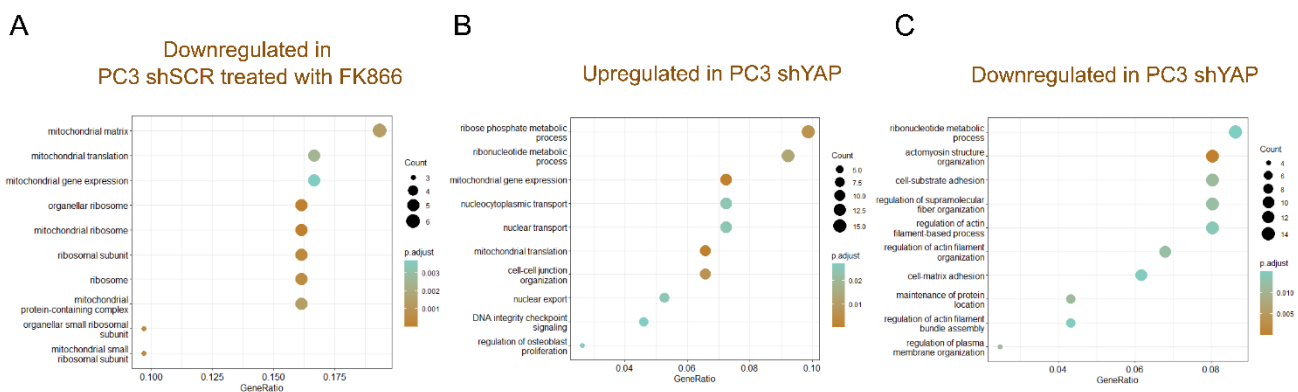


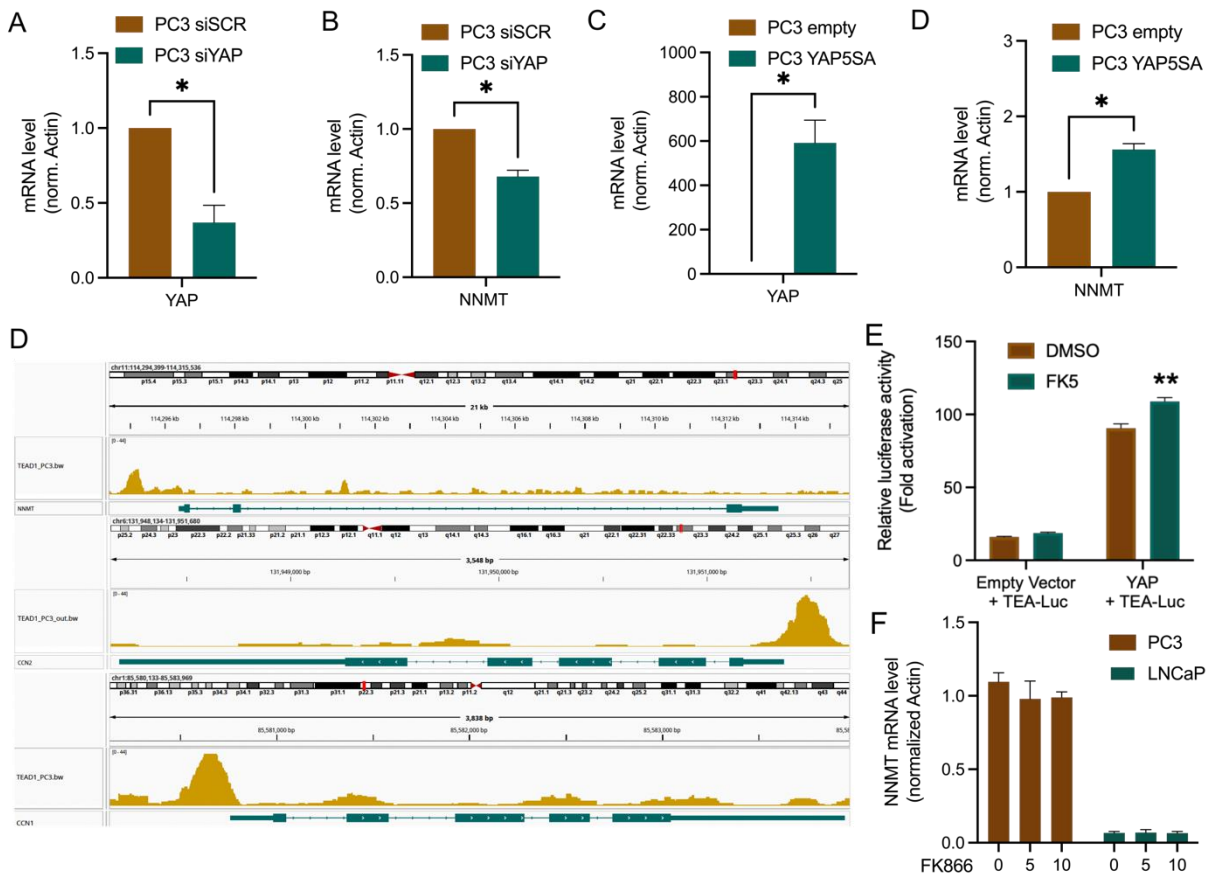
Nicotinamide N-methyl transferase (NNMT) sustains innate sensitivity to NAMPT inhibition in YAP-dependent stem-like/ mesenchymal prostate cancer

