

Supplement

NLRX1 Drives Prostate Cancer Progression Through Activation of AKT and ERK Signaling Pathways

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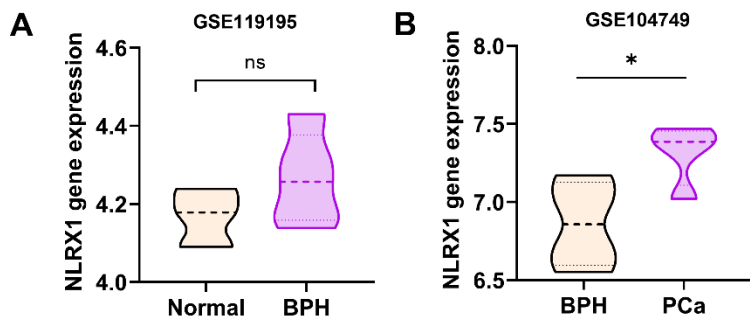
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14 Submission to: **International Journal of Biological Sciences**

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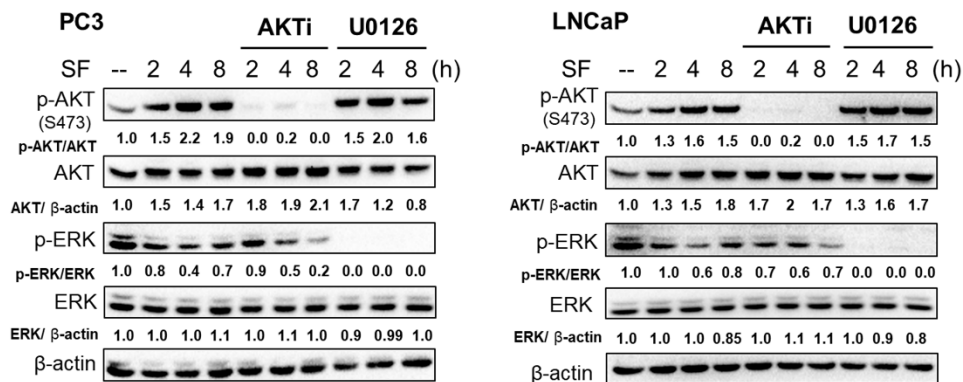
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20 **Fig. S1. NLRX1 expression in prostate tissues and clinical subgroups.**

21 (A) NLRX1 mRNA expression in normal prostate tissues and benign prostatic hyperplasia
 22 (BPH) samples derived from the GSE119195 GEO dataset. Differences between groups
 23 were evaluated using Student's t-test. (B) Comparison of NLRX1 expression between
 24 BPH and PCa tissues from a GSE104749 GEO dataset, with statistical significance
 25 assessed by Student's t-test.

