



Original Articles

Targeting AR-Beclin 1 complex-modulated growth factor signaling increases the antiandrogen Enzalutamide sensitivity to better suppress the castration-resistant prostate cancer growth

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ABSTRACT

While the recently developed antiandrogen Enzalutamide (Enz) can extend survival for 4.8 months in castration-resistant prostate cancer (CRPC) patients, eventually most of these CRPC patients may develop resistance to the Enz without a clear mechanism. Here we found the expression of Beclin 1 was decreased in both Enz-resistant (EnzR) cell lines (EnzR1-C4-2 and EnzR2-C4-2B) as compared to their parental Enz-sensitive (EnzS) (EnzS1-C4-2 and EnzS2-C4-2B) cells, and targeting the Beclin 1 could lead to increase the Enz-sensitivity in these two CRPC cell lines. Mechanism dissection revealed that Enz might function *via* altering the interaction between Beclin 1 and the androgen receptor (AR) to decrease the activity of Beclin 1/Vps15/Vps34 complex thus increasing the ERK-mediated growth factor signaling to alter the Enz sensitivity. Interrupting the AR-Beclin 1/ERK signaling with ectopic BECN1 or ERK inhibitor led to alter the Enz sensitivity in both EnzR1-C4-2 and EnzR2-C4-2B cells compared to EnzS1-C4-2 and EnzS2-C4-2B cells, respectively. Together, these results suggest that targeting this newly identified AR-Beclin 1 complex-mediated ERK growth factor signaling with small molecule ERK inhibitor may help potentially develop new therapies to better suppress the EnzR CRPC.

1. Introduction

Prostate cancer (PCa) is the most commonly diagnosed cancer in males in the western world and causes over 80 deaths in America per day [34]. Although the androgen-deprivation therapy (ADT) with the recently developed antiandrogen Enzalutamide (Enz) could extend patients survival an extra 4.8 months [6,31,38], most of these patients still succumb to the disease following development of Enz-resistance.

Recently, accumulating evidence has suggested that the androgen receptor (AR) splice variants, such as AR-v7, that lacks the androgen-binding domain, may play critical roles for the development of Enz-resistance [3,20,22,25,26]. However, results from human clinical sample surveys [23,32] and *in vitro* cell lines [35,37] indicated other non-AR-v7 mechanisms may also play roles in Enz-resistance. In

particular, enhanced growth factor signals, including AKT activation, might also promote the development of Enz-resistance [18,27,37]. Indeed AKT signaling is found to be constitutively activated after Enz treatment [14,17,37]. Results from Brett S. Carver et al. [8] indicated that AR could negatively regulate AKT activation through transcriptionally increasing PHLPP expression, and suppressing the AR *via* Enz could lead to reduced PHLPP expression, thus enhanced AKT activation. Separately, besides the function in regulating macro-autophagy, Rohatgia et al. [30] found suppressing Beclin 1 led to sustain the growth factor-stimulated activation of AKT and ERK signals *via* altering the PI3P (+) lipid levels, thus likely contributing to breast cancer cell progression [8].

Here, we found Enz might function *via* promoting AR interaction with the Beclin 1/Vps15/Vps34 complex to suppress the latter's activity

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thus enhanced ERK signaling. Targeting this newly identified AR-Becn1 1 complex-mediated ERK growth factor signaling with ectopic BECN1 expression or ERK inhibitor could lead to increase Enz efficacy.

2. Materials and methods

2.1. Cell lines, inhibitors and antibodies

The Enz resistant (EnzR) C4-2 cells (EnzR1-C4-2) were generated by culturing Enz sensitive (EnzS) C4-2 cells (EnzS1-C4-2) under increasing Enz concentrations from 10 μ M to 40 μ M (every 20 days) for 3 months. The EnzS2-C4-2B and EnzR2-C4-2B cells were a gift from Dr. Allen Gao (University of California, Davis, CA, USA). Both EnzR1-C4-2 and EnzR2-C4-2B cells were maintained in RPMI media supplied with 10 μ M and 20 μ M Enz, respectively. Antibodies used for immunoblotting including GAPDH (sc-47724), AR (sc-816), ATG5 (sc-133158), ERK 1/2 (sc-514302), EGFR (sc-71034), VPS34 (sc-365404), cleaved PARP (h215), BECN1 (sc-48341) and IGF-1R β (sc-9038) were purchased from Santa Cruz Biotechnology (Dallas, Tx, USA), The BECN1(cst-3495), AKT (cst-9272), p-AKT-T308 (cst-13038) and p-AKT-S473 (cst-4060) antibodies were purchased from Cell Signaling Technology (Danvers, MA, USA), the VPS15 (17894-1-AP) antibody was purchased from Proteintech Company (Rosemont, IL, USA) and the ERK inhibitor was purchased from Cayman (1049738-54-6, Cayman, MI, USA).

2.2. Plasmids and lentivirus packaging

The pLKO.1-ShBECN1^{#1}, pLKO.1-ShBECN1^{#2}, pLKO.1-ShAR, pLVTHM-ShBECN1, pLVTHM-Sh-ATG5, pWPI-oeBECN1 and pWPI-oeAR, the pMD2G envelope and psAX2 packaging plasmids were transfected into 293T cells, using calcium-chloride transfection method for 48 h to generate the supernatant virus. Details of the plasmid sequencing were described in Table S1. For pLKO.1 vector-infected cells, puromycin (1 μ g/mL) was used to select stable expression.

