

Protein Information Management System (PIMS)

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...and the PIMS development team

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Outline of talk

- Background to PIMS
 - What is PIMS?
 - Why develop PIMS?
 - Why am I here?!?
- Basic concepts of PIMS
 - Targets and Constructs
 - Samples
 - Experiments and Protocols
 - Workflows
- PIMS at the OPPF
 - A short demonstration of tracking science
- Come and play for yourselves
 - PIMS on Oracle



What is PIMS?

- BBSRC SPoRT funded two consortia:
 - Scottish Structural Proteomics Facility (SSPF)
 - Membrane Protein Structure Initiative (MPSI)
- PIMS is funded to develop a laboratory information management system (LIMS):
 - Funded by the BBSRC SPoRT initiative
 - Funding Jan 2005 Dec 2009
 - Supports SSPF, MPSI, OPPF & YSBL
 - Developers in Daresbury, EBI, OPPF & YSBL
 - Support from OPPF, Dundee & Daresbury
- http://www.pims-lims.org/



PIMS development laboratories

PIMS PIs

Kim Henrick, Dave Stuart, Keith Wilson, Colin Nave,
 Jim Naismith, Neil Isaacs

PIMS supported

Anne Pajon, Ed Daniel, Marc Savitsky, Susy Griffiths

CCP4 supported

Chris Morris (project manager), Bill Lin

Other support

- Robert Esnouf (scientific sponsor), Jon Diprose
- Petr Troshin, (Jo van Niekerk)
- (Ian Berry, Gael Seroul, Diederick De Vries)



Why develop PIMS?

- Longstanding need for rational data management for protein production
 - Complex, ever-changing workflow
 - To exploit higher throughput
 - To aid collaboration and make data public
- Academic LIMS (and industrial?)
 - LISA, HalX, SESAME, MOLE, (Beehive)
 - Specific to one site, hard to maintain
- PIMS is a collaborative effort to find a common solution
 - Most laboratories have some similar processes
 - All have some unique processes
 - PIMS is fully featured LIMS, not target tracking



Some ancient history

- Starts with 2001 Airlie House agreement
 - To share protein production data
 - TargetDB is limited implementation
 - Detailed specification of terms
- European-based projects
 - eHTPX: data exchange models (dictionaries)
 - HAL & HALX: LIMS concentrating on workflow
 - MOLE: LIMS built on generic data model
 - SPINE encouraged collaboration
- Loose PIMS consortium formed to seek a common solution
 - BBSRC SPoRT provided funding opportunity



