

Supplemental Information

SI Materials and Methods

Antisera used in immunoblots and CHIP assays

Antibodies used for immunoblots were as follows: AR (N-20, directed against amino acids 1-20 by Bethyl Laboratories, and validated by immunoblot, IP, IHC, and CHIP to perform identically to commercial antisera directed against this epitope), PARP1 (Active Motif, Cat#39559), PARP (for verification of knockdown)(Cell Signaling, Cat#9542S), PAR (Enzo Life Sciences, Cat#804-220-R100), CDK4 (Santa Cruz, Cat#sc-601), GAPDH (Santa Cruz, Cat#sc-25778), Histone H4 (Millipore Cat#07-108), Lamin B (Santa Cruz, Cat#sc-6217) and Ran (BD Transduction Labs, Cat#610340). Antibodies for CHIP assays were as follows: AR (N-20), PARP1 (Active Motif, Cat#39559), Histone H1 (Millipore, Cat#05-457), H3K4me2 (Millipore Cat#07-030), FoxA1 (Abcam, Cat#5089), GATA2 (Santa Cruz, Cat#sc-9008x), and H3K4me3 (Invitrogen, Cat#49-1005).

MEF transfection

Cells were seeded at 2×10^5 cells per well (in 6 well dishes), and allowed to grow 24 hours before transfection. For each well of cells, 4ug of DNA plus serum free media to 50ul were mixed. In a separate tube, 10ul Lipofectamine 2000 (Invitrogen, 11668)) plus 40ul serum free media were mixed. Separately the DNA and Lipofectamine were incubated 5 minutes at room temp, then mixed together and incubated at room temp for 20 minutes before added to the cells. The cells were grown an additional 24 hours before treatment(1).

Dose response/cell survival analysis

Prostate cancer cell lines were plated at 1×10^3 cells per well (Ab1, 22RV1 and C4-2) as previously described(2-4), except for LNCaP and LAPC4 cell lines which were plated at

2×10^3 cells/well in 100ul into 96 well flat bottom culture plates. These dilutions allowed for cells to be plated at 50-70% confluency, depending on the cell line. Once cells attached to the bottom of the 96-well plates (6 hr later), ABT-888 was added at the indicated doses in triplicate in 100ul of additional media. Plates were then incubated at 37C°, 5% CO₂ for 5-7 days. Cell culture media was discarded from the plates and the surviving cells were washed 3 times with PBS. To lyse the surviving cells, 100ul/well H₂O was then added and cells were lysed at 37C°. After 1 hour, cell survival was measured by staining for double-stranded DNA via Quant-iT Picogreen(Invitrogen, Carlsbad, CA, 1:200 dilution with H₂O) and the fluorescence was detected as previously described by a TECAN SpectraFluor. Survival rate was determined by averaging the fluorescence intensity of the treated wells (in triplicate) divided by the fluorescence intensity of the untreated wells.

LB1 cell reporter assay

LB1 cells as described previously (5) were seeded in steroid-proficient media, treated for 24h with either control or ABT888 (2.5uM). 36h after treatment, cells were harvested, lysed, and relative luciferase activity was assessed using a luminometer. Relative luciferase activity was compared to total protein present in the lysate (determined by modified Lowry assay).

