

**Supplementary information for  
Chemotherapy-induced adenosine A2B receptor  
expression mediates epigenetic regulation of  
pluripotency factors and promotes breast cancer  
stemness**

Jie Lan, Guangyao Wei, Jia Liu, Fan Yang, Rong Sun, Haiquan Lu

Corresponding Author: Haiquan Lu

Email: [lvhaiquan@sdu.edu.cn](mailto:lvhaiquan@sdu.edu.cn)

This file includes:

Supplementary figures S1-S7;

Supplementary figure legends S1-S7;

Supplementary tables S1-S6.

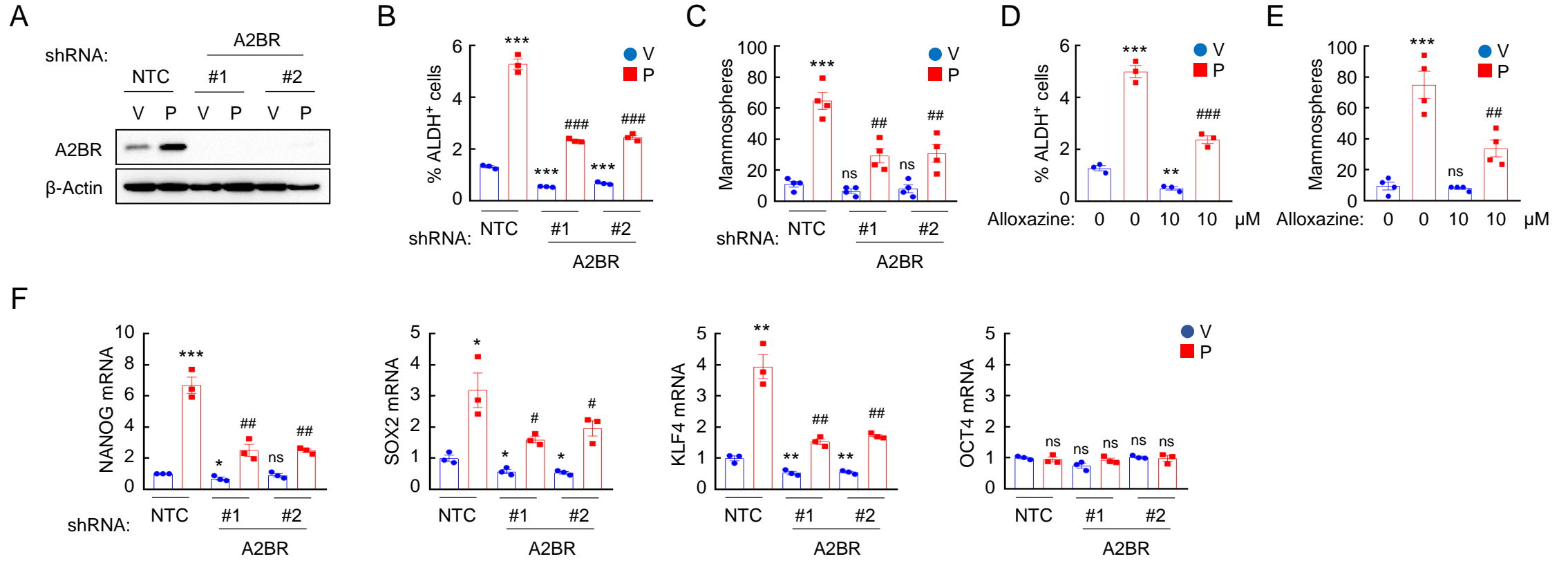


Figure S2

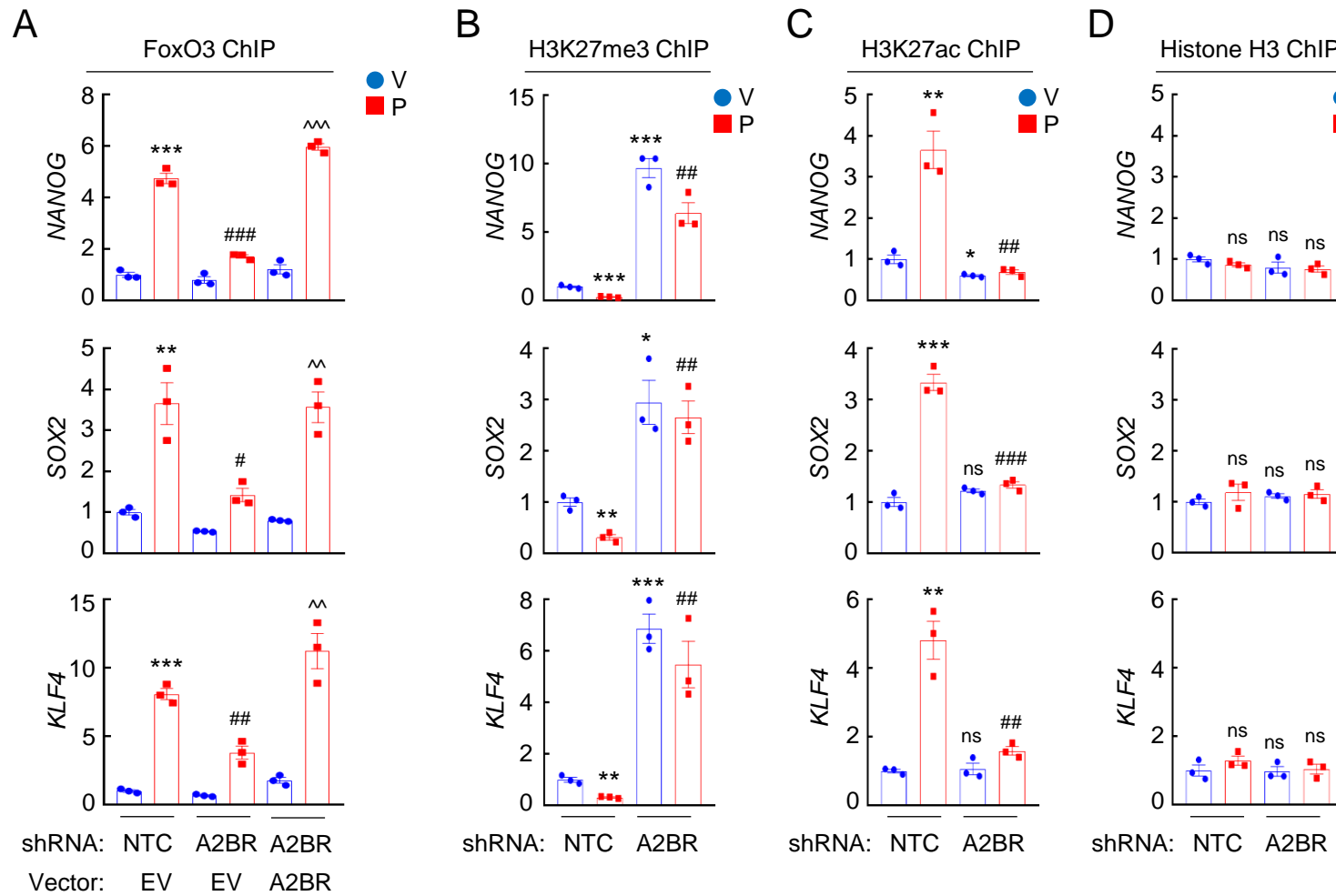
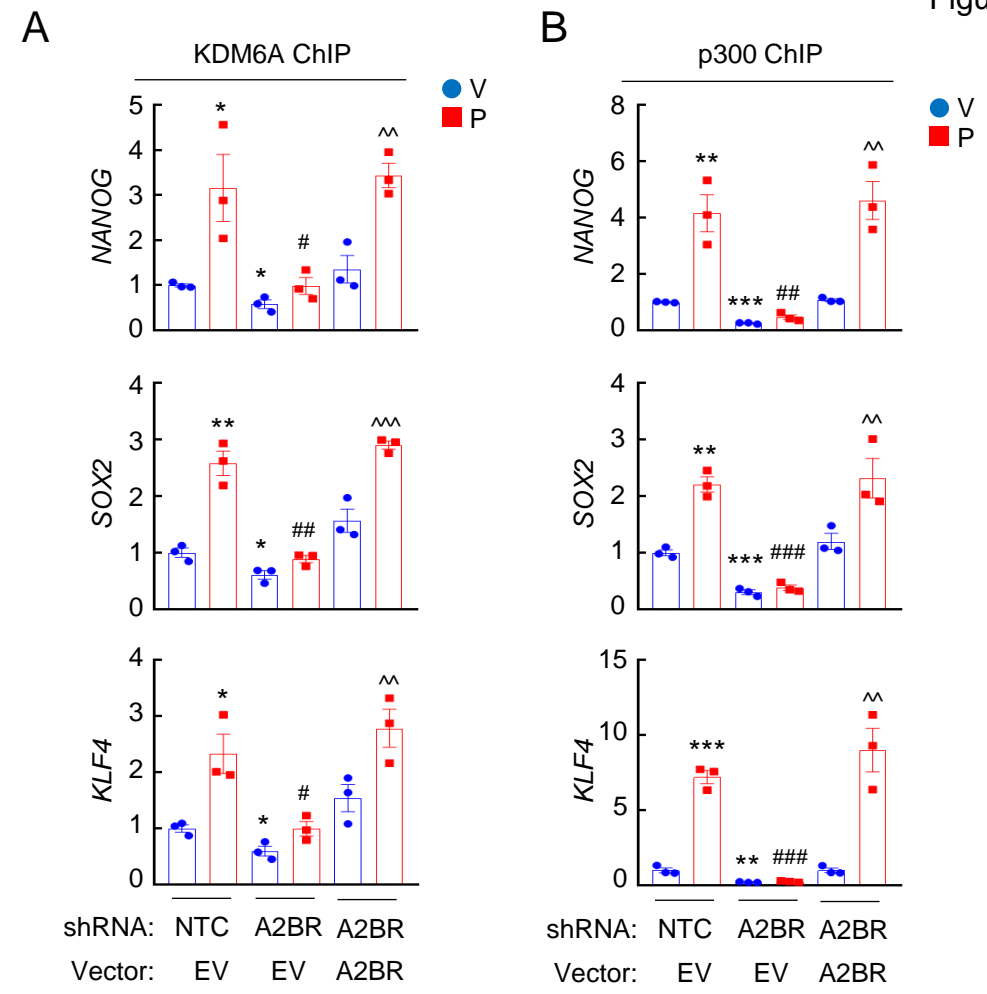


