# Lysophosphatidic acid stimulates epithelial to mesenchymal transition marker Slug/Snail2 in ovarian cancer cells via Gɑi2, Src, and HIF1ɑ signaling nexus

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### ABSTRACT

Recent studies have identified a critical role for lysophosphatidic acid (LPA) in the progression of ovarian cancer. Using a transcription factor activation reporter array, which analyzes 45 distinct transcription factors, it has been observed that LPA observed robustly activates the transcription factor hypoxia-induced factor-1a (HIF1a) in SKOV3.ip ovarian cancer cells. HIF1a showed 150-fold increase in its activation profile compared to the untreated control. Validation of the array analysis indicated that LPA stimulates a rapid increase in the levels of HIF1a in ovarian cancer cells, with an observed maximum level of HIF1a-induction by 4 hours. Our report demonstrates that LPA stimulates the increase in HIF1a levels via Gai2. Consistent with the role of HIF1a in epithelial to mesenchymal transition (EMT) of cancer cells, LPA stimulates EMT and associated invasive cell migration along with an increase in the expression levels N-cadherin and Slug/Snail2. Using the expression of Slug/ Snail2 as a marker for EMT, we demonstrate that the inhibition of Gai2, HIF1a or Src attenuates this response. In line with the established role of EMT in promoting invasive cell migration, our data demonstrates that the inhibition of HIF1a with the clinically used HIF1a inhibitor, PX-478, drastically attenuates LPA-stimulates invasive migration of SKOV3.ip cells. Thus, our present study demonstrates that LPA utilizes a Ggi2-mediated signaling pathway via Src kinase to stimulate an increase in HIF1g levels and downstream EMT-specific factors such as Slug, leading to invasive migration of ovarian cancer cells.

### **INTRODUCTION**

Ovarian cancer remains as the most fatal gynecological cancers in the world with a five-year survival rate of only approximately 45% [1]. This is primarily due to our poor understanding of the disease in addition to the asymptomatic nature of this cancer in the early stages. LPA is known to elicit its diverse cellular responses by stimulating various members of the G protein families Gi, G12, and Gq [2–6]. Importantly, studies from several laboratories, including ours [7–9], have shown that LPA plays a crucial role in the progression of ovarian cancer [10–12]. Indeed, our lab and others

have shown that LPA-mediated signaling stimulates proliferation, migration, and invasion of ovarian cancer cells [5, 7–9, 13]. Increased levels of LPA in the ascites of the ovarian cancer patients and a robust membrane-bound LPA-synthetic machinery quite adjacent to LPA-receptors in ovarian cancer cells raise the concentration of LPA in the tumor microenvironment to micromolar concentrations, which may not allow LPA-receptor antagonist to be used as an effective therapeutic agents. Therefore, defining a signaling node downstream of LPA-receptors has become critical to develop therapeutic strategy for inhibiting LPAmediated oncogenic pathway(s). With this overarching goal, our laboratory as well as others have shown that the oncogenic activity stimulated by LPA involves the *gep* oncogenes G $\alpha$ 12 and G $\alpha$ 13 [14] as well as the putative *gip2* oncogene G $\alpha$ 12 [8, 15]. However, the role of these oncogenic G $\alpha$ -subunits in the activation of specific LPA-mediated oncogenic responses is far from clear. Therefore, we focused on defining the signaling nodes involved in LPA-mediated activation of a specific transcription factor, if any, which can be correlated with a critical oncogenic response.

HIF1 $\alpha$  has been shown to play a critical role in ovarian cancer malignancy, especially ovarian cancer cells found in the hypoxic conditions of the peritoneal cavity [16–18]. While HIF1 $\alpha$  is rapidly degraded in normoxia, it is rapidly stabilized by hypoxia, thereby promoting its transcriptional activity [19, 20]. In addition to hypoxia, several growth factors including LPA have been shown to induce the expression/stability of HIF1 $\alpha$  [21–24]. However, the mechanisms by which LPA stimulates the increase in the levels of HIF1 $\alpha$  and its activation are not fully understood.

The activation of HIF1 $\alpha$  involves its dimerization with the constitutively expressed HIF1 $\beta$  [25]. This is followed by the translocation of HIF1 $\alpha$  and HIF1 $\beta$ dimers to the nucleus and subsequent HIF1a mediated transcription of a multiple genes that can promote angiogenesis, glucose metabolism, cell survival, proliferation, and metastasis in cancer [26]. Importantly, one of the critical oncogenic responses orchestrated by HIF1 $\alpha$  is epithelial-to-mesenchymal transition (EMT) process [27-29] in which the cancer cells switch expression of markers of epithelial cells, such as E-cadherin to mesenchymal markers such as N-cadherin, vimentin, and transcription factors Snail1, Slug (Snail2), ZEB1, ZEB2 and Twist thereby facilitating the invasive migration and metastasis of cancer cells [28, 29]. Cells suppress the expression of proteins such as E-cadherin that allow for cell-to-cell attachment and increase the expression of proteins such as N-cadherin and vimentin that promote cell-detachment and migration. Furthermore, expression of EMT-specific transcription factors has been shown to increase the expression of proteins that can degrade extracellular components, which allow the cancerous cells to invade neighboring tissues [30]. This change in cellular markers characterizes a specific shift in the phenotype of the cancerous cells from being stationary to markedly increased invasive phenotype [28, 29]. Accordingly, EMT has been well recognized as a critical mechanism underlying carcinogenesis, cancer progression, and metastasis. Therefore, identifying pathways that can inhibit EMT are of critical importance for cancer therapy.

In the present study, using a transcription array to identify transcription factors activated by LPA-mediated signaling, we demonstrate that LPA potently stimulates the activation of HIF1 $\alpha$  via a pathway involving G $\alpha$ i2 and Src. We further demonstrate that that the activation

of LPA-G $\alpha$ i2-Src-mediated signaling pathway induces EMT in ovarian cancer cells and subsequent invasive migration of ovarian cancer cells that can be inhibited by PX-478, a clinically tested inhibitor of HIF1 $\alpha$ . Thus, our current study demonstrates that LPA stimulates a signaling nexus involving G $\alpha$ i2, Src, and HIF1 $\alpha$  to induce EMT and migration of ovarian cancer cells. Furthermore, we show that G $\alpha$ i2 signaling is necessary and sufficient for hypoxia-mediated induction of HIF1 $\alpha$  expression, which has not been shown, to our knowledge, by any previous studies to date.

