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Extracellular Hsp90 mediates an NF- κ B dependent inflammatory stromal program: Implications for the prostate tumor microenvironment

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Abstract

Background—The tumor microenvironment (TME) plays an essential role in supporting and promoting tumor growth and progression. An inflammatory stroma is a widespread hallmark of the prostate TME, and prostate tumors are known to co-evolve with their reactive stroma. Cancer-associated fibroblasts (CAFs) within the reactive stroma play a salient role in secreting cytokines that contribute to this inflammatory TME. Although a number of inflammatory mediators have been identified, a clear understanding of key factors initiating the formation of reactive stroma is lacking.

Methods—We explored whether tumor secreted extracellular Hsp90 alpha (eHsp90 α) may initiate a reactive stroma. Prostate stromal fibroblasts (PrSFs) were exposed to exogenous Hsp90 α protein, or to conditioned medium (CM) from eHsp90 α -expressing prostate cancer cells, and evaluated for signaling, motility, and expression of prototypic reactive markers. In tandem, ELISA assays were utilized to characterize Hsp90 α -mediated secreted factors.

Results—We report that exposure of PrSFs to eHsp90 upregulates the transcription and protein secretion of IL-6 and IL-8, key inflammatory cytokines known to play a causative role in prostate cancer progression. Cytokine secretion was regulated in part via a MEK/ERK and NF- κ B dependent pathway. Secreted eHsp90 α also promoted the rapid and durable activation of the oncogenic inflammatory mediator signal transducer and activator of transcription (STAT3). Finally, eHsp90 induced the expression of MMP-3, a well-known mediator of fibrosis and the myofibroblast phenotype.

Conclusions—Our results provide compelling support for eHsp90 α as a transducer of signaling events culminating in an inflammatory and reactive stroma, thereby conferring properties associated with prostate cancer progression.

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INTRODUCTION

The host microenvironment is profoundly altered during tumor growth, acquiring a reactive and inflammatory phenotype inextricably linked to its tumor supportive activity. This is observed in prostate cancer (PCa) wherein tumors co-evolve with their reactive or inflammatory tumor stroma (1). Lesions indicative of reactive stroma appear early in PCa etiology, originating during prostatic intraepithelial neoplasia (PIN) (2), a pre-neoplastic condition affecting prostatic ducts. The reactive stroma initiated during PIN represents a critical step in PCa adenocarcinoma progression, a premise supported by its co-existence with advanced prostate cancer tissue (2), and as a predictor of reduced recurrence free survival (3,4). Although the key tumor supportive mechanisms of reactive stroma are not well defined, reactive cancer associated fibroblasts (CAFs) play a central role in this process (5–7). The conversion of stromal fibroblasts to a reactive myofibroblast or CAF-like phenotype is triggered either by direct contact with cancer cells or upon exposure to tumor secreted factors (8,9). As CAFs are prominent modifiers of tumor progression, it is essential to define factors driving this phenotype.

The similarity of reactive stroma in both wound repair and cancer (10,11) is in large part due to the secretion of angiogenic and inflammatory cytokines from reactive stromal fibroblasts (6–8). Interleukins, potent mediators of inflammatory processes, are commonly found in the tissue microenvironment of both wound repair and cancer. In particular, IL-6 and IL-8 are multifunctional chemokines associated with prostatic reactive stroma (9,12). The expression of IL-6 and IL-8 is regulated in part by NF-kappaB (NF-κB) (13,14), a transcriptional master regulator of inflammatory processes, many of which play widespread and causative roles in cancer progression (15,16). Importantly, NF-κB has also been identified as a prominent factor involved in conferring the reactive phenotype of CAFs (8). Similar to NF-κB, STAT3 is a transcriptional mediator of immune responses, inflammation and tumorigenesis (17). The NF-κB and STAT3 pathways collaborate in cancer progression and engage in extensive crosstalk, due in part to the production of IL-6, which serves as a major cytokine for STAT3 activation (18).

Emerging reports indicate that many tumor types secrete extracellular Hsp90α (eHsp90α), a normally intracellular molecular chaperone (19). We have shown that PCa cells with increased aggressiveness secrete more eHsp90α relative to their less aggressive counterparts (20), a trend mirrored in breast cancer cells (21). The presence of secreted eHsp90α in patients with tumor burden has been widely reported in diverse cancer types (21), including PCa (22), implicating a potential clinical role for eHsp90α secretion. Functionally, eHsp90α plays a role in sustaining cancer cell motility, invasion, and metastatic spread (21,23–26). We have previously demonstrated that eHsp90α signals in an autocrine pathway to initiate epithelial to mesenchymal transition (EMT) events in prostate epithelial cells (20). For this study, we explored whether eHsp90α may utilize paracrine signaling mechanisms to

modulate the tumor stroma. Several EMT-inducing growth factors, such as TGF β , exhibit autocrine and paracrine signaling modalities, and are potent modifiers of the tumor stroma. Although no prior reports demonstrate the ability of eHsp90 α to promote a tumor reactive stroma, eHsp90 α has been shown to stimulate the motility and subsequent wound healing of dermal fibroblasts (27,28) and initiate signaling in endothelial cells (29). We have also shown that eHsp90 α induces signaling in vascular endothelial cell types (30,31). Collectively, these findings indicate that eHsp90 has the potential to modify the properties of stromal cell types.

We now demonstrate that eHsp90 α initiates signaling events in prostate stromal cells (PrSCs), conferring properties of increased cell motility and elevated expression of molecular markers indicative of a reactive phenotype. Importantly, eHsp90 α increased the expression and secretion of the inflammatory mediators IL-6 and IL-8 in a pathway partially dependent upon NF- κ B. Evidence of NF- κ B activation was further validated by detection of eHsp90 α -dependent phosphorylation of the NF- κ B effector RelA/p65. An eHsp90 α -initiated inflammatory stromal phenotype was reinforced by the presence of STAT3 phosphorylation. Our findings support the notion that eHsp90 α transduces a cascade of signaling events to promote the conversion of normal PrSCs to reactive and inflammatory stromal cells with the characteristics of CAFs. Moreover, our findings indicate that tumor eHsp90 α utilizes paracrine signaling mechanisms to alter the TME, thereby enabling a niche favorable for subsequent cancer progression.

MATERIALS AND METHODS

Reagents and antibodies

Hsp90 α human recombinant protein was purchased from Enzo Life Sciences (ADI-SPP-776). The F-5 Hsp90 α and control peptide was provided by Wei Li (University of Southern California, Los Angeles, CA), and has been described elsewhere (32). The MEK/ERK inhibitor UO216 was purchased from Promega (V112A). MMP inhibitors GM6001 and the 2/9 inhibitor were obtained from EMD Millipore Chemicals (CC1010 and 444274). DMAG-N-oxide modified geldanamycin, (herein referred to as non-permeable GA, NPGA) was synthesized by Chris Lindsey and Craig Beeson (Pharmaceutical Sciences, Medical University of South Carolina, Charleston, SC). The NF- κ B inhibitor Bay11-7082 (B5556), and antibodies to Tenascin-C (HPA004823) and GAPDH (G9295) were from Sigma-Aldrich. Cell Signaling antibodies included P-AKT (4058), AKT (4685), P-ERK (4370), ERK (4695), P-p65 (3033), p65 (4764), P-STAT3 (9138), and STAT3 (4904). Antibodies to Vimentin (ab8978) and SMA (ab5694) were purchased from Abcam. FAP antibody (H00002191-M01) was purchased from Abnova. The cytokine array (ARY005 R&D) was from R&D Systems and the MMP array (ab134004) was from Abcam.