2.3. Growth factor stimulation assays and western blotting (WB)

Cells were cultured in serum-starved media overnight comprising 0.1% BSA and then stimulated with IGF-1 (100 ng/mL) or EGF (50 ng/mL) for the time periods indicated. To inhibit internalization, cells were incubated with chlorpromazine (10 μ g/ml) for 1 h before the stimulation. Cells were infected with lentiviruses containing pLKO.1-shBECN1 or pWPI-oeBECN1 for knock down and rescue experiments and stimulated after 48 h infection.

We used cell lysis buffer (10 mM Tris-HCl/pH 8.0, 100 mM NaCl, 1 mM EDTA, 0.1% Nonidet P40, 1 mM DTT and 1 mM PMSF) to extract the protein for each sample, which were then diluted and boiled with 5 x loading buffer for 10 min. After that, these proteins were separated and run on 10% SDS/PAGE gel, and transferred onto PVDF membranes (Millipore, Billerica, MA, USA). The membranes were blocked for 1 h in 5% milk, then they were incubated with specific primary antibodies overnight. Next day, the blots were incubated with HRP-conjugated secondary antibodies (mouse or rabbit, depended on primary antibodies), and then visualized using the ECL system (Thermo Fisher Scientific, Rochester, NY, USA).

In addition, P13P (+) levels were determined by using the PI3P Mass ELISA Kit (Catalog # K-3300, Echelon Biosciences, USA) according to the manufacturer's instructions.

2.4. MTT, Trypan blue and BrDU assays for determining cell proliferation

For MTT assay, cell growth curve was determined by applying an MTT proliferation assay. Briefly, 2000–5000 cells were seeded on several 24-well plates containing 600 μ L RPMI media. Cell growth was tested at the indicated time points. After Enz treatment, 50 μ L of MTT reagent (Amresco Inc., Solon, OH, USA) was added to the media and

then the cells were incubated at the incubator for a further 2 h at 37 °C. After that, the media was removed, and 1 mL of dissolving reagent DMSO (Amresco Inc.) was added to dissolve the formazan crystals. The optical density value (OD) was determined at wavelength of 570 nm on a microplate reader. For Trypan blue staining assay, 200,000 cells were seeded on several 6-well plates containing 2 mL RPMI media, after 5 and 6 days culturing (with Enz treatment), the cell number was determined by Trypan blue assay. Briefly, we resuspended the cells with 1 mL RPMI after spin, then 50 mL cell suspension was mixed with 50 μ L Trypan blue solution. And then 20 μ L mix was examined using automated cell counting slides (Bio-Rad, Hercules, CA, USA). For BrdU (BrdU; Sigma-Aldrich, St. Louis, MO, USA) staining assay, refer to BrdU staining and BrdU assay protocol on the official website of Abcam (MA, USA)(<https://www.abcam.com/protocols/brdu-staining-protocol>). The staining was examined by fluorescence microscopic analysis (Olympus, Tokyo, Japan).

2.5. Co-immunoprecipitation (CO-IP)

Confluent cells on 150-mm plates were harvested and solubilized in lysis buffer. Insoluble material was removed by centrifugation. The supernatants were incubated overnight at 4 °C with 2 mg of anti-IgG, BECN1 or AR antibodies (Santa Cruz Biotechnology, Santa Cruz, CA, USA). Following the addition of 10 μ L Protein-A/G-agarose beads (Santa Cruz Biotechnology, Santa Cruz, CA, USA), mixtures were incubated for 2 h at 4 °C with rotation. Immune complexes were washed 15 times with regular lysis buffer; the agarose beads were then boiled for 10 min with loading buffer. Immunoprecipitates were run on 8% SDS/PAGE gel, followed by western blot assay with antibodies against AR, BECN1, Vps15, and Vps34.

2.6. Statistical analysis

Statistical analyses between two groups were performed using the two-tailed unpaired Student's t-test. Data are presented as mean \pm standard error (SEM) unless mentioned otherwise. A *P*-value of 0.05 was considered statistically significant.

3. Results

3.1. Enzalutamide regulates growth factor signaling

Enzalutamide (Enz) may extend the survival of CRPC patients an extra 4.8 months before the development of Enz-resistance [7]. The underlying mechanisms, especially its linkage to the growth factor signals, however, remain unclear. To investigate whether Enz-resistance was partly due to dysregulation of growth factor signals, we focused on the insulin-like growth factor-1 receptor (IGF-1R) and the epidermal growth factor receptor (EGFR), as their higher expression have been frequently observed in PCa patients and their activations are often correlated with poor prognosis of PCa (Fig. S1A-D) [1,12,43].

We compared the EnzR1-C4-2 cells with their parental Enz-sensitive C4-2 cells (EnzS1-C4-2) for the activation of growth factor signals [10,42], via assaying the phosphorylation of key downstream signaling effectors of the IGF-1R and EGFR, including serine/threonine kinases AKT and ERK, after treating with either EGF or IGF (Fig. 1 and Fig. S2), which could be used to activate these two signals. Before that, the basal phosphorylation levels of AKT and ERK were measured and described in Fig. S1E.

