

Video Article

An *In Vitro* System to Study Tumor Dormancy and the Switch to Metastatic Growth

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URL: <https://www.jove.com/video/2914>

DOI: [doi:10.3791/2914](https://doi.org/10.3791/2914)

Keywords: Medicine, Issue 54, Tumor dormancy, cancer recurrence, metastasis, reconstituted basement membrane extract (BME), 3D culture, breast cancer

Date Published: 8/11/2011

Citation: Barkan, D., Green, J.E. An *In Vitro* System to Study Tumor Dormancy and the Switch to Metastatic Growth. *J. Vis. Exp.* (54), e2914, doi:10.3791/2914 (2011).

Abstract

Recurrence of breast cancer often follows a long latent period in which there are no signs of cancer, and metastases may not become clinically apparent until many years after removal of the primary tumor and adjuvant therapy. A likely explanation of this phenomenon is that tumor cells have seeded metastatic sites, are resistant to conventional therapies, and remain dormant for long periods of time¹⁻⁴.

The existence of dormant cancer cells at secondary sites has been described previously as quiescent solitary cells that neither proliferate nor undergo apoptosis⁵⁻⁷. Moreover, these solitary cells has been shown to disseminate from the primary tumor at an early stage of disease progression⁸⁻¹⁰ and reside growth-arrested in the patients' bone marrow, blood and lymph nodes^{1,4,11}. Therefore, understanding mechanisms that regulate dormancy or the switch to a proliferative state is critical for discovering novel targets and interventions to prevent disease recurrence. However, unraveling the mechanisms regulating the switch from tumor dormancy to metastatic growth has been hampered by the lack of available model systems.

in vivo and *ex vivo* model systems to study metastatic progression of tumor cells have been described previously^{1,12-14}. However these model systems have not provided in real time and in a high throughput manner mechanistic insights into what triggers the emergence of solitary dormant tumor cells to proliferate as metastatic disease. We have recently developed a 3D *in vitro* system to model the *in vivo* growth characteristics of cells that exhibit either dormant (D2.OR, MCF7, K7M2-AS.46) or proliferative (D2A1, MDA-MB-231, K7M2) metastatic behavior *in vivo*. We demonstrated that tumor cells that exhibit dormancy *in vivo* at a metastatic site remain quiescent when cultured in a 3-dimension (3D) basement membrane extract (BME), whereas cells highly metastatic *in vivo* readily proliferate in 3D culture after variable, but relatively short periods of quiescence. Importantly by utilizing the 3D *in vitro* model system we demonstrated for the first time that the ECM composition plays an important role in regulating whether dormant tumor cells will switch to a proliferative state and have confirmed this in *in vivo* studies¹⁵⁻¹⁷. Hence, the model system described in this report provides an *in vitro* method to model tumor dormancy and study the transition to proliferative growth induced by the microenvironment.

Video Link

The video component of this article can be found at <https://www.jove.com/video/2914/>

Protocol

1. Cell culture maintenance of dormant and metastatic tumor cell lines

1. Grow dormant (D2OR/ MCF7/K7M2-AS.46) and metastatic tumor cells (D2A1/ MDA-MB-231/ K7M2) in 10 cm culture plates containing Dulbecco's Modified Eagle's Medium (DMEM) high glucose and 10% Fetal bovine serum (FBS) and antibiotics. Once the cells reach 70-80% confluence, proceed to the following assays.

2. Cell proliferation assay of dormant (quiescent) and metastatic (proliferating) tumor cells cultured in a 3D-BME system

Culturing dormant/metastatic cells in the 3D system

1. Thaw Cultrex growth factor-reduced Basement Membrane Extract (BME) in 4°C refrigerator a night before carrying out the assay. Note the BME should be handled on ice at all times.

