

Batch Processing increasing lab effectiveness



Robin Freeman, M.S. Harris County Medical Examiner's Office



BATCH METHOD

As businesses (laboratories) grow and demand increases, the work flow process is often changed to a "batch method". Batch methods require that a group of items move through a process together, a stage at a time.



Advantages

The batch method can be an advantage when administered appropriately. It is cheaper to complete a number of items in one go because instruments can be used more effectively, and the staff can specialize in that task.

There are advantages of employees concentrating their skills.

- They become more expert at their tasks, which will in turn increase productivity (output per employee).
- Employees are more familiar with processes and so can find ways of improving them.
- ➤ Builds team work among sections and staff meetings become more interactive.



Disadvantages

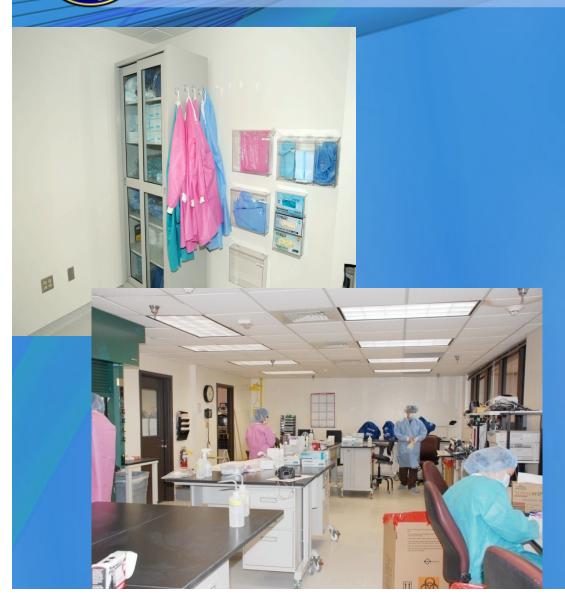
Batch methods can result in the build up of significant "work in progress" (i.e. completed batches waiting for their turn to be worked on in the next operation).

Managers must monitor workflow and foresee bottlenecks in the process. The batch method is not a hands-off process. Batching requires very careful planning to implement in the beginning.

Quality checks/verification steps must be put in place to monitor work product.



Preventative measures.



- Gowning required prior to entering an evidence exam area
- ➤ Monthly Swipe test
- ➤ Separate sectional Quality group
- ➤ In-House databases



Evidence Examination



Items are individually examined.

Areas of interest are identified, tested and documented.

- Presumptive tests
- Confirmatory tests

Samples are prepared for DNA testing and placed on appropriate extraction log.





DNA extraction

nalyst			Mi Co	icrocon Ver	ibe Verificat	n:	
extraction batch ID:						Hood #:_	
# CASE # & ITEM DESCRIPTION	Substrate COC From/To Date	Substrate COC From/To Date	MICROCON. Y/N	VOLUME in ul	AMT USED QUANT	AMT USED	AMT REMAINING
Verified by (date/initial)							
Verified by (date/mitial)							
			- 2				
Perified by (date/initial)		EVILLE					
				71-5	27-12		
Venified by (date/initial)		TP A					
				E. 1			10-4
Ventiled by (date/initial)				P P ST			
			-4-0		P 61,		100
Verified by (date/initial)				-			
vermed by (take nottal)							
Verified by (date/initial).					-		
Verified by (date/initial).							
Verified by (date/initial)							
		Je 1					
Verified by (date/mitial)					1		
147-1-1							
Verified by (date/initial)							
				Mark Y	and the		
Verified by (date/initial) Reagent Blank:	N/A	N/A					
Reagent Stank.	IN/A	19/24					
Extract storage: Box Q:	H W W		Note	s:			
	o K/exp						
	i cap						
						B07.4005-400	



