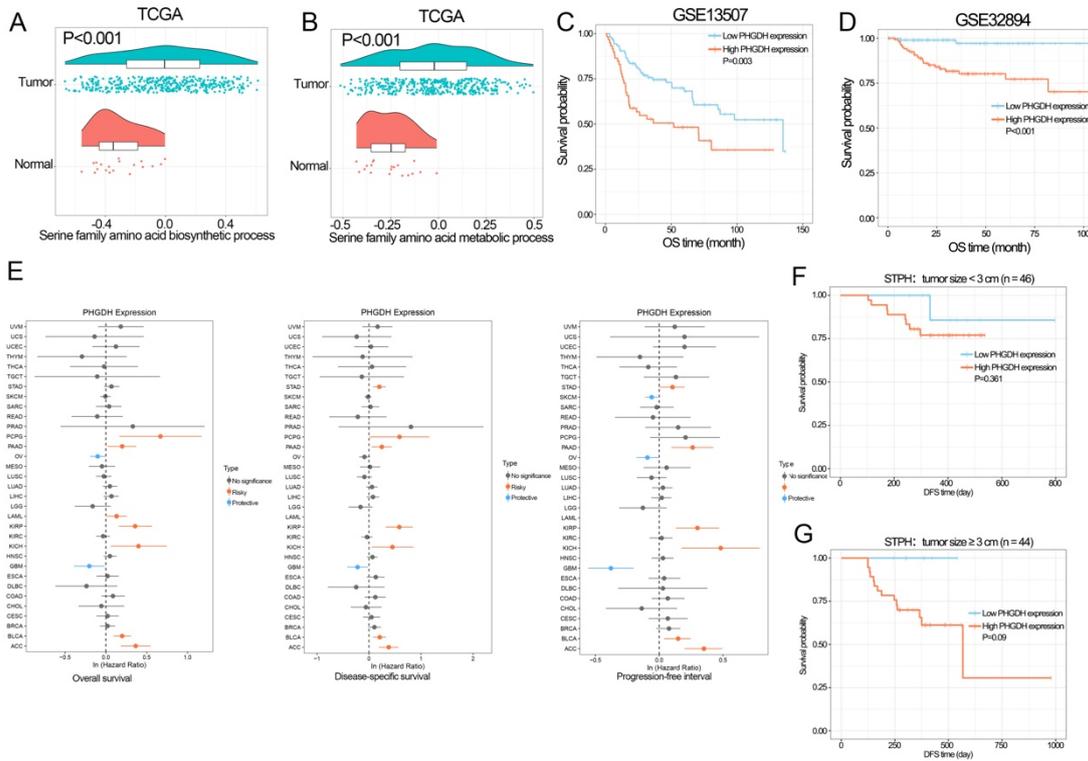
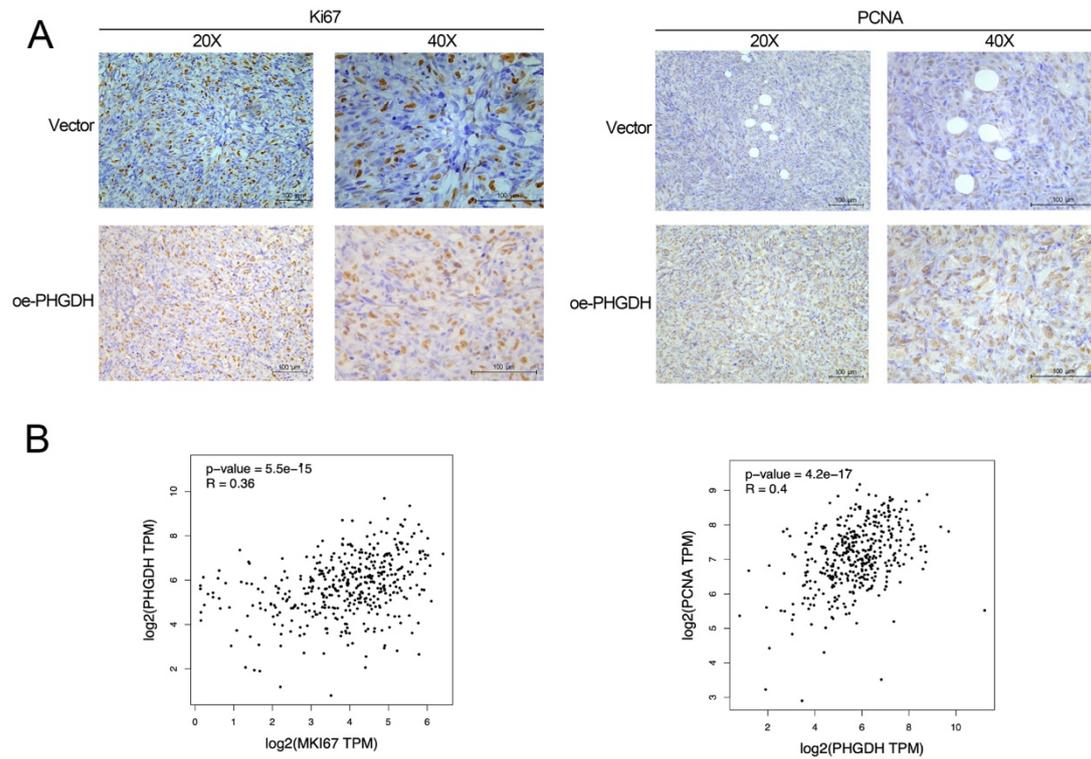


