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Title

Joint Genome Institute's Automation Approach and History

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Joint Genome Institute's Automation Approach and History

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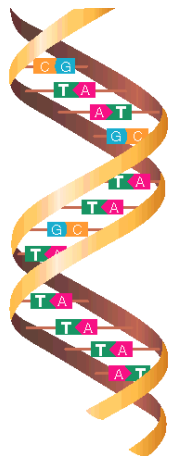
Simon Roberts

(Production Instrumentation Supervisor)

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Agenda

- **Brief overview of the how the Joint Genome Institute came into existence.**
- **Overview of DNA sequencing production line at the JGI.**
- **How our throughput has increased since 1999 to become a high through-put sequencing facility.**
- **Some instrumentation improvement highlights along the way and how they are used.**
- **Review our approach to successful selection & implementation of new instruments to meet our needs.**

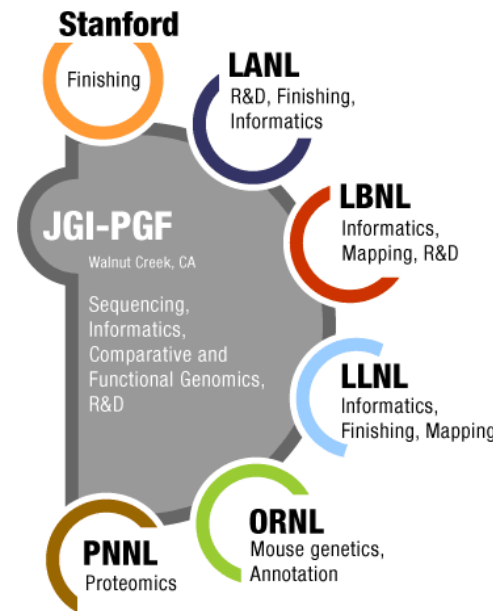




DOE JGI Production Genomics Facility



Opened in 1999
~240 UC Employees
60,000 sf
~\$66M Annual Budget



The DOE Joint Genome Institute (JGI) is a "virtual institute" that integrates the sequencing and analytical activities of six partner institutions:

Mission:

DOE JGI collaborates with DOE national laboratories and community users, to advance genome science in support of the DOE missions of clean bio-energy, carbon cycling, and bioremediation.

Important Dates in DOE Genomics

- **1986** DOE announces Human Genome Initiative. With \$5.3 million, pilot projects begin at DOE national laboratories to develop critical resources and technologies.
- **1990** DOE & NIH present their joint HGP plan to Congress. The 15-year project formally begins.
- **1997** DOE creates the JGI uniting activities at DOE human genome centers.
- **1999** JGI opens the Production Genomics Facility (PGF) in Walnut Creek, staff from LLNL & LBNL.
- **2000** HGP leaders & President Clinton announce the completion of a “working draft..the first great technological triumph of the 21st century.”
- **2003** HGP completed and published.



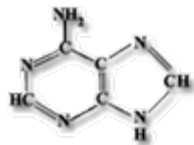
What IS a Genome???

A **GENOME** is all of a living thing's genetic material.

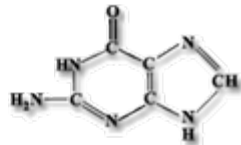
The genetic material is **DNA** (**D**eoxyribo**N**ucleic **A**cid)

DNA, a double helical molecule, is made up of four nucleotide "letters":

