Subependymal giant cell astrocytoma in the absence of tuberous sclerosis complex: case report

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The authors report the case of a 14-year-old male with a subependymal giant cell astrocytoma (SEGA) that occurred in the absence of tuberous sclerosis complex (TSC). The patient presented with progressive headache and the sudden onset of nausea and vomiting. Neuroimaging revealed an enhancing left ventricular mass located in the region of the foramen of Monro with significant mass effect and midline shift. The lesion had radiographic characteristics of SEGA; however, the diagnosis remained unclear given the absence of clinical features of TSC. The patient underwent gross-total resection of the tumor with resolution of his symptoms. Although tumor histology was consistent with SEGA, genetic analysis of both germline and tumor DNA revealed no TSC1/2 mutations. Similarly, a comprehensive clinical evaluation failed to reveal any clinical features characteristic of TSC. Few cases of SEGA without clinical or genetic evidence of TSC have been reported. The histogenesis, genetics, and clinical approach to this rare lesion are briefly reviewed.

http://thejns.org/doi/abs/10.3171/2015.1.PEDS13146

KEY WORDS mosaicism; subependymal giant cell astrocytoma; SEGA; tuberous sclerosis complex; TSC; oncology

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ties without pronator drift. No cutaneous stigmata of tuberous sclerosis, such as hypomelanotic macules (“ash-leaf spots”), shagreen patches, or sebaceous adenomas, were observed.

Neuroimaging demonstrated a cystic, multilobulated enhancing mass within the left lateral ventricle adjacent to the foramen of Monro, associated with contralateral ventriculomegaly. Marked cystic enhancement was seen on postcontrast CT, with obliteration of the left frontal horn of the lateral ventricle and perilesional edema throughout the left frontal lobe (Fig. 1). The lesion was isointense to the brain on T1-weighted MRI, with moderate edema throughout the left frontal lobe on noncontrast T2-weighted sequences (Fig. 2A and B). Postcontrast sequences demonstrated a mixed cystic and solid mass with avid enhancement and extension into the third ventricle (Fig. 2C and D). The radiographic differential diagnosis included SEGA, central neurocytoma, choroid plexus tumor, astrocytoma, and meningioma.

Operation

The patient underwent stereotactic left frontal craniotomy via a transcortical approach with gross-total excision of the lesion. A septostomy was performed to provide an additional CSF outflow tract given the marked involvement of the left lateral ventricle and foramen of Monro. An external ventricular drain was placed under direct visualization at the conclusion of the procedure. An intraoperative frozen section suggested a primary glioneuronal tumor.

Postoperative Course

The patient experienced no postoperative complications. His external ventricular drain was weaned without difficulty, and subsequently removed on postoperative Day 5. At the time of discharge from the hospital, the patient’s headaches had completely resolved. At his 4-month follow-up visit, his family described marked improvement in his inattention and impulsiveness. Follow-up neuroimaging at 1 year was without evidence of tumor recurrence or hydrocephalus (Fig. 3). A comprehensive clinical evaluation for TSC was performed including renal ultrasonography, echocardiography, electrocardiography, skeletal imaging, and ophthalmological examination. No features of TSC were identified.

Genetic Analysis

DNA sequencing of the TSC1 and TSC2 genes using peripheral blood failed to show mutations (Athena Diagnostics). At both loci, nonmutant wild-type alleles were observed. Tumor DNA was subsequently isolated from paraffin-embedded tissue and subjected to TSC1 and TSC2 sequencing. The patient was found to have nonmutant sequences in all exons of the TSC1 gene, except exons 8 and 16, which failed to demonstrate polymerase chain reaction products on multiple attempts (Center for Human Genetics). No pathogenic point mutations or deletions were seen in the TSC2 gene. Further analysis of the TSC1 gene was...
performed using multiple ligation probe amplification to detect whole-exon or whole-gene deletions or duplications. With this method, we confirmed amplification of exons 8 and 16 of the \textit{TSC1} gene, and both were nonmutant wild type. This was further confirmed with redesigned sequencing primers. Our analysis was expected to identify 85% of pathogenic \textit{TSC1/2} variants but would not detect mosaicism at the TSC locus. Further molecular studies were not possible given the depletion of all tumor specimens.