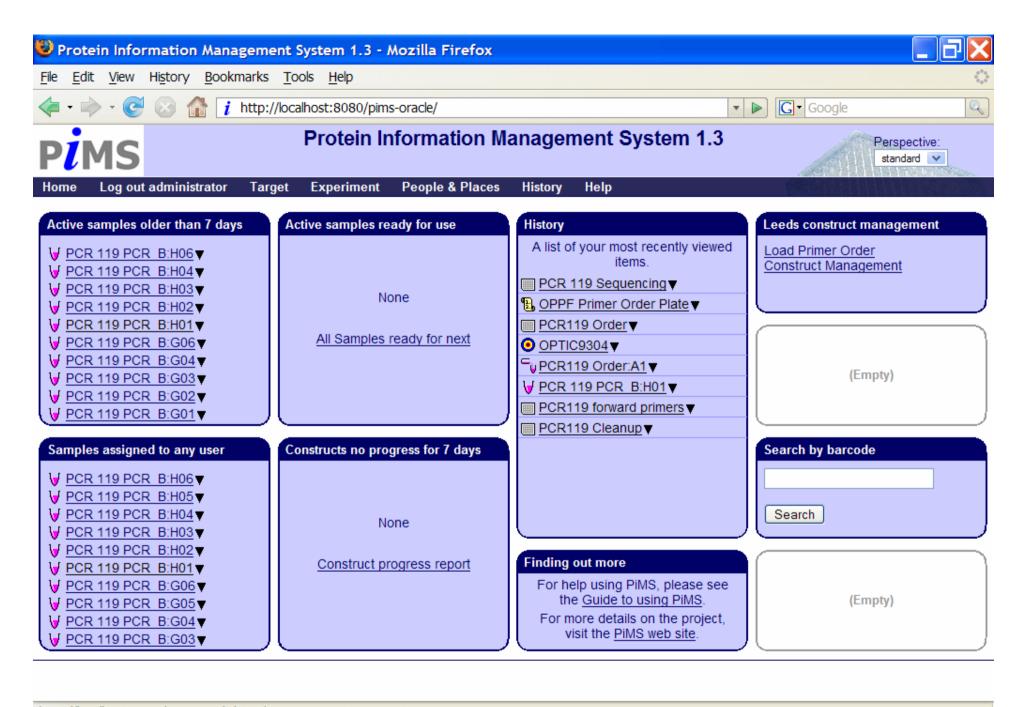
Technologies used

- PIMS is used from a web browser
 - Mozilla Firefox or Internet Explorer
 - No client software to install (perhaps plugins)
 - Windows, Macintosh and Linux clients
- PIMS requires a web and database server
 - Typically the same machine
 - Web server Apache Tomcat
 - Development on free PostgreSQL
 - Now available for Oracle
 - Windows and Linux servers
- Technologies used by developers
 - Java1.5, Hibernate, JUnit, BioJava, dot, batik, AJAX, ...



Why am I here?!?

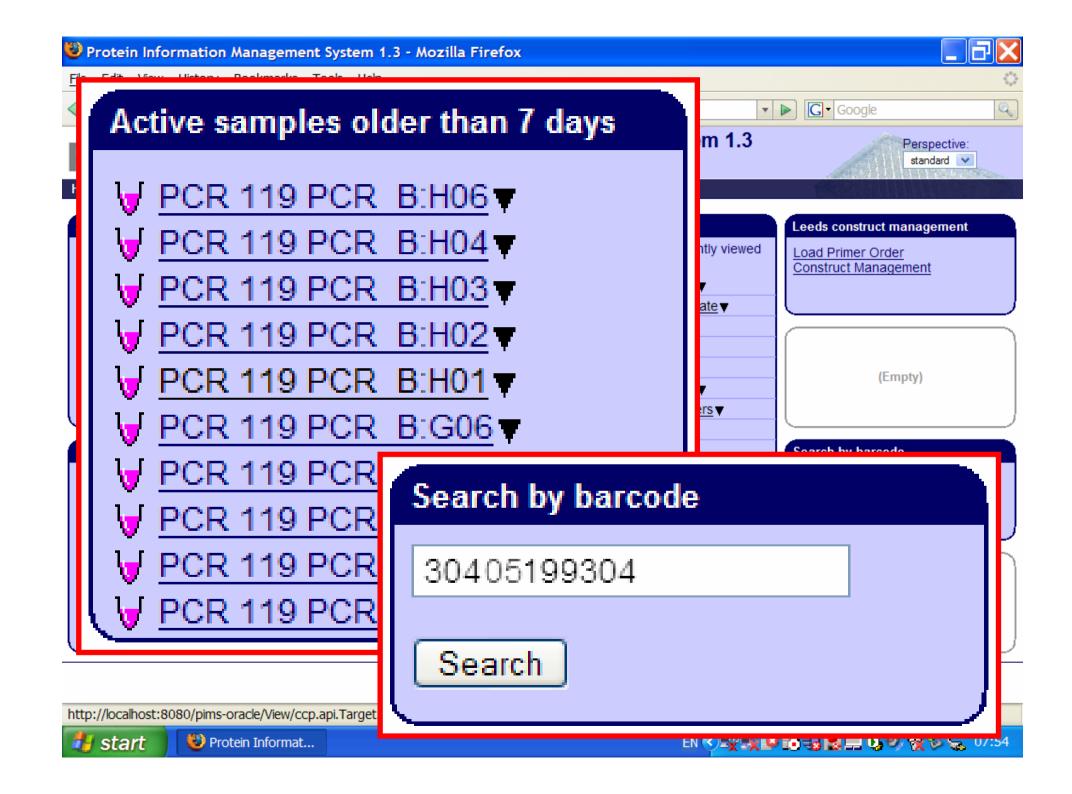
- PIMS is developing rapidly
 - Many problems overcome such as scalability and performance
 - Usability being addressed
- Academic LIMS developed to commercial standards
 - Multisite development makes it suitable for many sites
 - Industrial research laboratories not that different from academic ones
- Want to assess industrial interest in PIMS
 - Consortium-type agreements to fund continued development
 - Happy to offer on-site demonstrations

