

**26 AUG** 

11.30 A.M. ROOM A208 POVO 1

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A comparative analysis of G-Quadruplexes in viral genomes

> G-quadruplexes (G4s) are non-canonical nucleic acid structures that play a pivotal role in regulating essential molecular processes. Their potential as targets for antiviral therapies has garnered increasing attention. We performed a comprehensive analysis of putative Gquadruplex-forming sequences (PQS) across all viral genomes currently available in the NCBI database, including subviral agents. Using the **G4Hunter algorithm**, we screened over 11,000 viral genomes totaling approximately **350 Mbp**. Our findings reveal that PQS distribution varies significantly among viral evolutionary groups. G-quadruplexes may play a regulatory role in viral replication and lifecycle control. The distinct localization of PQS highlights their potential as molecular targets for therapeutic intervention, offering new avenues for antiviral strategy development.





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