Figure S3





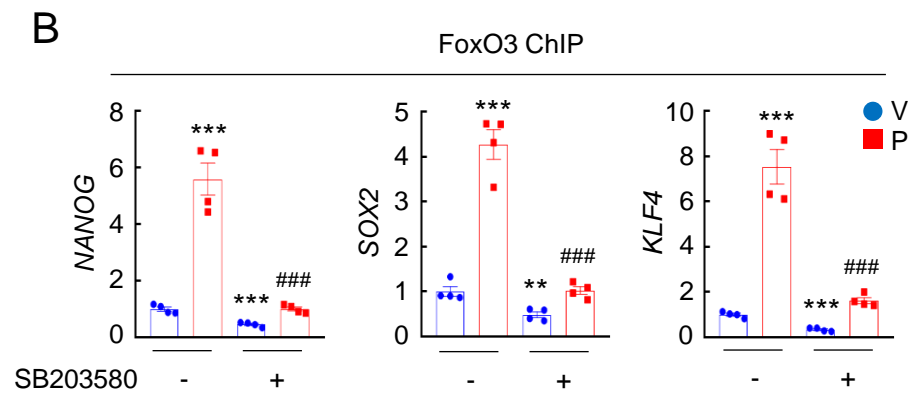
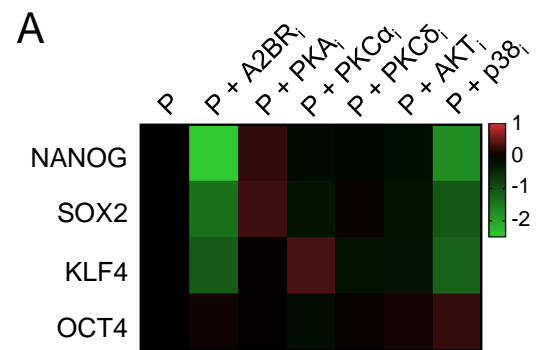
