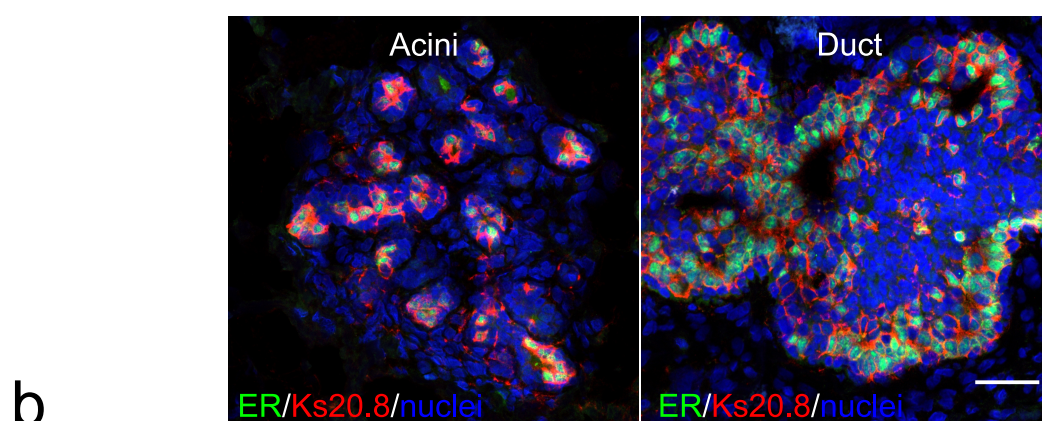
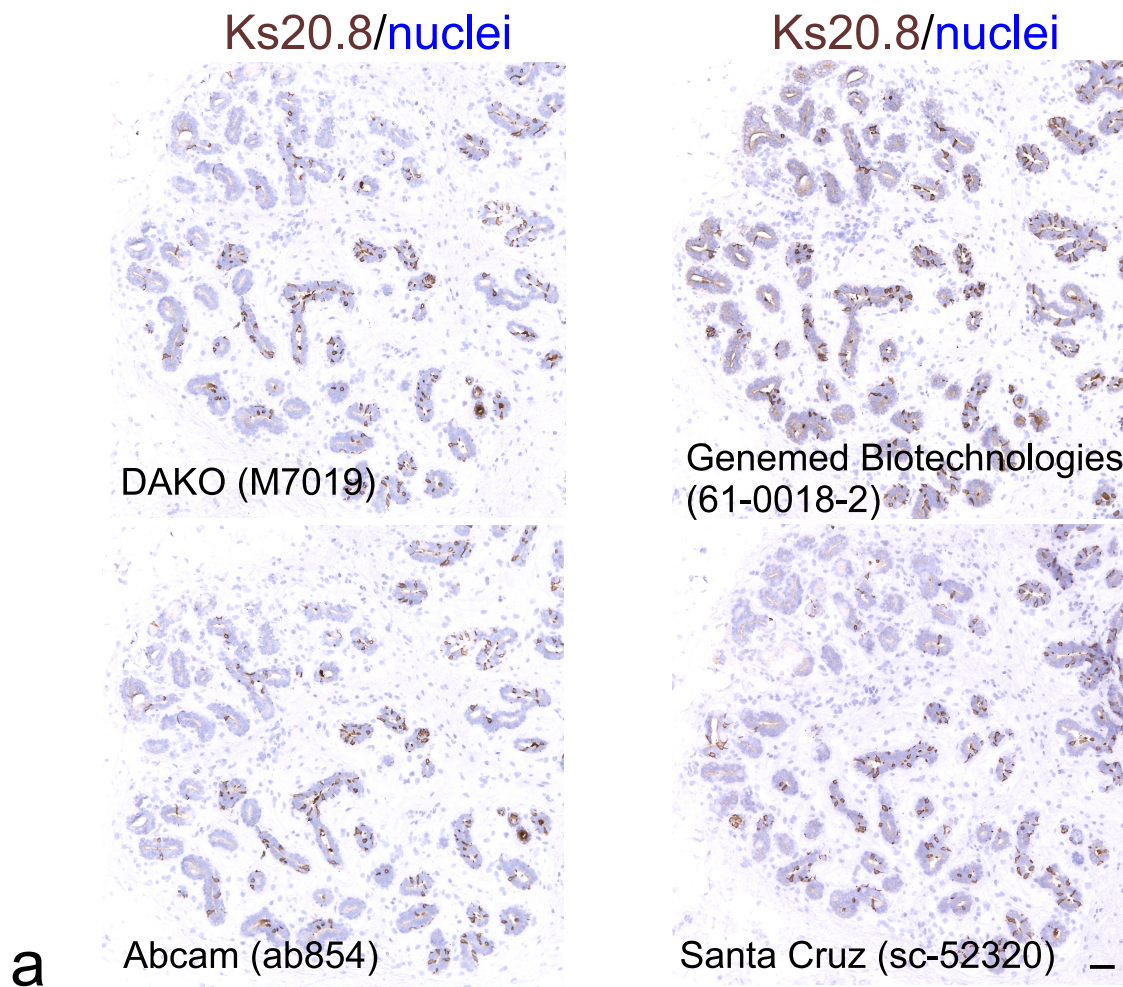
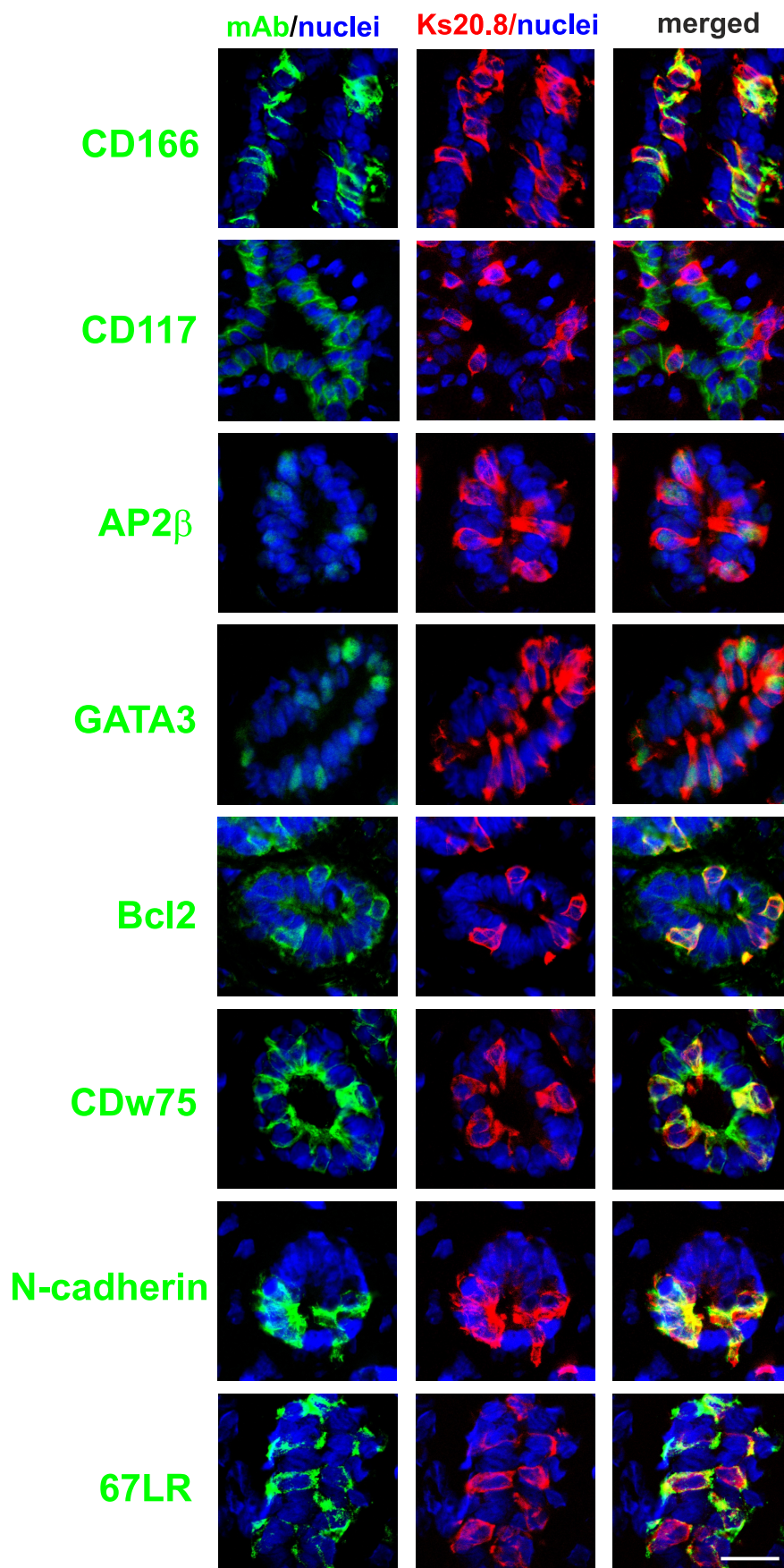


