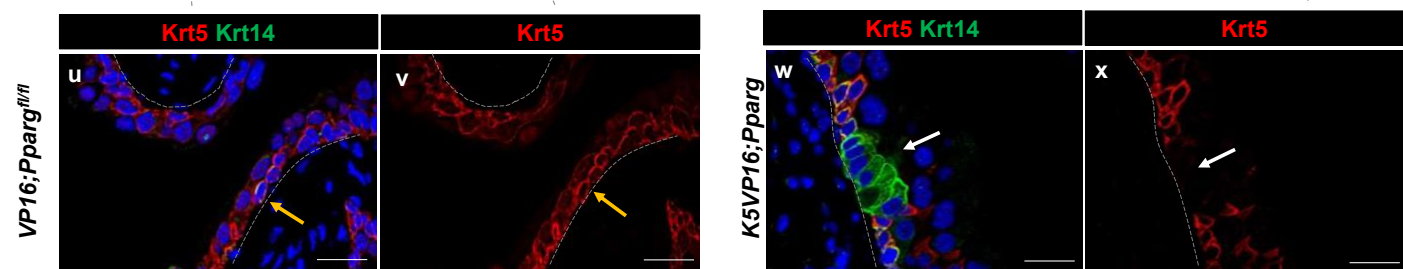
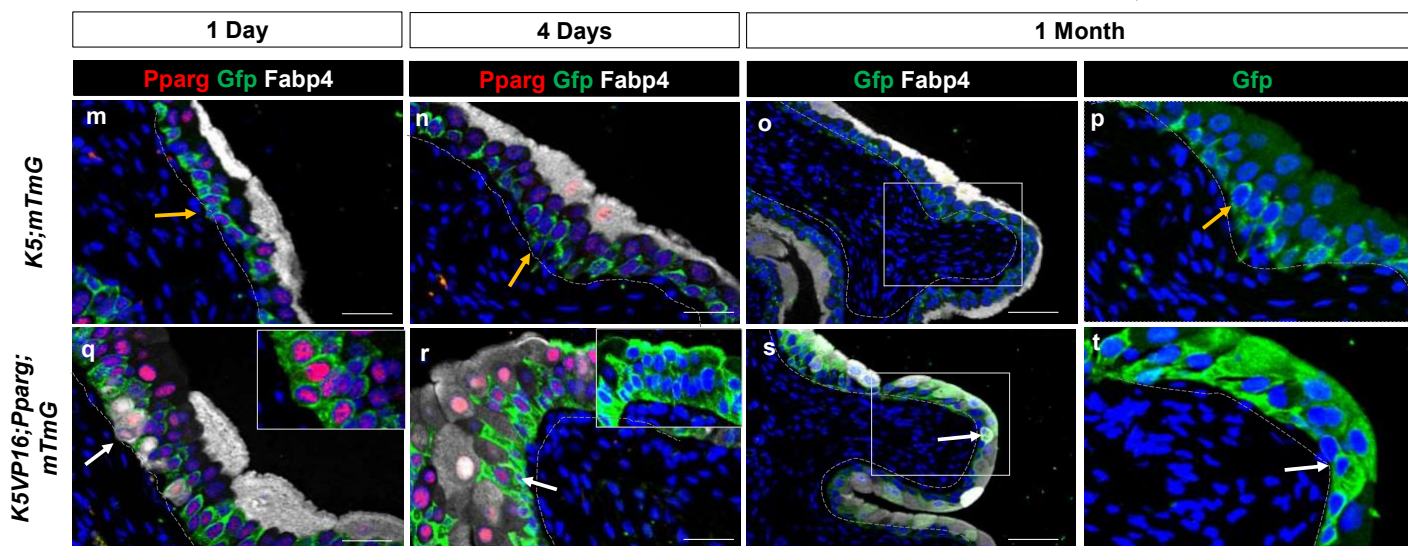
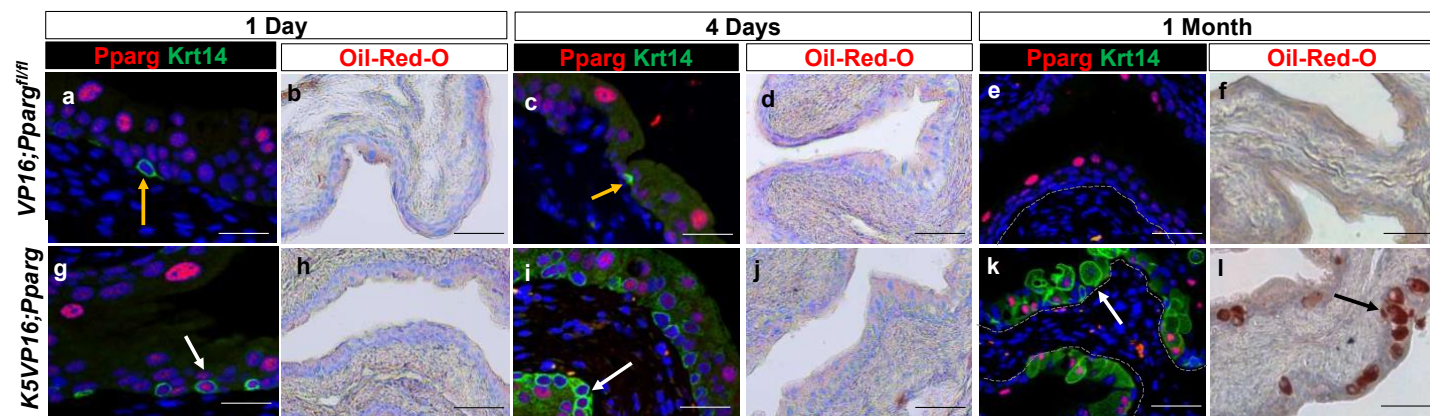