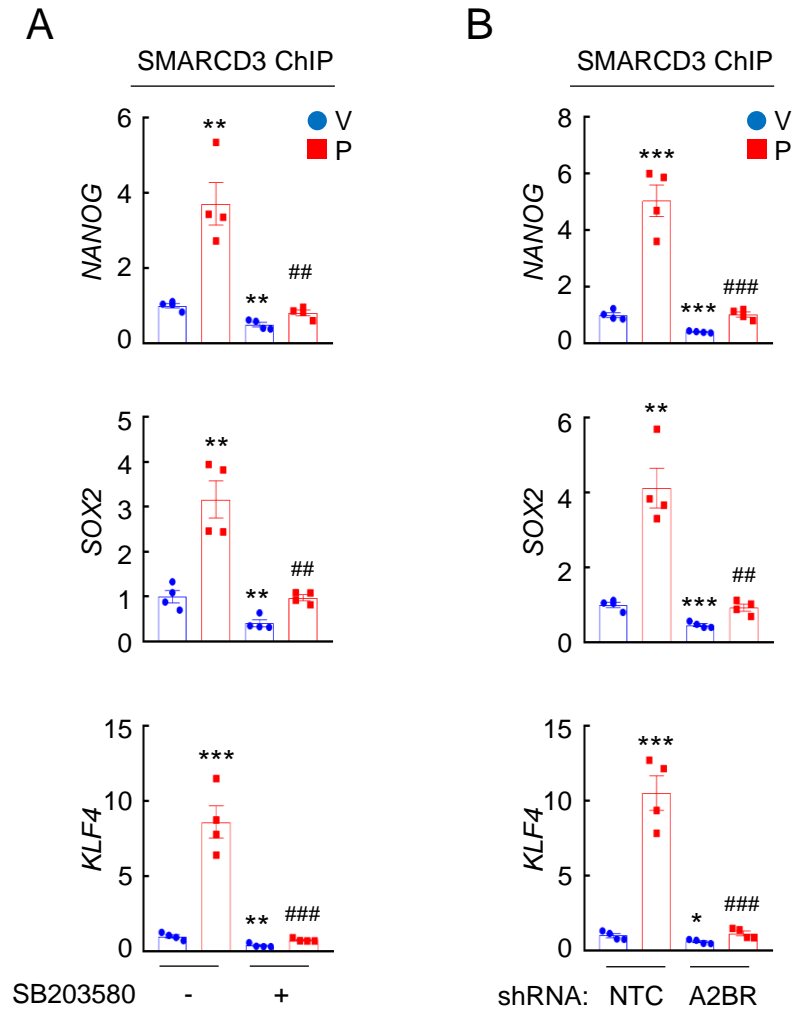
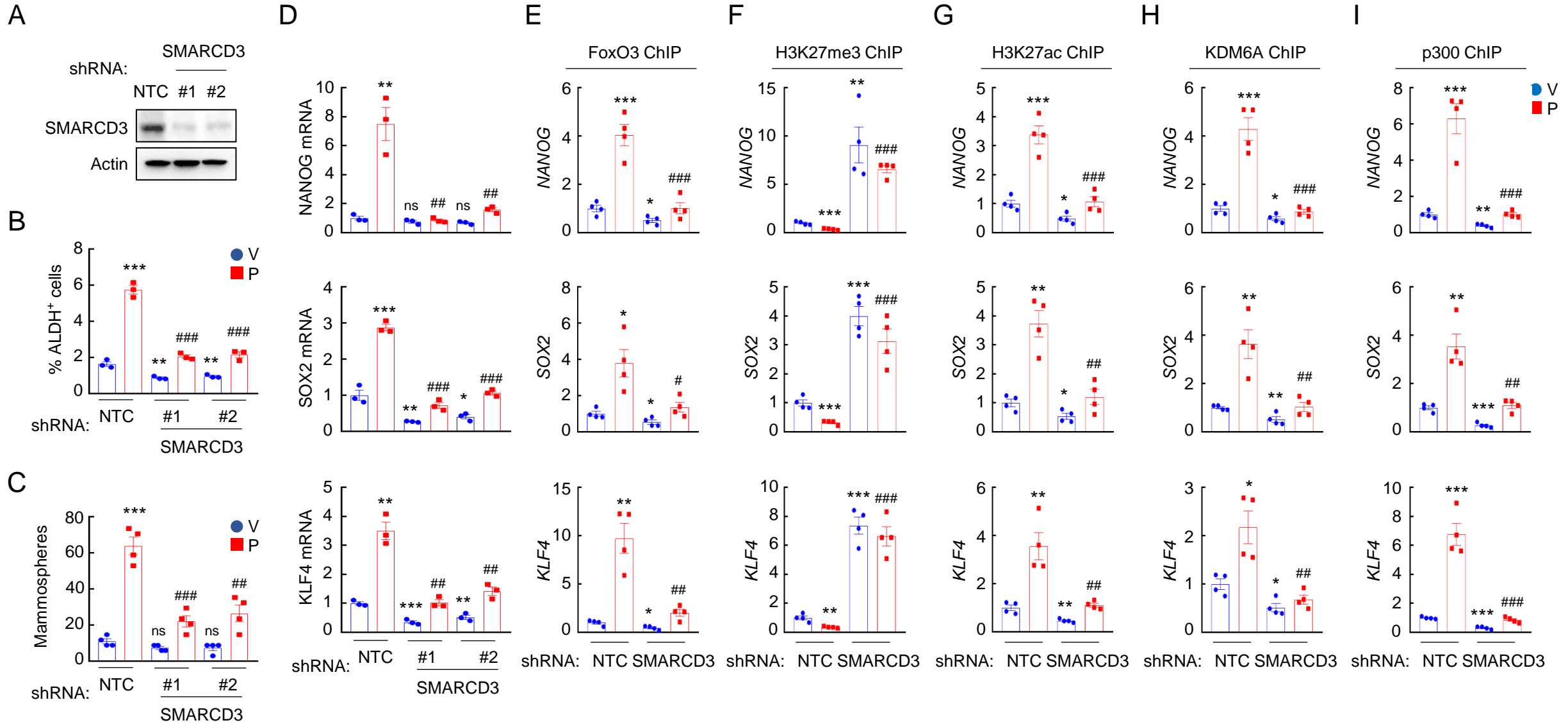