Extraction logs

Differentials

Low yield

Regular

Knowns



Multi-Step DNA Test Process



Purification
Quantization
Amplification







Multi-Step DNA Test Process

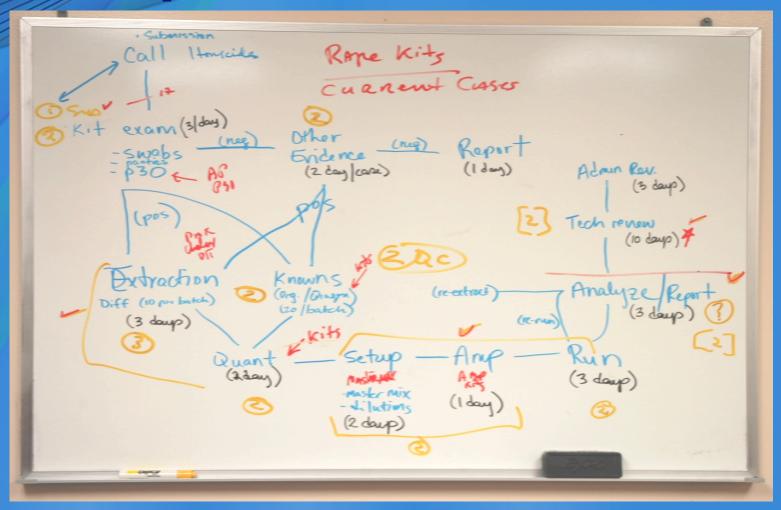


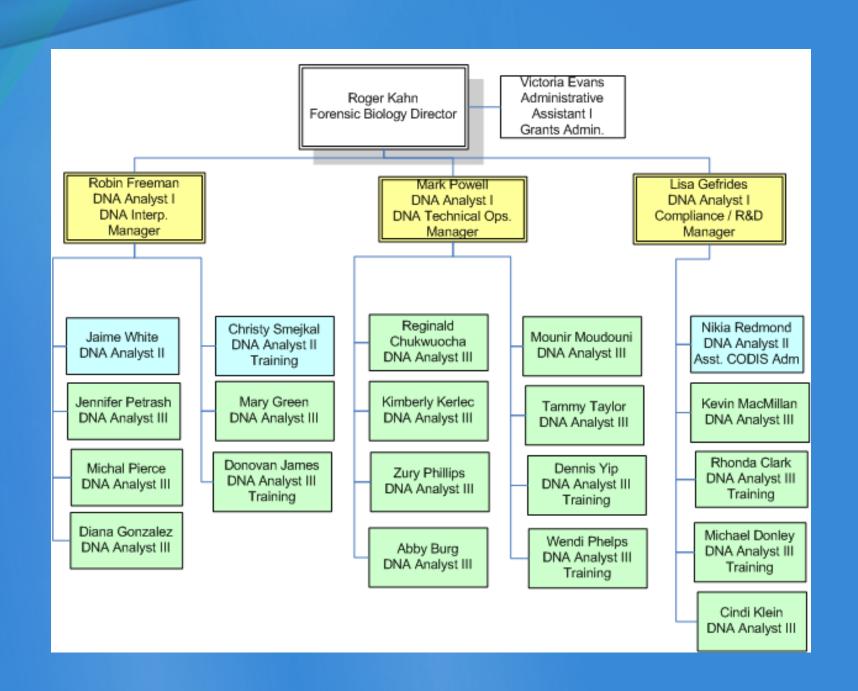
Analysis
Interpretation
Reporting

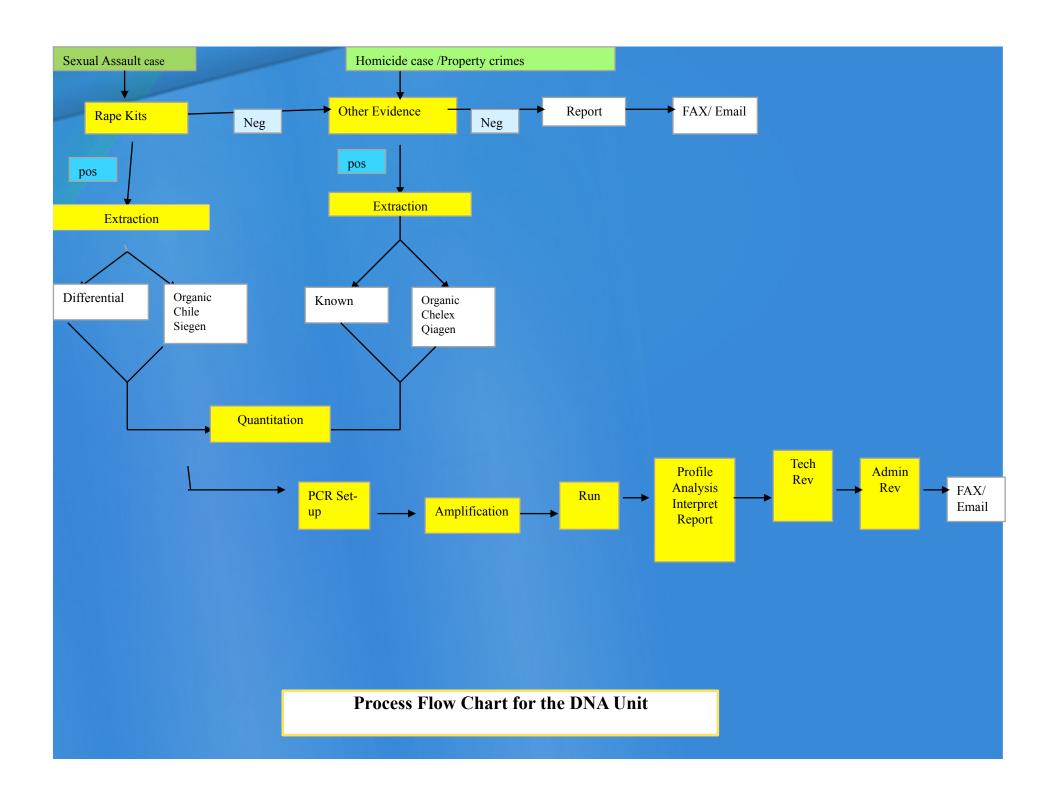




Planning and Implementation

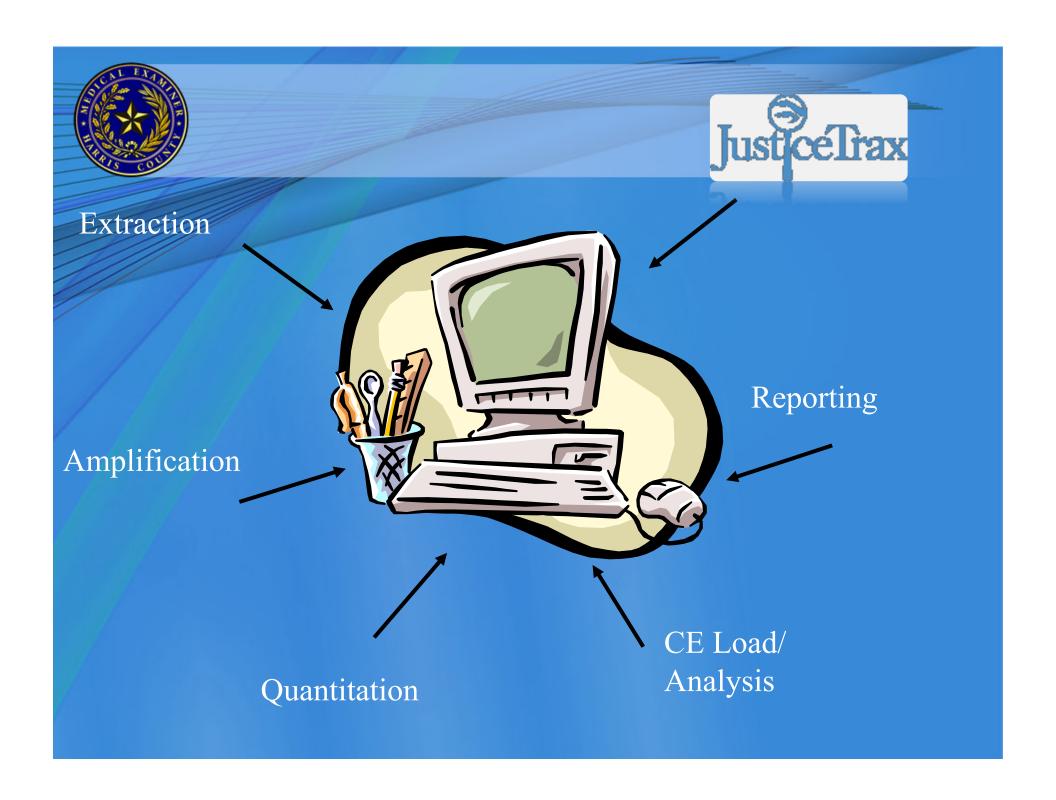






Organization of staff using Rotation schedules

SCHEDULE: Mon- Fri: 7-3:30 ART/DDH/LGS/ ACS/KDW LIG/MAD/DOJ (7-3:30) or (8-5:00) AT-TB	Lab Lock	Kits (1) Assigned Projects	Homicide (1) Burglary Property Cut Only (1)	Kit – (Other Evidence) (1)	Other Evidence 3 Minimum Overtime	Case Management	Ext.	Robot	Verificatio ns	Manual PCR/ Run Analyze	Report Writing Review
August 24	4 th fl. LGS LIG 5 th fl. LG	ACS LIG Bleach ART 5:00 Rotation LGS	DDH_ LGS MAD_ JBS 3:30-5	DOJ AT TB	ART KDW	JBS/SP Bleach/ Autoclave: Robot	ZP DCY	WP AB Float	Run: RF Paper: CS/DG/ KM	RUC	RPT =MLP// MG TR=RF/JP/JW AR =MP/ CS/ DG/KM
August 25	4 th fl. LGS LIG 5 th fl. LG	ACS Bleach ART 5:00 Rotation LGS	DDH_ LGS MAD_ JBS 3:30-5	DOJ AT TB	ART KDW LIG	JBS/SP Bleach/ Autoclave: Robot	WP TET	DCY ZP Float AB Float	Run: RF Paper: CS/DG/ KM	RUC	RPT =MLP// MG TR=RF/JP/JW AR =MP/ CS/ DG/KM
August 26	4 th fl. LGS LIG 5 th fl. LG	ACS Bleach ART 5:00 Rotation LGS	DDH_ LGS MAD_ JBS 3:30-5	DOJ AT TB	ART KDW LIG	JBS/SP Bleach/ Autoclave: Robot	WP TET	DCY ZP Float AB Float	Run: RF Paper: CS/DG/ KM	RUC	TR=RF/JP/ JWMLP/MG AR =MP/ CS/ DG/KM
August 27	4 th fl. LGS LIG 5 ^h fl. LG	ACS Bleach ART 5:00 Rotation LGS	DDH_ LGS MAD_ JBS 3:30-5	DOJ AT TB	ART KDW LIG	JBS/SP Bleach/ Autoclave: Robot	WP TET	DCY ZP Float AB Float	Run: RF Paper: CS/DG/ KM	RUC	TR=RF/JP/ JWMLP/MG AR =MP/ CS/ DG/KM
August 28	4 th fl. LGS LIG 5 ^h fl. LG	ACS Bleach ART 5:00 Rotation LGS	DDH LGS MAD JBS 3:30-5	DOJ AT TB	ART KDW LIG	JBS/SP Bleach/ Autoclave: Robot	WP TET	DCY ZP Float AB Float	Run: RF Paper: CS/DG/ KM	RUC	TR=RF/JP/ JWMLP/MG AR =MP/ CS/ DG/KM