*Pparg* signalling controls bladder cancer subtype and drives immune exclusion

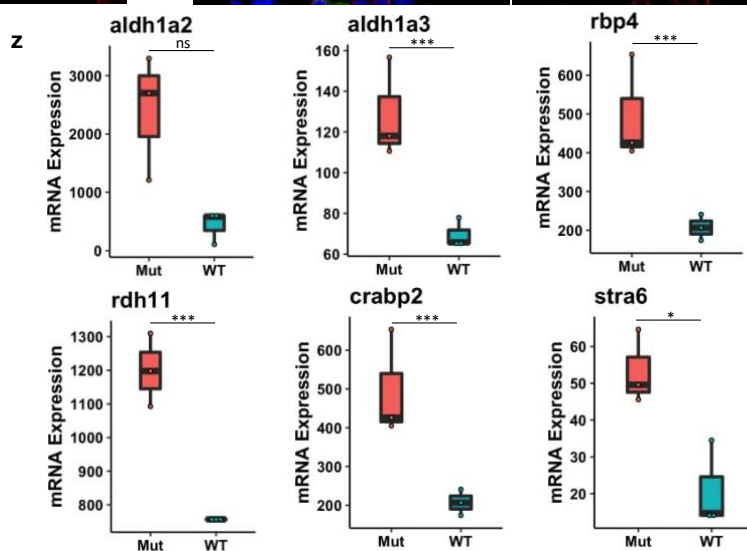
Tate, et al.