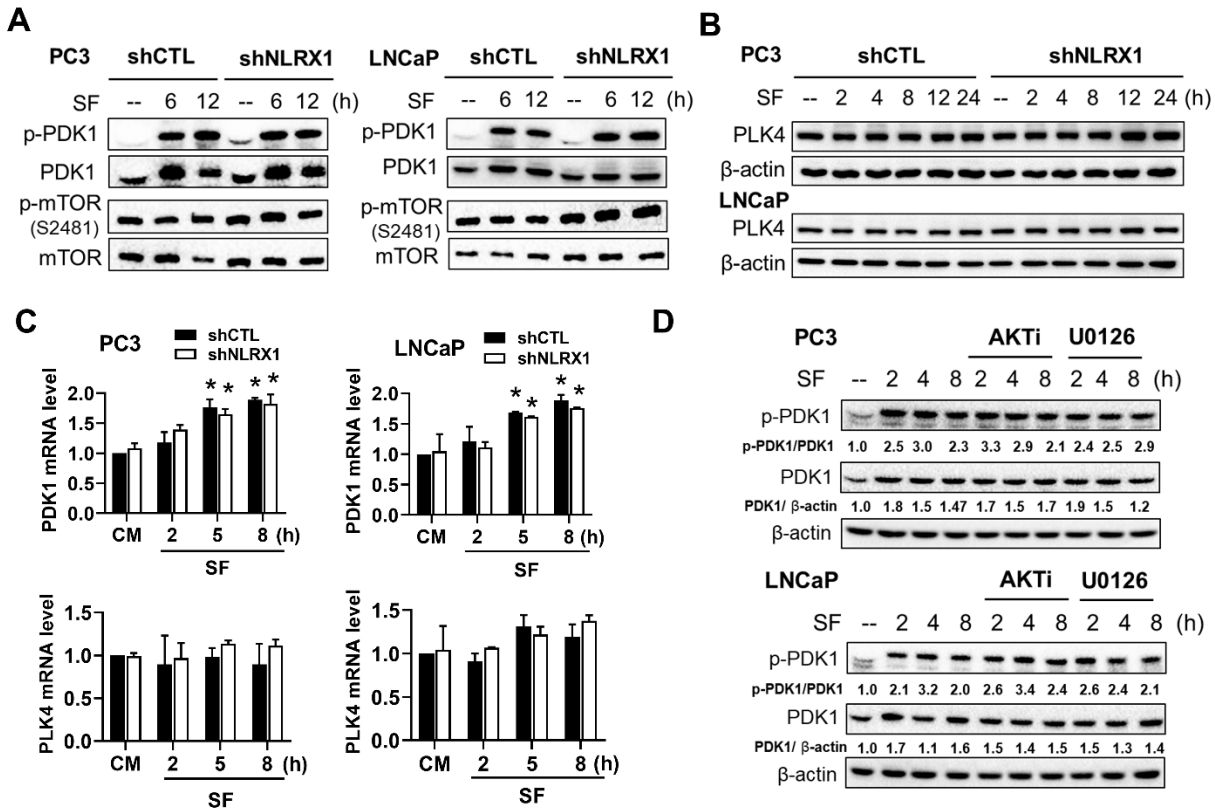


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27 **Fig. S2. AKT and ERK signaling dynamics following AKTi and U0126 treatment in**
 28 **PC3 and LNCaP cells.**

29 Cells were treated with AKTi or U0126 under SF conditions. p-AKT, total AKT, p-ERK,
 30 and total ERK were analyzed; β-actin served as a loading control. Densitometric analysis
 31 of Western blot bands was performed using ImageJ. Protein expression was normalized
 32 to β-actin, and values were expressed relative to the 0 h control (set to 1).