2. On the following day, place a 96 well plate on a tray of ice inside a laminar hood. Coat each well with 50-100µl of ice cold BME using a dispenser with a syringe. Make sure no bubbles are formed in the wells. Place the 96 well plate coated with BME in a humidified incubator with 5% CO₂ at 37°C for 30 minutes.
3. In the meantime aspirate the media from the dormant and or metastatic tumor cells (prepared in section 1). Rinse culture plates with 10 ml phosphate buffered saline, pH 7.4 (PBS). Aspirate the PBS and add 2ml trypsin pre-warmed at 37°C, to the culture plates. Incubate the plates in a humidified 5% CO₂ at 37°C, for 5 min.
4. Transfer cells to a 15 ml conical tube containing 5ml of DMEM high glucose supplemented with 10% FCS and antibiotics and count the cells.
5. Spin down the total cell number to be cultured in a tissue culture centrifuge at a speed of 1500g, at room temperature for 5 min. In our assays we prepare 2X10³ cells /well for each cell line or time point to be examined. However, this may vary depending upon the cell lines used.
6. Carefully aspirate the supernatant. Note, in most cases the pellet is not visible. Therefore, leave some media behind. Tap the bottom of the 15ml conical tube with your fingers to ensure that a suspension of single cells is obtained. Re-suspend the pellet with **DMEM low glucose with antibiotics supplemented with 2% FCS + 2% BME (assay media)**. 100 µl of the assay media should be added for every 2x10³ cells. Triturate the cells many times with a 5ml pipette to ensure that a single cell suspension is maintained.
7. Plate a 100µl of the cell mixture per well on top of the 96 well BME coated plate. For background evaluation (in section 2.8) plate in addition 100µl per well of only the assay media on top of the 96 well BME coated plate. Incubate the cultured 96 well plates in a humidified 5% CO₂ incubator at 37°C. Cells should be re-fed every 4 days with the assay media.

Proliferation assay:

8. Proliferation assay of the cells: add to the wells at the desired time points 20 µl of the Cell Titer 96 AQueous One Solution Cell Proliferation assay kit. Incubate in a humidified 5% CO₂ incubator at 37°C for 2h. Using an Elisa Plate Reader record the absorbance at 490nm. For background evaluation and subtraction, add 20µl of the Cell Titer 96 AQueous One Solution Cell Proliferation assay kit to wells pre coated with BME and only overlaid with assay media. Using an Elisa Plate Reader record the absorbance at 490 nm.

3. Immunofluorescent staining for cell signaling molecules in dormant (quiescent) tumor cells and/or metastatic (proliferating) tumor cells

Culturing dormant/metastatic cells in the 3D system for immunofluorescence staining

*The following protocol is a modification of a 3D culture protocol published by Debnath J *et al*¹⁸.

1. Prepare BME as described in section 2.1. The following day: place an 8-chamber glass slide system on a tray of ice inside a laminar hood. Coat each well with 50µl of ice-cold BME using a 200µl pipetman. Make sure BME is spread evenly and no bubbles are formed in the wells. Place the 8 chamber glass slide coated with BME in a humidified 5% CO₂ at 37°C for 20 min.
2. Harvest dormant and or metastatic cells from section 1 and prepare to culture as described in section 2.3- 2.4. Collect the total number of cells to be cultured in a 15 ml conical tube. We prepare 5 x10³ cells / well for each cell line and time point to be examined. Spin down the cells in a tissue culture centrifuge at a speed of 1500g, at room temperature for 5 min. Aspirate the supernate carefully. Note that the pellet is not visible; therefore leave some media behind. Tap the bottom of the 15ml conical tube with your fingers to ensure that a single cell suspension is obtained. Re-suspend the pellet with assay media. 400µl of the assay media should be added for every 5x10³ cells. Triturate the cells many times with 5ml pipette. This step is very important to ensure that the single cell suspension is maintained.
3. Plate 400µl of the cell mixture per well on top of each of the 8 chambers coated with BME. Incubate the cultured 8 chambers glass slide system in a humidified 5% CO₂ incubator at 37°C. Cells should be re-fed every 4 days with the assay media.