Supplementary Fig. 1



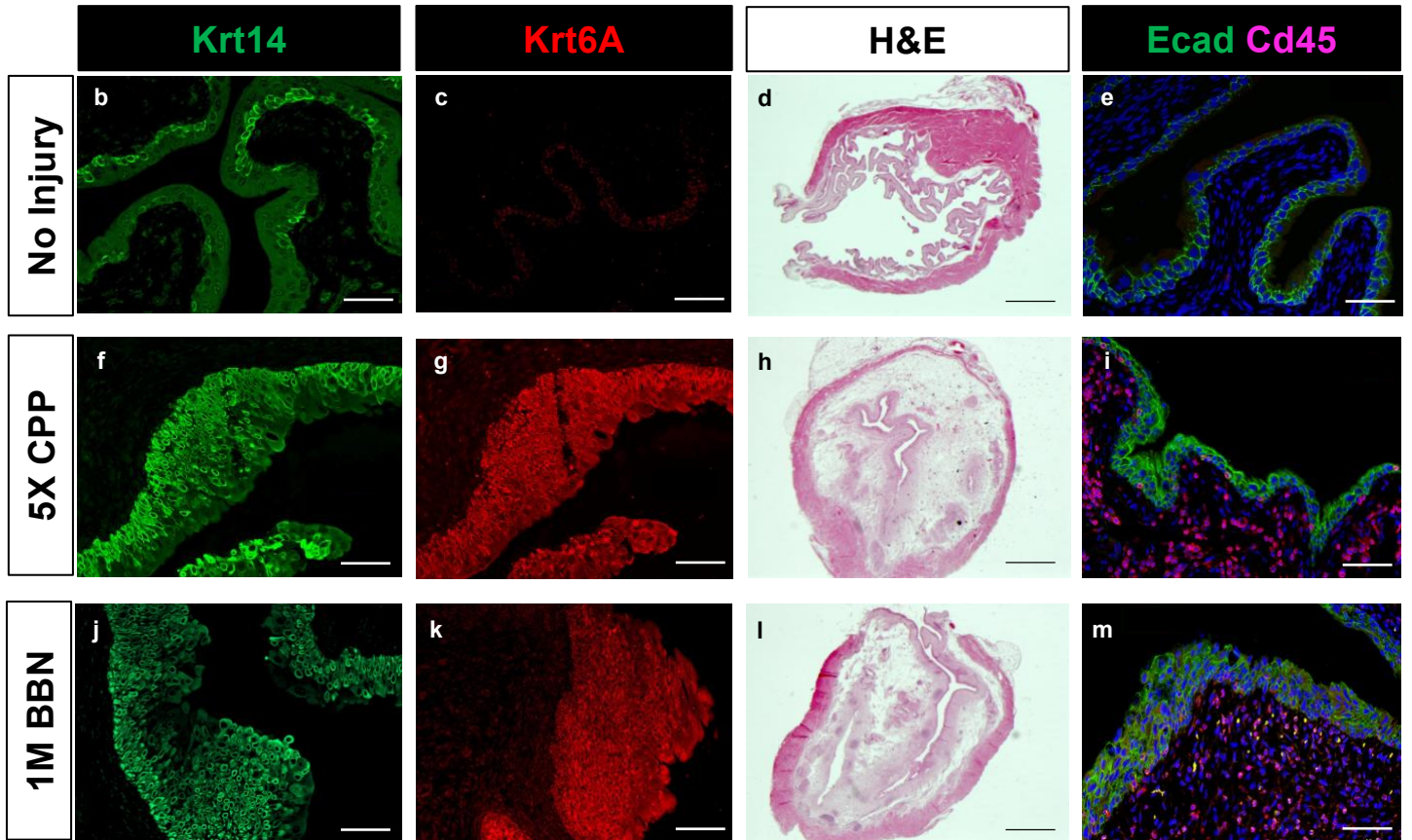
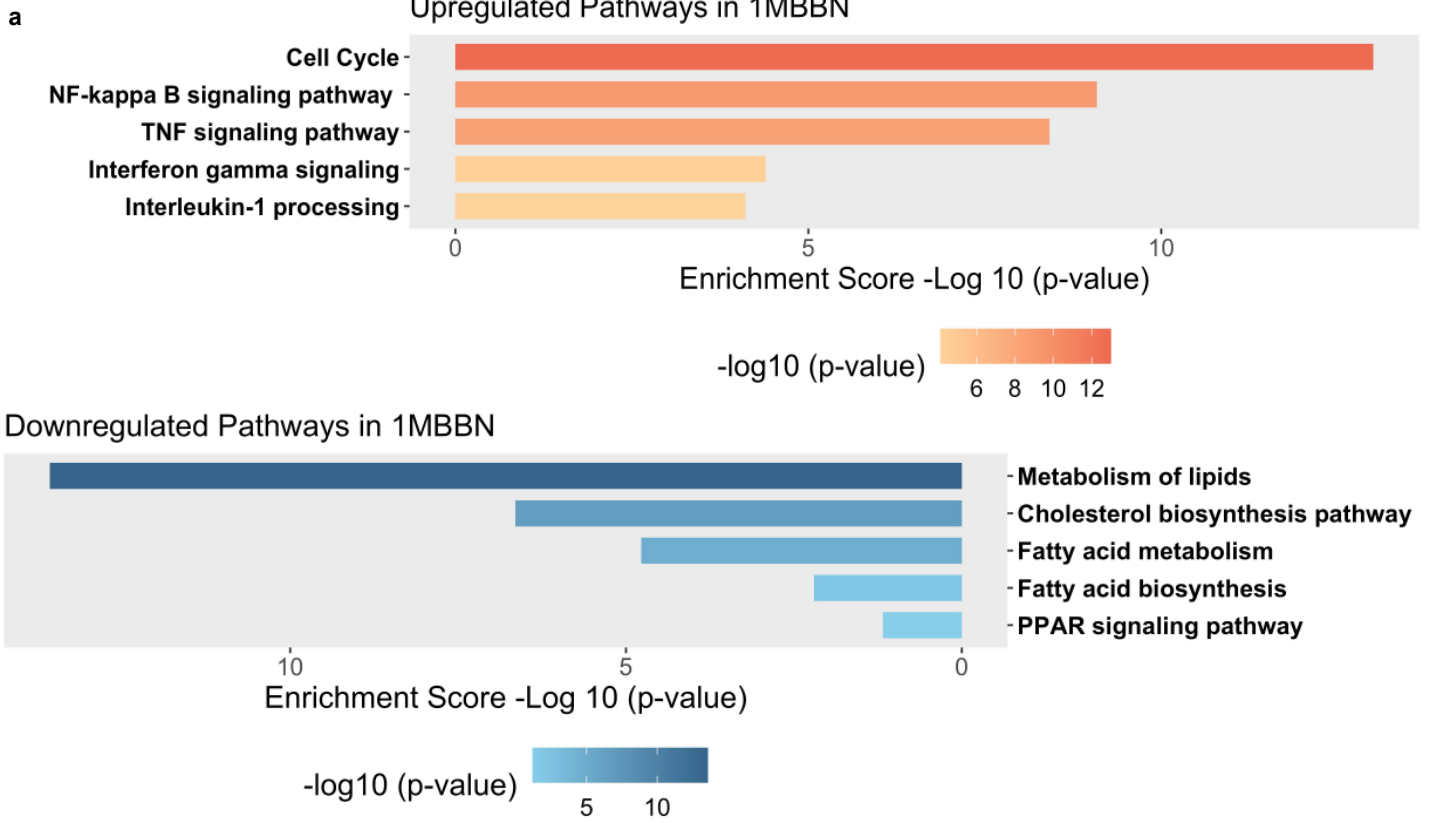
y

	Gene	FC	Pvalue
Luminal Markers	Upk1a	1.273411	0.101058
	Upk1b	1.771737	0.004768
	Upk2	1.831418	0.001319
	Upk3b	1.472945	0.021860
	Krt19	1.329701	0.041765
	Krt20	3.083968	2.04E-07
	Krt8	1.726677	0.000156
	Xbp1	1.424798	0.012264
	Cd24	1.618554	0.011801
	Snx31	1.57461	0.01846
	Fabp4	17.43987	2.90E-35
	Cldn8	1.370199	0.010296
Basal Markers	Krt5	0.829999	0.208742
	Col17a	0.303966	1.64E-10
	Trp63	0.579855	9.94E-07