A--



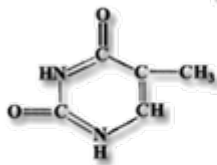
Adenine



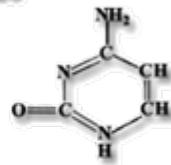
Guanine

Purines

T--



Thymine

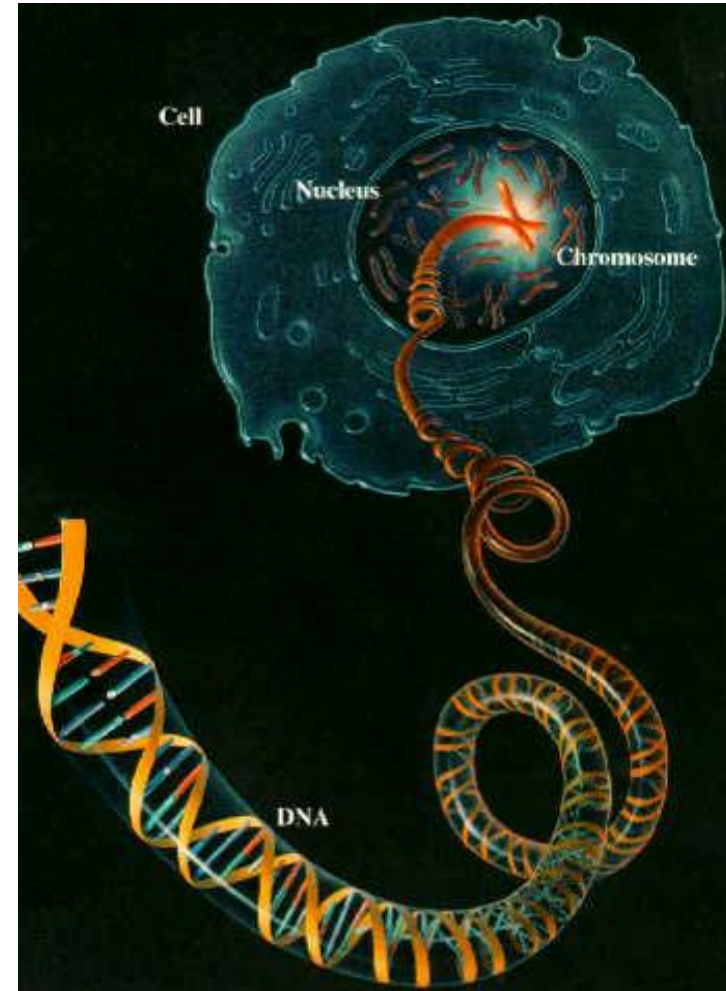
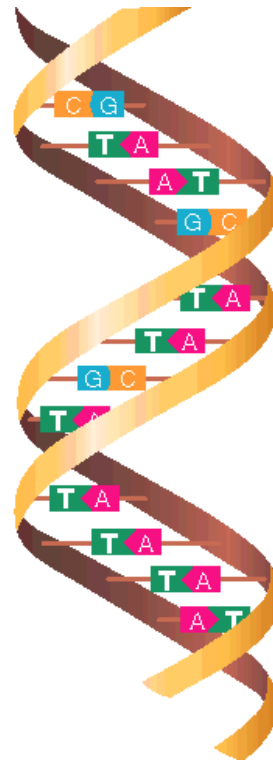


