This Manual is compiled of the following sections. If a section was revised during the year, each revision and date effective is listed. Ensure to use the appropriate effective date.

# **Serology Manuals in use for 2012**

Number		
order	Procedures	Effective Date
1	Amylase Diffusion Presumptive Test for Saliva	7/16/12
2	Amylase Diffusion Presumptive Test for Saliva	3/24/10
3	Acid Phosphatase Presumptive Test for Blood	3/24/10
4	Bloodstain Preparation of Whole Blood	7/16/12
5	Bloodstain Preparation of Whole Blood	3/24/10
6	Christmas Tree Stain for Spermatozoa	3/24/10
7	General Guidelines	3/24/10
8	Kastle-Meyer (KM) Presumptive Test for Blood	3/24/10
9	P30 ELISA	7/16/12
10	P30 ELISA	3/24/10
11	Processing of Postmortem Specimens	3/24/10
12	References	3/24/10
13	Slide Preparation for Spermatozoa Searches	3/24/10



# FORENSIC BIOLOGY SEROLOGY MANUAL

Approving Authority: Eugene Y. Lien, Technical Leader, Serology Operations

Procedure	Effective Date	Revision History
General Guidelines	3/24/2010	
Processing of Postmortem Spremens	7/16/2012	
Bloodstain Preparation from Whole Blood	7/16/2012	
Kastle-Meyer (KM) Presumptive Test for Blood	3/24/2010	
Acid Phosphatase Presumptive Test for Semen	3/24/2010	
Amylase Diffusion Presumptive Test for Saliva	7/16/2012	
Slide Preparation for Spermatozoa Searches	9/17/2012	<b>Initial Version</b>
Christmas Tree Stain for Spermatazoa	3/24/2010	
P30 ELISA	7/16/2012	
<u>References</u>		

	AMYLASE I	DIFFUSION PRESUMPTIVE TEST F	OR SALIVA
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# **Preparing Amylase Plate**

1. Prepare starch-containing agarose gel by adding the ingredients listed below:

100 ml batch size (enough for 2 plates)
1.0g Sigma Type I agarose
0.1g potato starch
100mL amylase gel buffer

To dissolve, mix and boil this solution. Allow to slightly cool. Pour 40mL each into a 10 x 10cm disposable Petri dishes. Avoid air bubbles as much as possible. Scale up batch size when necessary.

- 2. Punch wells in the gel using the suction tube apparatus leaving at least 1.5cm between wells. Use Amylase sheet as a template.
  - Make sure that the holes that you create a completely clean of agar debris and residual liquid. This can be ensured by punching each hole twice in succession.
  - Following this protocol and with an accurate dispersion of agarose will guarantee an adequate amount of space for the loading of 10uL each of standard, control, or sample into each well.
- 3. Use Parafilm® around the lightestom joint to seal the amylase plates.
- 4. Store in a 4°C refrigerator upside-down (resting on the lid) to avoid condensation on the gel.
- 5. Pre-made plates are good for one week.

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## **Preparing Standards/Samples**

- 1. Extract approximately a 5 x 5mm stain or a portion of a swab in 100uL deionized water for 30 minutes at room temperature using the pipette tip and test tube method. For samples that have been analyzed with P30 ELISA, use the extracts prepared in that procedure.
- 2. Prepare α-amylase standards containing 0.02 and 0.002 units each per 10 uL of deionized water (dH<sub>2</sub>O) from purchased amylase.
  - A. Prepare 1mL of 20 units/10uL amylase by adding the appropriate amount of amylase standard to dH<sub>2</sub>O. The appropriate amount of amylase standard to add is determined by the QC of the current lot of amylase. See example calculation below.
  - B. Continue to prepare the remaining 2, 2, 2.02, and 0.002 unit standards by doing ten-fold serial dilutions. This is easily accomplished by first adding 900uL of dH<sub>2</sub>O to each of 4 microcentrifuge tubes. Then transfer 100uL of your 20 unit standard into one of the tubes containing 900uL of dH<sub>2</sub>O. This is your 2 unit standard. Continue making the remaining dilutions in the same manner.

