

Special Issue

Herpesvirus Transcriptional Control

Message from the Guest Editor

For over four decades, researchers have studied the transcription of herpesvirus genes to understand their roles in lytic and latent infections and as a paradigm for eukaryotic transcription. The advent of new technologies, approaches and model systems, such as RNA sequencing (RNA-seq), chromatin immunoprecipitation sequencing (ChIP-seq), proteomics and organoid cultures, is providing greater insights into the regulation of herpesvirus transcription during lytic infection and reactivation, as well as the restrictions in viral transcription observed during the establishment and maintenance of latency. This Special Issue will highlight current research examining herpesvirus–host interactions that modulate viral and/or host transcription, which influence the life cycles and pathogenesis of herpesvirus family members.

Guest Editor

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