Cytosine

Pyrimidines

G--

C--

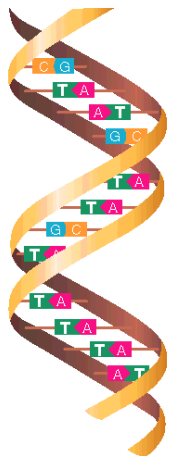


JGI Production Mission

**“Produce high quality cost efficient
‘assemble-able’ sequence in a safe
environment.”**

FY 2006 Goals

- 52 million lanes = 35 billion bases



How Sequencing is Done

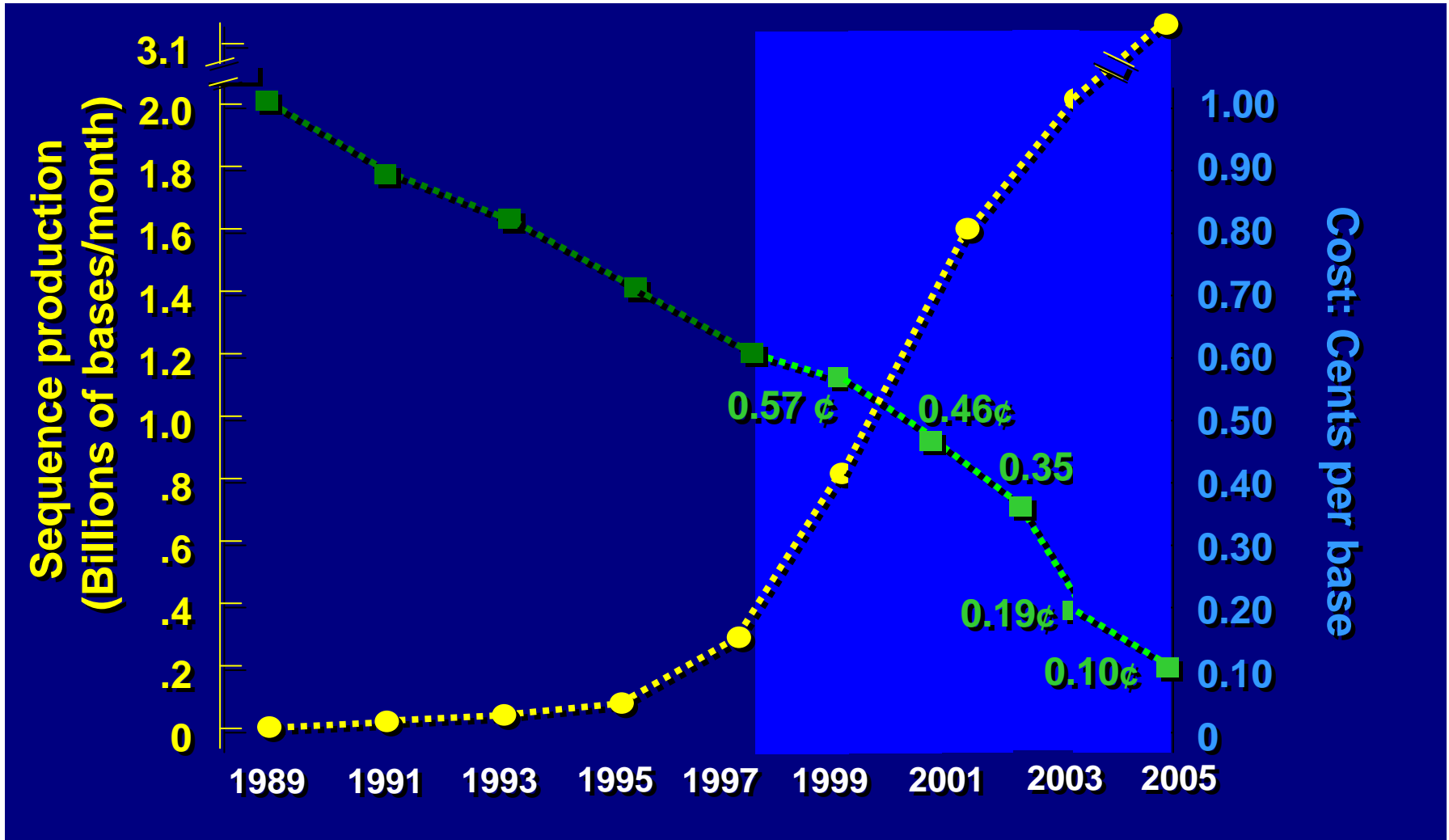
Production Line Overview

- **Library Creation**
 - Shearing the DNA (Genomic Solutions Hydroshear)
 - Insertion of Fragments into Plasmid (Ligation)
 - Transformation (Electroporation)
 - Subcloning the Sheared Fragment (Plating)
 - **Colony Picking (Genetix QPix)**
- **Production Sequencing**
 - Lysing the Cell (Matrix PlateMate)
 - Rolling Circle Amplification (MultiDrop Micro)
 - **Sequencing Chemistry (CyBio Vario)**
 - Post Sequencing Reaction Cleanup (BioMek FX)
 - **Capillary Sequencing (ABI 3730 & MB 4500)**
- **Assembly & QA**
 - Assembly (Phrap, JAZZ)
 - Quality Assessment (Phred Q20)





