## **Supplementary figures:**

#### **Supplementary Figure S1.**



**Supplementary Figure S1. The majority of Pb-Cre; PTEN<sup>fl/fl</sup> Trp53<sup>fl/fl</sup> mice are** *de novo* **resistant to ADT. (A-B)** Pb-Cre; PTEN<sup>fl/fl</sup> Trp53<sup>fl/fl</sup> mice with established prostate tumors were randomized to untreated or degarelix (ADT, 0.625 mg, single dose) for 4 weeks. Tumor volumes were non-invasively monitored by MRI, and % ORR at days 14 (R<sub>14</sub>) and 28 (R<sub>28</sub>) were determined as described in Methods. The % change in solid tumor volume is represented by waterfall plots for untreated (A) and ADT-treated groups (**B**). (**C**) Sera were collected from mice at 7 and 28 days post-treatment, and analyzed for testosterone levels by ELISA. n=6-8 mice per group.

Significances/p-values were calculated by Chi-square test (panel A and B, relative to untreated), one-way ANOVA (panel C) and indicates as follows, \*\*\*p<0.001.

#### **Supplementary Figure S2.**



Supplementary Figure S2. ADT/PI3Ki combination therapy halts prostate tumor growth up to 14 days, followed by development of resistance in majority of Pb-Cre; PTEN<sup>fl/fl</sup> Trp53<sup>fl/fl</sup> mice bearing established tumors were treated with copanlisib (14 mg/kg, *iv*, every alternate day), singly or in combination with ADT (degarelix, 0.625 mg, single dose). Tumor volumes were non-invasively monitored by MRI and % ORR at days 14 (R<sub>14</sub>) and 28 (R<sub>28</sub>) were determined as described in Methods. The % change in solid tumor volume is represented by waterfall plots for copanlisib (**A**) and ADT + copanlisib treated groups (**B**). n=6-8 mice per group. Statistical analysis was performed using Chi-square test. \*p<0.05 vs untreated. (**C**) Tumor extracts were harvested following 4 weeks of treatment with the indicated drug(s), and

analyzed by western blotting for PI3K activation status. n=6-8 mice per group. Significances/p-values were calculated by Chi-square test (panel A and B, relative to untreated) and indicates as follows, \*p<0.05.



Supplementary Figure S3. PI3Ki treatment with concurrent androgen depletion does not alter proliferation and survival of PTEN/p53-deficient murine PC cells *in vitro*. (A) PTEN/p53-deficient GEMM tumor-derived PC cell lines (AC1/SC1) were treated with copanlisib (C, 0-500 nM, 24 hours) in a dose-dependent manner and western blot analyses were performed for indicated proteins to determine IC<sub>90</sub>. (B-C) AC1/SC1 cells were treated with copanlisib (IC<sub>90</sub>=100 nM, for 24, 48 and 72 hours) in a time-dependent manner in normal and AD conditions.

(**B**) Proliferation index was calculated by counting cell number at the end of treatment, relative to their baseline seeding density. (**C**) Cells were stained with annexin V antibody/propidium iodide and % cell death was determined via flow cytometry. U=untreated group. n=3 independent experiments. Significances/p-values were calculated by one-way ANOVA.

# **Supplementary Figure S4.**



Supplementary Figure S4. ADT/PI3K inhibitor combination increases MHC-II and PD-1 expression on TAM within the TME of PTEN/p53-deficient murine PC. (A-K) Pb-Cre;

PTEN<sup>fl/fl</sup> Trp53<sup>fl/fl</sup> mice with established tumors were treated with degarelix (0.625 mg, single dose, ADT), copanlisib (14 mg/kg, *iv*, every alternate day, C) or their combination for 7 days. Tumor cell suspensions were collected and analyzed using flow cytometry. Bar graph demonstrates % frequency of TAM (suppressive (**A-C**), and activated (**D**)), Gr-MDSC (CD45<sup>+</sup>CD11b<sup>+</sup>F4/80<sup>-</sup>Ly6g<sup>+</sup>) cells (**E**)), CD4-T cells (CD45<sup>+</sup>CD3<sup>+</sup>CD4<sup>+</sup> cells, total (**F**), exhausted (**G**), and activated (**H**)) and CD8-T cells (CD45<sup>+</sup>CD3<sup>+</sup>CD8<sup>+</sup> cells, total (**I**), exhausted (**J**), and activated (**K**)). (**L-O**) Single cell suspensions were prepared from established PTEN/p53-deficient prostate tumors and treated *ex vivo* with either DMSO (negative control) or copanlisib (C, 100 nM) under normal and AD conditions for 24 hours. Bar graph demonstrates % frequency of MHC-II (**L**), PD-1 (**M**), PD-L1 (**N**) and CD86 (**O**) expressing TAM. U=untreated group. For *in vivo* experiment, n=2 mice per group and for *ex vivo* study, n=3 independent experiments. Significances/p-values were calculated by one-way ANOVA and indicates as follows, \*p<0.05, \*\*p<0.01 and \*\*\*p<0.001; ns = not statistically significant.

