

## SCIENTIFIC

#### **3500-Series instrument Overview**

Considerations for efficient implementation of the 3500 series genetic analyzer for Human Identification

HID University Seminar Series 2011

#### 3500 System Workflow and Feature Highlights





#### Dashboard



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#### **Redesigned Consumable Packaging**

- Pre-filled, quality-controlled reagents
- Radio Frequency Identification Tags
  - Fixed, passive tags
  - Limited read range
  - Memory only, cannot be read by the computer and cannot execute code
- Comparable per sample running cost to 31xx instruments





## Radio Frequency Identification (RFID)

- Three independent RFID readers track and record information on four primary consumables (with RFID labels)
- Consumables tracked
  - Polymer
  - Cathode Buffer
  - Anode Buffer
  - Array
- Information tracked
  - Lot numbers
  - Part numbers
  - Serial numbers
  - Dates (expiration and installation)
  - Capacity/Usage





#### Consumables: Polymer Pouch Volume





#### **RFID Tracking Limits for Consumables**

Consumable			On I	nstrument	RFID Tracking Limits		
		Shen Lhe	Lifetime <sup>1</sup>	Usage <sup>2</sup>	Warning	Hard Stop	
Polymer Pouch	960 Samples	) Expiry Date 7 D		960 Samples 50 Injections (24 cap) 120 Injections (8 cap)	7 Days	Expiry Date # Samples, Injections	
	384 Samples	Expiry Date	7 Days	384 Samples 20 Injections (24 cap) 60 Injections (8 cap)	7 Days	Expiry Date # Samples, Injections	
Array		Expiry Date	160 Injections	160 Injections	Expiry Date 160 Injections	None	
Buffer		Expiry Date	7 Days	50 Injections (24 cap) 120 Injections (8 cap)	None	Expiry Date On Instrument Lifetime	
Condition	ing Reagent	Expiry Date	24 Hours	Single Use	24 Hours	Expiry Date	

<sup>2</sup>On-instrument life/sample usage limit is based on whichever limit is reached first <sup>1</sup>Polymer validated for 7 days on instrument and array guaranteed for 160 injections

Reagent and consumable lifetime guarantees/ recommendations are designed to promote the production of high quality data, for all applications, in whichever laboratory

environment the instrument is located



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iii Calibrate	💽 Day		• •	Jar	nuary 2009		_
Spatial		Tuesday	Wednesday	Thursday	Friday	Saturday	Sur
Spectral		20	94	1	2	3	4
灯 Performance Check							
Sequencing Install Standard							
Fragment Install Standard		6	Perform Planned Ma	8	9	10 Clean Drip Tray	1
HID Install Standard			R Wash Pump Trap				
🍗 🦘 Maintenance Wizards			FR Defragment Hard Dr				
🖄 Planned Maintenance		13	14	15	16	17 R prepare bidi	1
Notifications Log							
Service Log							
Schedul		20	21	22	23	24	2
Return to Setup			FR Restart PC, Instrum				
Ya/							

FR or F = Applied Biosystems recommended tasks, user editable - specify dates, priorityR = User-specified repeating tasks



#### Calibration and Maintenance Reports and Export Files



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#### QC Analysis: QC Protocol

Edit QC Protoco	l GS_LS(8	1-400)+Norma	alization					9
Setup a QC Prot	tocol							
* Protocol Name:	G5_L5(80	400)+Normaliza	tion					
Description:	G5600L1Z	Default analysis	s Range set to Full					
Size Standard:	G5600_L12	+Normalization,	(80-400) 💌					
Sizecaller:	SzeCaller	v1.1.0 ¥						
Analysis Settings	QC Setting	5						
								0
Analysis Range:	Pul	*	Sizing Range:	Part	ial 💌	50	Calling Method:	Local Southern 💌
Analysis Start Po	oint: 0		Sizing Start Siz	e: 80				
Analysis Stop Po	int: 1000	000	Sizing Stop Size	400				
[								
		😥 Blue	👿 Green	E Y	ellow F	Z Red	Purple	😥 Orange
Peak Amplitude	Threshold	50	50	F	50	50	175	50
Common Settin								
	-		Use Sr	oothing	Linht Y			
		Use Baselir	ing (Baseline Windo	w (Pts))	R SI	-		
			Minimum Peak Ha	# width	2	-		
			Peak With	inu Sine	10	-		
			Polynomial	Deves	2	-		
			Sive Threshold De		0.0	_		
			Size Theodold P	aak Fod	0.0	_		
			Juge meanad P	ear cha	0.0			
L								
Close								Save

- QC protocol is the required primary analysis protocol for HID applications
- It defines peak detection, sizing, and quality values
- Factory-provided QC protocols are available in the software or can be created by the user