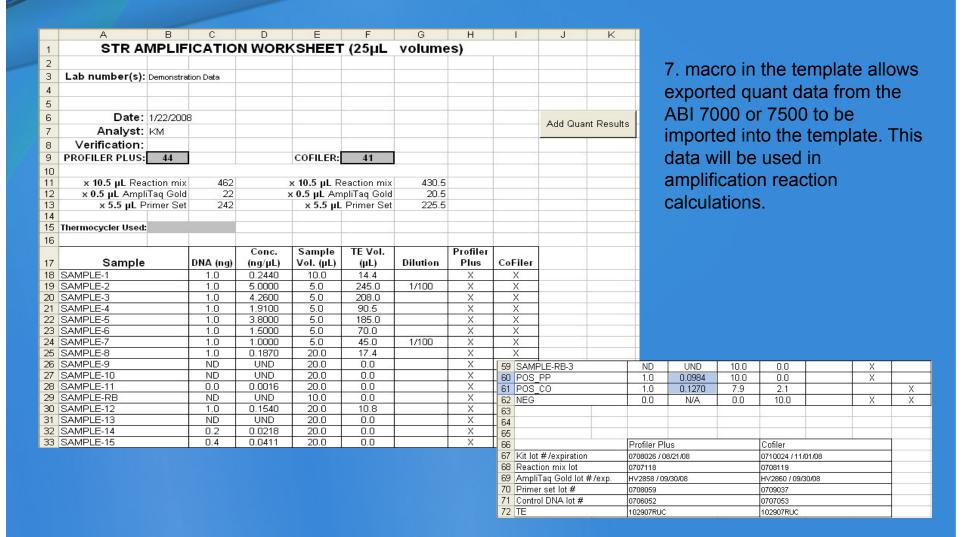


8	Α	В	С	D	E	F	G	Н	1	J	K	L		М					
	Lab	Numbers:	Demonstration	Data							Lot #/expi	ration d	ate '	Volume					
2			101							0 .:0 .0:									
3	V.	Analyst: erification:						D	CD Desetie	Quantifiler Kit n Mix (12.5ul)				852.5					
5	Ve	System:			+					er Mix (12.5ul)			-	716.1					
6			1/22/2008						1 (111)	Standards				1 10.1					
7		Reactions:	: 60							TE									
8		1	2	3	4	5	6	7	8	9	10	11		12					
9	Α	50	NEG	SAMPLE-8	SAMPLE-15	SAMPLE-RB- 2	SAMPLE-30	SAMPLE-38						50					
10	В	16.7	SAMPLE-1	SAMPLE-9	SAMPLE-16	SAMPLE-23	SAMPLE-31	SAMPLE-39						16.7					
11	С	5.56	SAMPLE-2	SAMPLE-10	SAMPLE-17	SAMPLE-24	SAMPLE-32	SAMPLE-RB- 3						5.56					
12	D	1.85	SAMPLE-3	SAMPLE-11	SAMPLE-18	SAMPLE-25	SAMPLE-33							1 oz A		В	С	D	Е
13	Е	0.62	SAMPLE-4	SAMPLE-RB	SAMPLE-19	SAMPLE-26	SAMPLE-34					16							
																	Conc		
	F	0.21	SAMPLE-5	SAMPLE-12	SAMPLE-20	SAMPLE-27	SAMPLE-35					17			nple		Conc. (ng/µL)	Dilution	RBs
14	F	0.21	SAMPLE-5 SAMPLE-6			SAMPLE-27 SAMPLE-28						18 S		LE-1	mple		555 1977 20	Dilution 1/100	RBs
14	F G		200 100 000 000 000	SAMPLE-13	SAMPLE-21	Manager Co. At the St. of St.	SAMPLE-36					18 S 19 S 20 S	AMP AMP	LE-1 LE-2 LE-3	mple		(ng/μL) 0.2440 500.0000 4.2600		RBs
14 15 16	F G H	0.068	SAMPLE-6 SAMPLE-7	SAMPLE-13	SAMPLE-21	SAMPLE-28 SAMPLE-29	SAMPLE-36 SAMPLE-37					18 S 19 S 20 S 21 S	AMP AMP	PLE-1 PLE-2 PLE-3 PLE-4	mple		(ng/μL) 0.2440 500.0000 4.2600 1.9100		RBs
14 15 16 17	F G H Sta	0.068 0.023 andards	SAMPLE-6 SAMPLE-7 (Expiration d	SAMPLE-13 SAMPLE-14 ate:	SAMPLE-21	SAMPLE-28 SAMPLE-29	SAMPLE-36 SAMPLE-37 Dispense 2			ito each reacti		18 S 19 S 20 S 21 S 22 S	AMP AMP AMP	PLE-1 PLE-2 PLE-3 PLE-4 PLE-5	mple		(ng/μL) 0.2440 500.0000 4.2600 1.9100 3.8000		RBs
14 15 16 17 18	F G H Sta	0.068 0.023 andards 50ul [stock	SAMPLE-6 SAMPLE-7 (Expiration d	SAMPLE-13 SAMPLE-14 ate: 50.0ng/ul	SAMPLE-21	SAMPLE-28 SAMPLE-29	SAMPLE-36 SAMPLE-37 Dispense 2 Add 2ul of	sample, star	dard, or co	ntrol to the app		18 S 19 S 20 S 21 S 22 S 23 S	AMP AMP AMP AMP	PLE-1 PLE-2 PLE-3 PLE-4 PLE-5 PLE-6	mple		(ng/µL) 0.2440 500.0000 4.2600 1.9100 3.8000 1.5000	1/100	RBs
14 15 16 17 18 19	F G H Sta	0.068 0.023 andards 50ul [stock 50ul [STD 1	SAMPLE-6 SAMPLE-7 (Expiration d	SAMPLE-13 SAMPLE-14 ate: 50.0ng/ul 16.7ng/ul	SAMPLE-21	SAMPLE-28 SAMPLE-29	SAMPLE-36 SAMPLE-37 Dispense 2 Add 2ul of		dard, or co	ntrol to the app		18 S 19 S 20 S 21 S 22 S 23 S 24 S	AMP AMP AMP AMP AMP	PLE-1 PLE-2 PLE-3 PLE-4 PLE-5 PLE-6 PLE-7	mple		(ng/µL) 0.2440 500.0000 4.2600 1.9100 3.8000 1.5000		RBs
14 15 16 17 18 19 20 21	F G H Sta 1 2 3 4	0.068 0.023 andards 50ul [stock 50ul [STD 1 50ul [STD 2 50ul [STD 3	SAMPLE-6 SAMPLE-7 (Expiration d 3 + 150ul TE 1) + 100ul TE 2] + 100ul TE 3] + 100ul TE	SAMPLE-13 SAMPLE-14 ate: 50.0ng/ul 16.7ng/ul 5.56ng/ul 1.85ng/ul	SAMPLE-21	SAMPLE-28 SAMPLE-29	SAMPLE-36 SAMPLE-37 Dispense 2 Add 2ul of	sample, star	dard, or co	ntrol to the app		18 S 19 S 20 S 21 S 22 S 23 S 24 S 25 S	AMP AMP AMP AMP AMP	PLE-1 PLE-2 PLE-3 PLE-4 PLE-5 PLE-6 PLE-7 PLE-8	mple		(ng/µL) 0.2440 500.0000 4.2600 1.9100 3.8000 1.5000	1/100	RBs