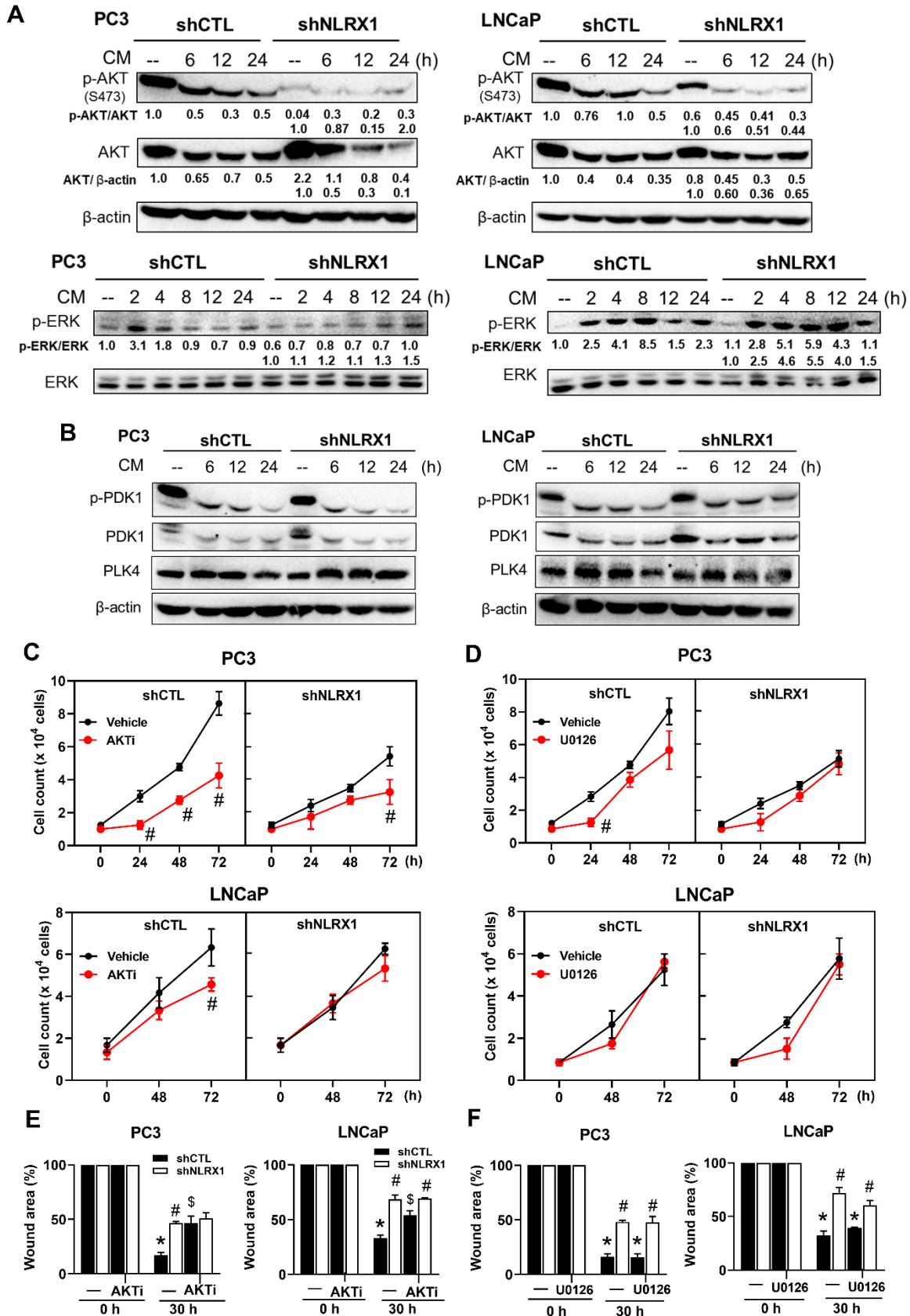
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35 **Fig. S3. NLRX1 silencing does not affect PDK1/mTOR/PLK4 signaling under serum-**
 36 **free conditions.**

37 (A, B) PC3 and LNCaP cells were cultured under serum-free (SF) conditions for the
 38 indicated times. Immunoblotting was performed to assess p-PDK1, p-mTOR, total mTOR,
 39 and PDK1 (A) and PLK4 (B). (C) PDK1 and PLK4 mRNA levels under SF were measured
 40 by qRT-PCR. (D) Cells were treated with AKTi or U0126 under SF for the indicated times.
 41 p-PDK1, total PDK1, and β-actin were analyzed by immunoblotting. Data are mean ±
 42 S.E.M. from independent experiments. *p < 0.05 indicates a significant effect of SF
 43 compared to the untreated control group.



45 **Fig. S4. NLRX1 silencing inhibits cell growth and migration via AKT suppression.**

46 (A, B) Cells were serum-starved (24 h), then stimulated with CM for the indicated times.
47 p-AKT (S473), total AKT, p-ERK, total ERK, and β -actin were analyzed by
48 immunoblotting. In (A), both quantification of p-AKT/AKT and AKT/ β -actin were
49 determined, and values were expressed relative to the 0 h control (set to 1). Moreover,
50 the quantification based on the 0 h response in shNLRX1 cells (set to 1) was also
51 determined. (C, D) Cells were serum-starved overnight, then co-treated with complete
52 medium (CM) and either AKTi (B) or ERKi (U0126) (C). Cell number was determined
53 using trypan blue exclusion. (E, F) Wound healing assays were performed in PC3 and
54 LNCaP cells pre-treated with mitomycin C (1 μ g/mL), followed by AKTi (E) or ERKi (F)
55 treatment. Wound closure was monitored microscopically. Data are mean \pm S.E.M. from
56 independent experiments. *, $p < 0.05$ indicates a significant effect on the control group
57 (shCTL) compared to the untreated shCTL. #, $p < 0.05$ indicates a significant effect of
58 shNLRX1 compared to shCTL. \$, $p < 0.05$ indicates a significant effect of the inhibitor
59 treatment under SF conditions.

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