levels peaked at 1 h and declined by 3 h, following HGF treatment as determined by quantitative real-time PCR (qPCR) (Fig. 1C). HGF-mediated upregulation of A20 mRNA was temporally comparable to that of TNF α (100U/mL, i.e. 5ng/mL), a bona-fide transcriptional activator of A20 (Kunter et al., 2005). Indeed, A20 mRNA levels also peaked at 1 h and declined by 3 h following addition of TNF α , (Fig. 1C). At these concentrations, A20 protein and mRNA levels were significantly higher following TNF α as compared to HGF (Fig. 1C&D). This reflects the use of TNF α concentration ranging 100-fold its effective dose (ED50). This TNF α dose is required to produce the pro-inflammatory effects of this cytokine in vitro (Gong et al., 2008; Lee et al., 2006). In fact, at comparable HGF (Rubin et al., 1991) and TNF α ED50 (Matthews et al., 1987) (2 \times ED50) TNF α (2 U/mL) failed to upregulate A20 or for the matter pro-inflammatory genes, as depicted in Figure 5, whereas HGF (50–60 ng/mL) led to maximal upregulation of A20 mRNA (Supplementary Fig. 1). Accordingly, all further experiments were performed using 50 ng/mL of HGF and 100 U/mL of TNF α .

HGF-induced upregulation of A20 is NF- κ B-dependent

NF- κ B activation is usually required to promote A20 transcription in response to pro-inflammatory stimuli (Bach et al., 1997; Krikos et al., 1992; Laherty et al., 1992; Laherty et al., 1993). However, activation of NF- κ B by HGF remains controversial (Giannopoulou et al., 2008). Accordingly, we gauged the impact of inhibiting NF- κ B by either overexpressing I κ B α using recombinant adenovirus (rAd.) mediated gene transfer, or silencing the p65/RelA subunit of NF- κ B, on HGF-mediated upregulation of A20. Our results showed that overexpression of I κ B α significantly inhibited HGF-induced A20 protein and mRNA levels, as compared to non-transduced (NT) and rAd. β gal transduced RPTEC (Fig. 2A&B). A

similar inhibition was also achieved in these conditions for TNF α -induced upregulation of A20. Moreover, knockdown of p65 using a specific siRNA, which decreased p65 mRNA levels by almost 80% and its protein levels by >90% (Supplementary Fig. 2), significantly decreased HGF-induced upregulation of A20 mRNA, while AllStars negative control siRNA did not (Fig. 2C). Altogether, these results indicate that HGF-induced upregulation of A20 in RPTEC requires NF- κ B activation involving the p65/RelA subunit.

HGF engages NF- κ B signaling in a unique quantitative and qualitative manner

NF- κ B activation by its primary activators, TNF α , IL-1 and LPS, occurs either through the canonical or alternative pathways. The former depends on activation of I κ B kinase (IKK) subunits IKK β and IKK α , I κ B α phosphorylation, ubiquitination and degradation, p65/RelA phosphorylation, and translocation of the classical p50/p65 NF- κ B heterodimer to the nucleus (Hacker and Karin, 2006). The latter requires specific activation of IKK α leading to p100 processing, followed by nuclear translocation of the p52:RelB NF- κ B heterodimer (Hacker and Karin, 2006). Having already demonstrated the role of p65/RelA in inducing A20 transcription, we investigated whether HGF activates this subunit in RPTEC. Accordingly, we stimulated RPTEC with HGF (50 ng/mL) or TNF α (100 U/mL) as a positive control, and evaluated I κ B α phosphorylation and degradation, and p65 phosphorylation by Western blot. Given our focus on early responses to HGF treatment, we did not evaluate its impact on phosphorylation and degradation of I κ B β . Indeed, I κ B β is classically responsible for delayed and persistent activation of NF- κ B, in a stimulus and cell-type dependent manner that excludes TNF α ; i.e TNF α does not usually cause degradation of I κ B β (Thompson et al., 1995). HGF stimulation resulted in a moderate, and transient phosphorylation of I κ B α and p65, peaking within 5 min of HGF addition. This led to incomplete I κ B α protein degradation within 15 min of adding HGF (Fig. 3). TNF α treatment, in contrast, led to significantly greater phosphorylation of I κ B α and p65, followed by a complete degradation of I κ B α within 15 min of adding TNF α (Fig. 3). Although the time course of phosphorylation/degradation of p65 and I κ B α were fairly similar in HGF and TNF α -treated cells, these results highlight a significant quantitative difference between these two stimuli in engaging NF- κ B signals. We also evaluated phosphorylation of RelB, a central player in the non-canonical pathway. Our results showed a robust phosphorylation of RelB at Ser-552 within 15 min of adding HGF, while we noted very mild phosphorylation of RelB following TNF α , at least within the studied time frame (Supplementary Fig. 3A). These latter results demonstrate a significant qualitative difference between TNF α and HGF in engaging NF- κ B activation in RPTEC. However, in contrast to RelA, silencing of RelB that led to 80% knockdown of its RNA levels and >95% knockdown of its protein levels (Supplementary Fig. 2) did not impact HGF-induced upregulation of A20, indicating that it was not likely required for this process (Supplementary Fig. 3B).

