CASE REPORT

# Successful treatment of a *TSC2*-mutant glioblastoma with everolimus

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Accepted 16 May 2019

#### **SUMMARY**

A 14-year-old boy with familial Li-Fraumeni syndrome presented with diplopia. Brain MRI revealed a right temporoparietal rim-enhancing mass. Following surgical resection and diagnosis of a gigantocellular-type glioblastoma multiforme (GBM), his family wished to avoid cytotoxic chemotherapy given the amplified risk of secondary malignancy. As such, we performed whole exome and transcriptome sequencing, which revealed germline TP53 and somatic TSC2 mutations. On completion of adjuvant radiotherapy, he was started on maintenance therapy with everolimus per recommendations from our multi-institutional brain tumour precision medicine tumour board. He has achieved a complete remission with resolution of visual symptoms and remains on everolimus therapy with concurrent electromagnetic field therapy, now 33 months from diagnosis. Our data highlight the benefit of precision medicine in children with GBM and offer insight into a targetable pathway that may be involved in similar cases.

#### **BACKGROUND**

Glioblastoma multiforme (GBM) is a rare and aggressive brain tumour, characterised as a WHO histologic grade IV tumour of glial origin. The vast majority of cases of GBM are observed in adults.<sup>2</sup> However, high-grade gliomas (HGGs) in general are the second most common malignant brain tumour type in the paediatric population, with an incidence of 0.8 per 100 000 for patients under the age of 19 years.3 Standard treatment for GBM at any age includes a combination of surgery, chemotherapy and radiation therapy. One-year overall survival for paediatric patients with HGG is estimated at 72%, and 1-year event-free survival is estimated at 49%, based on recent data from a cohort of 108 patients with anaplastic astrocytoma or glioblastoma; furthermore, 3-year overall survival is estimated at 28%, and 3-year event-free survival is estimated at 22%.

A histological subtype known as giant-cell GBM, characterised by the presence of multinucleated giant cells, abundant eosinophilic cytoplasm and a dense reticulin fibre network, has been well described. This histology is also believed to have a temporal lobe predilection. Giant-cell GBM only accounts for approximately 2%–5% of all GBM cases but appears to be more common in children and confers a superior, but still dismal, prognosis and life expectancy. Another similar histological

subtype, gigantocellular glioblastoma multiforme, characterised predominantly by bizarre, multinucleated giant cells, has been reported less frequently.<sup>9</sup>

Paediatric HGGs exhibit unique molecular genetics, and their phenotypes differ from their adult counterparts, despite histological similarities. These differences, taken together with the poor prognosis that is little changed over the past decade, motivate exploration of alternate treatment modalities to improve survival outcomes for paediatric patients with HGG in the future. The need for personalised, targeted treatment options for patients has led to the development of precision oncology with small molecular inhibitors directed at targets identified by tumour genomic sequencing. Here, we present an adolescent patient with a gigantocellular GBM and discuss our experience with targeted therapy.

#### **CASE PRESENTATION**

A 14-year-old previously healthy man presented with diplopia. Brain MRI (figure 1A) revealed a rim-enhancing right temporoparietal mass measuring  $2.7 \times 2.5 \times 3.2 \,\mathrm{cm}$  with significant oedema. He underwent gross total resection of the mass (figure 1B) and had near-immediate resolution of his diplopia. Pathology (figure 1C,D) revealed a glial neoplasm with gigantocellular morphology, no appreciable reticulin staining (not shown) and a Ki67 proliferation rate that approached 60% focally. favouring a diagnosis of GCG over pleomorphic xanthroastrocytoma. There was faint nuclear positivity for p53 and retained nuclear ATRX staining. Giant cells were weak to negative for glial fibrillary acidic protein (GFAP), while smaller, intermixed tumour cells were strongly positive for GFAP. The tumour was negative for BRAF V600E mutation by PCR. Total spine MRI 1 week postoperatively was negative for metastatic disease.

This patient's family history included a deceased father with leiomyosarcoma and a paternal grandfather with leiomyosarcoma and colorectal cancer. Both the patient and his brother were found to have a 5'UTR\_EX1 germline deletion in p53—the same as was identified in their father—confirming a diagnosis of familial Li-Fraumeni syndrome (LFS). Our central nervous system (CNS) tumour board recommended adjuvant radiation with consideration of adjuvant cytotoxic chemotherapy, according to Children's Oncology Group protocol ACNS0423. However, the patient's father had expressed his wish against the use of chemotherapy should either



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To cite: Zureick AH, McFadden KA, Mody R, et al. BMJ Case Rep 2019;12:e227734. doi:10.1136/bcr-2018-227734