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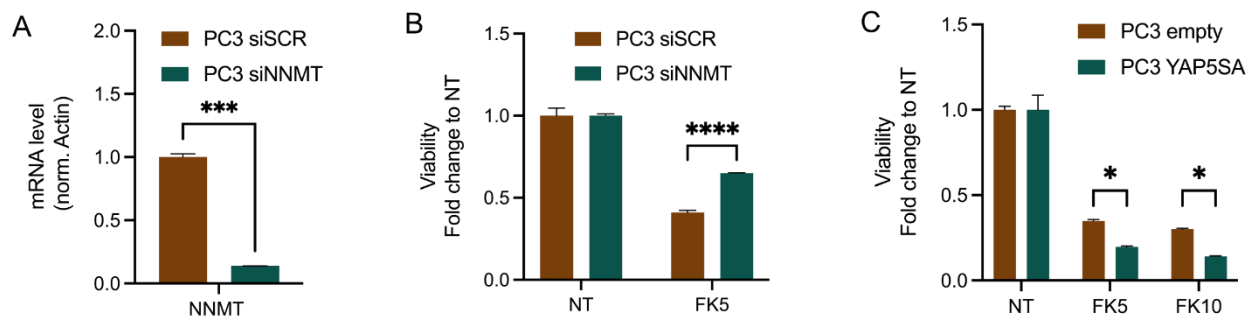
SUPPLEMENTARY FIGURES



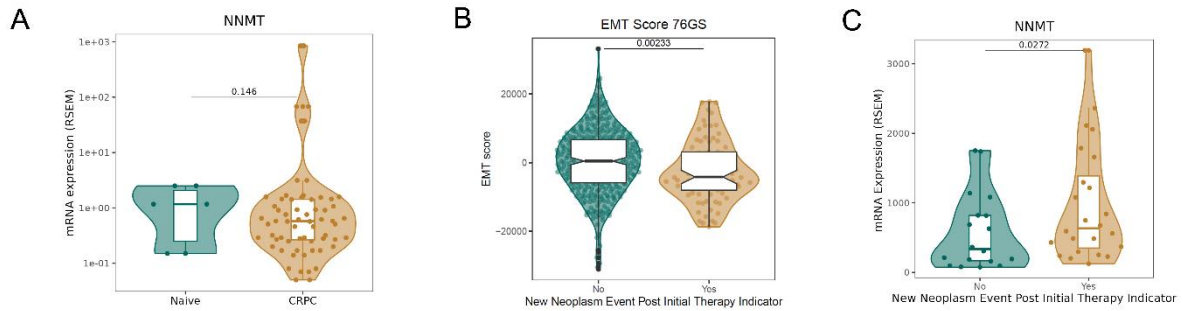
Supplementary Figure 1: Gene set enrichment analysis (GSEA) of proteomics data. **(A)** GSEA of downregulated proteins in PC3 shSCR cells treated with FK866 (5nM) for 48 hours **(B)** GSEA of proteins upregulated or **(C)** downregulated in PC3 shYAP, compared with the scramble condition (PC3 shSCR). Significantly modulated proteins presented a p-value above 0,05, calculated from three independent biological replicates.



Supplementary Figure 2: mRNA levels of (A), (C) *YAP* and (B), (D) *NNMT* in PC3 cells transfected either with siRNA targeting *YAP* or with plasmids overexpressing constitutively active *YAP* (YAP5SA) for 48 hours. Actin was used as the housekeeping gene. (D) ChIP-seq evaluation of TEAD1 binding sites in the promoter region of *NNMT* locus in PC3 cells. (E) Luciferase signal in PC3 cells transfected with either pRGFPN1 (empty vector) or *YAP*, as well as with the firefly TEA-Luc reporter (8xGTIIc-Luc reporter) and with the Renilla luciferase. Cells were treated with 5 nM of FK866 (FK5) or DMSO for 48 hours. Firefly luciferase signals were normalized to the ones of Renilla luciferase (F) *NNMT* mRNA levels in PC3 and LNCaP cells treated with 5 or 10 nM of FK866 for 48 hours. Actin was used as the housekeeping gene. Student t-test was used to calculate statistical significance (ns- not significant, * $P < 0.05$, ** $P < 0.01$) between control and experimental conditions.



