



hMENA is a key regulator in endothelin-1/ β -arrestin1-induced invadopodial function and metastatic process

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Edited by Robert J. Lefkowitz, Howard Hughes Medical Institute, Duke University Medical Center, Durham, NC, and approved January 12, 2018 (received for review September 13, 2017)

Aberrant activation of endothelin-1 receptors (ET-1R) elicits pleiotropic effects relevant for tumor progression. The network activated by this receptor might be finely, spatially, and temporarily orchestrated by β -arrestin1 (β -arr1)-driven interactome. Here, we identify hMENA, a member of the actin-regulatory protein ENA/VASP family, as an interacting partner of β -arr1, necessary for invadopodial function downstream of ET-1R in serous ovarian cancer (SOC) progression. ET-1R activation by ET-1 up-regulates expression of hMENA/hMENA Δ v6 isoforms through β -arr1, restricted to mesenchymal-like invasive SOC cells. The interaction of β -arr1 with hMENA/hMENA Δ v6 triggered by ET-1 leads to activation of RhoC and cortactin, recruitment of membrane type 1-matrix metalloproteinase, and invadopodia maturation, thereby enhancing cell plasticity, transendothelial migration, and the resulting spread of invasive cells. The treatment with the ET-1R antagonist macitentan impairs the interaction of β -arr1 with hMENA and inhibits invadopodial maturation and tumor dissemination in SOC orthotopic xenografts. Finally, high ET_AR/hMENA/ β -arr1 gene expression signature is associated with a poor prognosis in SOC patients. These data define a pivotal function of hMENA/hMENA Δ v6 for ET-1/ β -arr1-induced invadopodial activity and ovarian cancer progression.

endothelin receptor | β -arrestin | invadopodia | ovarian cancer | hMENA

High-grade serous ovarian cancer (HG-SOC) accounts for 70–80% of ovarian cancer (OC) deaths and is associated with aggressive metastasis (1). Invasive cancer cells utilize actin-rich membrane protrusions, invadopodia, to focally degrade the extracellular matrix (ECM) and metastasize (2, 3). These specialized structures, identified by the presence of cortactin and TKS5 proteins, serve as sites of physical convergence of adhesive, cytoskeletal, proteolytic, and membrane-trafficking signaling pathways. The invadopodia life cycle includes the formation of precursors, followed by invadopodium stabilization, maturation, and elongation to form active invasive protrusions, with the recruitment of matrix metalloproteinases (MMP), particularly membrane type 1 (MT1)-MMP (2–4). Although environmental cues and cell metabolism act as drivers of invadopodia, inputs from growth factors are now considered key players of the invadopodia life cycle (4). The endothelin-1 (ET-1) receptors (ET-1R) ET_AR and ET_BR, belonging to the G-protein-coupled receptor family, are well-recognized drivers of tumorigenesis and metastatic progression of human cancers, including SOC (5). Overexpression of ET_AR in patients with SOC is associated with a poor prognosis and chemoresistance (5–7). Autocrine activation of ET_AR is critically involved in controlling pleiotropic tumor-promoting activities, including epithelial-to-mesenchymal transition (EMT), invasion, and metastasis (5–7). The diversity of intracellular signaling pathways and cellular processes modulated by ET-1R is driven by the multifunctional protein β -arrestin1 (β -arr1), acting as a molecular hub, orchestrating active signaling complexes required for the engagement of several key pathways and transcriptional regulators, to promote persistent invasive signaling (9–14). We recently reported that ET-1 triggers the signaling to form invadopodia in OC cells (15, 16). However, a

detailed knowledge of how ET-1 could regulate critical effectors of invadopodia in the invasive process remains to be addressed.

Among the genes of the “invasion signature” up-regulated in metastatic tumor cells, the human enabled homolog (ENAH) encodes for the 570-aa actin regulatory hMENA protein (17). hMENA, belonging to the Ena/VASP family, is involved in cell adhesion and migration (18, 19). The hMENA gene undergoes a splicing process generating multiple isoforms that are expressed in specific tissues, and its pattern of expression has been proposed as a marker of tumor progression (20). Among these isoforms in breast (21), lung (22), and pancreatic cancer (23), it has been demonstrated that the two alternatively expressed splicing isoforms of hMENA—hMENA^{11a}, an epithelial-associated splice variant with an additional exon (exon 11a) (24), and hMENA Δ v6, a mesenchymal-associated isoform lacking exon 6 (21)—have opposite roles in cell invasion and migration. hMENA Δ v6 is expressed only in the invasive breast, pancreatic, and lung cell lines (21–23). A different splice variant isoform of hMENA, MENAINV, with an additional exon after the EVH1 domain, has been described as a regulator of invadopodial function and cell invasiveness in response to EGF signaling in breast cancer cells (25).

Significance

Discovering new targets and novel determinants of metastatic spread is an unmet need in ovarian cancer, which is plagued by high rates of recurrence. Endothelin-1 receptors (ET-1R), belonging to the G-protein-coupled receptor family, represent important targets critically involved in malignant progression. Here we identify a mechanistic link between ET-1R and the actin regulatory protein hMENA/hMENA Δ v6 through the specific interaction with the multifunctional protein β -arrestin1 (β -arr1), which initiates signaling cascades as part of the molecular complex crucial for invadopodial maturation and malignant dissemination. Targeting ET-1R by using macitentan, a Food and Drug Administration-approved antipulmonary arterial hypertension drug, can impair the β -arr1-mediated signaling network controlling ovarian cancer progression and therefore represents a therapeutic option for ovarian cancer patients.

