



Technology Transition Workshop | *Kristin S. Lowery, Ph.D.*

Ibis™ Assay Workflow Overview

Ibis™ Assay Workflow

- 1. Receipt and storage of assay kit components.**
- 2. Perform up-front sample processing.**
- 3. Register the experiment.**
- 4. Set up PCR plate(s).**
- 5. Seal the PCR plates.**
- 6. Thermocycle PCR plates.**
- 7. Prepare reagents for PLEX-ID™.**
- 8. Fill reagents on PLEX-ID™ and empty waste.**
- 9. Analyze the PCR plates on the PLEX-ID™.**
- 10. Review data.**

Ibis™ Assay Workflow – Step 1

- **Upon receipt, check for assay components**
 - **10 barcoded assay plates**
 - **1 bottle of magnetic beads**
 - **1 reservoir for magnetic beads**
 - **3 cleanup reagents (CR1, CR2 and CR3)**
 - **Instructions sheet**

Ibis™ Assay Workflow – Step 1

- **Storage: assay plates**
 - **Contain 35 μ L per well of PCR master mix**
 - **Primer pairs, modified dNTPs, PCR buffer, and enzyme**
 - **Each column is a separate sample**
 - **Eight wells comprised of three primer pairs per well**
 - **Profile is composite of all eight wells**
 - **User is required to add 5 μ L of template to each well of a sample**
 - **Store at -20°C – manual defrost freezer**

Ibis™ Assay Workflow – Step 1

- **Storage: magnetic bead bottle**
 - Store at 4° C and in upright position
 - DO NOT FREEZE
- **Storage: magnetic bead reservoir**
 - Store at room temperature
- **Storage: cleanup reagents**
 - Store at 4° C

Ibis™ Assay Workflow – Step 2

- **Up-front sample processing**
- **Extract DNA from samples (your choice of method)**
 - **Qiagen® columns**
 - **KingFisher® magnetic bead systems**
 - **Phenol/chloroform**
 - **Others**

Ibis™ Assay Workflow – Step 3

- **Register experiment**
 - **Import PCR barcodes**
 - **Use import wizard**
 - **Barcode file sent on CD with shipment**

Import Wizard Example

Use to
import
barcodes or
assay plans

Type of
import
selected
based on file

Import Assay Kit Configurations and Barcodes

Select Import File
Select a file to be imported below.

Assay Kit Barcodes
Imported barcodes will appear in the available barcode list for the assigned Assay Kit during the Sample Registration process.

Assay Kit Configuration
Imported Assay Kits configurations will appear in the Assay Kit selection field during the Sample Registration process.

File:
C:\Documents and Settings\klowery\Desktop\KitImport files\SERVICELAB_IBIS_KIT_20090105060354

Import Wizard Example

Barcodes to
be imported

Expiration
date

Barcode	Expiration Date
C05017850	11/9/2010
C05017851	11/9/2010
C05017852	11/9/2010
C05017853	11/9/2010
C05017854	11/9/2010
C05017855	11/9/2010
C05017856	11/9/2010
C05017857	11/9/2010

- Click **Import**
- “Importing has completed” message will display at bottom
- Click **Finish** to close wizard

Ibis™ Assay Workflow – Step 3

- **Register experiment (continued)**
 - For data analysis to work correctly, the plate must contain a positive control and a negative control
 - 10 samples can be run on the plate
 - Use different wizards depending on number of samples
 - Requires sample list – Microsoft® Excel spreadsheet
 - Can use tubes or plates for setup
 - Use control layouts
 - Define positions for negative and positive PCR controls
 - Do not need to put controls in sample list

Casework Wizard

Register a New Experiment

Assay Kit Properties
Assay Kit, Project and Experiment fields below are required.

Select an Assay Kit
CASEWORK FOR PCR BUFFER II TEST
Inventory: 59

Select a Project
FORENSIC-823

Select an Experiment
Casework registration for training

Enter a Comment

< Back Next > Cancel

Indicates number of barcodes available

- If “0” barcodes in inventory, import necessary assay barcodes before continuing

- Select an assay kit from the list
- Select a project from the list
- Select or enter an experiment title
- Click **Next**

Casework Wizard

Register a New Experiment

Select Sample Type
Choose a sample type from the options below.

Unquantified
Unquantified samples have not been characterized for the amount of genomic material present. No quantity is expected on input.