Activation of PI3K/AKT is required for the upregulation of A20 by HGF

To further analyze the molecular mechanisms upstream of I κ B α and p65 phosphorylation that support HGF-induced upregulation of A20, we investigated which signaling pathway triggered by HGF is responsible for mediating NF- κ B activation and increased A20 expression. The serine/threonine protein kinase, AKT, has been identified both as an activation target of the HGF/cMet signaling pathway in different cell types including RPTEC (Gong et al., 2005; Liu, 1999; Nakagami et al., 2001), and also as one of the possible upstream kinases in the NF- κ B pathway (Ozes et al., 1999; Romashkova and Makarov, 1999). To investigate the role of AKT in HGF-induced activation of NF- κ B and induction of A20, we treated RPTEC with 50ng/mL of HGF for 5 min to 2 h, and evaluated AKT phosphorylation at Ser-473 by Western blot. Our data indicate that HGF-induced phosphorylation of AKT was fast and sustained (Fig. 4A). Phosphorylation of GSK3 β at Ser9, downstream of AKT activation, paralleled AKT phosphorylation in both its fastness

and robustness in response to HGF (Fig. 4B). This result is in keeping with Gong et al's report implicating AKT-dependent phosphorylation/inactivation of GSK3 β in mediating the anti-inflammatory effects of HGF (Gong et al., 2008). In contrast, treatment of RPTEC with TNF α failed to phosphorylate AKT at Ser473, and GSK3 β at Ser9, further highlighting the engagement of differential signaling pathways by HGF and TNF α Fig. 4B. Pretreatment of RPTEC with the PI3K inhibitor LY-294002 abolished phosphorylation of p65, the indicator of NF- κ B activation (Fig. 4C), and significantly decreased A20 protein levels, downstream of p65 (Fig. 4D). These results indicate that AKT may directly activate IKK α and IKK β to phosphorylate I κ B α , and we confirmed by Western blot that HGF treatment of RPTEC phosphorylates IKK α/β , in an AKT-dependent manner, since this phosphorylation was blocked by pre-incubation with LY-294002 (Supplementary Fig. 4). Given low I κ B α and high RelB phosphorylation in response to HGF, we surmise that the phospho IKK α/β band detected by Western blot is likely to be mainly composed of IKK α , the upstream kinase of RelB and less IKK β , the upstream kinase of I κ B α (Hacker and Karin, 2006). HGF-induced phosphorylation of RelB, was also AKT-dependent, as it was significantly blunted by LY-294002 (Supplementary Fig. 3C). Together, these results indicate that HGF-induced A20 upregulation in RPTEC occurs via an AKT/NF- κ B-dependent pathway.

HGF-induced activation of NF- κ B in RPTEC does not cause pro-inflammatory responses

In most cell types, activation of NF- κ B in response to inflammation concomitantly triggers pro-inflammatory (Lawrence, 2009; Pasparakis, 2009) and anti-inflammatory responses, including A20, which secondarily suppresses NF- κ B activation (Bach et al., 1997; Lawrence and Fong, 2010). Accordingly, an inflammatory burst normally precedes accumulation of functional levels of A20. We questioned whether HGF-mediated activation of NF- κ B would be accompanied by such a burst. Therefore, we investigated the effects of HGF on expression of the NF- κ B dependent chemokine MCP-1, and adhesion molecules, VCAM-1 and ICAM-1 in RPTEC (Collins et al., 1995; Kunter et al., 2005). Cells were treated for 3h with varying concentrations of HGF (25–200 ng/mL) or TNF α (5–200 U/mL), and MCP-1, VCAM-1 and ICAM-1 mRNA levels were measured. HGF treatment did not significantly affect basal expression of these pro-inflammatory molecules, even at the highest HGF concentration (Fig. 5). In contrast, and in agreement with previously published data, TNF α increased MCP-1, VCAM-1 and ICAM-1 mRNA levels in a concentration-dependent manner (MCP-1, Pearson $r=0.8761$, $r^2=0.2351$, $p<0.05$; VCAM-1 Pearson $r=0.9516$, $r^2=0.61$ $P<0.0001$; ICAM-1 Pearson $r=0.9679$, $r^2=0.644$, $p<0.001$) (Fig. 5). These results suggest that HGF may activate NF- κ B in a manner that favors the transcription of anti-inflammatory and protective NF- κ B dependent genes such as A20, but not that of the pro-inflammatory, NF- κ B dependent genes MCP-1, VCAM-1 and ICAM-1. Our finding that HGF upregulates mRNA and protein levels of the NF- κ B dependent anti-apoptotic Bcl family member, A1/Bfl1, in RPTEC supports this hypothesis (Kunter et al., 2005) (Supplementary Fig. 5).

HGF inhibits TNF-induced pro-inflammatory response in RPTEC, in part via an A20-dependent mechanism

As stated, HGF's remarkable anti-inflammatory effect in RPTEC occurs partly through blocking NF- κ B activation (Gong, 2008). To determine whether HGF affects TNF α -induced inflammatory responses in RPTEC, we examined the effects of pre-treatment with HGF on TNF α -induced ICAM-1 upregulation. Our results indicate that 4 h pre-treatment of RPTEC with HGF significantly inhibited TNF α -induced upregulation of ICAM-1 mRNA (Fig. 6). In contrast, simultaneous treatment of RPTEC with HGF and TNF α failed to reduce ICAM-1 upregulation by TNF α (data not shown), suggesting that HGF-dependent inhibition of TNF α -induced inflammatory response may require *de novo* protein synthesis. Having demonstrated that maximal HGF-induced upregulation of A20 occurs at a time similar to the

HGF pre-treatment time required to achieve inhibition of TNF α -induced upregulation of ICAM-1, we questioned whether HGF-induced A20 contributed to the anti-inflammatory effect of this growth factor in RPTEC.

To test this hypothesis, we silenced A20 expression with a specific siRNA that knocked down HGF-induced A20 mRNA upregulation by >90% and TNF α -induced upregulation by almost 50% (Supplementary Fig. 6), and evaluated the impact of HGF upon TNF α -induced upregulation of ICAM-1 mRNA in both conditions. Our results indicated that A20 silencing significantly reduces HGF's inhibitory effect upon TNF α -mediated upregulation of ICAM-1 mRNA (Fig. 6). This result demonstrates that A20 actively mediates the anti-inflammatory effects of HGF in RPTEC. Remarkably, A20 silencing significantly upregulated ICAM-1 mRNA levels in basal condition, and following HGF and TNF α treatment of RPTEC (Fig. 6), which highlights A20's ability to maintain basal anti-inflammatory status, prevent an inflammatory response following HGF treatment, and control TNF α pro-inflammatory responses. In support of this latter, we demonstrate that A20 silencing in RPTEC significantly increases ICAM-1 and MCP1 basal but also TNF α -induced mRNA levels (Supplementary Fig. 7).