**Supplementary Figure 1. Ks20.8 staining is independent of antibody supplier and ER staining when detected by immunofluorescence reflects ER/PR staining**



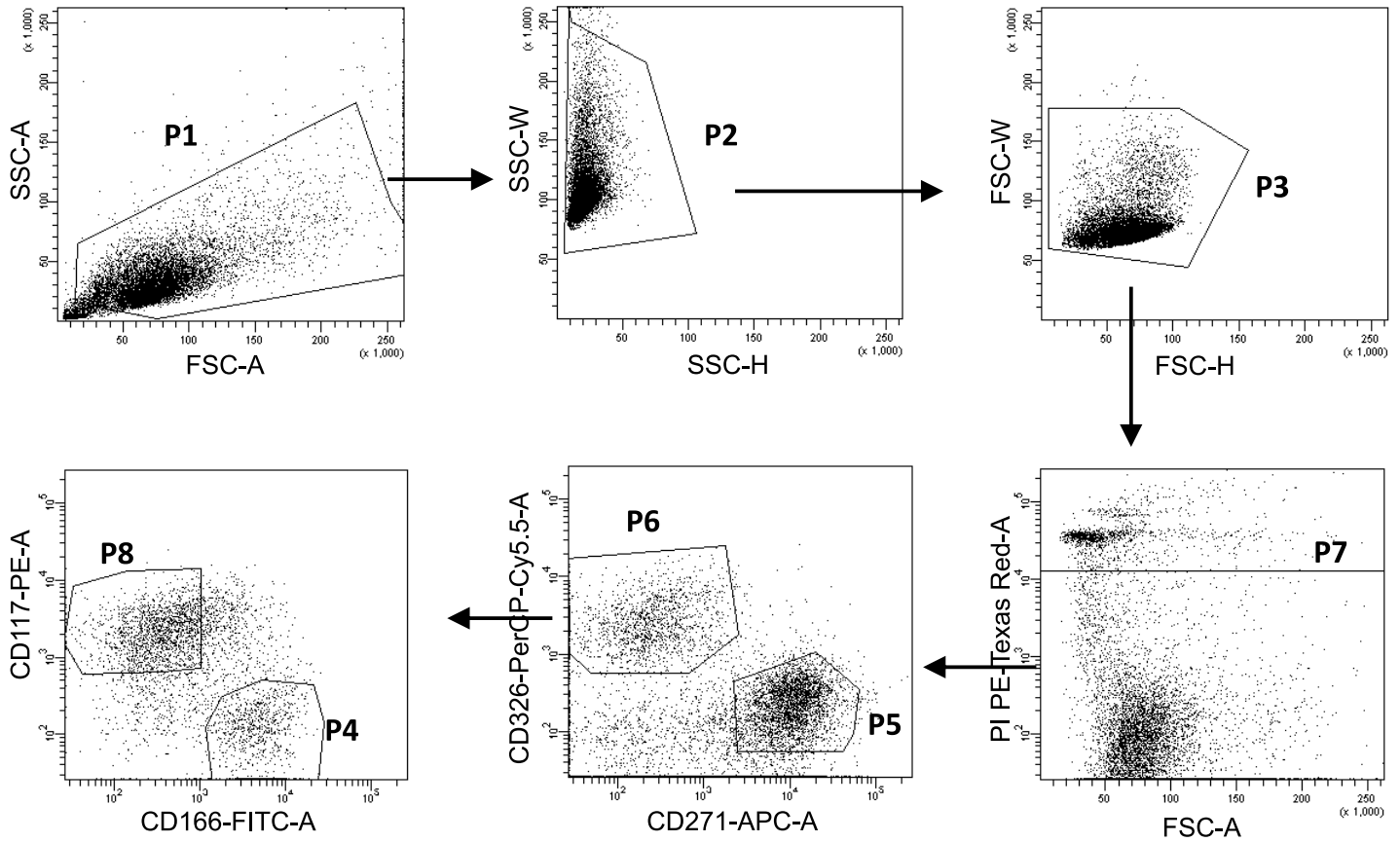
(a) Micrographs of serially sectioned human breast tissue immunoperoxidase stained with Ks20.8 purchased from four different suppliers (DAKO, Abcam, Genemed and Santa Cruz). All antibodies showed the same staining pattern. Nuclei are counterstained with hematoxylin. Bar = 50 $\mu$ m. (b) Multicolor imaging of cryostat sections from the same normal human breast tissue as used in figure 1b and stained with estrogen receptor (SP1; green), Ks20.8 (red), and nuclei (blue). ER protein localizes to essentially the same cells as Ks 20.8 and reflects ER/PR staining in tissues with high expression. Bar=25 $\mu$ m.

**Supplementary Figure 2. Ks20.8 positive cells exhibit an elaborate differentiation program**



Multicolor imaging of cryostat sections from normal human breast tissue stained with selected mAbs (green) from a library, Ks20.8 (red), and nuclei (blue). Note that selected markers except CD117 are colocalized with Ks20.8. Bar=25 $\mu$ m.

