


Targeting STAT3 by HO3867 induces apoptosis in ovarian clear cell carcinoma

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Advanced ovarian clear cell carcinoma (OCCC) carries a very poor prognosis in large part secondary to the extremely high rate of resistance to standard platinum and taxane chemotherapy. Signal transducer and activator of transcription 3 (STAT3) expression and activation has been shown to regulate tumor progression in various human cancers, though has not been well studied in OCCC. Preliminary work in our lab has demonstrated constitutive activation of STAT3 (pSTAT3^{Tyr705} or pSTAT3⁷²⁷) in OCCC cell lines as well as human OCCC tumor tissue samples. Significantly, pSTAT3 is expressed in the absence of other forms of activated STAT (pSTAT1, 2, 6). Therefore, this work was planned to investigate the role of STAT3 and examine the efficacy of a novel anti-cancer compound -HO-3867, which is an inhibitor of STAT3, using known OCCC cell lines. Results demonstrate that treatment with HO-3867 decreased expression of pSTAT3^{Tyr705} as well pSTAT3^{Ser727}, while total STAT3 remained constant. STAT3 overexpression increased the migration capability in OVTOKO cells *in vitro* and led to an increased tumor size when injected *in vivo*. The inhibitory effect of HO-3867 on cell proliferation and cell survival was accompanied by increased apoptosis, within 24 h post treatment. Treatment with HO-3867 resulted in a decrease in Bcl-2 and increase of cleavage of caspase 3, caspase 7, and PARP, confirming induction of apoptosis after treatment with HO-3867. In addition, HO-3867 significantly inhibited formation of human umbilical vein endothelial cells capillary-like structures and invasion at both 5 and 10 μ M concentrations. STAT3 expression plays an important role in the spread of OCCC *in vitro* as well as *in vivo*. Thus, we can exploit the STAT3 pathway for targeted drug therapy. Inhibition of pSTAT3 using HO-3867 in OCCC cell lines appears to be a promising therapy. This is of utmost importance given the poor response of OCCC to standard chemotherapy regimens.

Introduction

Ovarian clear cell carcinoma (OCCC) accounts for approximately 5–25% of epithelial ovarian cancers (EOCs) with varying incidence by population.^{1–3} Though rare, it presents a unique treatment challenge. Although OCCC tends to present at earlier stages, women with advanced disease have a poor prognosis compared with other EOC subtypes.^{3–5} This is illustrated by a meta-analysis of multiple clinical trials of advanced-stage EOC that noted a median overall survival of 21.3 months for women with OCCC, compared with 40.8

months for those with high grade serous ovarian cancer (HGSOC).⁵ It has been hypothesized that the discrepancy in outcomes may be at least in part secondary to tumor resistance to chemotherapy. Although platinum-based chemotherapy is standard first-line treatment for EOC, OCCC does not respond well to the regimen. In a multi-center retrospective study of patients with stage III/IV OCCC with measurable disease after surgery, response to platinum-based chemotherapy was 11.1%, as compared with 72.5% in patients with HGSOC.⁴ Another analysis of data from multiple clinical trials for advanced ovarian cancer found that overall response rates were 45% for OCCC versus 81% for HGSOC.^{6,7}

Although the relative chemoresistance of OCCC is felt to portend a worse prognosis, the precise mechanism of resistance remains unclear. Thus, to improve outcomes for patients diagnosed with OCCC, it is critical to find ways to overcome chemoresistance or identify new targets for therapy. Signal transducer and activator of transcription 3 (STAT3) is a transcription factor that mediates cytokine and growth factor signaling. STAT3 activation contributes to tumorigenesis by promoting cell proliferation, survival, angiogenesis, metastasis, inflammation, and immune evasion.^{8–11} STAT3 expression and activation has been shown to regulate

Key words: ovarian clear cell carcinoma, STAT3, HO-3867, curcumin analog, angiogenesis

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What's new?

The overall clinical response in chemotherapy-treated patients with recurrent ovarian clear cell carcinoma (OCCC) is only 25–40%. Development of new treatment approaches is urgently needed to improve the survival of patients afflicted with the disease. Here, the authors find high levels of expression of phosphorylated STAT3 (pTyr705 and pSer727) in OCCC and propose that this may represent a new chemoresistant pathway in these cancers.

tumor progression in various human cancers including HGSOC, though has not been well studied in OCCC. OCCC is characterized by a high rate of ARID1A and PI3K mutations which have been shown to cooperate in animal models to promote OCCC tumor growth through sustained IL-6 overproduction leading to increased activation of STAT3.¹² Additionally, Anglesio *et al.*¹³ demonstrated overexpression of IL6-STAT3-HIF pathway in OCCC. These findings support further investigation into STAT3 as a potential therapeutic target in OCCC.

We have previously shown that activation of STAT3 is necessary for HGSOC tumor progression/metastasis and targeting STAT3 with HO-3867 (a member of a novel class of anticancer drugs, diarylipenylpiperiden-4-ones (DAP's) significantly suppressed tumor growth and metastasis.^{10,14–16} Therefore, this study was designed (i) to investigate and establish the role of STAT3 in OCCC progression in OCCC cells and human OCCC patient tissue samples (ii) to examine the efficacy of our STAT3 inhibitor, HO-3867 *in vitro* and *in vivo* using the orthotopic tumor model.

Material and Methods**Culture of OCCC cells**

The OCCC cell lines OVTOKO, JHOC, OVISe and ES2 were a kind gift from Ikuo Konishi, Kyoto Medical University, Japan. The cells were cultured in T75 flasks in RPMI medium supplemented with FBS (10%) and Penicillin/streptomycin (1%).

Immunocytochemistry

Cells in RPMI medium were seeded onto sterile glass coverslips in 6-well plates with an average population of 50,000 cells/well. After 24 hr of culture, the cells were washed, fixed, and incubated with primary antibody according to a previously described protocol.

STAT3 overexpression/knockdown experiments

For downregulation of STAT3 in OVTOKO cells, a lentiviral system with a set of different siRNA – short-interfering Lentiviral Particles, Santa Cruz Biotechnology, TX) using Dharmafect Transfection Reagent (GE, Lafayette, CO) in OVTOKO cells. For STAT3 overexpression (STAT3 OE), we used EF. STAT3C.UbC.GFP, which was a gift from Linzhao Cheng (Addgene plasmid no. 24983), transfected into OVTOKO cells using Dharmafect Transfection Reagent (GE, Lafayette, CO).

