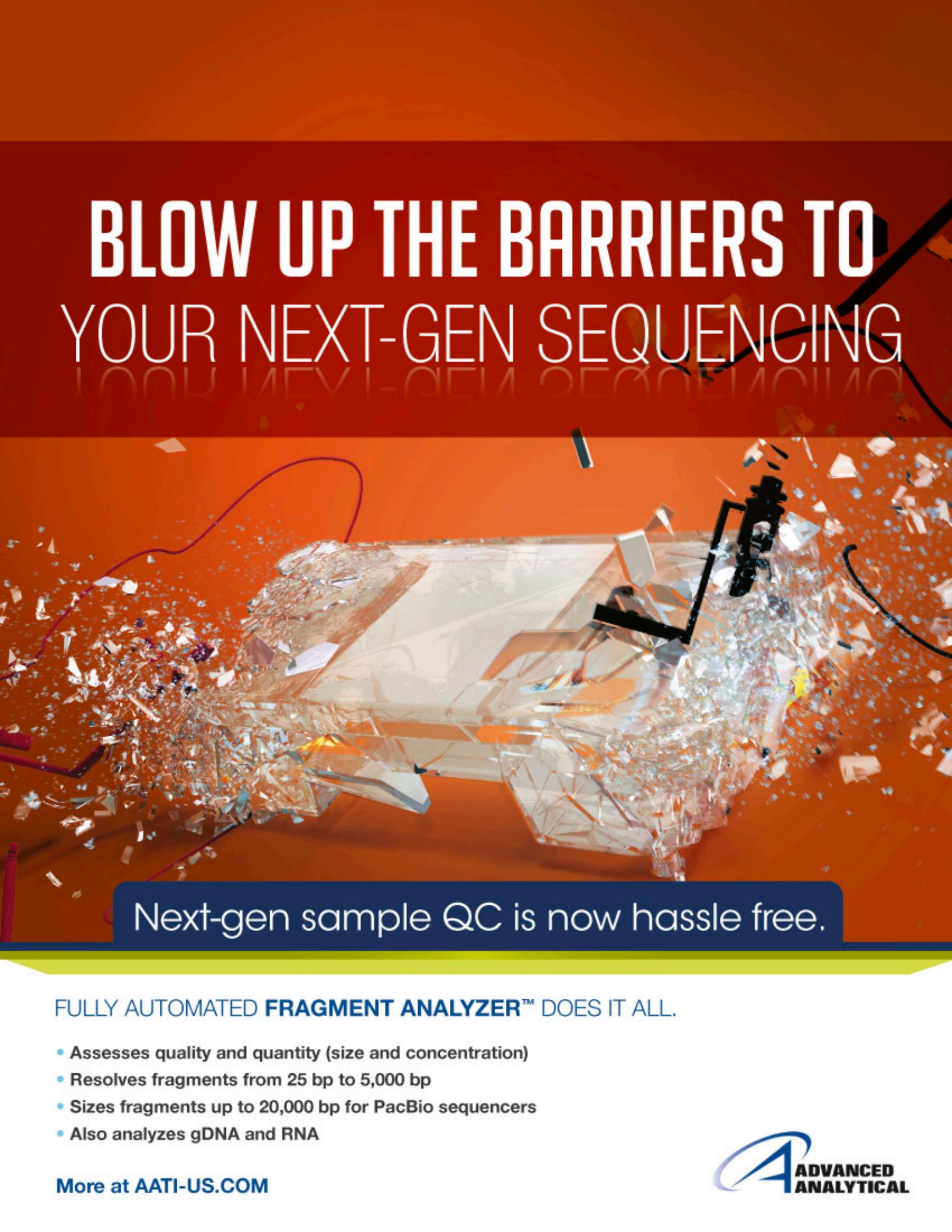


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# Next-Generation Genome Engineering Technologies & Services

One-Step Stable Transfection & Transgenesis



Targeted Knockouts



Gene Editing Base-Pair Changes



Knock-ins & Humanization

Targeted Selection Markers

## With So Many Opportunities What Are You Waiting For?

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**NextGEN™ CRISPR**

**XTN™ TALEN**

**Stable Transfection Kits &  
piggyBac™ Transposon**

**Cell Line & Rodent Model Engineering Services**

## Your Next Discovery Awaits



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[info@transposagenbio.com](mailto:info@transposagenbio.com)

[www.transposagenbio.com](http://www.transposagenbio.com)



# Break free from qPCR

Find gene targets faster



Validating gene targets with qPCR is often slow and costly due to lengthy primer design, experimental assay optimization, and sample processing time.

