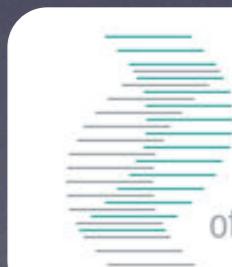


# Efficient tool deployment to the Galaxy Cloud: An RNA-seq workflow case study

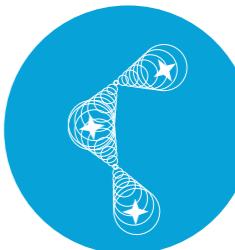
Sebastian J. Schultheiss <[sebi@tue.mpg.de](mailto:sebi@tue.mpg.de)>  
Machine Learning in Biology, Rätsch Lab, FML of the Max Planck Society  
Tübingen, Germany



Friedrich Miescher Laboratory  
of the Max Planck Society



**oqtans**  
online  
quantitative  
transcript  
analysis



# Web Services Availability

Schultheiss 2011, PLoS CB acc.  
Schultheiss et al. 2011, PLoS ONE i.r.

- ▶ 927 web services (NAR Web Server Issues)
- ▶ Collection of 45 data on every service
- ▶ Survey among authors
- ▶ Problems:
  - ▶ URL change
  - ▶ missing example data
  - ▶ program on server not functional
  - ▶ undocumented changes from updates

# Web Service Availability

Schultheiss 2011, PLoS CB acc.  
Schultheiss et al. 2011, PLoS ONE i.r.

- ▶ 927 web services (server issues)
- ▶ Collection of 100 services
- ▶ Survey analysis
- ▶ Problems with LIDI changes

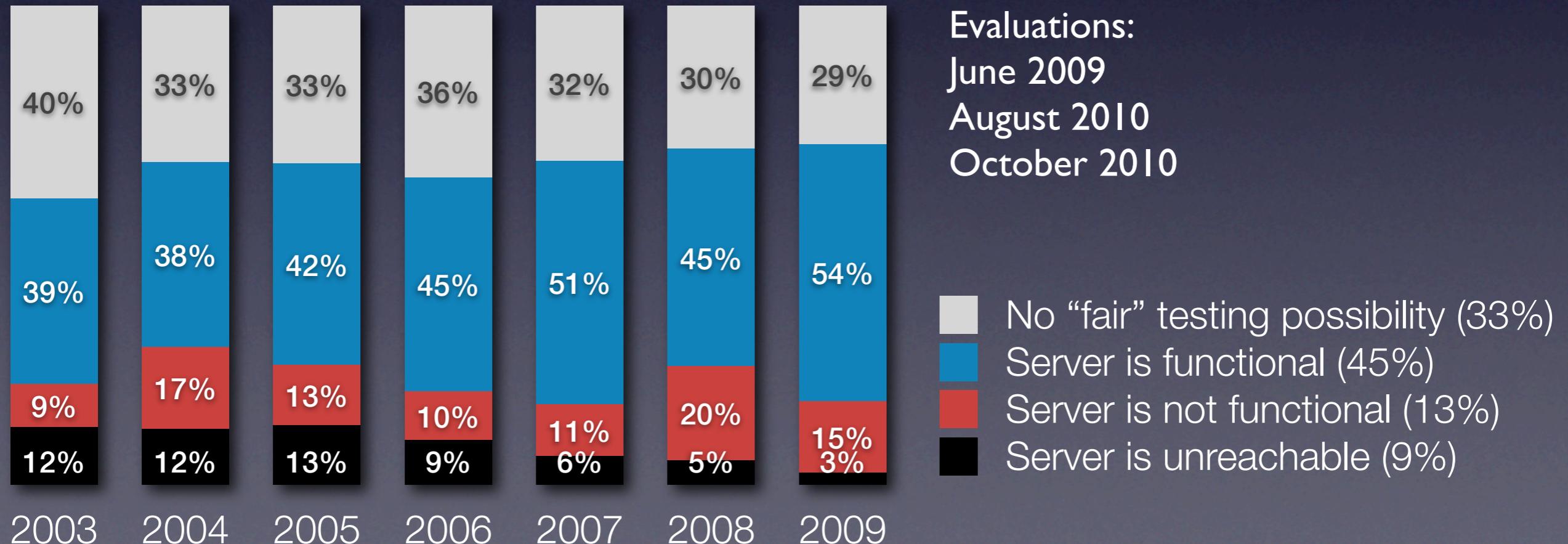


**Caution: Published results  
may not be reproducible**

# Web Services Availability

- ▶ Redirected from URL stated in abstract: 13%
- ▶ Unreachable, only via search engine: 7%
- ▶ Unreachable and not found: 9%
- ▶ Correct URL in abstract: 72%

Schultheiss et al. 2011, PLoS ONE i.r.