DOE JGI Production Sequencing Efficiencies



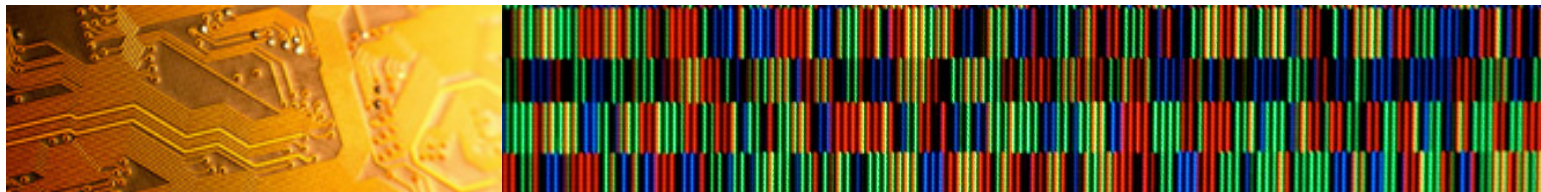
“Moore’s Law” of DNA Sequencing

- **April 2002:** 1 Gb/month
- **January 2004:** 2 Gb/month
- **July 2004:** 2.5 Gb/month
- **March 2005:** 3.1 Gb/month

(equivalent to 1 human genome/month)

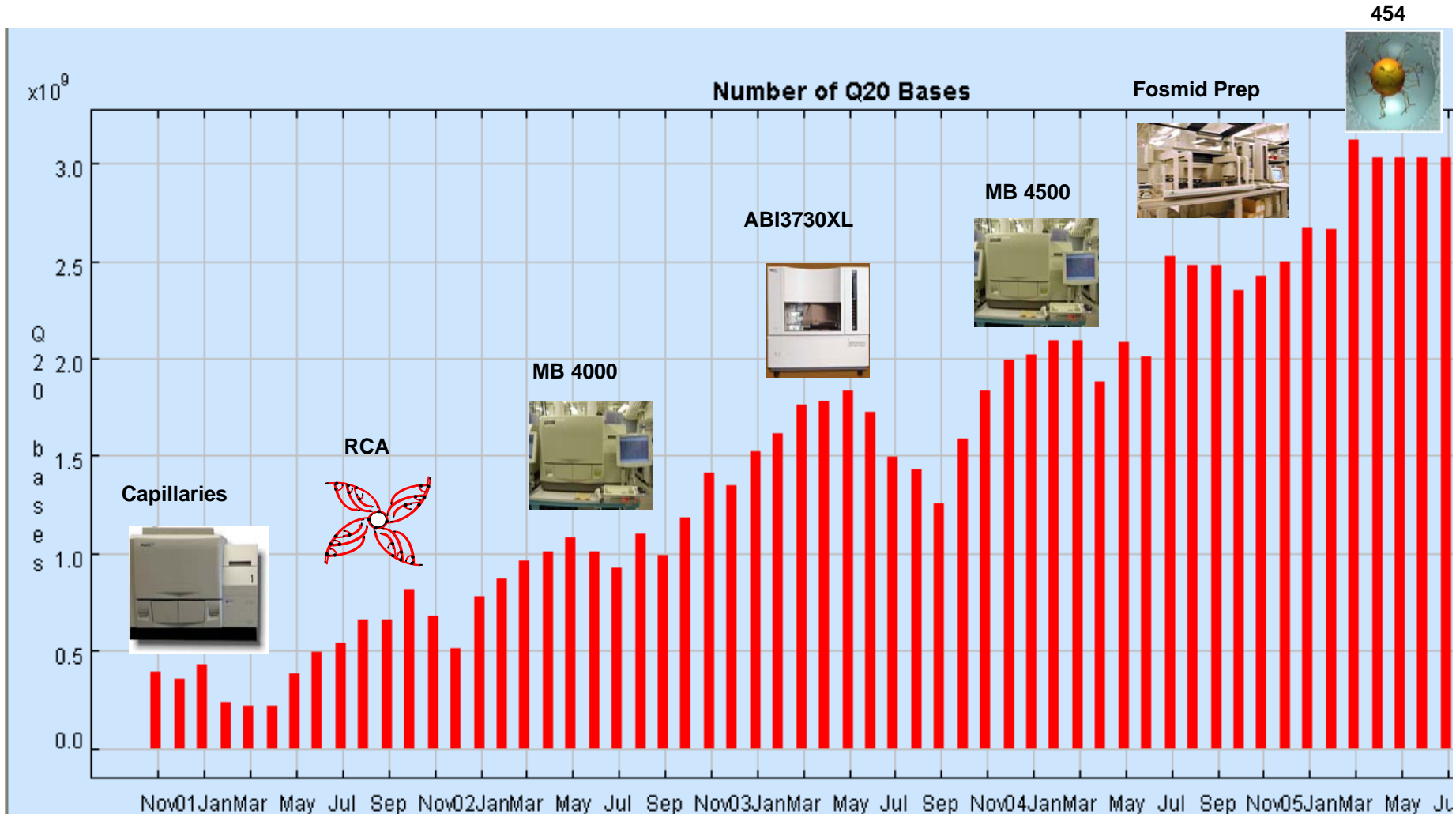
Total (3/99-4/06) 114 Gb

(equivalent of sequencing 38 human genomes)