Cell culture

Human primary prostate stromal cells (PrSC; Clonetics) were grown in stromal cell growth medium (SCGM; Clonetics) and were used between passages 5 to 10. Immortalized normal human prostatic fibroblasts (NPFs) were obtained from Dr. Simon Hayward (33), (Vanderbilt-Ingram Cancer Center); 6015N and 5905N PrSFs were provided by Dr. David

Rowley (Baylor College of Medicine); immortalized PSC27 PrSFs were provided by Dr. Peter Nelson (34), (Fred Hutchinson Cancer Research Center). ARCaPE-LacZ and ARCapE-eHsp90 cells were generated as previously described (20). All cells were maintained at 37°C with 5% CO₂.

Western blots

Nuclear preparations were prepared as previously described (35). For whole cell lysates, cells were washed with phosphate-buffered saline, and lysates prepared in lysis buffer as previously described (20,36). All blots are representative of a minimum of two independent experiments.

Immunofluorescent microscopy

Cells were treated as indicated, fixed with 4% paraformaldehyde and permeabilized with 0.1% Triton X-100 in PBS. Immunofluorescent microscopy was performed as described (36).

qRT-PCR analysis

PrSFs were pretreated (6 hr) with the indicated inhibitors, followed by treatment with Hsp90 α protein, F-5 peptide, or conditioned medium (CM). Total RNA was extracted, converted to cDNA and amplified as previously described (20). Primers were purchased from IDT and sequences are as follows: MMP3 F-AGGTGTGGAGTTCCTGATGTTGGT R-TACAGCCTGGAGAATGTGAGTGGA; IL-6 F-AAGCCAGAGCTGTGCAGATGAGTA R-GCTGCGCAGAATGAGATGAGTTGT; IL-8 F-CCTTGTTCCACTGTGCCTTGGTTT R-GTGCTTCCACATGTCCTCACAACA; GAPDH F-TCGACAGTCAGCCGCATCTTCTTT R-ACCAAATCCGTTGACTCCGACCTT

Immunoassays

Equal numbers of PrSFs or PrSCs were starved overnight in low serum (0.1% FBS), and treated as indicated, also under in low serum conditions. Prior to treatment with Hsp90 α protein, F-5 peptide, or conditioned medium, cells were rinsed with PBS and incubated with freshly prepared treatments of similar medium for 16 hr. Conditioned medium was collected following brief centrifugation, and IL-6, IL-8, and MMP-3 concentrations were measured by ELISA (R&D Systems) according to the manufacturer's instructions.

Cell Migration Analysis

Cell migration analyses were performed as previously described (20,36).

Statistical analysis

Statistical significance was determined using one-way ANOVA followed by one-tailed student's T-tests. All cell motility and quantitative real time-PCR experiments were performed in triplicate. Data shown are presented as mean % S.D.; differences in treatment groups are defined as statistically significant at $p < 0.05$ value, as calculated from Student's t test.

RESULTS

eHsp90 α promotes ERK-dependent prostate stromal fibroblast cell motility

To investigate a possible paracrine function for eHsp90 α within the context of PCa, we utilized human primary PrSCs and PrSFs as representative models of stromal cell types found in the TME. We have shown that eHsp90 α stimulates ERK activation in prostate epithelial cells, an event required for cell motility (20). Addition of Hsp90 α protein to PrSCs elicited the robust and rapid activation of ERK, which was blocked by the ERK inhibitor UO126 (Fig. 1A). Treatment of cells with either the pan-MMP inhibitor GM6001 or a non-permeable geldanamycin (GA) derivative (NPGA) that specifically blocks eHsp90 function (20,25,36), similarly attenuated eHsp90-mediated ERK activity. We then investigated whether eHsp90 affected PrSC cell motility. As shown, eHsp90 enhanced PrSC motility by over 50% (Fig. 1B, Supp Fig. 1A). PrSC motility was comparably blocked by NPGA, UO126, and GM6001. Given that both NPGA and GM6001 blocked ERK phosphorylation, our results indicate that ERK activity is required for the pro-motility function of eHsp90 α . eHsp90 α -mediated ERK activation was also observed in normal prostatic fibroblasts (NPFs) (Fig. 1C), consistent with its ability to stimulate cell motility (Fig. 1D, Supp Fig. 1B). Moreover, this pro-motility function of eHsp90 α was similarly inhibited by UO126, GM6001 and NPGA, thereby supporting a shared mechanism of action. We also observed eHsp90-mediated ERK activation in the immortalized stromal fibroblast cell line, PSC27 (34) (Supp Fig. 1C).

eHsp90 α initiates molecular changes associated with a reactive stromal phenotype

We next asked whether exposure of PrSFs to eHsp90 α conferred expression of molecular markers associated with a reactive phenotype. Given that TGF- β can convert fibroblasts into reactive myofibroblasts or CAFs (37,38), we used TGF- β as a positive control. As expected, TGF- β induced smooth muscle actin (SMA) and tenascin C, whereas eHsp90 α modestly upregulated vimentin, SMA, fibroblast activation factor (FAP) and tenascin C (Fig. 2A). Therefore, eHsp90 α stimulated the expression of markers associated with a CAF-like phenotype, although the profile exhibited some differences from TGF- β -treated cells. We next utilized fluorescent microscopy to evaluate the cellular distribution of SMA. As shown, addition of eHsp90 α to NPFs induced SMA (Fig. 2B), which also appeared to be polymerized and recruited to stress fibers, consistent with changes associated with reactive stroma (39). To evaluate the effects of eHsp90 α within a more physiological context, we exposed stromal cells to the conditioned medium (CM) of cells expressing eHsp90 α , using ARCAPE tumor cells with low basal secretion (20) that were transduced with lentivirus expressing either LacZ (control) virus or eHsp90 α virus as described (20). The CM from a clonal population of ARCAPE-eHsp90 α that exhibited a 6-fold increase in eHsp90 α secretion over LacZ cells (Supp Fig. 2) similarly upregulated SMA expression. A similar analysis was performed for PSC27 cells. In keeping with these trends, eHsp90 α protein and CM from ARCAPE-eHsp90 α induced polymerization of SMA (Fig. 2C). In addition, we evaluated the activity of an eHsp90 α peptide (F-5 peptide) representing an N-terminally truncated Hsp90 α protein deficient in the ATPase domain (32) critical for Hsp90 chaperone activity (19). The F-5 peptide elicited similar effects, indicating that eHsp90 α -dependent chaperone function was not required for these effects. A control peptide (27-mer) derived

from the middle domain of Hsp90 α (32) had no effect. Notably, Hsp90 α protein, F-5 peptide, and Hsp90 α derived from ARCaPE CM dramatically altered PSC cell morphology and promoted the formation of multiple protrusions.

eHsp90 α initiates a pro-inflammatory phenotype in stromal prostate

fibroblasts—We next investigated whether eHsp90 confers an inflammatory phenotype resembling the PCa TME. Towards this end, NPFs were treated with Hsp90 α protein, and resultant CM was analyzed via a cytokine and MMP array. As shown, Hsp90 α induced the secretion of inflammatory cytokines IL-6 and IL-8, as well as the proteolytic enzyme MMP-3 (Fig. 3A). These findings were subsequently validated by ELISA analysis (Fig. 3B). Similar trends of eHsp90 α action were observed in PSC27 cells (Fig. 3C). The eHsp90 α -mediated effects were generally sustained over a 3-day treatment period, indicating the ability of eHsp90 α to initiate a durable response.