**Supplementary Fig. 1. (a-l)** Lipid droplet formation one month after Tamoxifen induction in *VP16;Pparg<sup>fl/fl</sup>* mutants. Pparg and Krt14 expression 1 day after Tamoxifen induction in controls **(a)** and in *K5VP16;Pparg* mutants **(g)**. Oil-Red-O staining in controls **(b)** and in *K5VP16;Pparg* mutants **(h)** 1 day after Tamoxifen induction. Expression of Pparg and Krt14 in controls **(c)** and in *K5VP16;Pparg* mutants **(i)** 4 days after Tamoxifen induction. Oil-Red-O staining in controls **(d)** and in *K5VP16;Pparg* mutants **(j)** 4 days after Tamoxifen induction. Pparg and Krt14 expression in controls **(e)** and in *K5VP16;Pparg* mutants **(k)** 1 month after Tamoxifen induction. Oil-Red-O staining of controls **(f)** and mutants **(l)** 1 month after induction. Expression of Pparg, Gfp, and Fabp4 1 day after Tamoxifen induction in *K5;mTmG* controls **(m)** and in *K5VP16;Pparg;mTmG* mutants **(q)**. Expression of Pparg, Gfp, and Fabp4 4 days after Tamoxifen induction in *K5;mTmG* controls **(n)** and in *K5VP16;Pparg;mTmG* mutants **(r)**. Expression of Gfp and Fabp4 1 month after Tamoxifen induction in *K5;mTmG* controls **(o)** and *K5VP16;Pparg;mTmG* mutants **(s)**. Higher magnification image showing expression of Gfp 1 month after Tamoxifen induction in *K5;mTmG* controls **(p)** and *K5VP16;Pparg;mTmG* mutants **(t)**. Yellow arrows denote Gfp-positive control basal cells. White arrows denote Gfp-positive mutant cells expressing the transgene. Expression of Krt5 and Krt14 4 days after Tamoxifen induction in *VP16;Pparg<sup>fl/fl</sup>* controls **(u)** and *K5VP16;Pparg* mutants **(w)**. Expression of Krt5 4 days after Tamoxifen induction in controls **(v)** and in *K5VP16;Pparg* mutants **(x)** Yellow arrows denote control basal cells. White arrows denote mutant basal cells losing Krt5 expression. **(y)** Table showing the luminal/basal gene signature in based on RNA-seq analysis of urothelium from *K5;mTmG* controls and *K5VP16;Pparg* mutants 4 days post-Tamoxifen induction. **(z)** mRNA expression showing increased expression of genes involved in RA signaling in *K5VP16;Pparg;mTmG* mutants (n=3) compared to controls (n=3) 4 days after Tamoxifen induction. Box plots display minima, maxima, and interquartile range (IQR). Significance was calculated by one-sided Mann-Whitney U test (aldh1a2 p=0.67, aldh1a3 p=0.0059, rbp4 p=0.00061, rdh11 p=0.00032, crabp2 p=0.00032, stra6 p=0.036). ns=not significant; \*p≤0.05; \*\*0.05 ≤p≤0.01; \*\*\*0.01 ≤p≤0.0001. Scale bars, 50 μm.