Main Production Metrics; Read Length & Pass Rate

New Technologies Fuel Growth in Capacity and Reduce Costs



Capillary Sequencing

MegaBace

1000



Qty 10; 2000 to
Qty 84; 2002

4000



Qty 21; 2002 to Qty 36, 2003

4500



Qty 36, upgraded 2004

7 days a week operation

400 plates/day

ABI

377



Qty 28; 1997



Qty 5; 2001
Qty 35; 2002
Qty 55; 2003

3730s



Qty 70; 2004



7yr Throughput Increases Correlated with # Sequencers

ABI

28, ABI377s

5, ABI3730s

35, ABI3730s

55, ABI3730s

70, ABI3730s

MB

20, MB1000s

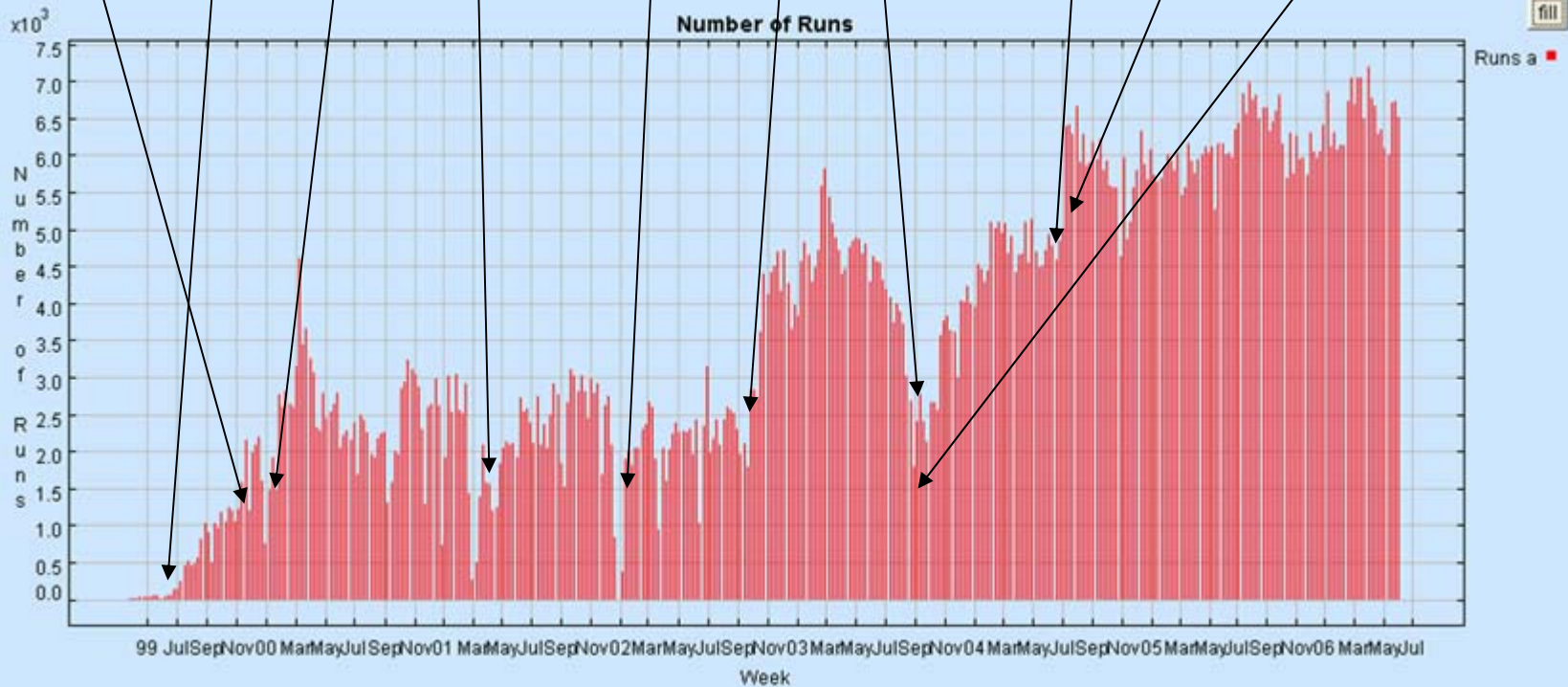
84, MB1000s

21, MB4000s

36, MB4000s

35, MB4500s

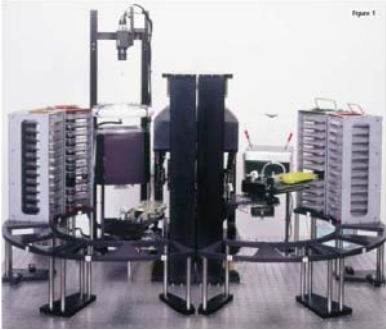
Weekly Number of Runs Summary



TOTAL, Sequencer Type

Colony Picking

Agar Colonies on BioAssay Plates into 384 Destination Plates



1999 Genomic Solutions & Hybaid Colony Pickers



OLD

Throughput ~Trays/day

96 pin picking head

Throughput (2 shifts = 16hrs)