We have demonstrated that more aggressive prostate cancer cells exhibit increased secretion of Hsp90 α (20), raising the question of whether these cell types may have an increased propensity to impact the behavior of adjacent stromal cells. To assess the ability of tumor-derived eHsp90 to elicit a pro-inflammatory response, stromal cells were exposed to CM derived from either ARCaPE-LacZ or ARCaPE-eHsp90 α . As shown, IL-6 and IL-8 secretion was absent or nominal in both ARCaPE-LacZ and ARCaPE-eHsp90 α (Fig. 3D, note -NPF data sets), indicating that eHsp90 does not appreciably induce these cytokines in an autocrine manner in this cell line. Remarkably, a dramatic increase in both IL-6 and IL-8 secretion was observed following co-incubation of NPFs with CM from eHsp90 α -expressing ARCaPE-eHsp90 (Fig. 3D, compare left and right columns for +NPF data sets). These data demonstrate that eHsp90 α initiates paracrine signaling mechanisms to elicit a robust pro-inflammatory phenotype in PrSFs. We next evaluated whether the increased cytokine secretion was controlled at the transcriptional level. As shown, ARCaPE-eHsp90 CM modestly induced IL-6 and IL-8 message levels in NPFs (Fig. 3E). Thus, although transcriptional mechanisms participate in modulating increased cytokine secretion, the magnitude of the increase implicates additional collaborating mechanisms.

eHsp90 α promotes the stromal secretion of inflammatory mediators in a chaperone-independent manner—Hsp90 α recombinant protein has been shown to promote the motility of dermal fibroblasts (27). Moreover, the truncated Hsp90 α peptide (F-5) retains comparable pro-motility activity (28,40), indicating that eHsp90 α activity may be driven by chaperone-independent signaling events. Given that the F-5 peptide elicited molecular and morphologic changes consistent with a reactive phenotype (Fig. 2), we next investigated whether it was sufficient to elicit a pro-inflammatory response. As shown, while a control Hsp90 α peptide (32) had no effect, exposure of NPFs to F-5 peptide induced secretion of IL-6, IL-8 and MMP-3 (Fig. 4A). This analysis was expanded to additional prostate stromal cell lines, including PSC27 cells (Fig. 4B) and two primary prostate fibroblast lines (Figs. 4C, D). In all instances, F-5 promoted IL-6 and IL-8 secretion, and also induced modest MMP-3 secretion. These results confirm that eHsp90 α signaling activity is responsible for the secretion of these inflammatory mediators.

Hsp90 α regulates stromal IL-6 and IL-8 secretion via an ERK1/2 and MMP-2/9 dependent pathway

—We next focused on potential mechanisms involved in eHsp90 α -initiated cytokine secretion. Inhibition of either ERK or MMP-2/9 attenuated the ability of Hsp90 α protein to induce cytokine secretion, with blockade of eHsp90 α eliciting the most robust suppression (Fig. 5A). Inhibition of either ERK or MMP-2/9 similarly diminished the Hsp90 α F-5-mediated increase in cytokine secretion (Fig. 5B). As expected, NPGA was much less effective at blocking the activity of the F5 peptide relative to its ability to suppress the effects of eHsp90 α protein, due to the absence of the ATP binding site in the F5 peptide that is targeted by NPGA. We confirmed that MMP-2/9 was required for eHsp90 α -mediated ERK activation (Supplemental Fig. 3A). Interestingly, although eHsp90 α -ERK had no effect upon MMP-2/9 transcription (not shown), eHsp90 α -ERK appears to regulate MMP-3 at the transcriptional level, in an MMP-2/9 dependent pathway (Supplemental Fig. 3B). Finally, we demonstrate that blockade of ERK or MMP-2/9 reduced the transcriptional message levels of IL-6 and IL-8 in response to the F-5 peptide (Fig. 5C). Mirroring trends for the secretion data, NPGA did not appreciably dampen IL-6 and IL-8 mRNA levels induced by F-5.

NF- κ B signaling participates in eHsp90 α -mediated cytokine secretion

—Given that both IL-6 and IL-8 are regulated in part by NF- κ B-dependent signaling mechanisms (41), we investigated whether NF- κ B served as a molecular effector for Hsp90 α -mediated cytokine secretion. In normal unstimulated cells, NF- κ B is maintained in an inactive cytoplasmic complex by its association with the I κ B kinases. In the canonical activation pathway, inflammatory cytokines such as TNF α promote the phosphorylation and degradation of I κ B proteins, resulting in the subsequent phosphorylation and nuclear translocation of p65/RelA. As shown, pharmacologic inhibition of NF- κ B activity attenuated the eHsp90 α -mediated induction of IL-6 and IL-8 mRNA and protein secretion (Figs. 6A, 6B), highlighting a role for NF- κ B in eHsp90 α -dependent cytokine expression. We next evaluated whether NF- κ B signaling participated in eHsp90 α -dependent NPF cell motility. As shown, NF- κ B inhibition (Bay11-7082) dramatically impaired the pro-motility function of F-5 peptide (Fig. 6C). These findings support the notion that NF- κ B plays a functional role in promoting the reactive inflammatory phenotype of prostate stromal cells.

To further confirm the connection between eHsp90 and NF- κ B activation, we examined the phosphorylation status of the NF- κ B subunit p65. Using TNF α as a positive control, we demonstrate the rapid phosphorylation of p65 (Fig. 6D). We then evaluated the ability of eHsp90 protein and F-5 peptide to activate p65. As shown, both treatments similarly elicited the rapid and durable phosphorylation of p65. The control peptide, which was not effective in promoting cytokine secretion, was also unable to induce p65 phosphorylation. Extensive crosstalk exists between IL-6 and STAT3 signaling (42), due in large part to the ability of IL-6 to function as a major effector for STAT3 (18,43). STAT3 activation requires phosphorylation (Tyr705), which mediates its dimerization and subsequent nuclear entry. Using IL-6 treatment as a positive control, we show the expected phosphorylation of STAT3 (Fig. 6E). Surprisingly, both eHsp90 α protein and the F-5 peptide rapidly induced STAT3 phosphorylation, and moreover, this activation was sustained over a 24 hr period, indicative of a durable and chronic inflammatory response. Treatment with UO126 did not attenuate

F-5-mediated STAT3 phosphorylation (not shown) indicating the involvement of additional pathways.

DISCUSSION

We report the novel finding that eHsp90 α activates and initiates crosstalk between proteolytic effectors (MMP-2/9) and signaling effectors (MEK/ERK) to activate NF- κ B and consequently induce expression of the inflammatory mediators IL-6 and IL-8. To our knowledge, our findings are the first to demonstrate the role of eHsp90 α as an initiator of NF- κ B signaling in prostate stromal fibroblasts. Our data also indicate that these inflammatory changes were inextricably linked with the conversion of stromal cells to a reactive phenotype. This was supported by the increased expression of molecular markers consistent with a reactive phenotype, coupled with profound changes in cell morphology, including the recruitment of SMA to stress fibers. Moreover, MMP-2/9, ERK and NF- κ B collaborated to regulate eHsp90 α -driven stromal cell motility. Thus, our collective findings indicate that an eHsp90-initiated MMP-2/9-ERK cascade is critical for the NF- κ B driven stromal inflammation and fibroblast activation.