Discussion

Inflammation is a major pathological finding in a variety of chronic kidney diseases and acute renal failure (Segeer et al., 2000). In the latter, it manifests primarily in ischemia/reperfusion injury, often following renal transplantation (Gueler et al., 2004). Rapid activation of NF- κ B follows ischemic events in the kidney, resulting in increased production of pro-inflammatory cytokines and chemokines, and culminating in vascular compromise and kidney damage (Cao et al., 2004; Linas et al., 1988). It is well appreciated that RPTEC are the most susceptible to this injury, which results in tubular necrosis. In fact, RPTEC contribute to their demise by acquiring a pro-inflammatory phenotype, promoting the local inflammatory milieu (Bonventre and Zuk, 2004). Therefore, controlling inflammation, particularly in RPTEC, is an attractive therapeutic strategy to reverse acute kidney injury, halt chronic disease, and protect renal allografts.

Our laboratory has explored the anti-inflammatory effect of A20 in different cell types. We have shown that this protein exerts potent NF- κ B inhibitory effects in RPTEC, shutting down the upregulation of pro-inflammatory molecules such as MCP-1, and ICAM-1 (Kunter et al., 2005). We surmised that A20-based therapies could reduce inflammation, ultimately preventing renal dysfunction. As evidence, overexpression of A20 in rat kidneys using rAd-mediated gene transfer protects from acute tubular necrosis following renal ischemia (Lutz et al., 2008). However, translation of A20 gene-therapies to the clinic has proven difficult given the toxicity of these viral vectors (2002). Our laboratory, therefore, has been exploring means to induce A20 in RPTEC without prompting a concurrent inflammatory response (Bach et al., 1997).

Our results presented herein demonstrate that this goal could be achieved by HGF. Treatment of RPTEC with HGF promoted A20 transcription and protein expression in a time and concentration-dependent manner. HGF-induced A20 transcription in human primary RPTEC, echoes similar results obtained in the mammary MCF7 tumor cell line (Leroy et al., 2006). Interestingly, HGF also induced A20 transcription in primary human coronary artery endothelial cell (EC) but not smooth muscle cells despite similar expression of the HGF receptor, c-Met, in these cells (Nakamura et al., 1995) (Supplementary Fig. 8). This indicates a cell-type specific effect of HGF that could be exploited for achieving differential therapeutic effects. Remarkably, our data also demonstrates that HGF upregulates A20 in RPTEC without upregulating the pro-inflammatory molecules MCP-1,

VCAM-1 and ICAM-1. This held true even at concentrations 4-fold higher than those upregulating A20 expression. In complete contrast, pro-inflammatory mediators like TNF α upregulated MCP-1, VCAM-1 and ICAM-1 concurrently with A20.

As stated earlier, A20 is a NF- κ B dependent gene activated through the canonical NF- κ B pathway in response to pro-inflammatory cytokines. We questioned whether NF- κ B activation was also required for HGF-mediated activation of A20. Our results showed unequivocally that it was NF- κ B dependent, since over-expression of the NF- κ B repressor I κ B α and silencing of p65 inhibited HGF-dependent A20 upregulation. However, HGF-triggered signaling differed qualitatively and quantitatively from that of TNF α . HGF treatment of RPTEC induced very mild I κ B α and p65 phosphorylation paralleling slower and significantly lower I κ B α degradation, compared to TNF α . These results are in agreement with previous results obtained in the MLP29 liver cell line (Muller et al., 2002), and with earlier studies implicating HGF/cMet signaling in activating the canonical NF- κ B pathway (Fan et al., 2005). In contrast, HGF promoted quicker and stronger phosphorylation of RelB Ser-552 than TNF α . HGF-mediated phosphorylation of RelB was likely not implicated in A20 upregulation, since RelB silencing did not affect it. This result demonstrates qualitative differences between HGF and TNF α engagement of non-canonical NF- κ B signaling molecules. Phosphorylation of Ser-552 of RelB is a potential tag for its degradation (Marienfeld et al., 2001), which could protect from kidney injury since RelB silencing in the kidney was shown to prevent renal ischemia/reperfusion injury in mice (Feng et al., 2009). However, we did not observe any loss of RelB following its phosphorylation by HGF in RPTEC, at least within the time limit of our experiments (data not shown).

Having established that HGF-induced upregulation of A20 relied on the activation of the canonical NF- κ B pathway, we searched for the pertinent upstream kinase (Dejardin, 2006; Scheiderei, 2006). AKT is a recognized activator of both canonical and non-canonical NF- κ B pathways (Gustin et al., 2006; Ozes et al., 1999; Sizemore et al., 1999). It is also an activation target of HGF/cMet in different cell types (Khwaja et al., 1998; Kroening et al., 2010; Muller et al., 2002; Suzuki et al., 2000). We identified AKT as key to NF- κ B activation and subsequent A20 upregulation in HGF-treated RPTEC. Indeed, inhibition of PI3K/AKT by LY294002 blocked phosphorylation of RelB and abolished HGF-mediated IKK α / β and p65-Ser536 phosphorylation, thereby inhibiting A20 upregulation. These novel data in primary human RPTEC parallel AKT's role in protecting breast cancer, glioma and canine kidney cell lines from apoptosis (Fan et al., 2005). However, they contradict results showing that ERK1/2 and p38 MAPK, but not AKT, are required for HGF-induced activation of NF- κ B in the MLP29 liver cell line (Muller et al., 2002), which again emphasizes cell type specificity. Interestingly, HGF treatment of RPTEC led to stronger and more prolonged AKT phosphorylation when TNF α almost did not impact AKT phosphorylation levels in RPTEC, further underlining the differences between HGF and TNF α signaling in these cells (Fig. 4B).

