

International 'A' team debuts brain cancer atlas

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It takes an "A" team to make headway against glioblastoma, a highly aggressive type of brain cancer. Glioblastoma is the most common type of malignant brain tumor in adults. In addition to the caliber of the researchers involved, in this case "A" also stands for atlas.

A key member of the team, Jill S. Barnholtz-Sloan, Ph.D., Sally S. Morley Designated Professor in Brain Tumor Research at Case Western Reserve University School of Medicine, and approximately 80 other internationally renowned neurologists, bioinformaticians, and pathologists from the United States and India recently published details of the Ivy Glioblastoma Atlas in *Science*.

The atlas maps out comprehensive, visually-rich information on the anatomic and genetic bases of [glioblastoma](#) at the cellular and molecular levels associated with the disease. It is aimed at helping researchers and physicians improve the diagnosis and [treatment](#) of glioblastoma, including finding new drug targets. While Atlas-based data have been previously available, the new paper in *Science* represents the first published research outlining the project in detail.

According to Barnholtz-Sloan, there are three aims to the venture. The first is to make all the relevant data fully transparent and available. The second is to create a road map of all of the different cells and potential [molecular changes](#) that occur in these cells in a glioblastoma. And the third, is to build a better understanding of how much heterogeneity can be found in these tumors.

"Currently we provide the same standard treatment of surgery, followed by radiation plus chemotherapy, for all glioblastoma patients," says Barnholtz-Sloan, who is also professor and associate director for bioinformatics/translational informatics at CWRU's School of Medicine. "But the tumors are not all the same. They have different molecular changes, which means that we may need to provide separate, tailored treatments to tackle each one. By identifying unique features of these tumors that may benefit from targeted therapies, the Atlas will enable patients to experience the benefits of precision medicine, increasing the chances for better response to treatment and hence better survival."

In virtually all cases of glioblastoma, excised tumors are assessed by highly specialized pathologists, who confirm the presence of malignancy and provide details about the unique characteristics of the tumors they examine. For this project, Barnholtz-Sloan assisted the team in developing a study design to assess the concordance between features outlined in the tumors by both these human experts and automated machine-learning techniques. "In many ways it's a belt and suspenders approach," she says. "The gold standard is the pathologist. Machine-driven analysis adds value by bringing further information to the findings of the pathologists. In the article we published, we found that the concordance rates were very high; 85-95 percent."

By way of comparison, Barnholtz-Sloan cites the case of medulloblastoma, a common type of [malignant brain tumor](#) mostly seen in children. In medulloblastoma, researchers have been able to group tumors into four distinct molecularly-defined groups, which have specific drugs that would work best within each subtype. The new Atlas hopes to replicate this success in glioblastoma.

"As more new information is added to the Atlas and the associated database for clinical and genomic data, we believe that it will continue to be very valuable research and treatment tool, allowing more refined diagnosis and treatment and enabling researchers to generate potentially life-extending hypotheses about causes and treatments," said Barnholtz-Sloan.

More information: Ralph B. Puchalski et al, An anatomic transcriptional atlas of human glioblastoma, *Science* (2018). [DOI: 10.1126/science.aaf2666](https://doi.org/10.1126/science.aaf2666)

Provided by Case Western Reserve University

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