

Brainstem Glioblastoma Multiforme in a Patient with NF1

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Abstract. *This case report presents the first known case of a brainstem glioblastoma multiforme (GBM) in a patient with neurofibromatosis type 1 (NF1). While research has proposed that larger germ-line mutations in NF1 may be the driving factor that predisposes patients with NF1 to high-grade astrocytomas, this patient had a nonsense mutation in the NF1 gene, suggesting a variant tumorigenesis. Limited data on targeted immunotherapy for NF1 patients with a GBM have been reported and more data are required before targeted therapies could be proven as second-line treatment options.*

While associations between neurofibromatosis type 1 (NF1) and CNS tumors, such as optic pathway gliomas and low-grade brainstem gliomas are well established, the prevalence of glioblastoma multiforme (GBM) in NF1 is less common (1-3). However, patients with NF1 are at a 10- to 50-fold increased risk of developing high-grade gliomas compared to those without NF1 (4). Typically, GBMs in NF1 patients are treated with concurrent radiotherapy and temozolomide (TMZ), which is standard of care for sporadic GBM patients (5, 6).

Few cases of NF1-associated GBMs have been reported. NF1-associated GBMs have been reported in the frontal lobe (5, 7), occipital lobe (8), parietal lobe (9), cerebellum (10, 11), and thalamus (12). We present the case of a 23 year-old male with NF1 and a midbrain GBM. To our knowledge, none of the NF1-associated GBMs have been reported in the brainstem.

Case Report

A 23 year-old male with NF1 presented with 3 months of progressively worsening dizziness, vertigo, headaches,

increased sleep, slurred speech, and confusion. He also had double vision for 3 months, requiring a right eye patch. A post-contrast brain magnetic resonance imaging (MRI) showed a ring enhancing mass lesion in the midbrain (Figure 1).

Biopsy demonstrated a grade IV astrocytoma (GBM) that was *IDH* wild-type, H3K27 negative, and had *ATRX* loss. FoundationOne© genetic testing of the tumor tissue revealed genetic alterations of the *NF1* gene c. 3628G>T (E1210*) and c. 4234A>T (R1412*), *ATRX* gene 569delC (P190fs*16), and *CDKN2A/B* copy number loss. The *NF1* alterations were both nonsense mutations; c.3628G>T was identified at an allele frequency of 51.5% in the tumor, suggestive of germ line origin.

The patient received concurrent radiotherapy plus oral TMZ for 6 weeks followed by 12 cycles of adjuvant TMZ. The patient also elected to receive treatment with Optune. He subsequently developed significant steroid side effects as well as adrenal insufficiency.

Discussion

Brainstem gliomas that occur in patients with NF1 are typically low-grade, indolent, and generally have a better prognosis compared to patients with sporadic brainstem gliomas (13-15). Further, most of these lesions are discovered in childhood (1, 4, 16). High-grade astrocytomas, while rare, have been reported in NF1, but are almost always supratentorial (5, 7-9, 12). Two cases of infratentorial GBMs in patients with NF1 have been reported, both occurring in the cerebellum (10, 11).

Molecular analysis of NF1-related low- and high-grade astrocytomas showed *NF1* inactivation in all NF1-associated astrocytomas (12). Molecular analysis of NF1-associated GBMs has shown both an *NF1* deletion and either *CDKN2A* loss or homozygous chromosome 10q losses, which are commonly seen mutations in sporadic GBMs. Since both low-grade and high-grade astrocytomas in NF1 have *NF1* inactivation, but only the latter has additional tumor markers that are commonly seen in GBMs, it is suspected that their respective tumorigenesis occurs *via* distinct pathways.