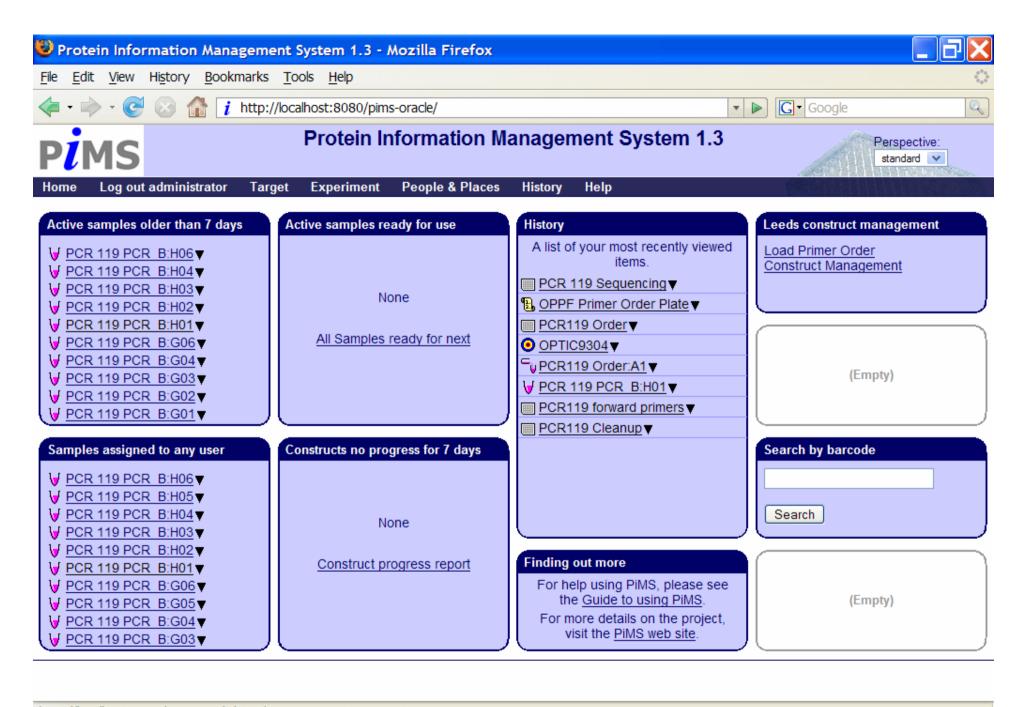








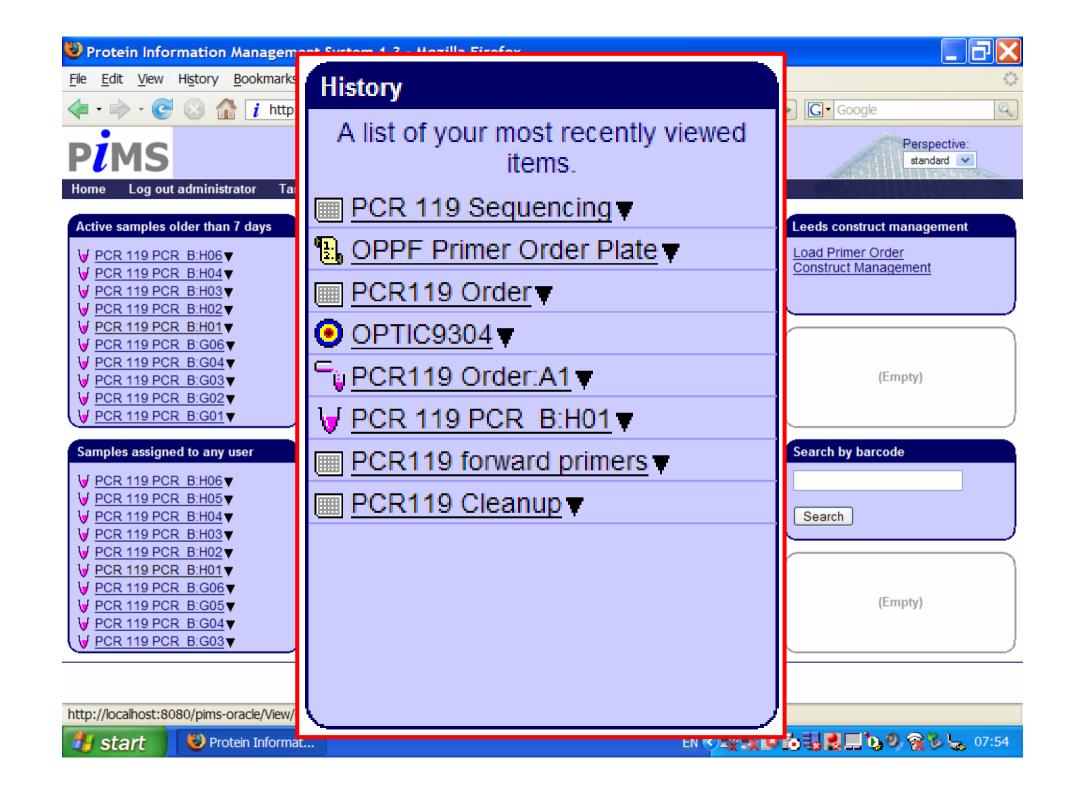


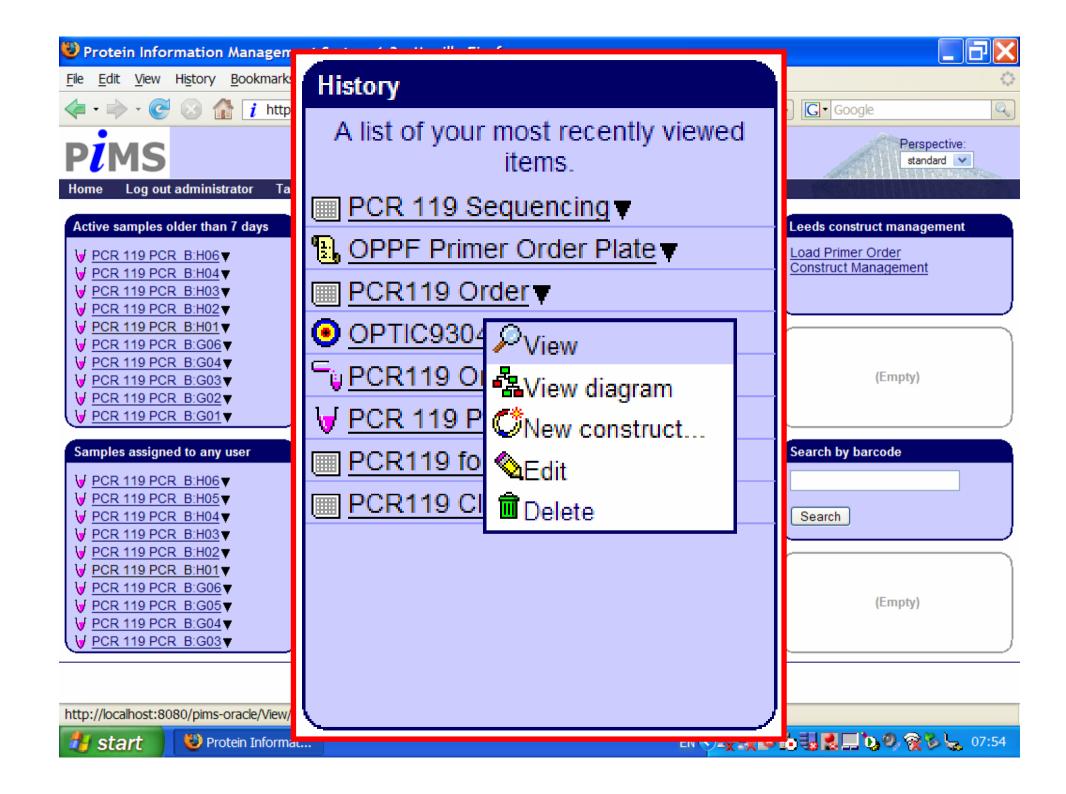














Basic concepts of PIMS

PIMS uses a few simple key concepts which can be linked together to model complex workflows

Targets

Description of sequences, store annotations

Constructs

Starting points for real experiments, link to targets

Samples

- Tracked samples made & used by experiments
- Samples have types, owners, locations etc.

Experiments

Take one (or more samples), produce new sample(s) as outputs



Experiments and protocols

A protocol is a reusable user-defined template describing what you record for your experiments.