Furthermore, pre-incubation of RPTEC with HGF for 4 h significantly prevented TNF α -induced upregulation of ICAM-1 expression, while simultaneous exposure to both HGF and TNF α failed to do so, likely indicating the need for *de novo* synthesis of an anti-inflammatory protein by HGF. These findings agree with previous studies showing that HGF decreases TNF α -mediated upregulation of the NF κ B-dependent pro-inflammatory chemokines, RANTES and MCP-1 and adhesion molecule E-Selectin in the human RPTEC cell line, HKC8, and in human umbilical vein EC (Gong et al., 2006b; Gong et al., 2008). However, in contrast with our data, this inhibitory effect of HGF could be achieved with short (30 min) or no pre-incubation period, suggesting an alternative mechanism. In fact, our results indicate that in primary human RPTEC, the anti-inflammatory effect of HGF depends

significantly on its ability to upregulate A20, given 4 h pre-treatment, *i.e.* to allow for *de novo* synthesis of the A20 protein. Whereas, Gong *et al.* showed that the anti-inflammatory effect of HGF in HKC8 cells correlated with an AKT-dependent inhibition of GSK3 β activity, which inhibits p65/RelA phosphorylation at the GSK3 β target residue Ser-468, an event required for the activation of NF- κ B dependent pro-inflammatory MCP-1 and RANTES, but not protective I κ B α and Bcl-2 (Gong *et al.*, 2006b; Gong *et al.*, 2008). Since this effect is post-translational, it understandably did not require prolonged pre-incubation with HGF. In accordance with these data, we confirm that HGF inactivates GSK3 β as a result of its phosphorylation by AKT, which could certainly contribute to the anti-inflammatory effect of HGF, *i.e.* prevent HGF-induced upregulation of MCP-1, VCAM-1, and ICAM-1. We are also exploring whether GSK3 β inactivation is implicated in HGF-induced A20 upregulation in RPTEC.

Furthermore, A20 silencing significantly increased peak and basal expression levels of ICAM-1, used a surrogate for inflammation, despite pretreatment of RPTEC with HGF prior to TNF α , further confirming the importance of A20 in maintaining a non-inflammatory homeostatic state in RPTEC. Altogether, these data stress the diversity of seemingly independent HGF targets, underscoring its profound anti-inflammatory effects.

HGF treatment of RPTEC also led to upregulation of A1, an NF- κ B dependent anti-apoptotic Bcl family member, that, as we have shown earlier, exerts potent anti-apoptotic effects in RPTEC (Kunter *et al.*, 2005). This result highlights HGF's ability to activate NF- κ B in a way that promotes transcriptional upregulation of anti-inflammatory and anti-apoptotic but not pro-inflammatory genes, creating an enduring protection against a later inflammatory insult. These results support data from Bendinelli *et al.* showing that 24 h treatment with HGF decreases NF- κ B transactivation in chondrocytes by increasing I κ B α expression (Bendinelli *et al.*, 2010). However, these authors did not show whether or not this was related to an initial activation of NF- κ B, as we have for A20.

In summary, this study unravels and characterizes a novel, HGF-induced pathway that upregulates the potent anti-inflammatory protein A20 in human RPTEC without concurrent up-regulation of pro-inflammatory molecules. Furthermore, it shows that this upregulation of A20 contributes to HGF's anti-inflammatory capacity. While it requires NF- κ B activation, this process differs qualitatively, quantitatively and kinetically from that initiated by TNF α . Work is underway to further delineate these differences, which could help us identify novel therapeutic targets. Importantly, these data reconcile conflicting literature demonstrating both activation and inhibition of NF- κ B by HGF/cMet signaling. We clearly show the bimodal ability of HGF in human primary RPTEC, to activate the protective arm of NF- κ B and enhance the anti-inflammatory defenses of the cell, preventing full-blown responses to later inflammatory insults.

Clinically, these data unveil the possibility of increasing renal levels of A20 without causing inflammation allowing for the prevention of acute and chronic kidney diseases. It would also be ideal for kidney preconditioning to avoid ischemic and inflammatory injury in the peri-transplant period, thereby reducing primary non-function and incidence of acute rejection. HGF-based therapies are already available, having been implemented in experimental kidney disease and renal transplant models (Gong *et al.*, 2004; Mizuno *et al.*, 2001; Mizuno and Nakamura, 2004; Yamada, 2005; Zhang *et al.*, 2008); and in early clinical trials treating kidney disease and ischemic ulcers of vascular diseases (Shigematsu *et al.*, 2010). The fact that HGF also induces A20 in EC is likely to further improve renal protection, especially in the context of ischemic injury. However, we must acknowledge the potential carcinogenic effects of HGF (Comoglio *et al.*, 2008) and its cell-type specificity. With these caveats, we aim to better delineate the molecular targets of HGF and isolate differences in affecting NF-