## **Supplementary Figure S5.**



**Supplementary Figure S5. PD-1 upregulation suppresses phagocytic capacity of activated TAM. (A)** Experimental schema for phagocytosis experiment, in which sorted TAM from PTEN/p53-deficient prostate GEMM tumors are co-cultured with CTV dye stained-AC1/SC1 cells for 24 hours. (**B**) Bar graphs demonstrate fold change (FC) in phagocytosis of AC1 (upper panel) and SC1 (lower panel) cells by MHC-II<sup>hi/lo</sup>/ PD-1<sup>hi/lo</sup> expressing TAM, relative to untreated group. PD-L1 and CD206 expression were analyzed in the TAM subsets following co-culture with AC1

(C) and SC1 (D) cells. n=2 independent experiments. Significances/p-values were calculated by one-way ANOVA and indicates as follows, \*p<0.05, \*\*p<0.01 and \*\*\*p<0.001.

#### **Supplementary Figure S6.**



Supplementary Figure S6. *Ex vivo* AD + PI3Ki + PD-1 antibody treatment activates MHC-II<sup>10</sup> TAM when co-cultured with PTEN/p53-deficient murine prostate tumor cells. (A-B) TAM subsets (MHC-II<sup>10</sup>/PD-1<sup>hi/lo</sup>) were sorted from PTEN/p53-deficient prostate tumors using flow cytometry, and co-cultured with PTEN/p53-deficient prostate tumor cells under normal and AD conditions. These co-cultures were treated with copanlisib (C, 100 nM), PD-1 antibody (P, 10  $\mu$ g/mL) or their combination for 24 hours. Bar graphs demonstrate MHC-II expression on MHC-II<sup>10</sup>/PD-1<sup>hi/lo</sup> TAM following co-culture with AC1 (A) and SC1 (B) cells in the presence of indicated treatments, relative to untreated (U). FC=fold change. n=2 independent experiments. Significances/p-values were calculated by one-way ANOVA and indicates as follows, \*p<0.05, \*\*p<0.01 and \*\*\*p<0.001; ns = not statistically significant.



Supplementary Figure S7. The addition of PD-1 blockade to androgen depletion/PI3Ki therapy does not alter phagocytic capacity of PD-1<sup>lo</sup> macrophages. (A) TAM subsets (MHC-II<sup>hi/lo</sup>/PD-1<sup>lo</sup>) sorted from PTEN/p53-deficient prostate tumors using flow cytometry were co-

cultured with CTV dye stained-AC1/SC1 cells under normal and AD conditions. These co-cultures were treated with copanlisib (C, 100 nM), PD-1 antibody (P, 10  $\mu$ g/mL) or their combination for 24 hours. Bar graphs demonstrate fold change (FC) in phagocytic activity of MHC-II<sup>hi/lo</sup>/PD-1<sup>lo</sup> expressing TAM relative to untreated group (U) in AC1 (**B**) and SC1 (**C**) cells. n=2 independent experiments. Significances/p-values were calculated by one-way ANOVA and indicates as follows, \*\*p<0.01 and \*\*\*p<0.001; ns = not statistically significant.

#### **Supplementary Figure S8.**



**Supplementary Figure S8. The combination of androgen depletion, PI3Ki and aPD-1 blockade does not alter phagocytic checkpoint expression on PTEN/p53-deficient prostate tumor cells.** (**A-B**) AC1/SC1 cells were treated with copanlisib (C, 100 nM), PD-1 antibody (P, 10 μg/mL) or their combination under normal and AD conditions for 24 hours. Bar graphs demonstrate phagocytosis inhibitory checkpoints (CD47 and PD-1) expression on (**A**) AC1 and (**B**) SC1 cells following indicated treatments. U=untreated group. FC=fold change. n=2 independent experiments. Significances/p-values were calculated by one-way ANOVA.