14 15 16 17 18 19 20 21 22	F G H Sta 1 2 3 4 5	0.068 0.023 andards 50ul [stock 50ul [STD 1 50ul [STD 2 50ul [STD 3 50ul [STD 3	SAMPLE-6 SAMPLE-7 (Expiration d (1 + 150ul TE 11 + 100ul TE 22 + 100ul TE 33 + 100ul TE 41 + 100ul TE	SAMPLE-13 SAMPLE-14 ate: 50.0ng/ul 16.7ng/ul 5.56ng/ul 1.85ng/ul 0.62ng/ul	SAMPLE-21 SAMPLE-22	SAMPLE-28 SAMPLE-29	SAMPLE-36 SAMPLE-37 Dispense 2 Add 2ul of	sample, star	dard, or co	ntrol to the app		18 S 19 S 20 S 21 S 22 S 23 S 24 S 25 S 26 S	AMP AMP AMP AMP AMP AMP	PLE-1 PLE-2 PLE-3 PLE-4 PLE-5 PLE-6 PLE-7 PLE-8 PLE-9	mple		(ng/µL) 0.2440 500.0000 4.2600 1.9100 3.8000 1.5000	1/100	RBs
14 15 16 17 18 19 20 21 22 23	F G H Sta 1 2 3 4 5 6	0.068 0.023 andards 50ul [stock 50ul [STD 1 50ul [STD 2 50ul [STD 3 50ul [STD 4 50ul [STD 4	SAMPLE-6 SAMPLE-7 (Expiration d () + 150ul TE 1) + 100ul TE 2) + 100ul TE 3) + 100ul TE 4) + 100ul TE 5] + 100ul TE	SAMPLE-13 SAMPLE-14 ate: 50.0ng/ul 16.7ng/ul 5.56ng/ul 1.85ng/ul 0.62ng/ul 0.21ng/ul	SAMPLE-21 SAMPLE-22	SAMPLE-28 SAMPLE-29	SAMPLE-36 SAMPLE-37 Dispense 2 Add 2ul of	sample, star	dard, or co	ntrol to the app		18 S 19 S 20 S 21 S 22 S 23 S 24 S 25 S 26 S 27 S	AMP AMP AMP AMP AMP AMP AMP	PLE-1 PLE-2 PLE-3 PLE-4 PLE-5 PLE-6 PLE-7 PLE-8 PLE-9 PLE-10	mple		(ng/µL) 0.2440 500.0000 4.2600 1.9100 3.8000 1.5000 100.0000 0.1870	1/100	RBs
14 15 16 17 18 19 20 21 22 23 24	F G H Sta 1 2 3 4 5 6 7	0.068 0.023 andards 50ul [stock 50ul [STD 1 50ul [STD 2 50ul [STD 3 50ul [STD 4 50ul [STD 4 50ul [STD 5 50ul [STD 5 50ul [STD 5 50ul [STD 5 50ul [STD 6	SAMPLE-6 SAMPLE-7 (Expiration d () + 150ul TE 1) + 100ul TE 2) + 100ul TE 3) + 100ul TE 5) + 100ul TE 6) + 100ul TE	SAMPLE-13 SAMPLE-14 ate: 50.0ng/ul 16.7ng/ul 5.56ng/ul 1.85ng/ul 0.62ng/ul 0.21ng/ul 0.068ng/ul	SAMPLE-21 SAMPLE-22	SAMPLE-28 SAMPLE-29	SAMPLE-36 SAMPLE-37 Dispense 2 Add 2ul of	sample, star	dard, or co	ntrol to the app		18 S 19 S 20 S 21 S 22 S 23 S 24 S 25 S 26 S 27 S 28 S	AMP AMP AMP AMP AMP AMP AMP	PLE-1 PLE-2 PLE-3 PLE-4 PLE-5 PLE-6 PLE-7 PLE-8 PLE-9 PLE-10 PLE-11	mple		(ng/µL) 0.2440 500.0000 4.2600 1.9100 3.8000 1.5000	1/100	
14 15 16 17 18 19 20 21 22 23 24	F G H Sta 1 2 3 4 5 6 7	0.068 0.023 andards 50ul [stock 50ul [STD 1 50ul [STD 2 50ul [STD 3 50ul [STD 4 50ul [STD 4 50ul [STD 5 50ul [STD 5 50ul [STD 5 50ul [STD 5 50ul [STD 6	SAMPLE-6 SAMPLE-7 (Expiration d () + 150ul TE 1) + 100ul TE 2) + 100ul TE 3) + 100ul TE 4) + 100ul TE 5] + 100ul TE	SAMPLE-13 SAMPLE-14 ate: 50.0ng/ul 16.7ng/ul 5.56ng/ul 1.85ng/ul 0.62ng/ul 0.21ng/ul 0.068ng/ul	SAMPLE-21 SAMPLE-22	SAMPLE-28 SAMPLE-29	SAMPLE-36 SAMPLE-37 Dispense 2 Add 2ul of	sample, star	dard, or co	ntrol to the app		18 S 19 S 20 S 21 S 22 S 23 S 24 S 25 S 26 S 27 S 28 S 29 S	AMP AMP AMP AMP AMP AMP AMP	PLE-1 PLE-2 PLE-3 PLE-4 PLE-5 PLE-6 PLE-7 PLE-8 PLE-9 PLE-10 PLE-11 PLE-RB	mple		(ng/µL) 0.2440 500.0000 4.2600 1.9100 3.8000 1.5000 100.0000 0.1870	1/100	RBs