~115 Bioassay Trays to produce 300 (384 well) destination plates/day based on ~1000 colonies/tray



QPix2 XT
(Qty 1; 2004)
2nd Due 7/06

NEW

QPix2 (Qty 1; 2000, Qty 3; 2001, Qty4; 2002)

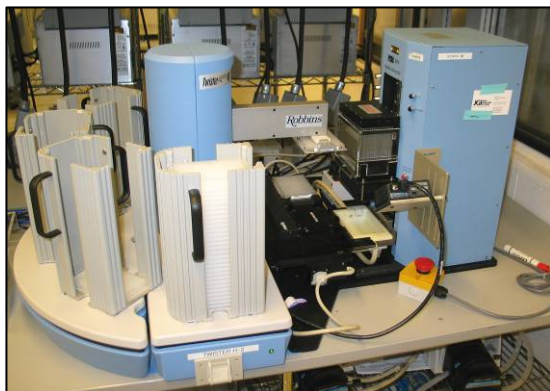


2 BioAssays & Stacked Dest Plates

Sequence Chemistry

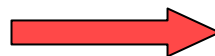
Aliquot RCA Product to Daughter Plates
& Dispense FWD & REV Chemistry

**Hydra-Twister
System**
Qty 2; 2001



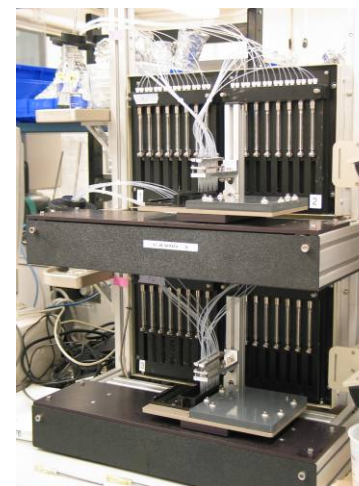
**21 source to 42
destination ~1hr/batch**

OLD



**Cavro Syringe
Pump Dispenser
System**

**42 plates
~40min/batch**



**Typical Aliquot 1.5uL RCA Product into
two destination plates followed by 3.5uL
Sequencing Chemistry Cocktail Dispense**

Throughput (2 shifts = 16hrs)

**6 batches, 42 source plates = 84
destination plates**

500+ plates/day

NEW

**CyBi-Well Vario
Integrated System**
Qty 2; 2006



2.5hr/batch



Why Automate?