# RESULTS

# LPA stimulates the activity and expression of HIF1a in ovarian cancer cells

In order to identify possible mechanism utilized by LPA to drive the progression of ovarian cancer we employed a transcription factor array that can analyze the activation profile of fortyfive different transcription factors. SKOV3.ip cells were stimulated with LPA for 20 minutes along with the appropriate vehicle control and the lysates were subjected to the transcription array analysis. Our results indicated that LPA stimulation activated several transcription factors that have previously been shown to be stimulated by LPA including STAT3 [7, 31, 32] and CREB [7, 33, 34], thus establishing the functional validity of our array analysis. In addition, we observed that LPA stimulated the activity of HIF1a by 150-fold compared to the untreated control cells and its activation far exceeded the activation of any other transcription factor on the array (Figure 1A). In light of the recent findings that HIF1 $\alpha$ plays a critical role of in ovarian cancer progression and malignancy [16–18], we sough to investigate the mechanism by which LPA stimulates the activity of HIF1 $\alpha$ . Since the expression levels of HIF1 $\alpha$  correlate with its activation [25, 35, 36], we first determined the expression levels of HIF1a following LPA stimulation in a panel of ovarian cancer cells. Our results indicated that LPA stimulated an increase in HIF1 $\alpha$  in three different ovarian cancer cell lines, namely OVCAR5, OVCAR2, and OVCA429 (Figure 1B). Thus, our results establish that the effect of LPA on HIF1 $\alpha$  is cell-line independent. Next, we carried out a time-course analysis for the expression of HIF1 $\alpha$  in SKOV3.ip cells in response to LPA. As shown in Figure 1C, LPA stimulated increase in the levels of HIF1a could be seen from 60 minutes onwards. Furthermore, it can be observed that the levels of HIF1a increases with time, reaching the maximum levels by 4 hours. Next, we carried out a dose-response curve with different concentrations of LPA. Our results indicated that LPA stimulated an increase in HIF1a levels in a dose-dependent manner (Figure 1C). Since the maximal increase of HIF1 $\alpha$ 

could be seen with 10  $\mu$ M LPA by 4 hours, the remainder of the experiments in this study involved the use of 10  $\mu$ M LPA.

### LPA signaling to HIF1a involves Gai2

Next, we sought to identify the downstream G protein that mediates LPA- signaling in this process. Previous studies from us [5, 7–9, 13, 37] and others [2, 38–41] have shown that LPA-stimulated oncogenic signaling is transduced by the heterotrimeric G protein  $\alpha$ -subunits, G $\alpha$ i2, G $\alpha$ q, and G $\alpha$ 12/13. Therefore, to

identify the G protein involved in LPA signaling to HIF1 $\alpha$ , we stably knocked out the expression of individual G $\alpha$ subunits, namely G $\alpha$ 12, G $\alpha$ 13, G $\alpha$ i2, or G $\alpha$ q in SKOV3. ip cells (Figure 2A) and stimulated these cells with 10  $\mu$ M LPA for 4 hours and monitored the expression levels of HIF1 $\alpha$ . Results from such analysis indicated that the G $\alpha$ i2-silenced cells showed a marked decrease in HIF1 $\alpha$  levels compared to the control cells. In contrast, the silencing of G $\alpha$ 12, G $\alpha$ 13, or G $\alpha$ q failed to have such an effect (Figure 2B). This was further corroborated using SKOV3.ip cells in which the expression of G $\alpha$ i2 was transiently silenced using G $\alpha$ i2-specific



**Figure 1:** LPA stimulates the activity and the expression of HIF1a. (A) LPA stimulates the activation of HIF1a. SKOV3.ip cells were stimulated with 20  $\mu$ M of LPA for 20 minutes or left untreated in serum-free condition. Activation profiles of 45 different transcription factors were analyzed with a Cignal<sup>TM</sup> 45-Pathway Reporter Array per manufacturer's protocol. (B) LPA-stimulated increase in HIF1a is independent of cell types. OVCAR5, OVCAR2, and OVCAR49 cells were serum-starved overnight and then stimulated with 20  $\mu$ M of LPA for 4 hours. The cells were lysed and the level of HIF1a was analyzed via Western blot. GAPDH was used as a loading control for each lane (n = 3). (C) Time-course Analysis of LPA stimulated increase in the levels of HIF1a. SKOV3.IP cells were serum-starved overnight for 16 hours following which they were stimulated with 20  $\mu$ M of LPA for the indicated time points. Lysates were subjected to immunoblot analysis using antibodies to HIF1a. The blot was stripped and re-probed with antibodies to GAPDH to ensure equal loading of proteins in each lane. (D) LPA-stimulated increase in the levels of HIF1a is dose-dependent. SKOV3.IP cells were serum starved overnight and then stimulated with the different concentrations of LPA for 4 hours. The cells were lysed and subjected to immunoblot analysis using antibodies to HIF1a. The blot was probed with GAPDH-antibodies to monitor equal loading of proteins.

siRNAs. As shown in Figure 2B, the ability to induce the expression levels of HIF1 $\alpha$  was drastically reduced cells in which the expression of G $\alpha$ i2 was silenced. To further confirm that G $\alpha$ i2 is involved in stimulating an increase in HIF1 $\alpha$  levels, we transiently transfected SKOV3.ip cells with a constitutively active form of G $\alpha$ i2 (G $\alpha$ i2Q205L). The expression of HIF1 $\alpha$  was monitored at 48 hrs following the transient expression of the constitutively active G $\alpha$ i2. As presented in Figure 2C, overexpression of constitutively active G $\alpha$ i2, without any exogenous LPA, resulted in an increase in HIF1 $\alpha$  levels, suggesting that G $\alpha$ i2-signaling is sufficient and responsible for mediating the effect of LPA in increasing the levels of HIF1 $\alpha$  levels in ovarian cancer cells.

