

Fueled by:



Bisque

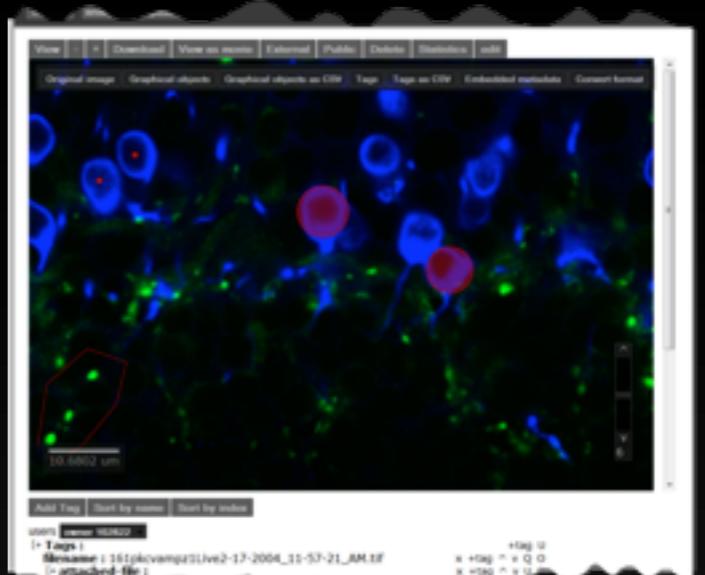
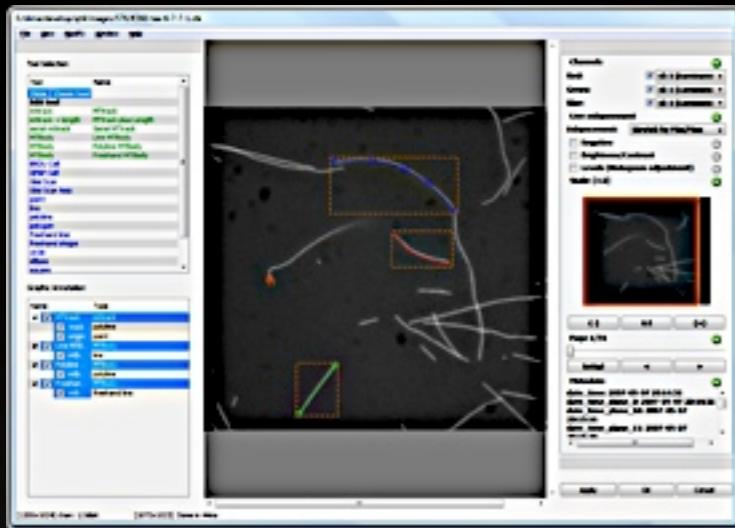
Bisque and iRods

Center for Bio-Image Informatics, UCSB

<http://bovary.iplantcollaborative.org>

Kris Kvilekval

Bisque - Image database



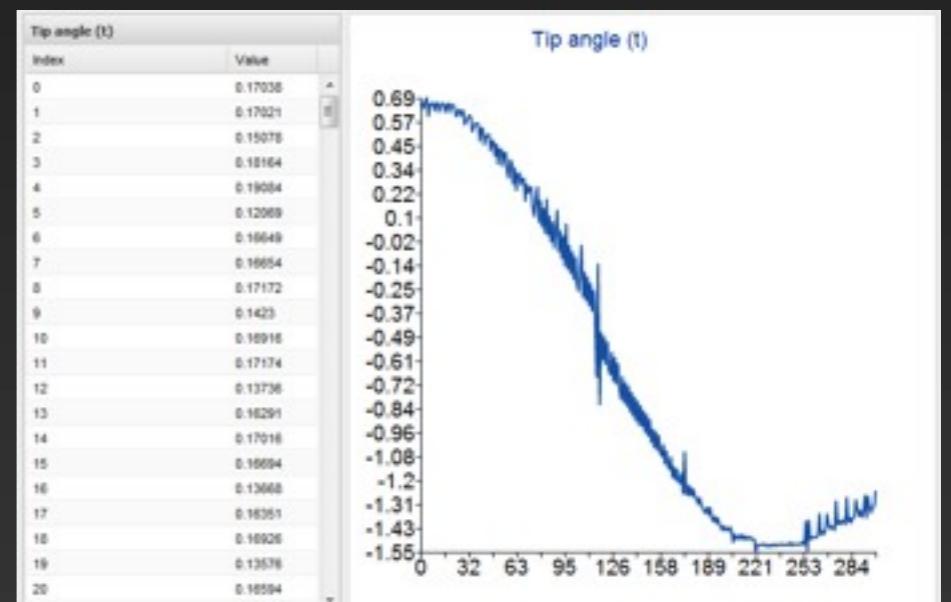
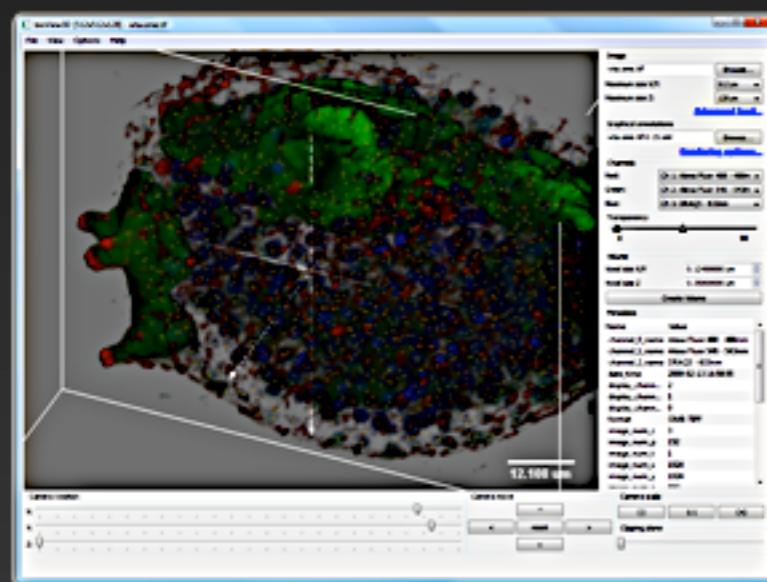
Ground truth