Figure S6







## Supplementary Figure Legends

### Figure S1. Chemotherapy-induced A2BR expression promotes pluripotency

**factor expression and the BCSC phenotype. A-C**, SUM159 subclones stably transfected with vector encoding non-targeting control shRNA (NTC) or either of two different shRNAs targeting A2BR (#1 and #2) were treated with vehicle (V) or 5 nM paclitaxel (P) for 72 hours. Expression of A2BR protein (A), the percentage of ALDH<sup>+</sup> cells (B; mean  $\pm$  SEM; n = 3) and the number of mammospheres formed per 1,000 cells seeded (C; mean  $\pm$  SEM; n = 4) were determined; \*\*\*p < 0.001 vs. NTC-V; ##p < 0.01, ###p < 0.001 vs. NTC-P; ns, not significant. **D and E**, SUM159 cells were treated with 5 nM paclitaxel, 10  $\mu$ M alloxazine, or both for 72 hours. The percentage of ALDH<sup>+</sup> cells (D; mean  $\pm$  SEM; n = 3) and the number of mammospheres formed per 1,000 cells seeded (E; mean  $\pm$  SEM; n = 4) were determined; \*\*\*p < 0.001 vs. V; ##p < 0.01, ###p < 0.001 vs. P; ns, not significant. **F**, SUM159 NTC or A2BR knockdown subclones were treated with V or 5 nM P for 72 hours, and RT-qPCR assays were performed (mean  $\pm$  SEM; n = 3); \*p < 0.05, \*\*p < 0.01, \*\*\*p < 0.001 vs. NTC-V; #p < 0.05, ##p < 0.01 vs. NTC-P; ns, not significant.