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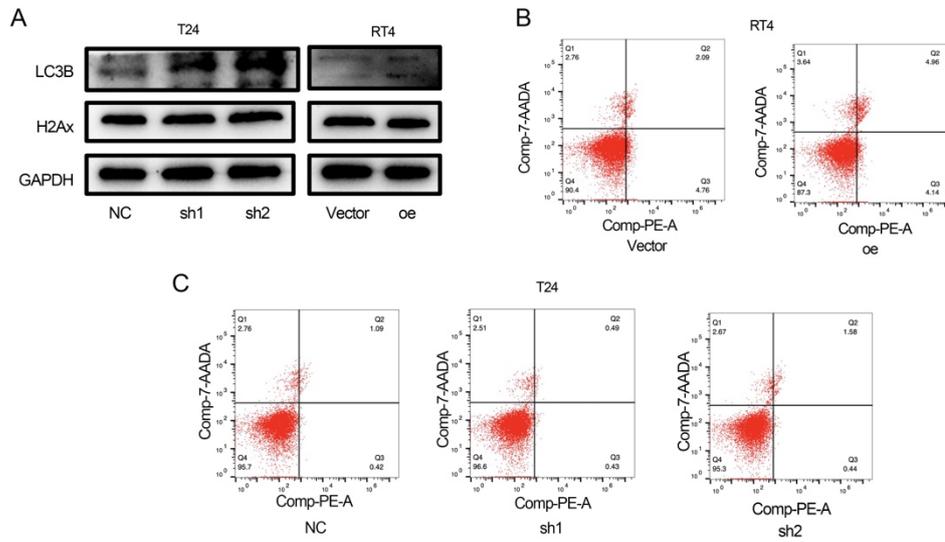


2 **Figure S1:** (A) The activities of serine family amino acid biosynthetic processes
3 between normal and BCa cancer in TCGA. (B) The activities of serine family amino
4 acid metabolic processes between normal and BCa cancer in TCGA. (C) Kaplan-
5 Meier plots of PHGDH in GSE13507. (D) Kaplan-Meier plots of PHGDH in
6 GSE32894. (E) Risk plot of correlation between PHGDH with overall survival,
7 disease-specific survival, and progression-free interval. (F) Kaplan-Meier plots of
8 PHGDH in BCa patient patients with tumor size < 3 cm in STPH. (G) Kaplan-Meier
9 plots of PHGDH in BCa patient patients with tumor size ≥ 3 cm in STPH.

10



12 **Figure S2** (A) Representative IHC images of Ki67 and PCNA in mouse subcutaneous
 13 tumor tissue (RT4 cell line), Scale bars: 50/25 μ m. (B) PHGDH and Ki67/PCNA were
 14 significantly correlated based on the TCGA database.



