

## **Supplementary Information for**

AR-negative prostate cancer is vulnerable to loss of JMJD1C demethylase

Yohei Yoshihama, Kyle A. LaBella, Eiru Kim, Lori Bertolet, Medina Colic, Jiexi Li, Xiaoying Shang, Chang-Jiun Wu, Denise J. Spring, Y. Alan Wang, Traver Hart, Ronald A. DePinho

Correspondence and communicated by Ronald A. DePinho and Traver Hart Email: <a href="mailto:rdepinho@mdanderson.org">rdepinho@mdanderson.org</a>, <a href="mailto:traver@hart-lab.org">traver@hart-lab.org</a>

## This PDF file includes:

Figures S1 to S4

## Other supplementary materials for this manuscript include the following:

Tables S1 and S2



Fig. S1. **Knockdown efficiency of JMJD1C in growth assays.** (*A* and *B*) Knockdown efficiency of JMJD1C mRNA in PC3 (A) and LNCaP (B). (C) Establishment of TP53 knockout LNCaP. (*D*) Knockdown efficiency of JMJD1C mRNA in LNCaP<sup>APIPC</sup>. (*E*) Knockdown efficiency of JMJD1C in MSKPCa1 prostate cancer organoid.







Fig. S3. GSEA analysis for JMJD1C depleted AR-positive LNCaP cells. (A) GSEA

analysis (Hallmark) showing enriched signatures by JMJD1C knockdown in LNCaP

(doxycycline treatment 7 days). Signatures with P < 0.05 and FDR q-value <0.25 are shown.

(B) Enrichment plot showing Hallmark TNF $\alpha$  signaling via NF $\kappa$ B by JMJD1C knockdown in

LNCaP.



Fig. S4. **Correlation of JMJD1C signature and TNF** $\alpha$  **signature.** (*A*) The Venn diagram showing commonly downregulated genes by JMJD1C-depletion in PC3 and AR negative LNCaP cells (LNCaP<sup>APIPC</sup>). (*B* and *C*) TNF $\alpha$  signature calculated with GSVA for prostate cancer patients with high (> 0) or low (< 0) AR or JMJD1C signature in TCGA (*B*) and MCTP (*C*) data set.

Table S1 (separate file). List of BF and fraction essentiality plotted in Figure 1 A.

Table S2 (separate file). List of genes downregulated by JMJD1C depletion in PC3 and LNCaP<sup>APIPC</sup> cells.