### Figure S2. Chemotherapy-induced A2BR expression promotes FOXO3 binding

#### on pluripotency factor genes through decreased H3K27me3 and increased

**H3K27ac chromatin marks. A**, SUM159 NTC or A2BR knockdown subclones were transfected with pLX304 (empty vector, EV) or pLX304 encoding A2BR. Cells were treated with vehicle (V) or 5 nM paclitaxel (P) for 72 hours and chromatin immunoprecipitation (ChIP) was performed with FOXO3 antibody followed by qPCR with primers flanking

FOXO3 binding sites in the *NANOG*, *SOX2* and *KLF4* gene (mean  $\pm$  SEM; n = 3); \*\*p < 0.01, \*\*\*p < 0.001 vs. NTC-V; #p < 0.05, ##p < 0.01, ###p < 0.001 vs. NTC-P; ^^p < 0.01, ^^p < 0.0001 vs. A2BR shRNA/EV-P; ns, not significant. **B-D**, SUM159 NTC or A2BR knockdown subclones were treated with V or 5 nM P for 72 hours. ChIP was performed using antibodies against H3K27me3 (B), H3K27ac (C), or Histone H3 (D) followed by qPCR with primers flanking FOXO3 binding sites in the *NANOG*, *SOX2* and *KLF4* gene (mean  $\pm$  SEM; n = 3); \*\*p < 0.01, \*\*\*p < 0.001 vs. NTC-V; #p < 0.01, ###p < 0.001 vs. NTC-P; ns, not significant.

**Figure S3. A2BR regulates recruitment of KDM6A and p300 at FOXO3 binding sites of pluripotency factor genes.** **A** and **B**, SUM159 NTC or A2BR knockdown subclones were transfected with pLX304 (empty vector, EV) or pLX304 encoding A2BR. Cells were treated with vehicle (V) or 5 nM paclitaxel (P) for 72 hours. ChIP was performed using antibodies against KDM6A (A) or p300 (B) followed by qPCR with primers flanking FOXO3 binding sites in the *NANOG*, *SOX2* and *KLF4* gene (mean  $\pm$  SEM; n = 3); \*p < 0.05, \*\*p < 0.01, \*\*\*p < 0.001 vs. NTC-V; #p < 0.05, ##p < 0.01, ###p < 0.001 vs. NTC-P; ^^p < 0.01, ^^p < 0.0001 vs. A2BR shRNA/EV-P; ns, not significant.

**Figure S4. A2BR decreases H3K27me3 and increases H3K27ac marks and promotes FOXO3 binding on pluripotency factor genes in vivo.** **A-F**,  $2 \times 10^6$  MDA-MB-231 NTC or A2BR knockdown subclone cells were implanted into the MFP of SCID mice. When tumor volume reached 200 mm<sup>3</sup> (day 0), mice were grouped randomly and treated with vehicle (V) or paclitaxel (P; 10 mg/kg, days 0, 5, and 10). Tumors were harvested on day 13, and ChIP was performed using antibodies against FOXO3 (A),

H3K27me3 (B), H3K27ac (C), Histone H3 (D), KDM6A (E), or p300 (F) followed by qPCR with primers flanking FOXO3 binding sites in the *NANOG*, *SOX2* and *KLF4* gene (mean  $\pm$  SEM; n = 4); \*p < 0.05, \*\*p < 0.01 vs. NTC-V; #p < 0.05, ###p < 0.01 vs. NTC-P; ns, not significant.