Author contributions: L.R. designed research; V.C., L.C., P.T., and F.S. performed research; F.D.M. contributed new reagents/analytic tools; F.D.M., F.S., G.F., A.S., G.B., P.N., A.B., and L.R. analyzed data; and A.B. and L.R. wrote the paper.

The authors declare no conflict of interest.

This article is a PNAS Direct Submission.

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This article contains supporting information online at www.pnas.org/lookup/suppl/doi:10.1073/pnas.1715998115/-DCSupplemental.

Published online February 8, 2018.

Here, to define how SOC cells acquire the plasticity for invadopodial formation and maturation driven by the ET-1 axis, and considering the role of hMENA in invadopodia, we delineate an important role for hMENA/hMENA Δ v6 as a critical regulator of ET-1/ β -arr1-induced cytoskeleton architecture changes and invasive protrusion function in malignant progression.

Results

Expression of hMENA in Human Ovarian Cancer. To assess the pathophysiological functions of hMENA (ENAH) in OC, two published mRNA expression profiles, obtained from the National Center for Biotechnology Information (<https://www.ncbi.nlm.nih.gov/geo/>), have been evaluated (26). ENAH mRNA is significantly up-regulated in OC tissues and, in particular, in HG-SOC tumors, compared with normal tissues (Fig. 1 *A* and *B*). Consistently, analysis of a clinically annotated patient cohort of HG-SOC reveals that all 24 patients express ENAH mRNA at different levels (Fig. 1 *C*), supporting the relevance of hMENA in this tumor type. We analyzed the expression of total hMENA and the specific hMENA^{11a} (antiinvasive) and hMENA Δ v6 (proinvasive) isoforms (21) in a panel of SOC cell lines, exhibiting a different ability to invade in the transwell Matrigel assay (Fig. S1 *A* and *B*). All SOC cells express hMENA, and only the highly invasive cell lines, HEY and SKOV3, also express hMENA Δ v6 associated with EMT-related markers, such as N-cadherin and vimentin, consistent with the association of hMENA Δ v6 expression with the EMT phenotype. On the contrary, the low invasive cell line CAOV3 expresses hMENA^{11a} along with the epithelial marker E-cadherin (Fig. 1 *D* and *E*). Immunofluorescent analysis indicates that hMENA is localized to the periphery of SOC cells in the tips of actin stress fiber-like structures (Fig. S1 *C*).

ET-1 Is a Regulator of hMENA Expression in SOC Cells. Given that ET-1 is a druggable pathway in OC (7), we explored whether ET-1 signaling is involved in the regulation of hMENA expression and function. In invasive HEY cells, all ENAH transcripts increase upon ET-1 treatment (Fig. 2 *A*). In HEY and SKOV3 cells, hMENA Δ v6 and hMENA protein expression enhances upon ET-1 addition in a time-dependent manner with a maximum increase after 48 h (Fig. 2 *B* and Fig. S1 *D*). On the contrary, in the epithelium-like CAOV3 cells, the ET-1 addition leads to a decreased expression of

hMENA^{11a} after 72 h (Fig. S1 *E*). These findings strengthen the idea that ET-1 regulates the balance between the mesenchymal-associated hMENA Δ v6 and the epithelial-associated hMENA^{11a} isoforms in determining SOC invasive behavior. The treatment with macitentan, a potent dual ET_AR/ET_BR antagonist, inhibits ET-1-induced hMENA/hMENA Δ v6 expression (Fig. 2 *C*). Moreover, considering that SOC cells express both β -arr1 and -2 (8), we evaluated the involvement of β -arr on ET-1-mediated hMENA isoform expression at the mRNA and protein levels. Silencing of β -arr1, but not of β -arr2, inhibits ET-1-induced hMENA/hMENA Δ v6 expression (Fig. 2 *A* and *D* and Fig. S4 *A–C*), indicating that ET-1 through β -arr1 up-regulates hMENA/hMENA Δ v6 expression at the transcriptional level. In untreated cells, hMENA is distributed to the cell periphery where it appears to be organized with F-actin at the tips of actin filaments (Fig. 2 *E* and Fig. S2 *A* and *B*), as shown by confocal scanning laser microscopy (CSLM) analysis. Upon ET-1 addition, hMENA relocates from the periphery to the cytosol in a time-dependent manner and, between 30 and 60 min, colocalizes in F-actin/cortactin-enriched dot-like structures in the ventral region of the cells, while treatment with macitentan, as well as silencing of β -arr1, abrogates the ET-1-induced relocation of hMENA (Fig. 2 *E* and Fig. S2). These data indicate that ET-1/ β -arr1 signaling up-regulates hMENA and hMENA Δ v6, as well as their recruitment to regions of the F-actin/cortactin-rich dots.

hMENA Is an Interactor with β -arr1 upon ET-1R Activation. Given the ability of β -arr1 to directly interact with cytoskeleton remodeling players, and to act as a mediator of ET_AR signaling (15, 27, 28), we reasoned that ET-1-dependent hMENA localization could arise from its association with β -arr1. Endogenous β -arr1 and hMENA readily coimmunoprecipitate (co-IP) in ET-1-stimulated HEY cells in a time-dependent manner (Fig. 2 *F* and Fig. S3 *A*). Macitentan treatment significantly impairs the interaction between β -arr1 and endogenous hMENA or exogenously expressed GFP-tagged hMENA Δ v6 induced by ET-1 (Fig. 2 *G* and *H*). In agreement with these findings, CSLM analysis shows that hMENA colocalizes with β -arr1 in ET-1-stimulated cells, but not upon treatment with macitentan (Fig. S3 *B* and *C*). Collectively, these results reveal that hMENA/hMENA Δ v6 is a β -arr1 interactor in ET_AR-dependent signaling.