# Galaxy Approach

J. Goecks et al. 2010

D. Blankenberg et al. 2010

E. Afgan et al. 2010

S. Koskovsky Pond et al. 2009

W. Miller et al. 2007

J. Taylor et al. 2007

D. Blankenberg et al. 2007

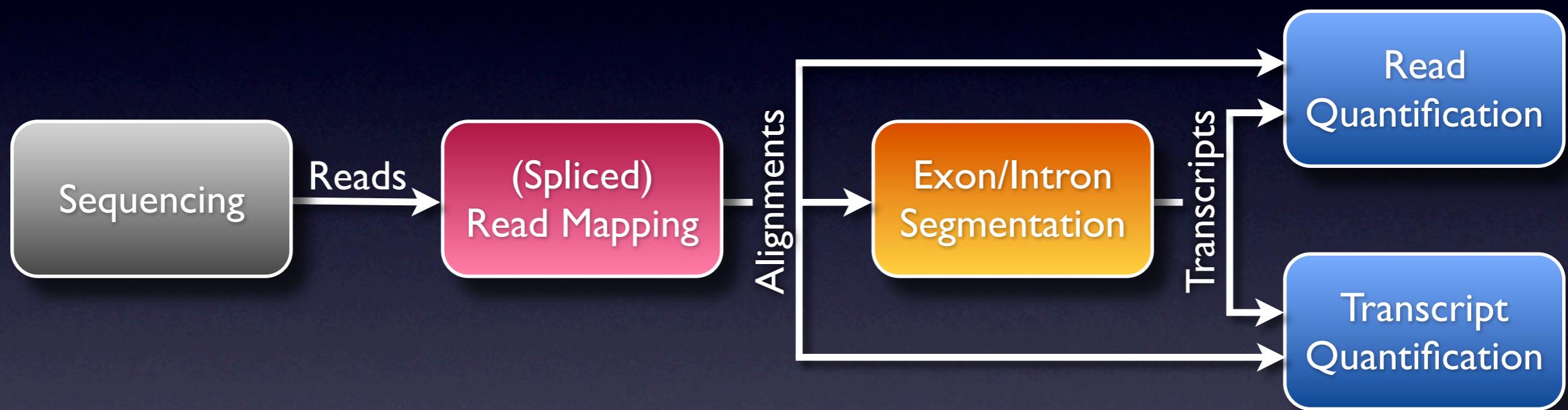
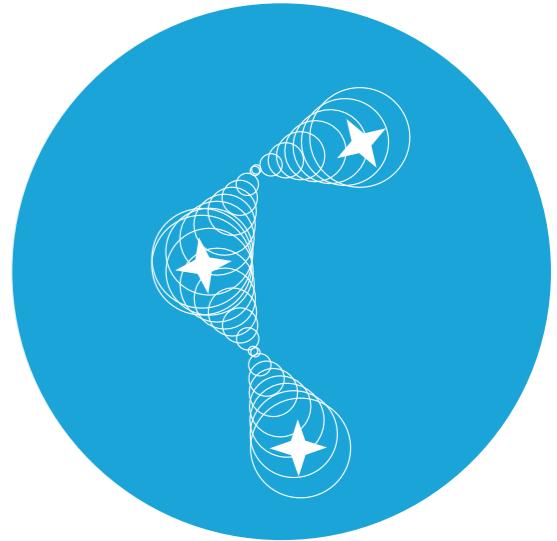
Giardine et al. 2005

- ▶ Persistent, reproducible approach to bioinformatics research
- ▶ Integration of tools made simple
- ▶ Source code release, VM/AMI, cloud instances, Galaxy pages