### Supplementary Figure 3. Representative FACS gating strategy of sorted cells

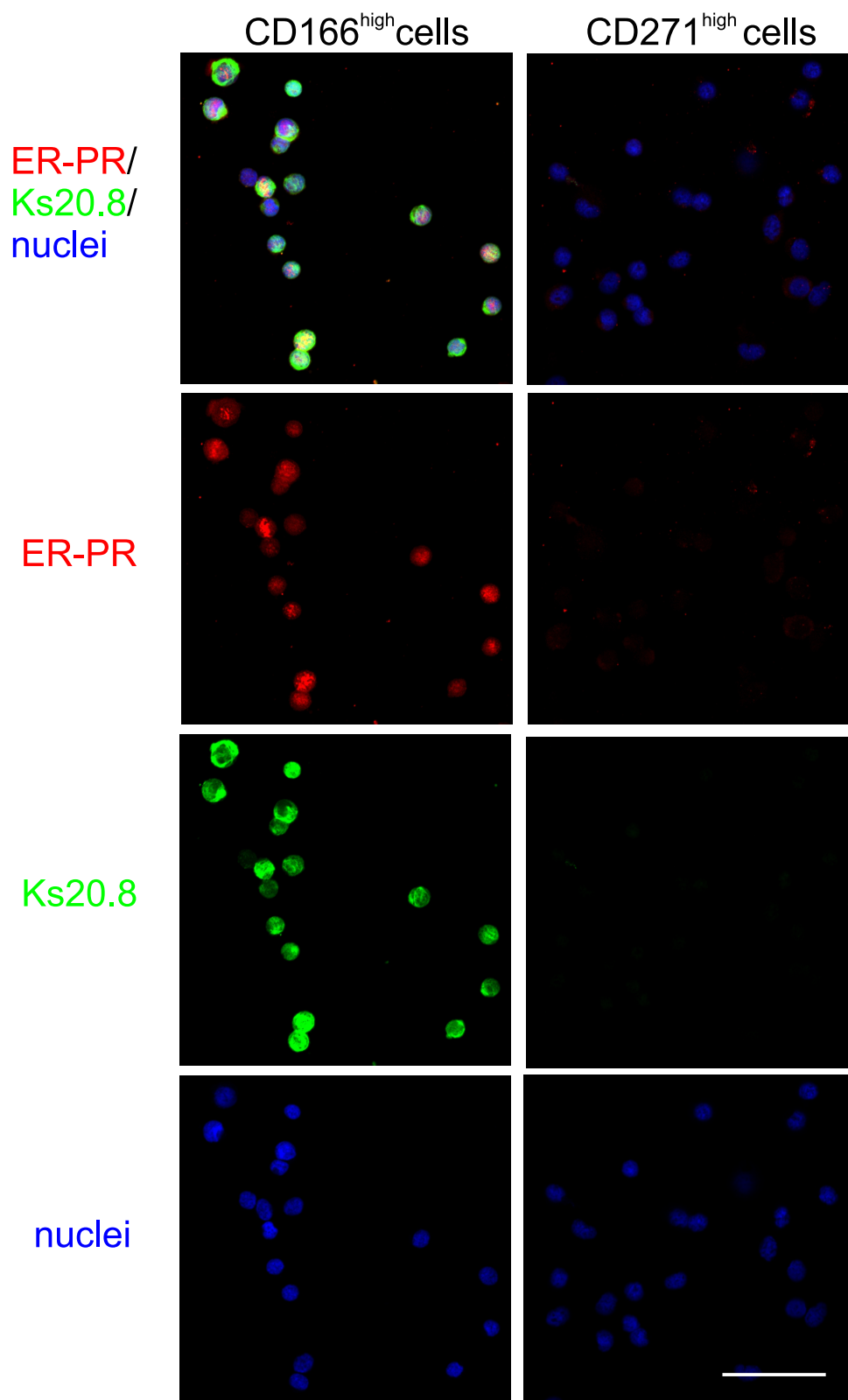


Tube: P790\

Population	#Events	%Parent	%Total
All Events	50,000	100.0	100.0
P1	44,803	89.6	89.6
P2	44,353	99.0	88.7
P3	44,284	99.8	88.6
P7	38,063	86.0	76.1
P5	15,144	39.8	30.3
P6	7,106	18.7	14.2
P4	1,737	24.4	3.5
P8	3,319	46.7	6.6

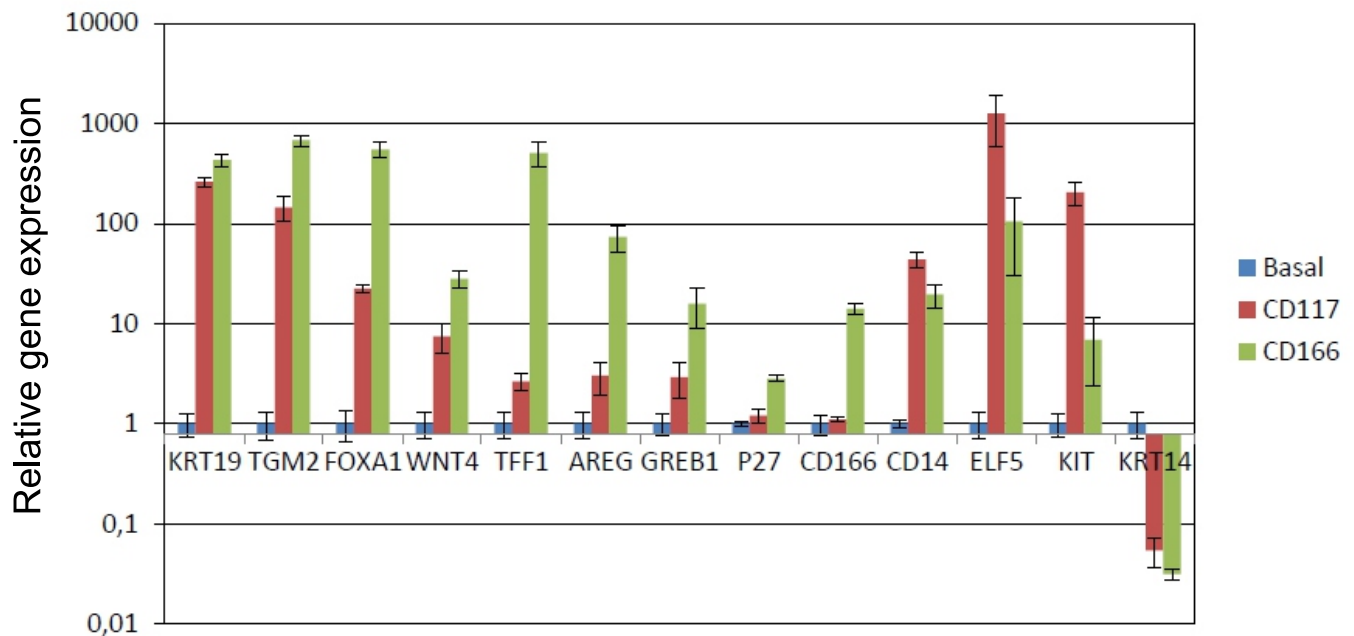
Cells were gated on forward scatter (FSC) and side scatter (SSC) to remove debris and select for single cells. Propidium iodide was used to select for live cells which were then further gated for luminal CD326<sup>high</sup> cells (P6) and basal CD271<sup>high</sup> cells (P5). CD326<sup>high</sup> luminal cells were further sub-gated as non-overlapping gates of CD166<sup>high</sup> (P4) and CD117<sup>high</sup> (P8).