Immunoblot analysis

Cells were treated with HO-3867 (5 or 10 μ M) for 24 hr. Following treatment, the cell lysates were prepared in non-denaturing lysis buffer as previously described.¹⁷

Cell migration assay

Cell migration assays were performed on both treated and nontreated cells using a wound-healing method.¹⁸

RNA isolation and reverse transcription PCR

OVTOKO cells were counted and plated in equal numbers in petridishes. The petridishes were treated with HO-3867 at 5 and 10 μ M concentrations, with at least three plates per treatment. At 24 hr posttreatment, the cells were collected and stored in the -80°C until further use. Total RNA was isolated using the RNeasy Mini Kit (Qiagen, Valencia, CA). RNA samples with an optical density A260/A280 ratio between 1.8 and 2.1 were used. Reverse transcription PCR (RT-PCR) was then performed using the Transcriptor First Strand Complementary DNA (cDNA) Synthesis Kit (Roche Applied Science) to synthesize cDNA. RT-PCR was performed with 1 mg of RNA template. The reaction was carried out using the Veriti Thermal Cycler (Applied Biosystems, Carlsbad, CA) and random hexamer primers.

Quantitative real time PCR

The genes studied for their relative genetic expression patterns are provided in Supplementary Table 1. LightCycler 480 SYBR Green I Master Mix (Roche Applied Science) was used to analyze 100 ng of cDNA from each experimental condition along with their respective primers (Supplementary Table 1). qRT-PCR was performed using the Light Cycler 480 System (Roche Applied Science). Each sample was normalized to the control gene glyceraldehyde 3-phosphate dehydrogenase (GAPDH).

STAT3 DNA-binding assays

After treatment with HO-3867 for 24 hr, a nuclear extract kit (Clontech Inc., Mountain View, CA) was used to prepare cell nuclear extracts following the manufacturer's protocol. Nuclear extracts were analyzed for STAT3 DNA binding activity using the TransFactor Universal STAT3-specific kits (Clontech Inc., Mountain View, CA) with an ELISA-based method.

Ubiquitin assay

To trace the ubiquitinated proteins in the cell lysates, agarose beads coated with domains having affinity to ubiquitin were incubated in the lysates at 4°C for 2 hr. After washing the beads, the ubiquitinated proteins were subjected to immunoblot for pSTAT3 and blotted by the ubiquitin antibody.¹⁹

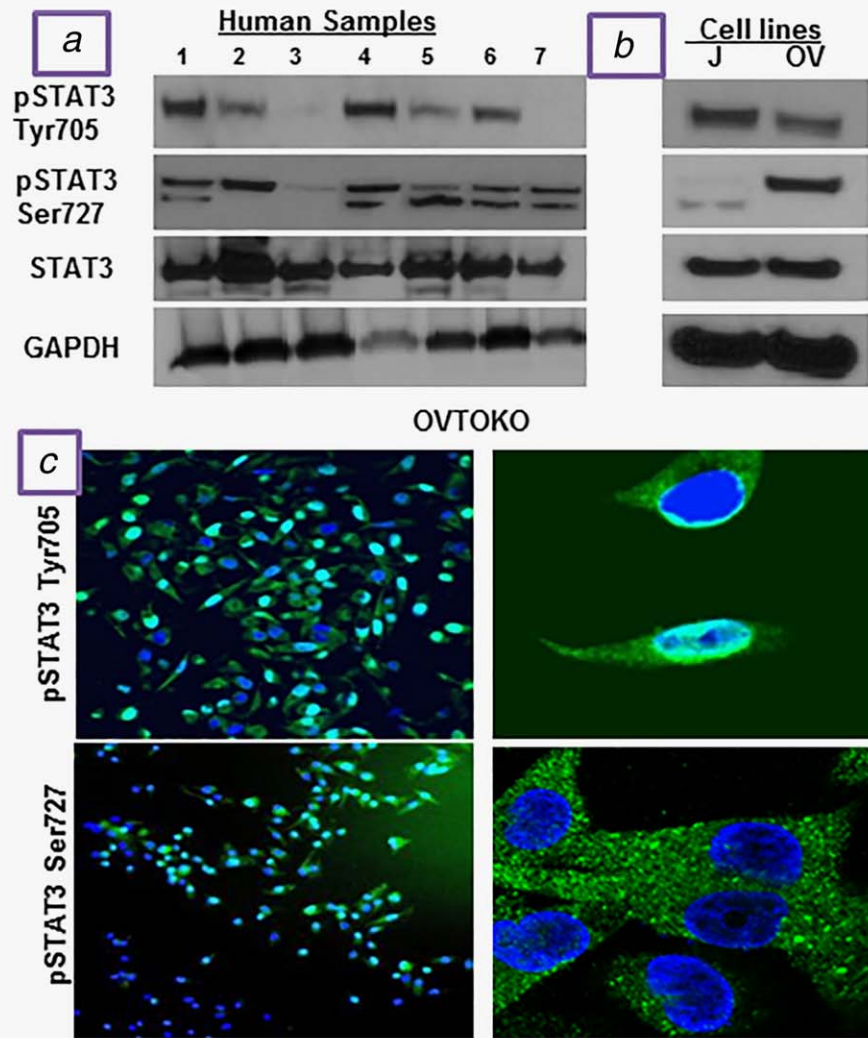


Figure 1. Validation of the expression of STAT3 in OCCC cell lines and human patient tissue samples. (a) Western blot for Human OCCC tumor samples obtained from seven consented OCCC patients. The extracted proteins blotted on a PVDF membrane were probed for pSTAT3 Tyr705 (topmost), pSTAT3 Ser727 (middle lane), total STAT3 (lower lane). All the samples show a medium to high pSTAT3 Tyr705 expression except for patient no. 3 and 7. All the patients showed pSTAT3 Ser727 and total STAT3 expression. (b) Western blot results for human OCCC cell lines-JHOC (J), and OVTOKO (OV). Both cell lines showed a high range of expression of pSTAT3 Tyr705 and pSTAT3 Ser727 (highly elevated in OV) and total STAT3. (c) Immunocytochemistry (ICC) of OCCC cell line OVTOKO with pSTAT3 Tyr705 (left 2 in the row) which is mostly nuclear and pSTAT3 Ser727 (right 2) which is predominantly cytoplasmic. [Color figure can be viewed at wileyonlinelibrary.com]

Evaluation of the bioabsorption of DAPs in ovarian cancer cells using electron paramagnetic resonance

Our previous study showed that cellular uptake of HO-3867 was significantly greater than curcumin.²⁰ We evaluated the bio absorption of HO-3867 compounds in ES2 and OVTOKO cells after 1, 3, and 6 hr posttreatment, using electron paramagnetic resonance (EPR), as previously described.²¹

Development of orthotopic tumor model

STAT3 OE OCC cells (3×10^6 cells in 100 μ L of PBS) were injected into the ovarian bursa of 6-week-old BALB/c nude mice from the OSU Transgenic mice core lab. In vivo MRI imaging was done periodically to check upon the tumor growth. After sacrifice, the tumor weight and volume was measured.