**Figure S5. Inhibition of p38 MAPK blocks paclitaxel-induced FOXO3 binding and expression of pluripotency factor genes.** **A**, SUM159 cells were treated with 5 nM paclitaxel (P), alone or in combination with 10  $\mu$ M alloxazine (A2BR<sub>i</sub>), 2.5  $\mu$ M H-89 (PKA<sub>i</sub>), 1  $\mu$ M Gö6983 (PKC $\alpha$ <sub>i</sub>), 1  $\mu$ M rottlerin (PKC $\delta$ <sub>i</sub>), 1  $\mu$ M MK-2206 (AKT<sub>i</sub>), or 5  $\mu$ M SB203580 (p38<sub>i</sub>), for 72 hours. RT-qPCR assays were performed and log<sub>2</sub> fold change of mRNA expression of pluripotency factors (vs. P) is presented as a heat map. **B**, SUM159 cells were treated with 5 nM paclitaxel, 5  $\mu$ M SB203580, or both for 72 hours. ChIP was performed using FOXO3 antibody followed by qPCR with primers flanking FOXO3 binding sites in the *NANOG*, *SOX2* and *KLF4* gene (mean  $\pm$  SEM; n = 4); \*\*p < 0.01, \*\*\*p < 0.001 vs. V; ###p < 0.001 vs. P.

**Figure S6. p38 MAPK inhibition or A2BR knockdown blocks paclitaxel-induced SMARCD3 binding on pluripotency factor genes.** **A**, SUM159 cells were treated with 5 nM paclitaxel (P), 5  $\mu$ M SB203580, or both for 72 hours. ChIP was performed using SMARCD3 antibody followed by qPCR with primers flanking FOXO3 binding sites in the *NANOG*, *SOX2* and *KLF4* gene (mean  $\pm$  SEM; n = 4); \*\*p < 0.01, \*\*\*p < 0.001 vs. V; #p < 0.01, ###p < 0.001 vs. P. **B**, SUM159 NTC or A2BR knockdown subclones were treated with V or 5 nM P for 72 hours. ChIP was performed using SMARCD3 antibody followed by qPCR with primers flanking FOXO3 binding sites in

the *NANOG*, *SOX2* and *KLF4* gene (mean  $\pm$  SEM; n = 4); \*p < 0.05, \*\*p < 0.01, \*\*\*p < 0.001 vs. NTC-V; ##p < 0.01, ###p < 0.001 vs. NTC-P.

**Figure S7. SMARCD3 knockdown blocks paclitaxel-induced FOXO3 binding on pluripotency factor genes and inhibits BCSC enrichment.** **A**, SUM159 cells were transfected with vector encoding NTC or either of two shRNAs targeting SMARCD3 (#1 and #2), and immunoblot assay was performed. **B-D**, SUM159 NTC or SMARCD3 knockdown subclones were treated with vehicle (V) or 5 nM paclitaxel (P) for 72 hours, and ALDH (B; mean  $\pm$  SEM; n = 3), mammosphere (C; mean  $\pm$  SEM; n = 4), and qPCR (D; mean  $\pm$  SEM; n = 3) assays were performed; \*p < 0.05, \*\*p < 0.01, \*\*\*p < 0.001 vs. NTC-V; ##p < 0.01, ###p < 0.001 vs. NTC-P; ns, not significant. **E-I**, SUM159 NTC or SMARCD3 knockdown subclones were treated with V or 5 nM P for 72 hours. ChIP was performed using antibodies against FOXO3 (E), H3K27me3 (F), H3K27ac (G), KDM6A (H), or p300 (I), followed by qPCR with primers flanking FOXO3 binding sites in the *NANOG*, *SOX2* and *KLF4* gene (mean  $\pm$  SEM; n = 4); \*p < 0.05, \*\*p < 0.01, \*\*\*p < 0.001 vs. NTC-V; ##p < 0.01, ###p < 0.001 vs. NTC-P; ns, not significant.