When doing serial dilutions, make sure to mix each standard well before each subsequent transfer. Use a fresh unplugged pipette tip for each transfer.

Sample calculation

Given a specific activity of 870 units amylase/mg total protein (from vendor) with a total protein concentration of 30 mg/mL, then:

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Use this value in the equation C1 x V1 = C2 x V2 where C1 and C2 are concentrations of solutions 1 and 2, while V1 and V2 are volumes of solutions 1 and 2. In this case, solution 1 is the vendor amylase stock solution while solution 2 is the 20 units standard in preparation:

(26.1 units amymase/uL)(x uL) = (20 units/10uL)(1000uL)

Solving for x = 77uL of vendor amylase stock solution 1000uL (total volume) - 77uL (amylase stock solution) = 923uL of  $Q_2O$ 

# **Loading/Incubation/Staining of Amylase Plates**

- 1. Have a witness verify the Amylase documentation with the tube labels.
- 2. Fill wells according to the Amylase Diffusion forumentation (10 uL each well) with standards, negative control (deionized water), and samples. The first two wells are reserved for the 0.02U and 0.002U amylase standard, the negative control is added to the third well, and the remaining wells are filled with samples.
- 3. Incubate 5-8 hours at 37°C or 12-16 hours at room temperature; keep the plate in a humid chamber to avoid drying.
- 4. Pour a 0.01N (100-fold dilution of a 1N stock) iodine solution onto the gel; clear areas indicate regions of amylase activity. **Do not over stain the plate.** Do this by monitoring the plate as it is staining; pour off the iodine solution when a sufficient amount of staining has occurred so that all the standards are clearly visible.
- 5. Photograph the results via the Mideo System. Ensure there is a scale in the photograph. Save the file as a JEPG and upload to the LIMS system for the related Amylase assay.
- 6. Measure the diameter of the clear areas and record on the documentation.

Entire items (blood spatter patterns, etc.) can be tested for amylase. Prepare a large plate on a bordered glass plate (scale up reagents) and allow to solidify; bring item (or area of item) into contact with the gel for 5 minutes. Follow steps 3-6 above to visualize any amylase pattern.

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# **Interpretation of Results**

The values of diffusion for the 0.02 and 0.002 unit standards should fall in the ranges of 7-15 and 4-10 mm, respectively. In addition, the amount of diffusion of the 0.02 unit standard must be greater than that of the 0.002 unit standard.

The interpretation of amylase results depends on the source of the sample:

- 1. Body cavity swabs (e.g., vaginal and anal) are positive if the diameter is equal to or greater than the diameter of the 0.02 U standard. Designate as or the ("O") on the amylase documentation.
- 2. Samples not from a body cavity (e.g., penile swabs, creater butts, cups, etc.) are positive if the diameter is equal to or greater than the 4002U standard. Designate as external ("E") on the amylase documentation.
- 3. The location from which a "dried secretion" swab is taken affects the interpretation. Swabs taken essentially from a body cavity or similar place (e.g., introitus, etc.) are interpreted as if the sample is from a body cavity. Other locations (e.g., breast, thigh, penis, etc.) may need to be interpreted differently.

Below is a general list of common sample types and designation as **body cavity**/**orifice** samples:

OUTSIDE/OUTER ANUS
OUTSIDE/OUTER VAGINA/L
PERIANAL
PERINEAL
PERINEUM
PERIORAL
PERIVAGINAL
VESTIBULE
VULVA

#### AMYLASE DIFFUSION PRESUMPTIVE TEST FOR SALIVA APPROVING AUTHORITY DATE EFFECTIVE **PAGE** 07-16-2012 QUALITY ASSURANCE MANAGER 5 OF 5

Below is a general list of common sample types and designation as **external** samples:

**BUTTOCKS INGUINAL CHEEK INNER THIGH** 

**CHIN** MONS VENEVIS (mons pubis)

**GROIN PENIS** 

**SCROTUM** 

Archived for 2012 Manuals

**Revision History:** 

March 24, 2010 – Initial version of procedure.