Statistical analysis

Data are presented as mean \pm 1 SD

Results

Validate the expression of STAT3 in OCCC cell lines and human patient tissue samples

We received 7 OCCC tissue samples from consented patients with OCCC at our institution and analyzed the expression of pSTAT3 Tyr705, pSTAT3 Ser727 and total STAT3. Five out of seven samples showed moderate to high expression of pSTAT3 Tyr705 on Western Blot (Fig. 1a). Expression was tested in OCCC cell lines (JHOC and OVTOKO). The studied cell lines showed moderate to high expression of pSTAT3 Tyr705, Ser727 and total STAT3, when analyzed by western

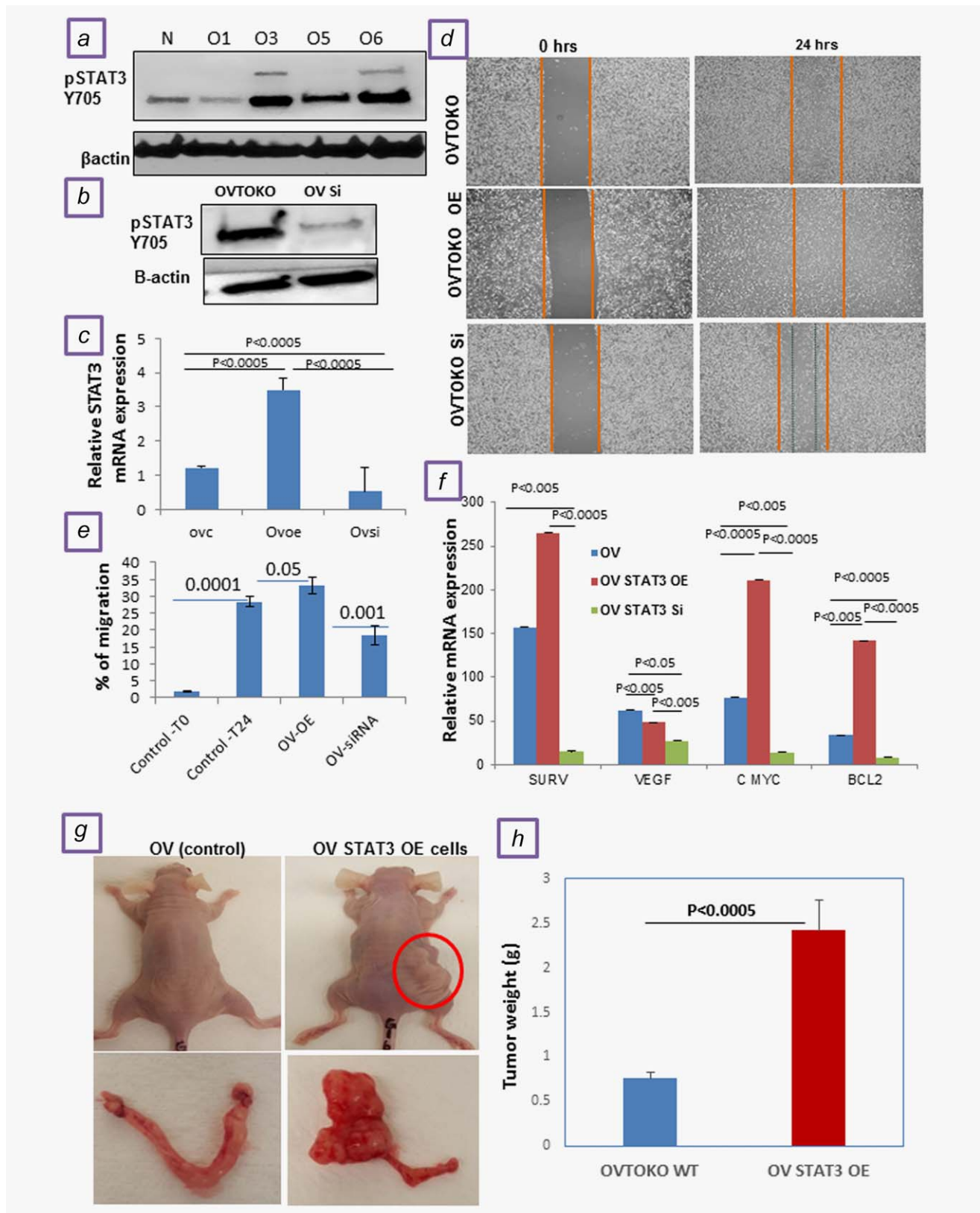


Figure 2.

blot (Fig. 1b). pSTAT3 Tyr705 showed high expressions in both tested cell lines (Fig. 1b). OVTOKO cells (90–95%) showed pSTAT3 Tyr705 localized in the nucleus as well as the cytoplasm while pSTAT3 Ser727 was mostly found in the cytoplasm of OVTOKO cells (Fig. 1c).

Effect of STAT3 OE or knockdown in OCCC cells *in vitro* as well as *in vivo*

In our previous studies with HGSOc, we found that the knockdown of STAT3 causes reduction in tumor growth and metastases *in vivo*.¹⁰ Based on this finding, we considered it imperative to determine if STAT3 plays a role in cell proliferation, migration, and survival of OCCC cell lines as well. Therefore, we developed OVTOKO STAT3 OE cells by transfecting OVTOKO cells with a plasmid construct constitutively expressing pSTAT3 Tyr705. Figure 2a shows the western blot expression of pSTAT3 Tyr705 in 4 of the tested OE clones. We found that clone O6 showed the highest expression of pSTAT3 Tyr705 so we moved forward with this clone for our future studies. Additionally we used siRNA for STAT3 to knockdown STAT3 from the OVTOKO cells, which was confirmed by western blot (Figs. 2b and 2c). Since an increased STAT3 expression is known to cause extensive migration as well as invasion,^{10,22} we analyzed the changes in the wound closure behavior of the cells. To confirm this, we performed a wound healing assay, where a monolayer of cells growing on a petri dish were scratched uniformly and observed at 24 hr for their ability to repair the damaged area for the control OVTOKO cells, OVTOKO STAT3 OE as well as OVTOKO STAT3 KO cells. The control OVTOKO and OVTOKO STAT3 OE cells migrated at about the same rate in the first 24 hr. However, OVTOKO STAT3 KO cells migrated to approximately 20–30% of the wounded area (Figs. 2d and 2e).