## QuantiGene® Plex Assays validate gene targets faster:

- No RNA purification or reverse transcription is required
- Multiplex panels of up to 80 genes are functionally validated
- One QuantiGene Plex® plate provides as much information as 20 qPCR plates

Download "The 7 best reasons to break free from qPCR" at [www.ebioscience.com/breakfree-genes](http://www.ebioscience.com/breakfree-genes)

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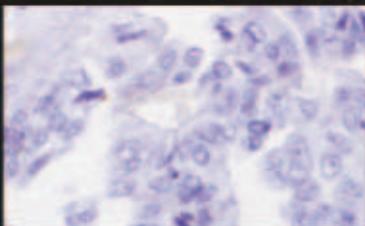
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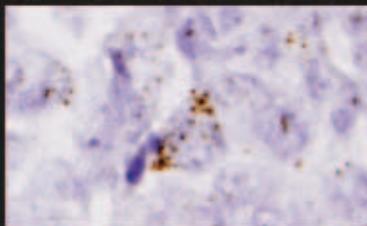


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RNAscope ISH detects Napsin A RNA molecules in the same specimen

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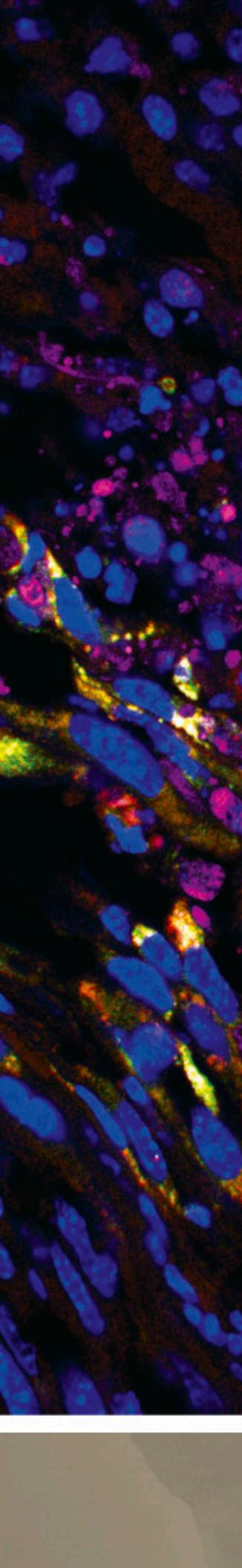
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# SCIENTIFIC CONFERENCES 2014-2015

Presenting the most significant research on cancer etiology, prevention, diagnosis, and treatment



## 7th AACR Conference on The Science of Cancer Health Disparities in Racial/Ethnic Minorities and the Medically Underserved

*Co-Chairpersons: Ethan Dmitrovsky, Rick A. Kittles, Electra D. Paskett, and Victoria L. Seewaldt*  
November 9-12, 2014 • San Antonio, TX

## The 10th Anniversary Personalized Medicine Conference

*Co-Chairpersons: Raju Kucherlapati and Scott Weiss*  
November 12-13, 2014 • Boston, MA

## EORTC-NCI-AACR International Symposium on Molecular Targets and Cancer Therapeutics

*Co-Chairpersons: Jean-Charles Soria, Lee J. Helman, and Jeffrey A. Engelman*  
November 18-21, 2014 • Barcelona, Spain

## Tumor Immunology: A New Chapter

*Co-Chairpersons: Robert H. Vonderheide, Nina Bhardwaj, Stanley Riddell, and Cynthia L. Sears*  
December 1-4, 2014 • Orlando, FL

## San Antonio Breast Cancer Symposium

*Co-Directors: Carlos L. Arteaga, Ismail Jatoi, and C. Kent Osborne*  
December 9-13, 2014 • San Antonio, TX

## Myc: From Biology to Therapy

*Co-Chairpersons: James E. Bradner, Martin Eilers, Dean W. Felsher, and Carla Grandori*  
January 7-10, 2015 • La Jolla, CA