Automated analysis

Visualization

Flexible and
hierarchical
annotations

Generic statistics



Bisque basics

- **Support varying data models**
 - Database supports dynamic data model
- **Everything is a web accessible resource**
 - Image, Metadata, Analysis, Index
- **Scalable and distributed**
 - Add servers
 - Combine and use multiple diverse collections
- **Rich web clients for interactive analysis**
 - Web based applications

Flexible and hierarchical

- Hierarchical structure
- Flexible name:value fields
- Textual and 5D graphical annotations
- Biologically meaningful objects and groups

```
<tag name="description" value="some text" >
    <tag name="nested" value="http://host/someplace" type="object" />
</tag>
<gobject type="mttrack" name="MTtrack" >
    <gobject type="polyline" name="track" >
        <vertex x="206" y="542" t="0" index="0" />
        <vertex x="206" y="512" t="1" index="1" />
        <vertex x="232" y="502" t="2" index="2" />
    </gobject>
    <gobject type="point" name="origin" >
        <vertex x="212" y="552" />
        <tag name="color" value="#FF0000" />
        <tag name="probability" value="96.7" />
    </gobject>
</gobject>
```

Bisque architecture

Client tools



Scalable services

Image Services



XML

Data Services



XML

Analysis Services



XML



XML,
JSON

HTML

Client Services

XML

TWIX

Analysis Challenges

- Data organization and access
- Using Computational Resources
- Development methods

- Datasets
 - Storage hierarchies
 - Movement
-
- Local machines
 - Local/Remote clusters
 - Environment
-
- Language Neutral
 - Local development
 - Distribution

HT-Imaging Challenges

Automated imaging can produce
large-scale data

- Minimize data movement
- Protection/Ownership
- Integration with current workflow
- Metadata collection and binding
- Automated analysis

HT-Imaging

Automated Imaging



Bisque



Image ingest

- ✓ Discovery: iRules based or Polling
- ✓ Pre-processing: unpacking/construction
- ✓ Initial analysis: resource based