Supplementary Figure S9. Androgen depletion, singly and in combination with aPD-1, did not alter phagocytosis activity of inactivated MHC-II<sup>lo</sup>/PD-1<sup>lo</sup> and MHC-II<sup>lo</sup>/PD-1<sup>hi</sup> TAM subsets. (A-B) Pre-treatment of FACS-sorted TAM subsets from PTEN/p53-deficient prostate tumors with copanlisib (C, 100 nM), PD-1 antibody (P, 10 μg/mL) or their combinations under normal and AD conditions for 24 hours. After PBS wash, treated TAM were co-cultured with CTV dye stained-AC1/SC1 cells for 2 hours. Bar graphs demonstrate fold change (FC) in phagocytosis

of AC1 (**A**) and SC1 (**B**) cells by MHC-II<sup>lo</sup>/PD-1<sup>hi/lo</sup> expressing TAM, relative to untreated group (U). n=2 independent experiments. Significances/p-values were calculated by one-way ANOVA.

**Supplementary Figure S10.** 



Supplementary Figure S10. Androgen depletion, not PI3Ki or aPD1, directly enhances TAM activation within the TME of PTEN/p53-deficient PC. (A-B) FACS-sorted TAM from PTEN/p53-deficient prostate tumors were directly treated with copanlisib (C, 100 nM), PD-1 antibody (P, 10  $\mu$ g/mL) or their combinations under normal and AD conditions for 24 hours. Bar graphs demonstrate MHC-II expression on TAM following co-culture with AC1 (A) and SC1 (B) cells for 2 hours, in the presence of indicated treatments. U=untreated group. FC=fold change. n=2 independent experiments. Significances/p-values were calculated by one-way ANOVA and indicates as follows, \*p<0.05.

**Supplementary Figure S11.** 



**Supplementary Figure S11. PI3Ki does not alter phagocytosis/histone lactylation status of MHC-II<sup>Io</sup>/PD-1<sup>Io</sup> TAM and MHC-II<sup>Io</sup>/PD-1<sup>hi</sup> TAM. (A-B)** Single cell suspension of PTEN/p53deficient prostate GEMM tumors were treated with copanlisib (C, 100 nM) and CM was collected 24 hours following treatment. FACS-sorted TAM from PTEN/p53-deficient prostate tumors were cultured *ex vivo* with CM for 24 hours in presence or absence of lactate (100 nmol/µL). After PBS wash, TAM were co-cultured with CTV dye stained-AC1/SC1 cells for 2 hours. Bar graphs demonstrate fold change (FC) in phagocytic activity and histone lactylation status of inactivated MHC-II<sup>Io</sup>/PD-1<sup>Io</sup> and MHC-II<sup>Io</sup>/PD-1<sup>hi</sup> TAM following incubation with AC1 (**A**) and SC1 (**B**) cells, relative to untreated group. n=2 independent experiments. Significances/p-values were calculated by one-way ANOVA. Supplementary Figure S12.



Supplementary Figure S12. PI3Ki inhibits lactate secretion from PTEN/p53-deficient prostate tumor cells within TME. Single cell suspensions of PTEN/p53-deficient prostate GEMM tumors or tumor-derived AC1/SC1 cells were treated with copanlisib (100 nM) for 24 hours. *Ex vivo* or *in vitro* CM were collected from these groups and analyzed for lactate content using colorimetry kits. n=3 independent experiments. Significances/p-values were calculated by one-way ANOVA and indicates as follows, \*\*p<0.01.



**Supplementary Figure S13.** 

Supplementary Figure S13. Direct *ex vivo* treatment of TAM with PI3Ki, singly and in combination with PD-1 antibody and/or androgen depletion does not alter their histone lactylation profile. FACS-sorted TAM from PTEN/p53-deficient prostate tumors were directly treated with copanlisib (C, 100 nM), PD-1 antibody (P, 10  $\mu$ g/mL) or their combinations under normal and AD conditions for 24 hours. Bar graphs demonstrate histone lactylation status of MHC-II<sup>hi/lo</sup>/PD-1<sup>hi/lo</sup> expressing TAM in response to indicated treatments. U=untreated group. FC=fold change. n=2 independent experiments. Significances/p-values were calculated by one-way ANOVA.