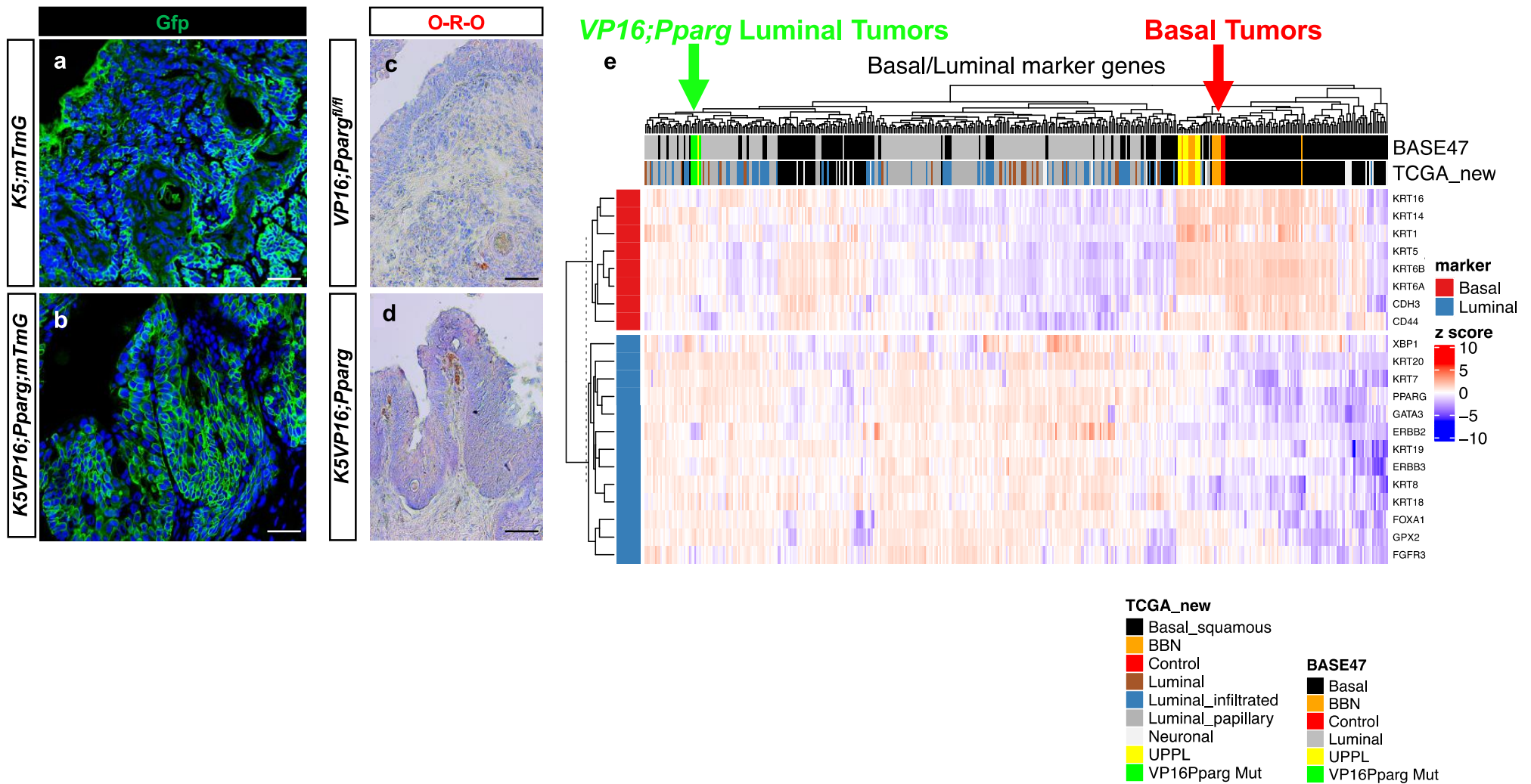
# Supplementary Fig. 2



**Supplementary Fig. 2. (a)** Up-regulated and down-regulated pathways from RNA-seq analysis of controls and bladders after 1MBBN treatment. p-values were calculated by hypergeometric test and corrected for multiple testing. **(b-m)** Repeated CPP treatment induces a similar activation program in the urothelium as that induced by 1M of BBN treatment. **(b,f,j)** Expression of Krt14 in controls **(b)**, animals treated with 5X CPP **(f)** and animals treated with BBN for 1 month **(j)**. **(c,g,k)** Expression of Krt6A in controls **(c)**, animals