**Supplementary Figure 4. Hormone receptor expression is restricted essentially to the Ks20.8<sup>pos</sup> compartment**



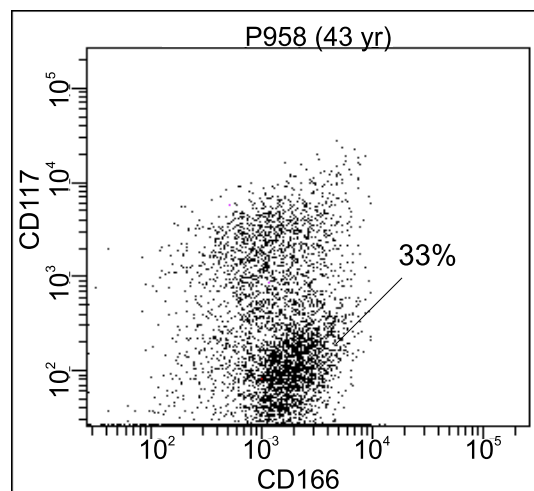
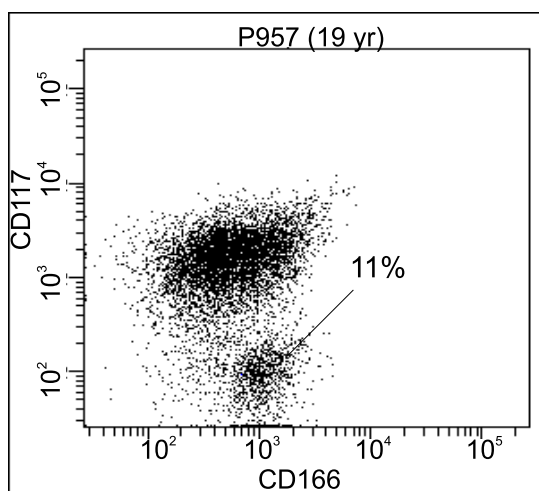
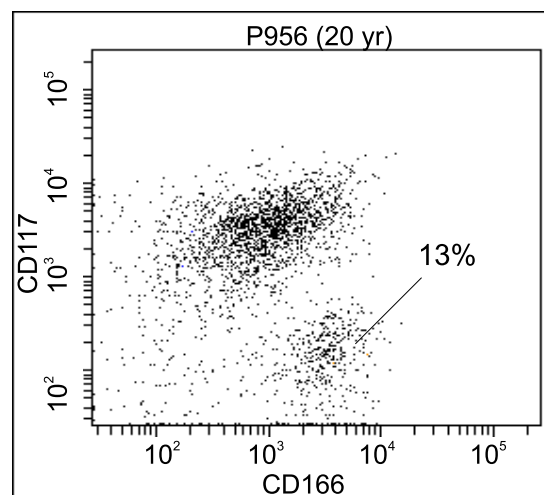
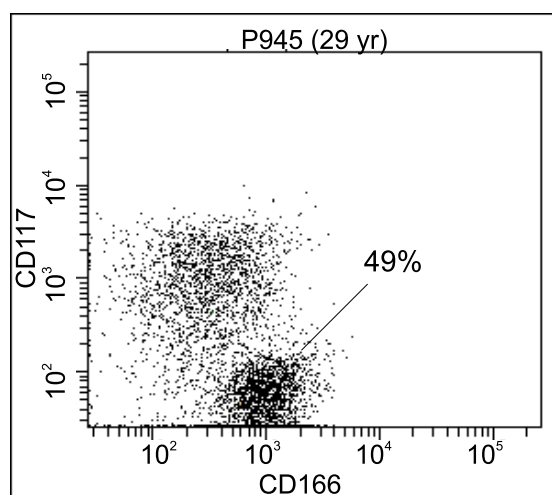
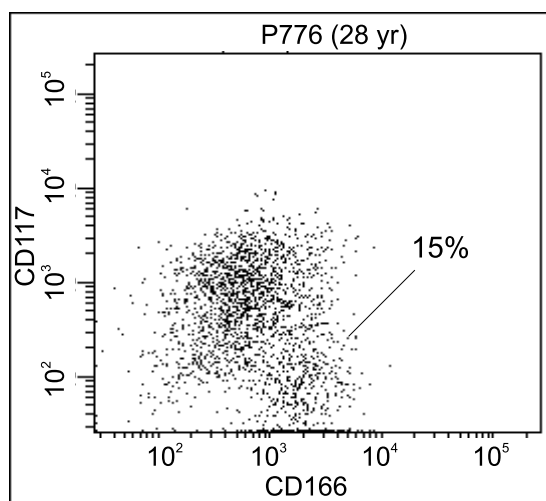
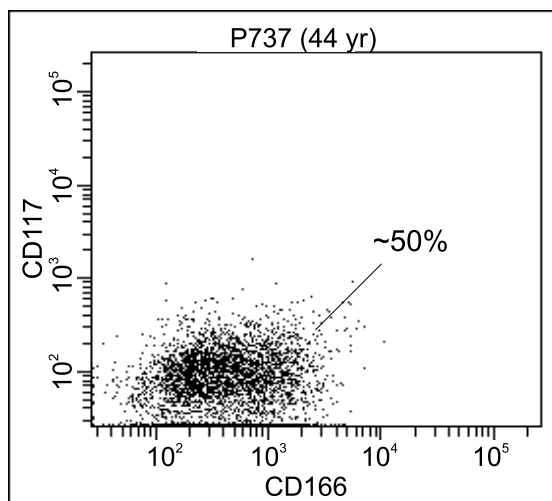
Multicolor imaging of smears of normal human CD166<sup>high</sup>/CD117<sup>low</sup> (CD166<sup>high</sup> cells) and basal cells (CD271<sup>high</sup> cells) purified by FACS and stained with ER-PR (SP1, SP2; red), Ks20.8 (green) and nuclei (blue). The vast majority of ER-PR<sup>pos</sup> cells are contained within the Ks20.8<sup>pos</sup> compartment. Bar=50  $\mu$ m.