This was further confirmed at the mRNA level using real time quantitative PCR to check the relative expression of

genes related to angiogenesis, cell survival and cell proliferation. The relative mRNA expression of STAT3 was highest in OVTOKO STAT3 OE cells followed by the control OVTOKO cells and OVTOKO STAT3 KO cells (Fig. 1f). The STAT3 target genes like Survivin, VEGF, c-myc and Bcl2 were highest in the OVTOKO STAT3 OE cells and least in the OVTOKO STAT3 KO cells (Fig. 1f). Taken together, these results suggest that deleting STAT3 inhibits the expression of a number of genes responsible for cell survival, proliferation and angiogenesis. These findings were further evaluated *in vivo*. The OVTOKO control and OVTOKO STAT3 OE cells were injected into the right ovarian bursal cavity of nude mice (Fig. 2g). It is noteworthy that the mice injected with control OVTOKO cells failed to develop any primary tumor in the first 5 weeks postinoculation. However, the mice injected with OVTOKO STAT3 OE cells developed significantly large tumors as demonstrated in Figure 2h. No metastases were observed in any of the cases. These results add support to the role of STAT3 in OCCC tumor development and growth, and its ability to regulate the expression of different genes involved in survival and proliferation.

H0-3867 targets STAT3 and its associated genes in OCCC cells

We used OVTOKO and JHOC cell lines to test the therapeutic efficacy of our previously developed novel DAP compound on OCCC cell lines. We treated OVTOKO and JHOC cells with two different concentrations (5 and 10 μ M) of H0-3867. Western blot expression in Figure 3a clearly shows that the expression levels of both pSTAT Tyr705 and pSTAT Ser727 drastically decreases even in the lowest dose of 5 μ M and is almost non-existent in the higher dose of 10 μ M for both the tested OCCC cell lines (Fig. 3b). To determine whether H0-3867 inhibited pSTAT3 DNA-binding activity, OVTOKO and JHOC cells were treated with 10 μ M of HO-

Figure 2. Effect of STAT3 OE or knockdown in OCCC cells *in vitro* as well as *in vivo*. (a) Confirmation of STAT3 OE in OVTOKO cells using a Western Blot where N is control OVTOKO, O1, O3, O5, and O6 are overexpression clones 1,3,5 and 6. The bottom panel is β -actin expression as loading controls. In order to create STAT3 OE cells, OVTOKO cells were transfected with a vector harboring STAT3 gene or with the empty vector backbone for control. Based on the strongest expression in clone O6, we proceeded with this clone for all our future studies. (b,c) OVTOKO cells were transfected with STAT3 si RNA and the knockdown was confirmed by western blot analysis and RT-PCR. Knockdown of pSTAT3 705 in OVTOKO cells is clearly evident. (d) OVTOKO cells with STAT3 OE and STAT3 knockdown were subjected to wound healing assays in order to understand the correlation between STAT3 expression level and migration capability in OVTOKO cells. The panel on the left shows the plates at 0 hr after scratch and the panel on the right show the migration pattern after 24 hr. The cells over-expressing STAT3 migrate and close the gap within the first 24 hr (top and middle picture on the right panel) while the STAT3 knockdown cells (bottom left) migrate to fill 65–70% of the scratched area. (e) The percentage of migration was quantified using Image J software in 0 hr, 24 hr compared with STAT3 OE and siRNA experiments; (f) Real time quantitative PCR was performed for the relative mRNA expression of STAT3 in OVTOKO control, OVTOKO STAT3 OE and OVTOKO STAT3 Ko cells. The values were calculated using the $2^{-(\Delta\Delta C(T))}$ method and were normalized to the housekeeping gene GAPDH. Each experiment was repeated thrice and each sample had four replicates. The horizontal line over the bars indicates that the results are significant with a p values ≤ 0.005 . RT. Quantitative real time PCR (RT-qPCR) of OVTOKO, OV STAT3 Si and OVTOKO STAT3 OE cells for STAT3 and regulatory genes- survivin, VEGF, c-myc, and Bcl2. The very high relative mRNA expression was observed in OVTOKO STAT3 OE cells for all the studied genes. (g) The panel shows orthotopic mice injected with OVTOKO control and OVTOKO STAT3 OE cells. Mouse injected with OVTOKO STAT3 OE cells developed a big tumor in 4 weeks postinoculation but no tumor could be seen in the OVTOKO control mice. Some of the mice injected with control OVTOKO cells, which were left for 3 months, developed a noticeable tumor; (h) The weight of the ovarian tumor was significantly higher in the STAT3 OE transfected OV cells transplanted mice when compared with the OV cells transplanted mice. [Color figure can be viewed at wileyonlinelibrary.com]

