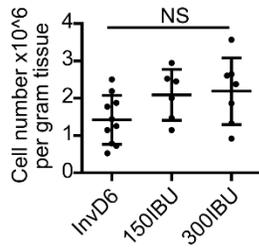
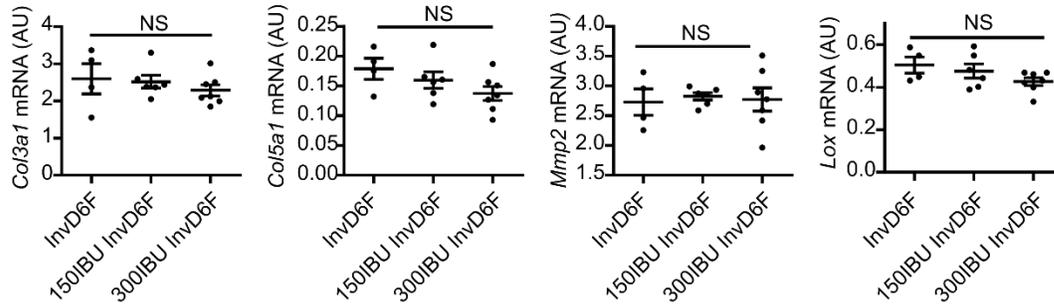


Supplemental figure 1. PDGFR α is expressed dominantly by stromal cells surrounding mammary ducts and alveoli. **A)** IHC staining of PDGFR α in nulliparous (left panel) and InvD6 mouse mammary glands (right panel). Black arrows: mammary ductal and alveolar structures; blue arrow: muscle; yellow arrows: blood vessel. **B)** Co-staining for PDGFR α (brown) and CD31 (green) in nulliparous Balb/c mammary gland shows lack of dual positive cells. Counterstain: pink. Scale bar: 30 μ m. **C)** Quantification of PDGFR α positive area surround CD31⁺ blood vessels (black) or elsewhere (gray) in three mammary glands from nulliparous mice (N 1-3) and three InvD6 mammary glands (InvD6 1-3). **D)** Flow cytometry for PDGFR α and PDGFR β in nulliparous mouse mammary glands reveals significant dual positivity. **E)** RT-qPCR for *Col1a1*, *Col3a1*, *Col5a1* and *LOX* gene expression in sorted PDGFR α ⁺ and PDGFR α ⁻ cells. N=6. Scale bar: 100 μ m. **P<0.01, ***P<0.001, #P<0.0001.

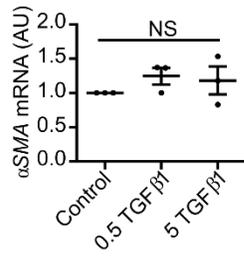
A.



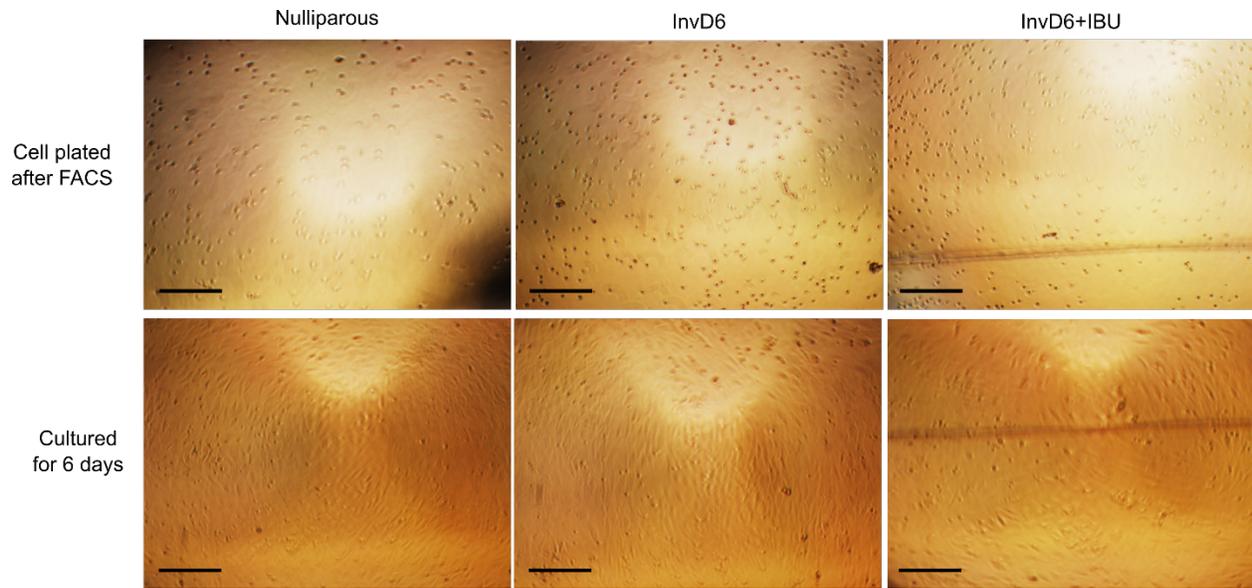
B.