The results revealed that AKT and ERK were phosphorylated in a transient manner in response to EGF stimulation after pretreated with Enz for three days in EnzS1-C4-2 and EnzR1-C4-2 cells (Fig. 1A). We found the basal phosphorylation levels of AKT (S473 and T308) were increased after pretreated with 10 μ M Enz for three days. After EGF stimulation, the phosphorylation levels of AKT (S473 and T308) could still be increased in Enz-treated and Enz-resistant groups compared to

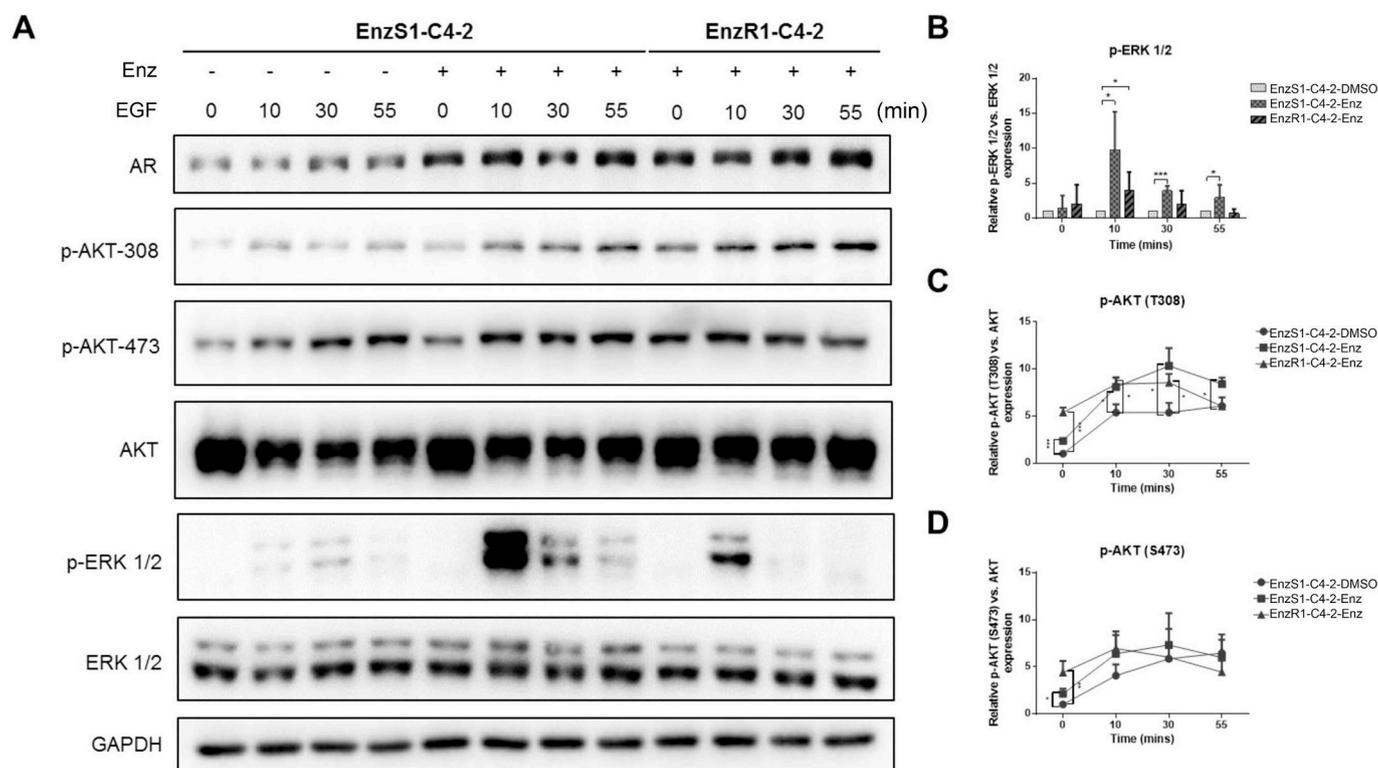


Fig. 1. Enzalutamide enhances growth factor signals. A. EnzS1-C4-2 cells were pretreated with Enz for three days, and serum-starved overnight in media containing 0.1% BSA, then stimulated with 50 ng/ml human EGF for the indicated time periods. The p-AKT-T308, p-AKT-T308 and p-ERK1/2 (T202/Y204) levels were detected by western blot. B-D. Quantification results of p-ERK 1/2, p-AKT-S473, and p-AKT-T308. The data shown represent the mean \pm SEM of three independent experiments. * $p < 0.05$, *** $p < 0.001$.

control group, but could not be sustained for an extended time, while the phosphorylation level of ERK was significantly increased and also sustained in response to EGF stimulation (quantified in Fig. 1B–D). Meanwhile, IGF was also used to stimulate the signaling cascade, and similar results for the AKT signal were found, but not ERK signal (Fig. S2A, quantified in Fig. S2B–C).

We then replaced the EnzS1-C4-2 cells with EnzS2-C4-2B cells, and used EGF to activate the growth factor signaling. We found the EGF stimulation could further increase the enhanced phosphorylation level of AKT (308T) by Enz, but still could not sustain this signal (Fig. S2D, quantified in Fig. S2E–F), while the ERK signal could be both increased and sustained in response to EGF after pretreated with Enz for three days, consistent with the earlier findings (Fig. S2D, quantified in Fig. S2E–F).

Although IGF and EGF both could activate AKT signal, only EGF could increase and sustain the ERK signal, and these results were consistent with previous findings that the ERK phosphorylation was more significant downstream of EGFR, while AKT phosphorylation was more significant downstream of the IGF-1R [30]. Together, these data suggest that Enz may increase the AKT and ERK growth factor signals, and Enz-resistance may be associated with enhanced growth factor signals in the PCa cells.