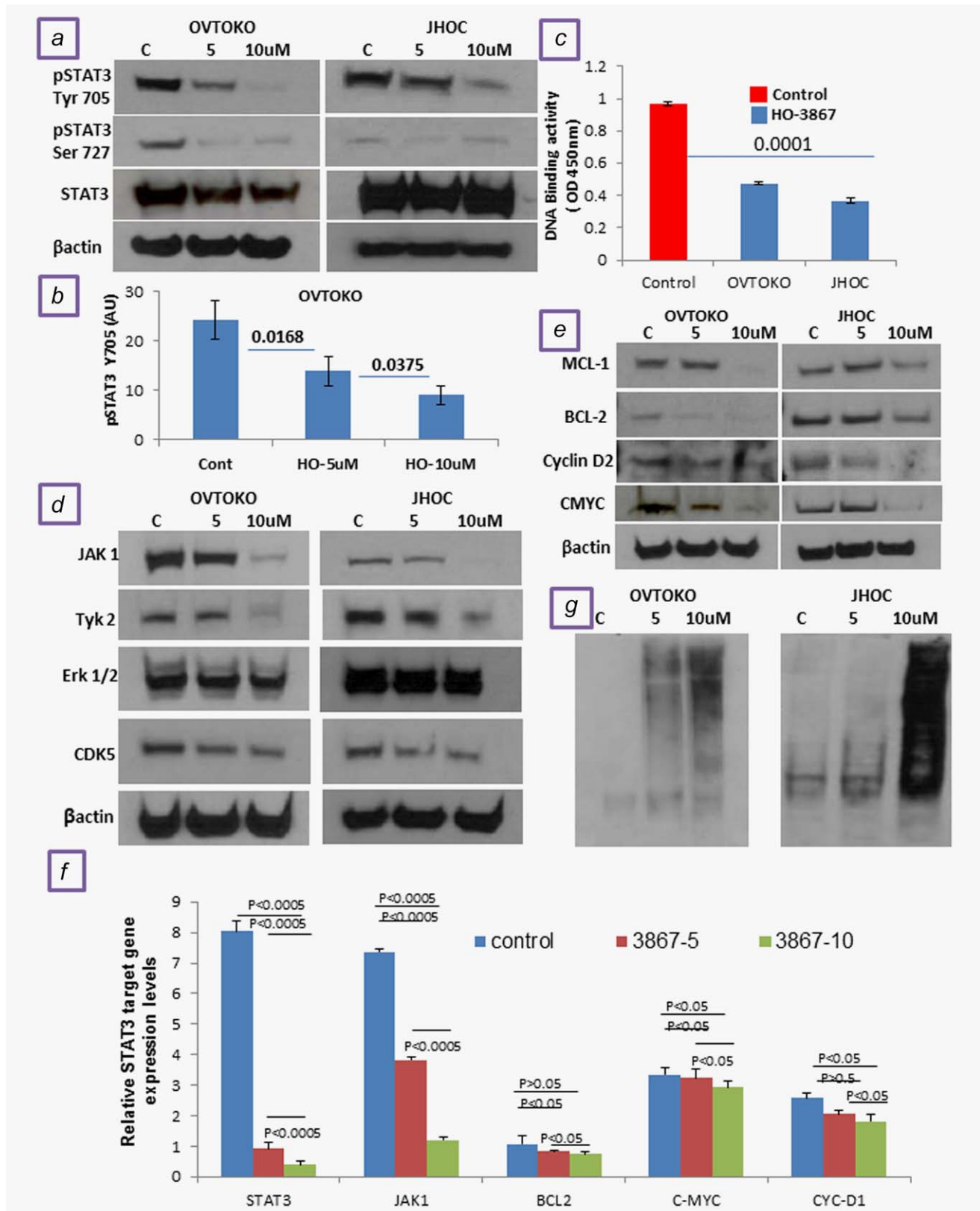


Figure 3. HO-3867 targets STAT3 and its associated genes in OCCC cells. (a) Western blot of proteins extracted from OCCC cells with/without treatment with HO-3867 at 5 or 10 μ M concentrations of HO-3867. The membrane was blotted for pSTAT3 705 (first row), pSTAT3 727 (second row), total STAT3 (third row) and β -actin (fourth row) for OVTOKO (left column) and JHOC cells (right column). (b) quantitative results of pSTAT3 Tyr705 bands by densitometry analysis in HO-3867 treated cells. (c) HO-3867 inhibit DNA-binding of STAT3, as indicated by ELISA assay results. (d) Western blot for genes associated with STAT3 upstream. The membrane was blotted for JAK1 (first row), TYK2 (second row), ERK 1/2 (third row), CDK5 (fourth row) for OVTOKO (left column) and JHOC cells (right column). E: Western blot for genes associated with STAT3 downstream. The membrane was blotted for MCL1 (first row), BCL2 (second row), cyclin D2 (third row), and c-myc (fourth row) for OVTOKO (left column) and JHOC cells (right column). (f) RT-qPCR of OVTOKO cells post treatment with HO-3867, with primers specific for STAT3 and regulatory genes- JAK1, BCL2, c-myc, and cyclin D1. (g) ubiquitination status of the OVTOKO (left panel) and JHOC (right panel) cells post treatment with HO-3867. Increased degradation in both OVTOKO and JHOC cells post treatment for both the concentrations. [Color figure can be viewed at wileyonlinelibrary.com]

3867, and samples were subjected to ELISA assay. HO-3867 compound inhibited STAT3 DNA binding activity in both cells when compared with control (Fig. 3c). Further, the STAT3 regulated genes like JAK1, Tyk2, and CDK5 also showed a diminished expression post treatment with HO-3867 for both the tested OCCC cell lines. MCL1, BCL2, Cyclin D2, and c-myc were also decreased post HO-3867 treatment for both JHOC and OVTOKO cells (Figs. 3d and 3e). This demonstrates that HO-3867 effectively targets STAT3 and its associated genes in OCCC cells. We sought to confirm these findings at the level of relative mRNA expression using qRT-PCR. The RNA obtained from the treated OVTOKO cells was converted to cDNA and subjected to real time PCR using the gene specific primers for STAT3, JAK1, BCL2, c-myc and cyclin D1. We found that mRNA expression for STAT3 and JAK1 was decreased post treatment with HO-3867 at both the doses of 5 and 10 μ M. However, the expression levels of BCL2, c-myc and cyclin D1 were not significantly decreased after treatment with HO-3867 (Fig. 3f).

Ubiquitination is the first step of the ubiquitin-proteasome pathway that regulates cells for their homeostatic functions and is an enzymatic, protein posttranslational modification process in which ubiquitin is transferred to a target protein substrate by a set of three ubiquitin enzymes.^{23–25} Given the importance of this process, it is plausible that HO-3867 has a function to regulate the ubiquitination. Therefore, we examined the ubiquitination status of the OVTOKO cells post treatment with HO-3867 and observed an enhanced degradation in both OVTOKO and JHOC cells post treatment for both the concentrations. Further, the degradation was concentration specific and increased with the increasing dosage of HO-3867 (Fig. 3g).