Immunofluorescence staining:

4. At the desired time points, aspirate the upper layer of the media and add 200µl of the fixative containing 4% Paraformaldehyde (PFA), 5% sucrose and 0.1% Triton X-100 and incubate at room temperature for 5 minutes. Aspirate the fixative and add 200µl of 4% PFA containing 5% sucrose and incubate at room temperature for 25 minutes.
5. Aspirate the fixative; add 400 µl of phosphate buffered saline (PBS) to each well. Incubate for 10 minutes at room temperature. Aspirate the PBS and add 400 µl PBS containing 0.05% Tween 20 for 10 minutes at room temperature.
6. Block the fixed cells at room temperature with 200µl of either 10% donkey serum or with 3% BSA for 1hour (blocking solution to be used should be determined empirically for each primary antibody).
7. Aspirate the blocking solution and add 200µl of the primary antibody (dilution should be determined empirically for each primary antibody to be used). Dilute the primary antibody in 10% donkey serum if 10% donkey serum was used for blocking or dilute primary antibody in 3% BSA if 3% BSA blocking solution was used. Incubate with the primary antibody overnight at 4°C.
8. Aspirate the antibody, wash the wells with 400µl PBS for 15 minutes and repeat twice. Aspirate the PBS and add 200µl of donkey anti-respective- IgG conjugated to rhodamine red (dilution should be determined empirically), cover the 8 chamber slide with aluminum foil and incubate for 1 hour at room temperature.
9. Wash the wells with 400µl PBS (3x15 minutes each wash). Aspirate PBS. Mounted with VECTASHIELD mounting medium with DAPI. Dry slides for 40 minutes at room temperature in the dark. Slides can be kept for 1 week at 4°C. Store the slides in the dark. Image slides by Confocal microscopy.

4. Representative Results:

An example of a proliferation analysis of the dormant D2.0R and metastatic D2A1 tumor cells in the 3D culture is shown in **Figure 1A**. D2.0R cells are dormant (quiescent) through the entire experimental 14 day culture period whereas the highly metastatic D2A1 cells remain dormant only for four to six days after which they begin to proliferate. During the initial dormant phase, many cells remain solitary in the 3-D culture (**Figure 1B**; day 4) whereas other non-proliferating cells form multi-cellular spheroids. The transition of D2A1 cells from a dormant to proliferative state in 3-D culture (**Figure 1B**; day 12) is associated with dramatic changes in cell morphology. Hence, this assay can be used to test what

factor/s may trigger dormant D2.0R cells to emerge from their dormant state and what factor/s may prevent D2A1 cells to transition from their dormant state. **Figure 2** is an example of an agent preventing D2A1 cells to transition from dormant to a proliferative state. As illustrated in **Figure 2**, treatments of D2A1 cells with a specific inhibitor of myosin light chain kinase (ML-7) maintained D2A1 cells in a dormant state.

Cell signaling in the dormant and proliferating tumor cells cultured in the 3D system can be studied by immunofluorescence staining for cell signaling molecules. As illustrated in **Figure 3** a significant increase in myosin light chain phosphorylation in D2A1 cells (red staining) followed by reorganization of the f-actin filaments forming actin stress fibers (green staining) occurs during their transition from dormancy (1-4 days) to proliferation (day 7). However, blocking myosin light chain kinase activity in D2A1 cells by shRNA or specific drug (ML-7) retains D2A1 cells in a dormant state and results in inhibition of myosin light chain phosphorylation and f-actin stress fiber organization (**Figure 4**).