Pathological Findings

Histopathological examination of the tumor revealed a low-grade neoplasm comprised of glial cells with phenotypes varying from gemistocytic-like cells, ganglion-like cells, and spindle cells (Fig. 4 left). The cells had moderate to abundant eosinophilic cytoplasm with vesicular nuclei and occasional prominent nucleoli. Immunohistochemically, the cells exhibited faint staining for synaptophysin and were strongly positive for glial fibrillary acidic protein (GFAP) (Fig. 4 right). The neurofilament stain supported a solid growth pattern, while the chromogranin stain was negative. The overall MIB-1 proliferation index was low, reaching 4.8% in the most active area. Overall findings in conjunction with imaging studies were most consistent with SEGA, WHO Grade I.\textsuperscript{17,21}

Discussion

Subependymal giant cell astrocytoma is a rare central nervous system tumor with mixed glioneuronal features, most frequently seen in the setting of TSC.\textsuperscript{17} Approximately 5%–20% of patients with TSC develop SEGA,\textsuperscript{1} although solitary lesions without clinical or radiographic evidence of tuberous sclerosis have been reported.\textsuperscript{7,10,15,17,20,21} It is important to note that none of the previously reported cases were subjected to \textit{TSC1/2} gene sequencing, leaving the histogenesis of these particular lesions unclear. Nevertheless, the absence of clinical features of TSC in patients with solitary SEGA has been attributed to somatic mosaicism.\textsuperscript{13} However, both loss of heterozygosity and allelic mutation of \textit{TSC2} have been reported in a single solitary SEGA, suggesting that the genetic etiology may also be the co-occurrence of 2 de novo mutations, as opposed to somatic mosaicism.\textsuperscript{11} In the present case, the patient had a solitary SEGA with neither clinical features of TSC nor a detectable mutation of the \textit{TSC1} or \textit{TSC2} genes, despite genetic analysis of both germline and tumor using 2 complimentary sequencing methods.

Approximately 38 cases of SEGA in the absence of clinical features of TSC have been described,\textsuperscript{7,10,11,13,17–19,21} with the first case reported by Halmagyi and colleagues in 1979.\textsuperscript{7} Bonnin et al. reported on 22 cases of SEGA, and only 5 patients among these cases carried a clinical diagnosis of TSC.\textsuperscript{3} In a cohort of 23 patients with SEGA, clinical manifestations of TSC were identified in only 39% of the patients during the follow-up period.\textsuperscript{17} Interestingly, 1 patient did not develop symptoms of TSC until 10 years postoperatively. There have been 2 additional reports of SEGA in the absence of TSC until 10 years postoperatively. There have been 2 additional reports of SEGA in the absence of TSC,\textsuperscript{12,20} The first included a small case series of 6 patients diagnosed with SEGA, whereas the second identified the disorder at autopsy in a 75-year-old woman without any history or clinical features suggestive of TSC. The diagnoses in these reports were based solely on histopathology, as genetic analysis for the \textit{TSC1/2} mutations was not performed.

Only 8 patients with solitary SEGA have been studied using molecular or genetic techniques, 7 of whom demonstrated \textit{TSC1} or \textit{TSC2} germline mutations. Chan et al.\textsuperscript{4} reported 100% prevalence of \textit{TSC1/2} mutations in the pe-
ripheral blood of 7 patients with a histological diagnosis of SEGA. Notably, 1 patient also demonstrated a somatic TSC2 mutation that was undetectable in the germline (peripheral blood, hair, nails, buccal mucosa). Similarly, there is an additional case report describing a solitary SEGA with an isolated somatic TSC2 mutation. In the present case, DNA sequencing of both peripheral blood and tumor tissue failed to reveal mutations or large deletions in the TSC1 or TSC2 genes. The case represents the third reported instance of the histopathological diagnosis of SEGA in which no mutation of TSC1/2 could be identified. Although our genetic analysis was expected to detect at least 85% of pathogenic TSC1/2 mutations, it is possible we failed to detect novel single-base variants. Furthermore, our study was not designed to detect somatic mosaicism, which has been proposed as an etiology for solitary SEGA. Subependymal giant cell astrocytoma results from inactivation of the tumor suppressor genes TSC1 on chromosome 9q34 and/or TSC2 on chromosome 16p13, encoding the proteins tuberin and hamartin, respectively. The tuberin/hamartin complex functions as a suppressor of Ras homolog enriched in brain (RHEB), which directly activates the mammalian target of rapamycin (mTOR). The complex also inhibits cyclin-dependent kinase inhibitor 1B, which regulates cell cycle progression. Loss of upstream inhibition allows constitutive activation of mTOR and cell cycle progression leading to protein translation, cell growth, and proliferation. Interestingly, a posttranslational mechanism has been reported, with tuberin inactivated by phosphorylation. Although not yet explored, a potential epigenetic etiology of SEGA is intriguing to consider, particularly in the setting of a solitary lesion in the absence of features of TSC. Whether by promoter methylation, histone modifications, or nucleosome remodeling, both the solitary cases of SEGA and mutation-negative TSC could result from epigenetic silencing of several tuberin/hamartin complex components. In the present case, the molecular etiology of this solitary SEGA remains unknown.

References

Author Contributions
Conception and design: Beaumont, Smyth. Acquisition of data: Beaumont, Godzik. Analysis and interpretation of data: all authors. Drafting the article: Beaumont, Godzik. Critically revising the article: Beaumont, Dahiya, Smyth. Reviewed submitted version of manuscript: all authors. Approved the final version of the manuscript on behalf of all authors: Beaumont. Study supervision: Smyth.

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