## Supplementary Figure 5. Profiling of basal cells, CD117<sup>high</sup> and CD166<sup>high</sup> cells



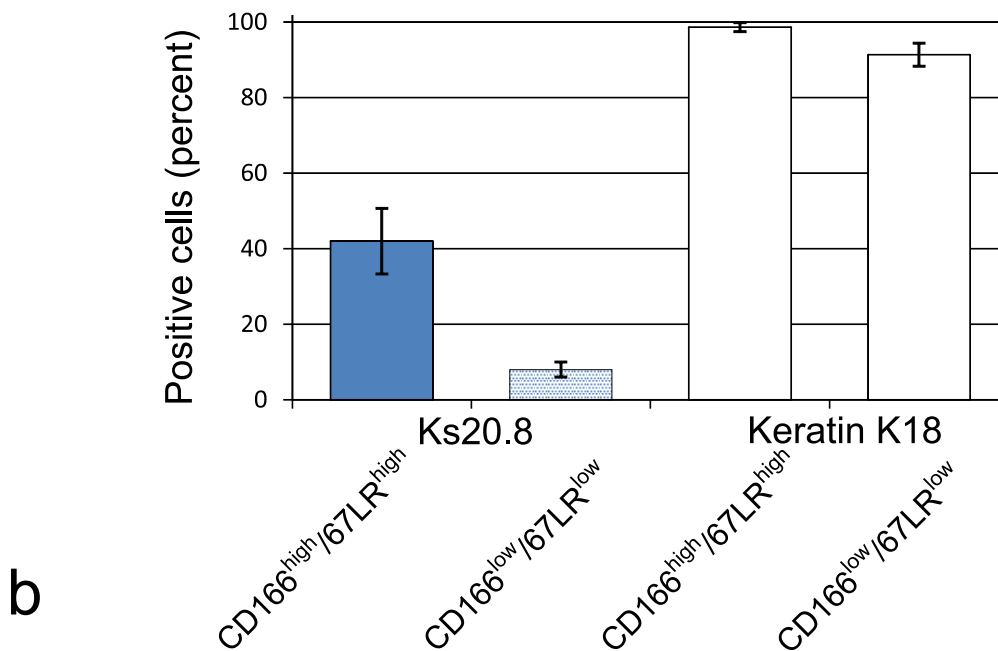
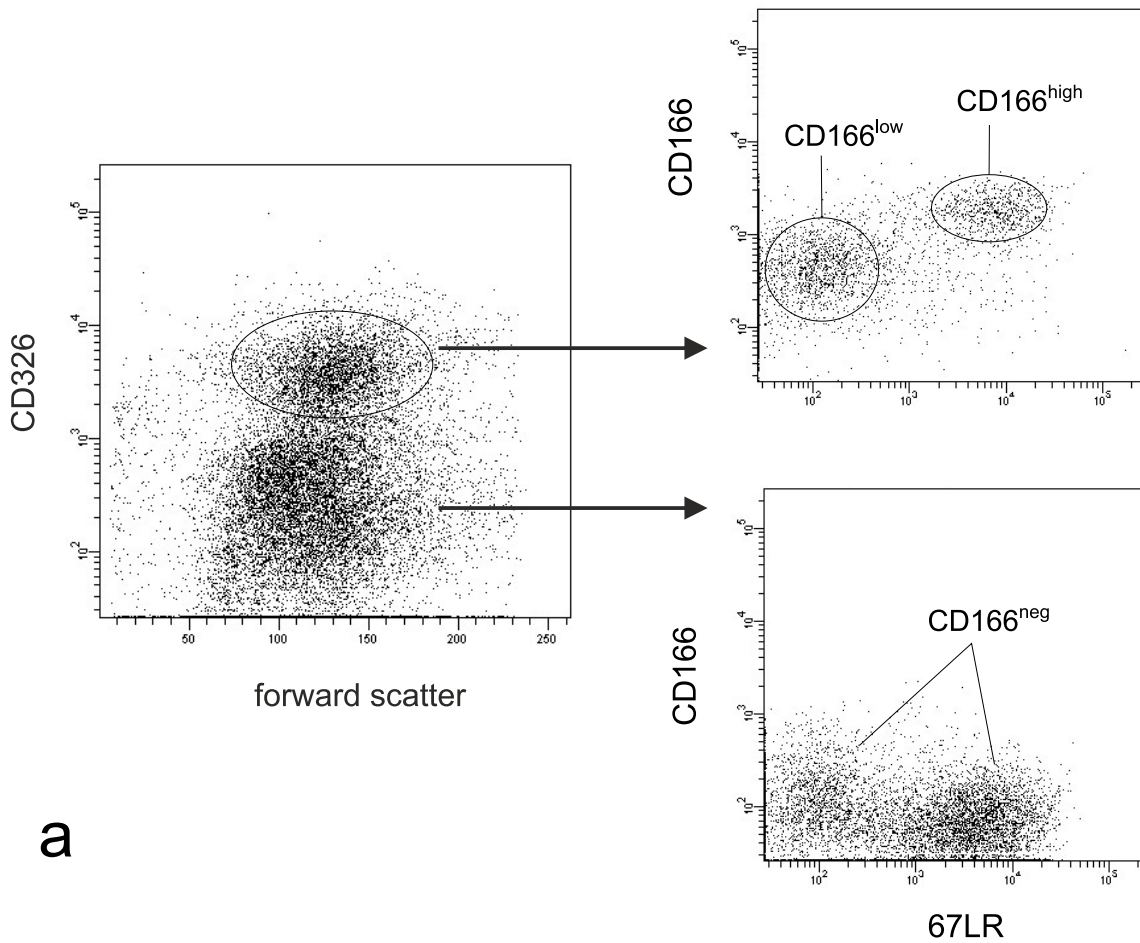
Bar graph representing qRT-PCR analysis of additional gene expression profiles in sorted basal, CD117<sup>high</sup>, and CD166<sup>high</sup> cells from three different biopsies. Each gene expression level was normalized by genomic mean of four reference gene expressions and then compared to normalized gene expression levels in basal cells. Error bars represent SEM of three biological samples which were each run in triplicate by qPCR.

**Supplementary Figure 6. Five out of six biopsies show a distinct CD166<sup>high</sup> population**



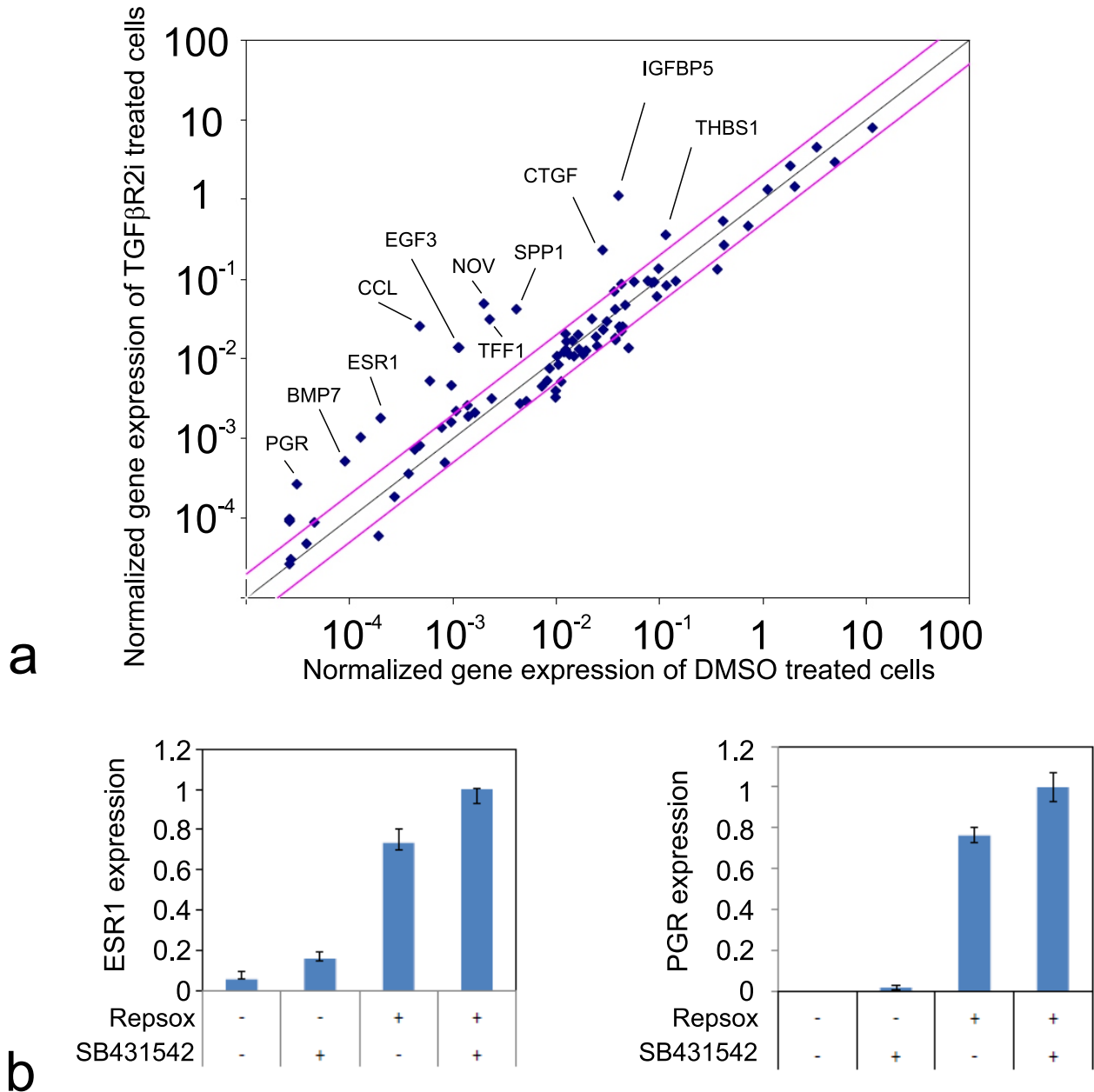
Representative FACS diagrams of uncultured HBEC from six biopsies sorted with CD166 and CD117.