HO-3867 inhibits cell survival and increases the caspase cascade pathway

To evaluate the effect of HO-3867 on OCCC survival, we treated cells with different concentrations (1, 2.5, 5, 10, and 20 μ M) of HO-3867 at time points (24 and 48 hr). Cell viability was measured by MTT assay and cell count. The proliferation and survival of OCCC cells were inhibited by HO-3867 in a dose-dependent manner (Fig. 4a). Caspases are aspartic acid-specific proteases and are the major effectors of apoptosis. Therefore, in order to confirm if the activity of HO-3867 also activates the caspase cascade, we performed western blot on OVTOKO, ES-2, OVISe, and JHOC cells and found that PARP, cleaved caspase 3 and cleaved caspase 9 were all upregulated post treatment with HO-3867 (Fig. 4b). As an additional proof of concept experiment, we further studied the caspases in ES2 and OVISe cells and found that the higher dose of 20 μ M was able to activate cleaved caspase 3 and cleaved caspase 7 and poly(ADP-ribose)-transferase (PARP) was activated even by the lower dose of 10 μ M (Fig. 4b). This was further confirmed by Annexin V staining assays using a flow cytometer. It is known that in early stages of apoptosis, the plasma membrane excludes

viability dyes such as propidium iodide (PI) and display only Annexin V staining. We observed an increase in the percent of Annexin V positive cells with 30–51% of apoptotic cells post treatment as compared with 5–10% cells undergoing apoptosis in the untreated cells (Figs. 4c and 4d). The HO-3867-treated STAT3-overexpressing cells showed increased cell proliferation and decreased apoptotic proteins, in comparison with HO-3867 alone treated cells (Figs. 4e and 4f), suggesting that HO-3867 might target, at least in part, STAT3 in OCCC. EPR spectrum obtained from ovarian clear cell carcinoma showing the presence on HO-3867 in the oxidized (nitroxide) form. HO-3867 levels in the cell samples were determined using EPR, clear, triplet signals indicating the presence of the $-NO_2$ radical (nitroxide) form of HO-3867 were obtained from both cell lines and quantified (Figs. 4g and 4h).

Effect of HO-3867 on human umbilical vein endothelial cells

In the ovarian clear cell tumor microenvironment, tumor cells secrete vascular endothelial growth factor (VEGF) and other angiogenic factors that promote the growth and assembly of neighboring endothelial cells. Since HO-3867 can block secretion of important angiogenic proteins, we next examined if HO3867 treatment suppresses the growth and differentiation of human umbilical vein endothelial cells (HUVEC) in response to treatment with VEGF. HO-3867 significantly inhibited formation of capillary-like structures at both 5 and 10 μ M concentrations (Fig. 5a), indicating that signaling through STAT3 is necessary for VEGF-stimulated proliferation and tube formation of these endothelial cells. A 3D angiogenesis model utilizing VEGF stimulated HUVEC spheroids embedded in a collagen matrix recapitulates the inhibition of tube-like formation by HO3867 seen in the 2D model. After successful stimulation, new blood vessels sprouted into the collagen matrix within 2–3 days and were quantified in the presence and absence of HO-3867. Fewer number and shorter length of the sprouts demonstrates the anti-angiogenic activity of HO-3867 in comparison to the control group (Fig. 5b). HO-3867 treatment also significantly blocked the invasion of VEGF-stimulated HUVECs through a Boyden chamber (Fig. 5c).

Cytosolic STAT3 is known to be a co-regulator of F-actin fiber²⁶ and microtubule²⁷ formation. The disruption of lamellipodia formation and microtubule breakdown at the trailing edge may explain the drastically reduced invasion of HUVEC cells. In order to confirm this, HUVEC cultures were treated with VEGF alone, VEGF plus DMSO (drug vehicle) or VEGF with HO-3867 for 18 hr followed by staining for F-actin and β -tubulin (Fig. 5d). Thin, uniform fibers spanning the length of the cells were characteristic of the F-actin in the control and VEGF-treated cultures, with greater localization at the peripheral lamellipodia and intercellular junctions. The microtubules extended to the periphery of the cells forming a dense lattice that emanated from

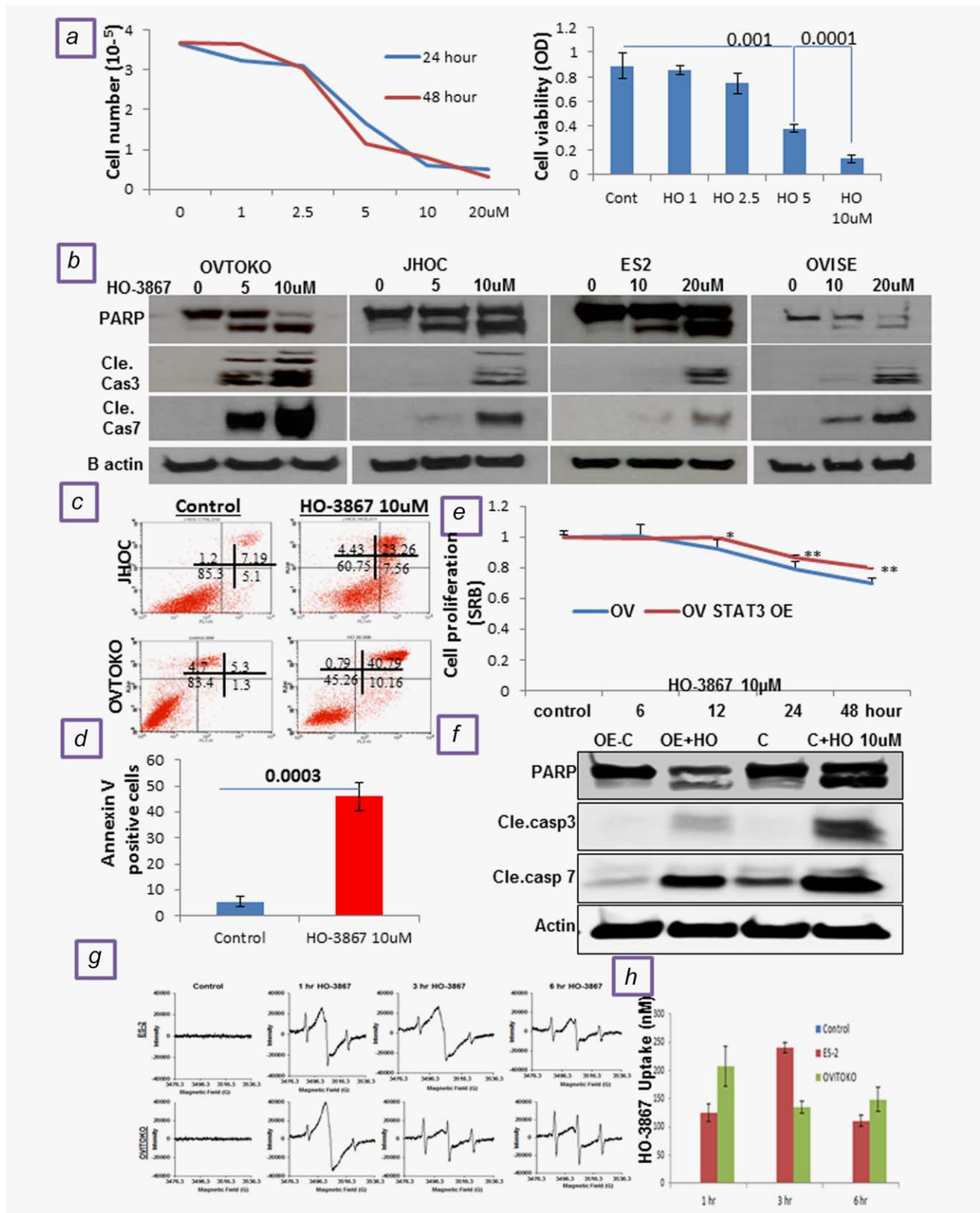


Figure 4.