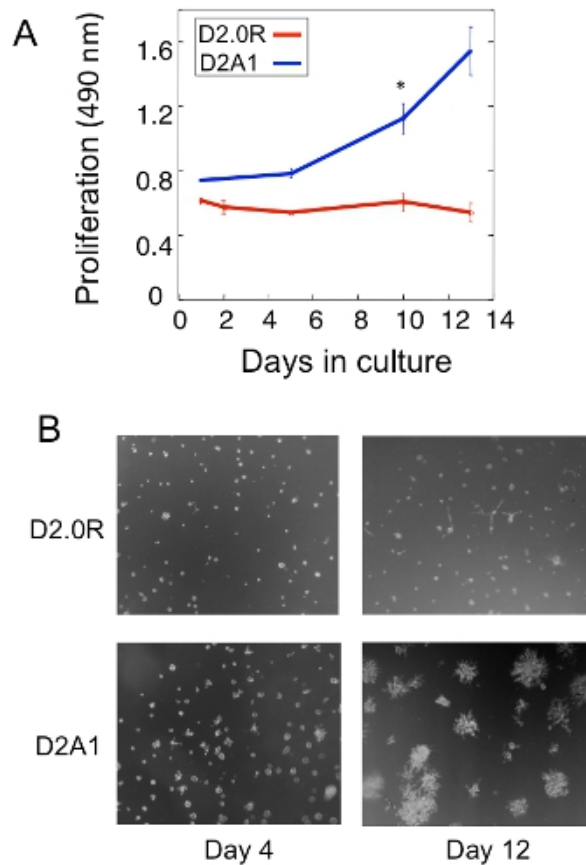


Figure 1. *in vitro* model to study solitary tumor cell dormancy and the switch to metastatic growth. A) Proliferation of dormant D2.0R and metastatic D2A1 in 3-D *Cultrex* BME, $n=8$ (mean \pm SE). Representative results of three experiments (* $p \leq 0.05$). B) Light microscopy images of D2.0R and D2A1 cells cultured in 3-D *Cultrex* BME magnification x20. Figure modified from Barkan et al ¹⁷.