Parameters

Numerical values, free text values, T/F. *E.g.* incubation temperature or the number of PCR cycles; details of incubation conditions; was reagent added?

Input Samples

Samples or reagents used when performing an experiment that you wish to track

Output Samples

 Samples or reagents produced when performing an experiment that you wish to track



OPPF TrialExpression

Perspective:

standard 🔻 Log out administrator Target Experiment People & Places Home Details of protocol: OPPF TrialExpression Name OPPF TrialExpression Remarks Standard OPPF plate-based trial expression Experiment type Small scale expression Design new protocol based on this Copy protocol Method ① Inputs Add new Parameters Add new: Number Text Yes/No ○ Output Samples Add new [™]Purified plasmid Name Purified plasmid 2.0uL Amount v Category Nucleic acid



Typing of PIMS items

Typing helps PIMS offer sensible choices: only a plasmid can be used for transfection experiments...

Samples

Typed to show what they are

Input/Output samples for protocols

State what type of sample can be used and what is produced

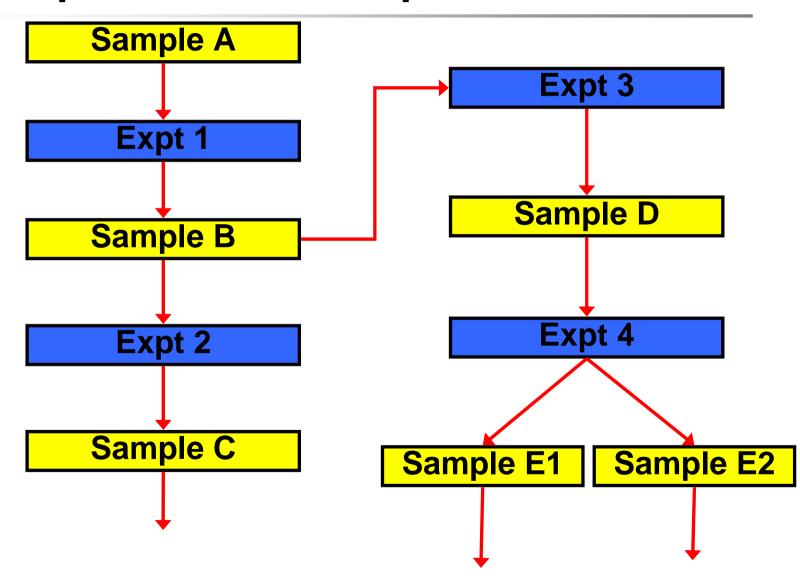
Experiments and protocols

 An experiment type is defined by its protocol. A protocol type links similar protocols together



