Diffuse Midline Glioma with H3-K27M Mutation: A Rare Case with GFAP-positive Anucleate Whorled Patterns

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Case Report

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Abstract

Background: Diffuse midline glioma with H3-K27M mutation is a new tumor type in the 2016 WHO classification of tumors of the central nervous system, with predominantly astrocytic differentiation and is characterized by K27M mutation in either H3F3A/-HIST1H3B/C. It occurs at the midline of the CNS. It is mainly found in children, and adult cases are relatively rare. Although its histopathological features have been widely described, GFAP-positive anucleate whorled patterns have not been reported to the best of our knowledge. Hence, we have reported this unusual case.

Case presentation—A woman presented with complaints of walking instability. MRI showed a mass shadow of isometric T1 and slightly longer T2 with mild mixed signals in the third ventricle of the suprasellar region, about 3.4 x 2.5 x 3.4 cm in size. The enhanced scan showed that the lesion was irregular and annular, with a clear edge. The lesion had spread to the right lateral ventricle, which dilated the ventricular system, especially the right lateral ventricle. Furthermore, the lamellar long T2 signal was seen in bilateral paraventricular white matter, the midline structure had shifted to the left, and the corpus callosum was thinner. Radiological findings suggested ependymoma. Subtotal resection craniotomy was performed. After surgery, the patient was routinely treated with temozolomide for chemotherapy and synchronous radiotherapy (Dt=60 Gy/30f). It has been 11 months now, and the patient is living well.

Result: We described an extremely rare case of diffuse midline glioma, H3-K27M mutation with GFAP-positive anucleate whorled patterns.

Conclusions: This case report provides information on the microscopic morphological features of diffuse midline glioma with H3K27M mutation, which can help pathologists to make a definitive diagnosis of this tumor.

Introduction

Diffuse midline glioma with H3-K27M mutation is a new tumor type in the 2016 WHO classification of tumors of the CNS [1,2], with predominantly astrocytic differentiation and is characterized by K27M mutation in either H3F3A/-HIST1H3B/C. It occurs at the midline (spinal cord, thalamus, brainstem, cerebellum, hypothalamus, third ventricle, pineal gland, etc.) [3,4,5,6] of the CNS. It is mainly found in children, and adult cases are relatively rare [3,5].

Although its histopathological features have been widely described, GFAP-positive anucleate whorled patterns have not been reported to the best of our knowledge. Hence, we have reported this unusual case.

Clinical Summary

A 54-year-old Chinese female developed memory loss for half a month, which was characterized by intermittent short-term memory loss, but her long-term memory remained normal. After 15 days, the patient's memory loss obviously worsened, and both long-term memory and short-term memory were

poor. Concurrently, she also developed walking instability. Magnetic resonance imaging (MRI) showed a mass shadow of isometric T1 and slightly longer T2 with mild mixed signals in the third ventricle of the suprasellar region, about 3.4 x 2.5 x 3.4 cm in size. The enhanced scan showed that the lesion was irregular and annular, with a clear edge. The lesion had spread to the right lateral ventricle, which dilated the ventricular system, especially the right lateral ventricle. Furthermore, the lamellar long T2 signal was seen in bilateral paraventricular white matter, the midline structure had shifted to the left, and the corpus callosum was thinner (Fig. 1). Radiological findings suggested ependymoma. Subtotal resection craniotomy was performed. After surgery, the patient was routinely treated with temozolomide for chemotherapy and synchronous radiotherapy (Dt = 60 Gy/30f). It has been 11 months now, and the patient is living well.

Materials And Methods

The resected specimens were fixed with 10% neutral buffer formaldehyde and embedded with paraffin. Thereafter, the tissue was cut into 4 μ m sections, decarbonized in xylene, rehydrated with graded alcohols, and immunostained with the following antibodies: ATRX, GFAP, IDH, NeuN, Olig-2, S-100, Syn, Vimentin, Ki-67, and P53 (MaiXin, China). All samples were mildly counterstained with hematoxylin, dehydrated and mounted. The negative controls were incubated with PBS instead of the primary antibody, using the same procedure.