Supplemental Table S1

mRNA target	Primer sequences
<i>KLK3/PSA</i>	Forward: TGTGTGCTGGACGCTGGA Reverse: CACTGCCCCATGACGTGAT
<i>TMPRSS2</i>	Forward: GGACAGTGTGCACCTCAAAGAC Reverse: TCCCACGAGGAAGGTCCC
<i>FKBP5</i>	Forward: GCAACAGTAGAAATCCACCTG Reverse: CTCCAGAGCTTTGTCAATTCC
<i>18S</i>	Forward: GCAATTATTCCCCATGAACG Reverse: GGCCTCACTAAACCATCCAA
<i>UBE2C</i>	Forward: TGGTCTGCCCTGTATGATGT Reverse: AAAAGCTGTGGGGTTTTTCC
<i>ERG</i>	Forward: CGCAGAGTTATCGTGCCAGCAGAT Reverse: CCATATTCTTTCACCGCCCACTCC
<i>PARP1</i>	Forward: TGTAGCCTGTACGGGCGCT Reverse: GCGTGAAGGCGAATGCAGC
<i>AR</i>	Forward: CAGTGGATGGGCTGAAAAAT Reverse: GGAGCTTGGTGAGCTGGTAG
<i>KLK3/PSA</i> (for <i>in vivo</i> work)	Forward: GAGCACCCCTATCAACCCCTATT Reverse: AGCAACCCTGGACCTCACACCTAA
<i>β-actin</i>	Forward: CGCGAGAAGATGACCCAGAT Reverse: GAGTCCATCACGATGCCAGT
ChIP primers	Primer sequences
<i>KLK3/PSA</i> promoter	Forward: CCTAGATGAAGTCTCCATGAGCTACA Reverse: GGGAGGGAGAGCTAGCACTTG
<i>KLK3/PSA</i> enhancer	Forward: TGGGACAACCTTGCAAACCTG Reverse: CCAGAGTAGGTCTGTTTTCAATCCA
<i>TMPRSS2</i> ARE V	Forward: TGGTCCTGGATGATAAAAAAGTTT Reverse: GACATACGCCCCACAACAGA
MNase primers	Primer sequences
<i>KLK3/PSA</i> set 1	Forward: CCACATTGTTTGCTGCACGTTGG Reverse: TGGTCCACAGATCCTCTAGCCCAG
<i>KLK3/PSA</i> set 2	Forward: TTCTGGGCTAGAGGATCTGTGGAC Reverse: CCAACCCAGAATCCAGCTCCA
<i>KLK3/PSA</i> set 3	Forward: TGTATCTGTGGAGCTGGATTCTGGG Reverse: TCTTTTCCTTGCACTCCCAACCC
<i>KLK3/PSA</i> set 4	Forward: TGGTCTCAGAGTGGTGCAGGGA Reverse: GCCCTAAGACACCAGCACTCAGG
<i>KLK3/PSA</i> set 5	Forward: AGGGCACACTGGGTCTTGGAGT Reverse: GCCTCACGTGCCTAGATCCTTTGC
<i>KLK3/PSA</i> set 6	Forward: GGGATCGTACCCACCCCTGTT Reverse: GCAGAGGAGACACGCCAGGA
<i>KLK3/PSA</i> set 7	Forward: TGGGCGTGTCTCCTCTGCCT Reverse: TGCACCAGGCCCTGTAGCTC
<i>KLK3/PSA</i> set 8	Forward: CTGGTGCATCCAGGGTGATCTAGT Reverse: TGTGGAAGGGGAGGGAGAGC
<i>KLK3/PSA</i> set 9	Forward: ACAGCTCTGGGTGTGGGAGGG Reverse: CCATGCTGCTGGAGGCTGGAC

<i>KLK3/PSA set 10</i>	Forward: TGGGTGTGGGAGGGGGTTGT Reverse: TGCTGGCACCCAGAGGCTGA
<i>KLK3/PSA set 11</i>	Forward: TTGGTCAGCCTCTGGGTGCC Reverse: TCATTCCCCAGGACTCCGCC
<i>KLK3/PSA set 12</i>	Forward: AGGGCTCCTGGGGGAGGCT Reverse: GCTCTCCGGGTGCAGGTGGT
<i>KLK3/PSA set 13</i>	Forward: TGCACCCGGAGAGCTGTGTC Reverse: CCACGTCACGGACAGGGTGA
<i>KLK3/PSA set 14</i>	Forward: CCCTGTCCGTGACGTGGATTGG Reverse: AGGGCTGGCTCCCTCTCCTG
<i>KLK3/PSA set 15</i>	Forward: GAGGGAGCCAGCCCTGACTGT Reverse: AGAGCCCAGCTCCCTGTCTGG
<i>KLK3/PSA set 16</i>	Forward: ACCCCAGCCCAGACAGGGAG Reverse: ATGGGCTTGGAGTGGGGCTG
<i>KLK3/PSA set 17</i>	Forward: GCCCATACCCCAGCCCCTC Reverse: GCGGGGACCTGGTGTGGGAG
<i>KLK3/PSA set 18</i>	Forward: CGCTCCCTCCCCTTACCCCA Reverse: AGCTGGGAGCAGGGAGCTGG
<i>KLK3/PSA set 19</i>	Forward: CACCAGGTCCCCGCTCCCTC Reverse: GGAGCTGGCTGGGCAATGGG

Supplemental References

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