Experiments & samples → Workflows







The PIMS holder (plate experiments)

A holder groups samples. This allows PIMS to perform plate experiments in groups

Samples

For plate experiments output samples of previous experiment are mapped to input samples of next. (Provided sample type matches!)

User interface for plate experiments

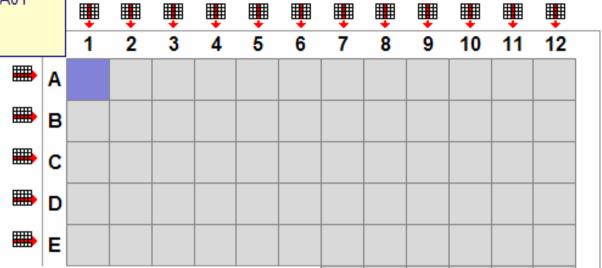
Gives graphical and spreadsheet views.
 Allows editing, reformatting and spreadsheet upload

Plate PCR119 Cleanup, well A1 Target OPPF3926

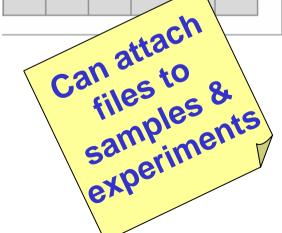
Kit: 90.0uL, Unspecified

PCR product: 50.0uL PCR119 PCR:A01

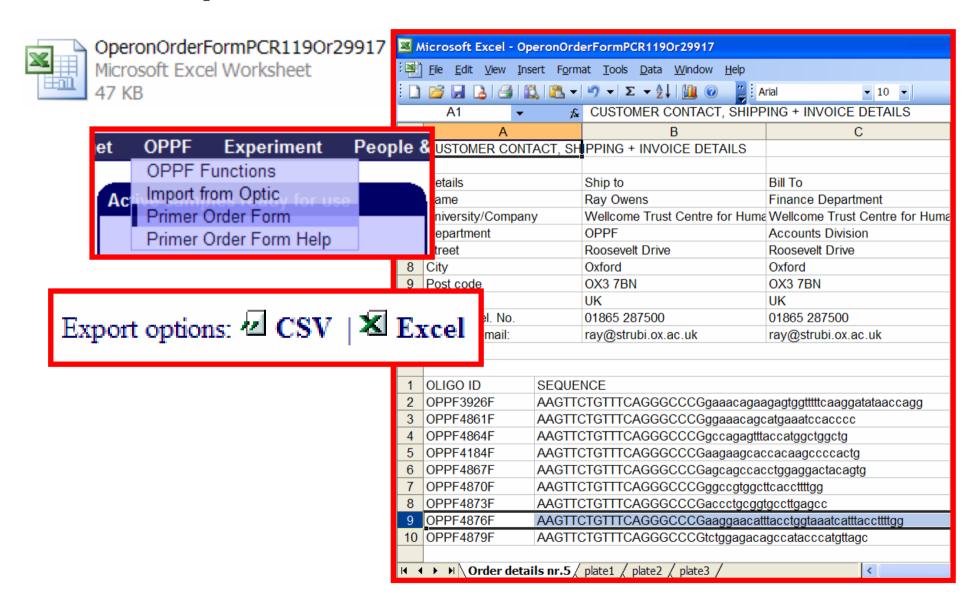
Checked on gel?: Yes Status: To be run