#### LPA induces EMT in ovarian cancer cells

It has been well established that the induction of HIF1 $\alpha$  expression and its subsequent dimerization with HIF1 $\beta$  to function as a transcription factor in hypoxic conditions is involved in EMT and migration of many different cancer cell types [25, 27, 42]. Taken together with the observation that LPA stimulates HIF1 $\alpha$ , it can be surmised that the activation of HIF1 $\alpha$  by LPA could promote EMT in ovarian cancer cells. Therefore, we

tested whether LPA could stimulate EMT in these cells. Likewise, it has also been well established that Slug, a critical EMT-specific transcription factor, can be used as a marker to monitor EMT [43, 44]. In addition, previous studies have shown that HIF1 $\alpha$  can induce the EMT and expression of Slug in many cancer cells [45-47]. Therefore, we monitored the expression of Slug in response to LPA in ovarian cancer cells to test if LPA activated Slug and induced EMT in these cells. SKOV3. ip cells were stimulated with increasing doses of LPA for 4 hours and the expression levels of HIF1 $\alpha$  and Slug were monitored by immunoblot analysis from the lysates derived from these cells. As shown in Figure 3A, the expression of HIF1 $\alpha$  as well as Slug increased in these cells in a dose-dependent manner. Next, we sought to confirm that the increased expression of HIF1 $\alpha$  and Slug by LPA leads to an increased activation of HIF1 $\alpha$ and Slug. Since activated HIF1a and Slug translocates to nucleus, the nuclear levels of HIF1 $\alpha$  and Slug are often used as indices of their activation status [48, 49]. Accordingly, to assess the activation of HIF1 $\alpha$  and Slug by LPA, we determined the nuclear levels of HIF1 $\alpha$ and Slug. SKOV3.ip cells were stimulated with 20  $\mu$ M LPA or vehicle control for 4 hours, following which the nuclear extracts were isolated from these cells. The levels



**Figure 2:** LPA stimulates an increase in the levels of HIF1a via Gai2. (A) Confirmation of knockdown of G proteins in SKOV3. ip stable clones. SKOV3.ip cells were stably transfected with shRNA against Ga12, Ga13, Gai2 and Gaq. Stable knockdown of these proteins was confirmed by Western blot. (B) Effect of silencing different Ga-subunits on LPA-sti\imulated increase in HIF1a levels. The stable cell lines were serum-starved overnight and then stimulated with 10  $\mu$ M for 4 hours the following day. Immunoblot analysis was carried out with the cell lysates using antibodies to HIF1a followed by stripping and re-probing with antibodies to GAPDH. (C) Silencing of Gai2 abrogates the LPA-stimulated increase in of HIF1a levels. Expression of Ga<sub>i2</sub> was silenced by siRNA targeting Gai2 or control siRNA for 48 hrs. Cells were serum starved for 4 hours and then stimulated with 10  $\mu$ M LPA. After 4 hours, the lysates from the cells were subjected to immunoblot analysis with antibodies to HIF1a. The blot was sequentially stripped and probed with antibodies to Gai2 and GAPDH equal loading. (D) Effect of transient expression of constitutively active mutant of on HIF1a levels. SKOV3.ip (2 × 10<sup>6</sup>) cells were transiently transfected with either pcDNA3 control vector or pcDNA3 vector encoding Gai2QL, an activated mutant of Gai2. After 48 hours, the cells were lysed and the lysates were subjected to immunoblot analysis. The blot was sequentially stripped and gai2. The blot was sequentially stripped and re-probed with antibodies. The blot was sequentially stripped and re-probed with antibodies. The blot was sequentially subjected mutant of Gai2. After 48 hours, the cells were lysed and the lysates were subjected to immunoblot analysis using HIF1a-antibodies. The blot was sequentially stripped and re-probed with antibodies to Gai2 and GAPDH to monitor Gai2QL-expression and equal loading respectively (*n* = 3).

of HIF1 $\alpha$  and Slug in these extracts were monitored by immunoblot analysis. Results indicated that the treatment of ovarian cancer cells with 10  $\mu$ M LPA led to a dramatic increase of HIF1 $\alpha$  and Slug in the nucleus of these cells, thus pointing to the strong activation of these transcription factors by LPA (Figure 3B. To establish that the observed effects of LPA is not cell type-dependent, we examined the ability of LPA to induce the expression of Slug in three different ovarian cancer cells lines. As shown in Figure 3C, stimulation with 20  $\mu$ M of LPA induced Slug expression in all the tested ovarian cancer cell lines: OVCAR3 and OVCAR5 (representing high-grade serous ovarian cancer cell lines [50, 51]) and OVCAR2 cell lines.

# LPA induced EMT in ovarian cancer cells is dependent on Gαi2 and HIF1α

Our findings presented above (Figures 1–3) indicating the ability of LPA to stimulate the activation of HIF1 $\alpha$  via G $\alpha$ i2 taken together with the established role of HIF1 $\alpha$  in the regulation of EMT [27–29] point to a signaling paradigm in which the activation of HIF1 $\alpha$  by LPA via G $\alpha$ i2 is involved in induction of EMT in ovarian cancer cells. To validate such a paradigm, we first analyzed whether the silencing of G $\alpha$ i2 abrogates LPA-induced

expressions of Slug. In addition to Slug, we monitored the anticipated increased expression of N-cadherin and decreased expression of E-cadherin as additional markers for EMT [28, 29]. As shown in Figure 4A, LPA stimulated an increase in the EMT markers N-cadherin and Slug along with a decrease in E-cadherin. The silencing of Gai2 drastically blunted the ability of LPA to stimulate the increase Slug and N-cadherin as well as its ability to decrease the levels of E-cadherin. The role of  $G\alpha i2$  in this process was further confirmed using the constitutively activated mutant of Gai2. SKOV3.ip cells were transiently transfected with a constitutively active Gai2Q205L and the expression levels of Slug, N-cadherin, and E-cadherin were monitored by immunoblot analysis using the lysates from these transfectants. Consistent with the mediatory role for Gai2 in this process, the expression of Gai2QL dramatically increased Slug levels as well as N-cadherin levels with a concomitant decrease in E-cadherin levels (Figure 4B). Since our data demonstrates LPA-Gai2 signaling axis is involved in the activation of HIF1 $\alpha$ (Figure 2), it can be reasoned that the induction of EMT by LPA through Gai2 involves HIF1a. To establish such a role for HIF1 $\alpha$  in LPA-induced EMT in ovarian cancer cells, we tested whether the silencing of HIF1 $\alpha$ attenuates the expression of EMT markers in these cells.





