Introduction to JAGS and rJAGS

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Section header

Methods for analyzing genomic count data have typically been oriented around RNA sequencing (RNA-Seq). Here, we explore how RNA-Seq methods can be extended to other types of count data, specifically phage immunoprecipitation sequencing (PhIP-Seq).

- Setup & Notation
- Background
    - RNA-Seq
    - Immunology and PhIP-Seq
    - Methods for evaluating differential expression
- HIV example
- Additional Comments