## Translation of the Cancer Genome

*Co-Chairpersons: William C. Hahn, Lynda Chin, and William R. Sellers*  
February 7-9, 2015  
The Fairmont • San Francisco, CA

## Computational and Systems Biology of Cancer

*Co-Chairpersons: Andrea Califano, Brenda Andrews, and Peter Jackson*  
February 8-11, 2015  
The Fairmont • San Francisco, CA

## AACR-Society of Nuclear Medicine and Molecular Imaging Joint Conference: Molecular Imaging in Cancer Biology and Therapy

*Co-Chairpersons: Carolyn J. Anderson, Christopher H. Contag, and David Piwnica-Worms*  
February 11-14, 2015 • San Diego, CA

## Tumor Angiogenesis and Vascular Normalization: Bench to Bedside to Biomarkers

*Co-Chairpersons: Rakesh K. Jain, Peter Carmeliet, Helen Chen, Harold F. Dvorak, and Napoleone Ferrara*  
March 5-8, 2015 • Orlando, FL

## AACR Annual Meeting 2015

*Program Committee Chairperson: Lewis C. Cantley*  
April 18-22, 2015 • Philadelphia, PA

## Advances in Brain Cancer Research

*Co-Chairpersons: Eric C. Holland, Franziska Michor, Martine F. Roussel, and Michael D. Taylor*  
May 27-30, 2015 • Washington, DC

## Metabolism and Cancer

*Co-Chairpersons: Ralph J. DeBerardinis, David M. Sabatini, and Almut Schultze*  
June 7-10, 2015 • Bellevue, WA

## Cancer Biostatistics Workshop

*Director: Steven Piantadosi*  
June 7-13, 2015 • Lake Tahoe, CA

## AACR Precision Medicine Series:

**Integrating Real-Time Genomics and Cancer Therapy**  
*Co-Chairpersons: Charles L. Sawyers, Elaine R. Mardis, and Arul M. Chinnaiyan*  
June 13-16, 2015 • Salt Lake City, UT

## EACR-AACR-SIC Special Conference Special Conference on Anticancer Drug Action and Drug Resistance: From Cancer Biology to the Clinic

*Organizing Committee: Moshe Oren, Riccardo Dolcetti, Richard M. Marais, Daniel S. Peeker, Pasi Jänne, Alice T. Shaw, Paola Chiarugi, and Silvia Giordano*  
June 20-23, 2015 • Florence, Italy

## Pediatric Oncology

*Co-Chairpersons: Scott Armstrong, Charles G. Mullighan, Kevin M. Shannon, and Kimberly Stegmaier*  
November 9-12, 2015 • Fort Lauderdale, FL

## Developmental Biology and Cancer

*Co-Chairpersons: Hans Clevers, Stuart Orkin, Suzanne Baker*  
November 30-December 3, 2015 • Boston, MA

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The Maverix Analytic Platform was designed by biologists for life science researchers, allowing you to directly analyze, visualize, explore and manage your own NGS data in a easy-to-use system without the need for bioinformatics expertise or large-scale computing infrastructure.