FISH was performed to check for deletions of chromosomes 1p and 19q. Dual color-probe hybridization was performed with Vysis 1p36/1q25 and 19q13/19p13 FISH probe kit (Abbott Molecular, Illinois, USA) according to the manufacturer's instructions. At least 100 non-overlapping nuclei were counted; samples were considered to be 1p- or 19q-deleted when > 30% of counted nuclei presented one target (red) signal and two reference (green) signals. Sanger sequencing was used to detect the IDH 1/2 mutation and TERT promoter mutations. K27M mutation and KIAA1549-BRAF fusion were performed by Sinomdgene company (Beijing, China) and Kanglu company (Hubei, China), respectively.

Pathological And Genetic Findings

Small grayish-red fragments of the resected lesion were sent for histological examination. The lesion was 1.2 × 0.7 × 0.3 cm in size. Histological examination with hematoxylin and eosin staining revealed that the tumor was primarily sheet-like, and the tumor cells showed moderate cellularity. Notably, there were many “anucleate areas” in the tumor, composed of long and thin fibrillary processes of the bipolar cells, which formed “whorls”. The neoplastic nuclei were ovoid and moderate in size. Moreover, the tumor showed brisk mitotic activity and vascular proliferation, with no necrosis (Fig. 2). Immunohistochemical staining revealed that the tumor cells were positive for GFAP, Olig-2, histone H3K27M mutant protein, ATRX, S-100 and Vimentin, but negative for IDH1 R132H, Syn and NeuN. The “anucleate areas” were positive for GFAP, but negative for Syn. The Ki-67 proliferation index was about 10% (Fig. 3). Molecular genetic analyses detected *H3F3A* K27M mutation (Fig. 4), while no mutations in *IDH1* or *IDH2*, *TERT* promoter mutations,

MGMT promoter methylation, *KIAA1549-BRAF* fusion or deletion of 1p/19q were found. Based on these findings, the patient was diagnosed with diffuse midline glioma with H3-K27M mutation in the third ventricle, corresponding to WHO grade 4.

Discussion

According to the 2016 WHO classification of the CNS and cIMPACT Working Committee 3 for diagnosing diffuse midline glioma with H3-K27M mutation, a tumor should be diffuse, midline glioma, with H3 K27M mutation [1,2]. The histological features of diffuse midline gliomas are diverse. The relatively rare morphological characteristics include epithelioid and rhabdoid cells, gliomas with giant cells, ependymal-like areas, primitive neuroectodermal tumor (PNET)-like foci, sarcomatous transformation, ganglionic differentiation and pleomorphic xanthoastrocytoma (PXA)-like areas, pilomyxoid features, and "rosette-like" nerve islands, neuropil-like islands [6,7]. Notably, Gao et al. reported a diffuse midline glioma with PNET-like and neuropil-like islands [7]. The neuropil-like islands were similar to the anucleate whorled structures in the current case, and both of them appeared as vortex-like islands. The difference was that the neuropil-like islands showed neuronal differentiation in the previous study and were positive for Syn, with scattered NeuN-positive neuronal cells [7]. In contrast, in our case, the nuclear-free zone was formed by the extension of slender cytoplasmic processes by GFAP-positive glial cells, but was negative for neuronal markers. Moreover, the tumor cells showed positive expression of H3K27M by immunohistochemistry, and gene detection further confirmed that they contained H3K27M gene mutation. Based on these findings, we diagnosed the tumor as diffuse midline glioma with histone H3-K27M mutation in the third ventricle. Given the varied histological features of diffuse midline gliomas, understanding the location and genetic alterations can help with the correct diagnosis.