treated with 5X CPP **(g)**, and animals treated with BBN for 1 month **(k)**. H&E staining of bladders from controls **(d)**, from animals treated with 5X CPP **(h)**, and from animals treated with BBN for 1 month **(l)**. **(e,i,m)** E-cadherin and Cd45 expression in controls **(e)**, in animals treated with 5X CPP **(i)** and in animals after 1 month of BBN treatment **(m)**. Scale bars in **(b,c,e,f,g,l,j,k,m)** 50  $\mu\text{m}$ . Scale bars in **(d,h,l)** 200  $\mu\text{m}$ .

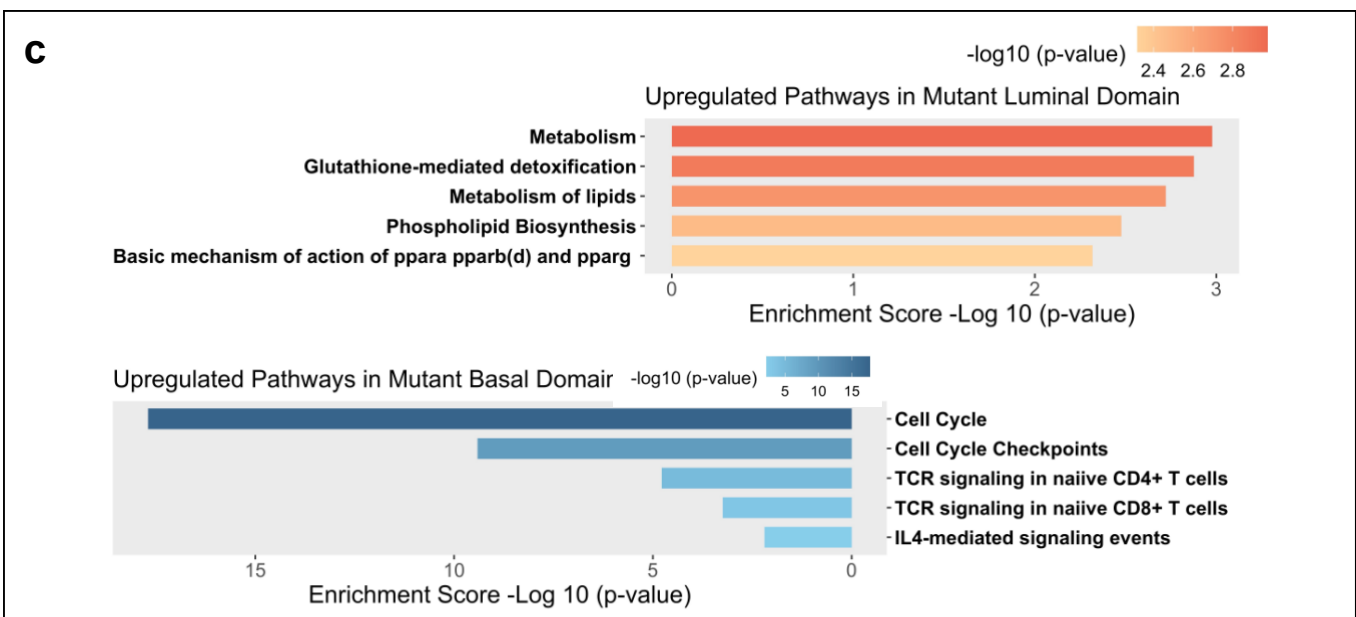
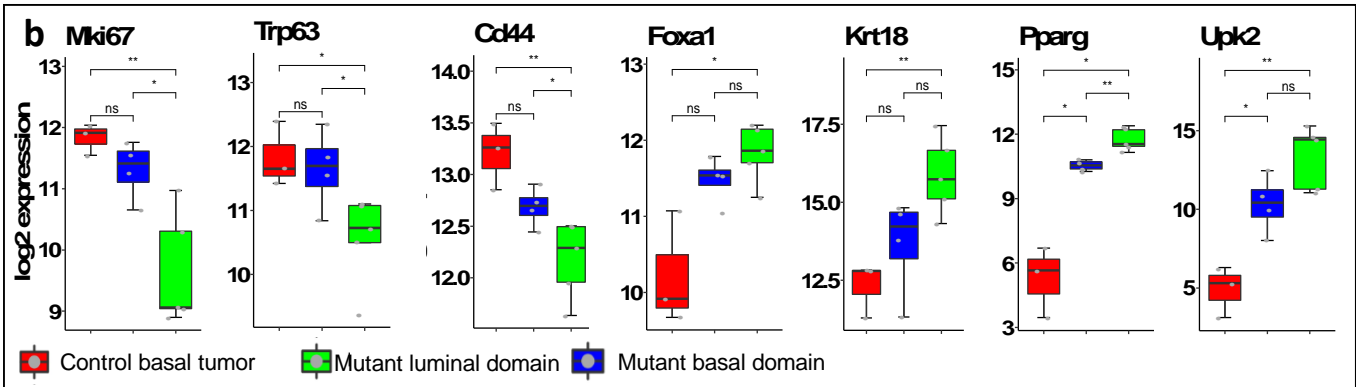
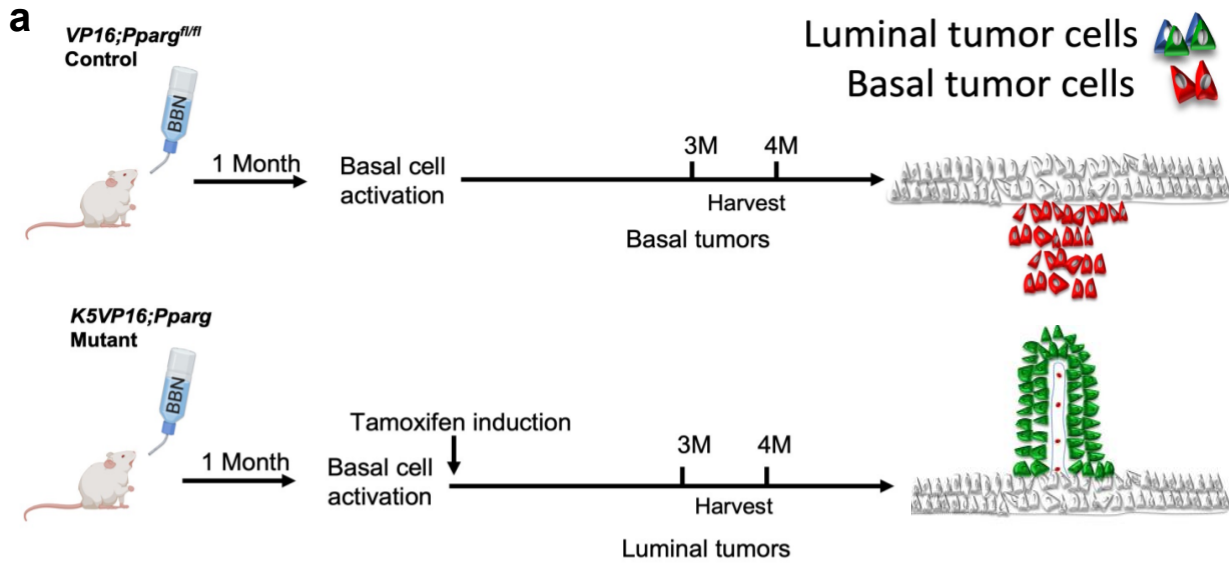