G5\_LS(80-400)+Normalization



#### QC Analysis: Result Review

🚹 Review Results

#### View Sequencing Results

View Fragment/HID Results

#### **Review QC Flag Details – Set Up Reinjections**

	Sample Name	Sample Type	Size Standard	Assay Name	PA Protocol	SA Protocol	Offscale	Broad Peak	Normalization Limit	Sizing Quality 1	Sample
1	1	Sample 🗾 💌	HID_GS600_LIZ	IF_POP4	G5_LS(80-400)			_		<b>~</b>	
2	9	Sample 💌	HID_GS600_LIZ	IF_POP4	G5_LS(80-400)					✓	
3	17	Sample 💌	HID_GS600_LIZ	IF_POP4	G5_LS(80-400)					✓	
4	2	Sample 💌	HID_GS600_LIZ	IF_POP4	G5_LS(80-400)					✓	
5	10	Sample 💌	HID_GS600_LIZ	IF_POP4	G5_LS(80-400)					✓	
6	18	Sample 💌	HID_GS600_LIZ	IF_POP4	G5_LS(80-400)					✓	
<b>1</b>			UTD COCCO 173	75 DOD4	CE 1 (00 400)			<b>—</b>			



Thermo Fisher

- Factory Standardization: Hardware-based calibration for more consistent instrument to instrument performance
- Internal Standard (IS) Normalization: Chemistry and Software based method for more consistent signal across capillaries, injections and instruments
  - User enabled
  - Utilizes re-designed GeneScan<sup>™</sup>600 LIZ Size Standard
  - Sample level peak heights are scaled relative to the intensity of the coinjected size standard compared to an optimized average size standard peak height (Normalization Target)



#### **Signal Normalization Tools**

- Forensic laboratories with multiple instruments have reported that signal variation between instruments can impact data interpretation
- Improved factory standardization and userselectable internal standard normalization minimize signal variation between instruments and from injection to injection







Instrument to Instrument Peak Height Consistency

Injection to Injection Peak Height Consistency



#### **Data Scaling Considerations**



- Peak heights on the 3500 ~3-4 x greater than 31XX platforms due to differences in data scaling
- Higher peak amplitude threshold required for 3500 analysis if using a static threshold
  - Use of the Global Cut-off function in the analysis method may allow for fine-tuning of analysis thresholds

#### Comparison of 3130xl and 3500xL Data



Identification of appropriate analysis thresholds promotes consistency of analysis outcome

#### PAT Definition for 3500 Data

PAT: 50rfu

PAT: 175rfu



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1000

#### PAT Definition for 3500 Data

- Peak Amplitude Threshold
  - Minimum suggestion of [Mean Baseline Noise] + [3-10 x SD]
    - Default Data Collection value based on G5 Dye Set noise evaluation is ~175 RFU (independent tests recommended on individual instruments and workflows)
      - Dependent upon multiple factors including choice of chemistry, injection conditions, injection recipe, instrument platform
  - Other considerations
    - Artifacts
    - Spectral pull-up
    - Stutter
- Stochastic Threshold
  - Determine level of confidence for homozygote calls

Consider differential thresholds for casework and database applications

#### Security, Audit and E-Sig

Settings Resources	🗮 Enable System Security 🛛 😭 🖓 Enable System Security	
Applied Biosystems	Account Setup	
💥 Manage Reports		
Audit Reports	User Names	User Passwords
E-Signature Reports	The length of user names must be between 8 and 32 characters.	The length of user passwords must be between 8 and 32
💥 Manage Users	Define name spacing	Define password spacing
Users	✓ Leading ✓ Trailing ✓ Consecutive	✓ Leading ✓ Trailing ✓ Consecutive
Kanage Settings	Define name characteristics	Define password characteristics
Security	🗹 Alpha 🕑 Numeric 🐨 Uppercase 🐨 Lowercase 🔲 Special	1 Alpha 1 Numeric 1 Uppercase 0
Audit		
E-Signature		User may not reuse the previous 5 passwords.
📄 Import		
🔛 Export		
Return to Setup	Security Policies	
Manage Settings Security Audit E-Signature Import Export Return to Setup	Define name characteristics          Image: Construction of the second	Define password characteristics           1         Alpha         1         Numeric         1         Uppercase           User may not reuse the previous         5         passwords.