Having demonstrated that the signal-integrating activity of β -arr1 on Rho GTPase is accomplished by its interaction with postsynaptic density protein 95/Disk-Large/Zonula occludens (PDZ)-RhoGEF, a member of the RhoGEF family (15, 16), we tested whether hMENA links the β -arr1/PDZ-RhoGEF complex supporting the ET-1 signaling. ET-1 induces the formation of a trimeric complex with PDZ-RhoGEF, β -arr1, and hMENA, as shown by co-IP analysis (Fig. 3 *A*). The interaction of β -arr1 with PDZ-RhoGEF or PDZ-RhoGEF with hMENA is not detectable when the cells are silenced for hMENA or β -arr1 (Fig. 3 *B* and *C* and Fig. S4 *A, D, and E*), indicating that ET-1R activation might provide β -arr1 for physical recruitment of PDZ-RhoGEF and hMENA, relocating hMENA to invadopodia, as evaluated by colocalization of hMENA/F-actin/cortactin (Fig. 2 *E* and Fig. S2).

hMENA Mediates ET-1R/ β -arr1-Induced RhoC Activity. Among Rho GTPase family members, RhoC activity is spatially confined in the area surrounding the actin core of invadopodia controlling restricted cofilin signaling (29). ET-1-induced RhoC activity significantly decreases in si-hMENA or sh-PDZ-RhoGEF or si- β -arr1-transfected cells (Fig. 3 *D* and Fig. S4 *F*), indicating that hMENA/hMENA Δ v6, within the protein complex with β -arr1 and PDZ-RhoGEF, influences the activity of RhoC. Of note, the overexpression of hMENA Δ v6 promotes activation of RhoC, which is further potentiated by the addition of ET-1 (Fig. S4 *G*), demonstrating that the formation of a β -arr1/PDZ-RhoGEF/hMENA Δ v6 molecular complex elicits ET-1 responses by increasing RhoC signaling. Altogether these findings untangle the interconnected roles of RhoC GTPases, a specific GEF, and hMENA activated by ET-1R/ β -arr1 for the formation of invadopodia.

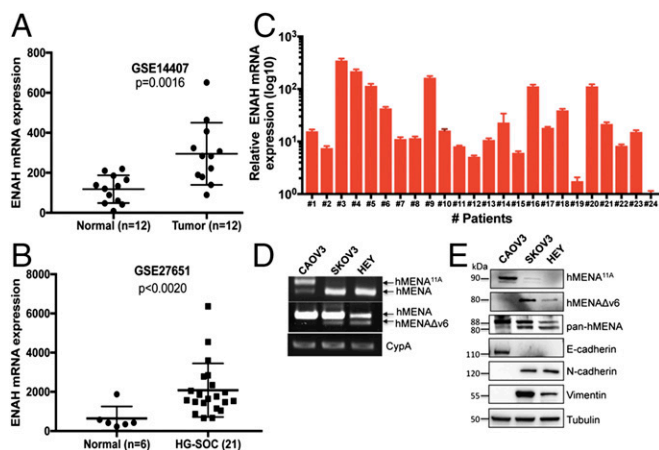


Fig. 1. HG-SOC expresses hMENA along with the hMENA Δ v6 isoform. (*A* and *B*) Expression profiling of ENAH mRNAs in human normal and tumor specimens from two Gene Expression Omnibus profile datasets. (*C*) Relative ENAH mRNA expression in 24 HG-SOC specimens normalized for GAPDH expression. Error bars: mean \pm SD ($n = 2$). (*D*) Representative RT-PCR analysis for hMENA, hMENA^{11a}, and hMENA Δ v6 mRNA in three different SOC cell lines. (*E*) Representative Western blotting (WB) analysis of hMENA (pan-hMENA), hMENA^{11a}, hMENA Δ v6, E-cadherin, N-cadherin, and vimentin in SOC cell lines. Tubulin was used as a loading control.