3.2. Beclin 1 may function via suppressing the ERK growth factor signaling to alter the Enz-resistance in PCa cells

Next, to dissect the molecular mechanism of how Enz alters these growth factor signals, we focused on the role of Beclin 1, as recent studies indicated that the Beclin 1 complex might function via regulating cellular macro-autophagy to control multiple growth factor signals [28,30,39]. For example, Beclin 1 complex may promote the conversion of PI3P to PI4P to terminate the signaling capacity of endosome-bound growth factor receptors to reduce intracellular signaling

transmission [28], thus loss of Beclin 1 could sustain growth factor stimulated activation of AKT and ERK signals resulting in cancer progression [30].

We first silenced the BECN1 expression via stably infecting BECN1-shRNAs (shBECN1^{#1} & shBECN1^{#2}) in the EnzS1-C4-2 cells (Fig. 2A), and results revealed that the phosphorylation levels of AKT (both T308 and S473) and ERK were significantly increased and sustained longer in the shBECN1-EnzS1-C4-2 cells compared with vector control in response to EGF stimulation (Fig. 2B, and quantification in Fig. 2C–E). However, ectopic Beclin1 expression (Fig. 2A) could only reverse the Enz-enhanced ERK growth factor signal rather than the AKT signal in response to EGF stimulation after pretreatment with Enz for three days (Fig. 2F, and quantification in Fig. 2G), suggesting that Beclin 1 loss could specifically regulate ERK growth signal to influence Enz-sensitivity.

We also found that Beclin 1 expression was significantly lower in EnzR1-C4-2 and EnzR2-C4-2B cells compared with EnzS1-C4-2 and EnzS2-C4-2 cells, respectively (Fig. 3A), and Enz-sensitivity was significantly decreased after silencing BECN1 expression in EnzS1-C4-2 and EnzS2-C4-2B cells (Fig. 3B–D). In addition, interruption approaches using ERK inhibitor (ERKI) treatment also demonstrated that inhibition of ERK activation could partially rescue the Enz-sensitivity in the shBECN1-EnzS1-C4-2 parental cells (Fig. 3E), as well as EnzR1-C4-2/EnzR2-C4-2B cells (Fig. S3A–B). Furthermore, we ectopically expressed BECN1 in EnzS1/R1-C4-2 and EnzR2-C4-2B cells, and found Enz-sensitivity was significantly increased (Fig. 3F–G and Fig. S3C).

In addition to MTT assay, we also conducted the Trypan blue, BrDU staining and western blotting assays to determine the effects of Enz on cell fate, and the results were consistent with the findings above that Enz mainly affects cell proliferation and not cell apoptosis (Fig. S3D–J).

Together, results from Fig. 2A–G, Fig. 3A–G, and Fig. S3A–J suggest that Enz may function via altering Beclin 1 to modulate the ERK-mediated growth factor signal and targeting these newly identified