**Table S1. Chemical information.**

<b>Chemical</b>	<b>Manufacture</b>	<b>Catalog #</b>
Paclitaxel	MilliporeSigma	T7402
Carboplatin	MilliporeSigma	C2538
Alloxazine	MilliporeSigma	A28651
H-89	Cayman Chemical	S1582
Gö6983	Santa Cruz	sc-203432
rottlerin	SelleckChem	S7862
MK-2206	SelleckChem	S1078
SB203580	Enzo	BML-EI286
LY2228820	APExBio	A5566

**Table S2. Oligonucleotide sequence of shRNAs.**

<b>shRNA</b>	<b>Clone ID</b>
ADORA2B #1	NM_000676.2-926s21c1
ADORA2B #2	NM_000676.2-1478s21c1
SMARCD3 #1	NM_003078.3-1190s21c1
SMARCD3 #2	NM_003078.3-1052s21c1

**Table S3. Primary antibody information for immunoblot assays.**

<b>Antibody</b>	<b>Manufacture</b>	<b>Catalog #</b>
A2BR	Novus Biologicals	NBP1-68953
NANOG	Novus Biologicals	AF1997
SOX2	Bioss	bs-0523R
KLF4	Novus Biologicals	NBP1-83940
OCT4	Novus Biologicals	NB100-2379
p-FoxO3 (S294)	Cell Signaling Technology	5538
FoxO3	Novus Biologicals	NBP2-16521
H3K27me3	Novus Biologicals	NBP2-16840
H3K27ac	Novus Biologicals	NBP2-54615
KDM6A	Novus Biologicals	NBP1-80628
p300	Novus Biologicals	NB100-616
p-p38 (Y182)	Novus Biologicals	NB100-82097
p38	Novus Biologicals	AF8691
SMARCD3	Santa Cruz	sc-101163
$\alpha$ -tubulin	Novus Biologicals	NB100-690
Histone H3	Novus Biologicals	NB500-171
Actin	Santa Cruz	sc-1616

**Table S4. Oligonucleotide sequence of RT-qPCR primers.**

<b>Gene</b>	<b>sequence (5' to 3')</b>
<i>NANOG</i> (human)	Forward: TTTGTGGGCCTGAAGAAACT
	Reverse: AGGGCTGTCCTGAATAAGCAG
<i>SOX2</i> (human)	Forward: GCCGAGTGGAACTTTTGTCTG
	Reverse: GGCAGCGTGTACTTATCCTTCT
<i>KLF4</i> (human)	Forward: CGGACATCAACGACGTGAG
	Reverse: GACGCCTTCAGCACGAACT
<i>OCT4</i> (human)	Forward: TTCAGCCAAACGACCATCTG
	Reverse: CACGAGGGTTTCTGCTTTGC
<i>Nanog</i> (mouse)	Forward: TCTTCCTGGTCCCCACAGTTT
	Reverse: GCAAGAATAGTTCTCGGGATGAA
<i>Sox2</i> (mouse)	Forward: GCGGAGTGGAACTTTTGTCC
	Reverse: CGGGAAGCGTGTACTTATCCTT
<i>Klf4</i> (mouse)	Forward: GTGCCCCGACTAACCGTTG
	Reverse: GTCGTTGAACTCCTCGGTCT
<i>Oct4</i> (mouse)	Forward: CGGAAGAGAAAGCGAACTAGC
	Reverse: ATTGGCGATGTGAGTGATCTG
18S	Forward: CGGCGACGACCCATTCGAAC
	Reverse: GAATCGAACCCCTGATTCCCCGTC

**Table S5. Primary antibody information for ChIP assays.**

<b>Antibody</b>	<b>Manufacture</b>	<b>Catalog #</b>
FoxO3	Novus Biologicals	NBP2-16521
H3K27me3	Novus Biologicals	NBP2-16840
H3K27ac	Novus Biologicals	NBP2-54615
Histone H3	Novus Biologicals	NB500-171
KDM6A	Novus Biologicals	NBP1-80628
p300	Novus Biologicals	NB100-616
SMARCD3	Santa Cruz	sc-101163

**Table S6. Oligonucleotide sequence of ChIP primers.**

<i>NANOG</i>	Forward: TGAGGCAAAGAGCATTGTTG
	Reverse: TGATGAAGGAATGGGCATAA
<i>SOX2</i>	Forward: GCAATGCTCTTTTAAGGGAAA
	Reverse: GACTTCCGGAATGACCAAAG
<i>KLF4</i>	Forward: CCCAGTGTCCCCTCTCTTCT
	Reverse: CGTGAATGTGGTGTTCGGTA