κ B activation pathways, to ultimately identify more selective and safer therapies. We envision preferential upregulation of the anti-inflammatory protein A20 as highly significant in this search.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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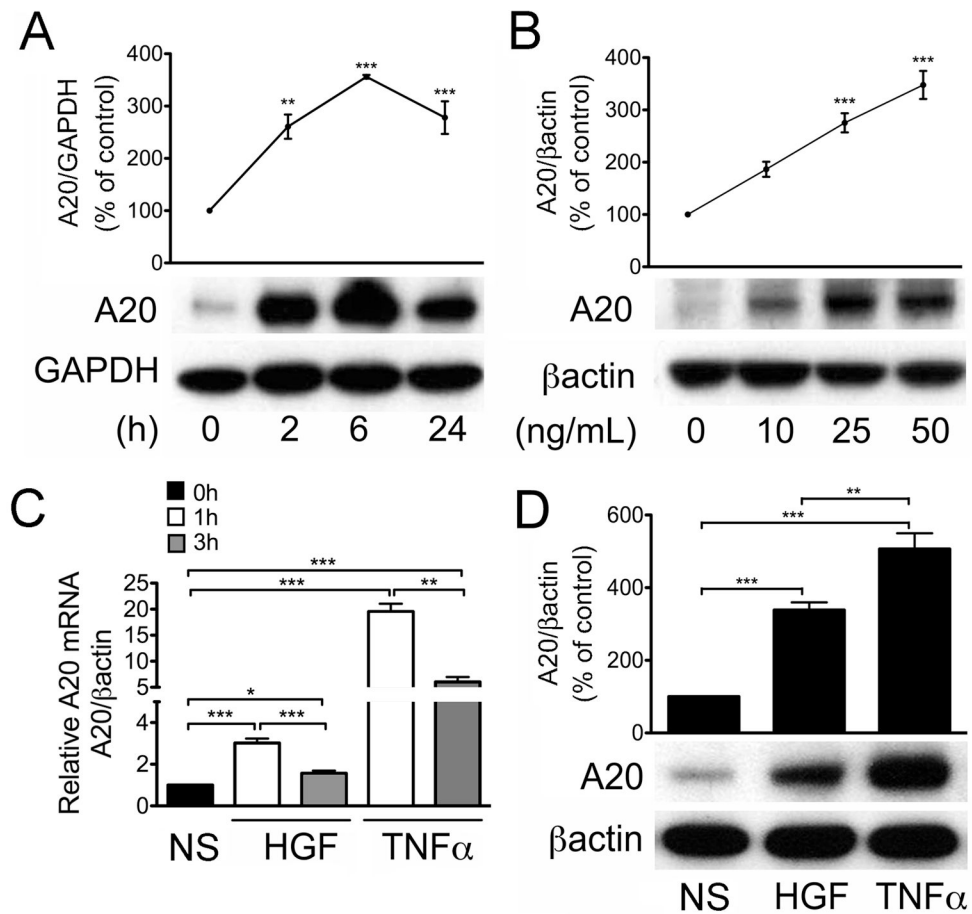


Figure 1. HGF upregulates A20 mRNA and protein levels in RPTEC

Western blots of RPTEC treated with **A.** HGF (50 ng/ml) for 2 to 24 h (time course); or **B.** HGF (10 to 50 ng/ml) for 6 h (dose curve). Cell lysates were run on PAGE, immunoblotted with anti-A20 antibody and re-probed with anti-GAPDH or anti- β actin antibody to correct for loading and allow semi-quantitative evaluation of the data by densitometry. Results of corrected densitometry, presented as percentage of control non-stimulated (NS) cells are expressed as mean \pm SEM of 4–5 independent experiments. **C.** Relative A20 mRNA levels measured by qPCR in human RPTEC cultures stimulated for 1 and 3 h with HGF (50 ng/ml) or TNF α (100 U/ml). Histograms represent the statistical analyzes of relative mRNA levels after normalization by β actin. Results are expressed as mean \pm SEM of 4 independent experiments. **D.** Corresponding Western blot of RPTEC treated with HGF (50 ng/ml) or TNF α (100 U/ml) for 6 h; Results of corrected densitometry, presented as percentage of control non-stimulated (NS) cells are expressed as mean \pm SEM of 4 independent experiments. * p <0.05, ** p <0.01 and *** p < 0.001.