Supplementary Figure 3: (A) mRNA level of NNMT in PC3 cells transiently transfected with siRNAs targeting *NNMT* for 48 hours. siSCR was used as a control of the transfection. Actin was used as a housekeeping gene. **(B)** Viability assay of PC3 cells treated with FK866 at 5 nM (FK5) concentration, for 48 hours, after 24 hours of *NNMT* silencing with siNNMT. **(C)** Viability assay of PC3 cells treated with FK866 at 5 (FK5) or 10 nM (FK10) of concentration, for 48 hours, after 24 hours of transfection with a constitutively active YAP plasmid (YAP5SA) or the empty control (empty). Repeated measures one-way ANOVA was used to calculate statistical significance (ns- not significant, * $P < 0.05$, ** $P < 0.01$, *** $P < 0.001$, **** $P < 0.0001$) between control and experimental conditions.



Supplementary Figure 4: (A) NNMT expression levels of PDX samples between naive and CRPC conditions. (B-C) Violin plots displaying EMT scores and NNMT mRNA changes, between prostate cancer patients displaying or not a new neoplastic event after treatment. The score was calculated using the 76GS scoring method. Wilcoxon rank-sum test was used to calculate statistical significance across the different conditions.

Supplementary Material 1: list of primers

Gene	Primers (5'-3')
<i>E-cadherin (CDH1)</i>	FW: ACACCCGGGACAACGTTTA RV: TGTGCAGCTGGCTCAAGT
<i>N-cadherin (CDH2)</i>	FW: CGGTTTCATTTGAGGGCACA RV: TTGGAGCCTGAGACACGATT
<i>Vimentin</i>	FW: GAGAACTTTGCCGTTGAAGC RV: GCTTCCTGTAGGTGGCAATC
<i>Snail1</i>	FW: AGTGGTTCTTCTGCGCTACT RV: GTAGGGCTGCTGGAAGGTAA
<i>ZEB1</i>	FW: GAAAATGAGCAAAACCATGATCCTA RV: CAGGTGCCTCAGGAAAAATGA
<i>NAMPT</i>	FW: TTATGGAACGAAAGATCCTG RV: CAAAAGCATCTTTTTCATGGTC
<i>CTGF</i>	FW: CCAATGACAACGCCTCCTG RV: TGGTGCAGCCAGAAAGCTC
<i>CYR61</i>	FW: AGCCTCGCATCCTATACAACC RV: TTCTTTCACAAGGCGGCACTC
<i>CHOP</i>	FW: AGAACCAGGAAACGGAACAGA RV: TCTCCTTCATGCGCTGCTTT
<i>NNMT</i>	FW: GTTTGGTTCTAGGCACTCTGCAG RV: AGAGCCGATGTCAATCAGCAGG
<i>CK8</i>	FW: CGGGGGATCCAACACTTTCA RV: GCTTCCCATCTCGGGTTTCA
<i>CK18</i>	FW: CCAGATTGCCAGCTCTGGAT RV: ATGTCCGCCATGATCTTGCT
<i>CK5</i>	FW: CAGAGCTGAGGAACATGCAG RV: CACAACTCATTCTCAGCCG
<i>CK14</i>	FW: GGCCCACTGAGATCAAAGAC RV: GGCCCACTGAGATCAAAGAC
<i>Twist</i>	FW: GGCCCACTGAGATCAAAGAC RV: GGCCCACTGAGATCAAAGAC
<i>Slug</i>	FW: GGCCCACTGAGATCAAAGAC RV: GGCCCACTGAGATCAAAGAC
<i>Snail1 (Figure 4)</i>	FW: GGCCCACTGAGATCAAAGAC RV: GGCCCACTGAGATCAAAGAC
<i>E-cadherin (CDH1) - Figure 4</i>	FW: GGCCCACTGAGATCAAAGAC RV: GGCCCACTGAGATCAAAGAC
<i>N-cadherin (CDH2) - Figure 4</i>	FW: GGCCCACTGAGATCAAAGAC RV: GGCCCACTGAGATCAAAGAC
<i>RPLP0</i>	FW: CATTCTCGCTTCCTGGAG RV: CTTGACCTTTTCAGCAAGTGG
<i>Beta-actin</i>	FW: AGTGTGACGTTGACATCCGT RV: CTTGCTGATCCACATCTGCT

Supplementary Material 2: list of antibodies

Antibodies	Company	Codice
β-actin	Cell Signaling	4970S
GAPDH	Cell signalling	5174S
NNMT	Origene	TA502624