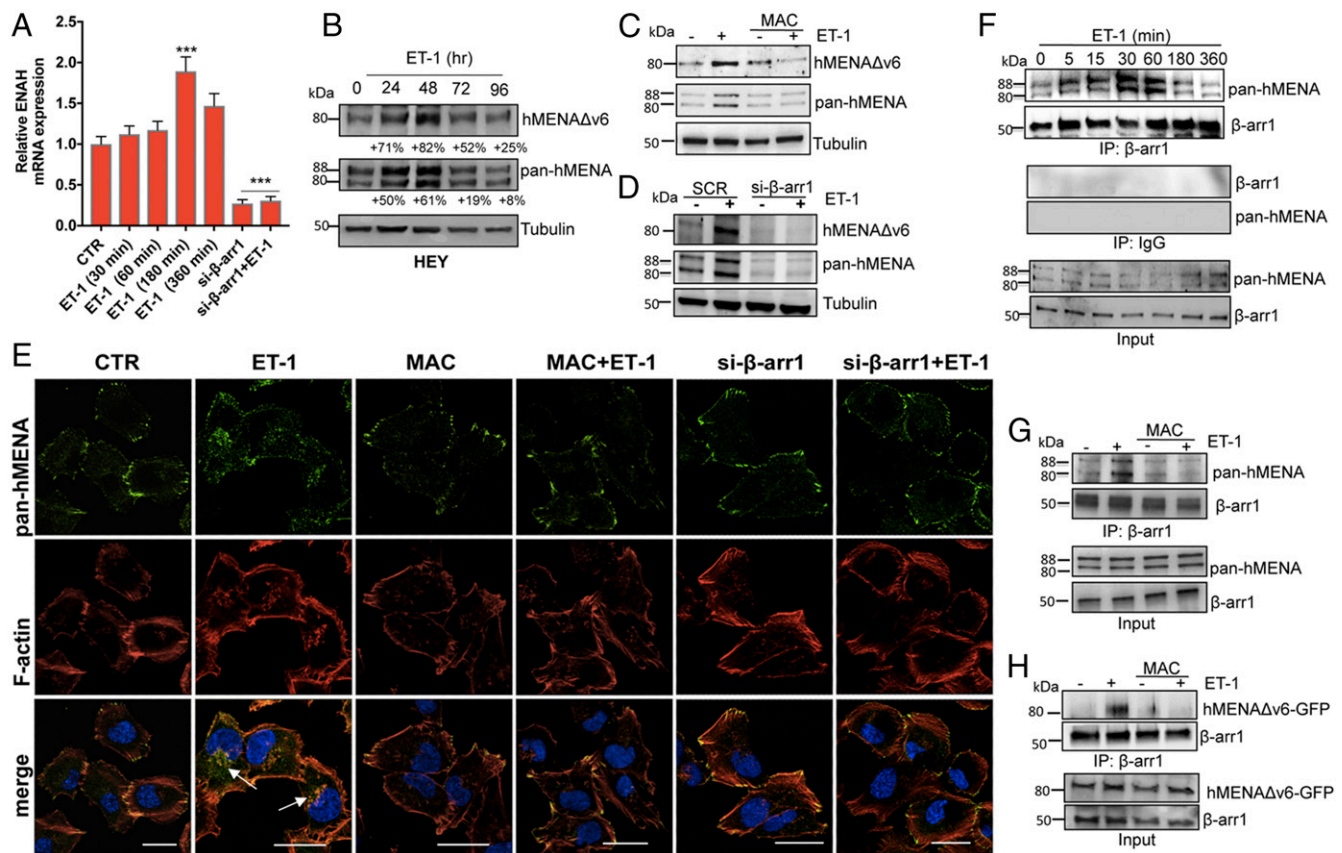


Fig. 2. ET-1R up-regulates the expression of hMENA and hMENA Δ V6 through β -arr1 and promotes the association of hMENA with β -arr1. (A) Expression of hMENA mRNA measured by qRT-PCR in HEY cells stimulated with ET-1 (100 nM) for the indicated times or silenced for β -arr1 and treated with ET-1 for 180 min. $***P \leq 0.001$ vs. CTR or ET-1-stimulated cells. Error bars: mean \pm SD. (B) WB analysis of hMENA Δ v6 and hMENA (pan-hMENA) expression in cells stimulated or not with ET-1 for different times or (C) treated with macitentan (MAC, 1 μ M) alone or in combination with ET-1 for 48 h. For B, percentage of band density compared with unstimulated cells (Lower). (D) WB analysis of hMENA Δ v6 and hMENA expression in HEY cells transfected with scramble (SCR)- or β -arr1-siRNA and stimulated with ET-1 for 48 h. (E) Representative immunofluorescent images of HEY cells, transfected with scramble (CTR)- or β -arr1-siRNA, stimulated with ET-1 (60 min) and MAC, and stained with phalloidin (to detect F-actin; red), pan-hMENA Ab (green), and DAPI (blue). hMENA/F-actin colocalization: white arrows. (Scale bar: 30 μ m.) (F) Protein lysates from HEY cells stimulated with ET-1 for the indicated times were immunoprecipitated with control IgG or anti- β -arr1 Ab. (G) Protein lysates from HEY cells stimulated with ET-1 and/or MAC for 60 min were immunoprecipitated with anti- β -arr1 Ab. (H) Protein lysates from HEY cells, transfected with hMENA Δ v6-GFP and unstimulated or treated with ET-1 and/or MAC for 60 min were immunoprecipitated with an anti- β -arr1 Ab. Input and immunoprecipitation were analyzed by WB for the indicated proteins. For B–H, representative blots are shown.

hMENA Is Recruited to Mature Invadopodia in ET-1/ β -arr1 Signaling.