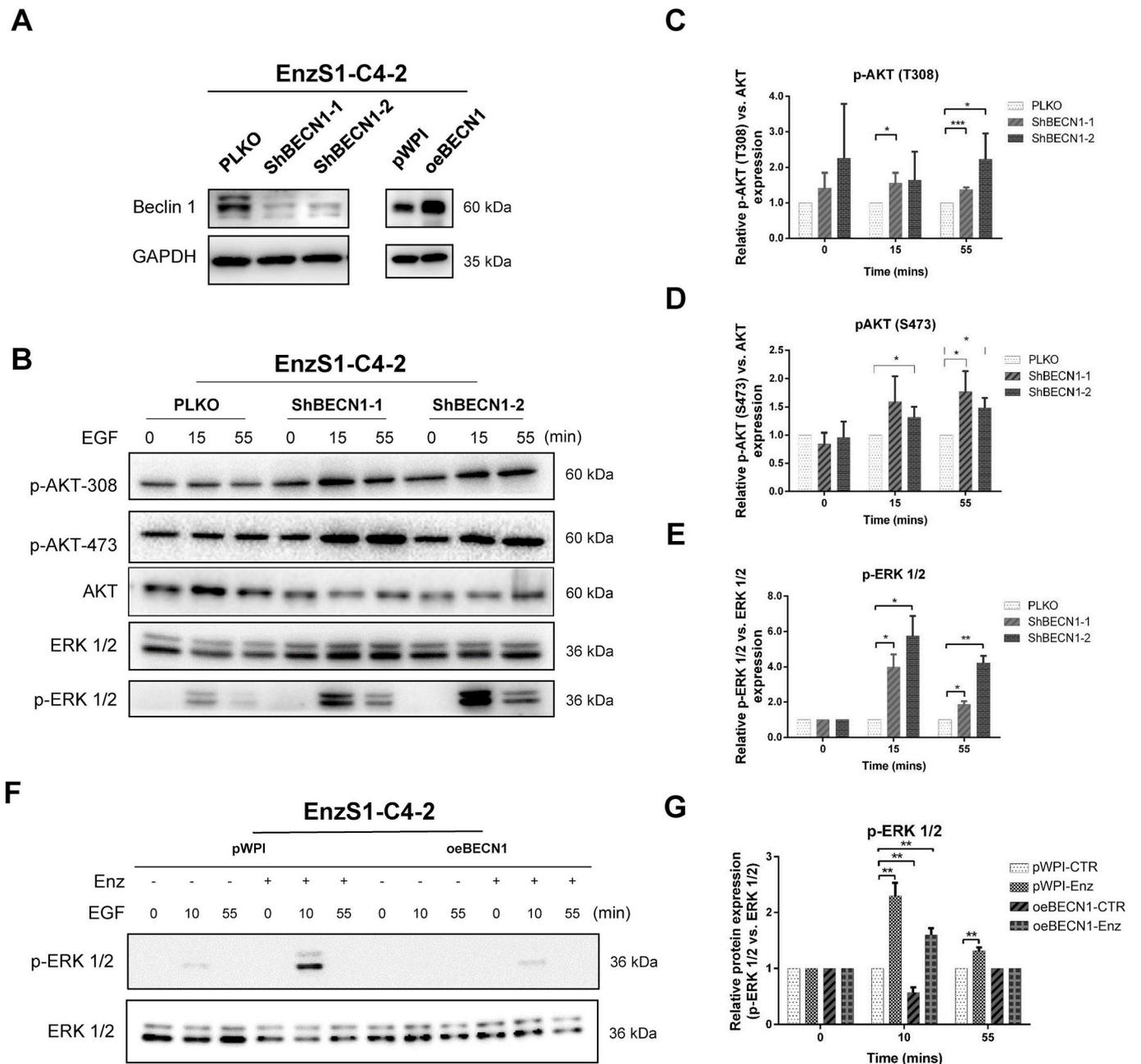


Fig. 2. Beclin 1 regulates ERK-growth factor signaling. A. Western blot of knock down and overexpression *BECN1* efficiencies in EnzS1-C4-2 cells. B. EnzS1-C4-2 cells were infected with *BECN1* shRNAs (ShBECN1-1/2), serum-starved overnight in media containing 0.1% BSA, then stimulated with 50 ng/ml human EGF for the indicated time periods. The p-AKT-S473, p-AKT-T308 and p-ERK1/2 (T202/Y204) levels were detected by western blot. C-E. Quantification data for p-AKT (S473), p-AKT (T308) and p-ERK1/2 (T202/Y204). F. Ectopic *BECN1* expression could reverse Enz enhanced ERK signaling, but not AKT signaling in EnzS1-C4-2 cells (The expression of Beclin 1 in EnzS1-C4-2 cells were slightly knocked down by lentivirus, then treated as the parental cells for viral overexpression of *BECN1*). We pretreated with Enz for three days, and serum-starved cells overnight in media containing 0.1% BSA, then stimulated with 50 ng/ml human EGF for the indicated time periods. G. Quantification data for p-ERK 1/2. The data shown represent the mean \pm SEM of three independent experiments. * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$.

Beclin 1-modulated ERK growth factor signal with a small molecule ERK inhibitor may increase the efficacy of Enz-treatment in PCa cells.

3.3. Mechanism dissection of how Enz can function via altering the AR-Beclin 1 interaction to modulate the ERK activation

Beclin 1 functions in a complex containing Vps34/Class III PI3K (PI3KC3), and Vps15/p150 to regulate the formation of endosome maturation via lipid phosphorylation in the transition of APPL1-containing phosphatidylinositol 3-phosphate-negative (PI3P-) endosomes

to PI3P positive (PI3P+) endosomes [30]. Interestingly, early studies indicated that proteins containing PolyQ, for example, Ataxin-3, might interact with Beclin 1 to promote the autophagy [5], and AR proteins with different PolyQ lengths have been reported to have different transactivation capacities to modulate AR target genes [9,11,45]. Therefore, we hypothesized that Enz might promote AR interaction with Beclin 1 to influence the activity of the Beclin 1 complex to regulate the intracellular ERK-growth factor signal.

We first examined the potential interaction between Beclin 1 and AR via co-immunoprecipitation (Co-IP) assay in the 293T cells, and