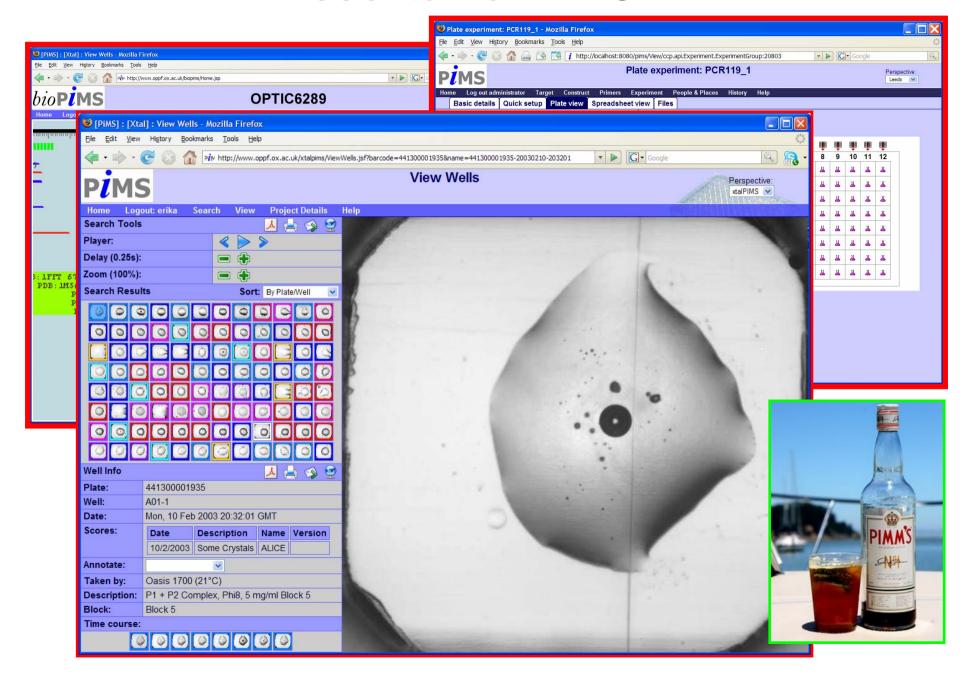
Basic deta	ils	Qui	ick setup	Plate view	Spreadsheet v	riew F	iles
Tray	Row	Col	Target	Status	Output	Kit	
DODAM	Α.	4	0000000	T. L	DODAM	Sample	
PCR119 Cleanup	Α	1	OPPF3926	To be run	PCR119 Cleanup:A01	(None)	50.0
PCR119 Cleanup	В	1	OPPF4861	To be run	PCR119 Cleanup:B01	(None)	90.0
PCR119 Cleanup	С	1	OPPF4864	To be run	PCR119 Cleanup:C01	(None)	90.0
PCR119 Cleanup	D	1	OPPF4184	To be run	PCR119 Cleanup:D01	(None)	90.0
PCR119 Cleanup	Е	1	OPPF4867	To be run	PCR119 Cleanup:E01	(None)	90.0
PCR119 Cleanup	F	1	OPPF4870	To be run	PCR119 Cleanup:F01	(None)	90.0
PCR119 Cleanup	G	1	OPPF4873	To be run	PCR119 Cleanup:G01	(None)	90.0



Experiments can read/write data



A cocktail of PIMS...



