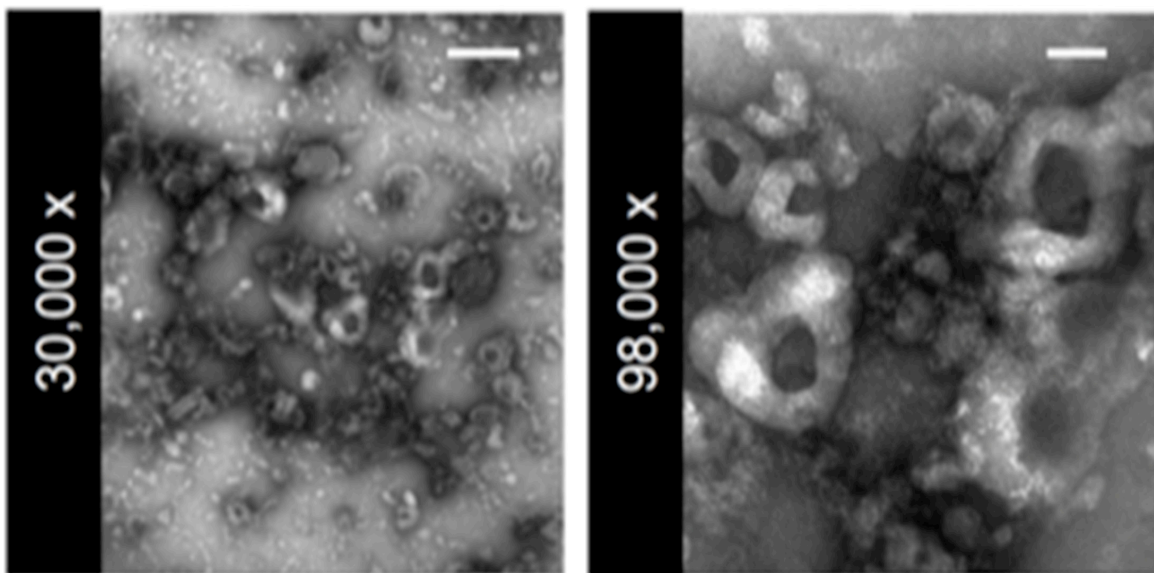
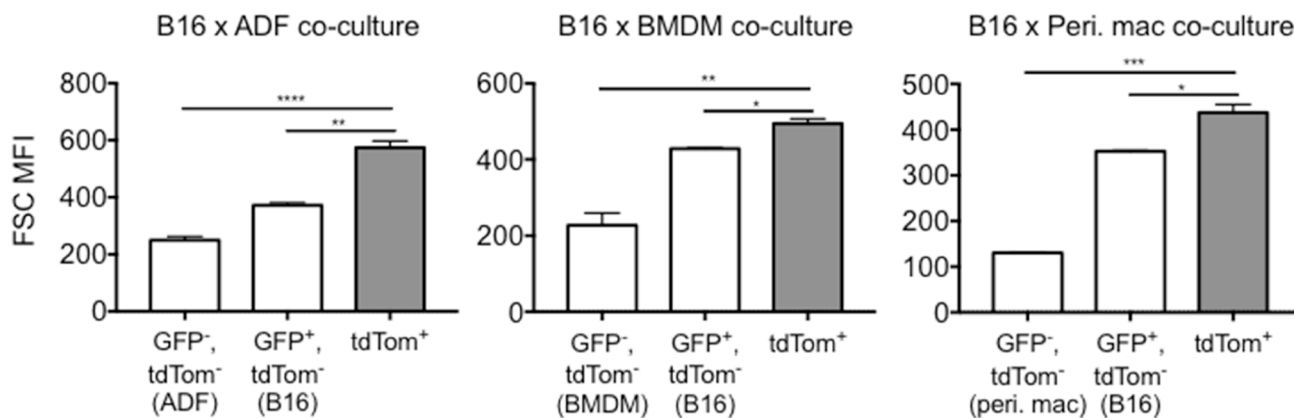