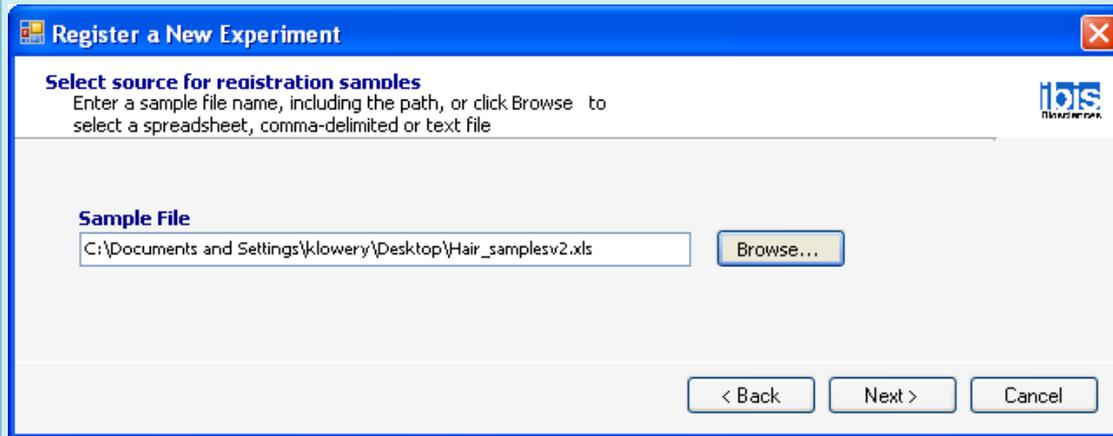
Quantified (Mass)
Samples quantified by mass indicate the amount of genomic material in nanograms. The amount is specified in the Quantity field on input and the value supplied is the mass in a single PCR well for the assay plate.

Quantified (Copies)
Samples quantified by copies specify the number of genomes or organisms present in the sample.. The amount is specified in the Quantity field on input and the number represents the copies present in a single PCR well for the assay plate.

< Back Next > Cancel

- Select sample type
 - Unquantified (enter in dilution factor (DF) for PCR plate)
 - Quantified (enter in picograms or copies for PCR plate)
- Click **Next**

Casework Wizard



	A	B	C
1	Sample	Quantity	
2	Hair.1	25	
3	Hair.2	25	
4	Hair.3	25	
5	Hair.4	25	
6	Hair.5	25	
7	Hair.6	25	
8	Hair.7	25	
9	Hair.8	25	
10	Hair.9	25	
11	Reagent.Blank	25	
12			

- Select file with sample information and click **Next**
- Sample file is validated
- Click **Next**

Casework Wizard

Select control layout

Automatically interleaves controls with samples based on control layout

Register a New Experiment

Control Layout
Choose a control layout below. The default layout is samples with no controls interleaved.

Control Layout
MITO STANDARD CONTROL

Plate	Position	Sample ID	Quantity	Unit	Type
1	1	HAIR.1	25	DF	Unquantified
1	2	HAIR.2	25	DF	Unquantified
1	3	HAIR.3	25	DF	Unquantified
1	4	HAIR.4	25	DF	Unquantified
1	5	HAIR.5	25	DF	Unquantified
1	6	HAIR.6	25	DF	Unquantified
1	7	HAIR.7	25	DF	Unquantified
1	8	HAIR.8	25	DF	Unquantified
1	9	HAIR.9	25	DF	Unquantified
1	10	REAGENT.BLANK	25	DF	Unquantified
1	11	PDB	1	DF	CTL
1	12	SC35495-4-POS	500	pg	POSCTL

< Back Next > Cancel

- Select control layout – predefined in a previous step
- Click **Next**

Casework Wizard

Number of barcodes required

Register a New Experiment

Selecting PCR Plate Bar Codes
For assay kit CASEWORK FOR PCR BUFFER II TEST

MITOTILING - PCR BUFFER II TEST → **1 PCR bar code required**

Available

BarCode
C00016820
C00016821
C00016822
C00016823
C00016824
C00016825
C00016826
C00016827
C00016828
C00016829
C00016830
.....

Assigned

Plate	BarCode
1	C00016819

< Back Next > Cancel

- Highlight barcodes and click **Left Arrow**
- Click **Next**

Casework Wizard

Register a New Experiment

Select Output File Paths
Default output file paths have been suggested. Enter a file path or use Browse to find a different location.