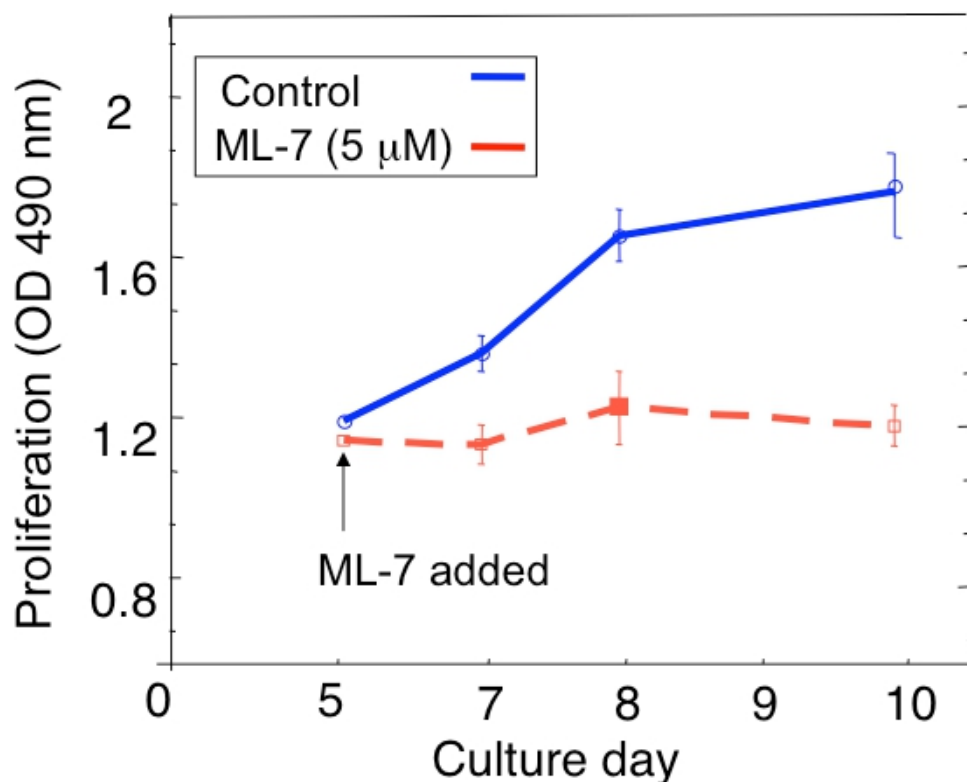


Figure 2. Preventing the switch of D2A1 cells from dormancy (quiescence) to proliferation in the 3D culture system by inhibition of myosin light chain kinase (MLCK). Time course of D2A1 cell proliferation cultured in 3-D *Cultrex* BME, $n=8$ (mean \pm SE). Cells were untreated (control), or treated with a specific inhibitor of MLCK (ML-7; 5 μ M) for 48 hr beginning on culture day 5. Figure modified from Barkan et al.¹⁷.

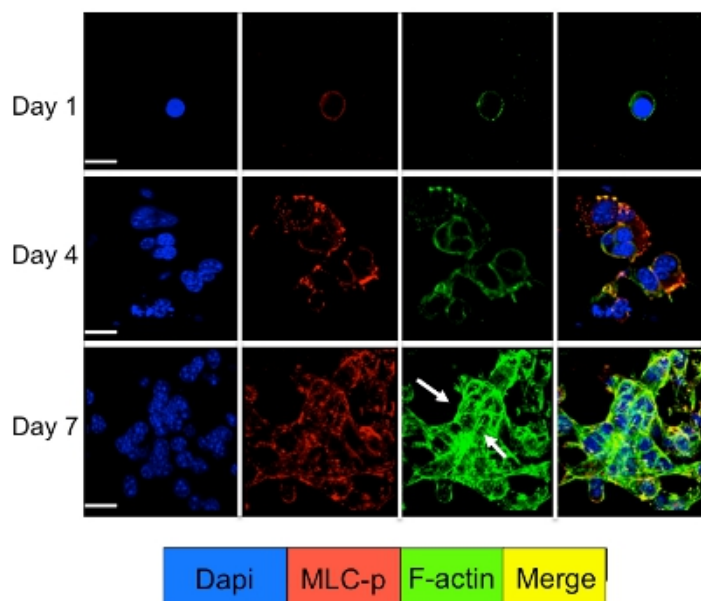


Figure 3. Myosin light chain phosphorylation followed by f-actin reorganization during the switch of D2A1 cells from dormancy to proliferative growth. D2A1 cells were cultured in 3-D *Cultrex* BME on 8 chamber glass slide. Cells were fixed and stained with DAPI (blue) for nuclear localization, phalloidin (green) for f-actin and with an antibody against the phosphorylated form of myosin light chain (MLC-p) (red), as indicated at various time points. Merge of f-actin, and MLC-p staining (yellow). Expression of MLC-p was increased during the transition of D2A1 cells from dormancy (days 1-4) to proliferative growth (day 7) followed by actin stress fiber formation (arrows). Confocal microscopy, magnification $\times 63$. White bar equals 20 microns. Figure modified from Barkan et al.¹⁷.

