

CHCHD2P9

Full name: Coiled-Coil-Helix-Coiled-Coil-Helix Domain Containing 2 Pseudogene 9

Gene type: [Pseudogene](#)

Related gene: [CHCHD2](#) (a protein-coding gene involved in mitochondrial function, apoptosis regulation, and oxidative phosphorylation)

Function: As a pseudogene, CHCHD2P9 is a non-functional genomic sequence that resembles the CHCHD2 gene but typically lacks the ability to produce a functional protein.

Location: Human genome, specific locus may vary by reference genome assembly.

Biological Relevance

While pseudogenes were traditionally considered “junk DNA”, growing evidence suggests some may have regulatory roles through:

Transcriptional interference

miRNA sponging

Epigenetic regulation

In a [translational research](#) integrating [single-cell RNA sequencing](#), [flow cytometry](#), and in vitro [functional assays](#). Ding et al. from the Anhui Medical University, Hefei; Shanghai Ninth People's Hospital, Shanghai; University of Science and Technology of China, Hefei. published in [Frontiers in Immunology](#) to elucidate the role of the pseudogene CHCHD2P9 in glioblastoma progression and tumor heterogeneity by leveraging single-cell RNA sequencing and functional assays. CHCHD2P9 is overexpressed in glioma and correlates with worse prognosis. It may influence glioma proliferation and migration and serve as a novel prognostic biomarker or therapeutic target ¹⁾.

Critical Review: This study attempts a multidimensional exploration of glioblastoma heterogeneity by integrating advanced single-cell [transcriptomics](#) with basic cellular assays. The identification of [CHCHD2P9](#) as a putative prognostic marker is intriguing, but the study lacks depth in mechanistic validation. While the correlation between CHCHD2P9 expression and [clinical outcome](#) is statistically supported, causality remains speculative. The study's reliance on a [pseudogene](#) raises biological plausibility concerns, especially without sufficient evidence for its [protein-coding function](#) or [epigenetic regulation](#). The model's [translational](#) utility is also not directly tested in patient-derived [xenografts](#) or [organoids](#). The work is [hypothesis-generating](#) rather than definitive, with significant promise for future mechanistic follow-up.

Final Verdict: Interesting early-phase study with a novel target but insufficient mechanistic or in vivo [validation](#).

Takeaway for the Practicing Neurosurgeon: [CHCHD2P9](#) may emerge as a biomarker of [glioblastoma prognosis](#), but clinical application is premature. No impact on current neurosurgical management.

Bottom Line: A promising [transcriptomic](#) marker in glioblastoma subpopulations, but further functional studies are essential.

Rating: 5/10

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¹⁾

Ding Y, Xiao L, Zhou X, Zhao J, Ke J, Cai H, Zhao M, Wang C, Yu T, Zhao Z, Wang Y, Ke J. Molecular insights into [glioblastoma progression](#): role of [CHCHD2P9](#) in [tumor heterogeneity](#) and [prognosis](#). Front Immunol. 2025 Jun 24;16:1581850. doi: 10.3389/fimmu.2025.1581850. PMID: 40630954; PMCID: PMC12234496.

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