To evaluate whether the recruitment of hMENA/hMENA Δ v6 is crucial in ET-1-induced invadopodia and ECM degradation, we plated SOC cells into fluorescently labeled gelatin. Cells stimulated with ET-1 readily degrade the gelatin substrate, and hMENA localizes to a proteolytically degraded area (Fig. S5). Moreover, hMENA colocalizes with cortactin within the degraded area in ET-1-stimulated cells, but not in cells treated with macitentan or silenced for β -arr1 (Fig. 4A), confirming the ET-1R/ β -arr1-mediated relocalization of hMENA within active, matrix-degrading invadopodial protrusions. Furthermore, ET-1-treated cells show colocalization of invadopodia markers, as cortactin, MT1-MMP, or TKS5 in the matrix degradation area, which is inhibited by macitentan or by hMENA or β -arr1 silencing (Fig. 4B and Fig. S6), confirming the role of β -arr1/hMENA in ET-1R-promoted maturation of invadopodia, which recruit MT1-MMP to fully degrade ECM.

Cortactin contains two tyrosine phosphorylation sites (Y421, Y466) to regulate invadopodium formation and maturation, triggered also by involvement of Src (30, 31). ET-1 stimulation produces an increase in cortactin phosphorylation (Y421), which is inhibited by treatment with macitentan or with the Src inhibitor PP2 (Fig. 4C and D). Moreover, a significant inhibition of ET-1-induced, Src-mediated cortactin phosphorylation is evident

upon silencing of hMENA (Fig. 4C), indicating that hMENA/hMENA Δ v6 is involved in tyrosine phosphorylation of cortactin to regulate invadopodium maturation downstream from ET-1R/ β -arr1 signaling.

hMENA Is Required for ET-1R/ β -arr1-Dependent Protease Secretion, Cell Plasticity, Transendothelial Migration, and Invasion. The temporal and spatial control of MMP activity is one of the important functions of invadopodia. Inhibition of ET-1-induced MMP-2/9 activation and cell invasion is detected upon macitentan treatment, as well as hMENA or β -arr1 silencing, implying a role for ET-1R-dependent β -arr1/hMENA interaction in enhanced proteolytic activity (Fig. 5A and B) and cell invasion (Fig. S7A).

Previous studies described vasculogenic mimicry (VM) as the functional plasticity of aggressive cancer cells that form vascular networks by regulating dynamic formation of actin-based structures (32). SOC cells form connective structures resembling a network-like appearance with long polygonal shape and cellular protrusions after addition of ET-1, as quantified with the number of nodes, junctions, and branches, which are markedly impaired by macitentan as well as by silencing of β -arr1 (Fig. 5C and Fig. S7B). Interestingly, the down-regulation of hMENA significantly disrupts the formation of VM channels, as evidenced by most tubes remaining open and an increase in dissociative cells (Fig. 5C

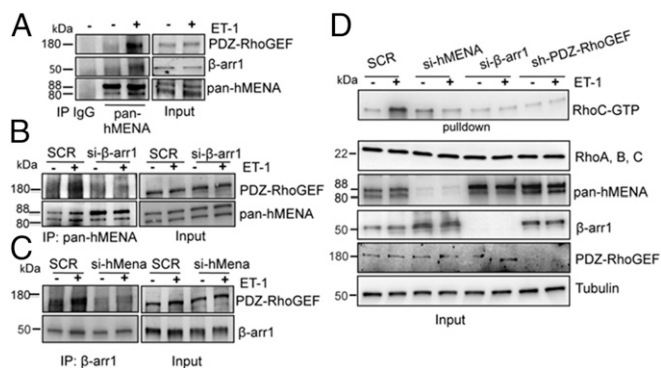


Fig. 3. ET-1 induces the association of hMENA with β -arr1 and PDZ-RhoGEF to regulate RhoC activation. (A) Protein lysates from HEY cells stimulated with ET-1 for 60 min were immunoprecipitated with control IgG or pan-hMENA Ab. (B) SCR- and β -arr1-siRNA-transfected HEY cells were stimulated with ET-1 for 60 min. Protein lysates were immunoprecipitated with pan-hMENA Ab. (C) SCR- or hMENA-siRNA transfected HEY cells were stimulated with ET-1 for 60 min. Protein lysates were immunoprecipitated with β -arr1 Ab. For A–C, input and immunoprecipitation were analyzed by WB for the indicated proteins, and representative blots are shown. (D) Rhotekin beads were used to pull down RhoC-GTP from SCR-, hMENA-, β -arr1-siRNA-, or PDZ-RhoGEF-shRNA-transfected HEY cells induced with ET-1 for 5 min. GTP pull-down and input were analyzed by WB for the indicated proteins and representative blots are shown.

and Fig. S7B). Moreover, hMENA Δ v6 overexpression mimics the addition of ET-1, indicating that hMENA/hMENA Δ v6 play an important role in VM induced by ET-1R/ β -arr1 activation, involved in ECM remodeling and protrusion formation.

To determine how SOC cells extravasate from the blood vessels upon ET-1 stimulus, we evaluated the ET-1-stimulated trans-endothelial migration capacity of cells, through a monolayer of human umbilical endothelial cells. This capacity is significantly enhanced compared with control cells, and is strongly reduced by macitentan or silencing of hMENA or β -arr1 (Fig. 5D and Fig. S7C). Together, these results imply that the integration with hMENA/hMENA Δ v6 is critical in ET-1R/ β -arr1-promoted cell plasticity and extravasation through the matrix degradative machinery.