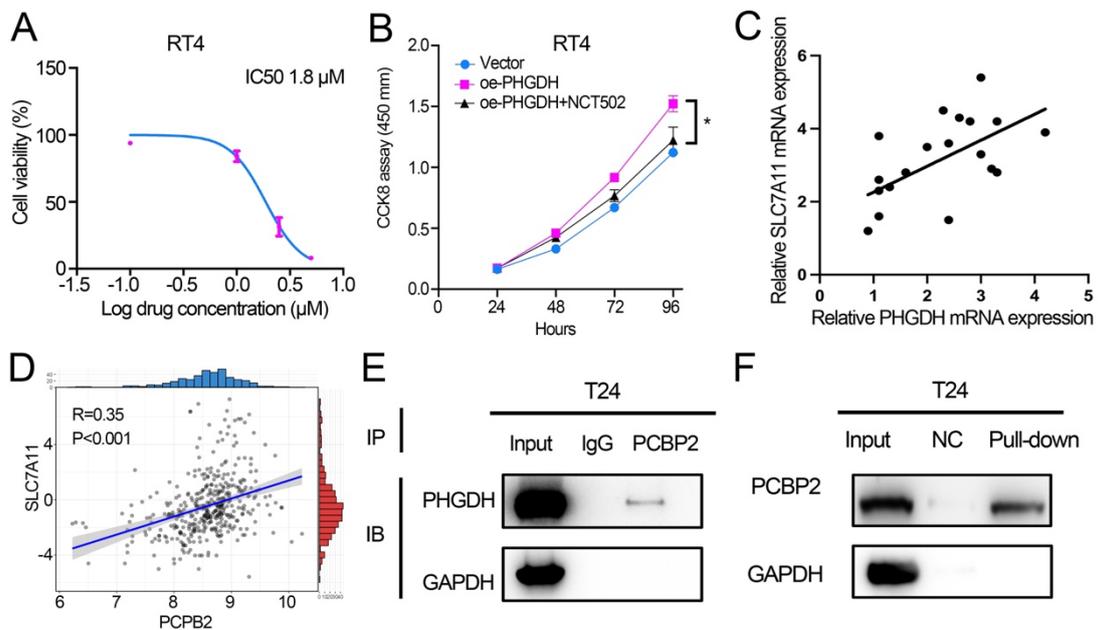
15

16 **Figure S3 (A)** Western blot was used to detect changes in autophagy-related proteins

17 after knockdown or overexpression of PHGDH. (B-C) Flow cytometry was used to

18 detect apoptosis after knockdown or overexpression of PHGDH.

19



G

The RBP-mRNA Interactions Supported by CLIP-seq Data

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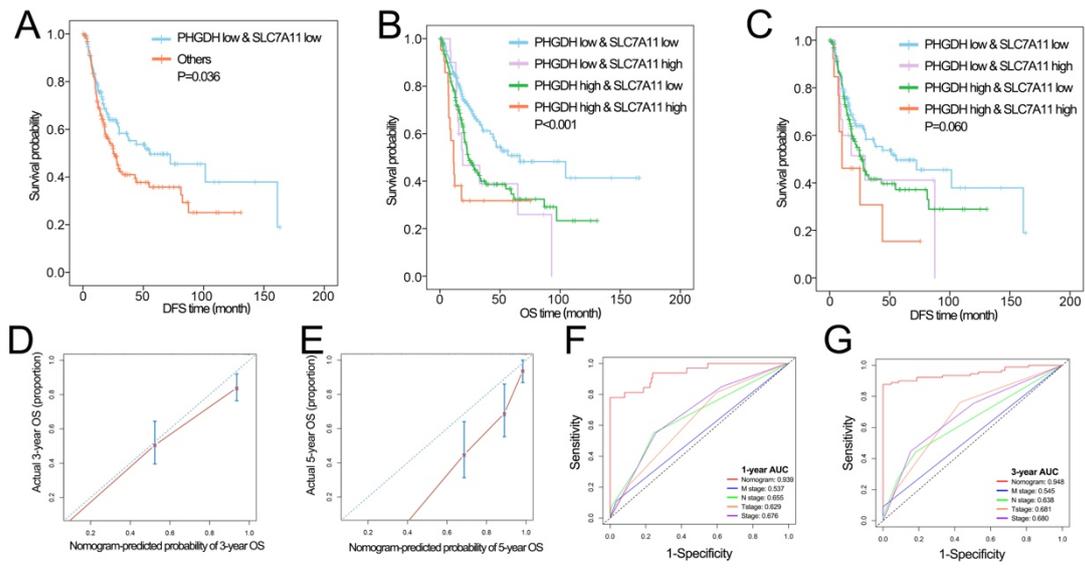
RBP	GeneID	GeneName	GeneType	ClusterNum	ClipExpNum	ClipSiteNum	HepG2(log2FC)	K562(log2FC)	Pan-Cancer
PCBP2	ENSG000001012	SLC7A11	protein_coding	14	2	16	1.843	-0.977	16

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20 **Figure S4** (A) IC50 values of NCT-502 in RT4 cell line. (B) CCK8 assays were used
 21 to validate the viability of NCT-502 in the RT4 cell line. (C) qPCR was used to detect
 22 the association of PHGDH with SLC7A11 in 20 bladder cancer patients. (D)The
 23 association of PHGDH with SLC7A11 in the TCGA database. (E) CO-IP assay was
 24 used to verify the binding of PHGDH to PCBP2 (F) RNA pull-down was used to
 25 verify the binding of PCBP2 to SLC7A11 mRNA. (G) Starbase database for
 26 prediction of PCBP2 binding to SLC7A11 mRNA.

27
 28



29 **Figure S5** (A-C) Kaplan-Meier survival analysis of patients' OS and DFS according
 30 to PHGDH+SLC7A11 score. (D-E) Calibration plots of the nomogram for predicting
 31 the probability of 3- and 5-year OS. (F-G) ROC curves by nomogram compare to
 32 TNM stage for 1-(F) and 3-year OS (G).

33

34 **Table S1** Clinicopathological characteristics of patients from Shanghai Tenth People's

35 Hospital

36

37 **Table S2** List of shRNA sequence; primer and antibodies

38

39 **Table S3** Data for mass spectrometry

40

41