Specify Report Folder:
C:\Documents and Settings\klowery\Desktop **Browse...**

Worklist Option

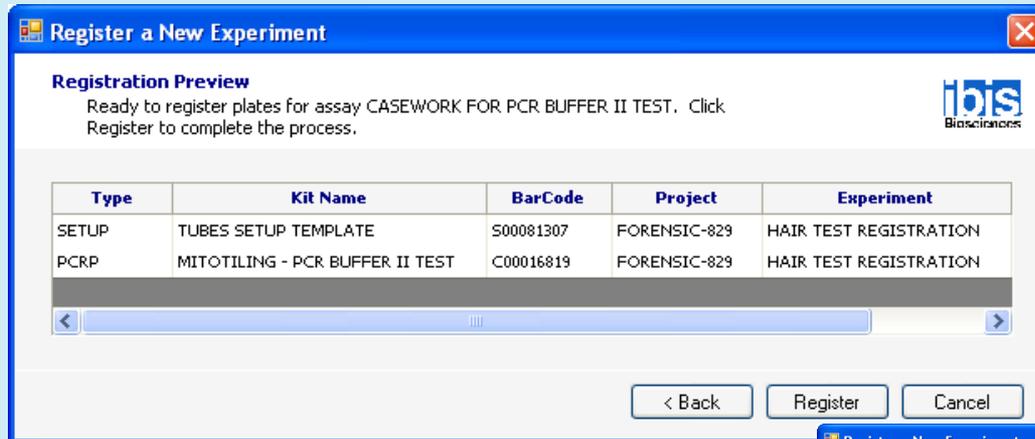
From Input Sample to Assay Plates (with Setup)
 From Input Sample to Assay Plates (without Setup)
 No Worklist

Specify Worklist Folder:
C:\Documents and Settings\klowery\Desktop **Browse...**

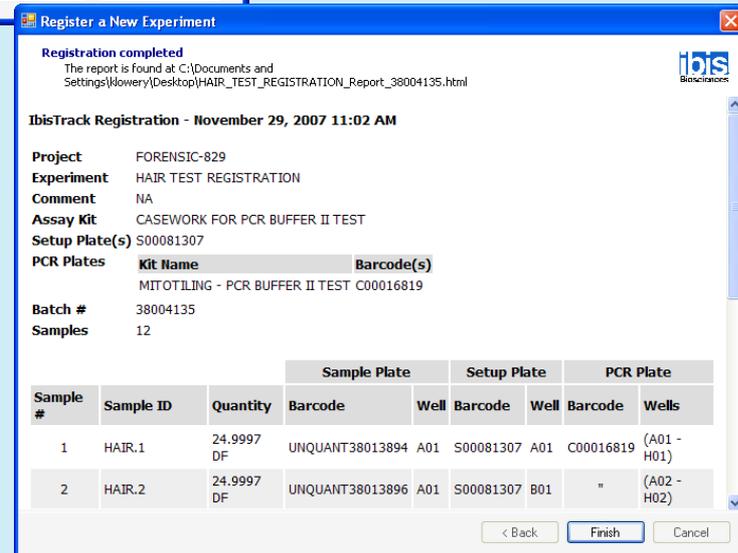
< Back Next > Cancel

- Select a folder for the report and worklist
- Select a worklist option
- Click **Next**

Casework Wizard



- Click **Register**
- Report generated with sample layout
- Go to file location to print out report and worklists



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Ibis™ Assay Workflow – Step 4

- Set up PCR plates either by hand or on reformatting robot



Ibis™ Assay Workflow – Step 4

- **Minimize contamination**
 - **Gowning procedure**
 - **Decontamination of PCR hood, pipettes and consumables**
- **5 μL of sample is added per well**
 - **A total of 50 μL is needed for each sample (40 μL required for sample + 10 μL for waste)**
 - **Use a new tip each time**

Ibis™ Assay Workflow – Step 4

- **Sample dilution**
 - For quantitated sample, dilute to 500 pg/5 µL with DNA-free water
 - For non-quantitated samples, dilute to set volume of 51 µL
 - For example, 17 µL stock + 34 µL of PDB

Ibis™ Assay Workflow – Step 4

- **Robotic setup**
 - **Barcode validation**
 - Ensures correct samples placed on plates
 - **Sort algorithm**
 - Reduces crossover contamination by ensuring that tips only pass on top of sealed wells



Ibis™ Assay Workflow – Step 5



Image courtesy of
<http://www.thermo.com/com/cda/product/detail/1,,10142764,00.html>

