

# Overview of Sample Management

Mansi Chovatia March 24, 2015



### **Functions**



- Sample Receipt
- Sample Tracking/Inventory
- Sample Quantification & Quality Assessment
- Sample Aliquots

Technical Support

### **Operation**



Sample Storage Sample QC

Sample Aliquot

# Sample Storage



-Coordinates shipping schedule -Ships out prelabled containers

Supply Chain Manager



#### **Sample Receipt**

- -Receive packages
- -Barcode association
- -Log sample into GLS



Project Manager

- -User Agreements
- -Project initiation
- -Sample metadata
- -Project tracking

JGI Users



Sample Management

#### Sample Return & Disposal

- -Completed projects returned to users or discarded
- Tracked in JIRA tickets

# Sample QC



- Measure sample volume
- Top-off low volume samples
- Quantify samples using fluorometric dyes
- Assess sample quality based on scope of work (SOW)
- Record measurements and analysis in LIMS

# Sample Aliquot

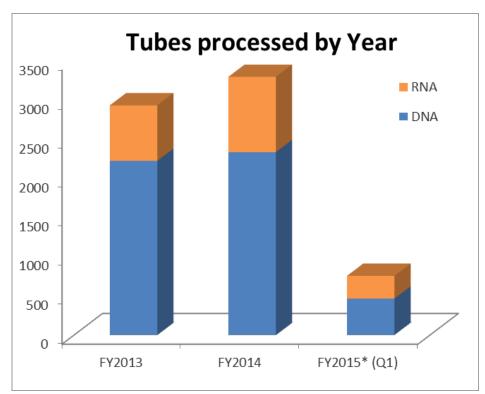


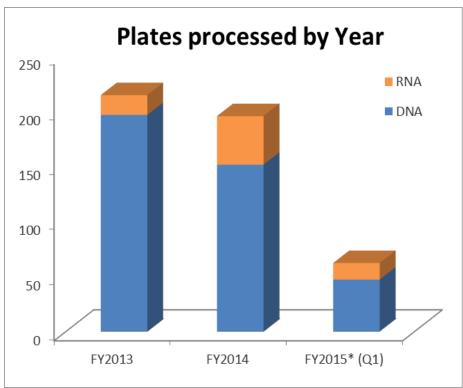
- Transfer required amount for library construction
- Top-off low volumes to required amount
- Create intermediate dilution for highly concentrated samples
- Re-array partial plates and tubes into a single final destination plate for library construction

Record sample transfer information in LIMS

# **Throughput**







### **Automation**



















#### Storage

#### Volume Check

QC

#### Aliquot

- Sample archival
- Manage in-progress containers
- Houses a total of 532 plates and 11,320 tubes.
- Barcode association Retrieves upon command
- Improved tracking
- **Provides security**
- Eliminates the human error

- Sonic pulses
- Non-contact measurement
- Compatible with plates and tubes
- Microplate reader for quantification
- NanoDrop for the purity
- Agarose gels for **DNA** quality
- 2 LabChips for RNA sample quality

- 2 STARlets in production
- Main workforce for both sample QC and aliquots
- Integrated tube decapper
- Re-array capability
- High consistency
- Reduces ergonomic issues

### Sample Receipt

















#### **Correct Packaging**



Example of tube WITH secondary containment (15mL conical tube on crushed dry ice).

Multiple sample tubes can be placed in a larger conical tube or secondary containment.



Properly labeled and tightly sealed with a FOIL seal. Ensure that the tab of the foil seal is NOT on the plate label side. Make sure label is on A12 side.



Example of a plate WITH secondary containment (cardboard box on crushed dry ice).

#### Incorrect Packaging





Example of tubes arriving WITHOUT secondary containment (placed directly in dry ice).



DO NOT use clear plate seals.



Example of a plate arriving WITHOUT secondary containment - plate seal has been damaged.