### Supplementary Figure 3



**Supplementary Fig. 3.** (a,b) Lineage tracing showing expression of Gfp-labeled basal cells and their daughters in a *K5;mTmG* control tumor (a) and in a *K5VP16;Pparg;mTmG* mutant tumor (b). Oil-Red-O staining of a *VP16;Pparg<sup>fl/fl</sup>* control tumor (c) and *K5VP16;Pparg* mutant tumor (d) 4 months after Tamoxifen induction. (e) Heatmap showing co-clustering of tumors from *K5;mTmG* controls and *K5VP16;Pparg* mutant tumors with luminal and basal tumors respectively. Data includes the TCGA BLCA data set, BASE47 base classifier data set, BBN treated tumors and UPPL tumors. Scale bars 50 $\mu$ m.

# Supplementary Fig. 4

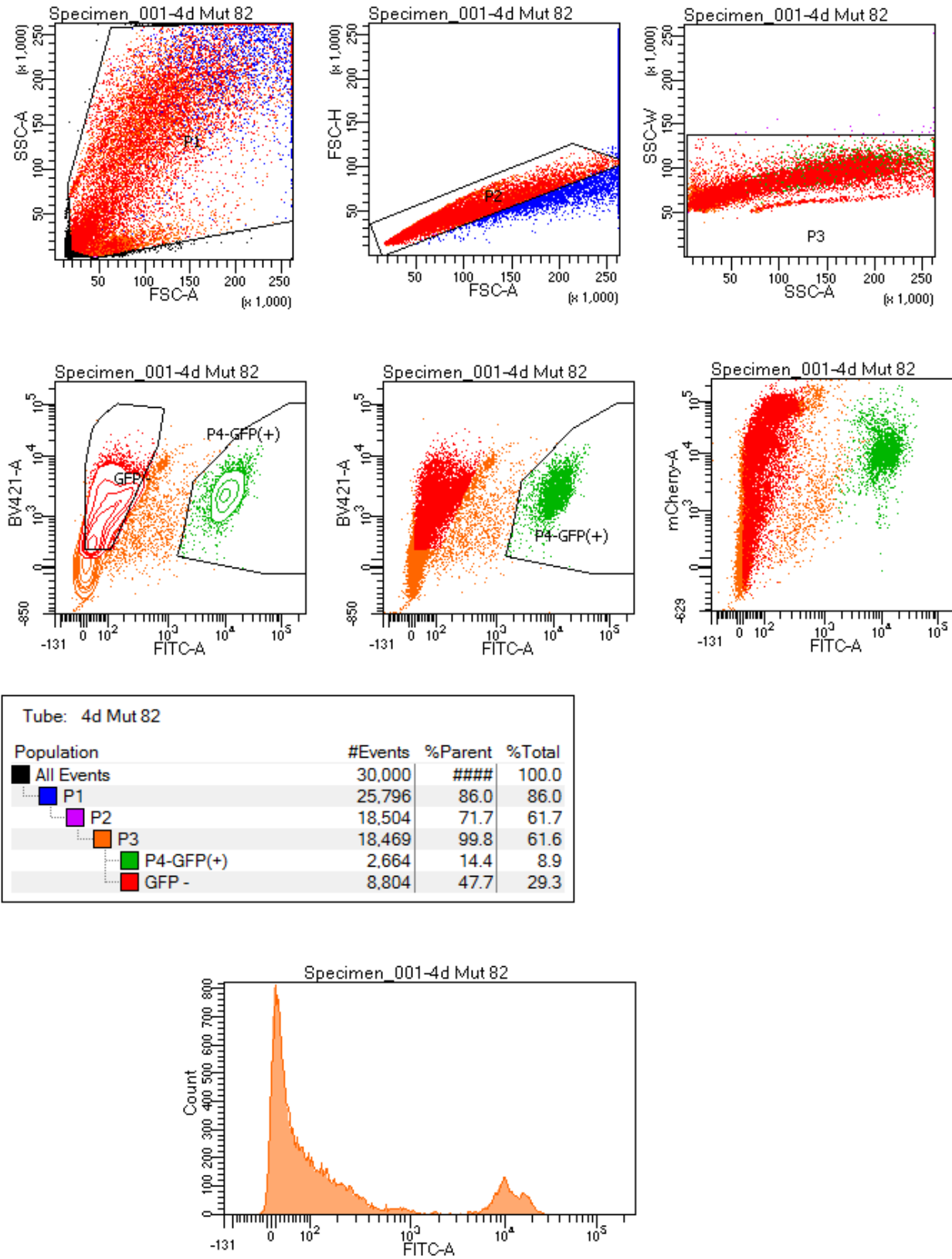


**Supplementary Fig. 4. (a)** Schematic of luminal and basal tumor formation in *K5;mTmG* controls and *K5VP16;Pparg;mTmG* mutants, respectively. **(b)** Expression levels mRNA encoding *Mki67*, *Trp63*, *Cd44*, *Foxa1*, *Krt18*, *Pparg*, and *Upk2* from *K5;mTmG* in basal control tumors (n=3) compared to luminal (n=4) and basal domains (n=4) of *K5VP16;Pparg;mTmG* mutant tumors at 4 months. Box plots display minima, maxima, and interquartile range (IQR). Significance was calculated by Mann-Whitney U test. ns=not significant; \* $p \leq 0.05$ ; \*\* $0.05 \leq p \leq 0.01$ . **(c)** Upregulated and downregulated signaling pathways based on RNA-seq analysis of *K5VP16;Pparg* mutant luminal domain compared to the *K5VP16;Pparg* mutant basal domain at 4 months. p-values were calculated by hypergeometric test and corrected for multiple testing.



# Supplementary Fig. 5

a



**Supplementary Fig. 5. (a)** Gating strategy used to collect GFP+ cells in homeostatic RNAseq.

**Supplementary Table 1. Key resources used in the study**

<b>ANTIBODIES</b>	<b>SOURCE</b>	<b>IDENTIFIER</b>	<b>CLONE</b>	<b>DILUTION</b>
Chicken Polyclonal Anti-Keratin 14	Biolegend	Cat#906001	N/A	1:400
Chicken Polyclonal Anti-Keratin 5	Biolegend	Cat#905901	N/A	1:400
Rabbit Polyclonal Anti-Krt6A	LSBio	Cat#LS-B12036-100	N/A	1:2000
Mouse Monoclonal Anti-Cytokeratin 20	AgilentDako	Cat# M701929-2		1:200
Rabbit Polyclonal Anti-PPARG	Cell Signaling Technology	Cat# 2435	N/A	1:200
Goat Polyclonal Anti-FABP4	R&D Systems	Cat# AF1443	N/A	1:1000
Mouse Monoclonal Anti-FOXA1	Seven Hills Bioreagents	Cat# WMAB-2F83	2F83	1:1000
Mouse Monoclonal Anti-p21/CDKN1A/WAF1	LSBio	Cat# LS-C389956	HJ21	1:200
Rabbit Polyclonal Anti-p63	GeneTex	Cat#GTX102425	N/A	1:300
Goat Polyclonal Anti-p63	R&D Systems	Cat#AF1916	N/A	1:200
Rat Monoclonal Anti-CD45	BD Bioscience	Cat#550539	30F11	1:100
Rabbit Polyclonal Anti-Cytokeratin 18	Abcam	Cat#ab52948	N/A	1:500
Rabbit Polyclonal Anti-Ki67	Abcam	Cat#ab15580	N/A	1:200
Chicken Polyclonal Anti-GFP	Aves Labs	Cat#GFP1020	N/A	1:300
Goat Polyclonal Anti-E-Cadherin	R&D Systems	Cat#AF748	N/A	1:400