# Our Galaxy Tools:

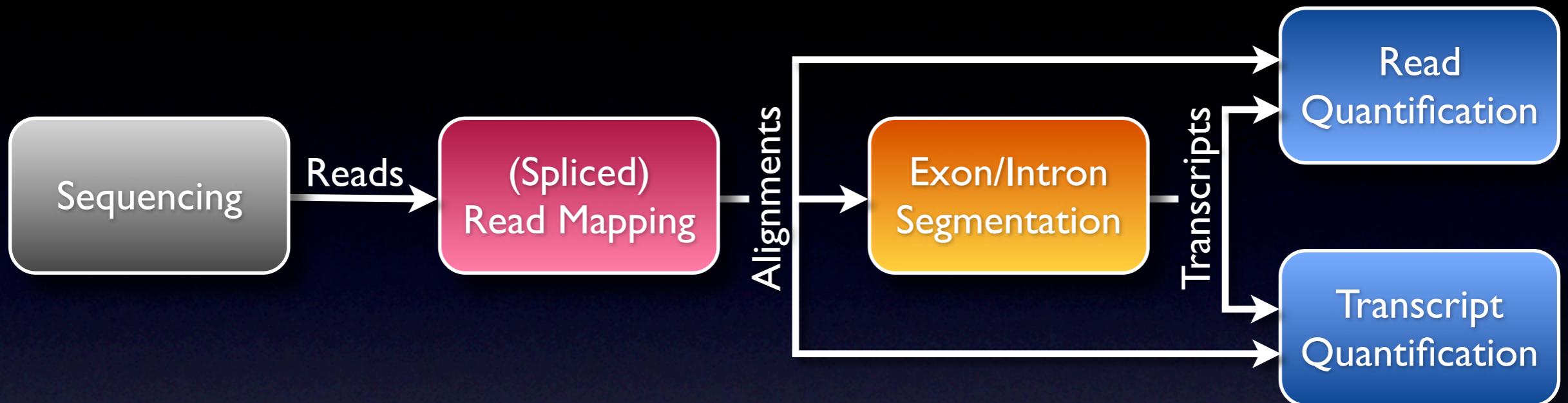
oqtans

online  
quantitative  
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analysis



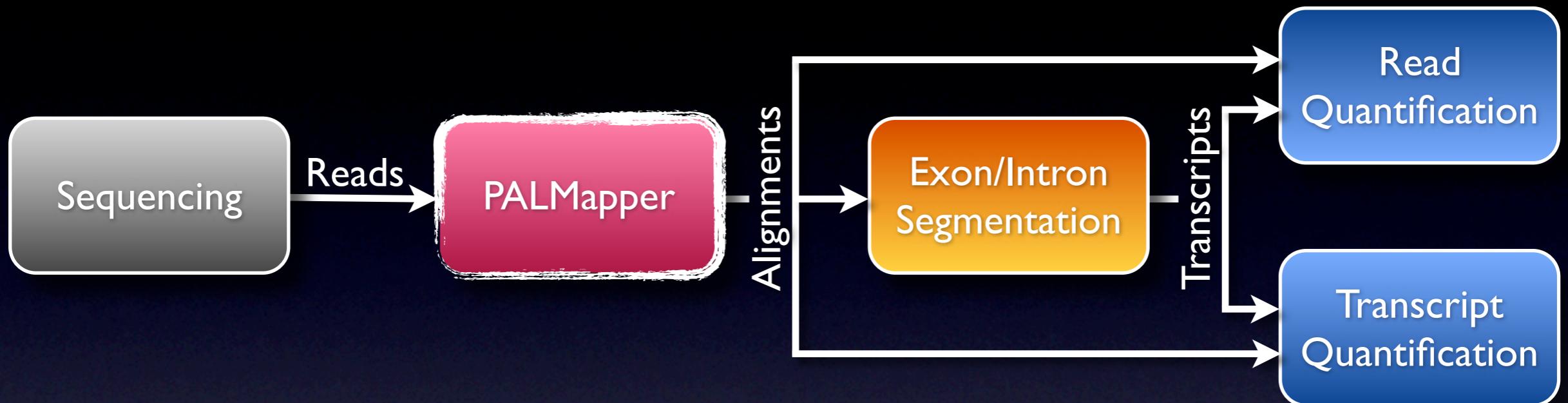
- ▶ Machine Learning-powered for quantitative analyses of RNA-seq experiments
- ▶ Workflow can be adjusted to your needs