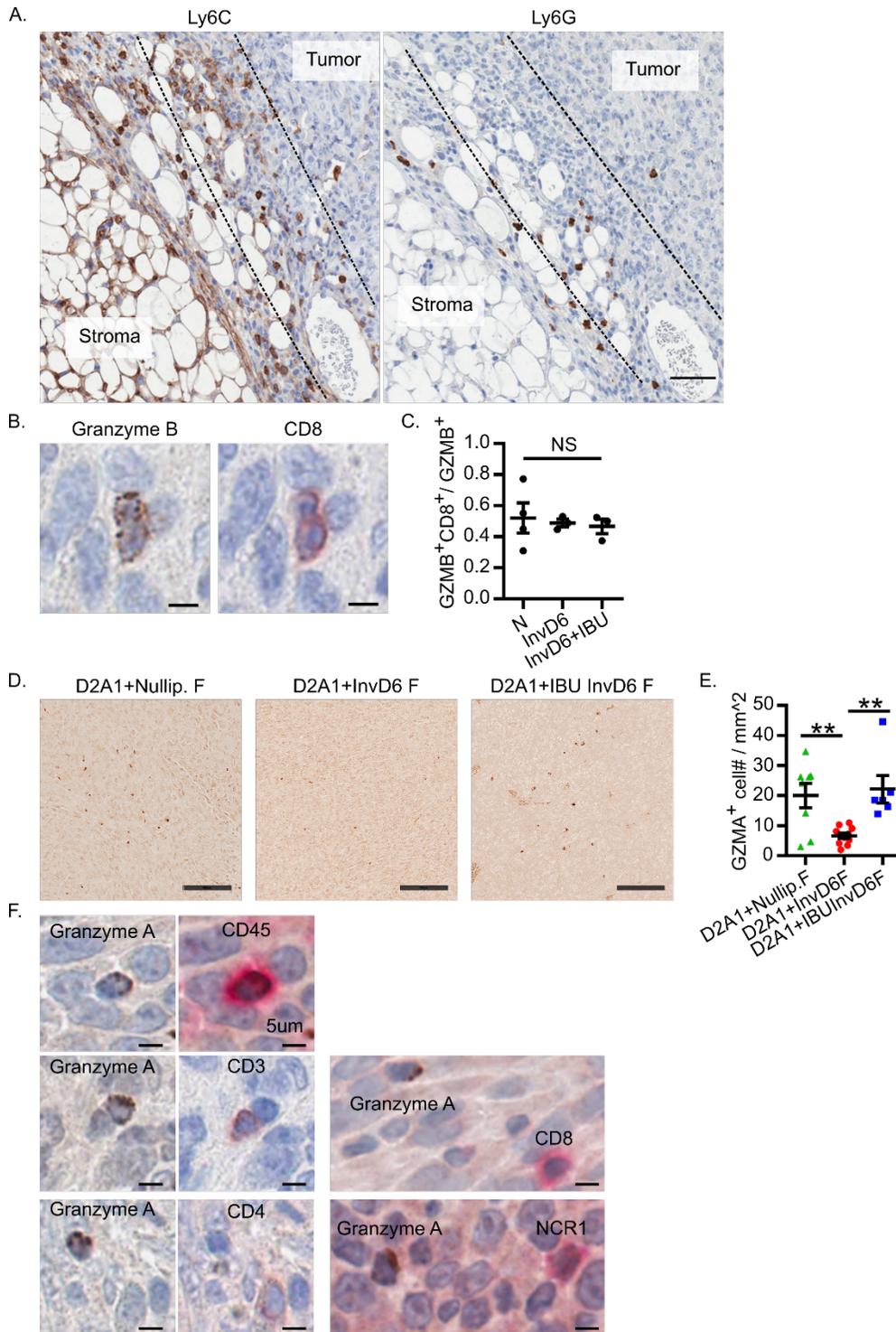
Supplemental figure 2. Effects of ibuprofen treatment on PDGFR α ⁺ cells in the InvD6 mammary gland. **A)** PDGFR α ⁺ cell number per gram mammary gland tissue quantitated by FACS analysis, in the absence or presence of two doses of ibuprofen treatment (150IBU:150 mg/kg and 300IBU: 300 mg/kg Ibuprofen in diet) for 6 days following weaning. N=6. **B)** RT-qPCR of *Col3a1*, *Col5a1*, *Lox* and *MMP2* gene expression in InvD6 fibroblasts in the absence or presence of ibuprofen. N=4-7. NS: not significant.