Poor prognosis of most H3K27M mutant gliomas has been confirmed in many studies [3,4,5,8,9,10], and less than 10% of patients survive for more than two years [1], so this tumor is classified as WHO grade 4. Some researchers noted that H3K27M mutant gliomas may also show histological grade 2/3 [3,8,10], and the median overall survival rate of H3K27M mutation cases with histological grade 2/3 is significantly longer than those with histological grade 4 [8,10]. Therefore, histological classification plays a very important role in prognosis. In this case, the density of the cells under the microscope was medium, and vascular proliferation was notable, but there was no obvious necrosis, and histology indicated WHO grade 3 tumor. The patient was routinely treated with temozolomide for chemotherapy and synchronous radiotherapy (Dt = 60 Gy/30f) after surgery. To date, she has not had a recurrence for 11 months. The association between prognosis and histological grade of diffuse midline glioma needs further validation.

Herein, we described an extremely rare case of diffuse midline glioma, H3-K27M mutation with GFAP-positive anucleate whorled patterns. This case report provides information on the microscopic morphological features of diffuse midline glioma with H3K27M mutation, which can help pathologists to make a definitive diagnosis of this tumor.

Abbreviations

CNS
central nervous system
MRI
Magnetic resonance imaging
FISH
Fluorescent in situ hybridization
ATRX
alpha-thalassemia/mental retardation syndrome X
GFAP
glial fibrillary acidic protein
IDH
isocitrate dehydrogenase
NeuN
neuronal nuclear antigen
Olig-2
oligodendrocyte transcription factor 2
Syn
synaptophysin

Declarations

Acknowledgments

Not applicable.

Authors' contributions

Yu JH acquired clinical data; Qiu XS and Wang EH performed the pathological examination; Gu J performed the image examination; Wang YH wrote the manuscript; Yu JH, Fu L and Li QC revised the manuscript. All authors issued final approval for the version to be submitted.

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Availability of data and materials

All data generated or analyzed during this case are included within the article.

Ethics approval and consent to participate

The need for ethics approval and consent was waived, since a consent for publication was obtained from the patient's kids.

Consent for publication

Written informed consent for publication of the clinical details and/or clinical images was obtained from the kids of the patient. A copy of the consent form is available for review by the Editor of this journal.

Competing interests

The authors declare that they have no competing interests.

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Figures

Figure 1

The third ventricle showed isometric T1, slightly longer T2 with mild mixed signal mass shadow, cystic solid. The boundary was clear, with no definite edema signal in the adjacent brain tissue. The main body of the lesion was located in the third ventricle and spread to the right lateral ventricle. Bilateral ventricular dilatation was observed, wherein the right ventricle was more significant. A long stripe T2 signal can be seen at the edge of the lateral ventricle (interstitial brain edema). Enhanced scanning of the mass sac wall and solid components showed irregular ring enhancement, and local lace-like enhancement (A, axial T2; B, axial T2 FLAIR; C, axial T1; D, axial T1 enhancement; E, sagittal T1; F, enhanced sagittal T1).

Figure 2

The resected tumor was primarily sheet-like (scale bars: 600µm)(A), and the spindle-shaped tumor cells with thin, elongated processes formed "whorls" (scale bars: 50µm)(B). Microvascular proliferation (scale bars: 50µm) (C) and brisk mitotic activity (scale bars: 50µm) (D) were observed.

Figure 3

(A) (scale bars: 100µm)The tumor cells were negative for IDH1 R132H. (B) (scale bars: 70µm)The tumor cells and "neuropil-like islands" were positive for GFAP. (C) (scale bars: 90µm)The tumor cells were positive for Olig-2. (D) (scale bars: 70µm)The tumor cells and "neuropil-like islands" were negative for synaptophysin. (E) (scale bars: 90µm)The tumor cells were positive for H3-K27M mutant protein. (F) (scale bars: 80µm)The Ki-67 proliferation index was about 20%.

Sequence analysis showed mutation at position K27M in the histone coding gene H3F3.