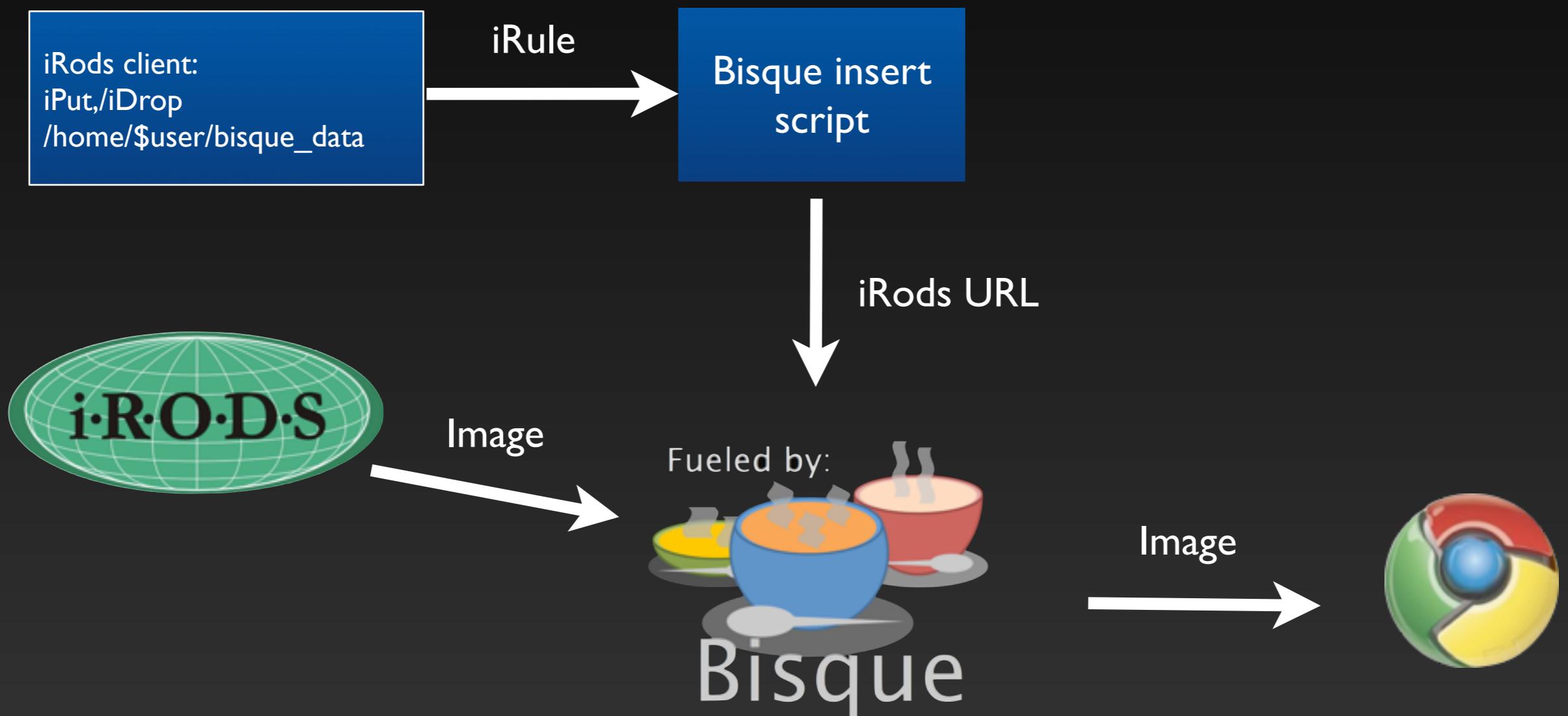
Data sources:

- iRods
- Local disk
- Uploads

Data types:

- Images/Files
- Metadata Records
- Experimental Objects

Discovery with iRods



iRodz Rule

irods://data.iplantcollaborative.org/home/<user>/
bisque_data/mybigfile.data

```
### server/config/reConfigs/core.re
### Example Rules for registering irods rules.
###

acPostProcForPut {
    ON ($userNameClient != "bisque" && $objPath like "/iplant/home/*/*bisque_data/*") {
        writeLine("serverLog","BISQUE: inserting object"++$objPath);
#        delay("<PLUSET>1s</PLUSET><EF>1s REPEAT UNTIL SUCCESS</EF>") {
        delay("<PLUSET>1s</PLUSET>") {
            msiExecCmd("insert2bisque.py", '$objPath $userNameClient', "winwood.iplantcollaborative.org", "null", "null", *cmdOut);
            writeLine("serverLog","BISQUE: inserted object"++$objPath);
        }
    }
}

acPostProcForCollCreate {
    ON ($collName like "/iplant/home/$userNameClient/bisque_data") {
        writeLine("serverLog","BISQUE: permitting bisque user RW on"++$collName);
        msiSetACL ('default', 'write', 'bisque', $collName);
        msiSetACL ('recursive', 'inherit', 'null', $collName);
    }
}

#### NEED acPreDelete
#### NEED acPreRename (in and out of bisque_data)
```