Macitentan Impairs Malignant Progression in Vivo by Interfering with the ET-1R/ β -arr1/hMENA Pathway. Next, we analyzed the effects of macitentan on SOC invasive behavior and molecular effectors of invadopodia in i.p. orthotopic OC xenografts. All mice in the control group show the spread to the surface of the peritoneum, intestines, omentum, small bowel, mesentery, liver, and spleen and ovaries (Fig. 6A). In mice treated with macitentan, the average of i.p. nodules is significantly reduced, associated with a well-tolerated toxicity profile (no weight loss in the treated mice; Fig. 6A). Consistent with results obtained in vitro in i.p. nodules from macitentan-treated mice, a significant inhibition of hMENA/hMENA Δ v6 expression at mRNA and protein levels and reduction of phosphorylation of cortactin are observed (Fig. 6B and C). These results indicate that the therapeutic efficacy of macitentan to control ET-1R/ β -arr1-driven OC progression relies also on its ability to reduce critical invadopodial components, such as hMENA/hMENA Δ v6 expression and cortactin activity.

Combined Expression of ET_AR, β -arr1, and hMENA as a Prognostic Gene Signature in SOC Patients. To determine the clinical relevance of our experimental data, we assessed the prognostic effect of EDNRA, ARBB1, and ENAH mRNA expression (Affymetrix

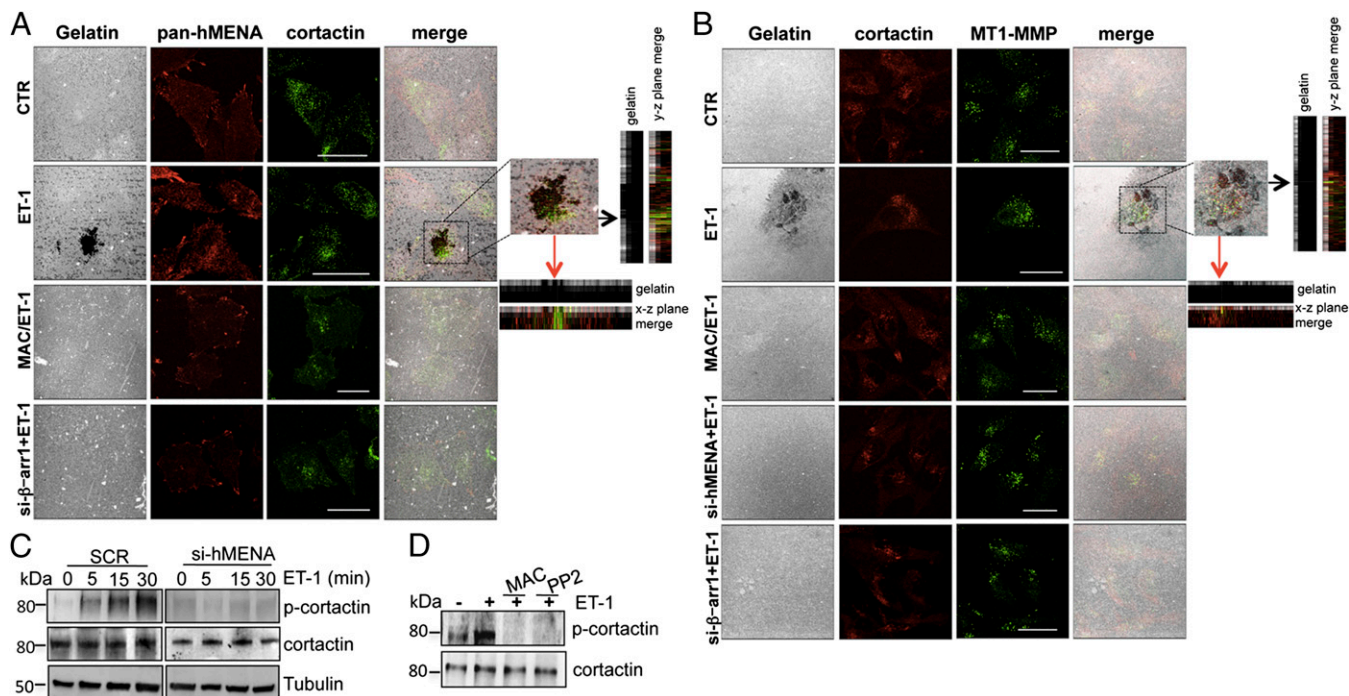


Fig. 4. hMENA is recruited to mature invadopodia in ET-1/ β -arr1 signaling. (A) CSLM analysis of CTR- or β -arr1-siRNA-transfected HEY cells, plated onto gelatin, and treated with ET-1 and/or MAC for 24 h. Representative images show gelatin (gray), cortactin (green), and hMENA (red). (Scale bar: 30 μ m.) (Inset) Higher-power magnification images of the degradation area. Orthogonal views (y - z plane; x - z plane) indicate (arrows) areas of gelatin degradation where hMENA and cortactin are colocalized. (B) CSLM analysis of HEY cells, transfected with SCR or hMENA-, or β -arr1-siRNA, were plated onto gelatin and treated with ET-1 and/or MAC for 24 h. Representative image show gelatin (gray), MT1-MMP (green), and cortactin (red). (Scale bar: 30 μ m.) (Inset) Higher-power magnification image of degradation areas. Orthogonal views (y - z plane; x - z plane) indicate (arrows) areas of gelatin degradation where cortactin and MT1-MMP are colocalized. (C) Protein lysates from SCR- or hMENA-siRNA-transfected HEY cells, unstimulated or stimulated with ET-1 for the indicated times, or (D) from HEY cells unstimulated or stimulated with ET-1 and/or MAC and/or PP2 for 30 min, were analyzed by WB by using anti-phospho-cortactin (Y421) and anticortactin Abs. Representative blots are shown.

