

**Phosphorylation of 14-3-3ζ links YAP transactivation to hypoxic glycolysis for tumorigenesis**

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**Supplemental Information**

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**Fig. S1 Hypoxia induces nuclear translocation of YAP in a HIF-1 $\alpha$ -independent manner.** **a** Subcellular fractionation analyses determining abundance of nuclear and cytoplasmic YAP protein in A498 RCC cells (A-4) and Huh-7 HCC cells (H-7) stimulated with hypoxia for 6h. GAPDH and Lamin B were used as internal control of cytoplasmic and nuclear extractions, respectively. Data are expressed as mean  $\pm$  s.d. of three independent experiments. \* $P$  < 0.05, \*\* $P$  < 0.01. Two-sided Student's t test was used to calculate the  $P$  value. **b** Subcellular fractionation analyses comparing levels of nuclear YAP protein expression in SW-1990 PDAC cells stimulated with the indicated concentrations of CoCl<sub>2</sub> for 12h. **c** Subcellular fractionation analyses comparing accumulation of nuclear and cytoplasmic YAP protein in hypoxia-stimulated SW-1990 PDAC cells with or without HIF-1 $\alpha$  siRNA transfection. **d** Coimmunoprecipitation assay evaluating the interaction between Myc-tagged wild-type TEAD4 and nuclear YAP in SW-1990 PDAC cells stimulated with hypoxia for 12h.

**Fig. S2 Hypoxia induces nuclear translocation of YAP in a 14-3-3 $\zeta$ -dependent fashion.** **a** Subcellular fractionation analyses assessing abundance of nuclear and cytoplasmic YAP protein expression in SW-1990 PDAC cells with or without 14-3-3 $\zeta$  siRNA transfection. Data are expressed as mean  $\pm$  s.d. of three independent experiments. \* $P$  < 0.05. Two-sided Student's t test was used to calculate the  $P$  value. **b** Left panel: Western-blotting comparing the levels of Flag expression in SW-1990 PDAC cells with or without Flag-tagged wild-type 14-3-3 $\zeta$  transfection. Right panel: Coimmunoprecipitation assay evaluating the interaction between Myc-TEAD4 and nuclear YAP in SW-1990 PDAC cells stimulated with hypoxia for 12h with or without Flag-tagged wild-type 14-3-3 $\zeta$  transfection. IP, immunoprecipitation. **c** Coimmunoprecipitation assay comparing the interaction between 14-3-3 $\zeta$  and YAP in SW-1990 PDAC cells treated with or without 1 mmol/L CoCl<sub>2</sub> for 12h. Data are expressed as mean  $\pm$  s.d. of three independent experiments. \* $P$  < 0.05. Two-sided ANOVA with Bonferroni post hoc t test correction was used to calculate the  $P$  value. **d** Coimmunoprecipitation assay assessing the interaction between endogenous WWTR1 and YAP in SW-1990 PDAC cells stimulated with or without hypoxia for 12h.

**Fig. S3 ERK1/2 is required for the hypoxia-stimulated YAP nuclear localization.** **a** Western-blotting comparing the levels of p-I $\kappa$ B $\alpha$  and p-ERK1/2 in SW-1990 PDAC cells stimulated with hypoxia in the presence or absence of 5  $\mu$ mol/L BAY 11-7085 (BAY) and 20  $\mu$ mol/L U0126 pretreatment, respectively. **b** Left panel: Western-blotting testing the levels of ERK2 expression in SW-1990 PDAC cells transfected with ERK2 siRNA. Right panel: Coimmunoprecipitation assay determining the interaction between Myc-tagged wild-type TEAD4 and nuclear YAP in SW-1990 PDAC cells with or without serum stimuli (SS) in the presence or absence of ERK2 siRNA transfection. **c** Alignment of the highly conserved Ser37 residue of ERK2 phosphorylation in 14-3-3 $\zeta$  from homo sapiens to drosophila.

**Fig. S4 Hypoxia phosphorylates 14-3-3 $\zeta$  at Ser37 via ERK2.** **a** Coimmunoprecipitation assay detecting the levels of Flag-tagged wild-type 14-3-3 $\zeta$  phosphorylation in hypoxia-stimulated SW-1990 PDAC cells in the presence or absence of 5  $\mu$ mol/L BAY 11-7085 (BAY) and 20  $\mu$ mol/L U0126 administration with an anti-phospho-serine antibody, respectively. **b** Sequencing analysis of Ser37 mutations in 14-3-3 $\zeta$  protein. **c** *Top panel:* Co-IP assay testing the specificity of 14-3-3 $\zeta$

pS37 antibody. *Bottom panel*: Representative immunofluorescence images testing the specificity of 14-3-3 $\zeta$  pS37 antibody in SW-1990 PDAC cells with Flag-tagged wild-type 14-3-3 $\zeta$  (WT) or mutant 14-3-3 $\zeta$  Ser37A (S37A) expression. Scale bar = 25  $\mu$ m. **d** RT-qPCR analyses of YWHAZ gene expression in 14-3-3 $\zeta$  shRNA (sh. 14-3-3 $\zeta$ )-expressed SW-1990 PDAC cells. Experiments were performed five times, each with quantitative RT-PCR in technical duplicate and real-time values were normalized to glyceraldehyde 3-phosphate dehydrogenase (GAPDH). Data are expressed as mean  $\pm$  s.d. \*\*\* $P$  < 0.001. Two-sided Student's t test was used to calculate the  $P$  value.