14 15 16 17 18 19 20 21 22 23 24	F G H Sta 1 2 3 4 5 6 7	0.068 0.023 andards 50ul [stock 50ul [STD 1 50ul [STD 2 50ul [STD 3 50ul [STD 4 50ul [STD 5 50ul [STD 6	SAMPLE-6 SAMPLE-7 (Expiration d () + 150ul TE 1) + 100ul TE 2) + 100ul TE 3) + 100ul TE 5) + 100ul TE 6) + 100ul TE	SAMPLE-13 SAMPLE-14 ate: 50.0ng/ul 16.7ng/ul 5.56ng/ul 1.85ng/ul 0.62ng/ul 0.21ng/ul 0.068ng/ul	SAMPLE-21 SAMPLE-22	SAMPLE-28 SAMPLE-29	SAMPLE-36 SAMPLE-37 Dispense 2 Add 2ul of	sample, star	dard, or co	ntrol to the app		18 S 19 S 20 S 21 S 22 S 23 S 24 S 25 S 26 S 27 S 28 S 29 S 30 S	AMP AMP AMP AMP AMP AMP AMP AMP	PLE-1 PLE-2 PLE-3 PLE-4 PLE-5 PLE-6 PLE-9 PLE-10 PLE-11 PLE-RB PLE-12	mple		(ng/µL) 0.2440 500.0000 4.2600 1.9100 3.8000 1.5000 100.0000 0.1870	1/100	
14 15 16 17 18 19 20 21 22 23 24	F G H Sta 1 2 3 4 5 6 7	0.068 0.023 andards 50ul [stock 50ul [STD 1 50ul [STD 2 50ul [STD 3 50ul [STD 4 50ul [STD 5 50ul [STD 6	SAMPLE-6 SAMPLE-7 (Expiration d () + 150ul TE 1) + 100ul TE 2) + 100ul TE 3) + 100ul TE 5) + 100ul TE 6) + 100ul TE	SAMPLE-13 SAMPLE-14 ate: 50.0ng/ul 16.7ng/ul 5.56ng/ul 1.85ng/ul 0.62ng/ul 0.21ng/ul 0.068ng/ul	SAMPLE-21 SAMPLE-22	SAMPLE-28 SAMPLE-29	SAMPLE-36 SAMPLE-37 Dispense 2 Add 2ul of	sample, star	dard, or co	ntrol to the app		18 S 19 S 20 S 21 S 22 S 23 S 24 S 25 S 26 S 27 S 28 S 29 S 30 S 31 S	AMP AMP AMP AMP AMP AMP AMP AMP	PLE-1 PLE-2 PLE-3 PLE-4 PLE-5 PLE-6 PLE-7 PLE-8 PLE-10 PLE-11 PLE-RB PLE-12 PLE-13	mple		(ng/µL) 0.2440 500.0000 4.2600 1.9100 3.8000 1.5000 100.0000 0.1870 0.0016	1/100	
14 15 16 17 18 19 20 21 22 23 24	F G H Sta 1 2 3 4 5 6 7	0.068 0.023 andards 50ul [stock 50ul [STD 1 50ul [STD 2 50ul [STD 3 50ul [STD 4 50ul [STD 5 50ul [STD 6	SAMPLE-6 SAMPLE-7 (Expiration d () + 150ul TE 1) + 100ul TE 2) + 100ul TE 3) + 100ul TE 5) + 100ul TE 6) + 100ul TE	SAMPLE-13 SAMPLE-14 ate: 50.0ng/ul 16.7ng/ul 5.56ng/ul 1.85ng/ul 0.62ng/ul 0.21ng/ul 0.068ng/ul	SAMPLE-21 SAMPLE-22	SAMPLE-28 SAMPLE-29	SAMPLE-36 SAMPLE-37 Dispense 2 Add 2ul of	sample, star	dard, or co	ntrol to the app		18 S 19 S 20 S 21 S 22 S 23 S 24 S 25 S 26 S 27 S 28 S 29 S 30 S 31 S 32 S	AMP AMP AMP AMP AMP AMP AMP AMP AMP	PLE-1 PLE-3 PLE-4 PLE-5 PLE-6 PLE-7 PLE-10 PLE-11 PLE-11 PLE-RB PLE-12 PLE-13 PLE-14	mple		(ng/µL) 0.2440 500.0000 4.2600 1.9100 3.8000 1.5000 100.0000 0.1870 0.0016 0.1540	1/100	
14 15 16 17 18 19 20 21 22 23 24 25	F G H Sta 1 2 3 4 5 6 7 8	0.068 0.023 andards 50ul [stock 50ul [STD 1 50ul [STD 2 50ul [STD 3 50ul [STD 4 50ul [STD 6 50ul [STD 6 50ul [STD 6	SAMPLE-6 SAMPLE-7 (Expiration d () + 150ul TE 1) + 100ul TE 2) + 100ul TE 3) + 100ul TE 5) + 100ul TE 6) + 100ul TE	SAMPLE-13 SAMPLE-14 ate: 50.0ng/ul 16.7ng/ul 5.56ng/ul 1.85ng/ul 0.62ng/ul 0.21ng/ul 0.068ng/ul 0.023ng/ul	SAMPLE-21 SAMPLE-22	SAMPLE-28 SAMPLE-29 1 2	SAMPLE-36 SAMPLE-37 Dispense 2 Add 2ul of NOTE: Sta	sample, star ndards shou	idard, or co Id be run in	ntrol to the app duplicate.		18 S 19 S 20 S 21 S 22 S 23 S 24 S 25 S 26 S 27 S 28 S 30 S 31 S 32 S 33 S	AMP AMP AMP AMP AMP AMP AMP AMP AMP	PLE-1 PLE-2 PLE-3 PLE-4 PLE-5 PLE-6 PLE-7 PLE-8 PLE-10 PLE-11 PLE-RB PLE-12 PLE-13	mple		(ng/µL) 0.2440 500.0000 4.2600 1.9100 3.8000 1.5000 100.0000 0.1870 0.0016	1/100	