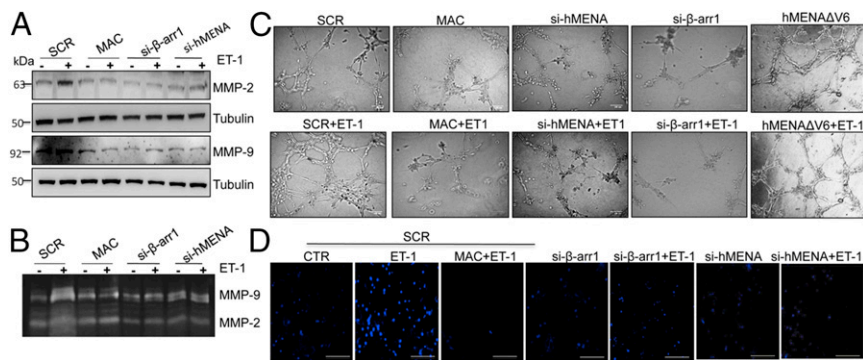


Fig. 5. hMENA is involved in ET-1R/β-arr1-induced protease activation, vasculogenic mimicry, and transendothelial migration. SCR-, hMENA-, or β-arr1-siRNA-transfected HEY cells were stimulated with ET-1 and/or MAC for 24 h. (A) Protein lysates were analyzed by WB for indicated proteins. Representative blots are shown. (B) Culture media of cells treated as above reported were analyzed by gelatin zymography to visualize active forms of MMP-9 and MMP-2. Representative gels are shown. (C) Representative tubule-like structure formation in SCR-, hMENA-, β-arr1-siRNA-transfected cells or hMENAΔv6-GFP-expressing SKOV3 cells, unstimulated or stimulated with ET-1 and/or MAC for 16 h. Original magnification: 20×. (Scale bar: 100 μm.) (D) Fluorescence-labeled HEY cells that had migrated through the endothelial layer were photographed. (Scale bar: 30 μm.)

ID is 204464_at, EDNRA; 218832_at, ARRB1; 222433_at, ENAH) using www.kmplot.com (33). Kaplan–Meier analysis and the log-rank test demonstrate that EDNRA is predictive of overall survival (OS) and progression-free survival (PFS) in SOC patients (Fig. S8). Interestingly, high EDNRA/ARRB1/ENAH gene expression significantly correlates with a worse prognosis for OS [hazard ratio (HR) = 1.44 (1.14–1.82), $P = 0.002$] and PFS [HR = 1.46 (1.19–1.8), $P = 0.0003$], with an enhanced HR (Fig. 6D), compared with the expression of each individual biomarker, or even when they are combined (Fig. S8). Overall, these observations indicate that the network-based EDNRA/ENAH/ARRB1 expression could be used as a prognostic signature and potential druggable pathway in SOC.

Discussion

The identification of molecular drivers of OC progression is critical for the development of therapeutic approaches in advanced-stage disease. Here, we identified hMENA/hMENAΔv6 as a signal transducer of ET-1R/β-arr1 signaling to induce SOC cell plasticity, invadopodia function, intravasation, and malignant dissemination. Several studies have demonstrated that ET-1 signaling enables OC cells to acquire the EMT phenotype, thereby increasing cell motility and invasion, and contributes to poor patient outcome (6–8). However, how the actin-cytoskeleton-mediated signaling network driven by ET-1R can regulate tumor progression is still unknown. Here we have identified hMENA/hMENAΔv6 as a binding partner of β-arr1 and as an interconnected regulator of the signaling platform activated in response to ET-1R in SOC cells.

In the present study, we provide an understanding of how ET-1/β-arr1 signals can confer malignant and invasive traits through the modulation of hMENA and the invasive isoform hMENAΔv6 (Fig. S9). Consistent with the opposite role of hMENA isoforms in tumor cell invasion and EMT (21–23), we demonstrate that the proinvasive hMENA/hMENAΔv6 is highly expressed in the most invasive SOC cells, expressing also EMT markers. These isoforms are up-regulated in highly aggressive and metastatic SOC cells by ET-1 signaling, which, conversely, down-regulates the antiinvasive hMENA^{11a} isoform. This is in line with previous data indicating that TGF-β1 up-regulates hMENAΔv6, but not hMENA^{11a} in pancreatic cancer (23), suggesting that the pattern of hMENA isoform expression mediates the functional effects of different proinvasive factors. In agreement, ET-1R/β-arr1 signaling promotes invadopodia maturation through a sequence of events that require the hMENA/hMENAΔv6/RhoC pathway and the recruitment of cortactin, TKS5, and MT1-MMP to invadopodia for pericellular matrix degradation.