Cell-cell fusion as a mechanism of DNA exchange in cancer

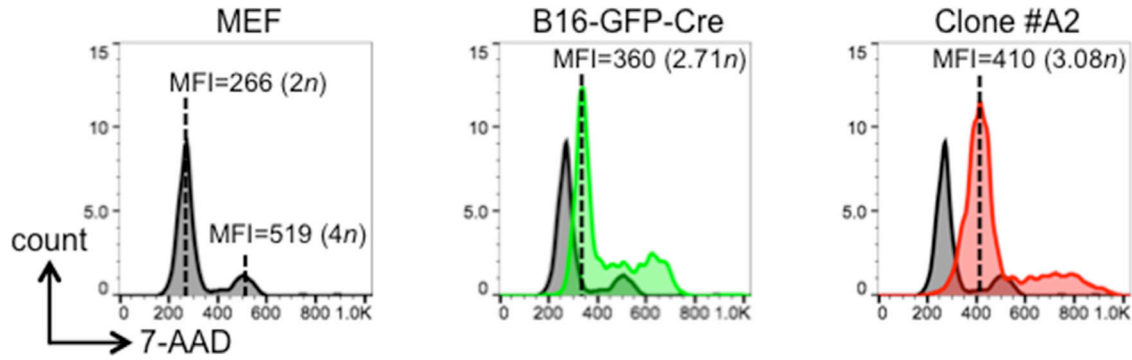
SUPPLEMENTARY MATERIALS



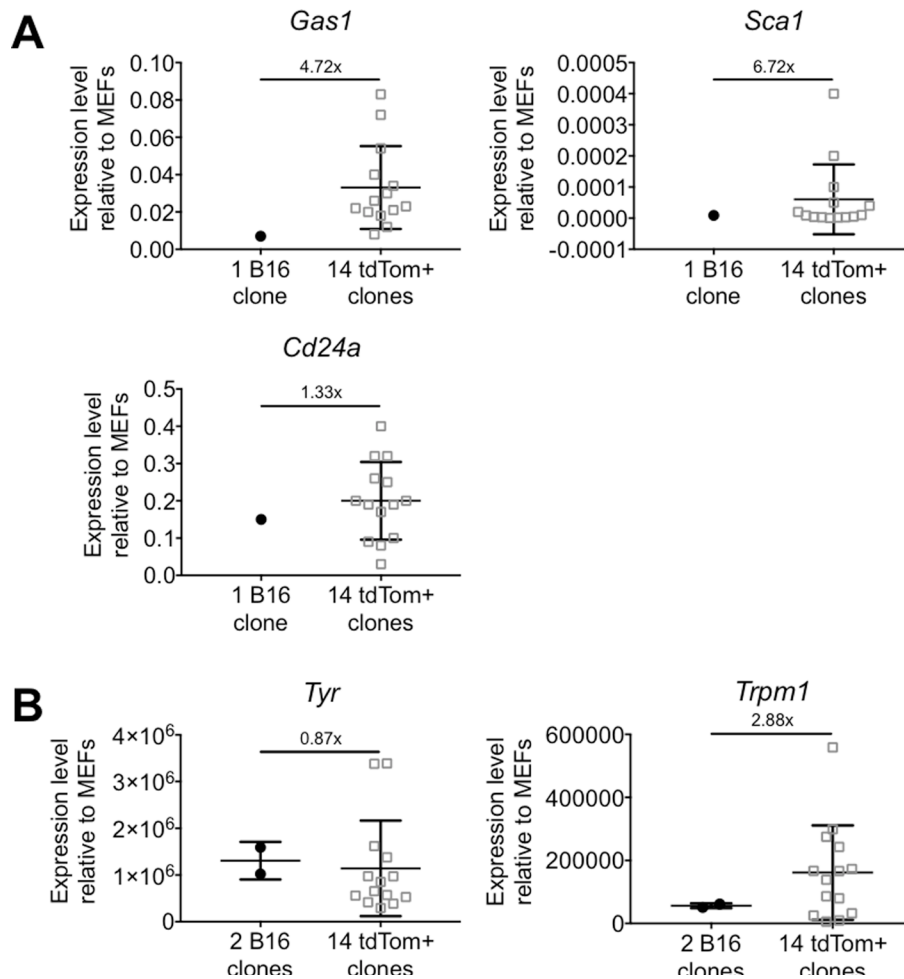
Supplementary Figure 1: Size and morphology of B16-GFP-Cre ECVs. Electron micrographs showing size and morphology of vesicles isolated from the conditioned media of B16-GFP-Cre cells via differential ultracentrifugation. Size bar equals 200 nm (30,000×) or 50 nm (98,000×).



