Supplementary Materials: CRISPR-mediated reactivation of DKK3 expression attenuates TGF-signaling in prostate cancer

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Figure S1. Methodological workflow of analyses performed

| Gene symbol | Oncomine gene expression | MSKCC 2010 Gene Expression (cBIO) |
|-------------|--------------------------|-----------------------------------|
| CDKN2A | 2 down | |
| HFE | 6 down | |
| PDX1 | | |
| PTGS2 | 5 down | |
| MSX1 | | |
| ANXA2 | 12 down | |
| NKX2-1 | 3 down | |
| EFEMP1 | 1 down | |
| PAX5 | | |
| SOX17 | | |
| LMX1B | | |
| PROM1 | 8 down | |
| SIX6 | 1 down | |
| CRHBP | 2 down | |
| HES5 | 1 down | |
| TBX15 | | |
| NKX2-2 | | |
| HIST1H4F | | |
| UCN | 1 down | |
| VSX1 | 2 down | |
| TNFRSF10C | 5 down | |
| VAX1 | | |
| CDX4 | 1 down | |
| CYP27A1 | 8 down | |
| RND2 | 5 down | |
| EN2 | 1 down | |
| HOXC4 | | |
| DAB1 | 3 down | |
| WNT16 | | |
| NKX2-3 | | |
| POU3F3 | | |
| LHX9 | | |
| ZNF154 | 2 down | |
| GSX2 | | |
| HS3ST1 | 1 down | |
| TMEM176B | 1 down | |
| ОТР | 2 down | |
| GPR50 | 1 down | |
| AKAP2 | 1 down | |
| RHCG | | |
| LAMP5 | | |
| SPATA6 | 3 down | |
| | 0 down | |
| SLC16A5 | 9 down | |
| | | |
| | 1 down (C1 or f114) | |
| TMEM106A | 2 down | |
| WDR86 | | |
| 110100 | | |

Figure S2. Analysis of the expression levels of the indicated genes in PCa datasets. Bars on right indicate upregulation (red), no change (grey) and downregulation (blue) in tumors.

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Figure S3. Location of CpG probes in the DKK3 promoter



Figure S4. Effects of TGF-b and CRISPR induction of DKK3 on *DNMT* mRNA expression. Q-RT-PCR analysis of the indicated genes in PC3 cells, either untreated and treated with TGF-b for 24 h (A) or transfected with dCas9-VPR and either control gRNA or DKK3 gRNAs (B). Graphs show mRNA expression fold change ± SD, versus control treatment, n = 3, *p < 0.05 two-tailed Student's t-test.

| Table S1. Sequences and genomic coordinates of gRNAs used to target the DKK3 promoter | | | | |
|---|---------------|---------------------|--|--|
| gRNA | gRNA sequence | Genomic Coordinates | | |
| | | | | |

| gRNA | gRNA sequence | Genomic Coordinates |
|---------|-----------------------|---------------------------|
| 1 | GTGGCGGTAAACAGTAATGTG | chr11:12030012-12030032 |
| 2 | GTCTGCCCGAAGTGACAAGCG | chr11:12030132-12030152 |
| 3 | GAGACGGGCCTGGGATGCCGC | chr11:12029712-12029732 |
| 4 | GCAACTCGGTCCAGTCGGGGT | chr11:12029472-12029492 |
| 5 | GTTCCCGCACCCGCCCGGAGA | chr11: 12029168 -12029187 |
| control | TACCAGAGCTAACTCA | - |

Table S2. Primer sequences used for qPCR AND CoBRA

| Com | | | Primer |
|---------|--|--|---------------|
| Gene | Forward Primer (F) $(5' \rightarrow 3')$ | Reverse Primer (R) $(5' \rightarrow 3')$ | concentration |
| name | | | (F/R) nM |
| DKK3 | TCATCACCTGGGAGCTAGAG | TTCATACTCATCGGGGACCT | 500/500 |
| 36B4 | GTGTTCGACAATGGCAGCAT | AGACACTGGCAACATTGCGGA | 500/500 |
| ANXA2 | CTCTACACCCCCAAGTGCAT | TCAGTGCTGATGCAAGTTCC | 300/300 |
| HFE | CCTTGTTTGAAGCTTTGGGC | CACGGCGACTCTCATGATCA | 300/600 |
| PTGS2 | TGCCATTCTTTGCCCAGCACT | AAAGGCGCAGTTTACGCTGT | 600/600 |
| SLC16A5 | GCCCTGCTTGAGTCTGGAATG | ACTGCCAATGTGGCTGCTG | 900/900 |
| ACGT2 | TACCCCATTGAACACGGCAT | TGCTCTTCAGGTGCTACACG | 600/600 |
| FZD8 | TAGAGCTAGAAATAGCAAGT | GCCACTTTTTCAAGTT | 600/600 |
| TGFBI | CACCAAGAGAACGGAGCAGA | GCCTCCGCTAACCAGGATTT | 300/300 |
| ECM1 | ATTTGGCTGTTGCTTCTGCT | TCTTGAAAGTGCTCTGGCCT | 600/600 |
| CYP27A1 | GTGTCTGGCTACCTGCACTT | TTGGATGTCGTGTCCACTCC | 600/600 |
| NKD1 | ACTTCCAGCCGAAAGTCGT | CACCATAGGCCGAAGCAC | 900/900 |
| DNMT1 | CGACTACATCAAAGGCAGCAACCTG | TGGAGTGGACTTGTGGGTGTTCTC | 300/900 |
| DNMT3A | CGAGTCCAACCCTGTGATGATTG | GCTGGTCTTTGCCCTGCTTTATG | 600/600 |
| DNMT3B | TTGGAATAGGGGACCTCGTGTG | AGAGACCTCGGAGAACTTGCCATC | 600/600 |
| DKK3 | | | 500/500 |
| (CoBRA) | IGGGIIGIIGIAAGIIIGAAGGI | CICACCCACCCIACIAAAC | 500/500 |



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