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# SMART

cDNA SYNTHESIS FOR NEXT GEN SEQUENCING

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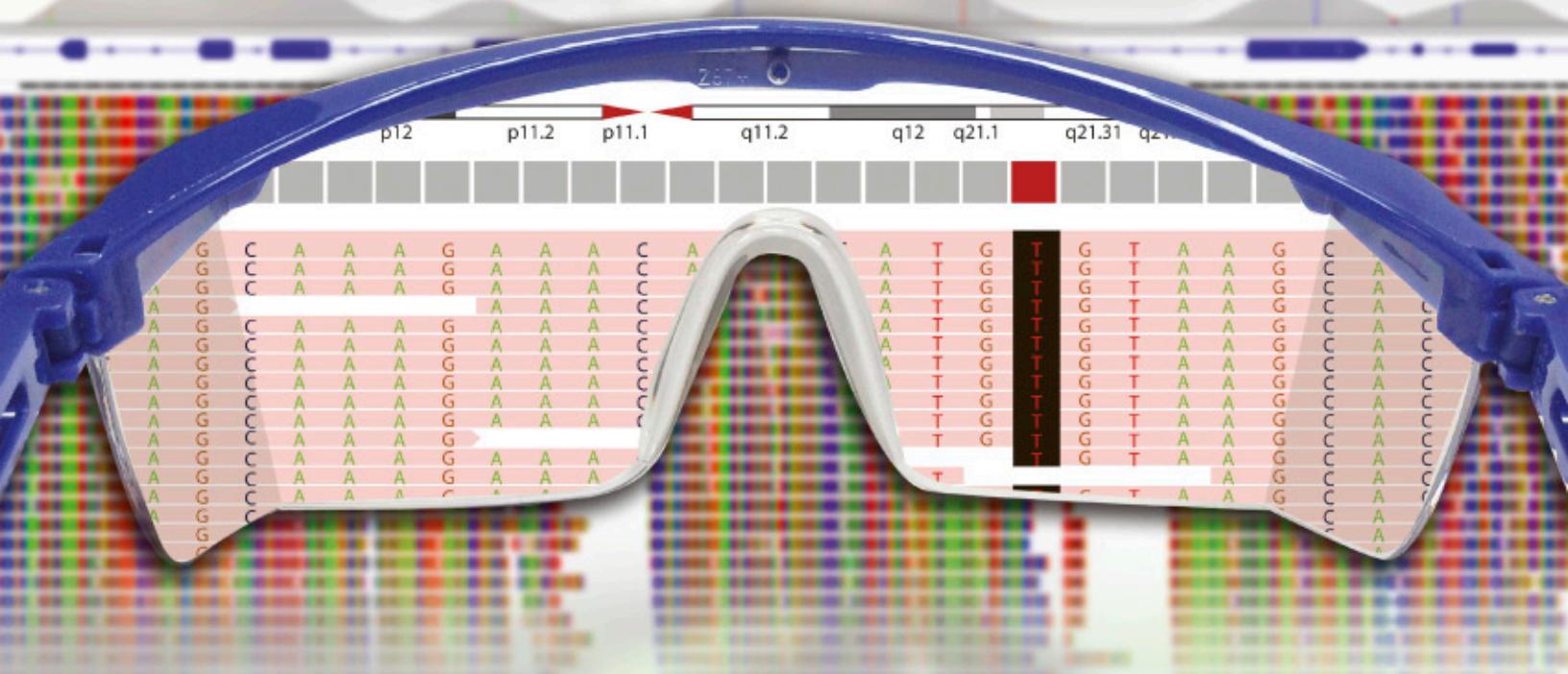
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# FOCUS ON GENOMICS



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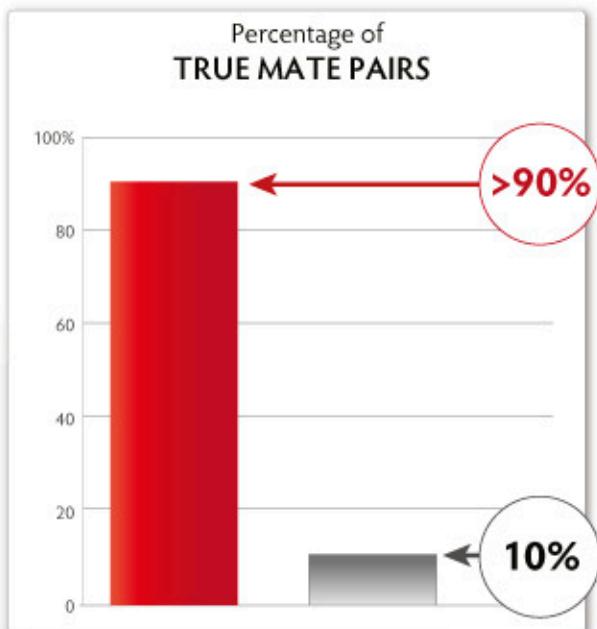
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- Cancer Panel Sequencing
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- Epigenomics

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- Variant (SNP/Indel) calling
- CNV & Breakpoints
- Expression Profiles
- DEGs/miRNA
- Enrichment Profiles
- Gene Annotation