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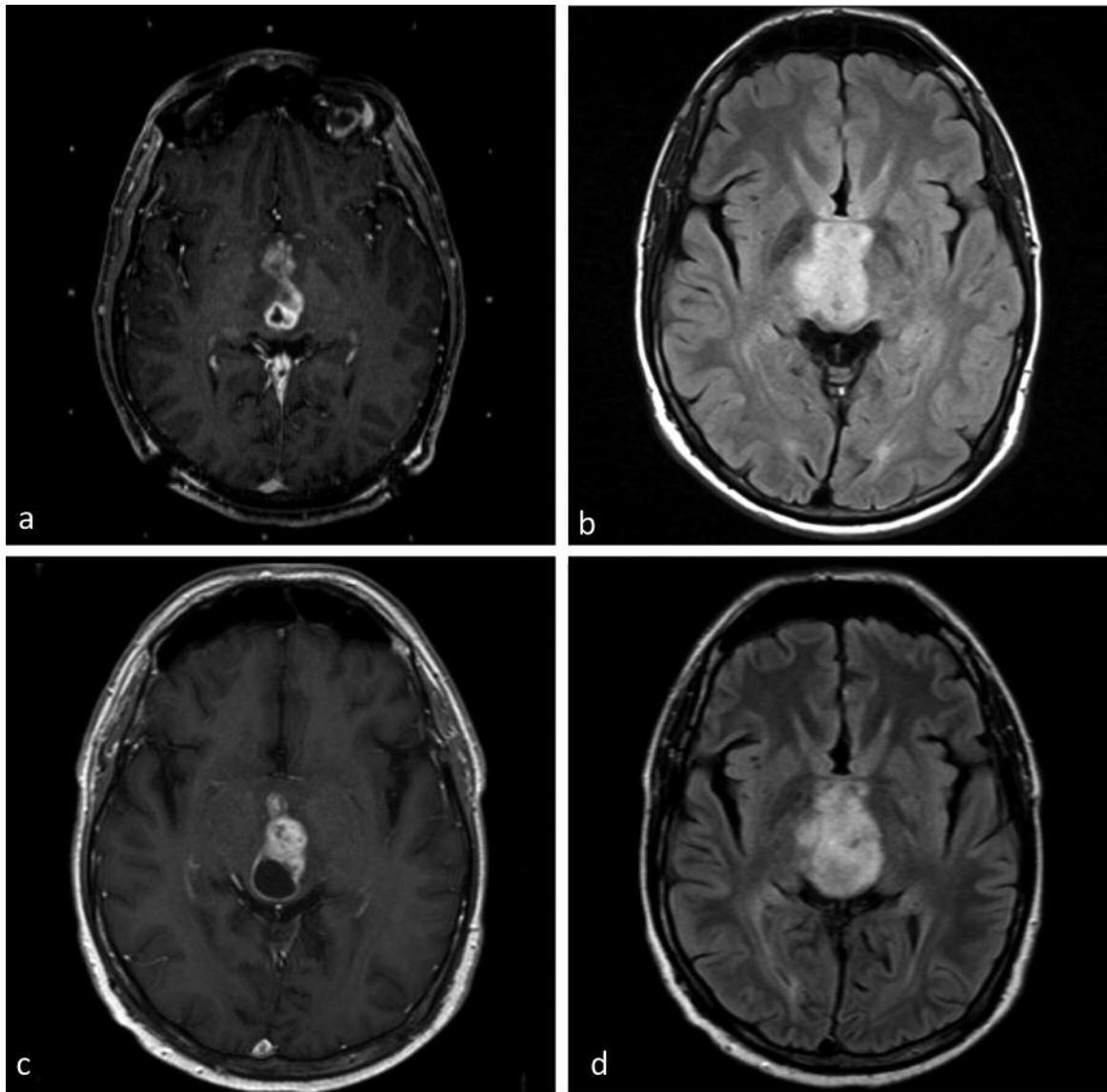


Figure 1. Brain MRI before and after treatment. A: Three months pre-treatment axial T1 post-contrast. B: Three months pre-treatment axial Flair. C: 5 months after treatment initiation axial T1 post-contrast. D: Five months after treatment initiation axial Flair.

Further, it has been proposed that large germ-line deletions of *NF1* may be the driving factor that places patients with *NF1* at an increased risk of malignant astrocytomas as opposed to low grade gliomas (12).

Our case has analogous findings of both *NF1* inactivation and loss of the *CDKN2A* tumor suppressor gene on chromosome 9. Our case had a nonsense *NF1* mutation rather than a large *NF1* deletion previously noted in other cases with high-grade astrocytoma. Thus, we propose protein-truncating mutations in *NF1* may yield similar

behaviors to large *NF1* deletions when occurring in conjunction with loss of *CDKN2A*.

Biopsy of a cerebellar GBM in 2009 revealed two distinct lesions: (1) a grade 4 astrocytoma (GBM) (+GFAP, -S-100, 30% immunopositivity for p53) and a diffuse neurofibroma (-GFAP, +S-100). First, this supports previous evidence that mutations in both *NF1* and p53 cooperate in the development of astrocytomas (12, 17, 18). Second, the authors hypothesized that the pre-existing neurofibroma may have created a favorable environment for tumor-initiating cells.

Thus they recommended close surveillance of existing lesions in *NF1* for the development of high grade astrocytomas (10). Our patient's GBM had no *p53* mutation. Though he refrained from MRI surveillance for the 4 years prior to his diagnosis, he had no previous MRI evidence of an infratentorial lesion; nor did his pathology or molecular reports indicate any separate cell type. Therefore, it is likely that his GBM developed *de novo*. Regardless, the utility of MRI surveillance for patients with *NF1* to monitor for both existing and *de novo* lesions is further reinforced.

While the molecular profiles of these tumors are of significant interest for the development of targeted treatment, more research is required. To date, treatment for *NF1* associated GBMs is identical to treatment for sporadic GBMs (5, 6). Targeted therapies, such as immune checkpoints inhibitors, are currently being assessed in clinical trials and could be beneficial for patients with NF1 associated GBMs (19). Of the cases discussed above, three received surgery followed by concurrent radiation and TMZ followed by monthly adjuvant TMZ. One patient lived 41 months after surgery (frontal GBM), (5) another 6 months after surgery (cerebellar GBM), (10) and the third was still alive at the time of publication (frontal GBM, no progression 9 months after surgery) (7). A separate case of an *NF1* associated cerebellar glioblastoma showed substantial improvement on MR imaging after 3 weeks of targeted immunotherapy with trametinib, a MEK inhibitor (11). Genomic testing for this tumor revealed no *IDH-1* mutation, with *NF1* splice site 3870 + 1G>T, *CDKN2A/B*. Trametinib was used after failure of the standard radiation and TMZ, carboplatin, and also progression on everolimus, an mTOR inhibitor. It should be noted that trametinib was chosen not based on a specific MEK amplification, but because of potential over-activity of the MEK pathway in *NF1* (11). More research is required to determine if additional NF1-associated GBM patients would benefit from similar immunotherapy.

Conclusion

In conclusion, we present a case of an *NF1*-associated midbrain GBM in young adulthood, which to our knowledge has not been previously reported. While data are limited, the molecular profile of this tumor is similar to other *NF1* related GBMs, having both an *NF1* mutation and *CDKN2A* loss. More research is required to determine if targeted therapies could provide benefit as second line treatment options.

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