**Supplementary Figure 7. Representative FACS profiles illustrating the separation of 67LR<sup>high</sup>/CD166<sup>high</sup> cells from uncultured HBECs**



(a) A single cell suspension of primary breast epithelial cells stained with CD326, CD166 and 67LR followed by flow cytometry. Ks20.8<sup>pos</sup> HBEC sorted with the CD166<sup>high</sup>/67LR<sup>high</sup> gate. (b) Smears of FACS sorted cells stained with Ks20.8 (left) or keratin K18 (right). Frequency of positive cells (n = 3x50 cells per slide ±SD).

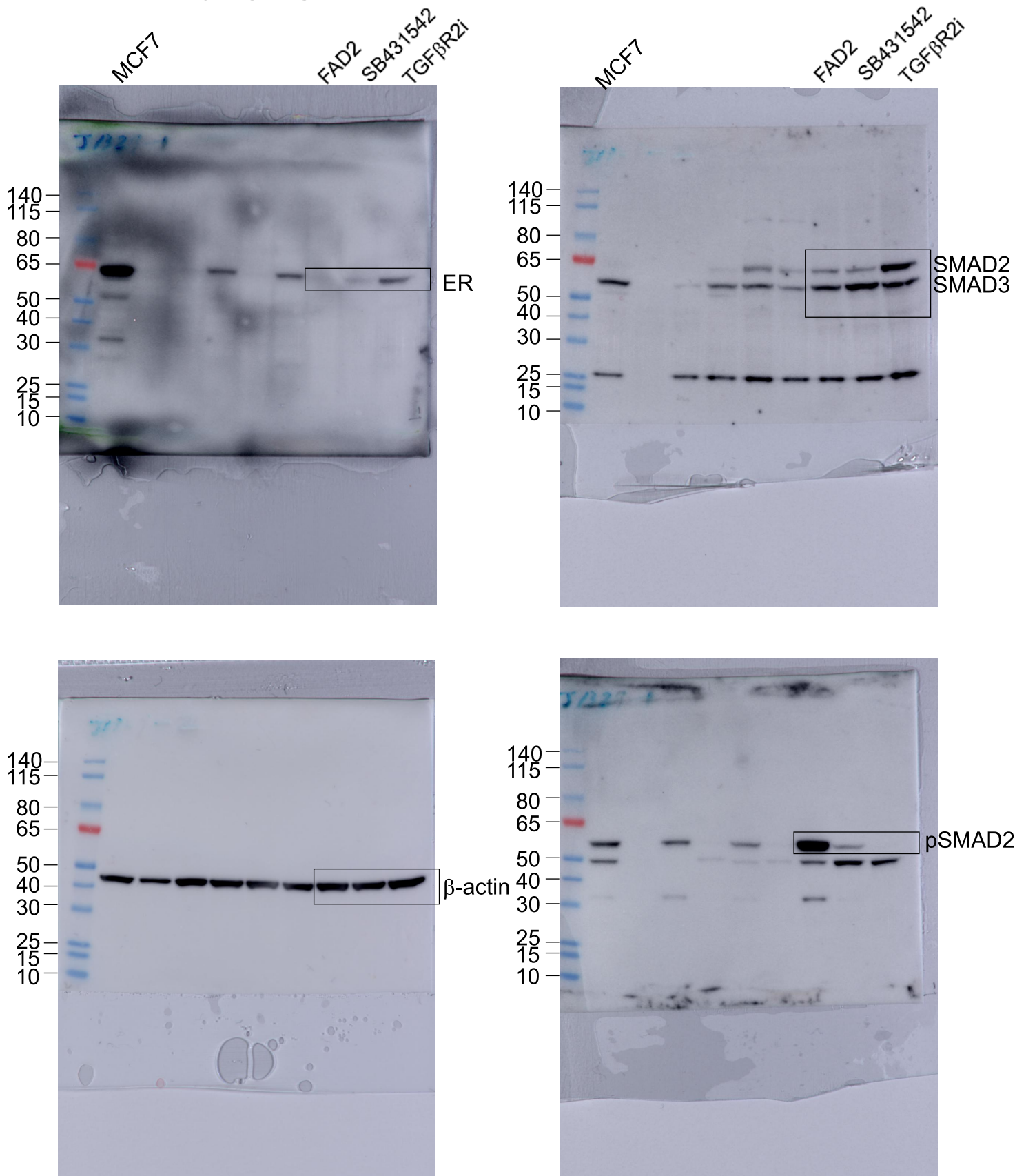
## Supplementary Figure 8. PCR array of human estrogen receptor signaling



(a) Using a commercially available PCR array containing 84 genes involved in ER activation and its response, qRT-PCR was conducted with RNA samples of ER<sup>pos</sup> HBEC with or without TGFβR2i. Among them, 20 genes (some of which are labeled in the plot) were highly expressed in the presence of TGFβR2i. Scatter plot graph shows the expression level of each gene in TGFβR2i treated CD166<sup>high</sup> cells versus DMSO treated CD166<sup>high</sup> cells. The black line indicates no difference between samples, while the pink lines indicate a two-fold change in gene expression. Two biological replicates were run in qPCR with duplicates of each. (b) qRT-PCR of ESR1 and PGR expression in hTERT/shp16 transduced ER<sup>pos</sup> HBECs exposed to vehicle(DMSO), SB431542 alone, Repsox alone or TGFβR2i for 7 days. Note that the highest levels of expression were obtained in presence of TGFβR2i. Colored bars indicate the fold difference of the relative gene expression and error bars represent SD of triplicates in qPCR.