#### **Common concerns:**

- Packaging
  - Dry ice
- Secondary containers
  - Container seals
  - Labels/documents

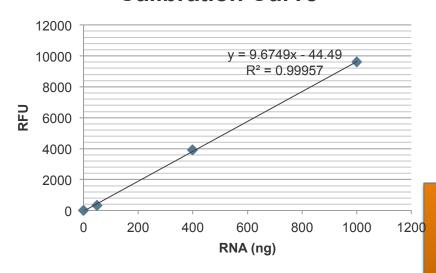
# **Sample Quantity**

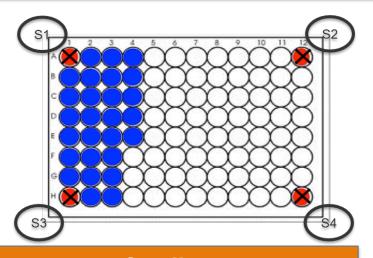


# DNA assays Quantifluor dsDNA system (0.4-20ng, 2-200ng)

RNA assays
Quant-iT RNA BR kit (50-1000ng)
Quantifluor RNA system (1.5-80ng)

#### **Calibration Curve**



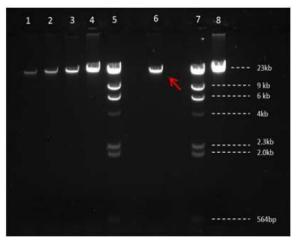


#### **Common cause for discrepancy:**

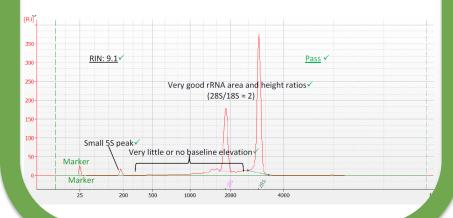
- Pipetting errors
- Calibration curve
  - Contaminants
    - Assays/kits
    - Standards

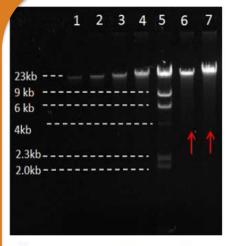
# **Sample Quality**



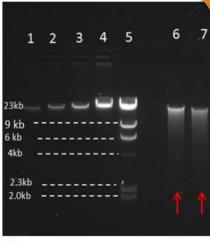


A. Lane 6 - High molecular weight DNA at 23kb

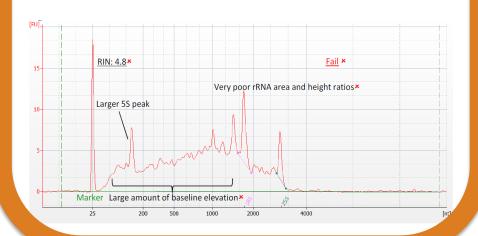




B. Lane 6 & 7 – Partially degraded DNA (smearing visible)



C. Lane 6 & 7 - Completely degraded DNA



### Sample QC Requirements per Library



Relaxed Stringent

Volume Check

All library types

#### Quantity

All library types except:

# Amplicon sequencing

Ribosomal iTags

#### Quality

#### **LMP**

- > 8kb CLRS
- > 4kb CLRS
- 2.5kb CLRS
- Nextera LMP

# PacBio Long reads Stranded RNA-seq

- Poly A selection
- > rRNA depletion
- > FFPE/LCM

#### **smRNA**

- Eukaryote
- Prokaryote

#### Full length

Iso-seq

#### Purity

#### Long reads

- PacBio 20kb
- PacBio >10kb
- PacBio 3kb
- PacBio 2kb

## **New Technology**



# Instrument & Method Evaluation

- Advanced Analytical Fragment
  Analyzer
- Agilent Tape Station
- Biotek Synergy
- STARlet method validation
- SAM validation



#### **Optimization**

- Chlamy DNA/RNA sample prep
- Microbial DNA sample prep
- Biomatrica RNAstable & DNAstable
- Promega Quantifluor DNA & RNA systems









# Technical Guidance

- Genomic DNA QC protocol
- Total RNA QC protocol
- Chlamydomonas RNA extraction
- Etc...



# Sample Management Team