**Fig. S5 YAP facilitates PKM2 transcription via physical interaction with HIF-1 $\alpha$  in an 14-3-3 $\zeta$  Ser37 phosphorylation-dependent manner under hypoxia.** **a** Coimmunoprecipitation assay examining the interaction between nuclear YAP and HIF-1 $\alpha$  in hypoxia-stimulated SW-1990 PDAC cells with or without ERK2 siRNA transfection. **b** Coimmunoprecipitation assay determining the interaction between nuclear YAP and HIF-1 $\alpha$  in 14-3-3 $\zeta$  shRNA (sh.14-3-3 $\zeta$ )-expressed SW-1990 PDAC cells in the presence or absence of Flag-tagged wild-type 14-3-3 $\zeta$  (WT) or mutant 14-3-3 $\zeta$  S37E (S37E) reconstitution. **c** Top panel: Western blotting results depicting the expression levels of YAP with scrambled shRNA (Scr) or YAP shRNA transfection. Bottom panel: CHIP analysis for HIF-1 $\alpha$  and HIF-2 $\alpha$  binding to PKM2 gene HRE in SW-1990 PDAC cells stimulated with hypoxia in the presence or absence of YAP shRNA transfection using the indicated antibodies. **d** Top panel: Western blotting results depicting the expression levels of HIF-1 $\alpha$  with control siRNA (Ctrl) or HIF-1 $\alpha$  siRNA transfection. Bottom panel: CHIP analysis for YAP binding to PKM2 gene HRE in SW-1990 PDAC cells stimulated with hypoxia in the presence or absence of HIF-1 $\alpha$  siRNA transfection using an anti-YAP antibody. **e** RT-qPCR analyses of PKM2 gene expression in YAP shRNA-expressed SW-1990 PDAC cells with hypoxia stimulation. Experiments were performed five times, each with quantitative RT-PCR in technical duplicate and real-time values were normalized to glyceraldehyde 3-phosphate dehydrogenase (GAPDH). Data are expressed as mean  $\pm$  s.d. \* $P$  < 0.05. Two-sided Student's t test was used to calculate the  $P$  value. **f** Western-blotting measuring levels of PKM2 protein expression in YAP shRNA-expressed SW-1990 PDAC cells with hypoxia stimulation. **g** Representative immunofluorescence images of Flag-tagged 14-3-3 $\zeta$  staining in 14-3-3 $\zeta$  shRNA (sh.14-3-3 $\zeta$ )-expressed SW-1990 PDAC cells with Flag-tagged wild-type 14-3-3 $\zeta$  (WT) or mutant 14-3-3 $\zeta$  Ser37A (S37A) reconstitution. Scale bar = 100  $\mu$ m. **h** Immunohistochemistry analysis of 14-3-3 $\zeta$  p-S37 antibody with or without 14-3-3 $\zeta$  peptide used for developing this antibody or the matched nonphosphorylated 14-3-3 $\zeta$  peptide. Scale bar = 50  $\mu$ m. **i** Representative cases stained by immunohistochemistry (*left panel*) and bar graph (*right panel*) showing the expression of 14-3-3 $\zeta$  p-S37 in 56 primary human PDAC specimens is positively correlated with nuclear localization of YAP. Scale bar = 50  $\mu$ m. The  $P$  value shown was calculated by Spearman order correlations.

**Table S1 Clinicopathological Features in PDAC Patients (n=87)**

<b>Variable</b>	<b>No. of patients</b>	<b>%</b>
<b>Gender</b>		
Male	50	57.5
Female	37	42.5
<b>Age (years)</b>		
<60	33	37.9
≥60	54	62.1
<b>Tumor size</b>		
≤40mm	56	64.4
>40mm	31	35.6
<b>Tumor location</b>		
Head	71	81.6
Body	9	10.3
Tail	7	8.1
<b>Lymphatic metastasis</b>		
Negative	49	56.3
Positive	38	43.7
<b>TNM stage</b>		
I	22	25.3
II	43	49.4
III	14	16.1
IV	8	9.2
<b>Chemotherapy</b>		
Yes	72	82.8
No	15	17.2
<b>p-14-3-3ζ expression</b>		
High	45	51.7
Low	42	48.3

**Table S2 Correlation Between Phosphorylation of 14-3-3 $\zeta$  and Clinicopathological Features in PDAC Patients**

Variable	p-14-3-3 $\zeta$		P value
	High (n = 45)	Low (n = 42)	
Gender			0.313
Male	27 (60.0%)	23 (54.8%)	
Female	18 (40.0%)	19 (45.2%)	
Age (years)			<b>0.015</b>
<60	22 (48.9%)	11 (26.2%)	
$\geq$ 60	23 (51.1%)	31 (73.8%)	
Tumor size			<b>&lt; 0.001</b>
$\leq$ 40mm	20 (44.4%)	36 (85.7%)	
>40mm	25 (55.6%)	6 (14.3%)	
Tumor location			0.103
Head	37 (82.2%)	34 (81.0%)	
Body	6 (13.3%)	3 (7.1%)	
Tail	2 (4.5%)	5 (11.9%)	
Lymphatic metastasis			<b>0.031</b>
Negative	21 (46.7%)	28 (66.7%)	
Positive	24 (53.3%)	14 (33.3%)	
TNM stage			0.200
I	14 (31.1%)	8 (19.0%)	
II	17 (37.8%)	26 (61.9%)	
III	8 (17.8%)	6 (14.3%)	
IV	6 (13.3%)	2 (4.8%)	
Chemotherapy			0.104
Yes	35 (77.8%)	37 (88.1%)	
No	10 (22.2%)	5 (11.9%)	