Rabbit Polyclonal NFkB p65	Abcam	Cat#AB19870	N/A	1:300
Rabbit Polyclonal Laminin	Sigma	Cat#L9393	N/A	1:100
Mouse Monoclonal SMA-CY3	Sigma	Cat#C6198	1A4	1:500
Alexa Fluor 488 Donkey Anti-Rabbit IgG	Jackson Immunoresearch	Cat#711-545-152	N/A	1:700
Alexa Fluor 488 Donkey Anti-Mouse IgG	Jackson Immunoresearch	Cat#711-545-150	N/A	1:700
Alexa Fluor 488 Donkey Anti-Chicken IgG	Jackson Immunoresearch	Cat#703-545-155	N/A	1:700
Alexa Fluor 488 Donkey Anti-Goat IgG	Jackson Immunoresearch	Cat#705-545-003	N/A	1:700
Cy3 Donkey Anti-Rabbit IgG	Jackson Immunoresearch	Cat#711-165-152	N/A	1:700
Alexa Fluor 594 Donkey Anti-Mouse IgG	Jackson Immunoresearch	Cat#715-585-151	N/A	1:700
Alexa Fluor 594 Donkey Anti-Chicken IgG	Jackson Immunoresearch	Cat#703-585-155	N/A	1:700
Alexa Fluor 594 Donkey Anti-Goat IgG	Jackson Immunoresearch	Cat#705-585-147	N/A	1:700
Alexa Fluor 647 Donkey Anti-Mouse IgG	Jackson Immunoresearch	Cat#715-605-150	N/A	1:400
Alexa Fluor 647 Donkey Anti-Rabbit IgG	Jackson Immunoresearch	Cat#711-605-152	N/A	1:400
Alexa Fluor 647 Donkey Anti-Chicken IgG	Jackson Immunoresearch	Cat#703-605-155	N/A	1:400
Alexa Fluor 647 Donkey Anti-Goat IgG	Jackson Immunoresearch	Cat#705-605-003	N/A	1:400

REAGENT or RESOURCE	SOURCE	IDENTIFIER
<b>Biological samples</b>		
Human bladder tumors samples	Columbia University Irving Medical Center	N/A

<b>Chemicals, Peptides, and Recombinant proteins</b>		
N-butyl-N-(4-hydroxybutyl) nitrosamine	Sigma	Cat#B8061-1G
Hank's Balanced Salt Solution (HBSS)	ThermoFisher Scientific	Cat#14170-112
Bovine Serum Albumin	Sigma	Cat#A2058
<i>Bacillus licheniformis</i> protease	Sigma	Cat#P5459
CaCl <sub>2</sub>	Sigma	Cat#21115
DNase I recombinant	Sigma	Cat#4716728001
RLT lysis buffer	Qiagen	Cat#1015750
Antigen unmasking solution	Vector Labs	Cat#H3300
Citrate buffer	Thermo Fisher Scientific	Cat#AP-9003-500
Horse Serum, heat inactivated	Gibco	Cat#26050070
Tween80	Sigma	Cat#P1754
DMEM/F12	Thermo Fisher Scientific	Cat#H7904
4-Hydroxytamoxifen	Sigma	Cat#11320033
Tamoxifen	Sigma	Cat#T5648
Histogene Staining Solution	TherFisher Scientific	Cat#KIT0425
<b>Critical commercial assays</b>		
SMART-Seq v4 Ultra Low Input RNA Kit for Sequencing	TaKaRa	Cat#634889
<b>Deposited data</b>		

Raw RNA-seq files	This paper	GEO: GSE172656
<b>Experimental Models: Organisms/Strains</b>		
Mouse: <i>VP16;Pparg</i>	This paper	
Mouse: <i>mTmGfl/fl</i> (Gt(ROSA)26Sortm4(ACTB-tdTomato,-EGFP)Luo/J)	Jackson Laboratory	Cat#007576
Mouse: FVB.Cg-Tg(KRT5-cre/ERT2)2lpc/JelJ (K5Cre <sup>ERT2</sup> )	D. Metzger P. Chambon	N/A
<b>Oligonucleotides</b>		
Primer: <i>VP16;Pparg</i>  Mutant forward: 5'-CTGCATTCTAGTTGTGGTTTGTCCA-3'  Mutant reverse:5'-ATCGGTAAACATCTGCTCAAACCTCG-3'  Wild type forward 5'-CCCAAAGTCGCTCTGAGTTGTTATC-3'  Wild type reverse 5'-AACTCGGGTGAGCATGTCTTTAATC-3'	This paper	N/A
Primer: <i>Krt5Cre<sup>ERT2</sup></i>  Forward: 5'-ATTTGCCTGCATTACCGGTC-3'  Reverse: 5'-ATCAACGTTTTGTTTTCGGA-3'	Indira et al, 1999	NA
Primer: <i>mTmG</i>	Muzumdar et al, 2007	NA



Common 5'-CTCTGCTGCCTCCTGGCTTCT-3'		
Mutant 5'-TCAATGGGCGGGGGTCGTT-3'		
Wild type 5'-CGAGGCGGATCACAAGCAATA-3'		
<b>Software and Algorithms</b>		
R version 4.0.4	R Core Team, 2016	<a href="http://www.r-project.org/">http://www.r-project.org/</a>
DESeq2	Love et al., 2014	<a href="https://github.com/mikelove/DESeq2">https://github.com/mikelove/DESeq2</a>
Gene Set Enrichment Analysis, v4.1.0	Subramanian et al., 2005	<a href="https://www.gsea-msigdb.org/gsea/index.jsp">https://www.gsea-msigdb.org/gsea/index.jsp</a>
<b>Other</b>		
VEVO 3100Ultrasound Imaging System	FUJIFILM VisualSonics	N/A
Zeiss AxioObserver.Z1 inverted microscope for laser capture microdissection	Zeiss	N/A