Bisque insertion

```
#!/usr/bin/env python
import sys
import shlex
import urllib
import urllib2
import urlparse
import base64
import logging

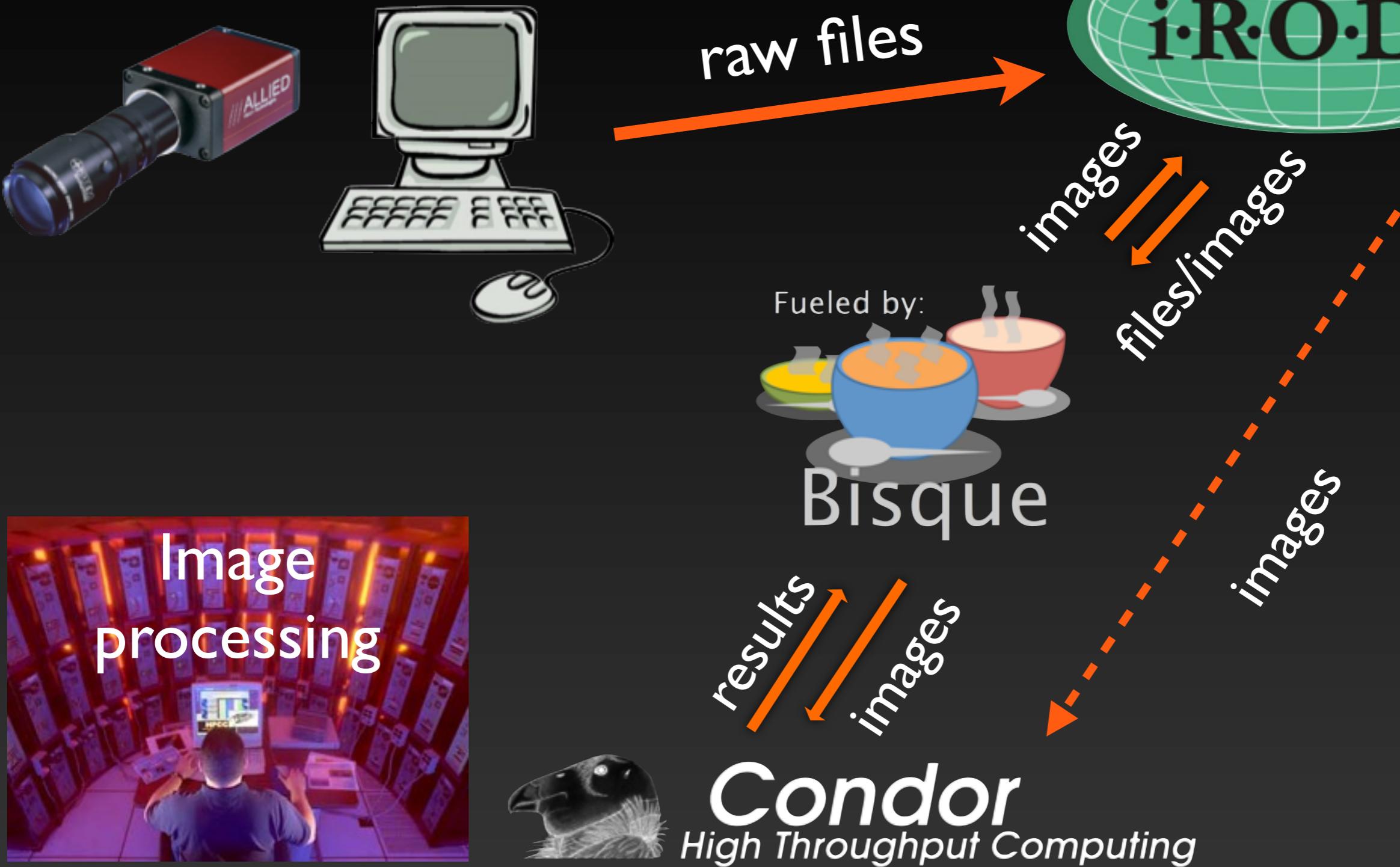
#####
# Config for local installation
LOGFILE='/tmp/bisque_insert.log'
BISQUE_HOST='http://bisque.ece.ucsb.edu'
BISQUE_ADMIN_PASS='guessme'
IRODS_HOST='irods://irods.ece.ucsb.edu'
# End Config

logging.basicConfig(filename=LOGFILE, level=logging.INFO)
log = logging.getLogger('i2b')

def main():
    log.debug( "insert2bisque recevied %s" % (sys.argv) )
    try:
        obj = sys.argv[1]
        user = sys.argv[2]
        url = "%s/import/insert%s" % (BISQUE_HOST, urllib.urlencode( { 'url': IRODS_HOST+obj, 'user': user} ))
        request = urllib2.Request(url)
        request.add_header('authorization', 'Basic ' + base64.encodestring("admin:%s" % BISQUE_ADMIN_PASS).strip())
        r = urllib2.urlopen(request)
        response = r.read()
        log.info( 'insert %s -> %s' % (url, response))
    except Exception,e:
        log.exception( "exception occurred %s" % e )
        raise e

if __name__ == "__main__":
    main()
    sys.exit(0)
```

HT image pre-processing/analysis



Challenges

- Pre-processing ops require local files
 - ex: unpacking and image construction
 - Should it move to iRod's server?
- Analysis also requires local files
 - Should it also move to iRod's server?
- Should an IS run on the iRod's server?
- iRod's on clusters?