## Novel treatment (new drug/intervention; established drug/procedure in new situation)

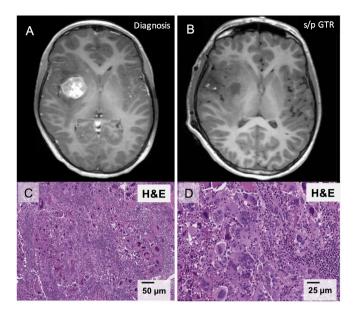
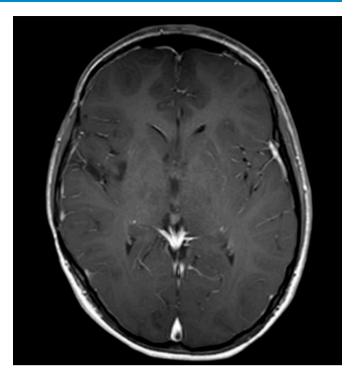


Figure 1 Gigantocellular glioblastoma imaging and pathology. (A) Brain MRI, axial image, T1 series postcontrast, illustrating right temporoparietal tumour prior to initial resection. (B) Brain MRI, axial image, T1 series postcontrast, showing the tumour bed following gross total resection. (C) H&E stain of the tumour at ×20 magnification reveals large cells with glassy eosinophilic cytoplasm and frequent multinucleation, bizarre nuclei and intranuclear inclusions. Numerous mitoses and significant lymphocytic infiltrates are also observed. (D) H&E stain at ×40 magnification shows multinucleated tumour 'giant cells' with atypical mitoses. GTR, gross total resection; s/p, status/post.

of his sons receive a cancer diagnosis. Given this wish, combined with the increased risk of second malignancy in the setting of LFS, he received 60.72 GyE of conformal proton radiotherapy to the right temporal region without concurrent or adjuvant chemotherapy.

We then performed whole exome (tumour and germline DNA) and transcriptome (tumour RNA) sequencing on his tumour to explore potential targeted therapy options. Sequencing revealed an R611W missense mutation in TSC2, homozygous deletion of PAK1 and copy-neutral loss of heterozygosity of TP53 in the tumour, as well as germline loss of exons 1 and 2 of TP53 (table 1). Additionally, the tumour was PTEN and IDH wild type. Results of his tumour sequencing were discussed in a CNS precision medicine tumour board teleconferenced with clinicians at multiple children's hospitals, which recommended everolimus, an mTOR inhibitor.

Table 1 Integrative sequencing results	
Mutation class	Gene/aberration
Copy number variation	Copy gain of chr1, 7, 9, 18 and X Copy loss of chr19 and 22 Homozygous deletion of <i>PAK1</i>
Somatic mutations	TSC2 (p.R611W), unknown significance
Gene fusions	No significant gene fusions
Outlier expression	N/A
Germline variants for disclosure	TP53, deletion of exons 1 and 2
Pathogens	N/A
N/A. not applicable.	



**Figure 2** MRI on maintenance everolimus brain MRI, axial image, T1 series postcontrast, taken 32 months after diagnosis of the patient's glioblastoma multiforme and after 24 months of maintenance therapy with everolimus.

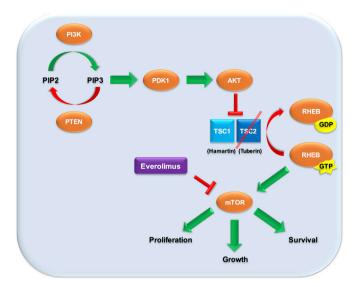
#### **OUTCOME AND FOLLOW-UP**

Five months after completion of radiation therapy, everolimus was initiated at 7.5 mg/day (patient weight=60.3 kg) and increased to 10 mg/day 4 months later. In addition, the patient was also approved for adjuvant electromagnetic field therapy with Optune (18 hours daily), based on the family's interest and promising results in adult GBM. The patient maintained complete remission and has survived 33 months since diagnosis (25 months on treatment with everolimus), with excellent quality of life and no adverse effects from everolimus other than occasional mouth sores. The patient's most recent brain MRI is shown in figure 2.