**Capillary Sequencing**

**Microarray**

**Oligonucleotide Synthesis**

**Genetically Engineered Mouse**

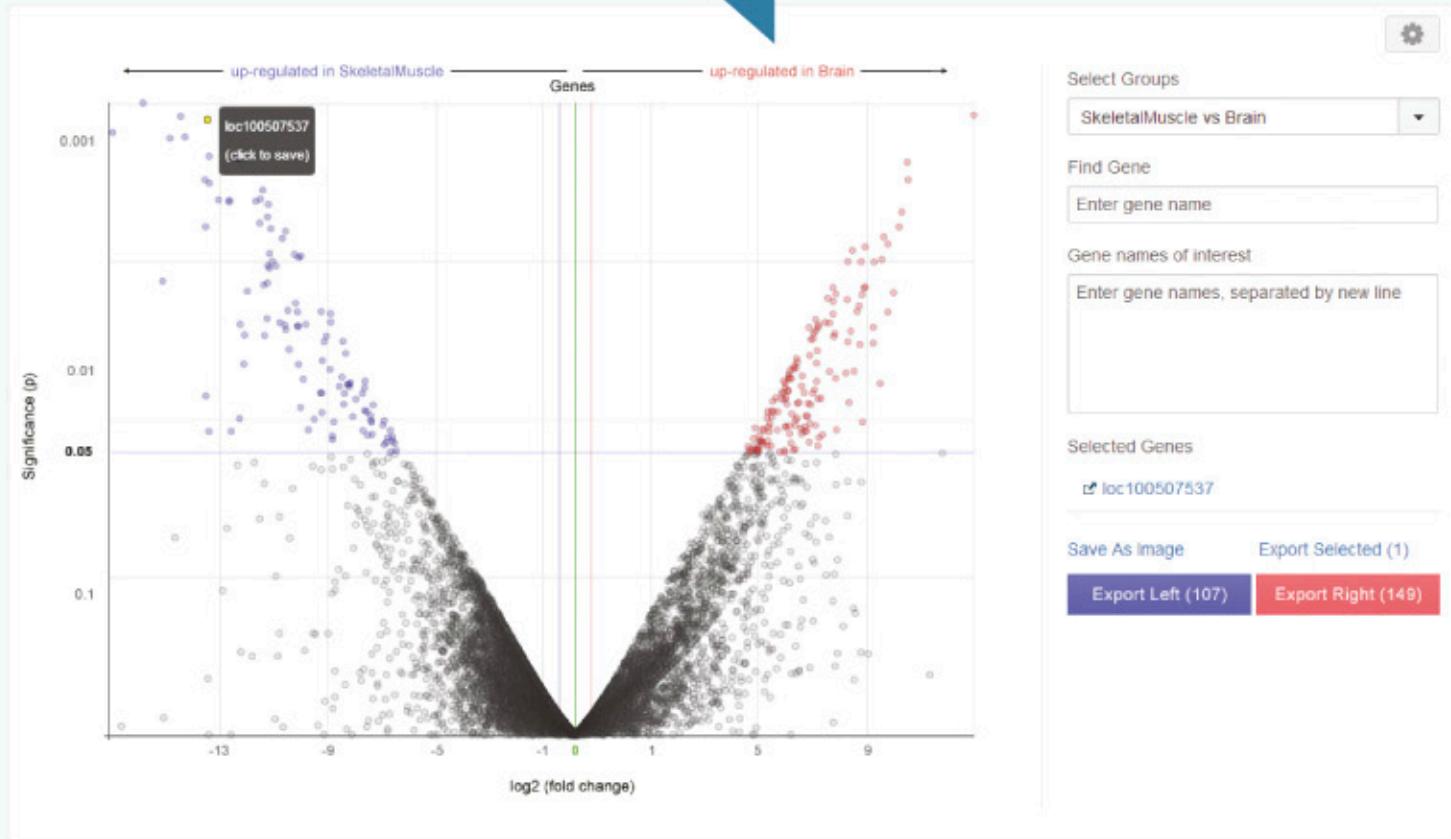


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## Web-based analysis

- Real-time integration with public databases; most up-to-date annotations
- Share analysis with collaborators, without duplicating files

## Advanced user interface

- Multiple views of the same data enable efficient transcript filtering and hypothesis generation
- Export views for presentations
- Search for genes using multiple identifiers

## Integrated analysis environment

- Preserves link from the final figure to the raw data, including all computational methods used



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