- 2. Kit lot information is stored in the template itself.
- 3. By selecting the correct kit, all other lot information is automatically entered.
- 4. Master mix volumes are calculated based on reaction count.

After Importing the Quant data:

- 5. The required dilutions are automatically calculated.
- 6. Reagent Blanks are also automatically recognized.



All calculations are performed automatically using the Quant and Dilution data (from the previous worksheets).

- 8. Kit lot information is stored in the template itself
- 9. By selecting the correct kit, all other lot information is automatically entered, including the quant value of the positive control

	Α	В	С	D	E	F	G	Н	1	J	K	L	М
1					P	CR An	nplificat	ion Pla	ate				
2													
3	Lab nu	mber(s):	Demonstratio	n Data									
4													
5	Amp	Tray ID:	012208KM										
6	3832	1 1000											
7		Date:	1/22/2008										
8													
9		Analyst:	KM										
10													
11		1	2	3	4	5	6	7	8	9	10	11	12
12	Α		SAMPLE- 6_PP	SAMPLE- 13_PP	SAMPLE- 21_PP	SAMPLE- 28_PP	SAMPLE- 36_PP		SAMPLE- 6_CO	SAMPLE- 13_CO	SAMPLE- 21_CO	SAMPLE- 28_CO	SAMPLE- 36_CO
13	В		SAMPLE- 7_PP	SAMPLE- 14_PP	SAMPLE- 22_PP	SAMPLE- 29_PP	SAMPLE- 37_PP		SAMPLE- 7_CO	SAMPLE- 14_CO	SAMPLE- 22_CO	SAMPLE- 29_CO	SAMPLE- 37_CO
14	С		SAMPLE- 8_PP	SAMPLE- 15_PP	SAMPLE-RB 2_PP	SAMPLE- 30_PP	SAMPLE- 38_PP		SAMPLE- 8_CO	SAMPLE- 15_CO		SAMPLE- 30_CO	SAMPLE- 38_CO
15	D	SAMPLE- 1_PP	SAMPLE- 9_PP	SAMPLE- 16_PP	SAMPLE- 23_PP	SAMPLE- 31_PP	SAMPLE- 39_PP	SAMPLE- 1_CO	SAMPLE- 9_CO	SAMPLE- 16_CO	SAMPLE- A	SAMPLE- B	SAMPLE- C
16	F	SAMPLE- 2 PP	SAMPLE- 10 PP	SAMPLE- 17 PP	SAMPLE- 24 PP	SAMPLE- 32 PP	SAMPLE-RB 3 PP	SAMPLE- 2 CO	SAMPLE- 10 CO	SAMPLE- 17 CO	1 2		

SAMPLE-

3_CO

SAMPLE-

SAMPLE-

5_CO

4_CO

SAMPLE-

11_CO

SAMPLE-

SAMPLE-

SAMPLE- 5

18_CO

19_CO

SAMPLE

20_CO

10. Simple links populate the PCR Amplification Plate Grid

SAMPLE-

SAMPLE-

SAMPLE-

25_PP

SAMPLE-

SAMPLE-

SAMPLE-

18

SAMPLE-

SAMPLE-

SAMPLE-

11_PP

RB_PP

SAMPLE-18_PP

SAMPLE-

SAMPLE-

19_PP

SAMPLE-

33_PP

34_PP

35_PP

SAMPLE-

SAMPLE-

11. Selecting the instrument for injections alters the plate import worksheet that follows

ь													
7		Date:	1/22/2008										
8													
9		Analyst:	KM										
10													
11		1	2	3	4	5	6	7	8	9	10	11	12
	_	Ladder_PP		SAMPLE-	SAMPLE-	SAMPLE-	SAMPLE-	Ladder_CO	SAMPLE-	SAMPLE-	SAMPLE-	SAMPLE-	SAMPLE-
12	Α		6_PP	13_PP	21_PP	28_PP	36_PP		6_CO	13_CO	21_CO	28_CO	36_CO
	_	POS_PP		SAMPLE- 14_PP	SAMPLE- 22 PP	SAMPLE- 29 PP	SAMPLE- 37 PP	POS_CO	SAMPLE- 7_CO	SAMPLE- 14 CO	SAMPLE- 22 CO		SAMPLE- 37_CO
13	В		_	_	_	_	_		_	_	22_00	_	_
14	С	NEG_PP		SAMPLE- 15_PP	SAMPLE- RB-2_PP	SAMPLE- 30_PP	SAMPLE- 38_PP	NEG_CO	SAMPLE- 8_CO	SAMPLE- 15_CO			SAMPLE- 38_CO
		SAMPLE-		SAMPLE-	SAMPLE-	SAMPLE-	SAMPLE-	SAMPLE-	SAMPLE-	SAMPLE-	SAMPLE-		SAMPLE-
15	D	_	_	16_PP	_	_	39_PP	_	9_CO	16_CO	23_CO	_	39_CO
16	Е	SAMPLE- 2_PP		SAMPLE- 17_PP	SAMPLE- 24_PP	SAMPLE- 32_PP	SAMPLE-RB- 3_PP	SAMPLE- 2_CO	SAMPLE- 10_CO	SAMPLE- 17_CO	SAMPLE- 24_CO	SAMPLE- 32_CO	
17	F	SAMPLE- 3 PP		SAMPLE- 18 PP	SAMPLE- 25 PP	SAMPLE- 33 PP	Ladder_PP		SAMPLE- 11 CO	SAMPLE- 18 CO	SAMPLE- 25 CO	SAMPLE- 33 CO	Ladder_CO
17		SAMPLE-	_	SAMPLE-	_	SAMPLE-	DOO DD	SAMPLE-	11_00	SAMPLE-	SAMPLE-	SAMPLE-	BOD 00
18	G			19_PP		34_PP	POS_PP	4_CO		19_CO	26_CO	34_CO	POS_CO
19	н	SAMPLE- 5_PP		SAMPLE- 20_PP	SAMPLE- 27_PP	SAMPLE- 35_PP	NEG_PP	SAMPLE- 5_CO	SAMPLE- 12_CO	SAMPLE- 20_CO	SAMPLE- 27_CO	SAMPLE- 35_CO	NEG_CO
20	Note	Samples	in bold	were inje	cted for	10 seco	nds						
21		•											
22						Rox Lot							
23						Formamide							
24						PP Allelic							
25						CO Allelic	Ladder Lot						

