

Modeling The Bone Marrow Microenvironment's Influence on Leukemic Disease

Evans R¹, Martin KH^{1,2}, Moses BS¹, Slone WL¹, Hare I^{1,3}, Piktel D¹, Thomas P¹ and Gibson LF^{1,3}

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The bone marrow microenvironment serves as both the site of initiation of the majority of hematopoietic malignancies and also contributes to maintenance of minimal residual disease by promoting biologically relevant changes in tumor cells. These functional alterations of leukemic cells include, but are not limited to modulation of cell cycle [1,2], regulation of anti-apoptotic signaling cascades [3-6], and influence on metabolic activity [7-9]. Of clinical relevance, these pathways impact on therapeutic response, making it critical to have robust *in vitro* systems to interrogate tumor cell interactions with stromal elements of the microenvironment to screen chemotherapeutic agents and inform the *in vivo* model design.

To generate a model of the marrow microenvironment niche human leukemic cells were co-cultured with human primary bone marrow stromal cells (BMSC). Frequently, applications require separation of tumor cells and stromal components for subsequent western blot, PCR, DNA or RNA based analysis. However, confocal microscopy provides the unique opportunity to study critical tumor: stromal cell interactions *in vitro* without physically detaching the tumor population prior to evaluation. This approach is valuable given the transient nature of cell signaling that may be immediately altered upon physical removal of leukemic cells from niche derived stroma. With this approach

- 1 Alexander B. Osborn Hematopoietic Malignancy and Transplantation Program of the Mary Babb Randolph Cancer Center, Robert C. Byrd Health Sciences Center, West Virginia University School of Medicine, USA
- 2 Department of Neurobiology and Anatomy, West Virginia University School of Medicine, USA
- 3 Department of Microbiology, Immunology and Cell Biology, Robert C. Byrd Health Sciences Center, West Virginia University School of Medicine, USA

Corresponding author: Gibson LF

✉ lgibson@hsc.wvu.edu

Mary Babb Randolph Cancer Center, Robert C. Byrd Health Sciences Center, West Virginia University School of Medicine, P.O. Box 9300, Morgantown, WV 26506-9300, USA.

Tel: 304 293 1547

Fax: 304 293 4667

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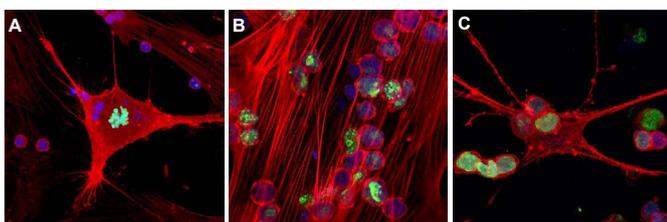


Figure 1 Ki67⁺ evaluation of human leukemic and bone marrow stromal cells (BMSC). Nalm-27 Ph⁺ ALL leukemic cells (Fujisaki Cancer Center) co-cultured with a dividing Ki67⁺ human primary BMSC (A), Nalm-27 cells co-cultured with confluent BMSC representing one structural component of the bone marrow microenvironment (B) and Nalm-27 tumor cells in a 3-dimensional static co-culture with BMSC (C) to model architecture of the niche.

a variety of targets are amendable to evaluation, with Ki67 shown in Figure 1 as just one example.

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