CASE REPORT

Molecularly Targeted Treatments for NF1-Mutant Diffuse Intrinsic Pontine Glioma

Nicholas A. Vitanza,^{a,b,*} Hedieh Khalatbari,^c Ralph Ermoian,^d Jay Sarthy,^{a,b} Christina M. Lockwood,^e Bonnie L. Cole,^{f,g} and Sarah E.S. Leary^{a,b}

CASE DESCRIPTIONS

The first patient presented at 2 years of age when an MRI brain/spine revealed a nonenhancing, T2-hyperintense, expanded pons (Fig. 1A). A biopsy revealed a diffuse midline glioma, H3 K27M-mutant (DMG) (Fig. 2A). UW-Oncoplex[™], a targeted sequencing platform (1), identified a H3F3A p.K28M (i.e., H3.3 p.K27M) mutation, a novel mutation in NF1 (NF1 p.N2071D), and mutations in PIK3CA (PIK3CA p.E545K and p.G1050S). Without a well-suited clinical trial option, Seattle Children's Hospital (SCH) institutional Pediatric Brain Tumor Board recommended treatment per our Best-Available Therapy (BAT): focal radiation $(\sim 54 \text{ Gy})$ with concurrent temozolomide followed by maintenance bevacizumab, irinotecan, and temozolomide. The patient had near resolution of symptoms until progression 9 months from diagnosis (Fig. 1B).

The second patient presented at 11 years of age when an MRI brain/spine showed a

heterogeneously enhancing mass arising from the medulla as well as intracranial/spinal leptomeningeal metastases (Fig. 1D). A biopsy revealed a DMG (Fig. 2B). Cerebrospinal fluid cytology was negative for malignant cells. UW-Oncoplex[™] identified mutations in H3F3A p.K28M and NF1 (NF1 p.K1423E) along with a subclone mutation in KRAS (KRAS p.G13D). The patient received 36 Gy craniospinal radiation with a 54 Gy boost to the cervicomedullary junction and foci of spinal disease. An MRI 4 weeks following radiation revealed resolution of supratentorial/infratentorial leptomeningeal disease, but persistent medulla T2 signal abnormality. Only 4 weeks later, the patient's weakness worsened. An MRI revealed increased leptomeningeal disease with high spinal T2 signal (Fig. 1, E and F).

After progression of disease in each patient, the families were not interested in the early phase clinical trials for which the patient qualified, but the families did want to continue with a cancerdirected therapy that would minimally impact

^aDivision of Pediatric Hematology/Oncology, Department of Pediatrics, Seattle Children's Hospital, University of Washington School of Medicine, Seattle, WA; ^bFred Hutchinson Cancer Research Center, Seattle, WA; ^cDepartment of Radiology, Seattle Children's Hospital, University of Washington School of Medicine, Seattle, WA; ^dDepartment of Radiation Oncology, University of Washington, Seattle, WA; ^eDepartment of Laboratory Medicine, University of Washington and Seattle Children's Hospital, Seattle, WA; ^fDepartment of Laboratories, Seattle Children's Hospital, Seattle, WA; ^gDepartment of Anatomic Pathology, University of Washington School of Medicine, Seattle, WA.

^{*}Address correspondence to this author at: Seattle Children's Hospital, M/S MB.8.501, 4800 Sand Point Way, Seattle, WA 98105. Fax 206-987-3946; e-mail nicholas.vitanza@seattlechildrens.org.

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quality of life. Therefore, we were tasked with providing a tolerable, potentially disease-stabilizing or life-extending therapy in the setting of progressive DIPG. Considering the potential benefit of MEK inhibition in the setting of *NF1* mutations, trametinib, in combination with other therapy, was prescribed in each case. At progression, the first patient was prescribed focal re-irradiation (~20 Gy) (2)along with the MEK inhibitor trametinib (0.025 mg/kg daily). This combination resulted in symptom improvement (Lansky score increasing from 70 to 90) (Fig. 1C). She received



trametinib for 5 months with the only adverse effect of grade 1 rash, but died 16 months from diagnosis. At progression, the second patient began trametinib (0.025 mg/kg daily) and oral etoposide (50 mg/m² daily, every 14/28-days), after which there was a reduction in back pain and hemiparesis. An MRI 6 weeks later confirmed significant reduction in leptomeningeal disease (Fig. 1G). While she had neurologic improvement during treatment with trametinib and etoposide, she experienced a shunt infection and lower extremity edema. Several weeks later she developed progressive hydrocephalus and died 8 months from diagnosis.