- **Heat seal plate**
- **Centrifuge**

Ibis™ Assay Workflow – Step 6



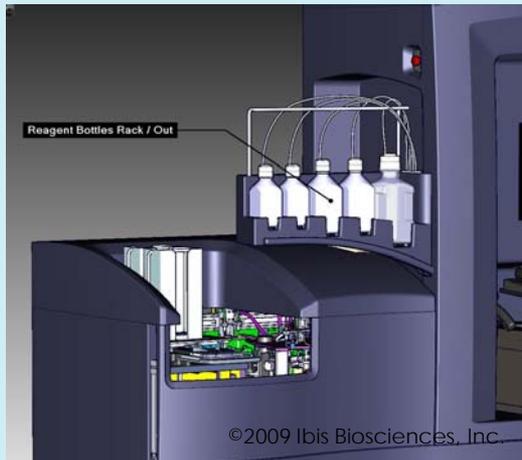
- Thermocycle the PCR plates
- Programs contain a 10 minute step at 99° C to minimize enzyme activity
- After thermocycling, centrifuge the plate for ~15 seconds at ~800 rpm
- Plate may be frozen until put on PLEX-ID™

Ibis™ Assay Workflow – Step 7

- **Prepare magnetic bead reservoir**
 - Dispense beads into reservoir
 - Seal with tape provided
- **Prepare clean-up reagents**
 - Add volume of methanol indicated on bottle
 - Use Burdick and Jackson™ HPLC grade
- **Recommend tracking lot numbers of reagents**

Ibis™ Assay Workflow – Steps 8 and 9

8



**Insert magnet
bead reservoir, fill
reagents and
empty waste**

9



**Run PCR plates on the
PLEX-ID™**

Ibis™ Assay Workflow – Step 10

View results
in IbisTrack

IbisTrack

File View Help Home Casework Databasing Import Analysis

Tasks

- Register Assays
 - Register Casework Plates
 - Register Database Population Samples
 - Register Service Lab Samples
 - Register Repeat Samples
 - Register Quality Control Plates
 - Generate Plate Setup Reports
- Manage Inventory
- View Plates
 - View Recent Plates
 - Find All Plates
- mDNA Analysis
 - Analyze mDNA
 - Import Mass Data
 - Add Database Items
 - Remove Database Items
 - Move Samples to Populations
 - Rebuild Unique Products
 - Build Amplified BC Database
- STR Analysis
 - Analyze STRs

Tasks

- Find Results
- Advanced Tasks
- In-House Tasks

Tasks-Analyze mDNA

Plate: P05010295 From: 13-Aug-2008 Analyze Generate report for P05010295

Sample: 12 To: 27-Sep-2008 New only

Register

- 00115-C
- 00012-C
- 00083-B

Comments for analysis report

Analysis Plate Profiles Database Mass Data Preferences Scenarios Monthly report Report archive Databasing plate setup

P05010295-00148-B (8) P05010295-P-31 (9) P05010295-P-39 (10) P05010295-PDB (11) P05010295-SC35495-10-POS (12)

P05010295, E12 P05010295, F12 P05010295, G12 P05010295, H12 P05010295-SC35495-10-POS composite

Filter ambiguous assignments Remove primers from profile

Spectrum viewing mode

Raw Deconvolved

P05010295-SC35495-10-POS composite

Well 12 (A12)
2906 + 2901 + 2892

Well 24 (B12)
2925 + 2891 + 2908

num	error	exp. mass	obs. i
2901:	15893..16012:	A47 G18 C25 T30	
2925:	15937..16041:	A35 G14 C24 T32	
2899:	15985..16073:	A25 G16 C21 T27	
2898:	16025..16119:	A25 G18 C27 T25	
2897:	16055..16155:	A31 G13 C29 T28	
2896:	16102..16224:	A44 G14 C41 T24	

Comments for analysis report

2.6.0510 MTDNA MTDNA KLOWERY

Workflow Timeline

- **Sample preparation**

- **Isolation: customers are free to use any method they choose**
 - **KingFisher®: 30 minutes setup plus 30 minutes run time (up to 96 samples); setup can be performed manually or on a Tecan Freedom EVO® liquid handler**
 - **Qiagen® columns**
 - **Phenol/chloroform**