The ability of full length eHsp90 α protein and ATP-ase deleted F-5 eHsp90 α peptide to comparably activate ERK and NF- κ B, and to elicit IL-6/IL-8 cytokine secretion, supports the notion that eHsp90 α 's pro-inflammatory action is elicited via signaling modalities independent of chaperone function. A working model of our findings is presented (Fig. 7). eHsp90 α plays a major role as a facilitator of stromal ERK activation. This eHsp90-ERK axis is likely reinforced by several mechanisms, including signaling via the LRP1 receptor (20). Herein, we demonstrate that MMP-2/9 is essential for eHsp90-mediated ERK activation. Our data are consistent with the current model wherein eHsp90 α activates MMP-2/9 proteolytic activity via protein-protein interactions (20,23,26,44,45). Although possible that eHsp90 α may similarly modulate MMP-3 proteolytic activity (46), our findings indicate that eHsp90-ERK signaling induces MMP-3 transcription. While MMP-3 inhibition had minimal impact upon IL-6/IL-8 secretion (not shown), MMP-3 is a well-known mediator of fibrosis and the myofibroblast phenotype (47). Our findings indicate that an eHsp90 α -MMP-2/9-ERK cascade plays a dominant role in NF- κ B-mediated induction of IL-6/IL-8. Although further studies are required to discern the precise mechanism of collaboration, MEK/ERK and p65/NF- κ B have well-documented roles as collaborators of inflammatory signaling and tumorigenesis (48) and MEK/ERK has been identified as an upstream regulator of NF- κ B (49). Additional mechanisms may also collaborate, such as eHsp90 α -LRP1 dependent interaction with I κ B kinases (50). We also demonstrate that eHsp90 induces the rapid and durable phosphorylation of STAT3. STAT3 may be activated by a diverse array of signaling mediators (51,52). As indicated, nuclear STAT3 and NF- κ B may collaborate to maintain the inflammatory milieu.

Secretion of pro-inflammatory cytokines IL-6 and IL-8 have been shown to drive a CAF-like phenotype in PrSFs (1,9). Given the importance and prevalence of these cytokines in the reactive stroma, and the emerging role of eHsp90 α in cancer progression, our findings support the notion that tumor-derived eHsp90 α may be a key factor in the maintenance of an inflammatory phenotype. The ability of eHsp90 α to activate stromal NF- κ B and STAT3,

well established master regulators of inflammatory responses, offers additional support for this premise. IL-6 is a major transducer of STAT3 signaling (18,42,43), and eHsp90 α elicited a durable activation of STAT3. Whereas transient STAT3 activation is a component of normal physiological responses, sustained STAT3 hyperactivity is associated with chronic inflammation and malignant transformation (53). Thus, the ability of eHsp90 α to sustain STAT3 activation may have implications for transformation and progression. The rapid activation of STAT3 by eHsp90 α argues against transcriptional mechanisms driving the initial activation process.

The co-activation of NF- κ B and STAT3 provides convincing evidence that eHsp90 α initiates a complex interplay of potentially reinforcing signaling events. For example, although IL-6 is a downstream target of NF- κ B, it also participates in a feed-forward mechanism to sustain NF- κ B and STAT3 activation (54,55). The co-activation of these pathways would be expected to amplify and sustain a number of downstream events. Indeed, STAT3 activation is a known driver of MMP-3 expression, herein also shown to be an eHsp90 α transcriptional target. The complexity of this signaling is expected to play a prominent role in altering tumor-stromal signaling. Our findings support this premise, as eHsp90 α -dependent signaling in epithelial cells resulted in a cytokine milieu with the ability to robustly increase stromal secretion of IL-6 and IL-8. The CM from eHsp90 α -expressing epithelial cells was also shown to confer morphological and molecular changes to prostate stromal fibroblasts consistent with adoption of a reactive phenotype. These data illustrate that eHsp90 dramatically modifies stromal-tumor signaling events to reinforce an inflammatory tumor microenvironment. This inflammatory function for eHsp90 α has potentially broad implications for its role in prostate cancer progression. Nuclear NF- κ B is found in almost half of all prostate cancers and is associated with progression, metastasis and recurrence (56–58), with similar functional roles bestowed upon the NF- κ B gene products IL-6 and IL-8 (59–61). STAT3, which is activated in a majority of human PCa (62), also appears to play a prominent role. STAT3 collaborates with IL-6 in PCa progression (63,64), and is sufficient to execute many of the tumorigenic effects of IL-6 (65).

Our findings highlight a novel role for eHsp90 α as a potent initiator of stromal inflammatory responses, exemplified by its robust activation of the master regulators NF- κ B and STAT3. Given that eHsp90 α expression was nominal in prostate stromal cells (data not shown), our data strongly support a paracrine role for tumor eHsp90 α in conferring a reactive stromal phenotype. We have recently demonstrated that more aggressive PCa cells secrete elevated levels of eHsp90 α , (20) indicating that PCa cells are a likely source of eHsp90 α . This premise is further supported by the stromal inflammatory responses elicited by CM derived from eHsp90 α -expressing PCa cells. This tumor-stromal co-culture model demonstrates that within a physiological context, modest increases in eHsp90 α secretion are capable of dramatically altering the prostate stroma. Thus, the elevated eHsp90 α expression observed in more aggressive PCa cells may portend their ability to more effectively initiate a profoundly reactive stroma. In summary, our findings indicate that eHsp90 is poised as a critical effector of tumor-stromal crosstalk to synergistically fuel cancer progression. Delineation of the mechanisms by which eHsp90 modifies the cancer niche to create a permissive and supportive microenvironment for tumor progression will be required to

establish the potential therapeutic value of targeting eHsp90 α as a means of preventing PCa progression.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

Acknowledgments

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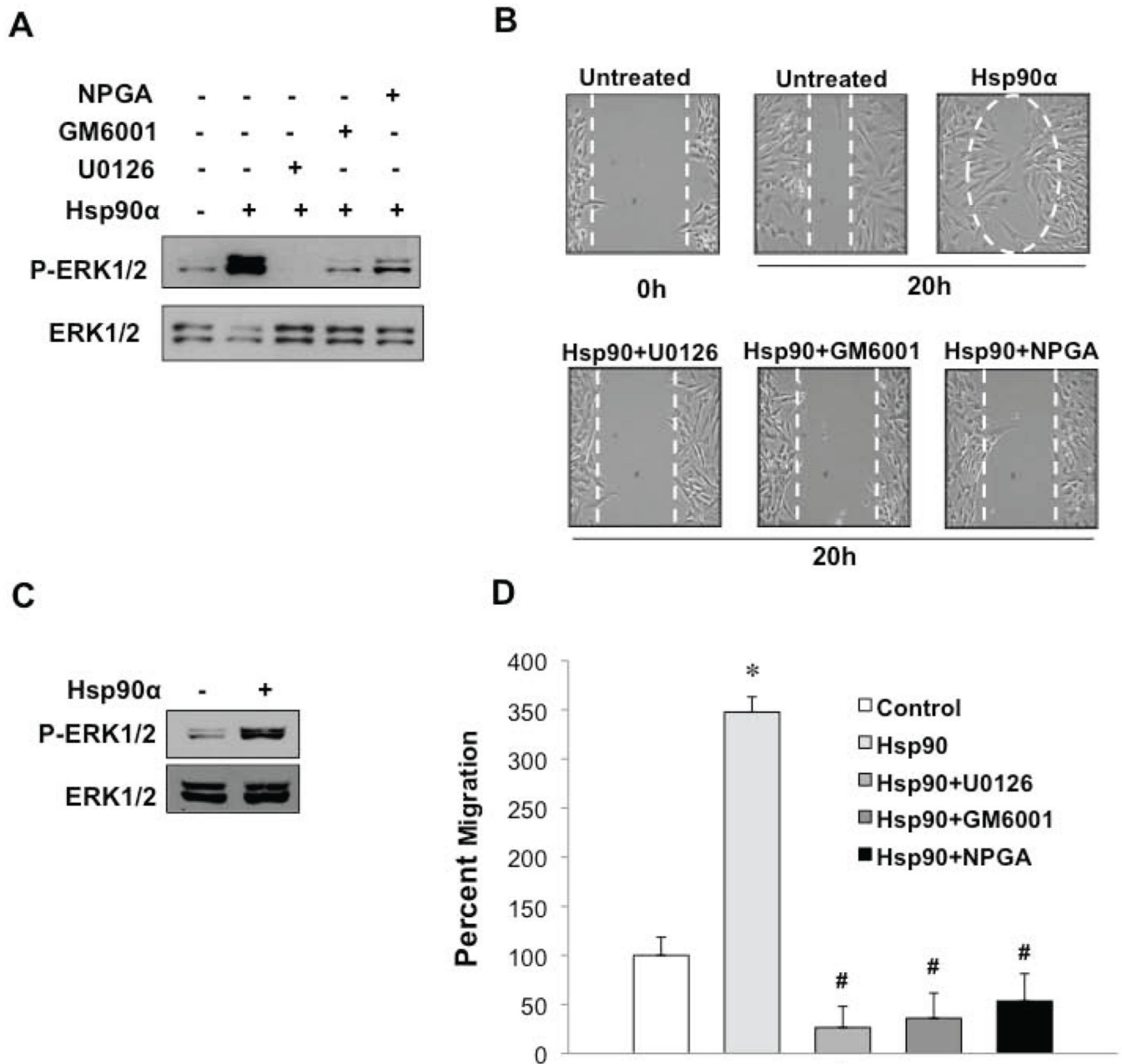