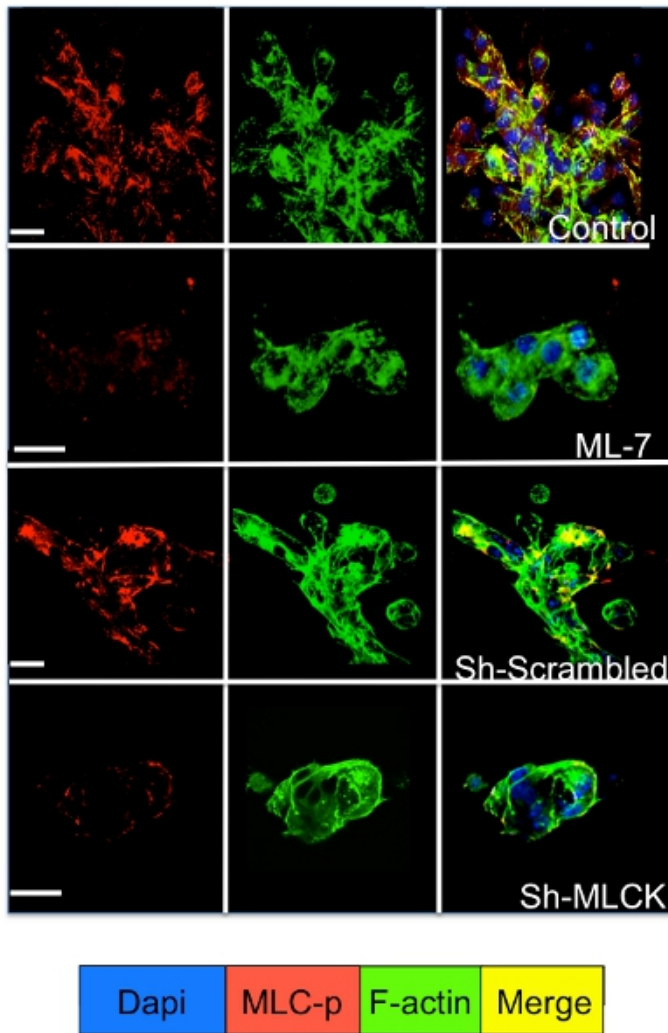


Figure 4. Inhibition of myosin light chain kinase (MLCK) mediated f-actin stress fiber formation in D2A1 cells. D2A1 Cells were untreated (control), or treated with inhibitor for MLCK (ML-7; 5 μ M), for 48 hr beginning on culture day 5, or treated with scrambled or MLCK shRNA and stained for the phosphorylated form of myosin light chain (MLC-p) (red), f- actin (green), and nuclei (blue). Merge of f-actin, and MLC-p staining (yellow). Confocal microscopy, magnification x63. White bar equals 20 microns.

Discussion

The underlying mechanisms that maintain disseminated tumor cells in a dormant state or result in their transition to metastatic growth remain largely unknown. This phenomenon has been extremely difficult to study in human patients^{4,12} and few preclinical models have been developed to address this issue. Nevertheless, some *in vivo* and *ex-vivo* model systems for tumor dormancy have been characterized (reviewed in^{1,12}). However, the *in vivo* models for tumor dormancy can primarily be utilized to validate potential mechanisms regulating tumor dormancy, but are not amenable to explore in real time the biology of a single disseminated dormant tumor cell.

The 3-D system presented here models for the first time solitary tumor dormancy and the switch to metastatic growth in an *in vitro* model system. The proliferation assay for modeling tumor dormancy (quiescence) and the transition to proliferation in the 3D system can be followed over time. This assay can be utilized to study in a high throughput manner and in a relatively short time frame potential factors/genes that maintain dormant tumor cells in their quiescent phase or may activate them to proliferate. Furthermore, the proliferation assay can be utilized as a high throughput platform to screen for additional tumor cell lines that may have a dormant phenotype.

Studying the molecular mechanisms which governs tumor dormancy or its switch to metastatic growth in the 3D system by conventional biochemical methods (RT-PCR/ western blots) is difficult given the low amount of RNA/protein that can be extracted for biochemical studies from the dormant tumor cells. However, immunofluorescence staining of cell signaling molecules of the dormant and outbreking tumor cells in the 3D model system, as presented in Figures 3 and 4 can be applied. Hence, the model system presented here can serve as an efficient tool to begin to explore the molecular mechanisms regulating solitary tumor cell dormancy and the transition to metastatic growth.

Disclosures

No conflicts of interest declared.

Acknowledgements

This research was supported in part by the Intramural Research Program of the National Cancer Institute.