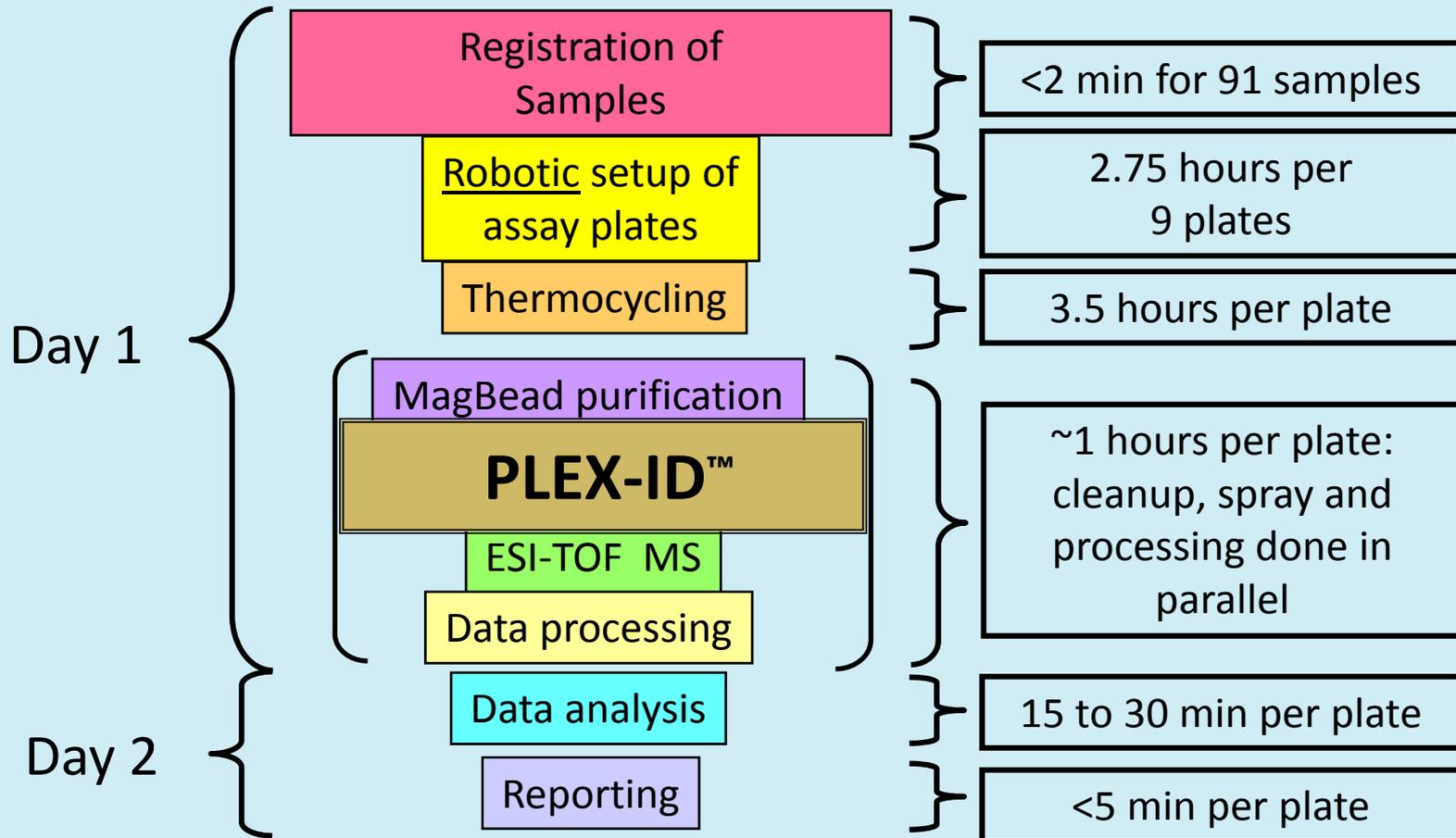
- **Plate setup**

- **Manual: 10 to 20 minutes per plate**
 - **Reproducibility is an issue**
- **Robotic: 15 minutes per plate**

Workflow Timeline

- **PCR**
 - PLEX-ID™ mtDNA Assay – 3.5 hours
 - PLEX-ID™ STRs Assay – ~3 hours
 - PLEX-ID™ SNPs Assay – 2.5 hours
- **PLEX-ID™**
 - Initial flushing and system startup: 20 minutes
 - Clean-up: 10 minutes for first well, then one well cleaned every minute
 - Spray on TOF: ~50 minutes per plate
 - 30 seconds per well – 4 minutes per eight well sample
 - Data processing: 15 to 20 minutes per plate

Workflow Timeline



Workflow Throughput

- **Assuming manual PCR setup and two cyclers, four plates a day**
 - 40 samples per day
 - 200 samples per week
 - 10,400 samples per year
- **Assuming robotic PCR setup and five cyclers, 10 plates a day**
 - 100 samples per day
 - 500 samples per week
 - 26,000 samples per year
- **Limiting factor – number of thermocyclers**

Questions?

Contact Information

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Note: All images are courtesy of Dr. Kristin S. Lowery unless otherwise noted.