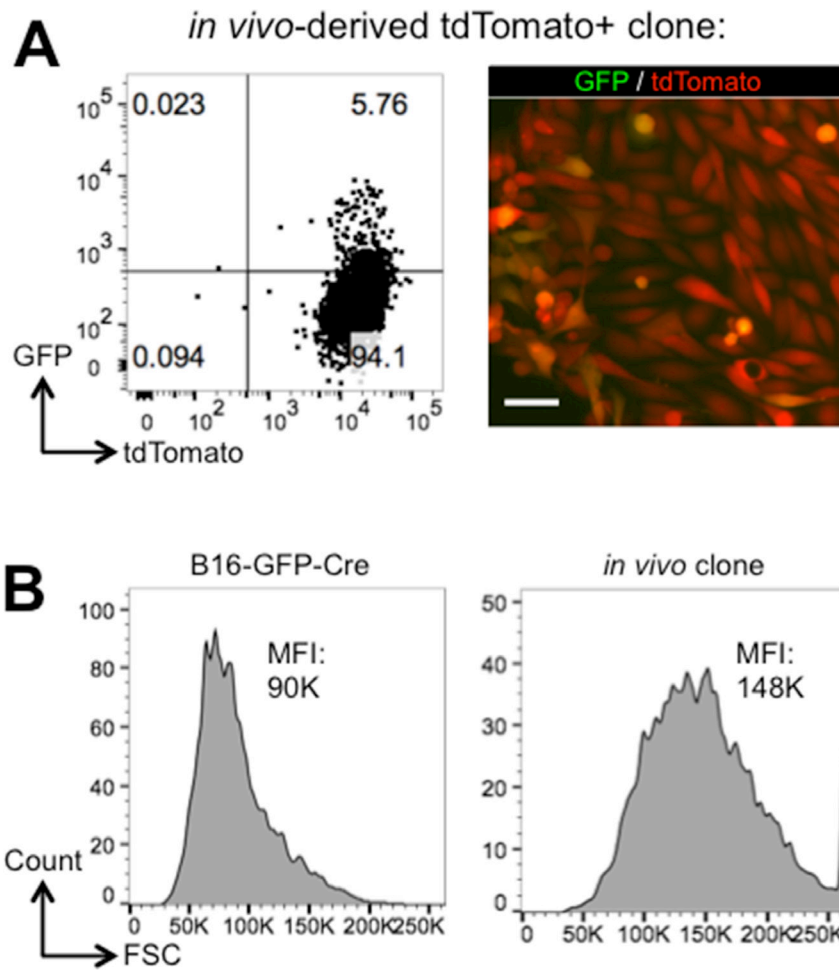
Supplementary Figure 2: (Related to Figure 3B): Reporter⁺ cells are larger than B16 cells and Reporter⁻ cells. FSC analysis of three populations of cells (GFP⁻/tdTomato⁻, GFP⁺/tdTomato⁻, tdTomato⁺) from 48 hr co-cultures of B16-GFP-Cre cells with various reporter cells (indicated).



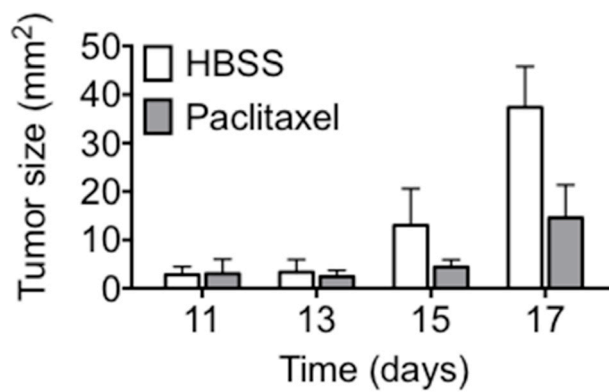
Supplementary Figure 3: (Related to Figure 4D): FACS-based quantification of DNA content. Representative histograms showing DNA content of MEF, B16-GFP-Cre, and one of the twenty tdTomato+ clones (#A2). The two peaks in each histogram correspond to cells in the G1 (left peak) or G2 (right peak) phase of the cell cycle. For B16-GFP-Cre and each of the twenty tdTomato+ clones, relative ploidy was calculated by normalizing the MFI of the G1 peak to MEF, which were set as $2n$.



Supplementary Figure 4: (Related to Figure 5C): B16xMEF hybrids express both B16- and MEF-restricted genes. (A) Comparison of the average expression level of three candidate “MEF genes” in B16-GFP-Cre cells and fourteen tdTomato+ clones ($n = 2$ independent experiments per data point). For the tdTomato+ clones, data is represented as mean \pm SEM. (B) Comparison of the average expression level of two candidate “B16 genes” in two B16-GFP-Cre clones and fourteen tdTomato+ clones ($n = 2$ independent experiments per data point). Data is represented as mean \pm SEM.



Supplementary Figure 5: (Related to Figure 7E): Cell-cell fusion occurs *in vivo*. (A) Representative FACS plot and fluorescent micrograph showing GFP and tdTomato expression in the *in vivo*-derived tdTomato⁺ clone after four passages. Scale bar = 100 μ M. (B) Representative histograms showing FSC (as a read-out for cell size) of B16-GFP-Cre and the *in vivo*-derived clone.



