

Letter: Is Developmental Venous Anomaly an Imaging Biomarker of *PIK3CA* Mutated Gliomas?

To the Editor:

Phosphoinositide 3-kinase catalytic subunit alpha (*PIK3CA*) gene encodes the p110 α catalytic subunit of phosphoinositide 3-kinase (PI3K), which phosphorylates phosphatidylinositol to generate phosphatidylinositol 3,4,5-trisphosphate (PIP3). *PIK3CA*, a component of the PI3K/protein kinase B (AKT)/mammalian target of rapamycin (*mTOR*) signalling pathway, is considered as a major oncogene in human cancers. Missense mutations include especially the E542K and E545K substitutions in the helical domain and the H1047R change in the kinase domain. Cell-based analyses confirmed that these hotspot mutations confer transformation via constitutive activation of p110 α . In cerebral gliomas, *PIK3CA* mutations have been observed in 17% to 27% of adult cases and in 21% of pediatric cases.^{1,2} *PIK3CA* postzygotic mutations have also been reported in congenital mosaic overgrowth syndromes: 1) congenital lipomatous overgrowth with vascular, epidermal, and skeletal anomalies (CLOVES) syndrome; 2) megalencephaly syndromes; 3) Cowden-like syndromes; and 4) macrodactyly. In addition, somatic mutations of the *PIK3CA* gene in lymphatic or vascular endothelial cells are associated with the development of lymphatic³ and venous malformations.⁴ The latter may explain the clinical observation of the association between the presence of facial venous malformations and cerebral developmental venous anomalies (DVAs).⁵

We have recently observed an increased prevalence of DVAs in adult patients with diffuse gliomas (21.5%-23.5%)⁶ and in pediatric patients with diffuse intrinsic pontine gliomas (24.1%), in contrast to other cerebral neoplasms.⁷ This association suggests a possible common underlying role of *PIK3CA* and related mutations in the development of DVAs and gliomas. Interestingly, the observed increase in DVA frequency is 2-fold in association with gliomas and facial venous malformations. This can be of great clinical and practical relevance since the identification of a brain DVA in the MRI workup of a glioma patient may serve as an imaging biomarker of a mutation in the *PIK3CA* gene.

To this end, we propose to perform accurate RNA-sequencing and/or DNA-sequencing (targeted panels or whole exome sequencing) on formalin-fixed and paraffin-embedded or frozen cerebral DVA-associated glioma samples to uncover somatic genetic changes and especially to look for postzygotic mutations in *PIK3CA*.

Disclosures

The authors have no personal, financial, or institutional interest in any of the drugs, materials, or devices described in this article.

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