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Use R viruses and hosts for their survival. The structural and functional information on the interactions between viruses and hosts should provide a molecular and biological basis to understand infection,

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Use R replication, cell/host-tropism, immune escape, pathogenesis, and direction of evolution of viruses. The information is also essential to develop methods to control transmission and replication of pathogenic

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Use R. However, the integrated information on the structure, function, and evolution of viruses and hosts has remained poorly accumulated, partly due to the limitation of analytical methods. Recent progress in

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Use R genome science and computational approach may open up a new avenue of research of the interactions between viruses and hosts by integrating information on the structures, functions, and evolution. In this

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Use R Research Topic, we welcome papers concerning the computer-assisted structural and functional studies based on genomic information, with theoretical or in combination with experimental approaches, for

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Understanding molecules,
infection, replication,
cell/host-tropism, immune
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nature.

Imprinted genes, many of

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Use R which generally control growth and development, frequently lose their imprints during cancer progression, a loss that then plays a substantial role in uncontrolled tumor growth. Imprint instability

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also R appears to be a major limitation to the success of mammalian cloning experiments. In Genomic Imprinting: Methods and Protocols, Andrew Ward and a team of experienced researchers have brought

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Use R together a collection of optimized classic and vanguard techniques for the identification and analysis of imprinted genes. The majority of protocols describe molecular techniques that allow

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Examination of gene structure or expression in an allele-specific manner. Protocols are included for identifying and cloning imprinted genes, for analyzing imprinted gene expression, for the study of

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DNA methylation and methylation-sensitive DNA-binding proteins, and for examining chromatin structure. There are also methods for the manipulation of mouse embryos to produce monoparental embryos and

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embryonic stem cells, and
for the generation of
transgenic mice with BAC,
PAC, and YAC constructs.
Each technique is described
in step-by-step detail to
ensure successful results.
Incorporating a wealth of

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Use R knowledge from leading
exponents in the field,
Genomic Imprinting: Methods
and Protocols brings
together all the essential
molecular, genetic, and
embryological methods
commonly used in today's

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laboratories for the
identification and analysis
of imprinted genes.

Genomic imprinting, a parent-
of-origin-specific

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Epigenetic modification of the genome, is observed in placental mammals and some plants (angiosperms). Since its discovery in the early 1980s, this intriguing phenomenon has been intensively investigated and

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Use R has spawned an exciting new research area. Imprinting is relevant to a variety of biological events such as embryogenesis, growth and behavior, and its disruption can lead to congenital malformation syndromes and

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Use R. Furthermore,
imprinting has been serving
as an excellent model system
to study epigenetic
mechanisms of gene
regulation and helping us to
understand how gene clusters
and genome domains are

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Use R regulated. This special issue provides an overview of the current knowledge of genomic imprinting.

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information provided in each
chapter

The application of homology
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