

# MiSeq<sup>™</sup>

## Better inside. Outside. Upside.

**Inside** you'll find a fully integrated sequencing solution delivering >85% Q30 bases and run times as fast as 8 hours from sample to data.

**Outside** you'll find "push button" sequencing, with a streamlined user experience, individually tracked load-and-go reagents, intuitive software interface, and minimal hands-on time.

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# ABRF 2012

LEARNING FROM BIOMOLECULES: THE TECHNOLOGY BEHIND THE STORY

MARCH 17 - 20, 2012 • DISNEY'S CONTEMPORARY RESORT • ORLANDO, FLORIDA

✓ **EARLY-BIRD REGISTRATION**  
will save you \$50 - Register by February 4th and save!

✓ **SATELLITE WORKSHOPS**  
focusing on lab management, pathways analysis, RNA & methyl sequencing and manual interpretation of ETD mass spectra of peptides.

✓ **SCIENTIFIC SESSIONS**  
and technical workshops will focus on genomics, proteomics, imaging and other technologies.

✓ **PLENARY SPEAKERS**

- Dr. George Church, Harvard University
- Dr. Kris Gevaert, Flanders Institute for Biotechnology
- Dr. Trisha Davis, University of Washington
- Dr. Ram Sasisekharan, Massachusetts Institute of Technology

✓ **ABSTRACTS**  
are currently being accepted - submit by January 17<sup>th</sup> for consideration.

✓ **EXHIBIT and SPONSORSHIP**  
opportunities are available - drive business, capture leads and make your mark at this year's meeting.

✓ **HOUSING**  
reservations made by February 10<sup>th</sup> receive the discounted room rate.  
Visit [www.mydisneymeetings.com/abrf2012](http://www.mydisneymeetings.com/abrf2012) to reserve your room at Disney's Contemporary Resort today!

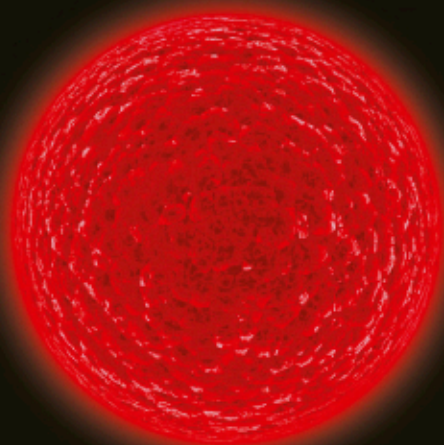
✓ **NETWORK**  
with fellow attendees at a roundtable discussion, reception, vendor seminar, meet the speaker or the not-to-be-missed closing social being held at **Universal Studios!**

**The Association of Biomolecular Resource Facilities** is an international society dedicated to advancing core and research biotechnology laboratories through research, communication, and education.

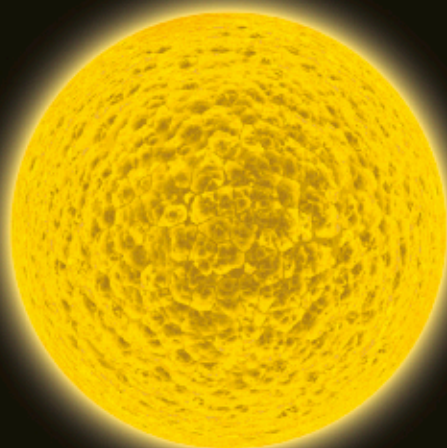
Visit us on the web for information, schedules, registration and more!  
**ABRF2012.ABRF.org**

For more information, please contact:

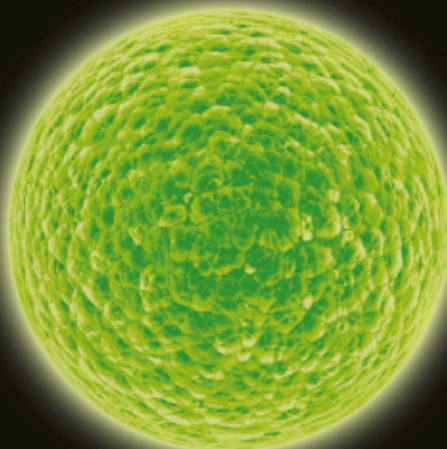
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## Standard Method

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Make cDNA



Ligate adaptors



Gel-purify



PCR



Clean up

>10 hours

## ScriptSeq v2

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and make cDNA



PCR



Clean up

4 hours