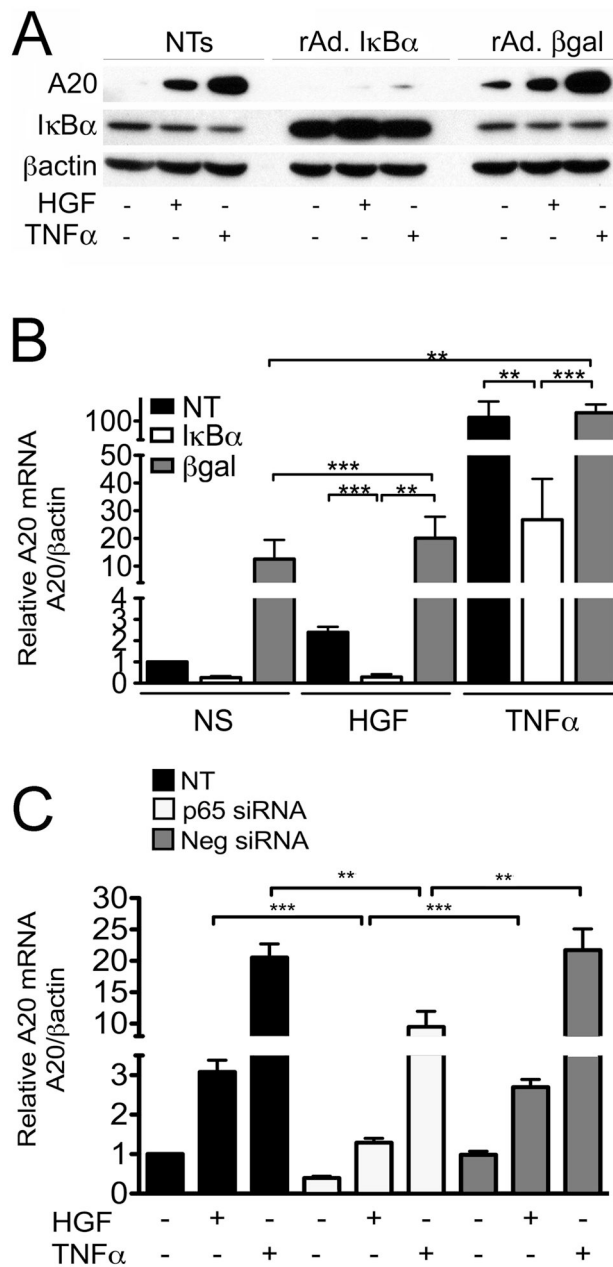


Figure 2. NF- κ B is required for HGF-induced upregulation of A20 in RPTEC
 Overexpression of I κ B α by means of rAd. mediated gene transfer (rAd.I κ B α) significantly inhibits HGF (50 ng/ml), and TNF α (100 U/ml)-induced upregulation of **A.** A20 protein in human RPTEC, as shown by Western blot analysis 6 h following treatment. Cell lysates were immunoblotted with antibodies to A20, I κ B α (to confirm transgene expression) and β actin (as loading control). Results shown are representative of 3 independent experiments.; and **B.** A20 mRNA levels, as evaluated by qPCR 1 h following treatment. Graphs represent the statistical analyzes of relative A20 mRNA levels after normalization by β actin. Results are expressed as mean \pm SEM of 4 independent experiments. A20 mRNA levels of non-stimulated (NS) RPTEC served as basal values. In both **A** and **B**, control cells were either non-transduced (NT) or transduced with the control adenovirus, rAd. β gal. * p <0.05 ** p <0.01, and *** p < 0.001. **C.** p65/RelA silencing in RPTEC cultures significantly inhibits

HGF (50 ng/ml) and TNF α (100 U/ml)-induced upregulation of A20 mRNA as shown by qPCR, 1 h following treatment. Graphs represent the statistical analyses of relative A20 mRNA levels after normalization by β actin. Results are expressed as mean \pm SEM of 3–5 independent experiments. Non-transfected (NT) or AllStars negative siRNA (Neg siRNA) served as controls. ** $p < 0.01$ and *** $p < 0.001$.

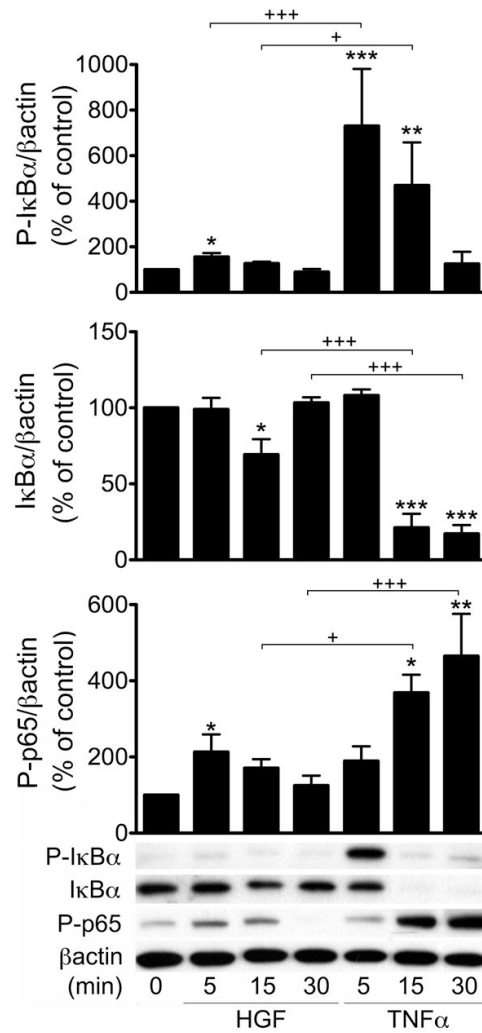


Figure 3. HGF and TNF α differently activate NF- κ B in RPTEC

Cell lysates from RPTEC treated with HGF (50 ng/ml) or TNF (100 U/ml) for 5–30 min were immunoblotted with antibodies to phospho-I κ B α (Ser32; P-I κ B α), I κ B α , phospho-65 (Ser536; P-p65). Results demonstrated significantly less I κ B α phosphorylation and degradation, and less p65 phosphorylation in HGF-treated versus TNF α -treated RPTEC. Immunoblotting with the housekeeping protein β actin was used to correct for loading and allow semi-quantitative evaluation of the data by densitometry. Results of corrected densitometry, presented as percentage of control non-stimulated (NS) cells are expressed as mean \pm SEM of 3–5 independent experiments. * p <0.05, ** p <0.01 and *** p < 0.001 when comparing with non-stimulated RPTEC, and + p < 0.05, +++ p <0.001 when comparing HGF vs. TNF α .

