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The current WHO definition of GBM relies on the presence of high-grade astrocytic neoplasm with the presence of either microvascular proliferation and/or tumor necrosis. High-throughput analyses have identified molecular subtypes and have led to progress in more accurate

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investigators in finding novel
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Glioblastoma is the most malignant brain tumor that still remains incurable. It is such a deadly disease that patients do not survive more than a few months after diagnosis. Our

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26;34(11):182. doi:
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Common and detrimental forms of solid brain tumor, with over 10,000 new cases reported every year in the United States. Despite aggressive multimodal treatment approaches, the overall survival period is reported to be less than 15 months after diagnosis. A widely used ...

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The differentially expressed genes (DEGs) in gliomas and glioblastoma were identified using FDR and ttests, and protein-protein interaction (PPI) networks for these DEGs were

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constructed using the protein interaction network analysis. The GeneTrail2 1.5 tool was used to identify potentially enriched biological processes among the DEGs using gene ontology (GO) terms and to identify the related pathways using the Kyoto Encyclopedia of Genes and Genomes,

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~~Molecular mechanisms underlying gliomas and glioblastoma ...~~

The continuously evolving field of understanding the molecular pathogenesis of GBM prompted to a

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more rational use of targeted molecular therapies. Inhibitors that target receptor tyrosine kinases (EGF, PDGF, VEGF, HGF, IGF receptors) and signal transduction inhibitors targeting mTOR, AKT/PI3K and farnesyltransferase are being investigated.

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The next molecular phenomenon in GBM pathogenesis to be discussed is the 1p/19q codeletion. This results from an unbalanced translocation between chromosomes 1p and 19q

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and leads to LOH. This molecular signature has been found to have tremendous significance and clinical utility in the evolving paradigm of molecular based prognostication and treatment of high grade glioma.

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Abstract. Glioblastoma (GBM) is one of the most lethal human cancers.

Genomic analyses are defining the

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molecular architecture of GBM, uncovering relevant subsets of patients whose disease may require different treatments. Many pharmacological targets have been revealed, promising to transform patient care through targeted therapies.

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Glioblastoma, the deadliest of all primary brain tumors, is heterogenic and has very complex pathology and multiple mechanisms of cell proliferation and growth. This book

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describes all the latest concepts about the molecular mechanisms of pathogenesis of glioblastoma and new therapeutic opportunities. The contributors to this book include basic scientists and clinicians, who are experts in the field, and dedicated investigators in finding novel

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