Review of: "Radiation-induced gliomas represent H3-/IDH-wild type pediatric gliomas with recurrent PDGFRA amplification and loss of CDKN2A/B"

Zhifeng Shi¹

¹ Fudan University

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This work is very valuable for RIGs study because set-up of such cohort with long term follow is difficult. In the paper, we can get to know the underlying genomic or genetic alterations of RIGs, and methylation profiling was also performed to precisely define the features of this kind of brain lesions. The results provide useful information for diagnosis of RIGs and therapeutic targets information.