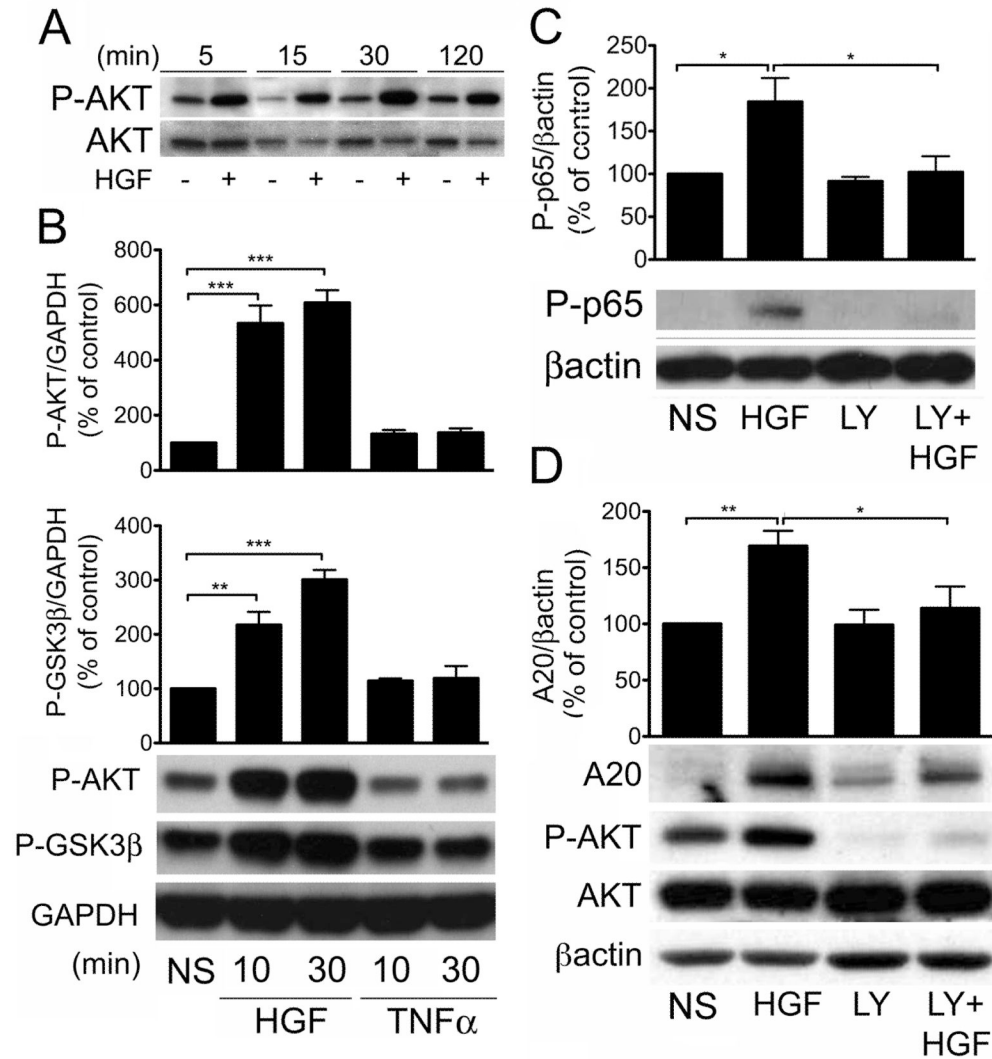


Figure 4. HGF-induced upregulation of A20 in RPTEC is AKT-dependent

A. Cell lysates from RPTEC treated with HGF (50 ng/ml) for 5–120 min were immunoblotted with antibodies to phospho-AKT or total-AKT. Results showed fast, robust, and sustained phosphorylation of AKT following treatment with HGF. Results shown are representative of 3 independent experiments. **B.** HGF-induced phosphorylation of GSK3β, marking its inactivation by AKT, was significantly upregulated 10 and 30 min following treatment with HGF but not TNFα, as shown by Western blot analysis using a phospho-GSK3β antibody (P-GSK3β). **C.** HGF-induced phosphorylation of p65 was significantly reduced when RPTEC were pre-incubated with the 10 μM of the PI3K inhibitor, LY294002, for 30 min prior to adding 50 ng/ml HGF for 15 min, as shown by Western blot analysis using a phospho-p65 specific antibody (P-p65). **D.** Similarly, HGF-induced upregulation of A20 was significantly inhibited when RPTEC were preincubated with 10 μM of LY294002 for 30 min prior to adding 50 ng/ml HGF for 6 h, as shown by Western blot analysis using an anti-A20 antibody. Cells lysates were also immunoblotted with anti P-AKT and AKT antibodies to control for LY294002 activity. In B, C, and D GAPDH or βactin immunoblots were used to correct for loading and allow semi-quantitative evaluation of the data by densitometry. Results of corrected densitometry, presented as percentage of control non-

stimulated (NS) cells, are expressed as mean \pm SEM of 4 (B&C), and 6–8 (D) independent experiments * $p < 0.05$, ** $p < 0.01$. *** $p < 0.001$.