# oqtans



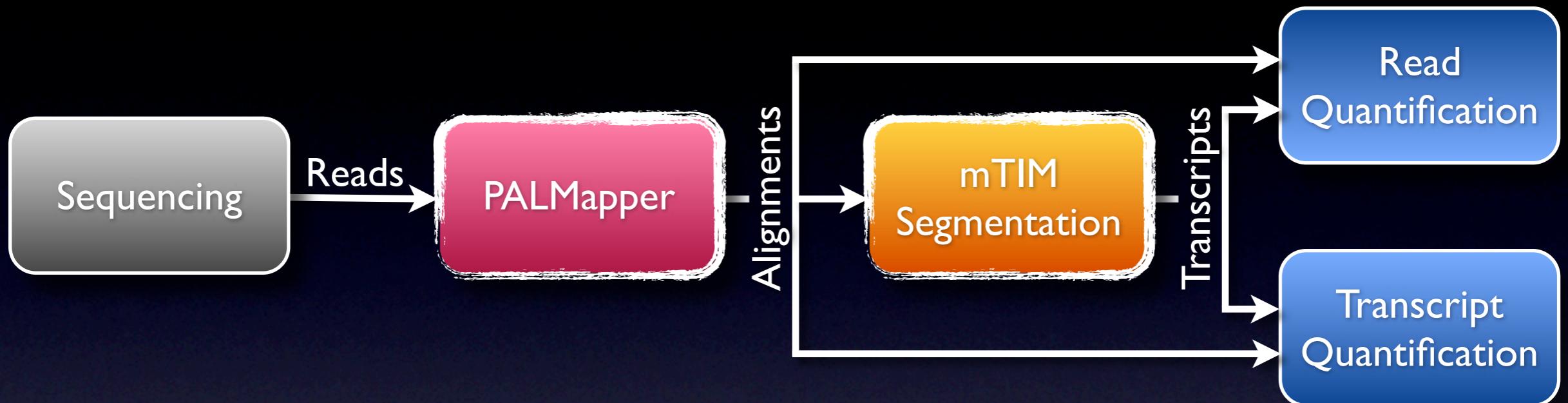
- ▶ Common experimental setups:
  - ▶ Identification of new transcripts
  - ▶ Comparison of samples

# oqtans



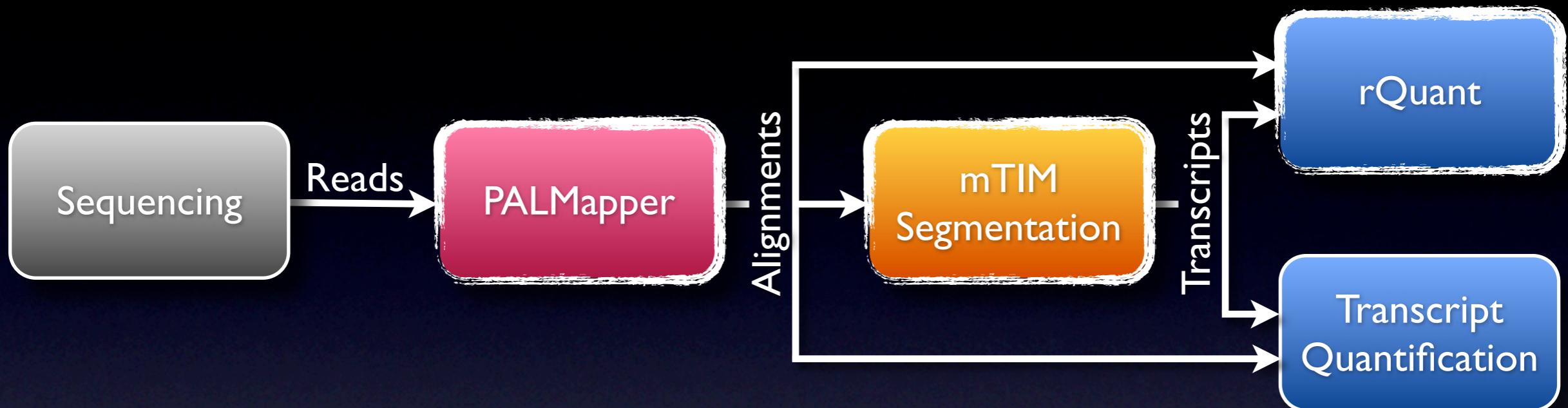
- ▶ **PALMapper:** highly accurate short-read mapper using base quality and splice site predictions