BQPhytomorph

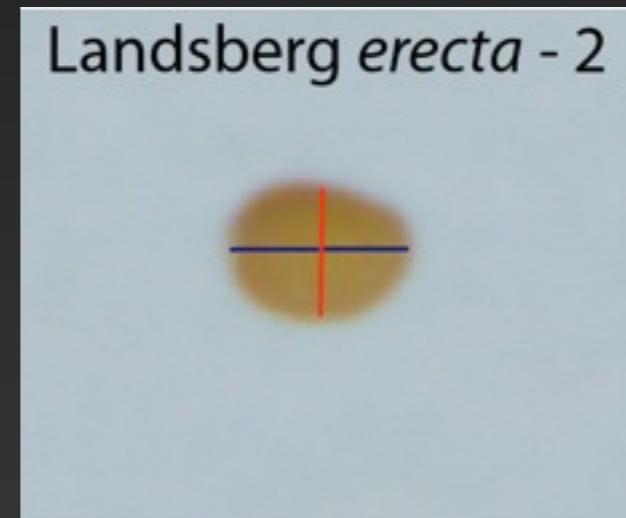
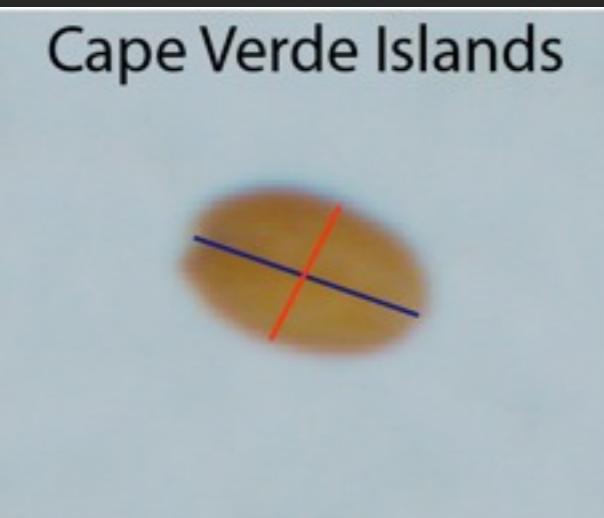
Bisque