The expression of HIF1 $\alpha$  was silenced in SKOV3.ip cell using HIF1 $\alpha$ -specific siRNA. The cells were stimulated with LPA for 4 hours and the expression levels of Slug, N-cadherin, and E-cadherin in the lysates were monitored by immunoblot analysis. As shown in Figure 4C, the silencing of HIF1 $\alpha$  in the cells inhibited the increased expression of N-cadherin and Slug along with the decreased expression of E-cadherin, thus validating the conclusion that HIF1 $\alpha$  in LPA-induced EMT of ovarian cancer cells is dependent on HIF1 $\alpha$ . Together, these results provide strong evidence that the induction of EMT by LPA involves G $\alpha$ i2-dependent pathway that utilizes downstream transcription factor HIF1 $\alpha$  to mediate the induction of EMT.

# LPA enhances hypoxia-induced activation of HIF1a via a Gai2-dependent pathway

Ovarian cancer cells are often found in the hypoxic environment of the peritoneal cavity and the core of the primary tumor [52]. A previous report has demonstrated that exogenous LPA stimulation synergistically enhanced hypoxia-induced stabilization of HIF1 $\alpha$  and hypoxia, which in turn, could enhance the oncogenic responsiveness of ovarian cancer cells to LPA [52]. However, the signaling mechanism and the role of G protein(s) in enhancing HIF1 $\alpha$  activation remain to be elucidated. Therefore, we set out to determine if LPA signaling could enhance the levels of HIF1 $\alpha$  and subsequent EMT of ovarian cancer cells in a hypoxic environment. To test, ovarian cancer cells incubated in 1% oxygen environment were stimulated with 10 µM of LPA for different lengths of time along with untreated controls. As anticipated, hypoxia alone increased the stabilization of HIF1 $\alpha$  (Figure 5A). However, HIF1 $\alpha$  levels were markedly increased when these cells were stimulated with LPA (Figure 5A). Similar to the results found in normoxic conditions, HIF1 $\alpha$  was maximally stabilized at 4 hours in hypoxic conditions. Next, we examined the effect of hypoxia alone or LPA plus hypoxia on the levels Slug, N-cadherin and E-cadherin. While hypoxic conditions alone induced HIF1 $\alpha$  stabilization and up-regulation of Slug, stimulation of these cells with exogenous LPA dramatically enhanced the up-regulation of Slug and HIF1α compared to hypoxic condition alone (Figure 5B). Furthermore, exogenous LPA drastically down-regulated the expression of E-cadherin. Overall, these data points to the synergistic role of LPA in enhancing the responsiveness of ovarian cancer cells to hypoxia and inducing EMT. Hypothesizing that the synergistic effect elicited by LPA could involve the Gai2-dependent mechanism, identified in normoxic conditions, we investigated whether the silencing of Gai2



**Figure 4: LPA stimulates the expression of EMT markers in ovarian cancer cells via Gai2 and HIF1a.** (A) Silencing of Gai2 inhibits LPA-mediated changes in EMT markers. SKOV3.IP cells were transiently transfected with siRNA specific for Gai2 or with scrambled control siRNA for 48 hours. The cells were serum-starved for 16 hours and then treated with 20  $\mu$ M of LPA for 4 hours. Lysates from these cells were subjected to immunoblot analysis with antibodies specific to Slug, E-cadherin, and N-cadherin. Silencing of Gai2 was confirmed by probing the blots with an antibody specific for Gai2. The blots were stripped and re-probed with antibodies to GAPDH to monitor equal loading of proteins. (B) Constitutively active Gai2 increases the levels of EMT-markers in ovarian cancer cells. SKOV3.IP (2 × 10<sup>6</sup>) cells were transiently transfected with either pcDNA3 vector control or vector encoding the activated mutant of Gai2, Gai2QL. After 48 hours, the cells were lysed and the lysates were subjected to immunoblot analysis using antibodies to Slug, E-cadherin, and N-cadherin. The blot was stripped and re-probed with antibodies to Gai2 and GAPDH to monitor Gai2QL-expression and equal loading respectively. The experiment was repeated thrice and the results from a typical experiment are presented. (C) Silencing of HIF1a inhibits LPA-mediated changes in EMT markers. SKOV3.IP cells were transiently transfected with siRNA directed against HIF1a or with non-targeting siRNA for 48 hours. The cells were serum-starved for 16 hours, following which they were stimulated with 10  $\mu$ M of LPA for 4 hours. Lysates derived from these cells were subjected to immunoblot analysis using antibodies specific Slug, E-cadherin, and N-cadherin. Silencing of HIF1a was confirmed by using an antibody specific to HIF1a. Levels of GAPDH were assessed to ensure equal loading of each lane. Results are from a typical experiment (*n* = 3).

abrogates such LPA-stimulated synergistic effect on the hypoxic response involving HIF1a and Slug. SKOV3.ip cells in which the expression of Gai2 was silenced were incubated in hypoxic condition and stimulated with 10 µM LPA along with untreated controls. Our results indicated that the silencing of Gai2 blunted the ability of LPA to enhance the expression of both Slug and HIF1a in hypoxic conditions (Figure 5C). Remarkably, silencing Gai2 alone, with no exogenous LPA present, led to decreased levels of HIF1 $\alpha$  and Slug compared to controls in hypoxic conditions with no exogenous LPA. This suggests that Gai2 is necessary for the induction of HIF1 $\alpha$  in hypoxic conditions (Figure 5C). Finally, to confirm that Gai2 is needed for hypoxia-induced EMT, both in the presence and absence of exogenous LPA, we analyzed the levels of E-cadherin and N-cadherin in response to LPA in ovarian cancer cells incubated in hypoxia or normoxia. As shown

in Figure 5D, silencing of Gai2 inhibited down-regulation of E-cadherin and up-regulation of N-cadherin indicating that Gai2 is necessary for inducing EMT in hypoxic conditions both with and without exogenous LPA.