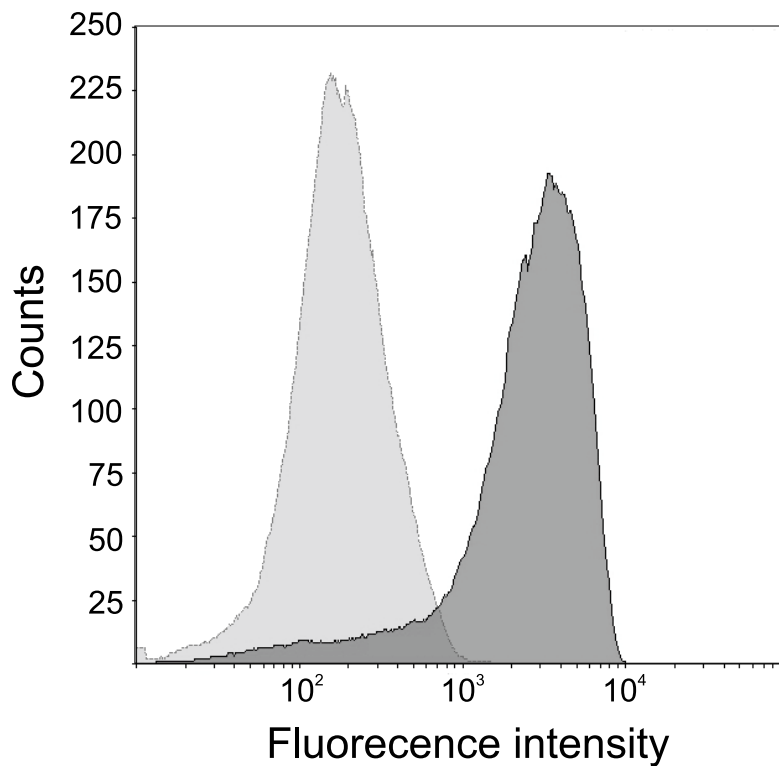


**Supplementary Figure 9. Full scale Western blots underlying Figure 4d**



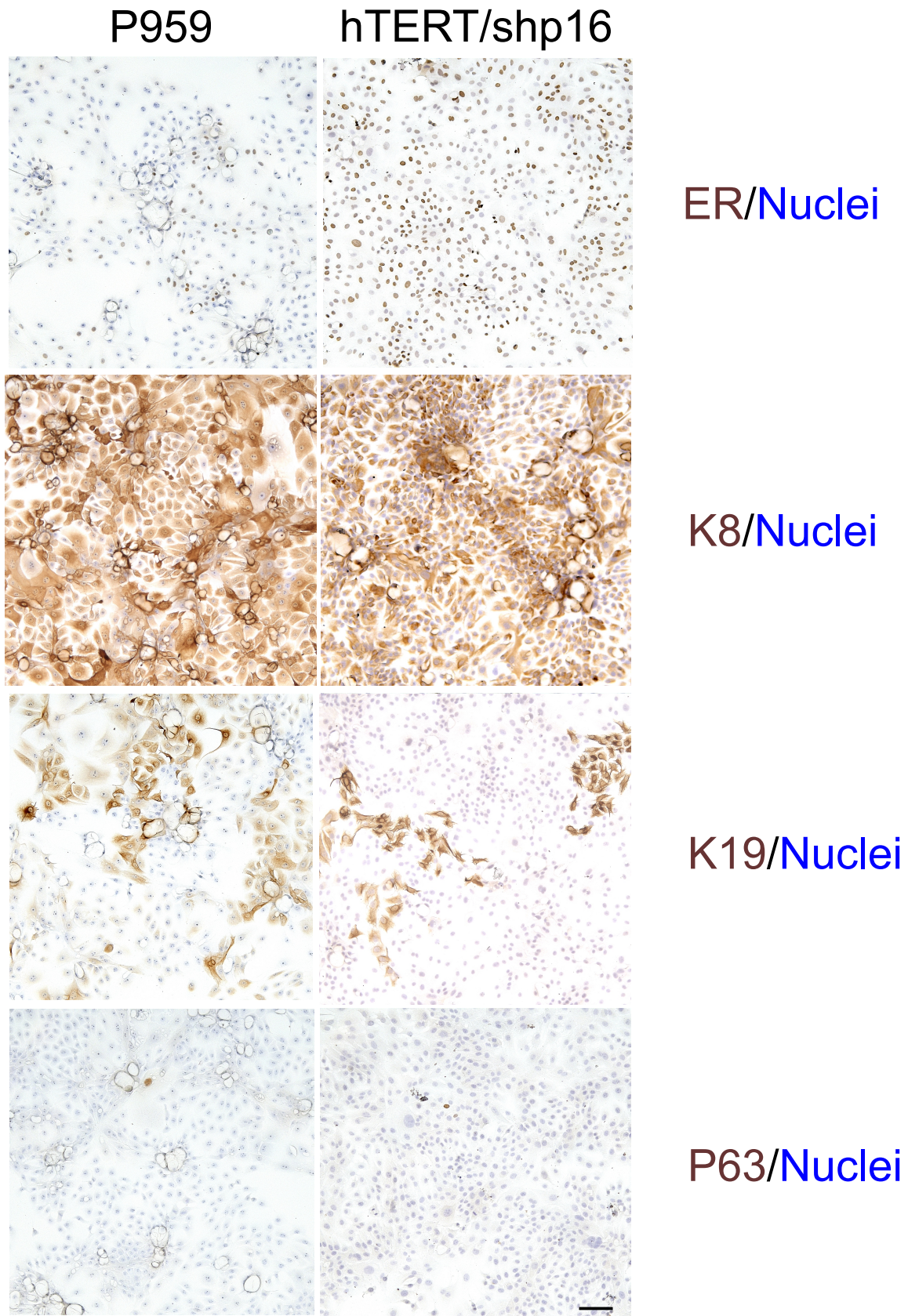
The frames indicate areas of the original Western blots selected for making the composite Figure 4d.

**Supplementary Figure 10.  $TGF\beta RII$  expression in  $ER^{pos}$  HBEC**



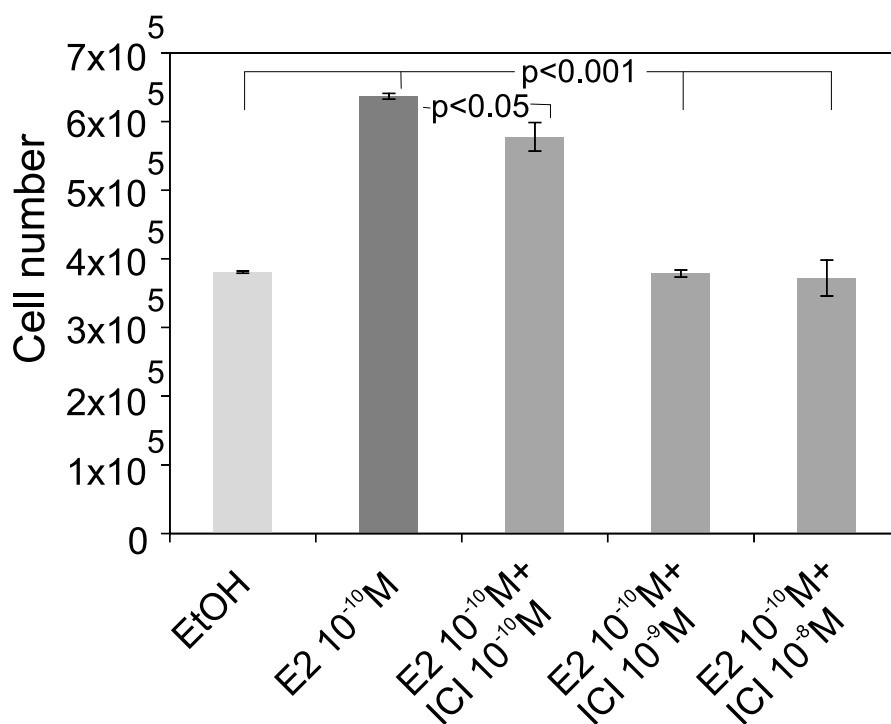
FACS analysis of  $TGF\beta RII$  surface expression in short-term cultured  $ER^{pos}$  HBEC. FITC- $TGF\beta RII$  (dark grey). FITC conjugated antibody in the absence of  $TGF\beta RII$  primary antibody (light grey).