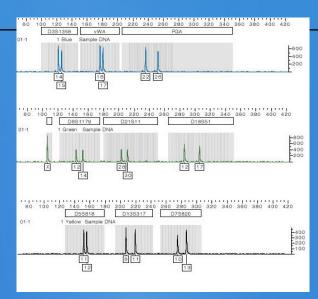
ABI CE Sample loading plate

Instrument:

Lab number(s): Demonstration Data

Run ID: 012208KM

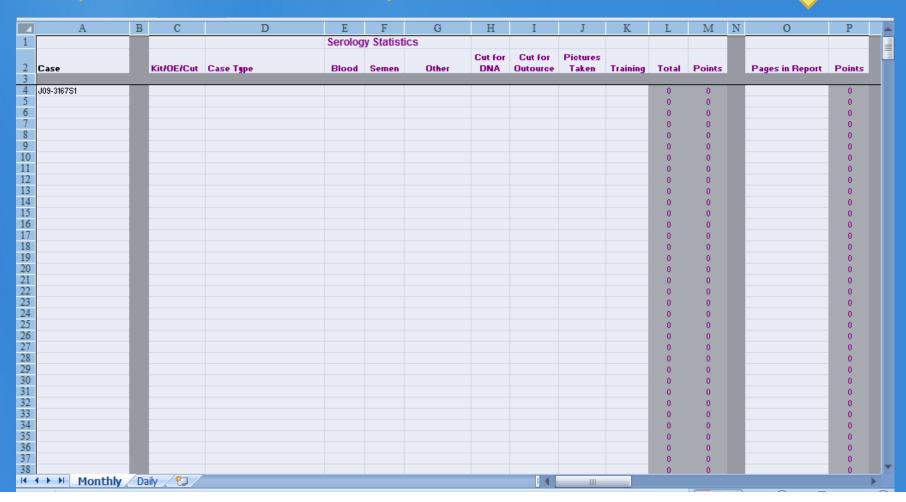
	Α	В	С	D	E	F	G	Н	1	J	K	L	М
1	Container Name	Description	ContainerType	AppType	Owner	Operator							
2	012208KM		96-Well	Regular	KM	KM							
3	AppServer	Applnstance											
4	GeneMapper	GeneMapper_d7dd4	151ab2a311d993	86000d56	d61fe6								
5	Well	Sample Name	Comment	Priority	Sample Type	Snp Set	Analysis Method	Panel	Size Standard	User-Defined	Results Gro	Instrument Pro	rtocol 1
6	A01	Ladder_PP		100	Allelic Ladder		HCMEO3100	Profiler_Plus_v1	RED 75-400	5_sec_inj	3100_run	5_sec_inj	
7	B01	POS_PP		100	Positive Control		HCMEO3100	Profiler_Plus_v1	RED 75-400	5_sec_inj	3100_run	5_sec_inj	
8	C01	NEG_PP		100	Negative Control		HCMEO3100	Profiler_Plus_v1	RED 75-400	10_sec_inj	3100_run	10_sec_inj	
9	D01	SAMPLE-1_PP		100	Sample		HCMEO3100	Profiler_Plus_v1	RED 75-400	5_sec_inj	3100_run	5_sec_inj	
10	E01	SAMPLE-2_PP			Sample		HCMEO3100	Profiler_Plus_v1	RED 75-400	5_sec_inj	3100_run	5_sec_inj	
11	F01	SAMPLE-3_PP		100	Sample		HCMEO3100	Profiler_Plus_v1	RED 75-400	5_sec_inj	3100_run	5_sec_inj	
	G01	SAMPLE-4_PP			Sample		HCMEO3100	Profiler_Plus_v1	RED 75-400	5_sec_inj	3100_run	5_sec_inj	
	H01	SAMPLE-5_PP		100	Sample		HCMEO3100	Profiler_Plus_v1		5_sec_inj	3100_run	5_sec_inj	
	A02	SAMPLE-6_PP		100	Sample		HCMEO3100	Profiler_Plus_v1		5_sec_inj	3100_run	5_sec_inj	
	B02	SAMPLE-7_PP			Sample		HCMEO3100	Profiler_Plus_v1		5_sec_inj	3100_run	5_sec_inj	
	C02	SAMPLE-8_PP			Sample		HCMEO3100	Profiler_Plus_v1		5_sec_inj	3100_run	5_sec_inj	
	D02	SAMPLE-9_PP			Sample		HCMEO3100	Profiler_Plus_v1		10_sec_inj	3100_run	10_sec_inj	
	E02	SAMPLE-10_PP			Sample		HCMEO3100	Profiler_Plus_v1			3100_run	10_sec_inj	
	F02	SAMPLE-11_PP		100	Sample		HCMEO3100	Profiler_Plus_v1		10_sec_inj	3100_run	10_sec_inj	
20	G02	SAMPLE-RB_PP		100	Sample		HCME03100	Profiler_Plus_v1	RED 75-400	10_sec_inj	3100_run	10_sec_inj	



- 12. Sample names are automatically imported from the previous worksheet.
- 13. Sample type and panel are determined by the sample name.
- 14. Analysis method, size standard and results group are determined by the instrument chosen on the proceeding worksheet.
- 15. Instrument protocol is decided based on Quantification values.