B.S. Manjunath
Kris Kvelikval
Dmitry Fedorov

Phytomorph

Edgar Spalding
Nathan Miller
Logan Johnson

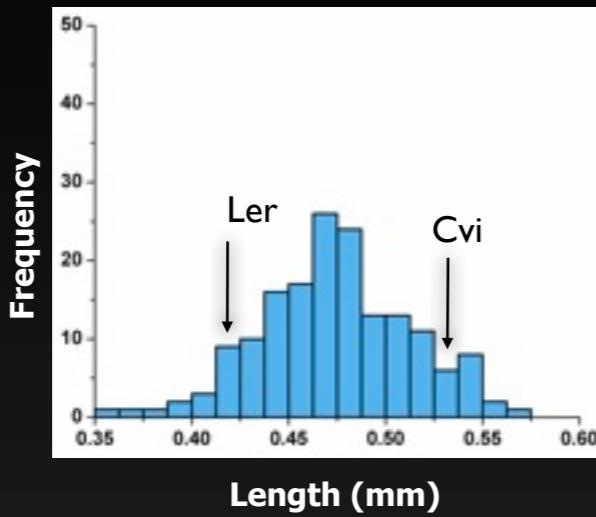
Whole Seedling-size Analysis



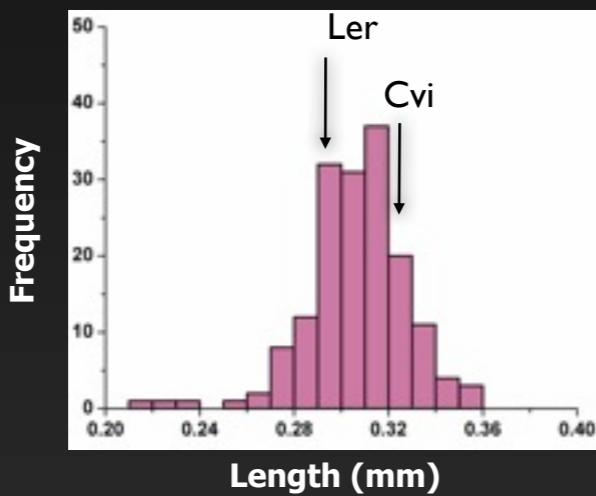
— Major Axis
— Minor Axis

Seed Size Features