**Supplementary Figure 11. hTERT/shp16 transduced cells remain ER<sup>pos</sup> and luminal-like**



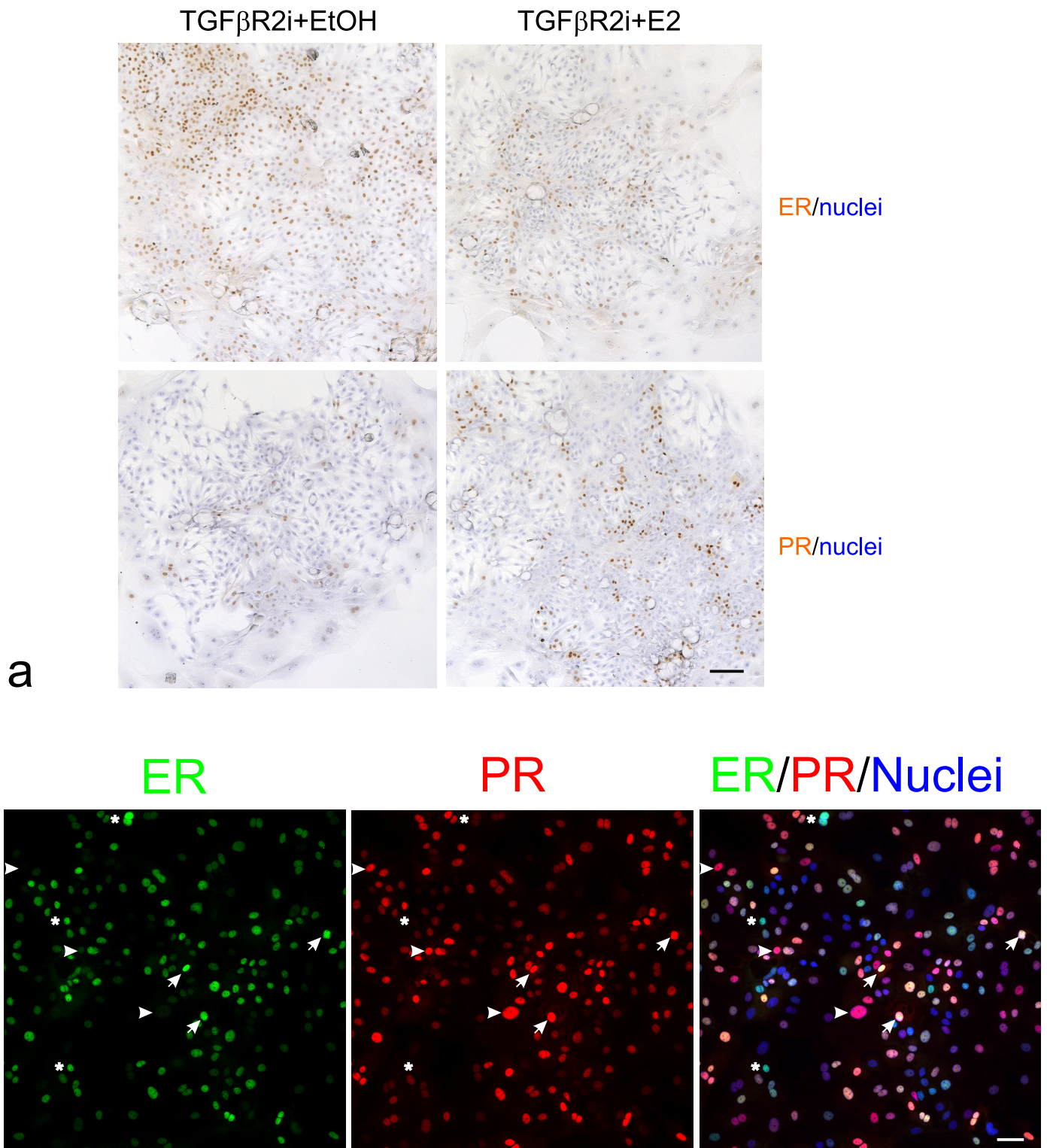
Immunoperoxidase staining of finite life span (left column) and hTERT/shp16 transduced (right column) ER<sup>pos</sup> HBEC with ER (SP1 prediluted), keratin K8 and K19 and P63. Cells were counterstained with hematoxylin. (Bar = 100µm)

**Supplementary Figure 12. Estrogen induced growth is completely abolished by pure antiestrogen**



ER<sup>pos</sup> HBEC with extended lifespan (hTERT/shp16) exposed for six days to vehicle (EtOH), E2 or E2 plus increasing concentrations of ICI182,780/Fulvestrant/Faslodex. Note that the pure antiestrogen abolishes the growth promoting effect of estrogen. Significance was assessed by ANOVA followed by Tukey's HSD test. Error bars represent SD of duplicates in EtOH and E2 and triplicates in ICI treated samples.

**Supplementary Figure 13. Augmented effect of estrogen on ER<sup>pos</sup> HBEC by long-term exposure and omission of EGF**



**a** Immunoperoxidase stainings of ER<sup>pos</sup> HBEC kept on TGFβR2i and in the presence of estrogen for two passages prior to omission of estrogen in the third passage (left column). Note the marked difference in staining with ER (SP1, prediluted) and PR (SAN27). Cells were counterstained with hematoxylin. (Bar = 100μm). **(b)** Multicolor imaging of ER<sup>pos</sup> HBEC in the presence of estrogen stained with ER (SP1; green), PR (PgR636; red) and nuclei (blue). Cells are either ER<sup>pos</sup> (asterisk), PR<sup>pos</sup> (arrowhead), ER<sup>pos</sup>/PR<sup>pos</sup> (arrows), or ER<sup>neg</sup>/PR<sup>neg</sup> (not labeled). (Bar = 50μm).

## Supplementary Table 1. List of primers used for qRT-PCR

Gene symbol	Assay ID	Category
<i>ACTA2</i>	Hs00426835_g1	Basal marker
<i>CD200</i>	Hs01033303_m1	Basal marker
<i>CD271</i>	Hs00609977_m1	Basal marker
<i>IGFBP3</i>	Hs00365742_g1	Basal marker
<i>KRT14</i>	Hs00265033_m1	Basal marker
<i>KRT5</i>	Hs00361185_m1	Basal marker
<i>MME</i>	Hs00153510_m1	Basal marker
<i>NGR1</i>	Hs00247620_m1	Basal marker
<i>SNAI2/SLUG</i>	Hs00950344_m1	Basal marker
<i>ST3GAL2</i>	Hs00199480_m1	Basal marker
<i>AREG</i>	Hs00950669_m1	Luminal marker
<i>BIM</i>	HS00708019_s1	Luminal marker
<i>CD166</i>	Hs00977640_m1	Luminal marker
<i>CDKN1B/P27</i>	Hs00153277_m1	Luminal marker
<i>ESR1</i>	Hs00174860_m1	Luminal marker
<i>FOXA1</i>	Hs00270129ml	Luminal marker
<i>GATA3</i>	Hs00231122_m1	Luminal marker
<i>GREB1</i>	Hs00536409_m1	Luminal marker
<i>ID2</i>	Hs04187239_m1	Luminal marker
<i>KRT18</i>	Hs02827483_g1	Luminal marker
<i>KRT19</i>	Hs00761767_s1	Luminal marker
<i>KRT8</i>	Hs01595539_g1	Luminal marker
<i>MYB</i>	Hs00920556_m1	Luminal marker
<i>NGRN</i>	Hs04185079_g1	Luminal marker
<i>PGR</i>	Hs01556702_m1	Luminal marker
<i>SORT1</i>	Hs00361760_m1	Luminal marker
<i>TFF1</i>	Hs00907239_m1	Luminal marker
<i>TGM2</i>	Hs00190278_m1	Luminal marker
<i>WNT4</i>	Hs01573504_m1	Luminal marker
<i>DPP4</i>	HS00175210_m1	Luminal progenitor marker
<i>ALDH1A3</i>	Hs00167476_m1	Luminal progenitor marker
<i>CD14</i>	Hs00169122_g1	Luminal progenitor marker
<i>CYP24A1</i>	Hs00167999_m1	Luminal progenitor marker
<i>DAPP1</i>	Hs00183937_m1	Luminal progenitor marker
<i>ELF5</i>	Hs01063022_m1	Luminal progenitor marker
<i>KIT</i>	Hs00174029_m1	Luminal progenitor marker
<i>KRT15</i>	Hs00267035_m1	Luminal progenitor marker
<i>NCALD</i>	Hs00230737_m1	Luminal progenitor marker
<i>PIGR</i>	Hs00922561_m1	Luminal progenitor marker
<i>Sox9</i>	Hs01001343_g1	Luminal progenitor marker
<i>GAPDH</i>	Hs02758991_g1	Reference
<i>HPRT1</i>	Hs99999909_m1	Reference
<i>TBP</i>	Hs00427621_m1	Reference
<i>TFRC</i>	Hs00951083_m1	Reference