Inputs derived from critical growth factors secreted by tumor cells can create an invasive milieu, acting as drivers of invadopodia in the most aggressive human tumors (3, 4). Although the basic components of invadopodia have been characterized, the complex interactome of invadopodia components and the strongest inputs derived upstream from receptors are poorly understood, suggesting that deciphering the regulators is an essential step to fully understanding invadopodia function and to providing

novel druggable pathways. The data presented here emphasize a unique mechanism by which ET-1R/β-arr1 increases expression of the invadopodial components hMENA/hMENAΔv6 and engages them in discrete regions at invadopodia, promoting their formation and subsequent ECM degradation. Of note, ET-1R/β-arr1 mobilizes hMENA to actin/cortactin dots, sustaining its presence in the ventral protrusions. These findings support the hypothesis that hMENA/hMENAΔv6 might favor the efficacy of the protrusive force to form the degradative protrusions, conferring also a fitness advantage to OC cells able to breach the endothelial barrier and start the transendothelial migration process. The tyrosine phosphorylation of cortactin at invadopodia is necessary for the release of cofilin from cortactin, supporting cofilin-dependent actin polymerization and the maturation of invadopodia into matrix-degrading structures (30, 31). In this regard, we demonstrated that ET-1 regulates cortactin phosphorylation at tyrosine 421 through hMENA/hMENAΔv6 and Src. It has been recently demonstrated that the MenaINV isoform supports invadopodium maturation by increasing cortactin phosphorylation through suppression of the specific phosphatase

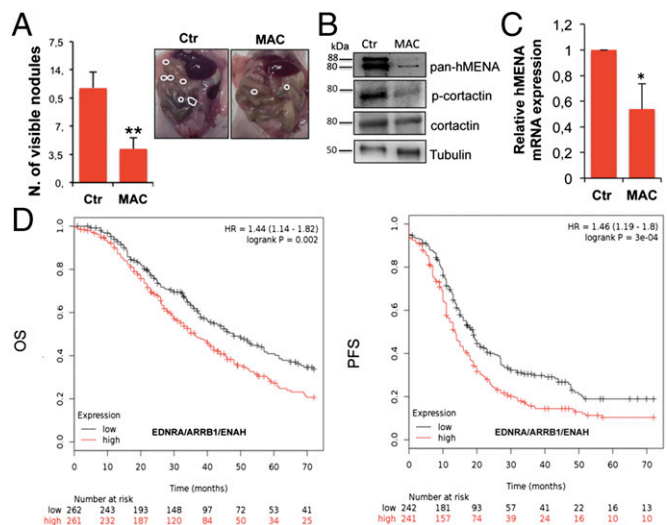


Fig. 6. ET-1R blockade by macitentan inhibits i.p. spreading and hMENA expression. (A) Female nude mice (eight mice for group) were i.p. injected with SKOV3 cells and treated with vehicle [control (Ctr)] or macitentan (MAC, 30 mg/kg oral daily) for 5 wk. $**P \leq 0.01$ ($n = 2$). Error bars: mean \pm SD. Representative i.p. nodules are indicated by white circles (Right). (B) Protein lysate from i.p. nodules of control- or MAC-treated xenografts as in A were analyzed by WB for the indicated proteins. (C) RT-qPCR analysis for the levels of hMENA mRNA from xenografts as in A; $*P \leq 0.05$. Error bars: mean \pm SD. (D) Combined expression of ET_AR, β-arr1, and hMENA as a prognostic signature in SOC patients. Kaplan–Meier curves of OS or PFS in SOC patients with low or high EDNRA/ARRB1/ENAH expression.

PTP1B activity within invadopodia (34, 35). These results likely reflect that different hMENA isoforms concur with nonredundant functional competence in invadopodial maturation. Indeed, our data, uncovering a role for hMENA/hMENA Δ v6 in invadopodial maturation, highlight the role of hMENA isoforms at the crossroads of several growth factor signaling receptors, such as ET-1R, to sensitize tumor cells to form invadopodia and to functionally contribute to the invasive process.

Mounting evidence suggests a direct correlation between the propensity of cancer cells to form invadopodia and a poor prognosis in cancer patients (3, 4). By using a public databases, we found that ENAH is overexpressed in OC patients compared with normal tissues. Clinically, in a large cohort of SOC clinical specimens, the integrated high expression levels of EDNRA/ARRB1/ENAH positively correlate with poor prognosis, validating the clinical implication of this druggable pathway and providing a predictive signature in OC progression.

Targeting ET-1R activity by using the small molecule macitentan, which the Food and Drug Administration has approved for nononcological indications, empowering the β -arr1-driven actin-related networks shuts down the expression of hMENA and impairs malignant progression of SOC xenografts. The therapeutic advantage of macitentan is not only to target OC cells expressing ET_AR by interfering with the mechanism utilized by β -arr1 to regulate hMENA-signaling networks, but also to target tumor-associated environmental elements, such as vascular, lymphatic, and inflammatory cells and fibroblasts, which all express ET_BR (5, 36–38).

Together, this present study highlights a hitherto unappreciated role of the ET-1R/ β -arr1 pathway, which integrates hMENA/hMENA Δ v6 to regulate invadopodial function and OC progression.

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