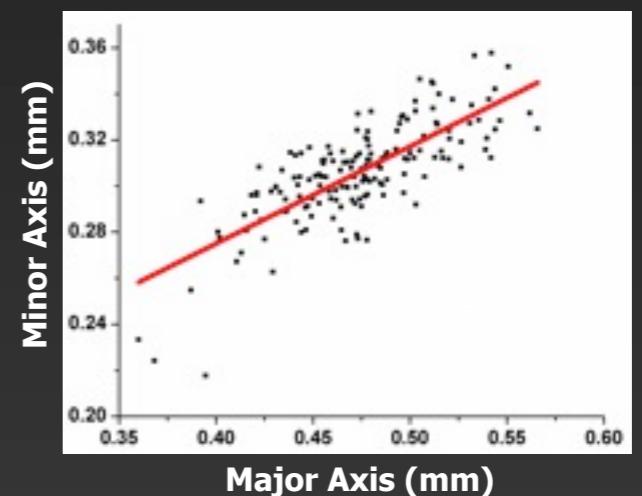
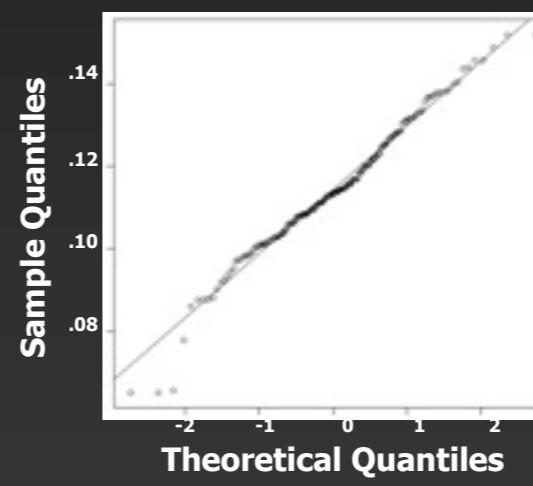
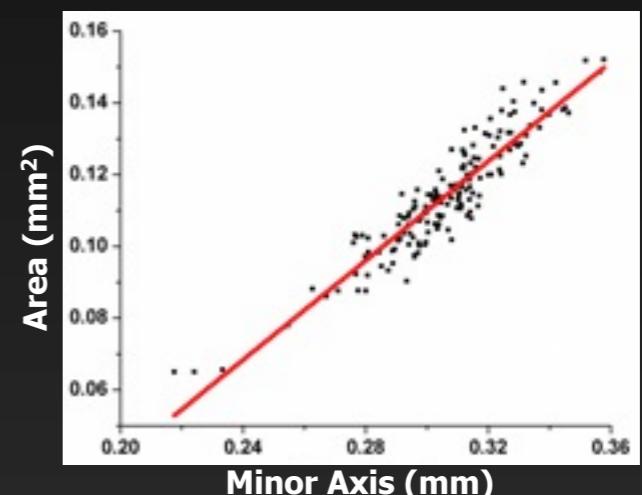
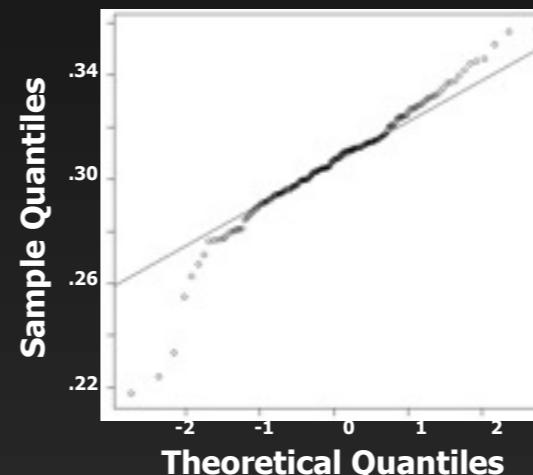
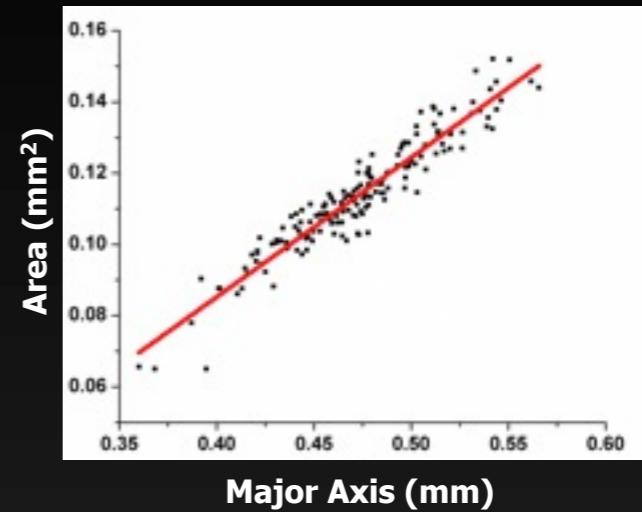
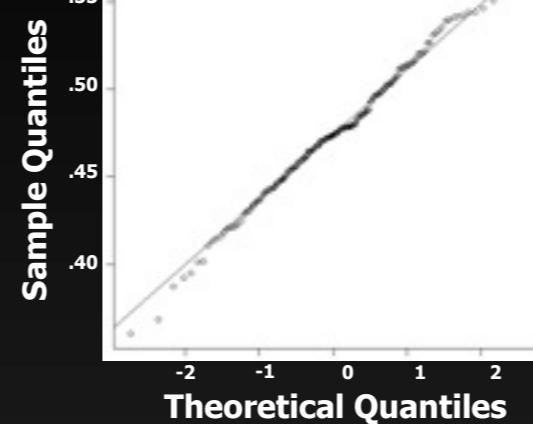
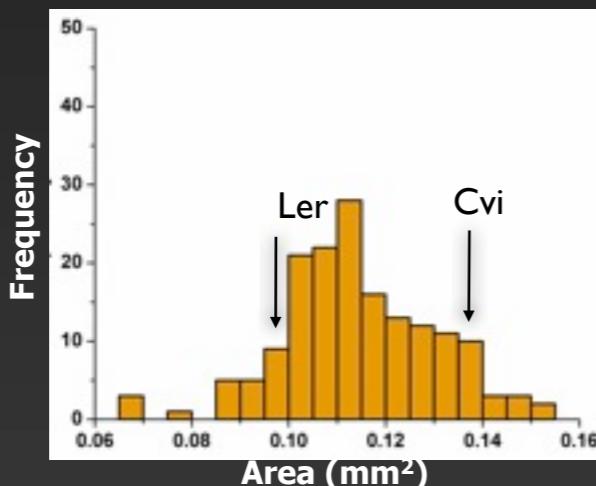
Major Axis