Supplemental figure 3. TGF β 1 does not induce fibroblast α SMA expression in floating collagen cultures. RT-qPCR of α SMA in primary mammary fibroblasts without (Control) or with the treatment of 0.5 ng/ml TGF β 1 (0.5 TGF β 1) and 5 ng/ml TGF β 1 (5 TGF β 1). N=3. NS: not significant.

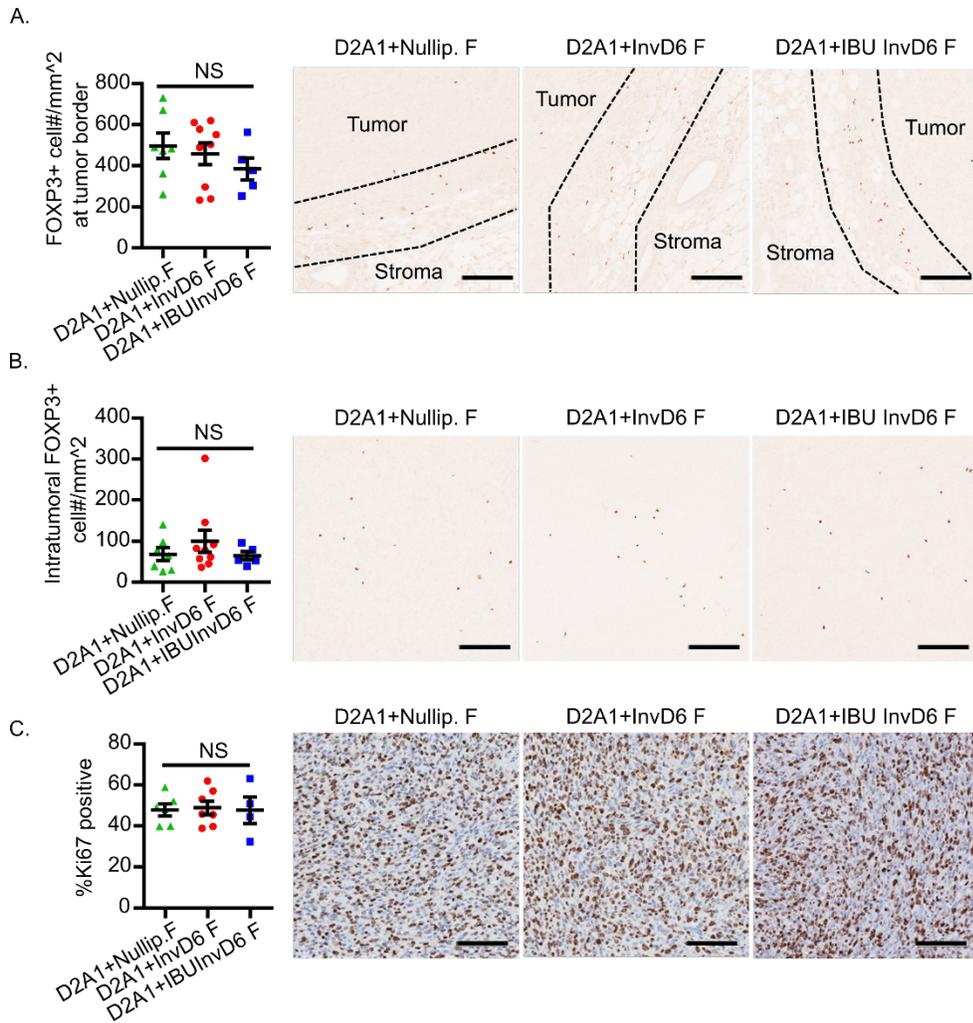


Supplemental figure 4. Fibroblast ex vivo adhesion and cell culture survival are independent of reproductive state. Top panel: Sorted PDGFR α ⁺ mammary fibroblasts used in the D2A1 tumor cell co-injection experiments and plated at equal concentration. Bottom panel: cells in the top panel after 6 days of culture. Scale bar: 100 μ m.

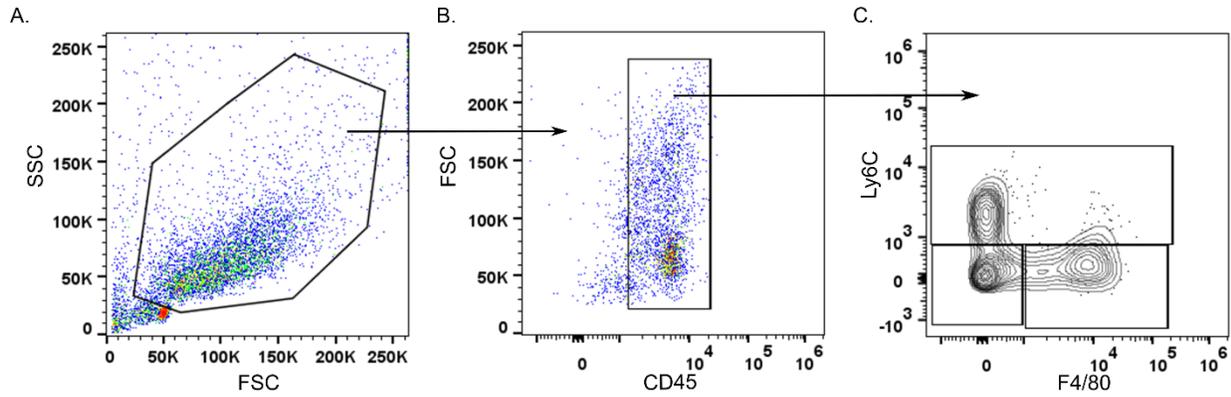


Supplemental figure 5. Immunohistochemical characterization of tumor infiltrating immune cells. **A)** Ly6C and Ly6G staining in serial sections of representative D2A1+ InvD6 F tumor. Dashed lines show tumor border. Scale bar: 50 μ m. **B)** IHC staining of granzyme B (left panel) and CD8 (right panel) of the same tumor section. **C)** Quantification the percentage of granzyme B⁺ cells that are CD8⁺. **D)** Representative images of granzyme A⁺ cells within tumors from each

group. Scale bar: 100 μm . **E)** Quantification of IHC staining for intratumoral granzyme A⁺ cells by group. **F)** IHC co-staining for granzyme A and immune cell markers CD45, CD3, CD8, CD4, and NCR1, as indicated by labels. Scale bar: 5 μm .



Supplemental figure 6. FOXP3 and Ki67 IHC tumor analyses demonstrate no differences between fibroblast groups. **A)** Left panel: Quantification of FOXP3⁺ cells at the tumor border. Right Panel: representative images of FOXP3⁺ cells at the tumor border. Scale bar: 100 μ m. **B)** Left panel: Quantification of intratumoral FOXP3⁺ cells. Right Panel: representative images of intratumoral FOXP3⁺ cells. Scale bar: 100 μ m. Counterstaining is not shown in A) and B) to enhance visualization of positive cells. **C)** Left panel: Quantification of Ki67⁺ cells in tumor. Right Panel: representative images of Ki67⁺ cells. Scale bar: 20 μ m.