- User Configurable according to laboratory's needs
- Security controls user access to system functions and data
- Auditing tracks changes and provides history reports
- E-signature requires user-authentication before changes are saved



#### System Preferences





### GeneMapper<sup>™</sup> *ID-X* Software v1.2 and higher

- Provides comprehensive support for files generated by the 3500 Series Genetic Analyzers
  - Analyzes both .fsa and .hid 3500 files
    - Both file types cannot be analyzed with earlier versions of GeneMapper ID-X Software
  - Records RFID information regarding consumable lot numbers, usage, and expiration dates
  - Records instrument status and electrophoresis and analysis parameters
  - Supports signal normalization
  - Compatible with Windows® XP and Windows® Vista operating systems
- Includes printing patch update for increased speed
  - 30-180% reduction of printing time vs. version 1.1
- Can be updated to v1.2.1 using the CODIS Export Patch
  - Enables the user to configure the list of markers that can be included in CODIS Export tables
- Continues to support all existing sample QC and data chain of custody features to facilitate compliance with ISO17025

# GeneMapper<sup>™</sup> *ID-X* Software Sample Info Tab

	Info Raw Data EPT Data	
🖻 🧀 Inj 2 2009-0	Stop Slope Threshold Sa	Sample Information
1_A01_01R	Size Calling Method	omplo Filo . 1 101 01Dup 2000-01-10 10 22-24-212
1_A02_02R	- 20	ampie file : 1_A01_01Rui 2009-01-19-10-23-34-312.
-1_A03_03R	Sal Sal	ampie Name : 1
- 1_B01_04R	Kun Information Sa	ample Origin Path : C:\Applied Biosystems\3500\Data\Run
1_802_05R	User Name : Admin St.	tatus Message : Size standard <gs600_liz_normalizati< th=""></gs600_liz_normalizati<>
1_B03_06R	Instrument ID : 3500 Fi.	ile Source : Disk media
1_C01_07R	Instrument Type : CE 3. Re-	e-Injection : NA
	Data Collection Ver : 3500	eserr Nerre : HTD36 DODA
1_C03_09R	Run Date & Time : 2009	ssay Name . Hibbo_for4
1_D01_10R	Run Duration : 1 min AS:	Normalization
1_D02_11R	Total Data Points : 6017 No:	ormalization Factor : 0.803
- 1 E01 13R		Factor Value
	Data Collection Settings	)C Information
1_E03_15R	Module File	: HID36 POP4x1
-1_F01_16R	Pre-Run Voltage	: 15 kVolts
	Pre-Run Time	: 180 sec.
1_F03_18R	Run Voltage	: 16400
	Injection Voltage	: 1200
1_GUZ_ZUR	Injection Duration	: 5
1 H01 22P	Temperature	: 60
1 H02 23R	Laser Power	: 20
1 H03 24R	Run Module Name	: HID36_POP4x1
	Instrument/ Run Protocol	: HID36_P0P4_G5_standard_5sec CONSUMAble
	Dye Set Name	: <sup>G5</sup> information
	Polymer Type	: POP4
	Polymer Lot Number	: 51A007 tracked via REID
	Polymer Expiration Date	: 2009-08-30 23:59:00.0
	Polymer Time on Instrument	: 2008-01-12 08:34:00.0
	Results Group Name	: JB_2009-01-19-11-03-57
	Anode Buffer Type	: AB 3xxx Buffer
	Anode Buffer Lot Number	: 51B007
	Anode Buffer Expiration Date	: 2009-08-30 23:59:00.0



#### Consumable Cost Comparison

Osarsakla	Instrument Platform								
Consumable	310	3130	3500	3130 <i>xl</i>	3500xL	3730			
Capillary/Array	\$0.78	\$1.26	\$0.96	\$0.59	\$0.43	\$0.25			
Polymer	\$0.52	\$0.50	\$0.48	\$0.50	\$0.48	\$0.44			
GA/Anode Buffer	\$0.07	\$0.07	\$0.03	\$0.07	\$0.02	\$0.01			
Cathode Buffer			\$0.04		\$0.03				
Plates/Tubes	\$0.10	\$0.06	\$0.06	\$0.06	\$0.06	\$0.06			
Septa	\$0.27	\$0.18	\$0.18	\$0.18	\$0.18	\$0.18			
Total Cost/ Sample	\$1.74	\$2.07	\$1.74	\$1.40	\$1.20	\$0.95			

• The cost/sample of 3500 consumables are comparable to the 3130 series

- Quality control of reagents coupled with pre-packaging saves operator time, eliminates sources of human error and promotes data optimum data quality
- The RFID tags automatically record lot numbers, part numbers, instrument serial numbers, reagent expiration and installation dates, and Capacity/Usage

- The 3500 Series Genetic Analyzers incorporate new hardware and software features designed specifically to support and enhance the Human Identification workflow
- Pre-packaged, quality controlled reagents deliver the same or improved on-instrument lifetime compared to previous platforms and help to maximize data quality
- RFID labelling of key consumables allow the user to monitor consumable status and all part and lot numbers are automatically recorded and transferred to the .fsa/.hid files
- GeneMapper<sup>™</sup> ID-X Software v1.2 and higher supports analysis of the .fsa/.hid files including signal normalization
- The 3500 Genetic Analyzers represent a strong collaboration between Thermo Fisher Scientific and forensic laboratories worldwide



#### THANK YOU!!

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