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Next Generation Sequencing (NGS) is the latest high throughput
 technology to revolutionize genomic research. NGS generates
 massive genomic datasets that play a key role in the big data
 phenomenon that surrounds us today. To extract signals from
 high-dimensional NGS data and make valid statistical inferences

and predictions, novel data analytic and statistical techniques are needed.

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Statistical Methods for Next Generation Sequencing

Next generation sequencing (NGS) is a revolutionary technology for biomedical research. One highly cost-efficient application of NGS is to detect disease association based on pooled DNA samples. However, several key issues need to be addressed for pooled NGS.

Design and Statistical Analysis of Pooled Next Generation

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The boom of next generation sequencing (NGS) technology and its applications to a wide range of biomedical fields has brought about many computational and statistical challenges. The NHGRI funded small conference "Statistical Analysis for NGS" was held September 26th-27th 2011 in Birmingham, AL, USA, to discuss these statistical challenges and strategies to tackle them.

Statistics for Next Generation Sequencing - Meeting Report

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Statistical Analysis of Next Generation Sequencing Data

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This proposal is to develop advanced statistical methods for analyzing large next generation sequencing data in genetic cancer epidemiological studies. The genomic era provides an

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Project 3: Statistical Methods for Analysis of Next ...

Abstract We develop a statistical tool SNVer for calling common and rare variants in analysis of pooled or individual next-generation sequencing (NGS) data. We formulate variant calling as a hypothesis testing problem and employ a binomial-binomial model to test the significance of observed allele frequency against sequencing error.

SNVer: a statistical tool for variant calling in analysis ...

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