Figure 1. eHsp90 promotes ERK-dependent prostate stromal fibroblast cell motility

A) Prostate stromal cells (PrSCs, Clonetics) were serum starved (0.1% FBS) overnight, treated with the indicated inhibitors for 6 hr, and exposed to Hsp90 α protein (3 ug/ml) for 10 min. Resultant cell lysates were analyzed for phosphorylated and total ERK1/2 by SDS-PAGE and Western blot. **B)** Scratch wound assays of serum starved PrSCs. PrSCs were pre-treated for 4 hours in the absence or presence of NPGA (1 μ M), U0261 (10 μ M), or GM6001 (1 μ M), followed by stimulation with Hsp90 α protein. Cell migration into the wound area was determined at 20 hours. Cell motility was quantified using ImageJ and significance (*) determined by ANOVA and Student's *t*-test ($p < 0.05$). **C)** Immortalized NPFs were treated as in A and lysates analyzed for activated and total ERK1/2. **D)** NPFs were treated as in B

and eHsp90 α -mediated cell motility similarly determined. (#) denotes $p < 0.05$ with respect to inhibitory effects of treatments relative to Hsp90 treated cells.

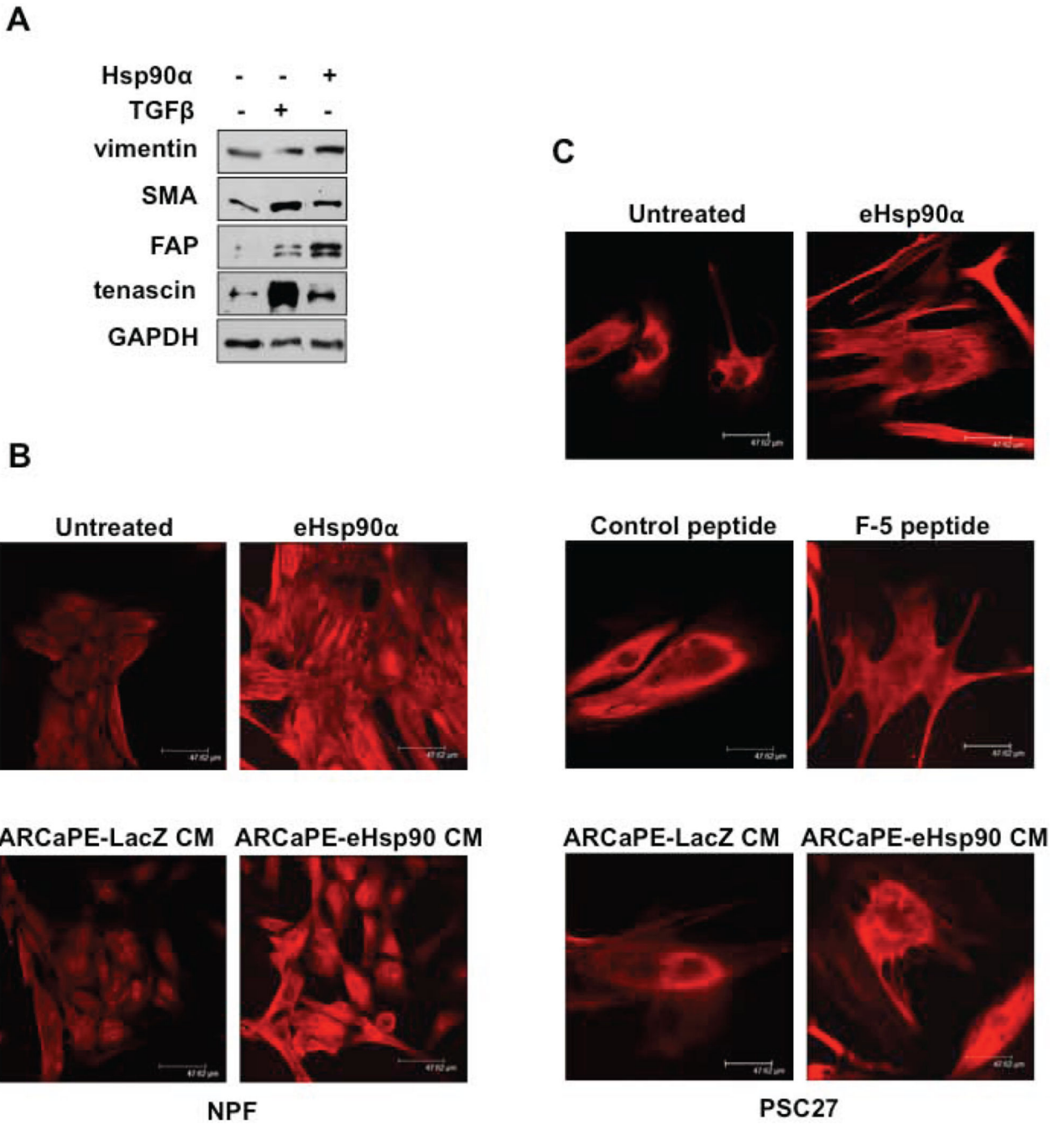


Figure 2. eHsp90α initiates molecular changes associated with a reactive stromal phenotype

A) Serum starved (0.1%) NPFs were treated with TGFβ (5 ng/mL) or Hsp90α (3 μg/mL) for 16 hr. Whole cell lysates were collected and analyzed for vimentin, SMA, FAP, and tenascin. **C. B)** NPF cells were serum starved overnight as in A and subsequently incubated (3 days) with either Hsp90α protein, or with conditioned media (CM) from transduced ARCaPE-LacZ or ARCaPE-eHsp90 cells. **C)** PSC27 cells were treated as in B. Comparative effects of Hsp90α F-5 peptide are shown. The control peptide is derived from the middle domain of Hsp90α.