References

1. Aguirre-Ghiso, J. A. Models, mechanisms and clinical evidence for cancer dormancy. *Nat Rev Cancer*. **7**, 834-846 doi:nrc2256 [pii] 10.1038/nrc2256 (2007).
2. Pantel, K. & Woelfle, U. Micrometastasis in breast cancer and other solid tumors. *J Biol Regul Homeost Agents*. **18**, 120-125 (2004).
3. Naumov, G. N. et al. Ineffectiveness of doxorubicin treatment on solitary dormant mammary carcinoma cells or late-developing metastases. *Breast Cancer Res Treat*. **82**, 199-206 (2003).
4. Klein, C. A. Framework models of tumor dormancy from patient-derived observations. *Curr Opin Genet Dev*. doi:S0959-437X(10)00189-9 [pii] 10.1016/j.gde.2010.10.011 (2010).
5. Naumov, G.N., et al. Persistence of solitary mammary carcinoma cells in a secondary site: a possible contributor to dormancy. *Cancer Res*. **62**, 2162-2168 (2002).
6. Townson, J.L. & Chambers, A.F. Dormancy of solitary metastatic cells. *Cell Cycle*. **5**, 1744-1750, doi:2864 [pii] (2006).
7. Chambers, A. F., Groom, A.C., & MacDonald, I.C. Dissemination and growth of cancer cells in metastatic sites. *Nat Rev Cancer*. **2**, 563-572 (2002).
8. Pantel, K., et al. Differential expression of proliferation-associated molecules in individual micrometastatic carcinoma cells. *J Natl Cancer Inst*. **85**, 1419-1424 (1993).
9. Demicheli, R. Tumour dormancy: findings and hypotheses from clinical research on breast cancer. *Semin Cancer Biol*. **11**, 297-306 (2001).
10. Braun, S., et al. A pooled analysis of bone marrow micrometastasis in breast cancer. *N Engl J Med*. **353**, 793-802 (2005).
11. Pantel, K., Alix-Panabieres, C. & Riethdorf, S. Cancer micrometastases. *Nat Rev Clin Oncol*. **6**, 339-351 doi:nrclinonc.2009.44 [pii] 10.1038/nrclinonc.2009.44 (2009).
12. Goss, P. E. & Chambers, A. F. Does tumour dormancy offer a therapeutic target? *Nat Rev Cancer*. **10**, 871-877 doi:nrc2933 [pii] 10.1038/nrc2933 (2010).
13. Mendoza, A., et al. Modeling metastasis biology and therapy in real time in the mouse lung. *J Clin Invest*. **120**, 2979-2988 doi:40252 [pii] 10.1172/JCI40252 (2010).
14. Naumov, G.N., et al. A model of human tumor dormancy: an angiogenic switch from the nonangiogenic phenotype. *J Natl Cancer Inst*. **98**, 316-325 (2006).
15. Barkan, D., et al. Metastatic growth from dormant cells induced by a col-I-enriched fibrotic environment. *Cancer Res*. **70**, 5706-5716, doi:0008-5472.CAN-09-2356 [pii] 10.1158/0008-5472.CAN-09-2356 (2010).
16. Barkan, D., Green, J.E., & Chambers, A.F. Extracellular matrix: A gatekeeper in the transition from dormancy to metastatic growth. *Eur J Cancer*. doi:S0959-8049(10)00154-1 [pii] 10.1016/j.ejca.2010.02.027 (2010).
17. Barkan, D., et al. Inhibition of metastatic outgrowth from single dormant tumor cells by targeting the cytoskeleton. *Cancer Res*. **68**, 6241-6250 doi:68/15/6241 [pii] 10.1158/0008-5472.CAN-07-6849 (2008).
18. Debnath, J., Muthuswamy, S.K., & Brugge, J.S. Morphogenesis and oncogenesis of MCF-10A mammary epithelial acini grown in three-dimensional basement membrane cultures. *Methods*. **30**, 256-268 (2003).
19. Morris, V.L., et al. Mammary carcinoma cell lines of high and low metastatic potential differ not in extravasation but in subsequent migration and growth. *Clin Exp Metastasis*. **12**, 357-367 (1994).
20. Khanna, C., et al. The membrane-cytoskeleton linker ezrin is necessary for osteosarcoma metastasis. *Nat Med*. **10**, 182-186 (2004).