# oqtans



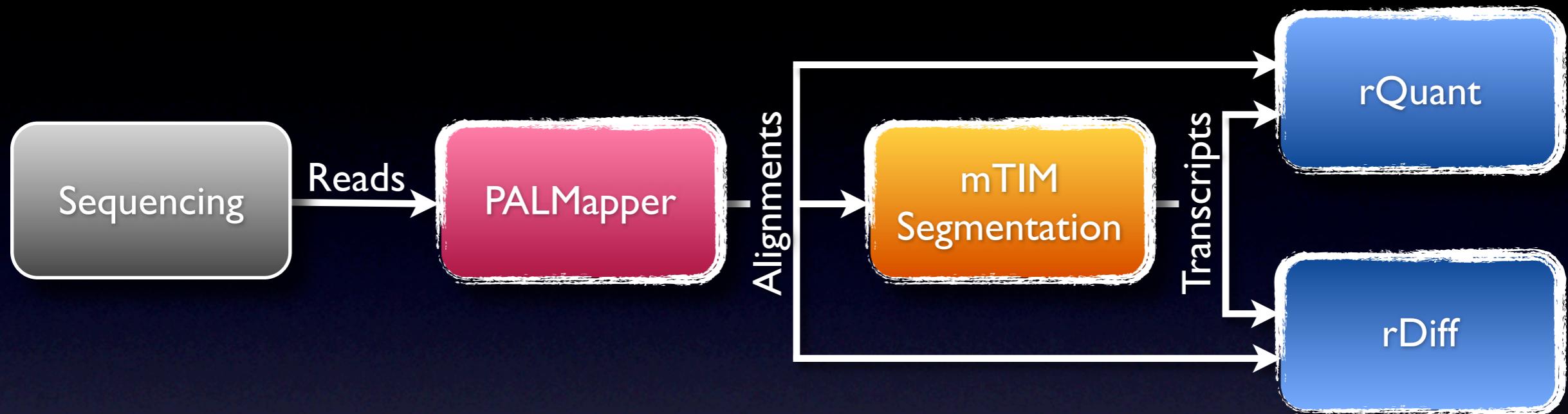
- ▶ mTIM: reconstructs exon-intron structure from alignments and splice site predictions
- ▶ SplAdder: adds isoforms to known annotation based on splice graph

# oqtans



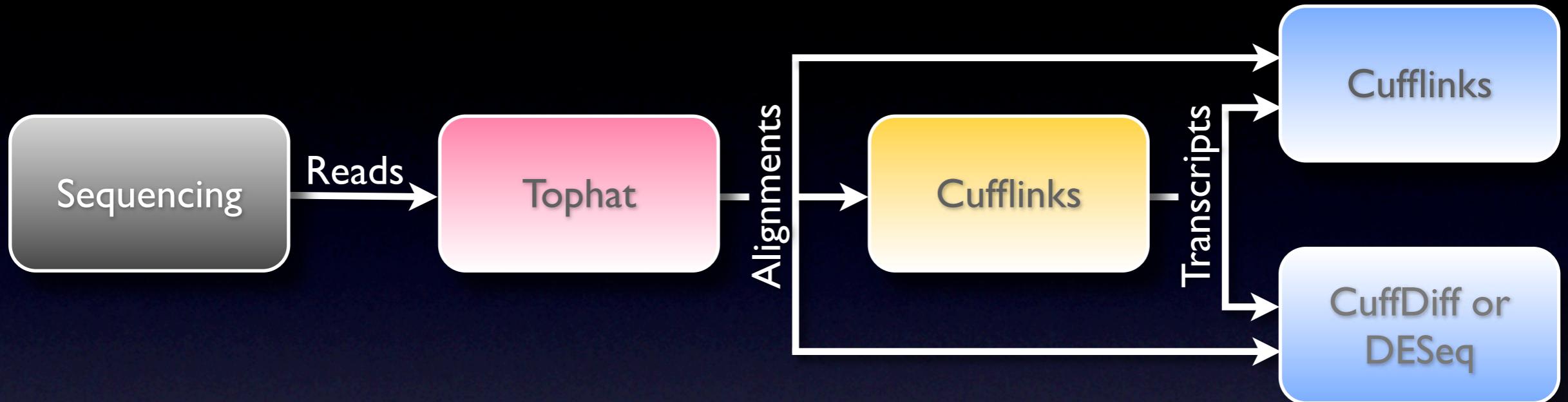
- ▶ **rQuant:** estimates biases in library prep, sequencing, and read mapping; accurately determines the abundances of transcripts