#### LPA stimulates the activation of Src via Gai2

Recent reports from our lab [8, 15] have indicated that Src, via G $\alpha$ i2, is involved in initiating invasive migration of ovarian cancer cells. Additionally, Src has been shown to activate HIF1 $\alpha$  by diverse pathways involving both direct as well as indirect mechanisms [53–55]. Therefore, we investigated if Src is involved in activating HIF1 $\alpha$  and Slug. First, to confirm that G $\alpha$ i2 is needed for the activation of Src by LPA, we transiently silenced G $\alpha$ i2 in these cells and stimulated with 10  $\mu$ M LPA. Activation of Src was monitored by





the phosphorylation status of Src on tyrosine residue 419, which has been established as a good indicator of Src activation [8]. As shown in Figure 6A, silencing Gai2 inhibited the activation of Src, indicating that Gai2 is necessary for the activation of Src by LPA. To confirm that Gai2 is sufficient to activate Src, we transiently transfected ovarian cancer cells with constitutively active Gai2, without any exogenous LPA stimulation, and checked the phosphorylation of Tyr-419 of Src. As shown in Figure 6B, constitutively active Gai2 led to the activation of Src indicating the ability of Gai2 to activate Src.

# LPA induces EMT of ovarian cancer cells via Gai2 and Src

Next, we investigated whether Src is involved in activating HIF1a and Slug. To test SKOV3.ip cells were treated with PP2, an inhibitor of the Src family of kinases, [53, 56], or Bosutinib, a clinically used Src inhibitor [57]. These cells were stimulated with LPA and the expression levels of HIF1a and Slug were monitored. The efficiencies of the inhibitors were monitored by he phosphorylation status of Tyr-419 of Src by immunoblot analysis. As shown in Figure 7A, treatment with PP2 and Bosutinib significantly inhibited the increase in the levels of Slug and HIF1 $\alpha$ , suggesting that Src is required for the effect of LPA on HIF1a- and Slug-levels. Next, we investigated whether Gai2 and Src were involved in the activation of Slug and HIF1 $\alpha$ . The translocation of Slug and HIF1 $\alpha$  to the nucleus has been used as an indicator of their respective activation. Therefore, we carried out immunofluorescence microscopic analysis to demonstrate the translocation of Slug (Figure 7B) and HIF1 $\alpha$  (Figure 6E) to the nucleus. Treatment of SKOV3.ip cells with 10 µM of LPA led to translocation of Slug and HIF1 $\alpha$  to the nucleus, indicating the activation of these transcription factors. More importantly, our results demonstrated that the stable knockdown of Gai2 or treatment of cells with either of the Src-inhibitors, PP2 or Bosutinib, significantly reduced the translocation of Slug (Figure 6D) and HIF1a (Figure 7B) to the nucleus in reponse to LPA. Together, these results indicate that LPA stimulates Gai2, which then utilizes Src to activate both Slug and HIF1 $\alpha$ , the mediators of EMT. To demonstrate the role of Src in LPA-induced EMT, we utilized immunofluorescence analysis of OVCA432 in response to exogenous LPA. OVCA432 cells have previously been shown to be a good model system for studying EMT in ovarian cancer cells due to fact that these cells express high levels of E-cadherin and have more of an epithelial phenotype compared to other ovarian cancer cell lines [58]. As shown in Figure 7A, OVCA432 cells in serum-free media demonstrated a distinct epithelial phenotype with strong E-cadherin staining between the cells. Conversely, treatment of OVCA432 cells with LPA overnight significantly decreased the overall expression levels of E-cadherin (Figure 7A, middle panel). In addition a drastic reduction in the presence of E-cadherin levels in cellular junctions could be observed in these cells. Interestingly, treatment of OVCA432 cells with Bosutinib drastically reduced the ability of LPA to down-regulate E-cadherin localization in inter-cellular junctions as well as its overall expression, indicating that the blockade of Src activation prevented LPA-induced EMT and that Src is needed for induction of EMT of ovarian cancer cells.



**Figure 6: LPA stimulates the activation of Src via Gai2.** (A) LPA-mediated activation of Src involves Gai2. SKOV3.ip cells were transiently transfected with either nonsense shRNA or shRNA that targeted Gai2. After 24 hours, the transfected cells were serum starved for 16 hours and treated with 10  $\mu$ M of LPA for 20 minutes along with untreated control groups. Lysates were subjected to immunoblot analysis using antibodies specific to Src phosphorylated on Tyr-419. The blot was stripped and probed for total Src to monitor expression levels and equal loading of proteins. A parallel blot was used to monitor for the silencing of Gai2 using antibodies specific to Gai2. The blot was stripped and re-probed with antibodies specific to GAPDH to monitor equal loading of proteins. (B) Constitutively active Gai2QL. After 48 hours, the transfectants were lysed and subjected to immunoblot analysis with antibodies specific to tyrosine-419 phosphorylated Src. The blot was stripped and re-probed for total Src. Expression of constitutively active Gai2 was confirmed with antibodies specific to Gai2. The blots were stripped and re-probed for total Src. Expression of constitutively active Gai2 was confirmed with antibodies specific to Gai2. The blots were stripped and re-probed with antibodies to GAPDH to monitor equal loading of proteins. Presented results are from a typical experiment (n = 3).

# Inhibition of HIF1α attenuates LPA-induced EMT and cell migration

Since EMT has been shown to promote invasive migration of cancer cells [28, 59], our results would suggest that the stimulation of HIF1 $\alpha$  by LPA is required for such invasive migration of ovarian cancer cells. It has been shown that PX-478, a clinically used inhibitor of HIF1 $\alpha$ , attenuates the activity of HIF1 $\alpha$  by lowering its expression levels [60, 61]. Therefore, we tested whether LPA-stimulated invasive migration of ovarian cancer cells could be attenuated by the inhibition of HIF1a by PX-478. As shown in Figure 7, LPA potently stimulated the invasive migration of SKOV3.ip cells. However, treating these cells with the escalating doses of PX-478 led to a concentration-dependent inhibition of invasive migration (Figure 8A and 8B). Even at the lowest tested dose of 25  $\mu$ M concentration at which the PX-478 markedly reduced the cellular levels of HIF1α (Figure 8C) -, PX-478 attenuated the invasive

migration of SKOV3.ip cells (Figure 8A and 8B). Together, our findings establish the functional role for LPA-Gai2-Src stimulated HIF1 $\alpha$  in promoting EMT phenotype in ovarian cancer cells involving the overexpression of Slug and increased invasive migration.

