



Emerging Diagnostic Techniques for Meningioma Variants: Integrating Imaging and Molecular Insights

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DESCRIPTION

Meningiomas, the most common primary brain tumors originating from the meninges, exhibit a broad range of biological behaviors and histological patterns. While many meningiomas follow a benign course, some display aggressive tendencies, including recurrence and invasion of surrounding brain tissue. This variability has prompted a growing interest in refining diagnostic approaches to better characterize meningioma subtypes. Advances in imaging technologies and molecular diagnostics have begun to enhance our ability to distinguish tumor subtypes, predict clinical outcomes, and guide treatment decisions more effectively.

Traditional imaging methods, such as Magnetic Resonance Imaging (MRI) and Computed Tomography (CT), remain essential for initial tumor detection and evaluation of anatomical involvement. MRI, in particular, offers detailed soft tissue contrast that helps determine tumor size, location, and effects on adjacent brain structures. However, conventional MRI sequences sometimes lack the specificity needed to differentiate between meningioma subtypes or to accurately predict tumor grade.

Recent progress in advanced MRI techniques has improved diagnostic accuracy. Diffusion Weighted Imaging (DWI) measures the movement of water molecules within tissue, providing insights into tumor cellularity. High-grade meningiomas tend to exhibit restricted diffusion due to increased cell density, while lower-grade tumors show less restriction. Similarly, perfusion MRI evaluates blood flow within the tumor, which correlates with angiogenesis and tumor aggressiveness. Elevated perfusion parameters have been associated with higher-grade meningiomas and may predict recurrence risk.

Another valuable tool is Magnetic Resonance Spectroscopy (MRS), which identifies metabolic profiles of brain tumors. Meningiomas display distinctive spectral patterns compared to other brain lesions, with increased choline levels indicating

cellular proliferation and reduced N-acetylaspartate suggesting neuronal loss nearby. Variations in metabolites may also help differentiate subtypes, although this remains an area of active investigation.

Positron Emission Tomography (PET) imaging has introduced further possibilities for meningioma evaluation. Radiotracers such as ^{68}Ga -DOTATATE bind to somatostatin receptors, which are frequently overexpressed in meningiomas. This receptor-targeted PET scan enhances tumor visualization, especially for lesions difficult to distinguish on MRI alone, such as those involving the skull base or recurrent tumors. PET imaging also aids in planning surgical approaches and radiation therapy by identifying metabolically active tumor regions.

Beyond imaging, molecular diagnostics have revolutionized the classification of meningiomas. Genetic and epigenetic alterations correlate with tumor behaviour and subtype identity, offering prognostic and therapeutic insights. For instance, mutations in the *NF2* gene are common in meningiomas, especially those located in the cerebral hemispheres. Loss of *NF2* function disrupts cell growth regulation, contributing to tumor development.

In addition to *NF2*, recent studies have identified mutations in genes such as *TRAF7*, *KLF4*, *SMO*, and *POLR2A*, which tend to cluster with specific meningioma subtypes and anatomical locations. For example, *TRAF7* and *KLF4* mutations often co-occur in tumors arising from the skull base and are generally associated with benign histology. In contrast, alterations in *TERT* promoter regions or homozygous deletion of *CDKN2A/B* are linked with more aggressive tumors and poorer prognosis.

Emerging diagnostic platforms combining radiomics and artificial intelligence show potential to further refine meningioma subtype identification. Radiomics involves extracting quantitative features from imaging data, such as texture, shape, and intensity, which can then be correlated with molecular and clinical characteristics. Machine learning algorithms can analyze these complex datasets to predict tumor

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grade, subtype, and likelihood of recurrence with increasing accuracy.

Despite these advances, several challenges remain. Standardization of molecular testing and imaging protocols is necessary to ensure consistent application across institutions. Additionally, some molecular alterations are rare or occur in small patient subsets, making it difficult to draw firm conclusions about their clinical significance. Longitudinal studies and larger cohorts will be essential to validate emerging diagnostic criteria and establish their role in routine care.

Collaboration between neurosurgeons, radiologists, pathologists, and molecular biologists is vital to translate these evolving diagnostic tools into improved patient outcomes. As

understanding of meningioma biology expands, diagnostic approaches will continue to evolve, allowing more precise characterization of tumor behaviour and better guidance for treatment planning.

In summary, combining advanced imaging techniques with molecular analysis enhances the capacity to distinguish meningioma subtypes. This integrated approach provides valuable information beyond what traditional methods offer, supporting more informed decisions throughout the patient's course. Continued innovation and clinical evaluation promise to refine diagnostic capabilities, ultimately benefiting individuals affected by this diverse group of brain tumors.