## Supplementary Figure S14.



Supplementary Figure S14. ADT + PI3Ki + aPD-1 induces tumor control in 60% of Pb-Cre; PTEN<sup>fl/fl</sup> TP53<sup>fl/fl</sup> mice. (A-D) Pb-Cre; PTEN<sup>fl/fl</sup> Trp53<sup>fl/fl</sup> mice with established tumors were treated with PD-1 antibody (aPD-1, 100  $\mu$ g/mouse, *ip*, every alternate day) alone or in combination with ADT (degarelix, 0.625mg, single dose) or copanlisib (14 mg/kg, *iv*, every alternate day) or ADT + copanlisib. Tumor volumes were non-invasively monitored by MRI and % ORR at days 14 (R<sub>14</sub>) and 28 (R<sub>28</sub>) were determined, as described in Methods. The % change in solid tumor

volume are represented by waterfall plot for the following treated groups: aPD-1 (**A**), ADT + aPD-1 (**B**), copanlisib + aPD-1 (**C**) and ADT + copanlisib + aPD-1 (**D**). n=6-10 mice per group. Significances/p-values were calculated by Chi-square test and indicated as follows, \*p<0.05 and  $^{\text{@}}$ p<0.05 (relative to untreated and aPD-1 treated groups, respectively).



Supplementary Figure S15. Depletion of activated TAM abrogates anti-cancer response elicited by ADT + PI3Ki + PD-1 antibody treatment in the PTEN/p53-deficient murine

**prostate GEMM tumors.** (**A-B**) Pb-Cre; PTEN<sup>fl/fl</sup> Trp53<sup>fl/fl</sup> mice with established prostate tumors were treated with clodronate (which depletes activated macrophages, 200 µg/mouse, *ip*, every week) alone or in combination with ADT (degarelix, 0.625mg, single dose) + copanlisib (14 mg/kg, *iv*, every alternate day) + PD-1 antibody (aPD-1, 100 µg/mouse, *ip*, every alternate day). Tumor volumes were non-invasively monitored by MRI and % ORR at days 14 (R<sub>14</sub>) and 28 (R<sub>28</sub>) were determined as described in Methods. The % change in solid tumor volume are represented by waterfall plot for clodronate (**A**) and ADT + copanlisib + aPD-1 + clodronate (**B**) treated groups. (**C**) Flow cytometry plot represents frequency of MHC-II expressing TAM within GEMM tumors from mice treated with the indicated drugs, relative to untreated mice. n=5-6 mice per group. Significances/p-values were calculated by Chi-square test and indicates as follows, #p<0.05 (relative to ADT/copanlisib/aPD-1 combination treated group).

**Supplementary Figure S16.** 



Supplementary Figure S16. Long-term treatment of ADT + PI3Ki + aPD-1 activates Wnt/ $\beta$ catenin pathway in murine PTEN/p53-deficient GEMM-derived SC1 cells. (A-C) SC1 cells were treated with copanlisib (C, 100 nM) + PD-1 antibody (P, 10 µg/mL) under AD condition for 24 and 72 hours. Western blot analyses (A) were performed for indicated protein markers on SC1 cell lysates, quantified using Image J software (B), and lactate levels (C) were determined by fluorimetry in the indicated SC1 supernatant groups. U=untreated group. n=3 independent experiments. Significances/p-values were calculated by one-way ANOVA and indicates as follows, \*p<0.05, \*\*p<0.01 and \*\*\*p<0.001; ns = not statistically significant.

## Supplementary Figure S17.



Supplementary Figure S17. Feedback Wnt/ $\beta$ -catenin-pathway activation within murine PTEN/p53-deficient GEMM-derived PC cells following long-term ADT + copanlisib + aPD-1 treatment suppresses phagocytosis via increased histone lactylation within bone marrow derived macrophages (BMDM). (A) Schema illustrating phagocytosis experiment on *in vitro* CM. AC1/SC1 cells were treated with copanlisib (C, 100 nM) + PD-1 antibody (P, 10 µg/mL) under AD condition for 24 and 72 hours. *In vitro* CM were collected from these groups and cocultured with BMDM for 24 hours. After PBS wash, these BMDM were co-cultured with CTV dye stained-AC1/SC1 cells for 2 hours. (B) Bar graphs demonstrate fold change (FC) in phagocytic activity and histone lactylation status of activated (MHC-II<sup>hi</sup>) BMDM when co-cultured with

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#### In vitro CM co-culture with BMDM

AC1/SC1 cells, relative to untreated group (U). n=3 independent experiments. Significances/p-values were calculated by one-way ANOVA and indicates as follows, p<0.05 and p<0.01; ns = not statistically significant.