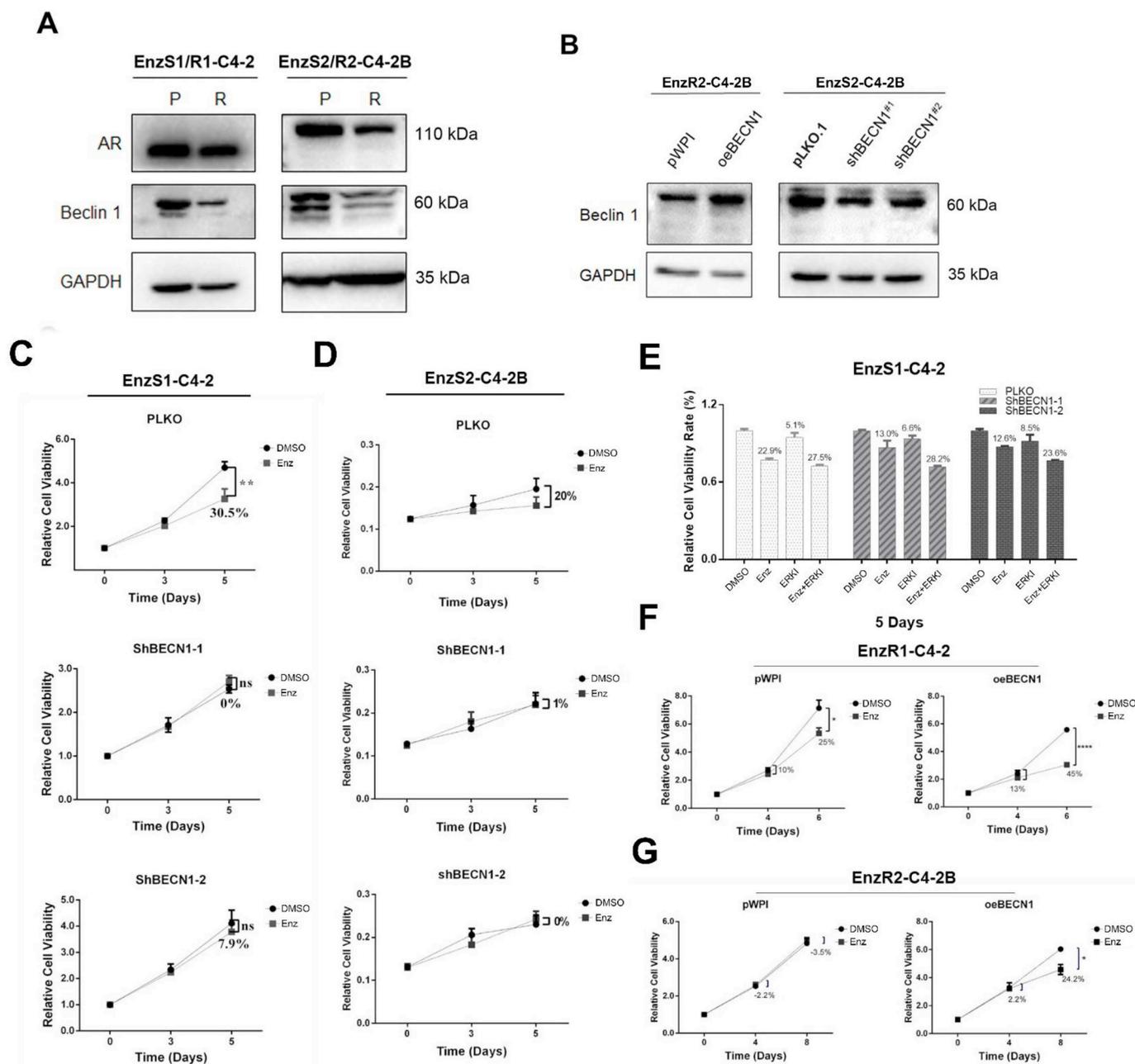


Fig. 3. Beclin 1 regulates Enzalutamide sensitivity. A. Comparing Beclin 1 expression between EnzS1-C4-2 & EnzS2-C4-2B and EnzR1-C4-2 & EnzR2-C4-2B cells (P, parental; R, resistant). B. Knock down and overexpression of BECN1 efficiencies in EnzS2-C4-2B and EnzR2-C4-2B cells, respectively. C-D. MTT assay demonstrated knocking down *BECN1* expression (with 2 Sh-BECN1s) decreased Enz sensitivity in EnzS1-C4-2 (C) and EnzS2-C4-2B (D) cells. E. MTT assay demonstrated that combining Enz with ERK inhibitor treatment could partially reverse ShBECN1 decreased Enz-sensitivity in EnzS1-C4-2 cells. F-G. MTT assay demonstrated ectopic *BECN1* expression in EnzR1-C4-2 (F) and EnzR2-C4-2B (G) cells re-sensitized Enzalutamide resistance.

results revealed that AR could directly interact with Beclin 1 (Fig. 4A). Next, to test the effects of Enz on the AR-Beclin 1 interaction, we treated EnzS1-C4-2 cells with 10 μM Enz and results revealed that Enz treatment increased the interaction between AR and Beclin 1, as well as the formation of the Beclin 1/Vps15/Vps34 complex (Fig. 4B). Consistent with the role of AR, there is no increased Beclin 1/Vps15/Vps34 complex formation in AR-negative cell line Du145 in response to Enz treatment (Fig. 4C). Furthermore, the Co-IP assay using AR antibody also found that AR could interact with Beclin 1 and demonstrably more so with Vps34 in EnzS1-C4-2 in response to Enz (Fig. 4D).

To examine whether Enz-regulated AR-Beclin 1 interaction could lead to the modulation of Beclin 1 complex's function on ERK growth factor signal activation, we applied the ELISA assay to measure the PI3P

generation, as the total cellular PI3P lipid level may represent the degree of growth factor activation [30]. We found that Enz significantly decreased PI3P generation (Fig. 4E), which was consistent with the ELISA data (Fig. 4F) showing a prolonged suppression of PI3P in cells with BECN1 knock down, supporting the conclusion that AR may interact with Beclin 1 complex to negatively impact the latter's activity. Data from the Cancer Genome Atlas (TCGA) database suggested that AR was negatively associated with Beclin 1 protein expression, which was also negatively associated with phosphorylation levels of EGFR (Fig. S4A-B).

To distinguish this AR-Beclin 1 interaction from the regulation of AR/Enz on cellular macroautophagy where Beclin 1 plays a significant role, we knocked down ATG5 expression in EnzS1-C4-2 cells. The

