

FAM3C encapsulated in circulating tumor-derived extracellular vesicles promotes distant growth in lung cancers

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Tumor-derived EVs containing FAM3C trigger aggressiveness of lung carcinoma cells in vivo. (A) Tail vein injection of A549 lung carcinoma cells in mouse model after 2 weeks pre-educated with PBS, EVs^{Control} or EVs^{FAM3C}. Animals were sacrificed after week 7. (B) Representative images of cytokeratin (red) stained tissues. Tumor nodules are stained pink with lacy appearance. Black circle, small tumor cell aggregates highlighted as dark red clusters; N=normal lung parenchyma. Counts for tumor aggregates were tabulated as median \pm interquartile range (n=4 for PBS, n=5 for EVs^{Control} and EVs^{FAM3C}). (C) Representative images of lung sections stained for FAM3C (brown) and Ki67 (red) positivity. Original images were captured at 20x magnification; box showed cropped image of the indicated area prior to image size reduction. Credit: *Theranostics* (2023). DOI: 10.7150/thno.72297

Metastasis, the process by which cancer spreads beyond its site of origin, is associated with poor clinical prognosis, cancer mortality, and resistance to therapy. Non-small cell lung cancer (NSCLC) in particular, has a high metastasis potential, and accounts for the highest cancer mortality worldwide. Hence, it is important to understand the molecular underpinnings driving metastasis—which up until now remains obscure—in order to design therapies for early intervention.

In this study, a group of researchers and clinicians, led by Professor Goh Boon Cher from the Cancer Science Institute of Singapore at the National University of Singapore, discovered that extracellular vesicles (EVs) released from tumors may be responsible for introducing growth and [metastasis](#) potential to recipient cells.

Through profiling tumor-derived EVs with unbiased proteomic mass spectrometry, the research team identified an enriched set of cargo molecules which include FAM3C protein encoded by Interleukin-like

EMT inducer (ILEI) was overexpressed in NSCLC tumors, which was strongly correlated with adverse clinical outcomes and [cancer](#) metastasis.

Abnormal expression of FAM3C in lung cancer cells augmented cellular transformation and stimulated distant lung tumor colonization in preclinical models. More importantly, the researchers identified a previously unknown mode of microenvironmental crosstalk involving FAM3C in EVs, in which the delivery and uptake of FAM3C via tumor-derived EVs (TDEs) promotes oncogenic signaling in recipient cells. Their findings were published in *Theranostics*.

The researchers' findings help to clarify the previously poorly-characterized mechanisms of carcinogenesis and metastasis driven by FAM3C. Furthermore, the study expanded the current knowledge on cell autonomous oncogenic signaling by demonstrating the cell-to-cell crosstalk via EVs. Although this study concentrated on [lung cancer](#), it may be pertinent to other cancers in which FAM3C is highly expressed, especially in the EVs of cancer patients.

Further investigations into the role of FAM3C in other carcinomas would be highly beneficial towards planning treatment strategies, as stratification of patients based on FAM3C expression in plasma EVs may pave the path for developing therapeutic strategies against widespread tumor metastasis.

More information: Win Lwin Thuya et al, FAM3C in circulating tumor-derived extracellular vesicles promotes non-small cell lung cancer growth in secondary sites, *Theranostics* (2023). [DOI: 10.7150/thno.72297](#)

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