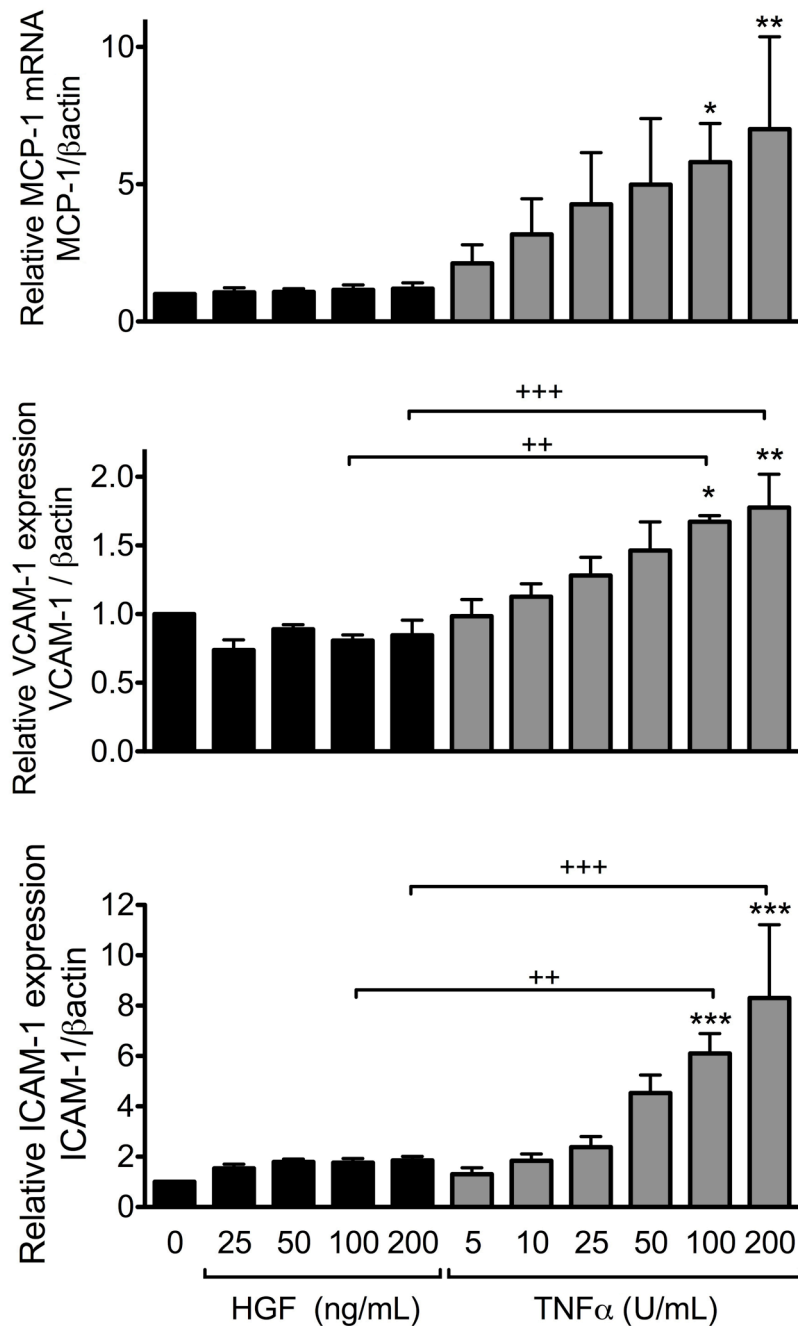


Figure 5. HGF does not upregulate pro-inflammatory cell surface adhesion molecules ICAM-1 and VCAM-1 and chemokine MCP-1 in RPTEC

qPCR analysis was performed on total mRNA isolated from RPTEC cultures treated with HGF (25–200 ng/mL) or TNF α (5–200 U/mL) for 3h. Results demonstrate a significant concentration-dependent upregulation of MCP-1, VCAM-1 and ICAM-1 following treatment with TNF, while no significant upregulation of these genes was noted following treatment of RPTEC with HGF. Graph shows the statistical analyses of relative MCP-1, VCAM-1 and ICAM-1 mRNA levels after normalization by the housekeeping gene β actin. Results are expressed as mean \pm SEM of 3–6 independent experiments. * p <0.05, ** p < 0.01

and *** $p < 0.001$ when comparing with non-stimulated RPTEC; and ++ $p < 0.01$, +++ $p < 0.001$ when comparing RPTEC stimulated with HGF vs. $\text{TNF}\alpha$.

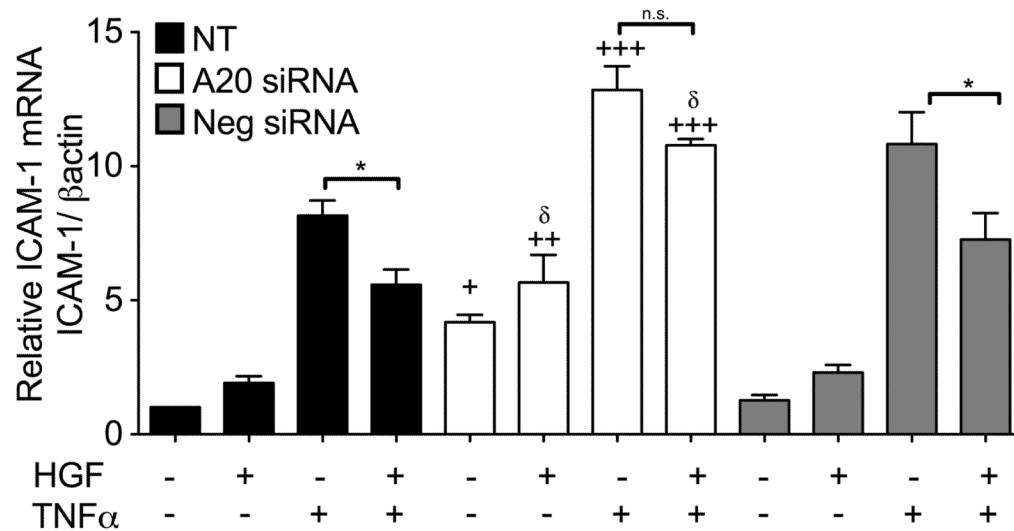


Figure 6. HGF-induced upregulation of A20 contributes to the anti-inflammatory effect of this growth factor in RPTEC

RPTEC were transfected with A20 silencing RNA (A20 siRNA) or AllStars negative control siRNA (Neg siRNA). Twenty-four h later, cells were pre-treated with HGF (50 ng/mL) for 4 h, then stimulated with TNFα (100 U/mL) for 3h. RNA was extracted and ICAM-1 mRNA levels were determined by qPCR. Our results indicate that pre-treatment of RPTEC with HGF significantly attenuates TNFα-induced upregulation of ICAM-1 in non-transfected (NT) and AllStars negative control siRNA transfected RPTEC. A20 silencing significantly decreases the ability of HGF to inhibit TNFα-induced upregulation of ICAM-1. Moreover, A20 silencing significantly increases basal, as well as HGF, and TNFα-induced upregulation of ICAM-1 mRNA. Graphs shown represent the statistical analyses of relative ICAM-1 mRNA levels after normalization by βactin. Results are expressed as mean ± SEM of 4–6 independent experiments. * $p < 0.05$ when comparing RPTEC stimulated with TNFα vs. cells pre-incubated with HGF prior to TNFα. + $p < 0.05$, ++ $p < 0.01$, +++ $p < 0.001$ when comparing with corresponding values in NT RPTEC, and δ $p < 0.05$ when comparing with corresponding values in AllStars negative control siRNA transfected RPTEC.