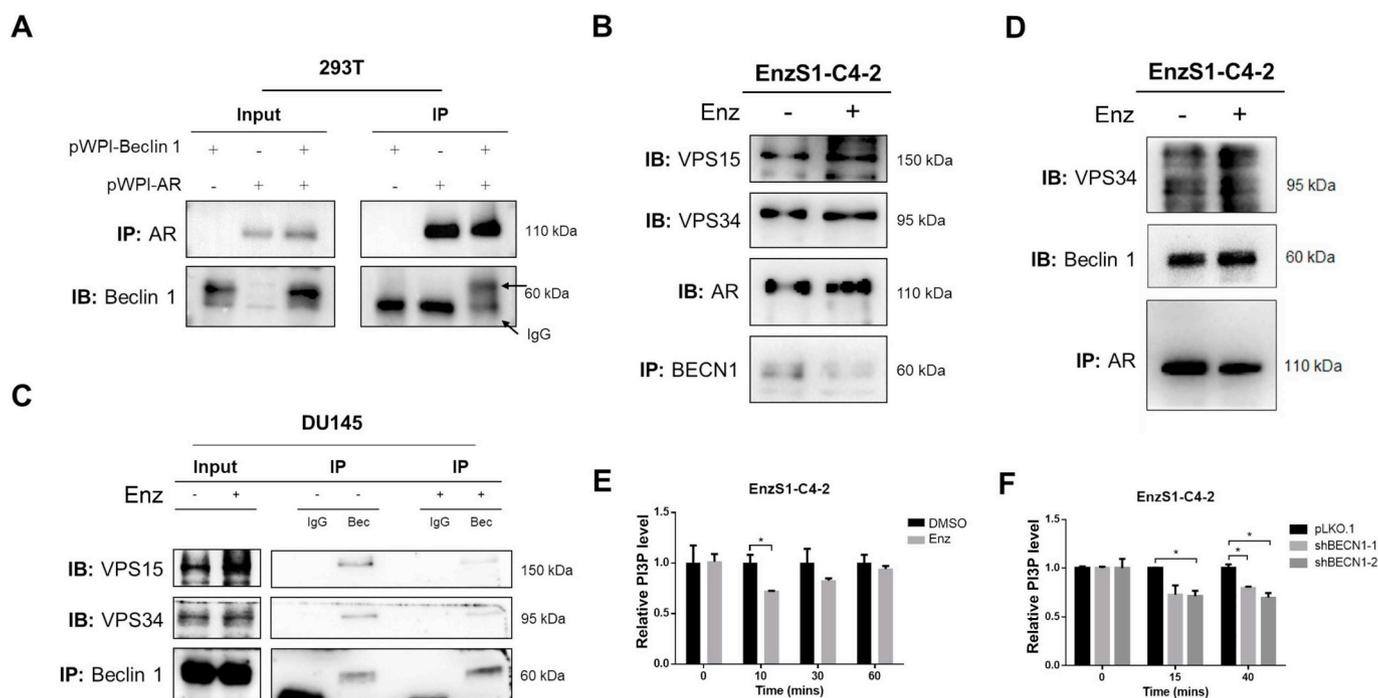


Fig. 4. Mechanism (s) why Enzalutamide can maintain growth factor signaling pathway. A. Exogenous Co-Immunoprecipitation (Co-IP) data proved the interaction between AR and Beclin 1 in 293T cells. B. Endogenous immunoprecipitation (IP) by pull-down of Beclin 1 suggested that Enz increased the interaction between Beclin 1 and AR, as well as the complex formation of Beclin 1/Vps15/Vps34 in EnzS1-C4-2 cells. C. Endogenous IP by pull-down of Beclin 1 suggested that Enz failed to increase the Beclin 1/Vps15/Vps34 complex formation in AR negative Du145 cells. D. Endogenous IP by pull-down AR suggested that AR could interact with Beclin 1 and Vps34 in EnzS1-C4-2 cells. E and F. ELISA data suggested that Enz (E) or ShBECN1 (F) expression could suppress P13P (+) generation in EnzS1-C4-2 cells. For E-F, the data shown represent the mean \pm SEM of three independent experiments. * $p < 0.05$.

results suggested that the early endosome formation involved Beclin 1's function that was independent of its role on autophagy, a result consistent with previous work in breast cancer [30], further suggesting that AR-Beclin 1 interaction for endosome maturation is a distinct regulatory step in response to Enz treatment, separate from the potential regulation of Enz on cellular autophagy (Fig. S5A). Moreover, we also analyzed our unpublished whole genome transcript sequencing work related to Enz treatment as well as other publicly available GEO datasets, and found that genes involved in cellular autophagy were not significantly changed in response to Enz (Fig. S5 B-F).

Together, results from Fig. 4A-G, Fig. S4A-B and Fig. S5A-F suggest that Enz may function via promoting the AR/Beclin 1 interaction to influence the activity of Beclin 1 complex, then positively regulate the growth factor signaling activation. The consequences of such alterations in the AR/Beclin 1 complex-modulated ERK growth factor signaling may lead to compromised Enz sensitivity as illustrated in Fig. 5.

4. Discussion

Several mechanisms have been involved in the development of Enz resistance in CRPC, including induction of altered glucocorticoid receptor [4], AKT signal [24,36,37], AR-v7 [2] and ARF876L mutation [15,19,41]. Although AR-v7 expression has the strongest clinical supports showing CRPC patients with detectable AR-v7 in circulating tumor cells had poorer responses to ADT-Enzalutamide [3], there are still many patients, who are resistant to Enz, but were found to be AR-v7-negative.

Previous publications indicated the growth factor signals, such as PI3K/AKT pathway, play critical roles during Enz-resistance development and serve as key potential therapeutic targets for CRPC patients [3,8,17]. In this study, we found that Enz could induce and maintain the phosphorylation of ERK in response to growth factor stimulation, which likely contributed to the development of Enz-resistance.

