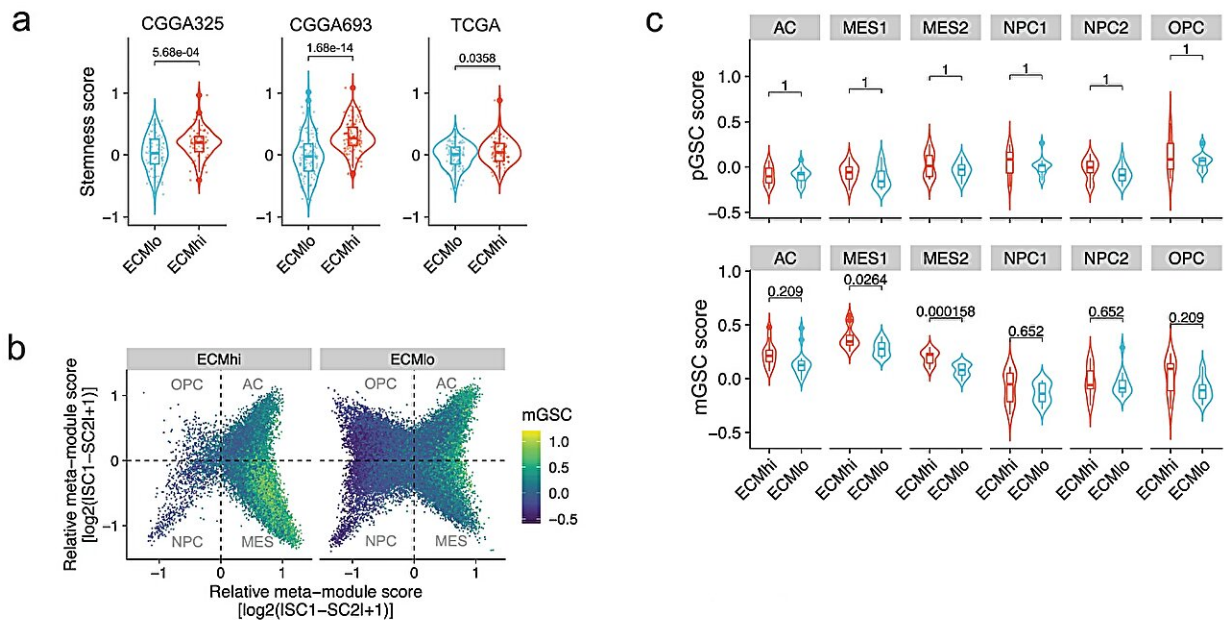


Understudied cell in the brain could be key to treating glioblastoma

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Perivascular fibroblasts secrete factors that upregulate stem-like programs in ECM^{hi} glioblastoma. **a** Stemness signature scores in ECM^{hi} and ECM^{lo} tumors in TCGA ($n = 175$), CGGA325 ($n = 325$) and CGGA693 ($n = 693$) GBM cohorts. **b** Two-dimensional butterfly plot of glioma cell states in ECM^{hi} and ECM^{lo} tumors, colored by enrichment of the mesenchymal glioma stem cell (mGSC) signature. OPC—oligodendrocyte progenitor cell-like; AC astrocyte-like; NPC—neural progenitor cell-like; MES—mesenchymal-like. **c** Proneural (top) and mesenchymal (bottom) glioma stem cell scores in ECM^{hi} and ECM^{lo} tumors. Two-sided t-test. Benjamini-Hochberg adjusted P values are shown. Credit: *npj Genomic Medicine* (2023). DOI: 10.1038/s41525-023-00381-w

Glioblastoma is one of the most treatment-resistant cancers, with those diagnosed surviving for less than two years. In a [new study](#) published in *npj Genomic Medicine*, researchers at the University of Notre Dame have found that a largely understudied cell could offer new insight into how the aggressive, primary brain cancer is able to resist immunotherapy.

"A decade ago, we didn't even know perivascular fibroblasts existed within the brain, and not just in the lining of the skull," said Meenal Datta, assistant professor of aerospace and [mechanical engineering](#) at Notre Dame and senior author on the study. "My lab's expertise is examining tumors from an engineering and systems-based approach and looking at the novel mechanical features in rare cancers that may have been understudied or overlooked."

Using standard bioinformatics and newer AI-based approaches, Datta's TIME Lab began analyzing different genes expressed in the [tumor](#) microenvironment related to the [extracellular matrix](#)—or the scaffolding cells create to support future cell adhesion, migration, proliferation and differentiation—and other various cell types.

What they found was a surprising, fairly new cell type: perivascular fibroblasts. These fibroblasts are typically found in the blood vessels of a healthy brain and deposit collagen to maintain the [structural integrity](#) and functionality of brain vessels.

"It was a serendipitous discovery," said Maksym Zarodniuk, graduate student in the TIME Lab and the bioengineering doctorate program, and first author on the study. "We started in a completely different direction and stumbled upon this population of cells by using a combination of both bulk and single-cell RNA sequencing analyses of patient tumors."

In their data, researchers were able to identify two groups of patients: those with a higher proportion of perivascular fibroblasts and those with

significantly less. They found that brain [cancer](#) patients with more perivascular fibroblasts in their tumors were more likely to respond poorly to immunotherapies and have poor survival outcomes.

When exploring how this is possible, the researchers found that perivascular fibroblasts support the creation of an immunosuppressive [tumor microenvironment](#), allowing the cancer to better evade the immune system. The fibroblasts may also help the cancer resist therapies—such as chemotherapy that targets dividing cells—by promoting stem-like cancer cells that rarely divide, which are believed to be a major source of tumor relapse and metastasis.

"Moving forward, we want to do new experiments to confirm what we found in this paper and provide some good ground to start thinking about how to improve response to immunotherapy," Zarodniuk said.

Because perivascular fibroblasts are a part of a healthy brain's vasculature, Datta believes that these cells are breaking off and getting close to or infiltrating the glioblastoma tumor. However, instead of supporting healthy brain function, these fibroblasts are getting reprogrammed and helping the tumor instead.

"Most people think about the brain as being very soft, with soft cells and a soft matrix. But by putting down these fibroblasts and making these very fibrous proteins, it gives us an entirely different perspective on the structure of the [brain](#) and how it can be taken advantage of by cancer cells originating in the same organ," Datta said.

More information: Maksym Zarodniuk et al, CNS tumor stroma transcriptomics identify perivascular fibroblasts as predictors of immunotherapy resistance in glioblastoma patients, *npj Genomic Medicine* (2023). [DOI: 10.1038/s41525-023-00381-w](https://doi.org/10.1038/s41525-023-00381-w)

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