# DISCUSSION

Previously, we have shown the critical role of LPA in promoting cell proliferation and migration in ovarian cancer cells [7–9, 15] in addition to establishing the role of downstream Gai2, Ga12, and Ga13 in ovarian cancer xenograft growth *in vivo* [14]. In the present study, we demonstrate the role of LPA in the potent activation of HIF1 $\alpha$  in ovarian cancer cells. Using a transcription factor reporter array, we show that LPA stimulates the activation of HIF1 $\alpha$  by 150-fold within 20 minutes (Figure 1). Further analysis of the underlying mechanism indicates that the activation of HIF1 $\alpha$  by LPA involves Gai2-dependent signaling mechanism (Figure 2) that



**Figure 7:** Src is required for the expression and activation of HiF1a and Slug. (A) Src is required for LPA-mediated increase in the levels of HIF1a and Slug. SKOV3.ip cells were stimulated with 10  $\mu$ M LPA with or without the incubation with 10  $\mu$ M PP2 or 1  $\mu$ M Bosutinib for 4 hours. Lysates from these cells were subjected to immunoblot analysis using antibodies to HIF1a, Slug, pY419 Src and total-Src. (B) Knockdown of Gai2 or inhibition of Src ablates LPA-stimulated activation of HIF1a and Slug. Parental SKOV3.ip cells, SKOV3.ip cells in which Gai2 using specific shRNA, SKOV3.ip cells treated with 10  $\mu$ M PP2, or SKOV3.ip cells treated with 10  $\mu$ M Bosutinib were stimulated with 10  $\mu$ M LPA for 4 hours along with the unstimulated control. Cells were stained with an antibody against Slug or HIF1a and counterstained with DAPI. (C) Inhibition of Src attenuates LPA-induced EMT. OVCA432 cells were stimulated with 10  $\mu$ M LPA for 4 hours or pre-treated 1  $\mu$ M Bosutinib prior to stimulation with 10  $\mu$ M LPA for 4 hours along with untreated control group. At 4 hours cells were stained with an antibody against E-cadherin and counterstained with DAPI (n = 3). can be correlated with the increased expression of Snail, which is involved in promoting the transcriptional activation of EMT-specific genes (Figures 3 and 4). Our results further demonstrate that the presence of LPA synergistically increases the expression levels of HIF1 $\alpha$ through a Gαi2-dependent signaling pathway in hypoxic conditions, such as those found in the ascites fluid of ovarian cancer patients. In this context, we demonstrate here that silencing Gai2 alone, completely of abrogates hypoxia-induced expression of HIF1 $\alpha$  even in the absence of exogenous LPA. These results indicate that Gai2 is required for the hypoxia-induced expression of HIF1a in ovarian cancer cells (Figure 5). Interrogating further, we establish that Gai2-dependent signaling involves Src to activate HIF1 $\alpha$  (Figure 6). LPA-stimulated signaling nexus involving Gai2 and Src, thus formed, induces

EMT in ovarian cancer cells as indicated by the nuclear translocation of Slug and up-regulation of N-cadherin expression levels and loss of E-cadherin between cells (Figure 7). Finally, we demonstrate that the inhibition of this pathway using PX-478, a HIF1 $\alpha$  inhibitor, drastically decreases the migration of ovarian cancer cells (Figure 8). Thus, Our results presented here demonstrate for the first time that LPA signaling in normoxic conditions activates a G $\alpha$ i2-Src-dependent pathway to up-regulate the transcription factors HIF1 $\alpha$  and Slug and the demonstrated G $\alpha$ i2-Src pathway is critical for induction of EMT by LPA. Importantly, we show that G $\alpha$ i2 is necessary for hypoxia-induced activation of HIF1 $\alpha$  and Slug.



**Figure 8: Inhibition of HIF1a attenuates LPA-induced invasive-migration of ovarian cancer cells.** SKOV-ip cells were stimulated with LPA or LPA plus varying concentration of HIF1a inhibitor PX-478 for 16 hours. A transwell invasive migration was carried out as detailed under Materials and Methods following our previously published procedure. Representative micrograph images of Hemacolor stained invaded cells were obtained at  $100 \times$  for each of the experimental groups are presented (**A**). Migrated cells were quantified and presented as fold change over untreated control values (**B**). Immunoblot analysis with antibodies to HIF1a was carried out to verify the inhibitory effect of PX-478 on HIF1a expression levels (**C**). Representative data from a typical experiment is presented (n = 3; \*\*\*p < 0.0001).

Previous studies have shown that the levels of HIF1 $\alpha$  is regulated at multiple levels such as the inhibition of degradation of HIF1 $\alpha$ , increased translation of HIF1 $\alpha$ mRNA, and enhanced transcription of HIF1 $\alpha$  gene [62]. Although these mechanism are not mutually exclusive, the observation that the effect of LPA on the activation of HIF1 $\alpha$  can be seen by as early as 20 minute, points to the role of LPA in the stabilization of HIF1a through Gai2. A novel and yet another critical observation reported here is the finding that the stabilization of HIF1 $\alpha$  in hypoxic condition - independent of exogenous LPA treatment is also dependent on Gai2. Previous findings from our laboratory have indicated that LPA-Gai2 signaling could rapidly stimulate Rac via p130Cas/Src dependent pathway [15]. It has also been shown that Src can stimulate an increase in the levels and subsequent activation of HIF1a involving Rac-stimulated ROS generation via NADPH oxidase [54]. Connecting these two independent observations, our data presented here points to a signaling paradigm in which the signaling by LPA propagates through Gai2 and Src to HIF1a (Figure 9).