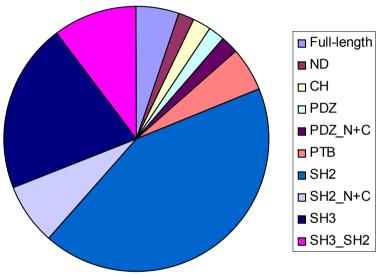


OMC Oxford Module Consortium

PCR107: OMC Plate 1

pOPINF (N-His) 96-well plate

94% cloned78% soluble expression72% OVERALL SUCCESS

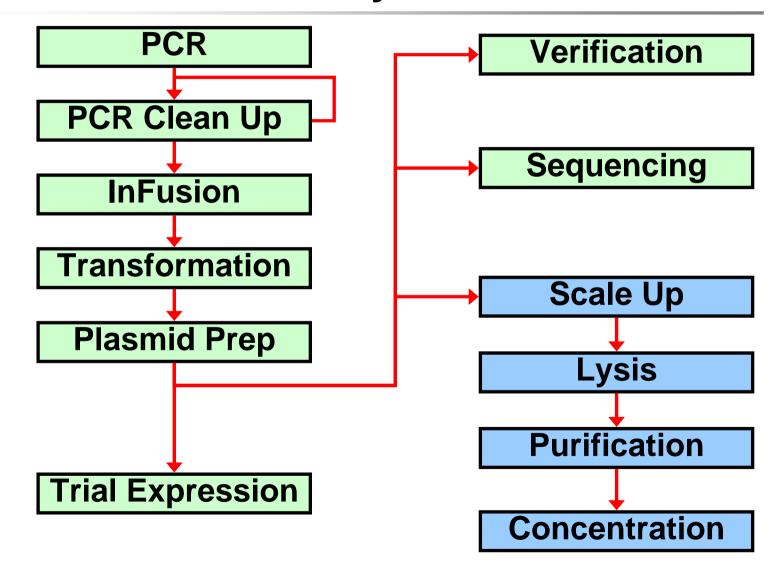


Now working on PCR119: OMC Plate 2 Real dataset used for demo



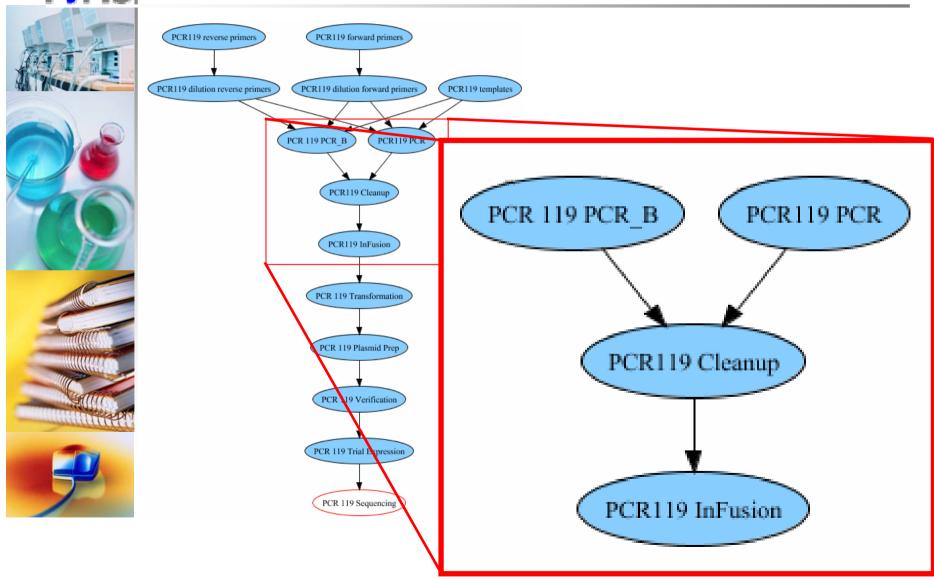
Protocols Used By PCR119







PCR119 workflow derived from PIMS





PIMS demonstration hardware

- Running on an average laptop
 - 1.7 GHz Pentium M Processor
 - 1GB memory
 - Windows XP SP2
- No external connection
 - Apache, Tomcat and Oracle all installed as services on this machine
 - Demonstrated using Mozilla Firefox
- Screen resolution limited to 1024x768
 - To suit projector



PIMS demonstration



- PIMS version 1.3
- Released last Friday (26/10/07)
- Running on Tomcat and Oracle
- Data for series of OPPF experiments (PCR119)