Productivity Index

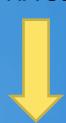




Productivity Index

DNA Stats







	W	X	Y	Z	AA	AB	AC	AD	AE	AF	AG	AH	AI	AJ	AK	AL	AM	Al .
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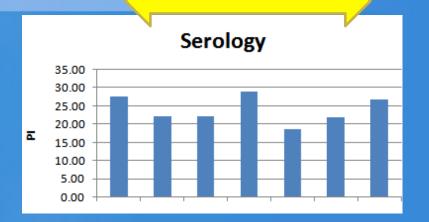
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			Print/Coll-	45-90 samples 1-44 samples total ate/Photocopy Data 45-90 samples 1-44 samples sued DNA Serology	0 0 0 0	0	0	
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			Print/Coll Reports Is	45-90 samples 1-44 samples total ste/Photocopy Data 45-90 samples 1-44 samples sued DNA Serology Rework DNA Tech Review DNA Admin Review Serology Tech Review Serology Tech Review Closed Report Outsource Review 10% Review	0 0 0 0 0 0 0 0 0	0	0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0	
			Print/Coll Reports Is	45-90 samples 1-44 samples total ate/Photocopy Data 45-90 samples 1-44 samples Steed DNA Serology Rework DNA Tech Review DNA Admin Review Serology Admin Review Serology Admin Review Closed Report Outsource Review Outsource Batches	0 0 0 0 0 0 0 0 0 0 0 0	0	0 0 0 0 0	
			Print/Coll. Reports Is Reviews	45-90 samples 1-44 samples total ate/Photocopy Data 45-90 samples 1-44 samples sued DNA Serology Rework DNA Tech Review DNA Admin Review Serology Tech Review Serology Tech Review Closed Report Outsource Review 10% Review Outsource Batches CODIS Review	0 0 0 0 0 0 0 0 0 0 0 0 0 0	0	0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0	
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			Print/Coll. Reports Is Reviews	45-90 samples 1-44 samples total ate/Photocopy Data 45-90 samples 1-44 samples 1-44 samples Seued DINA Serology Rework DINA Admin Review DINA Admin Review Serology Admin Review Closed Report Outsource Review 10% Review Outsource Batches CODIS Review Forensic Hit Offender Hit Conviction/Benchwork	0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0	0	0 0 0 0 0 0 0 0 0 0	
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			Print/Coll. Reports Is Reviews CODIS	45-90 samples 1-44 samples total ate/Photocopy Data 45-90 samples 1-44 samples 1-44 samples sued DNA Serology Rework DNA Tech Review DNA Admin Review Serology Tech Review Serology Tech Review Closed Report Outsource Review 10'x Review Outsource Batches CODIS Review Forensic Hit Offender Hit Conviction/Benchwork Weekly up/download Daily backup Biweekly media exchange Monthly stats	0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0	0	0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0	
			Print/Coll. Reports Is Reviews CODIS	45-90 samples 1-44 samples total ate/Photocopy Data 45-90 samples 1-44 samples 1-44 samples Seved DINA Serology Rework DINA Admin Review DINA Admin Review Serology Admin Review Closed Report Outsource Review 10% Review Outsource Batches CODIS Review Forensic Hit Conviction/Benchwork Weekly up/download Daily backup Bilweekly media exchange	0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0	0	0 0 0 0 0 0 0 0 0 0 0 0 0	
			Print/Coll. Reports Is Reviews CODIS CODIS Case Content of the content of th	45-90 samples 1-44 samples total ate/Photocopy Data 45-90 samples 1-44 samples sued DNA Serology Rework DNA Tech Review DNA Admin Review Serology Tech Review Serology Tech Review Closed Report Outsource Review 10'x Review Outsource Batches CODIS Review Forensic Hit Offender Hit Conviction/Benchwork Weekly up/download Daily backup Biweekly media exchange Monthly stats respondence onferences stiffed	0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0	0	0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0	
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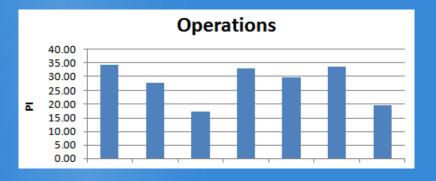


Metrics

Normalized

Serology	Hours Worked	Time Spent	% Time on		
Analyst	in Lab	on Points	Points	Points	PI
	166.75	148.05	89%	410.8	27.75
	151.1	142.8	95%	317.6	22.24
	157.5	107.2	68%	239.5	22.34
1	163.75	153.25	94%	445.9	29.10
	153.5	142.25	93%	265.7	18.68
	156.5	141.05	90%	309.5	21.94
	171.45	157.53	92%	421.9	26.78
Operations	Hours Worked	Time Spent	% Time on		
Analyst	in Lab	on Points	Points	Points	PI
	153	106.75	70%	366.4	34.32
	176.2	162.2	92%	451.59	27.84
	157.5	88.25	56%	153.3	17.37
	132.5	100.25	76%	330.4	32.96
	144	90.75	63%	271.1	29.87
	149.25	126.75	85%	428.6	33.81
	158.1	138.1	87%	272.75	19.75



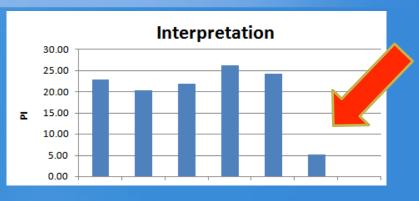




Metrics

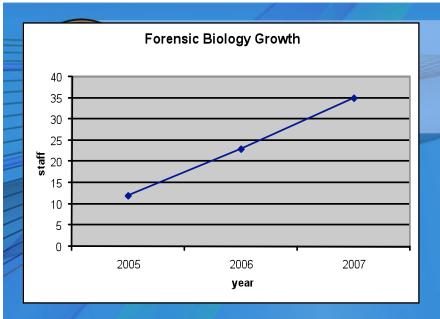
Outlier, low PI

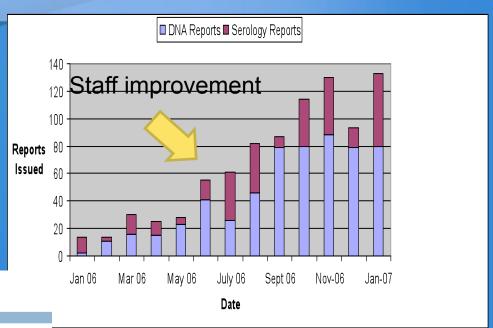
7						
	Interpretation	Hours Worked	Time Spent	% Time on		
	Analyst	in Lab	on Points	Points	Points	PI
		135	105.25	78%	240.4	22.84
		174	164	94%	333.2	20.32
		163	146.1	90%	320.2	21.92
		168.75	149	88%	392.15	26.32
		150	115	77%	278.2	24.19
		98.5	92.5	94%	47.4	5.12
	Compliance	Hours Worked	Time Spent	% Time on		
	Compliance Analyst	Hours Worked in Lab	Time Spent on Points	% Time on Points	Points	PI
	•		•		Points	PI 27.82
	•	in Lab	on Points	Points		
	•	in Lab 160.5	on Points 41.84	Points 26%	116.4	27.82
	•	in Lab 160.5 177	on Points 41.84 66.5	Points 26% 38%	116.4 204.3	27.82 30.72
	•	in Lab 160.5 177 127.5	on Points 41.84 66.5 26	Points 26% 38% 20%	116.4 204.3 85.5	27.82 30.72 32.88
	•	in Lab 160.5 177 127.5 178.8	on Points 41.84 66.5 26 64.9	Points 26% 38% 20% 36%	116.4 204.3 85.5 285.8	27.82 30.72 32.88 44.04



Outlier, High PI







DNA Reports



no increase in staff number since 2007



Process improvement

September 2008-2009:

- ~50% Increased monthly output
- ~85% Decreased backlog
- ~25% Increased CODIS monthly hits



Process improvement

- Increased monthly output
 - DNA cases reported September 2008 = 130
 - DNA cases reported September 2009 = 267
- Decreased backlog (>60 days)
 - Backlog as of September 2008 = 1077
 - Backlog as of September 2009 = 163
- Increased CODIS monthly hits
 - CODIS matches September 2008 = 21
 - CODIS matches September 2009 = 86



Thank You

Contact information:

Robin Freeman, M.S.

Forensic Biology-Interpretation Manager Harris County Medical Examiner's Office (713) 796-6974

robin.freeman@meo.hctx.net

Dr. Roger Kahn
Forensic Biology Director
Harris County Medical Examiner's Office
(713) 796-6978

roger.kahn@meo.hctx.net