# oqtans



- ▶ **rDiff:** determines significant differences in transcript expression between experiments using statistical tests

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- ▶ Performance at or above state-of-the-art
- ▶ Tophat, DESeq, Cufflinks, CuffDiff for comparison

A. Roberts et al. 2011  
S. Anders and W. Huber 2010  
C. Trapnell et al. 2010

# Galaxy Tool Installers

Joint work: Enis Afgan & Galaxy Team  
James Taylor, Anton Nekrutenko  
AG Rätsch

- ▶ MLB group tools into any Galaxy installation
- ▶ Python Fabric scripts: used to manage automation of a remote server

```
install_cmd("wget %s" % self.tool_env['url'])  
install_cmd('chown -R %s %s' % (env.user, install_dir_root))
```
- ▶ Available from Galaxy Tool Shed
- ▶ Adjusted to Ubuntu on Galaxy Cloud Image

# Tools with Fabric Scripts

Joint work: Enis Afgan & Galaxy Team  
James Taylor, Anton Nekrutenko  
AG Rätsch

- ▶ Machine Learning Toolbox *Shogun*
- ▶ EasySVM tools for easy-to-use SVM classifications
- ▶ All oqtans tools
- ▶ GFF Toolkit for pre-processing annotations
- ▶ KIRMES regulatory modules identification
- ▶ WebLogo interface

# oqtans Availability

Joint work: Enis Afgan  
Galaxy Team  
AG Rätsch

- ▶ External cluster (21 nodes, 168 CPUs) handles requests to our Galaxy instance
- ▶ Fabrics to install tools from the shed
- ▶ AMI instance created
- ▶ ‘Instant on’
- ▶ Cloudman to launch as many as you need

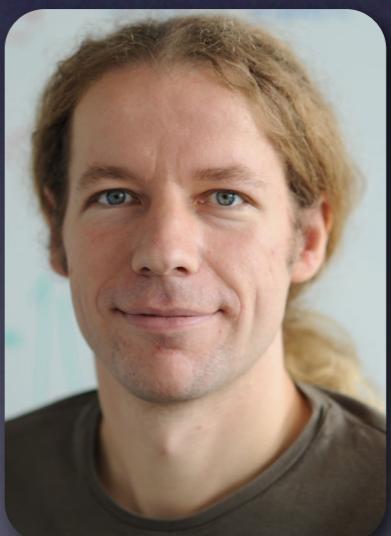
# oqtans Availability

- ▶ **MLB Group Galaxy Instance**  
<http://galaxy.fml.mpg.de>
- ▶ **EC2 Cloud Instance & AMI**  
ami-228a7e4b “Oqtans-Galaxy”
- ▶ **Fabric scripts install Oqtans on your Galaxy**  
<http://community.g2.bx.psu.edu/>
- ▶ **Source code releases of all tools, email info**  
<http://fml.mpg.de/oqtans>

<http://fml.mpg.de/oqtans>



Jonas Behr, Regina Bohnert, Philipp Drewe, Nico Goernitz, Géraldine Jean



André Kahles, Pramod Mudrakarta, Vipin T. Sreedharan, Georg Zeller, Gunnar Rätsch