- **In high throughput environment;**
 - automation does not necessarily increase productivity of workers,
 - does increase repeatability
 - increase equipment reliability
 - frees up operator to perform other tasks
 - reduce risk to operator of ergonomic issues
- **“Islands of Automation”**
 - modular instruments with stacks of micro-titer plates transported & loaded by operators
- **Semi-Automated Approach**
 - minimizes cost and maximizes flexibility
 - compared to fully robotic systems
 - presents a unique set of challenging issues when moving large volumes of plates



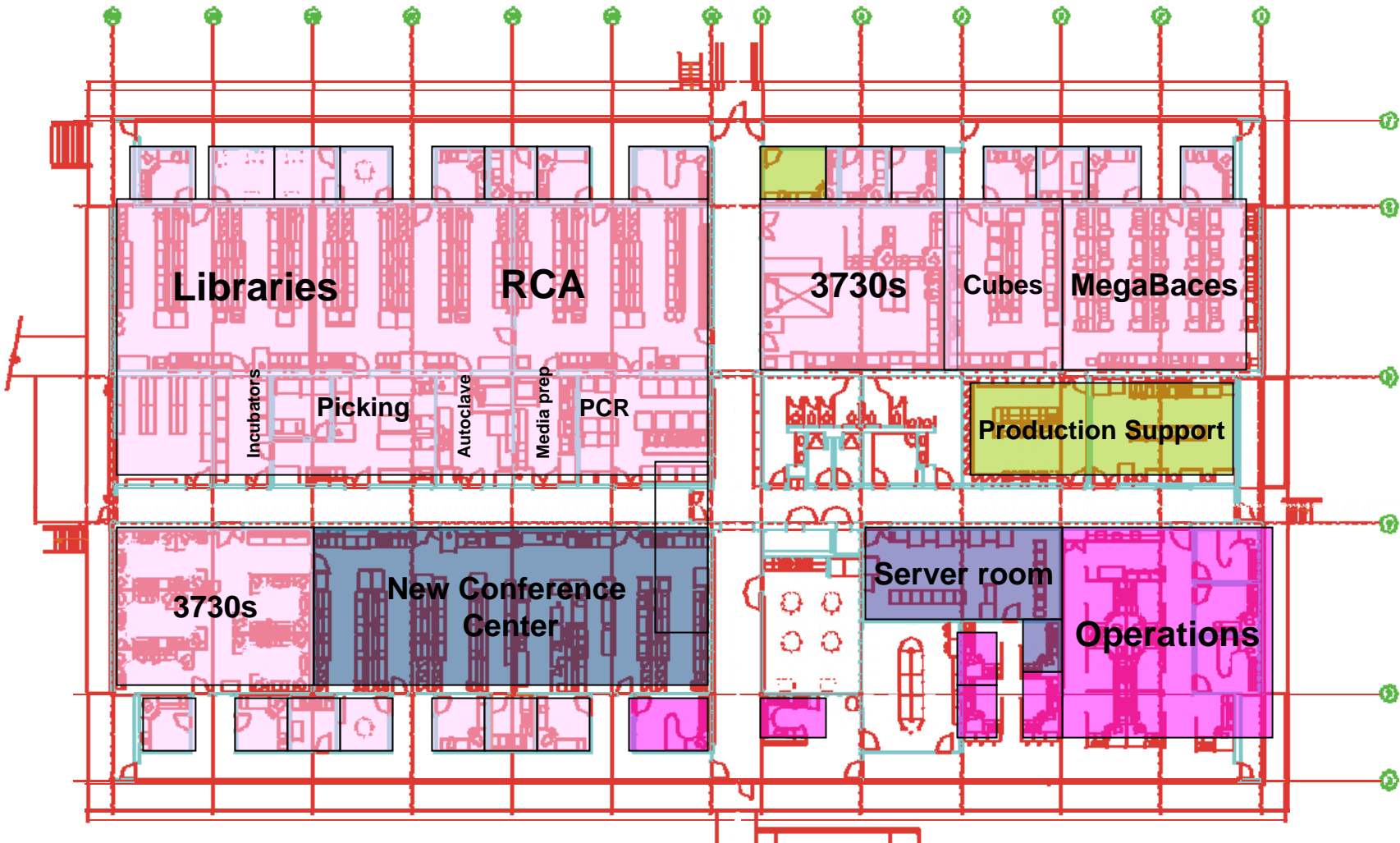
Why Automate?