HIF1 $\alpha$  signaling has been linked to EMT and cancer progression. There is voluminous reports that HIF1 $\alpha$  and hypoxic conditions are linked to EMT [27]. Indeed, HIF1 $\alpha$ has been linked to directly up-regulating the expression of Twist [63, 64] and Snail [65]. Similarly, several very recent reports have shown that HIF1 $\alpha$  can induce expression of Slug [45–47]. A recent has shown that knockdown of HIF1 $\alpha$  resulted in decreased mRNA levels of Slug, indicating that HIF1 $\alpha$  is directly or indirectly involved in Slug expression [45]. It has also been shown that Slug is involved in the transcriptional repression of E-cadherin [66, 67]. In this context, our current study defines the upstream signaling mechanism involving a specific G protein in the activation of HIF1 $\alpha$  and subsequently Slug. Future studies should define the mechanism by which HIF1 $\alpha$  increases the transcription of Slug. Nevertheless, it is clear that HIF1 $\alpha$  and Slug are two transcription factors whose levels are directly increased by LPA via the Gai2-Src signaling node. These findings provide evidence that this signaling node can be targeted directly to inhibit expression of Slug and stabilization of HIF1 $\alpha$ . Since these two transcription factors have been shown to be important in EMT and drug resistance in a multitude of cancers, our findings underscore the possibility that the pathway we have identified here is directly contributing to ovarian cancer progression and potentially drug-resistance. Recent finding that the expression of Gai2 increases in advanced stage ovarian cancers [68], further points to critical role of  $G\alpha i2$  and the identified pathway as a potential therapeutic signaling node in advanced ovarian cancer. Moreover, besides contributing to cell migration and EMT, it is highly likely that HIF1 $\alpha$  activation via the G $\alpha$ i2-Src pathway is also involved in other effects such as resistance to apoptosis, enhanced glucose uptake, and angiogenesis, all of which have been shown to be critically involved in tumor growth and progression.

Of critical importance, we show here that  $G\alpha i2$  is also necessary for HIF1 $\alpha$  activation independent of exogenous LPA signaling. Thus, there is a distinct possibility that LPA and/or other ligands that utilize  $G\alpha i2$ , such as CXCL12, could be responsible for activation



**EMT/Migratory Phenotype** 

Figure 9: Schematic representation of Gai2–Src-HIF1a nexus in the regulation of EMT in ovarian cancer cells. Stimulation of LPA receptors leads to the activation of Gai2 and the subsequent activation of Src, as we have shown previously [8, 15]. Src-dependent signaling, in turn, stimulates the upregulation and activation of HIF1a. HIF1a, once activated, stimulates the expression and resultant activation of Slug and other EMT-specific factors including N-cadherin, thereby promoting EMT and associated invasive migration of cancer cells.

HIF1a in hypoxic conditions via autocrine/paracrine signaling. However, this needs to be investigated further. Nonetheless, our report demonstrates that Gai2 and Src are needed for HIF1 $\alpha$  activation in hypoxic conditions, indicating that inhibition of this pathway can suppress hypoxia-induced resistance in ovarian cancer patients. This is also the first report to our knowledge that has shown the importance of LPA-signaling via Gai2 in inducing EMT. Although a recent study reported the ability of LPA to induce EMT via Wnt/β-catenin signaling pathways [69], the underlying mechanism was not fully clarified. In this regard, our study presented here firmly establishes the role of Gai2-Src-HIF1a signaling nexus in promoting LPA-stimulated induction of Slug which is involved in EMT. Finally, this report adds to our previous findings [8, 15] that demonstrated the role of Gai2-Srcp130Cas-dependent mechanism in LPA-induced invasivemigration of ovarian cancer cells. It should be noted here that PX-478 has been shown to enhance the antitumor effects of both radio- as well as chemotherapeutic modalities [61, 70–72]. Based on the potent inhibitory effect of PX-478 on HIF1a levels, one can speculate that the treatment with PX-478 will downregulate the multitudes of HIF1a-regulated genes, including those involved in EMT phenotype such as Slug and resultant ddecrease in the expression of E-cadherins. In this context, our present observation that the clinically relevant dose of PX-478 (25 µM) potently inhibits the invasive migration of ovarian cancer cells (Figure 8) further establishes the therapeutic potential of the LPA-Gai-HIF1a signaling node (Figure 9), especially in HIF1 $\alpha$  overexpressing ovarian cancers.

# **MATERIALS AND METHODS**

### **Cells and reagents**

The ovarian cancer cell lines OVCAR2, OVCAR3, OVCAR5 and OVCA432 were kindly provided by Susan K. Murphy (Duke University, Durham, NC). SKOV3. ip1 cells (SKOV3.ip), an in vivo passaged variant of SKOV3 cells established by Yu et al., [73] were kindly provided by Dr. Robert C. Bast (MD Anderson Cancer Center, Houston, TX). OVCAR2, OVCAR3, OVCAR5, OVCA429, OVCA432 and SKOV3.ip cells were maintained in Roswell Park Memorial Institute (RPMI) 1640 media (Mediatech, Manassas, VA) containing 10% FBS (Gemini Bio-Products, West Sacramento, CA), 50 µ/mL penicillin, 50 mg/mL streptomycin (Mediatech, Manassas, VA) at 37°C in a 5% CO<sub>2</sub> incubator. For serumstarvation, the media used was RPMI 1640 with 0.1% BSA (Roche, Indianapolis, IN), 50 U/mL penicillin and 50 mg/mL streptomycin (Mediatech). Lysophosphatidic acid (1-oleoyl-2-hydroxy-sn-glycero-3-phosphate) was obtained from Avanti Polar Lipids (Alabaster, AL) and dissolved into 10 mM stock solutions in PBS with 0.1% BSA and stored at -80°C until use. Non-target control shRNA pLKO.1 vector construct was purchased from Sigma-Aldrich, St. Louis, MO (SHC002) whereas pLKO.1 vector constructs targeting Gai2 (RHS3979-9596925) was purchased from Open Biosystems (Lafayette, CO). The siGENOME Non-targeting siRNA (D-001206-13-05), siGENOME SMARTpool Gai2 (M-003897-00-0005) and HIF1a (M-004018-05-0005) were purchased from Dharmacon (Lafavette, CO). Peroxidase-conjugated antirabbit IgG was purchased from Promega (Madison, WI), and peroxidase-conjugated anti-mouse was purchased from GE Healthcare (Little Chalfont, UK). E-cadherin antibody was purchased from Santa Cruz Biotechnology (Santa Cruz, CA). HIF1a antibody was purchased from BD Biosciences (San Jose, CA). Alexa 568 anti-mouse and Alexa 488 anti-rabbit antibodies were purchased from Invitrogen (Eugene, OR). DAPI was purchased from Life Technologies and used at a working concentration of 0.25  $\mu g/mL$ .