Supplementary Figure 6: (Related to Figure 7F): Paclitaxel treatment shrinks B16-GFP-Cre tumors *in vivo*. (A) Bar graph showing the size of B16-GFP-Cre tumors that were grown in mice treated with paclitaxel or HBSS (control). Data are represented as mean \pm SEM ($n = 6$ mice per group).

Supplementary Video 1: (Related to Figure 3): Cell-cell fusion mediates the transfer of Cre from B16 melanoma cells to MEF. (A) Live-cell imaging of B16-GFP-Cre cells co-cultured with CellTracker Blue-labeled reporter MEF between 2 and 18 hours. Scale bar equals 50 μ M. Green represents GFP, blue represents CellTracker Blue, and red represents tdTomato. See Supplementary_Video_1

Supplementary Video 2: (Related to Figure 3): Cell-cell fusion mediates the transfer of Cre from B16 melanoma cells to MEF. (A) Live-cell imaging of B16-GFP-Cre cells co-cultured with CellTracker Blue-labeled reporter MEF between 2 and 18 hours. Scale bar equals 50 μ M. Green represents GFP, blue represents CellTracker Blue, and red represents tdTomato. See Supplementary_Video_2

Supplementary Video 3: (Related to Figure 3): Cell-cell fusion mediates the transfer of Cre from B16 melanoma cells to MEF. (A) Live-cell imaging of B16-GFP-Cre cells co-cultured with CellTracker Blue-labeled reporter MEF between 2 and 18 hours. Scale bar equals 50 μ M. Green represents GFP, blue represents CellTracker Blue, and red represents tdTomato. See Supplementary_Video_3

Supplementary Video 4: (Related to Figure 3): Reporter⁺ cell undergoing apoptosis. (A) Live-cell imaging of B16-GFP-Cre cells co-cultured with CellTracker Blue-labeled reporter MEF between 18 and 40 hours. Scale bar equals 50 μ M. Green represents GFP, blue represents CellTracker Blue, and red represents tdTomato. See Supplementary_Video_4

Supplementary Video 5: (Related to Figure 3): Cell-cell fusion mediates the transfer of Cre from B16 melanoma cells to MEF. (A) Live-cell imaging of B16-GFP-Cre cells co-cultured with unlabeled reporter MEF between 2 and 20 hours. Scale bar equals 50 μ M. Green represents GFP and red represents tdTomato. See Supplementary_Video_5

Supplementary Table 1: Primer sequences

Table of primers	
Name	Sequence (5' to 3')
<i>Bmp4</i> (forward)	GGCTGGCCATTGAGGTGACT
<i>Bmp4</i> (reverse)	TCGGCTGATTCTGACATGCTG
<i>Cd24a</i> (forward)	CTGCTGGCACTGCTCCTAC
<i>Cd24a</i> (reverse)	GGTGGTGGCATTAGTTGGAT
<i>CRE</i> (forward)	GCCAGCTAAACATGCTTCATC
<i>CRE</i> (reverse)	ATTGCCCTGTTTTCACTATCC
<i>Fgf2</i> (forward)	TCCAGTTGGTATGTGGCACTGA
<i>Fgf2</i> (reverse)	CAGTATGGCCTTCTGTCCAGGTC
<i>Gapdh</i> (forward)	GGAGCCAAAAGGGTCATCAT
<i>Gapdh</i> (reverse)	GTGATGGCATGGACTGTGGT
<i>Gas1</i> (forward)	CCCTCTTCTGTGCGGTTTTTC
<i>Gas1</i> (reverse)	CACCGTTCAGCACTCTGAGTC
<i>Hprt</i> (forward)	GCTTGCTGGTGAAAAGGACCTCTCGAAG
<i>Hprt</i> (reverse)	CCCTGAAGTACTCATTATAGTCAAGGGCAT
<i>Met</i> (forward)	GCATGTCAGCATCGCTCAA
<i>Met</i> (reverse)	TGCAGGCCAGCTGTTTTC
<i>Mitf</i> (forward)	GCCTTGTTTATGGTGCCTTC
<i>Mitf</i> (reverse)	GTCCTCCTCCCTCTACTTTCTGT
<i>Scal</i> (forward)	CTCTGAGGATGGACACTTCT
<i>Scal</i> (reverse)	GGTCTGCAGGAGGACTGAGC
<i>Trpm1</i> (forward)	GAGCTGAAGGAGGCTAGGCTG
<i>Trpm1</i> (reverse)	CTTGGTGTCTCTCTCTGTTGT
<i>Tyr</i> (forward)	CCTCCTGGCAGATCATTTGT
<i>Tyr</i> (reverse)	ATCGCATAAAACCTGATGGC