#### **DISCUSSION**

The giant-cell variant of GBM confers a similar, but slightly improved, dismal prognosis to the classic GBM histology, with a 5-year overall survival of 12.3% versus 3.4%.<sup>7</sup> A gigantocellular variant of GBM has also been reported, with one small paediatric series reporting a longer but not statistically significant difference in mean survival time for patients with gigantocellular GBM compared with patients with classic GBM.<sup>9</sup>

The mutations observed in this patient's GCG included germline *TP53* in the setting of familial LFS, as well as a somatic mutation in *TSC2*. *TP53* is a tumour suppressor gene that codes for the protein p53, which has multiple roles in genomic stability and antineoplastic progression. <sup>14</sup> LFS is a genetic disorder classically caused by germline mutations in *TP53* (75% of patients), and it is known to increase susceptibility to a wide array of cancers, including HGG. <sup>10 15</sup> Pollack *et al* reported in a series of HGG cases (anaplastic astrocytoma and GBM) that *TP53* mutations were observed in 40.5% of tumours, further affirming this association. <sup>16</sup>



**Figure 3** mTOR inhibition for *TSC2* mutant-driven oncogenesis loss of function of tuberin, the gene product of *TSC2*, prevents inhibition of downstream signals promoting cellular proliferation and oncogenesis. Downstream inhibition of the PI3K/AKT/mTOR pathway can be achieved with everolimus, an mTOR inhibitor. This is hypothesised to be the driving mutant in the presented patient's tumour. Original illustration by authors AHZ and CK.

TSC1 and TSC2 are two tumour suppressor genes implicated in tuberous sclerosis complex (TSC), a syndrome characterised by widespread growth of benign and malignant tumours in the brain (particularly subependymal giant-cell astrocytomas (SEGAs)), heart and kidneys, as well as by seizures, skin findings and intellectual disability. The gene products, hamartin and tuberin, respectively, function as cell cycle inhibitors within the PI3K/AKT/mTOR pathway (figure 3), and the TSC2 R611W point mutation observed here has also been previously reported. While most TSC-associated brain tumours are WHO grade I SEGAs, a limited number of cases of TSC-associated GBM have been previously reported.

The identification of a *TSC2* mutation in this patient's tumour led to an actionable change in management with the selection of everolimus, after consideration of multiple factors as a part of our institution's CNS Targeted Agent Prediction programme. <sup>19</sup> Everolimus inhibits the oncogenic pathway activated by a *TSC2* mutation, there was positive preclinical data, and paediatric phase I and CNS clinical data were available. <sup>20</sup> <sup>21</sup> Additionally, CNS penetration with everolimus is promising. In animal models, everolimus achieves a brain:plasma concentration ratio of 1.5%, and a brain concentration above the IC<sub>50</sub> for GBM cell lines for 12–24 hours postdose. <sup>22</sup> While everolimus is a relatively large molecule, the moderate lipophilicity and inhibitory effects on P-glycoprotein also result in accumulation in the CNS over time. <sup>22</sup>

In summary, we have presented a paediatric patient with a GCG and associated *TP53* and *TSC2* mutations who remains in complete remission 33 months after diagnosis, with surgical resection, radiation therapy, everolimus and electromagnetic therapy. We are hopeful that this treatment approach for a poor prognosis tumour will contribute to a better outcome for this patient and will further endorse the value of personalised, targeted therapy for patients with GBM in the future.

### **Learning points**

- ► Precision oncology is a needed area of research for personalised treatment of paediatric patients with cancer.
- ► We demonstrate here a successful outcome for a paediatric patient with glioblastoma treated with a targeted agent selected through results of tumour sequencing.
- Everolimus has good central nervous system penetration and may be efficacious in targeting the cellular pathway implicated by a TSC2 driving mutation.

**Acknowledgements** The authors thank the Michigan Center for Translational Pathology for whole exome and transcriptome tumour sequencing analysis through the Peds-MiOncoSeq programme. Lastly, the authors thank Bernard L Marini, PharmD, and Patricia L Robertson, MD, for their contributions to the manuscript and for serving as scientific advisors. The authors also thank the patient and his family.

**Contributors** AHZ: investigation, methodology, writing—original draft preparation, and writing—review and editing. KAM: resources and writing—review and editing. RM: writing—review and editing, and funding acquisition. CK: conceptualisation, supervision, writing—review and editing, project administration and funding acquisition.

**Funding** CK is supported by National Institutes of Health (NIH)/National Institute of Neurological Disorders and Stroke grant K08-NS099427-01, the UM Department of Pediatrics Chad Carr Pediatric Brain Tumor Center, the Chad Tough Foundation, the Catching Up with Jack Foundation, the Prayers from Maria Foundation, and the U CAN-CER VIVE Foundation. The University of Michigan Peds-MiOncoSeq study was supported by grant 1UM1HG006508 from the NIH Clinical Sequencing Exploratory Research Award (private investigator: Arul Chinnaiyan, coinvestigator: Rajen Mody).

Competing interests None declared.

Patient consent for publication Obtained.

Provenance and peer review Not commissioned; externally peer reviewed.

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## Novel treatment (new drug/intervention; established drug/procedure in new situation)

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