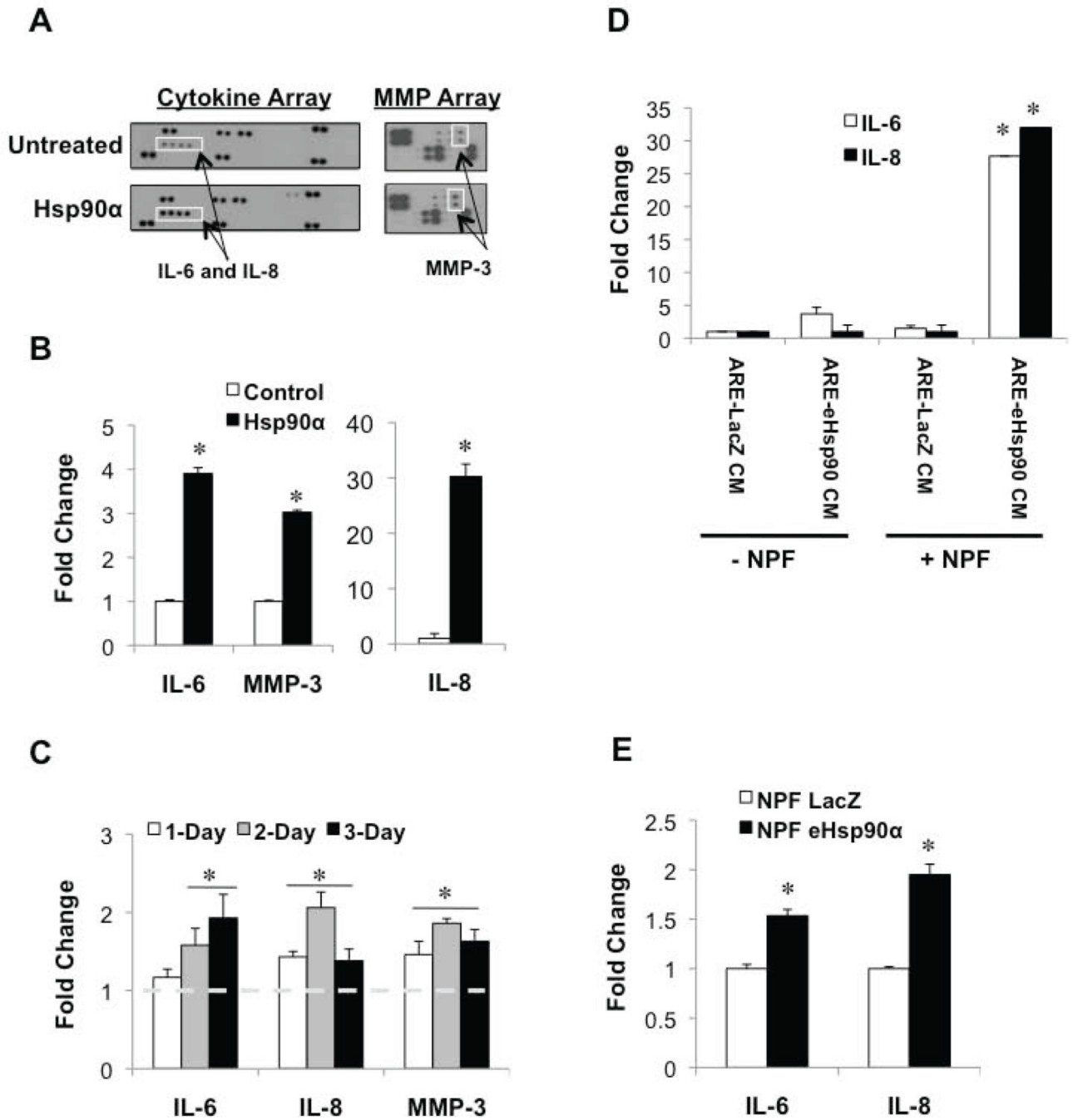


Figure 3. eHsp90α initiates a pro-inflammatory phenotype in stromal prostate fibroblasts
A) Serum starved (0.1%) NPFs were treated with Hsp90α protein overnight. CM was collected and analyzed via a cytokine and MMP-based array. **B)** ELISA confirmation of the Hsp90α-mediated secretion of IL-6, IL-8, and MMP3 in NPFs. **C)** Similar analysis of Hsp90α-mediated secretion of IL-6, IL-8, and MMP3 in PSC27 following the indicated treatment times. **D)** Analysis of IL-6 and IL-8 secretion in ARCaPE-LacZ and ARCaPE-eHsp90 in CM, and effects of addition of this CM (1:1) to NPFs (16 hr). Secreted protein values were normalized and are presented as fold change over untreated control. Hsp90α

protein significantly induced IL-6, IL-8, and MMP3 secretion (*) as determined by ANOVA and Student's *t*-test ($p < 0.05$). E) RNA was isolated from cells in D and the indicated targets analyzed via qPCR in triplicate.

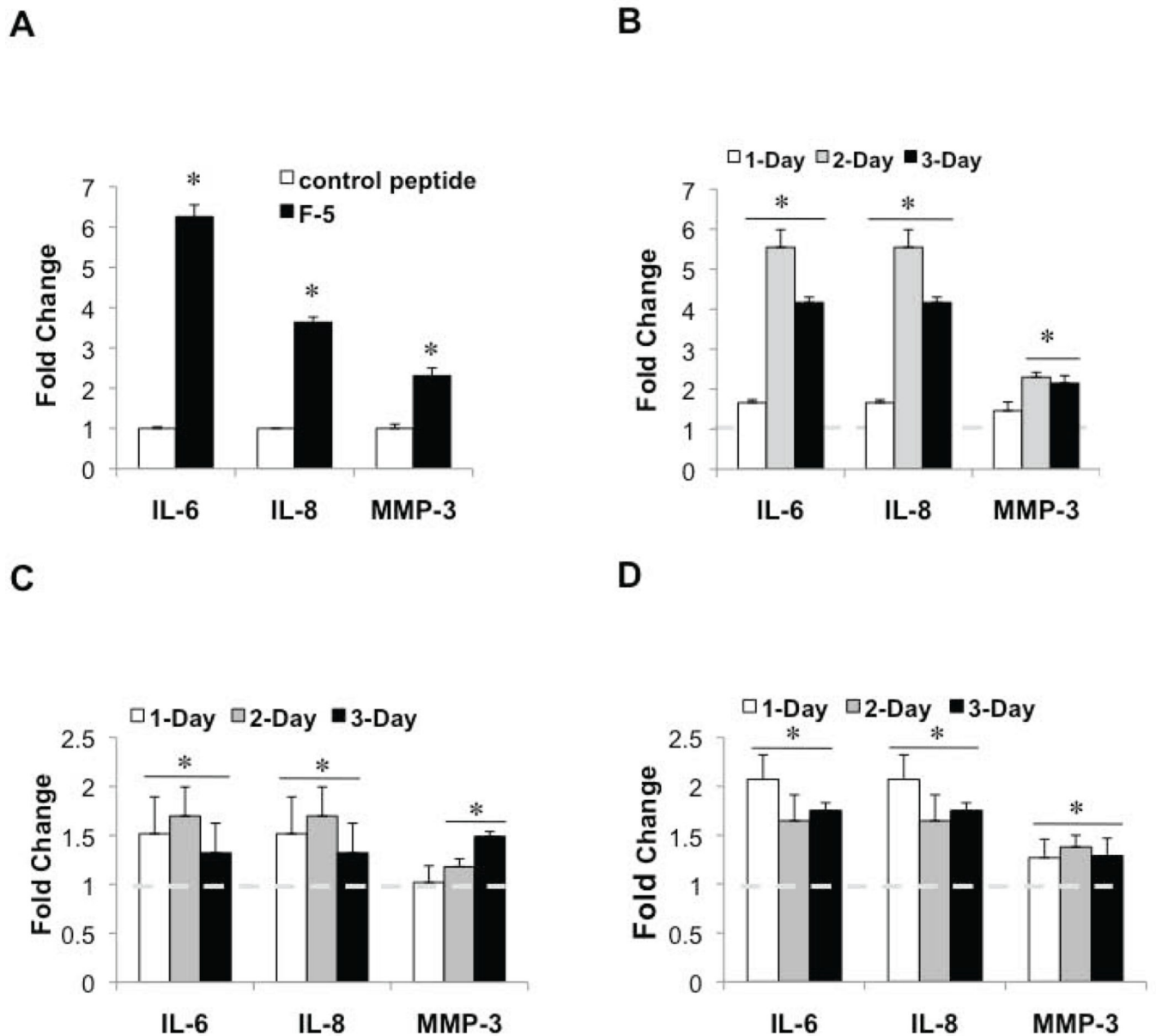


Figure 4. eHsp90 α promotes the stromal secretion of inflammatory mediators in a chaperone-independent manner

