

C17orf96

Chromosome 17 Open Reading Frame 96

Uncharacterized Protein C17orf96

The **PFA** molecular subgroup of **posterior fossa ependymomas** (PF-EPNs) shows poor outcomes. **H3K27me3** (me3) loss by **immunohistochemistry** (IHC) is a surrogate marker for **PFA** wherein its loss is attributed to **overexpression** of **Cxorf67/EZH2** inhibitory protein (**EZHIP**), **C17orf96**, and **ATRX** loss. Nambirajan et al. aimed to subgroup PF-EPNs using me3 IHC and study correlations of the molecular subgroups with other histone-related proteins, 1q gain, Tenascin C, and outcome. IHC for me3, acetyl-H3K27, H3K27M, ATRX, EZH2, EZHIP, C17orf96, Tenascin-C, and fluorescence in-situ hybridization for chromosome 1q25 locus were performed on an ambispective PF-EPN cohort (2003-2019). H3K27M-mutant gliomas were included for comparison. Among 69 patients, PFA (me3 loss) constituted 64%. EZHIP overexpression and 1q gain were exclusive to PFA seen in 72% and 19%, respectively. Tenascin C was more frequently positive in PFA ($p = 0.02$). H3K27M expression and ATRX loss were noted in one case of PFA-EPN each. All H3K27M-mutant gliomas ($n = 8$) and PFA-EPN ($n = 1$) were EZHIP negative. C17orf96 and acetyl-H3K27 expression did not correlate with me3 loss. H3K27me3 is a robust surrogate for PF-EPN molecular subgrouping. EZHIP overexpression was exclusive to PFA EPNs and was characteristically absent in **Diffuse midline glioma H3 K27M-mutants** and the rare PFA harboring H3K27M mutations representing mutually exclusive pathways leading to me3 loss ¹⁾.

¹⁾

Nambirajan A, Sharma A, Rajeshwari M, Boorgula MT, Doddamani R, Garg A, Suri V, Sarkar C, Sharma MC. EZH2 inhibitory protein (EZHIP/Cxorf67) expression correlates strongly with H3K27me3 loss in posterior fossa ependymomas and is mutually exclusive with **H3K27M** mutations. Brain Tumor Pathol. 2020 Nov 1. doi: 10.1007/s10014-020-00385-9. Epub ahead of print. Erratum in: Brain Tumor Pathol. 2021 Jan 9;; PMID: 33130928.

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