Mechanism dissection indicated Enz could enhance the interaction between AR and Beclin 1 complex, thus decreasing Beclin 1 complex's activity in mediating ERK-growth factor signaling inactivation. This post-translational mechanism was different from a transcriptional regulation by AR of the PHLPP phosphatase that inactivates AKT by removing activating phosphorylation [8]. Indeed, this difference might also underlie the different response towards the rescue by Beclin 1 overexpression, which appeared only for ERK signal, but not for AKT signal (data not shown). These findings were consistent with a similar mechanistic role of Beclin 1 in enhancing growth factor signaling in breast cancer [30]. Moreover, these results were consistent with our finding that Beclin 1 was expressed lower in both EnzR1-C4-2 and EnzR2-C4-2B cells compared with their parental cells while knocking down *BECN1* expression significantly decreased Enz sensitivity by increasing and maintaining ERK growth factor signal. Therefore, increasing Beclin 1 expression, or targeting its downstream ERK growth factor signal with small molecule ERK inhibitor, may reduce Enz-resistance in PCa cells and enhance its efficacy.

Beclin 1 is the mammalian homolog of the yeast Atg6/Vacuolar protein sorting 30 (Vps30) protein that plays an essential role in macroautophagy and vacuolar protein sorting [16,33]. Beclin 1 is required for normal mammalian development and it has been shown to play a critical role in pathogenesis, including cancers [29,46]. Reduction of Beclin 1 was observed in many tumors [13,40], including PCa [21]. In addition, there is reciprocal regulation between Beclin 1 and growth factor receptors such that posttranslational modifications of Beclin 1 result in its degradation while its loss can impact growth factor trafficking and signaling. Wei et al. [44] found that EGFR activation promotes the receptor binding to Beclin 1, leading to its multisite tyrosine phosphorylation, and decreased Beclin 1-associated Vps34 lipid kinase activity. Hyperactivation of growth factor signaling pathways in cancers likely enhance their oncogenic signaling potential through regulating Beclin 1-mediated regulation of receptor trafficking and

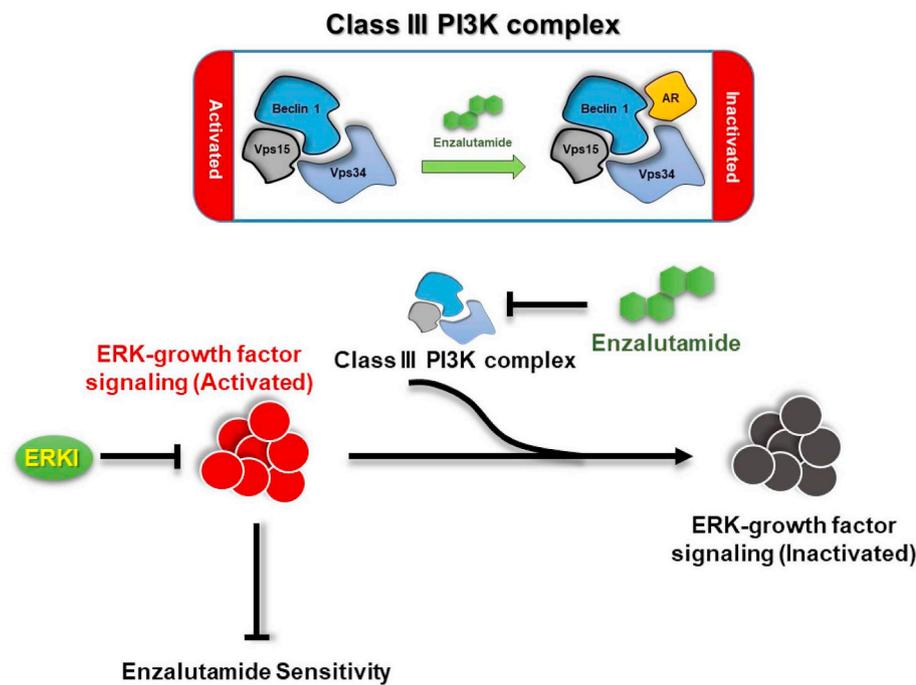


Fig. 5. Hypothesis. Enzalutamide promotes AR interacting with Beclin 1 complex, to inactivate its activity in mediating growth factor ERK signaling transit via decreasing PI3P generation. In the upper panel, Enzalutamide could increase AR interact with Beclin 1 complex. In the lower panel Beclin 1 complex could transfer growth factor from activation to inactivation status. Because Enzalutamide suppresses Beclin 1 complex's activity, therefore, the ERK growth factor signal could be continually activated, and cells become more “resistant”, while ERKI combined with Enzalutamide could re-sensitize cells to Enzalutamide treatment. ERKI: ERK Inhibitor.

duration of signal activation. In our work, we found that AR/Beclin1 complex interaction inactivated the complex's activity in mediating PI3P (+) generation, and therefore, suppressed ERK growth factor transit, consequently contributing to Enz sensitivity decreasing.

Our study reveals an alternative mechanism for how Beclin 1 loss, in addition to its function in regulating autophagy, may impact PCa cell Enz-resistance that involves enhancing the magnitude and duration of ERK growth factor signal. This study also suggests that targeting Beclin 1 or downstream ERK growth factor signal by an ERK inhibitor could effectively suppress Enz resistant CRPC cell growth, serving as potential therapeutic targets for PCa patients.

Conflicts of interest

The authors declare that they have no conflicts of interest.

Acknowledgments

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Appendix A. Supplementary data

Supplementary data to this article can be found online at <https://doi.org/10.1016/j.canlet.2018.11.008>.

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