A) Serum starved NPFs were treated with either control or Hsp90 α F-5 peptide (at a concentration equimolar to 3 μ g/mL Hsp90 α protein) overnight, and CM analyzed as in Fig. 3. **B–D)** Immortalized PSC27 (**B**) or human primary PrSF cells 5905N (**C**) and 6105N (**D**) were treated with F-5 peptide as in A, for the indicated times, and CM similarly analyzed. Statistics were performed as in Fig. 3.

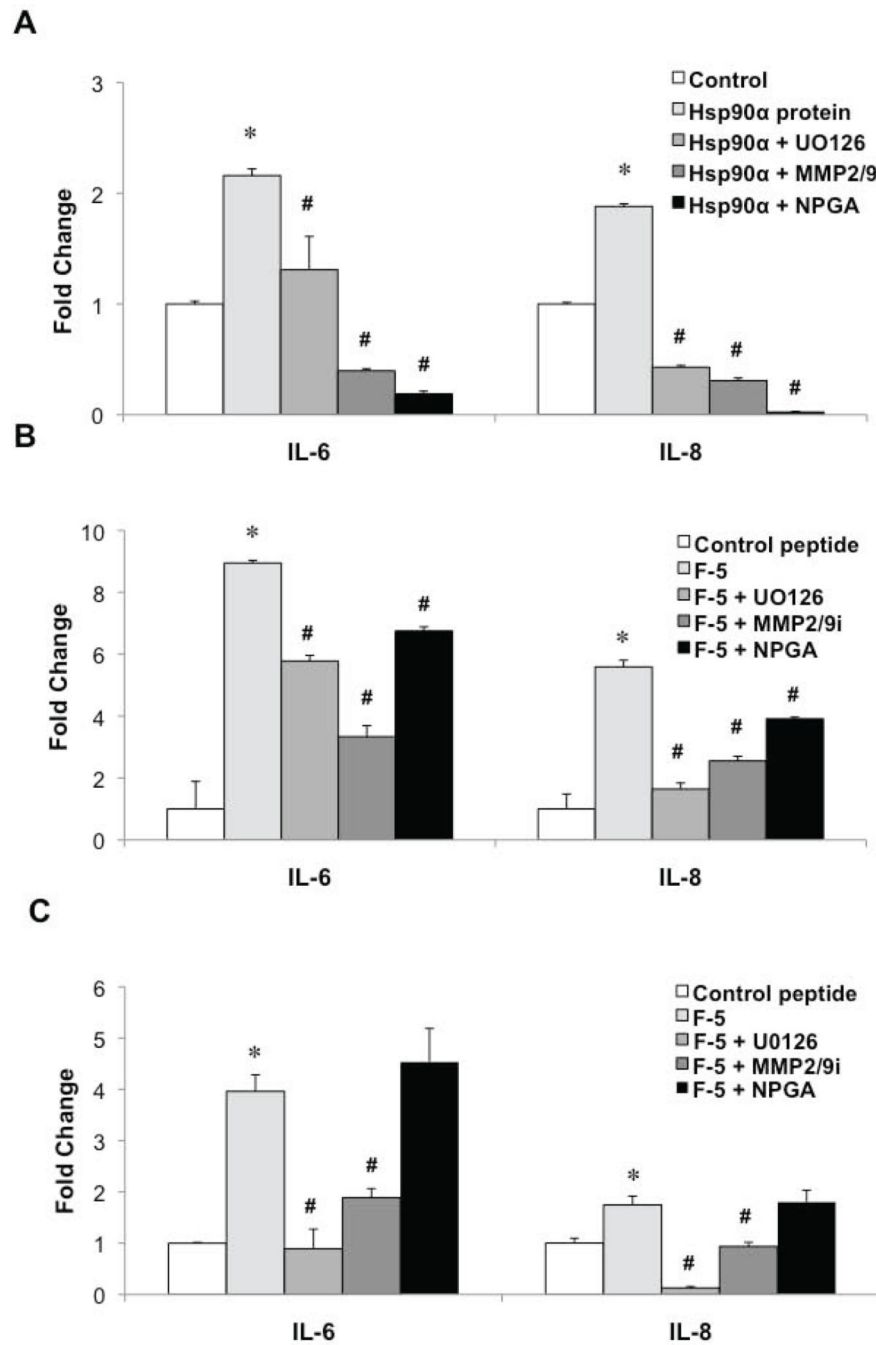


Figure 5. Hsp90 α regulates stromal IL-6 and IL-8 secretion via an ERK1/2 and MMP2/9 dependent pathway

A, B) Serum starved NPFs were pre-treated (6 hr) with either UO126 (10 μ M), MMP-2/9 inhibitor IV (1 μ M) or NPGA (1 μ M) prior to incubation (16 hr) with either Hsp90 α protein (**A**) or F-5 peptide (**B**). ELISA analysis was used to evaluate secreted levels of IL-6 and IL-8. **C)** RNA was harvested from the treated cells and transcript levels corresponding to IL-6 and IL-8 were determined by qRT-PCR. Values were normalized to untreated control and significance (*, #) was determined by ANOVA and Student's *t*-test ($p < 0.05$). (#)

denotes significance of inhibitory effects of treatments relative to Hsp90 protein or F-5 peptide treated cells.

