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

Computational analysis of next generation sequencing data ...

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Next-generation computational tools for interrogating ...

In summary, computational prediction methods are essential for the implementation of NGS into clinical decision-making. 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 the prediction of variant functionality rather than pathogenicity and are calibrated specifically on pharmacogenetic data.

Computational Methods for the Pharmacogenetic ...

Second, our lack of a systematic comparison of error-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 computational error-correction methods for ...

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