Supplemental figure 7. Characterization of monocyte derived cells by flow cytometry. **A)** Cells are gated by forward scatter and side scatter. **B)** Immune cells are further gated by CD45 and forward scatter. **C)** CD45⁺ cells are further gated by F4/80 and Ly6C. Three populations are indicated. Monocyte: Ly6C⁺F4/80⁻. Macrophages: Ly6C⁻F4/80⁺. Immature myeloid progenitor cells: Ly6C⁻F480⁻

Supplemental method

Immunohistochemistry. 4 μm sections of formalin-fixed, paraffin-embedded mouse mammary glands were used for IHC. Slides were pretreated with Target Retrieval Solution (Dako S1699 or DAKO S2367). Then, slides were treated with 3% H_2O_2 in Methanol for 10 minutes, followed by protein block (Dako, X0909) for 1 hour. Primary and secondary antibody information are as follows:

Targets	Primary Antibody vendor and catalog #	Dilution	Secondary antibody vendor and catalog #	Dilution	Chromogen and color
PDGFR α	R&D, AF1062	1:300	Santa Cruz, SC2922	1:100	DAKO, K3468, brown
CD31	Abcam, ab28364	1:100	ThermoFisher, 31461	1:100	Bio SB, BSB0129, green
Ly6G	BD Biosciences, 551459	1:1000	Biocare, RT517H	N/A	DAKO, K3468, brown
Granzyme A	Santa Cruz, SC5515	1:100	Santa Cruz, SC2922	1:100	DAKO, K3468, brown
CD45	BD pharmingen, 550539	1:400	Santa cruz, SC2021	1:100	Biocare, WR806, red
CD3	Abcam ab16669	1:100	Histofine, Simple Stain, anti-rabbit	N/A	Vector, SK-4200, brick red
CD4	eBioscience, 14-9766-82	1:100	Histofine, Simple Stain, anti-rat	N/A	Vector, SK-4200, brick red
NCR1	Abcam ab199128	1:50	Histofine, Simple Stain, anti-rabbit	N/A	Biocare, WR806, red
FOXP3	eBioscience, 14-5773-82	1:100	Biocare, RT517H	N/A	DAKO, K3468, brown
Ki67	Neo-markers, RM9106s	1:100	ThermoFisher, 31461	1:100	DAKO, K3468, brown

All primary antibodies were incubated for 1 hour at room temperature, and secondary antibodies were incubated for 30 minutes in room temperature. For dual co-staining slides, three methods were used based on the primary antibodies combination. Method-1: a combination of DAB (DAKO K3468, brown) with Warp Red (Biocare, WR806, red). Method-2: a combination of DAB (DAKO K3468, brown) with PolyDetector HRP Green kit (Bio SB, BSB0129, green). Method-3: AEC (Vector SK-4200) was used with antigen retrieval (BioGenex HK086) between each staining targets. For PDGFR α and CD31 dual staining, tissues were counterstained with Nuclear Fast Red (American MasterTech, STNFRPT) for 1 minutes. For other staining, tissues were counterstained with hematoxylin (DAKO, S3301) for 6 minutes.

Flow cytometry. 1×10^6 cells per sample were blocked with CD16/32 (eBioscience, 14-0161-81) for 30 minutes and cell surface markers were stained for 20 minutes. PDGFR α (BioLegend, 135910): 1:100 dilution, PDGFR β (BioLegend, 136007): 1:100 dilution, CD45 (eBioscience, 25-0451-82): 1:800 dilution, Ly6C (BioLegend, 128012): 1:400 dilution, F4/80 (BioLegend, 123108): 1:200 dilution. Cells were stained for 20 minutes. Experiments were performed on an

LSRFortessa (BD Biosciences; Oregon Health and Science University Flow Cytometry Shared Resource) and analysis was performed using FlowJo (FlowJo, LLC Data Analysis Software, Ashland, OR).