- **Need to automate driven by;**
 - Lowering Reagent Volumes of expensive reagents <1uL & increments of 0.1uL
 - Dilutions only go so far
 - Wet & dry dispense
- **1536 not going to work in the near term**
 - sequencers in 384 format
 - no thermal cycling
- **Physical constraints of laboratory space, big issue currently (next slide)**

2006 Community Sequencing Program



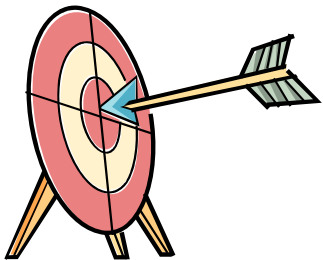
JGI Facility Layout



Approach to Automation Selection

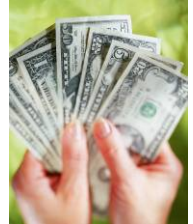
- **Performance**

- Configuration, plate types (source & destination, conditioned (thermal cycled & incubated)), reagent types, throughput, software compatibility, GUI ease of use, barcode scanning, ancillary services
- Error Rate (defining major & minor), how going to test stacker, pipettor, dispenser, barcode reader, tip wash.
- Measurement device, volumes, wet or dry dispense
- Safety – seismic restraint, ergonomics, hazards, “E” stop
- Integrated system test – method outline
- Operational testing
- Precision %CV, accuracy, reliability, ergonomic design



Approach to Automation Selection

- **Cost**
 - Expense warranted, efficient use of tax payer money
- **Delivery**
 - time, how, installation, training
- **Service**
 - field support, location of service engineers, response time, loaner parts, consumables supply



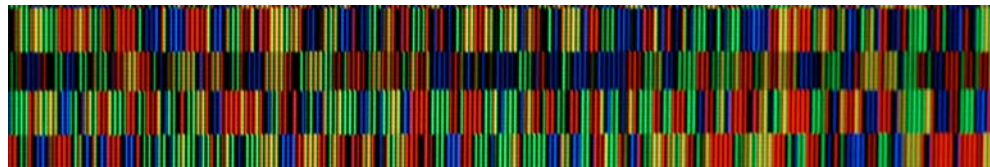
Approach to Automation Implementation

- **Technology Transfer**
 - New or upgrading instrument technology
 - Internal process improvements
 - New processes introduced to production from R&D
- **Performance**
 - Acceptance criteria
 - Operational testing
 - Duplicate partial batch
 - Single production batch
 - Statistical Analysis of Results
- **Training**
 - SOP written during development phase, working draft
 - Operator needs to know what to expect

Summary


- Brief history of the Joint Genome Institute
- DNA sequencing production line
- Increased throughput since 1999
- Instrumentation improvement highlights
- Our approach to successful selection & implementation

AGTCCGCGAATACAGGCTCGGT





More Information?



JGI
DOE JOINT GENOME INSTITUTE
US DEPARTMENT OF ENERGY
OFFICE OF SCIENCE

search jgi
contact us
site map
internal

[GO]

JGI brings the expertise of four national laboratories, [Lawrence Berkeley](#), [Lawrence Livermore](#), [Los Alamos](#), and [Oak Ridge](#), and the [Stanford Human Genome Center](#) to bear on the frontiers of genome sequencing and related biology. Our sequencing targets encompass a rapidly expanding range of microbes, animals, and plants. The new [Community Sequencing Program \(CSP\)](#) aims to broaden the range still further. JGI is operated by the [University of California](#) for the [U.S. Department of Energy](#).

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- JGI SCIENCE
- MEETINGS
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- EMPLOYMENT



[genomes]

[Microbial](#), [Eukaryotic](#)

[Integrated Microbial Genomes \(IMG\)](#)
[IMG w/ Microbiome Samples \(IMG/M\)](#)

[latest news]

[JGI Finishes 100th Microbial Genome](#)

[DOE BER call for bioenergy sequencing targets](#)

[sequencing]

This fiscal year : 23.407 billion base pairs sequenced
[More statistics](#)

[Sequencing for researchers](#)



More Information www.jgi.doe.gov

Acknowledgements

- **Martin Pollard, JGI Instrumentation Manager**
- **Susan Lucas, JGI Production Manager**
- **David Gilbert, JGI Public Relations Manager**



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