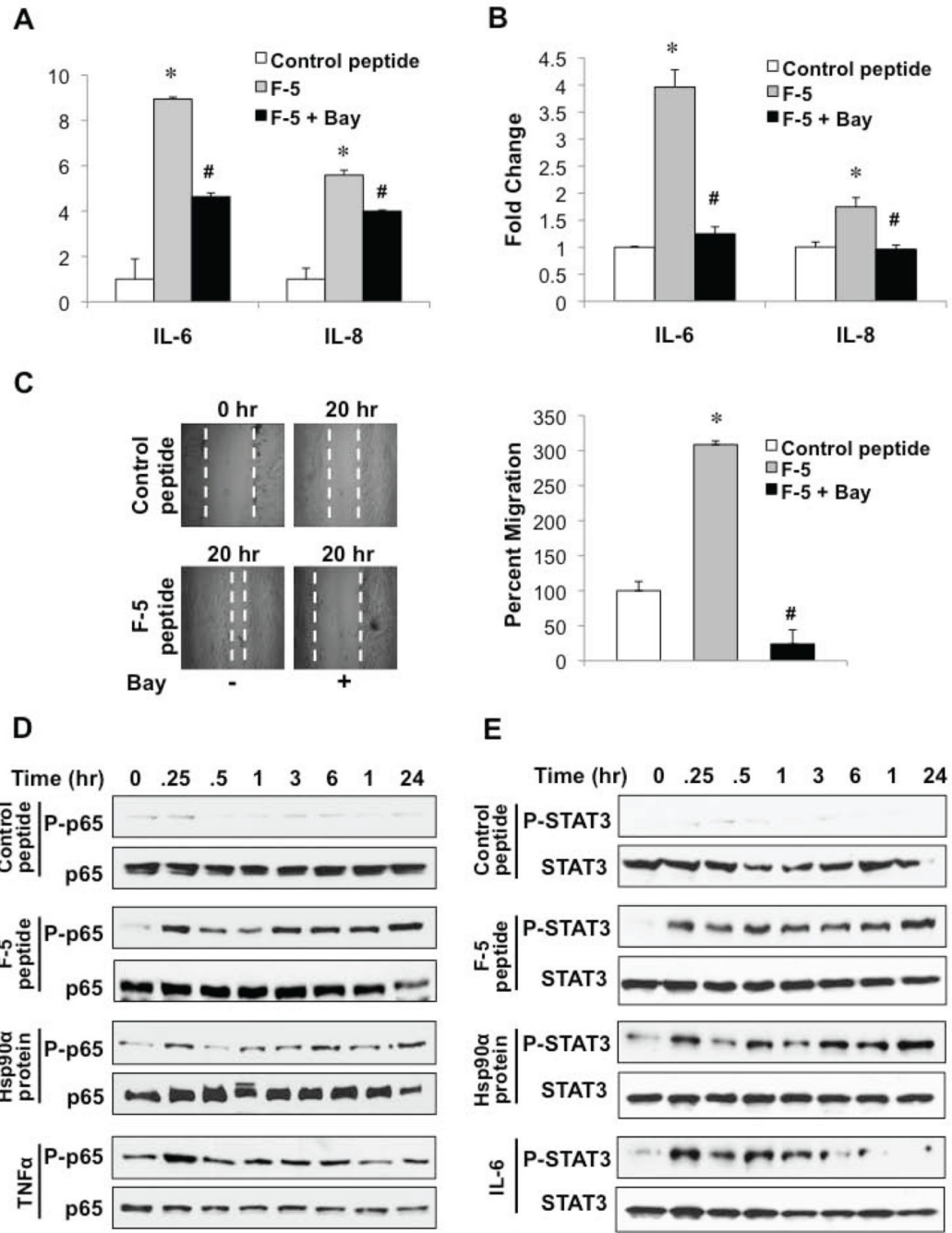


Figure 6. NF-κB signaling participates in eHsp90α-mediated cytokine secretion

A) Serum starved NPF cells were pre-treated with the NF-κB inhibitor Bay11-7082 (2 μM), for 6 h prior to stimulation with control or Hsp90α peptide (16 h). ELISA was used to assess levels of IL-6 and IL-8 in CM. **B)** NPFs were treated as in A and transcript levels determined by qRT-PCR. **C)** NPFs were treated as in A and subjected to a scratch wounding assay. Representative images and quantified values of migration are indicated. As in Fig. 5, (#) denotes significance of inhibitory effects of treatments relative to F-5 peptide treated cells. **D, E)** Serum starved NPFs were stimulated with Hsp90α protein or F5 peptide for the

indicated times. TNF α (20 ng/mL) and IL-6 (20 ng/mL) were used as positive controls for P-p65 and P-STAT3, respectively. Immunoblot analysis was performed on whole cell lysates for phosphorylated and total p65 (**D**) and phosphorylated and total STAT3 (**E**).

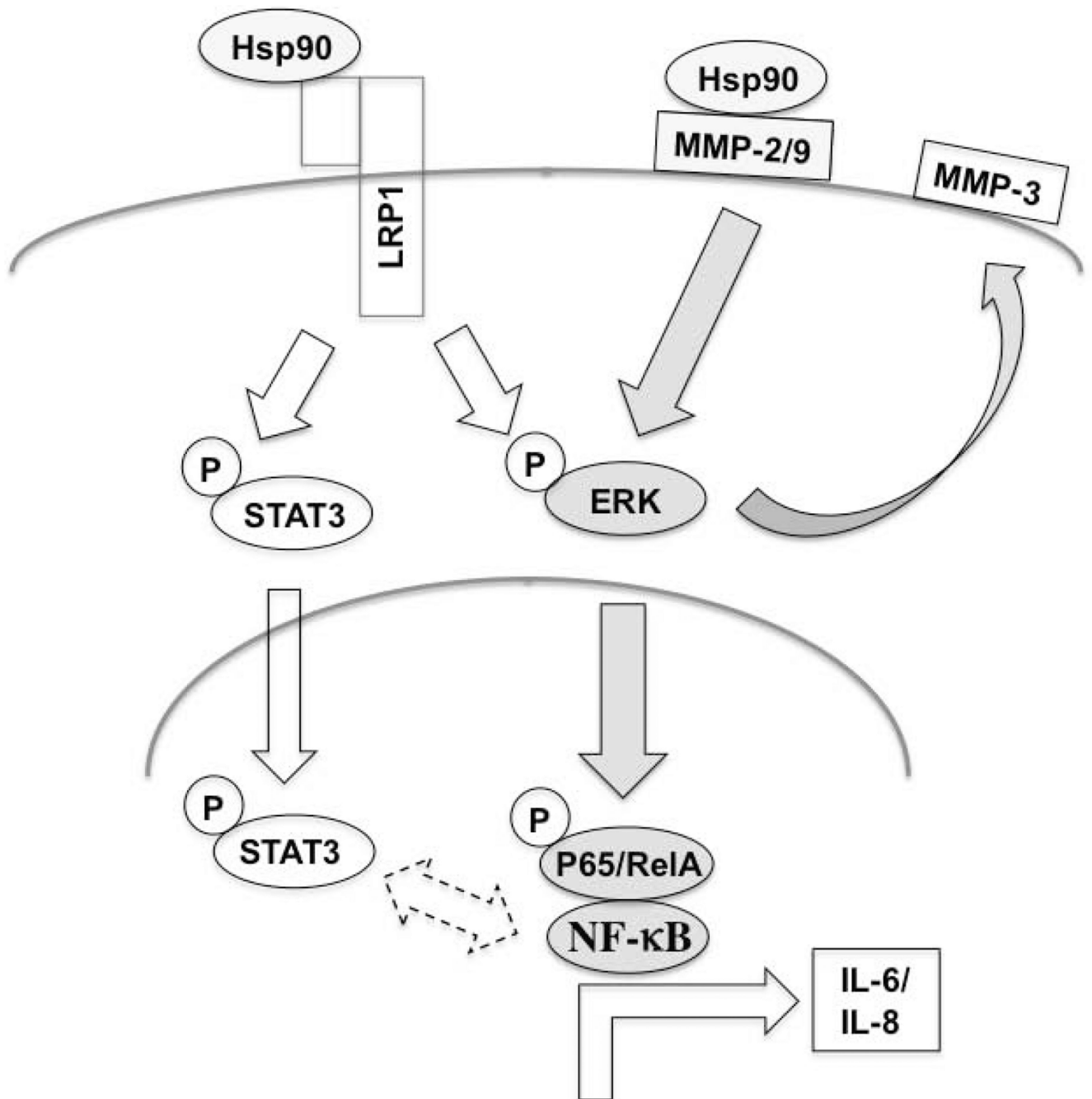


Figure 7. Model for eHsp90 α -mediated induction of an inflammatory stromal milieu
 eHsp90 α initiates ERK signaling, herein demonstrated via MMP-2/9 activation. An eHsp90 α -MMP-2/9-ERK pathway also upregulates MMP-3 transcription. The eHsp90 α -MMP-2/9-ERK axis is required for NF- κ B-dependent transcription of IL-6/IL-8. eHsp90 also stimulates STAT3 activation, albeit in an ERK-independent pathway. STAT-3 and NF- κ B may collaborate to promote tissue inflammation.