CASE DISCUSSION

DIPG is a fatal pediatric central nervous system (CNS) tumor affecting ~300 children per year in the United States with a median overall survival of 11 months (*3*). As 85% of DIPG harbor mutations in genes encoding histone 3 also present in other gliomas, the 2016 World Health Organization (WHO) reclassification included a novel diagnosis: diffuse midline glioma, H3 K27M-mutant (*4*). Other

mutations, such as ACVR1, TP53, PDGFRA, KIT, KDR, and MYC, are not infrequent, although the less common mutations such as in NF1 or FGFR1 may represent targetable vulnerabilities (5). NF1 is a tumor suppressor gene and regulator of the RAS/ MAPK pathway. A MEK inhibitor was prescribed for our patients despite multiple targetable members of the pathway, as inhibiting MEK-driven phosphorylation of ERK in BRAF- and NF1-mutant tumors has been preclinically and clinically efficacious, particularly against low-grade glioma (LGG) (6). Of note, the specific NF1 mutation, NF1 p.N2071D, has not previously been described in patients with neurofibromatosis, type 1, or DIPG. While single-agent molecularly-targeted therapies may not be curative in high-grade CNS tumors, they may play a role in multi-agent regimens. The median time from progression to death for children with DIPG is 4.8 months (3). While our first patient survived 6.1 months after progression and our second patient survived 4.6 months after progression, the overall survival benefit of trametinib cannot be assessed in such a small patient cohort.

Our patients highlight several critical factors in the care of patients with DIPG/DMG. DIPG/DMG may disseminate metastatically/leptomeningeally, so patients should undergo an MRI of the spine at diagnosis and then as clinically indicated. We also recommend, when reasonable, to biopsy suspected DIPG/DMG as molecular characterization of these tumors is routine for clinical trials and allows for identification of targetable mutations. We also advocate for obtaining CSF cytology, when clinically reasonable, to evaluate for disseminated disease. Ultimately, CSF-derived circulating tumor DNA (ctDNA) could provide a less invasive molecular characterization of DIPG at diagnosis, as well as a route to track its molecular evolution during treatment via serial CSF monitoring. Critically, ctDNA may also provide an avenue to offer targeted therapies to children clinically unable to undergo biopsy or who are being treated at institutions without that neurosurgical capability (7). Seattle Children's has partnered with Tgen[®] to investigate serial sequencing of CSF and plasma ctDNA, as well as interrogation of extracellular vesicle cargo to not only quantify but to understand the functional state of DIPG cells during treatment (8). While the discovery of histone mutations amongst DIPG/DMG has been critical, their role in tumor initiation and treatment response remains elusive. While concurrent mutations in NF1 are less common, they are potentially targetable as MEK inhibition has become routine in Raspathway aberrant LGG after selumetinib demonstrated $69 \pm 9.8\%$ 2-year progression-free survival (6). Interestingly, there may be a correlation between a high/responsive MEK signature and concurrent Ras mutations (9). The presence of a KRAS mutation in our second patient, therefore, may have led to a response-predictive gene expression pattern that could be evaluated in future patients. Along with combining molecular agents with cytotoxic chemotherapies, focal re-irradiation is tolerable and effective (2) and could be evaluated with new targeted agents. Recently, another DIPG case series demonstrated the tolerability of histone deacetylase (HDAC) inhibition in conjunction with re-irradiation, supporting the report of novel approaches even when in small series (10). While we believe the best system for evaluating novel therapeutics is within a clinical trial, some disease subsets, such as NF1-mutant DIPG, are so rare that even with international collaboration their molecular targeting could not be studied within a clinical trial. Ultimately, whether evaluated in a clinical trial or by experienced clinical teams with careful consideration, molecularly targeted agents may be beneficial in combination with other novel therapies, conventional chemotherapy, or radiation for children with DIPG/DMG.

Nonstandard Abbreviations: BAT, Best available therapy; ctDNA, Circulating tumor DNA; DIPG, Diffuse intrinsic pontine glioma; DMG, Diffuse midline glioma, H3 K27M-mutant; Gy, Gray; HDAC, Histone deacetylase; LGG, Low-grade glioma; SCH, Seattle Children's Hospital; WHO, World Health Organization.

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Human Genes: ACVR1, activin A receptor type 1; H3-3A, H3.3 histone A; KDR, kinase insert domain receptor; KIT, KIT proto-oncogene, receptor tyrosine kinase; KRAS, KRAS proto-oncogene, GTPase; NF1, neurofibromin 1; PDGFRA, platelet derived growth factor receptor alpha; PIK3CA, phosphatidylinositol-4,5-bisphosphate 3-kinase catalytic subunit alpha; TP53, tumor protein p53; MYC, MYC proto-oncogene, bHLH transcription factor; FGFR1, fibroblast growth factor receptor 1.

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