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~~4) Next Generation  
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| Tim Urban  
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## Sequencing

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*Methods for Next  
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*Sequencing Data ...*

Introduces readers to  
core algorithmic  
techniques for next-  
generation

sequencing (NGS)

data analysis and



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discusses a wide range of computational techniques and applications. This book provides an in-depth survey of some of the recent developments in NGS and discusses mathematical and computational challenges in various application areas of NGS technologies.

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Next-generation  
sequencing (NGS)  
provides accurate and  
parallel sequencing

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Methods for  
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thereby allowing  
researchers to  
comprehend  
epigenomic profiling.

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In this review, we  
summarize different  
computational  
methods such as  
machine learning and  
other bioinformatics  
tools, publicly  
available databases  
and resources to  
identify key

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Next generation sequencing (NGS) has made great strides in sequencing technology as it enables sequencing of genes in a high

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throughput manner with low cost. Various NGS platforms such as Illumina, Roche, ABI/SOLiD are used for wet-lab analysis of NGS data and computational tools such as BWA, Bowtie, Galaxy, SanGeniX are used for dry-lab analysis of NGS data.

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computational  
methods and software  
tools are required.  
Here, we review  
computational tools

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cancer immunity,  
discuss advantages  
and limitations of the  
various methods and  
provide guidelines to  
assist in method  
selection.

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In summary,  
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Methods for  
prediction methods  
are essential for the  
implementation of  
Next Generation  
Sequencing  
NGS into clinical  
decision-making.

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While much progress  
has been made and a  
plethora of

conceptually diverse  
tools is already  
available, there is a  
need to develop  
specialized methods  
that are optimized for



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Methods For  
the prediction of  
variant functionality  
rather than  
Next Generation  
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pathogenicity and are  
Data Mining  
calibrated specifically  
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Second, our lack of a  
systematic  
comparison of error-

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Methods For

correction methods  
impedes the optimal  
integration of these  
tools into

standardized next-  
generation

sequencing data  
analysis pipelines.

Fig. 1 Study design  
for benchmarking  
computational error-  
correction methods.

*Benchmarking of*

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methods | Ram U.

Verma (2013)

invented several

classes of the next

generation

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methods, which are

also referred to as

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