Minor Axis



Area



Root Tip Angle

Cvi Ler



9 h after
gravistimulation

164 lines
X 10 seedlings/line
1640 movies



QTL of gravitropism

Source: Edgar Spalding

Experimental Setup

Seeds Planted



1 mM KCl, 1 mM CaCl₂, 5 mM MES,
pH 5.7

2-4 day Stratification



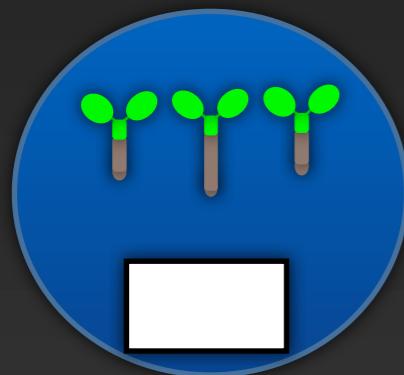
2-4° C

3 days in growth chamber



constant white light,
22.5° C

Record Initial
Root Lengths



Place in front of camera in
CCW orientation

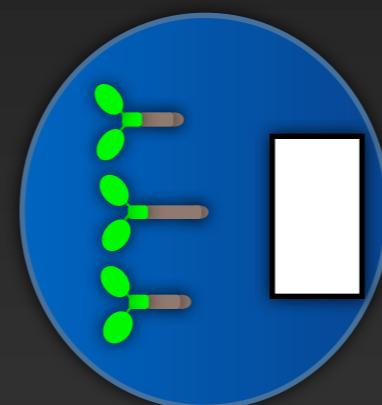


Image at 100 px/mm every
2 min for 8 hr

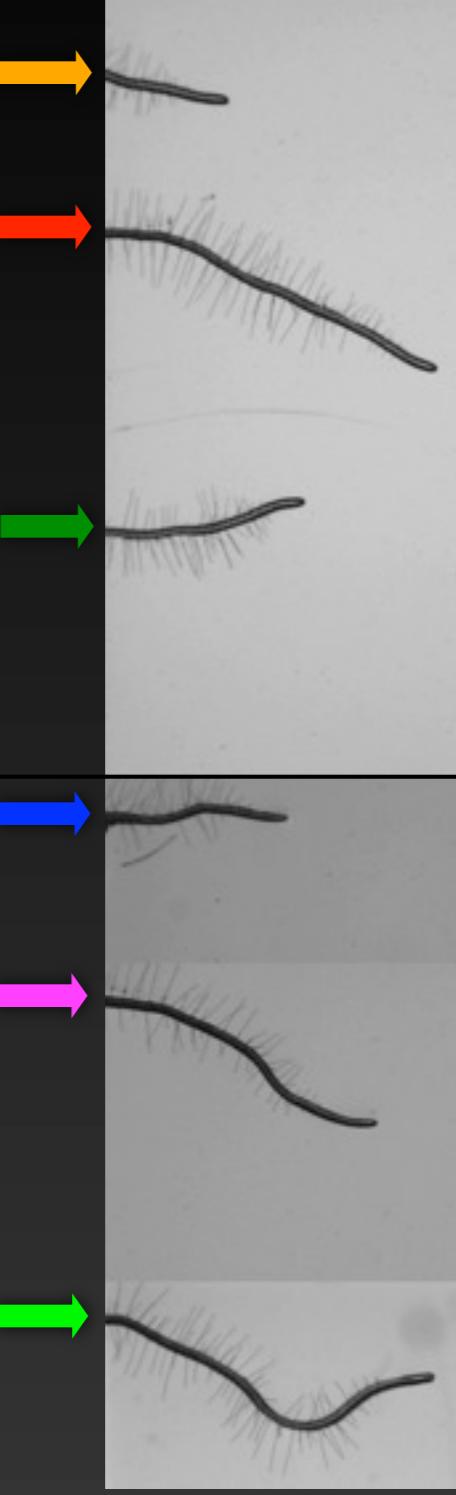


Source: Edgar Spalding

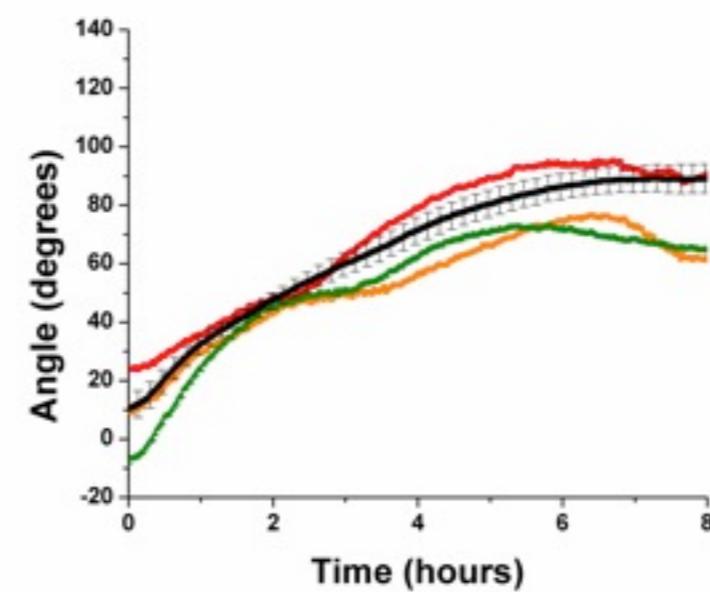


Multi-Tip and Growth-Rate

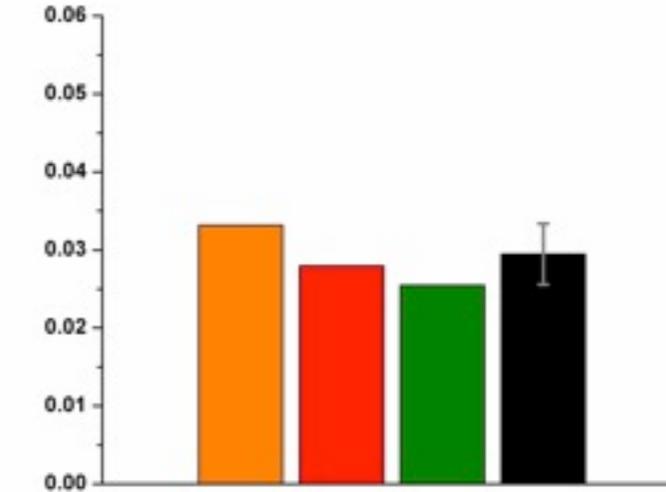
Cape Verde
Islands



Tip Angle



Average Growth Rate



Landsberg
erecta