# **Cell transfection**

All cells were transfected with a Nucleofector II system from Lonza (Allendale, NJ) using the provided transfection protocol for SKOV3 cells as published previously [8, 15]. SKOV3.ip cells were trypsinized and counted using a Countess automated cell counter (Life Technologies).  $2 \times 10^6$  cells per transfection cuvette were transfected with either non-targeting siRNA (100 nM), siRNA targeting Gai2 (100 nM), Gai2QL (2 µg) or pcDNA3 vector (2 µg) as indicated. After transfection, the cells were plated on 60 mm plates and allowed to adhere overnight. The following day the media was changed and the cells were allowed to grow until the end of the day. The cells were then were re-plated at a density of  $5 \times 10^5$  cells per 100 mm plate and allowed to adhere overnight. For stable transfection, SKOV3.ip cells transfected as previously described with shGai2 or control, nonsense shRNA and selected for the expression of shGai2 or the nonsense vector with puromycin [9].

# Hypoxia treatment

Hypoxia treatments  $(1\% O_2)$  were performed in  $INVIVO_2 400$  hypoxia workstation (Baker, Sanford, ME). Cells were incubated with 5% CO<sub>2</sub> and 1% O<sub>2</sub> at 37°C for the indicated lengths of time. After incubation, cells were collected and western blot analysis was carried out.

### Transcription factor reporter assay

Cignal<sup>™</sup> 45-Pathway Reporter Arrays (Qiagen, CA) was used to screen for different transcription factors upon LPA stimulation of SKOV3.ip ovarian cancer cells. Cells were seeded into wells (50,000 cells/well) of the Cignal<sup>™</sup> Finder 96-well plates (Qiagen, CA) to transfect the

reporters into cells via reverse transfection according to manufacturer's protocol. Briefly, reporter DNA constructs resident in each well of the plate were resuspended with 125  $\mu$ l Opti-MEM and complexed with 25  $\mu$ l of Lipofectamine 2000 (ThermoFisher, CA) transfection reagent. Each well is added with 5 × 10<sup>4</sup> cells suspended 25  $\mu$ l of Opti-MEM media. Transfection was allowed to happen by incubating the plate for 24 h at 5% CO<sub>2</sub> and 37°C. Following transfection, the cells were serum deprived for 16 h and treated with either vehicle (0.1% BSA in PBS) or LPA (20  $\mu$ M) for 20 min. Differential activation of the transcription factors were determined by lysing the cells and measuring the luminescence intensity following the manufacturer's protocol.

### **Fluorescence imaging**

OVCA432 and SKOV3.ip cells were plated at density of  $1 \times 10^5$  in 6-well plates with glass coverslips at the bottom. The cells were allowed to adhere overnight in a 37°C incubator with 5% CO<sub>2</sub>. The cells were washed 3× with sterile PBS and then serum-starved for 4 hours. After serum-starvation, the cells were treated with 10 µM of LPA for 4 hours. After LPA treatment, the cells were washed with ice-cold PBS one time and then treated with 4% paraformaldehyde for 15 minutes while rocking. The cells were then washed with PBS  $1 \times$  and then stored at 4°C until they were stained. All treatment groups were lysed with 0.25% Triton X-100 for 10 minutes and then washed with PBS 3×. After washing, the coverslips were blocked with 1% BSA in PBS for 30 minutes at room temperature while rocking. After blocking, the coverslips were washed with PBS 1×. After washing, the primary antibody was applied in 1%-BSA in PBS and rocked for 10 minutes at room temperature. The coverslips were then transferred to 4°C and incubated overnight while rocking. The following day the primary antibody was removed and the coverslips were washed  $3 \times$  for 5 minutes each. After washing, the coverslips were incubated with fluorescently tagged secondary antibody for 45 minutes at room temperature while rocking and covered with aluminum foil. After incubation with the secondary antibody, the coverslips were washed 1× with PBS and then stained with DAPI for 5 minutes. The coverslips were then washed  $3 \times$  with PBS for 5 minutes each wash and then allowed to dry. Once dry, the coverslips were mounted with ProLong Gold antifade from Life Technologies (Grand Island, NY) on glass slides. The coverslips were allowed to dry overnight at room temperature in the dark and then imaged the following day with a Nikon Eclipse Ni-U (Melville, NY) at  $600 \times$ .

#### **Collagen-1 transwell migration assay**

The Collagen-1 migratory invasion assay was performed as previously published [8]. Collagen type 1 was coated overnight onto 8-mm pore transwells at 4°C. The following day, the collagen-coated cell culture inserts containing  $5 \times 10^4$  SKOV3.ip cells were suspended in 200 µL serum-free media were placed in the well of a 24-well companion plate. Each well contained 500 mL media containing serum-free media control or serum-free media containing 10 µM of LPA. The cells were incubated for 20 hours. Non-migrating cells on the proximal side of the inserts were removed with a cotton swab, and the migrated cells on the distal side of the inserts were fixed and stained with Hemacolor (EMD Chemicals). Images were obtained at  $100 \times$  in 5 random fields for each group. The experiments were repeated 3 times. SKOV3.ip cells were transfected with the indicated plasmid (shRNA) and plated into 6-well plates for a total of 48 hours. Twentyfour hours after transfection, the cells were serum starved for an additional 24 hours, trypsinized, counted, and placed into the transwell.

# Western blotting

Immunoblot analysis with the indicated antibodies were carried out following previously published procedures [8, 74] and developed with a Kodak Image Station 4000 MM.

### Statistical analysis of data

An unpaired two-tail *t*-test with Welch's correction was performed to determine statistical significance.

# Abbreviations

LPA: lysophosphatidic acid; HIF1: hypoxia-induced factor-1; EMT: epithelial-to-mesenchymal transition.

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# **CONFLICTS OF INTEREST**

None.

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