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# The Sybil prophecy: Searching for predictors of response to bevacizumab in glioblastoma Get access

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#### **Extract**

We read with great interest the work by Hiller-Vallina et al. on the role of a sexual-dependent necroinflamed pattern in predicting prognosis and response to anti-angiogenic drugs in glioblastoma. The work has huge strong points, including its elegant and meticulous design, and also opens a number of issues. Female gender has not traditionally been recognized as a positive prognostic factor in glioblastoma, yet evidence is rising that this assumption is true. In the present paper, Hiller-Vallina et coworkers demonstrate that, in a subset of male patients, low levels of estrogen receptor-α (encoded by ESR1 gene) determine a necroinflamed phenotype, with activation of VEGF/hypoxia-related pathways and infiltration of S100A9-positive myeloid cells. Male patients harboring low-ESR1 necroinflamed glioblastomas have a worse prognosis than male and female patients harboring high-ESR1 non-necroinflamed tumors. On the other hand, necroinflamed tumors show a better response to anti-angiogenic treatment. Much work remains to be done to dissect into detail the relationship between sex hormones, tumor immune infiltration and hypoxic/angiogenic microenvironment. Our research group recently investigated the prognostic role of the immune infiltrate in glioblastoma, with focus on the tissue-resident memory (Trm) CD8 + CD103 + T lymphocytes, finding that high levels of Trm expressing PD1 or other immune checkpoints are correlated to worse prognosis, likely due to immune cell exhaustion. We reanalyzed survival data of patients included in that study and, interestingly, we found that the proportion of exhausted CD8 + CD103 + PD1 + Trms portends a worse prognosis only in male patients (Figure 1). Such finding corroborates the assumption by Hiller-Vallina et al. on the existence of two distinct subsets of GBM in males, with different prognosis. Notably, PD1 expression on Trm has been linked to persistent neuroinflammation in encephalitis,<sup>3</sup> further supporting the assumption that immune-exausted Trms can be enriched in necroinflammed glioblastoma. The issue of the improved response of male patients with necroinflamed glioblastoma to anti-angiogenic treatment with bevacizumab is more puzzling. Interestingly, no consistent predictors of response to bevacizumab in glioblastoma have been ever established. A post-hoc analysis of the AVAglio trial identified the proneural signature as associated with better response to bevacizumab. However, it is well known that the expression of inflammation-related players is downregulated in proneural and upregulated in mesenchymal glioblastoma, in contrast with the findings of the present study. <sup>5</sup> The role of VEGF expression is also debated. Hiller-Vallina et al<sup>1</sup> reported that necroinflamed glioblastoma upregulated VEGF, yet VEGF levels as assessed with RT-PCR were not associated with improved survival after bevacizumab treatment. Instead, in a work on 25 recurrent GBM patients, we had shown that high VEGF levels were associated with worse survival after bevacizumab treatment in recurrent glioblastoma. To conclude, the work here

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discussed has the merit to enliven the interest of the neuro-oncologic community on the interplay between gender and immune infiltrate in glioblastoma and on the identification of prognostic factors for response to anti-angiogenic treatment, which are long awaited.

Issue Section: Letter to the Editor

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