the center of the cells. However, in STAT3 inhibited cultures, the cells displayed a condensed and rounded morphology. The actin had condensed into fewer fibers, and most strikingly, was completely absent from the leading edges of the cells (white arrows, Fig. 5d). The microtubule structures were additionally affected by the HO-3867 treatment. As highlighted by the arrowheads in Figure 5d, tubulin staining still showed that the microtubules emanated from the nuclear region of the HUVEC cells, but at the periphery, they curled over.

Discussion

Our results indicate high levels of expression of phosphorylated STAT3 (pSTAT3^{Tyr705} and Ser727) in OCCC which may prove to be an important pathway for OCCC cell survival, proliferation, and angiogenesis. Further, this is the first report demonstrating efficacy of HO-3867, a curcumin analog, in OCCC. Here we show that treatment with HO-3867 results in inhibition of OCCC cell survival and proliferation, induces apoptosis, and inhibits angiogenesis.

Recent studies have identified potential driver genes and aberrant signaling pathways involved in development of OCCC.^{7,28,29} As described earlier, ARID1A and PI3K mutations are detected in >55 and 30% of OCCC cases, respectively.^{2,12} Co-existent ARID1A and PI3K mutations have been shown to promote OCCC tumor growth through overproduction of IL-6 and activation of STAT3 lending STAT3 an attractive target for therapy.^{12,13} Although, the expression of STAT3 oncogene is well known in HGSOE,^{9,10,30,31} moderately little is known about the STAT3 signaling pathways in OCCC; its involvement in chemo-resistance and poor survival or its potential as a therapeutic target. This study identified (i) the expression of STAT3 in OCCC patient tissues and cell lines and (ii) STAT3 expression can up-regulate its target genes such as c MYC (proliferation), and Bcl-2 (cell survival and anti-apoptotic) thus playing a key role in proliferation and tumor progression of OCCC. Recent studies in renal clear cell carcinoma have associated higher expression of pSTAT3 with the tumor grade and

poor patient survival.^{32,33} It is clearly evident from our study that the expression of STAT3 might affect patient survival in OCCC, and targeting STAT3 using small molecule inhibitors could possibly inhibit OCCC tumor growth and help overcome the chemotherapeutic resistance in OCCC tumors.

We have recently reported four promising DAP compounds which are STAT3 inhibitors- H-4073, HO-3867, H-4318, and HO-4200.^{15,16,34} HO-3867 demonstrating high cytotoxicity towards human ovarian cancer cells, including those known to be chemo-resistant, under both *in vitro* and *in vivo* conditions.¹⁴ We have selected one of our best DAP compound STAT3 inhibitor-HO-3867, based on the STAT3 DNA-binding activity and selective cytotoxicity in cancer cells rather than normal epithelial cells, for the current study. HO-3867 significantly inhibits pSTAT3 in OCCC cells, through the upstream pathways of STAT3 such as JAK and TYK2. In addition, the current study showed an enhanced poly-ubiquitination of pSTAT3 by HO-3867 treatment in OCCC. These results suggest that in OCCC, HO-3867 targets STAT3 through dual pathways—the ubiquitin-dependent degradation of pSTAT3 and the inhibition of upstream regulators of pSTAT3, such as JAK1 and TYK2, eventually inhibiting cell proliferation and inducing apoptosis via modulating its target genes (Cyclin D1 and Bcl-2).

Targeting angiogenesis and VEGF is another therapeutic approach used in treatment of several solid tumors, OCCC included.^{13,35} VEGF was reported to be strongly expressed in OCCC cells and patient samples, both in early- and advanced-stage disease. High levels of VEGF expression are additionally associated with shorter survival than those with lower levels of VEGF expression in OCCC.^{6,35} In this study we document the direct effects of HO-3867, which include: suppression of HUVEC cell migration and invasion, tube formation of endothelial cells, and inhibition of microtubule assembly. This proves that STAT3 signaling closely governs these processes and affect STAT3 downregulated protein—VEGF in OCCC. Further, clinical evidence showed that

Figure 4. The survival-inhibitory effect of HO-3867 on OCCC. (a) The cell number was counted at days 1 and 2 after HO-3867 incubated. A dosage-dependent inhibition of cell survival is shown. Cell viability assay in HO-3867 treated OCCC. OCCC were seeded on a 6-well plate, and incubated with each concentration of HO-3867 up to 2 days. Activation of the caspase cascade pathway by HO-3867. (b) Western blot of proteins extracted from OCCC cells with/without treatment with HO-3867 at 5 or 10 μ M concentrations of HO-3867. The membrane was blotted for PARP (first row), cleaved caspase 3 (second row), cleaved caspase 7 (third row), and β -actin (fourth row) for OVTOKO (left column) and JHOC cells (right column). (c) Cytograms obtained from Annexin V assays of HO-3867 (10 μ M)-treated OVTOKO (bottom row) and JHOC (top row) cells. Each cytogram consists of data showing live cells (PI and Fluorescein isothiocyanate (FITC) negative) bottom left; early apoptotic population (FITC positive) bottom right; midlate stage apoptosis (PI and FITC positive) top left; necrotic/end stage apoptotic cells (PI positive, FITC negative) top right. Left column: control untreated cells; Right column: HO-3867-treated cells. The live cell population is indicated for each sample and is representative of typical data obtained. (d) Data shown are representative of triplicate experiments and the numbers refer to the mean (and standard deviation) live (PI and FITC negative) populations (expressed as a percentage of the untreated controls). (e) STAT3 OE transfected OCC cells are resistant to HO-3867 10 μ M treatment. (f) The STAT3 gene was transiently transfected into OV cells. At 24 hr after the transfection of the STAT3 gene, HO-3867 10 μ M was added into the culture medium of the cells. The untransfected cells with vehicle (DMSO) alone, the untransfected cells with the HO-3867 exposure. The STAT3 gene-transfected OV cells inhibit the HO-3867-induced cleavages of caspase-3, -7, and PARP. (g,h) Cellular absorption of HO-3867 10 μ M using EPR spectrometry to quantify the intracellular content of HO-3867 in various time points in OVTOKO and ES2 OCCC cell lines. Results showed that HO-3867 rapidly entered the cells within the first hour of treatment period. [Color figure can be viewed at wileyonlinelibrary.com]

