

**Title: Exosomes released by highly migratory premalignant lung epithelial cells facilitate epithelial-mesenchymal transition and migration of unselected slow migratory cells**

**Authors: Tieyi Li <sup>1</sup>, Steven M. Dubinett <sup>2,3,4,5,6</sup>, Ken Yoneda<sup>7</sup>, Davis Gandara<sup>8</sup>, Suzanne Miyamoto<sup>8</sup>, Manash K. Paul <sup>2\*</sup>, Ya-Hong Xie <sup>1\*</sup>**

<sup>1</sup> Materials Science and Engineering Department, Henry Samueli School of Engineering, University of California Los Angeles, Los Angeles, California 90024, United States.

<sup>2</sup> Department of Medicine, Division of Pulmonary and Critical Care, David Geffen School of Medicine at UCLA; 10833 Le Conte Avenue, 43-229 CHS, Los Angeles, CA 90095-1690, USA

<sup>3</sup> Department of Medicine, VA Greater Los Angeles Healthcare System; 11301 Wilshire Boulevard, Los Angeles, CA 90073, USA

<sup>4</sup> Department of Molecular and Medical Pharmacology, David Geffen School of Medicine at UCLA; 650 Charles E. Young Drive South, 23-120 CHS, Box 951735, Los Angeles, CA 90095-1735, USA

<sup>5</sup> Department of Pathology and Laboratory Medicine, David Geffen School of Medicine at UCLA; 757 Westwood Plaza, Los Angeles, CA 90095, USA

<sup>6</sup> Jonsson Comprehensive Cancer Center, UCLA; 8-684 Factor Building, Box 951781, Los Angeles, CA 90095-1781, USA

<sup>7</sup> Pulmonary Medicine, University of California, Davis, Sacramento, CA, 95817, USA

<sup>8</sup> Division of Hematology and Oncology, University of California, Davis, Sacramento, CA, 95817, USA

\* Corresponding authors

**Abstract:**

Non-small cell lung cancer (NSCLC) recurs in 30%-55% of patients, and the metastatic process occurs early in the development of the disease. Exosomes are cell-derived nanovesicles that have the ability to induce a process called epithelial-mesenchymal transition (EMT), which may be involved in the development of field cancerization. Thus, we propose that using exosome-based liquid biopsy has the potential to identify early metastases in NSCLC.

We used human bronchial epithelial cells (HBEC) that exhibit activated Kras-G12D and p53 knockdown, representing a state of premalignancy. Using a novel "constricted migration"-based selection method, we have identified a distinct subgroup of premalignant, high-risk HBECs (highly motile or HM) that exhibit exceptional mobility both in vitro and in vivo. Thereby providing a distinctive platform for investigating premalignant cell migration and early metastasis. Exosomes derived from HM-HBECs have unique molecular characteristics and induce the unselected (UN or low migratory)-HBECs to acquire an HM phenotype characterized by heightened EMT, accelerated migration, and augmented invasive potential in vitro.

We used Surface-enhanced Raman spectroscopy (SERS) to derive exosomal spectroscopic signals to distinguish between HM-HBEC and UN-HBEC exosomal signatures. Then, we used machine learning (ML)-based approaches to differentiate the Raman fingerprints of HM and UN exosomes. These fingerprints were then integrated into a training dataset and evaluated using ML to assess their distinguishability, resulting in a high accuracy of over 85% for separating the two datasets. We are now in the process of enlarging our training dataset and optimizing the hyperparameters to train the ML model using human pleural effusion (metastatic)-derived exosomes. ML-based spectroscopic analysis has the potential to aid in the early detection and interception of metastatic lung cancer.