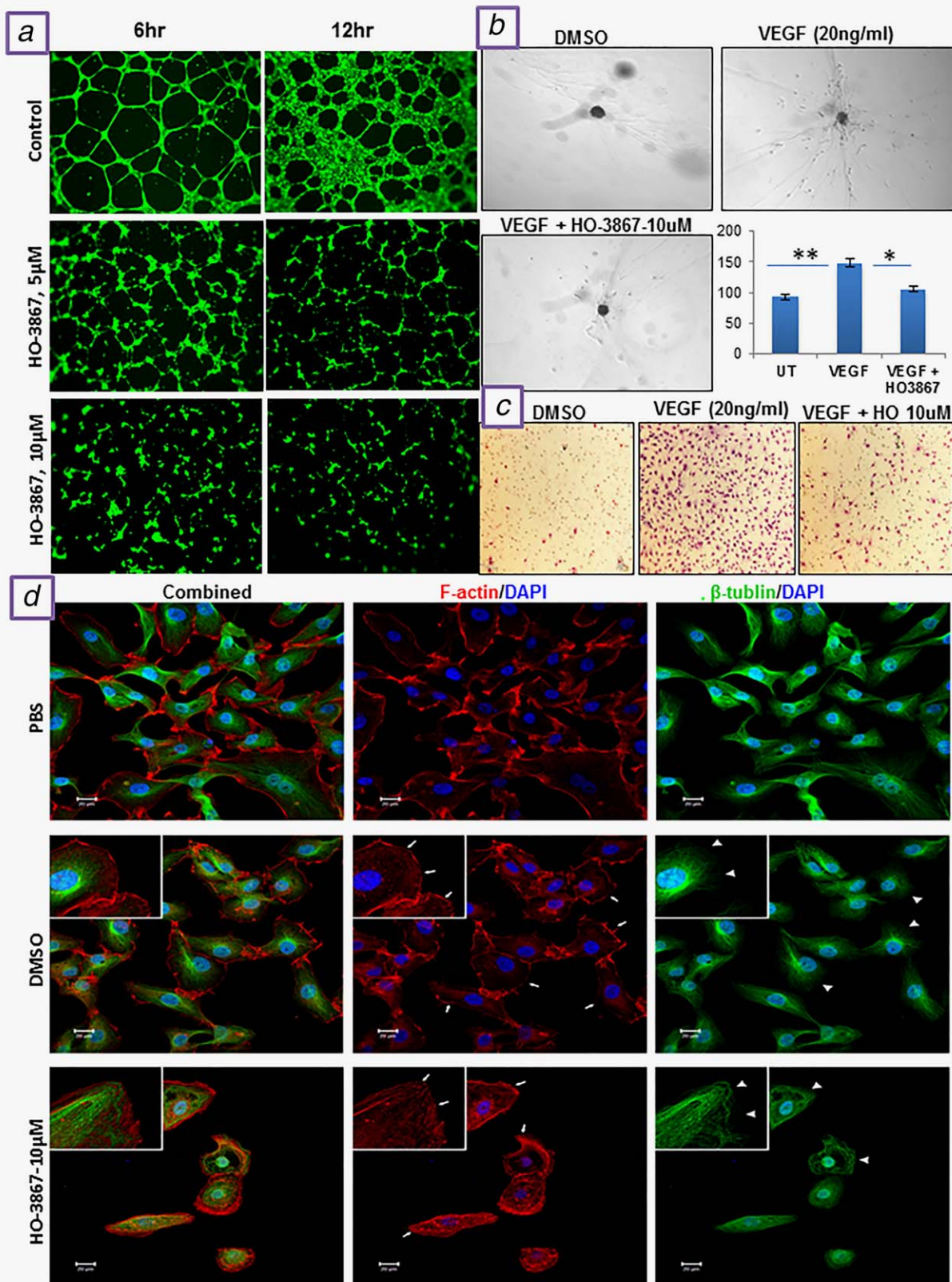


Figure 5. Inhibition of angiogenesis by HO-3867 correlates with inhibition of STAT3 phosphorylation. (a) Suppression of angiogenesis formation by HO-3867 in HUVEC cells after VEGF (10 ng/mL) stimulation. HUVEC cells were grown and stimulated with VEGF (10 ng/mL) for 10 min and treated with HO-3867 for 6 and 12 hr. (b) 3 D-Angiogenesis Assay (PromoCell) which demonstrates the entire angiogenesis process *in vitro*—proteolytic activity, migration, proliferation, and lumen formation. The number and length of the sprouts demonstrates the anti-angiogenicity of HO-3867 in comparison to the control group. HO-3867 inhibits HUVEC tube formation. HUVECs were grown in M200, and incubated with PBS or VEGF (10 ng/mL) for 18–20 hr in the absence or presence of HO-3867 (10 µM). Tube formation was quantified (bottom right panel) as described in Materials and Methods. (c) Invasion was determined using Matrigel coated membranes (photomicrographs show representative fields). Each data set represents the mean SE for at least three independent experiments. (d) The STAT3 inhibitor, HO-3867, induces cytoskeletal changes in cultured HUVEC cells. HUVEC cells cultured in 4-well chamber slides were treated with PBS, VEGF (10 ng/mL) alone, VEGF with DMSO or HO-3867 (10 µM) for 18 hr. The cultures were then probed using antiβ tubulin primary antibodies (green), and F-actin was stained using phalloidin (red). White arrows highlight F-actin localization at the leading edge, while white arrowheads indicate the curling of microtubules at the cell periphery. 2006×. Slice depth = 1 mm. Scale bar = 20 mm. Inset 400×.

sunitinib, a potent VEGF inhibitor, is more responsive in the treatment of clear cell carcinoma patients than serous tumors.^{6,36,37}

In conclusion, we demonstrate that pSTAT3 expression is highly elevated in OCCC tumors and is a potential therapeutic target for this disease. Our small molecule STAT3 inhibitor of HO-3867 efficiently inhibits pSTAT3 and its target

proteins, thereby impeding cell proliferation, angiogenesis and cytotoxicity in OCCC and seems promising in the future treatment of OCCC.

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