July 16, 2012 - Specific names of worksheets were removed and replaced with generic terminology to accommodate LIMS.

AMYLASE D	OFFUSION PRESUMPTIVE TEST	FOR SALIVA
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# **Preparing Amylase Plate**

1. Prepare starch-containing agarose gel by adding the ingredients listed below:

100 ml batch size (enough for 2 plates)
1.0g Sigma Type I agarose
0.1g potato starch
100mL amylase gel buffer

To dissolve, mix and boil this solution. Allow to slightly cool. Four 40mL each into a 10 x 10cm disposable Petri dishes. Avoid air bubbles as much as possible. Scale up batch size when necessary.

- 2. Punch wells in the gel using the suction tube apparatus leaving at least 1.5cm between wells. Use Amylase sheet as a template.
  - Make sure that the holes that you create a completely clean of agar debris and residual liquid. This can be ensured by punching each hole twice in succession.
  - Following this protocol and with an accurate dispersion of agarose will guarantee an adequate amount of space for the loading of 10uL each of standard, control, or sample into each well.
- 3. Use Parafilm® around the lightestom joint to seal the amylase plates.
- 4. Store in a 4°C refrigerator upside-down (resting on the lid) to avoid condensation on the gel.
- 5. Pre-made plates are good for one week.

AMYLASE DI	IFFUSION PRESUMPTIVE TEST	F FOR SALIVA
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# **Preparing Standards/Samples**

- 1. Extract approximately a 5 x 5mm stain or a portion of a swab in 100uL deionized water for 30 minutes at room temperature using the pipette tip and test tube method. For samples that have been analyzed with P30 ELISA, use the extracts prepared in that procedure.
- 2. Prepare α-amylase standards containing 0.02 and 0.002 units each per 10 uL of deionized water (dH<sub>2</sub>O) from purchased amylase.
  - A. Prepare 1mL of 20 units/10uL amylase by adding the appropriate amount of amylase standard to dH<sub>2</sub>O. The appropriate amount of amylase standard to add is determined by the QC of the current lot of amylase. See example calculation below.
  - B. Continue to prepare the remaining 2, 2, 2.02, and 0.002 unit standards by doing ten-fold serial dilutions. This is easily accomplished by first adding 900uL of dH<sub>2</sub>O to each of 4 microcentrifuge tubes. Then transfer 100uL of your 20 unit standard into one of the tubes containing 900uL of dH<sub>2</sub>O. This is your 2 unit standard. Continue making the remaining dilutions in the same manner.

When doing serial dilutions, make sure to mix each standard well before each subsequent transfer. Use a fresh unplugged pipette tip for each transfer.

Sample calculation

Given a specific activity of 870 units amylase/mg total protein (from vendor) with a total protein concentration of 30 mg/mL, then:

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Use this value in the equation C1 x V1 = C2 x V2 where C1 and C2 are concentrations of solutions 1 and 2, while V1 and V2 are volumes of solutions 1 and 2. In this case, solution 1 is the vendor amylase stock solution while solution 2 is the 20 units standard in preparation:

(26.1 units amymase/uL)(x uL) = (20 units/10uL)(1000uL)

Solving for x = 77uL of vendor amylase stock solution 1000uL (total volume) - 77uL (amylase stock solution) = 923uL of  $U_2O$ 

# **Loading/Incubation/Staining of Amylase Plates**

- 1. Have a witness verify the Amylase Worksheet with the labels.
- 2. Fill wells according to the Amylase Diffusion Worksheet (10 uL each well) with standards, negative control (deionized water), and samples. The first two wells are reserved for the 0.02U and 0.002U amylase standard, the negative control is added to the third well, and the remaining wells are filled with samples.
- 3. Incubate 5-8 hours at 37°C or 12-16 hours at room temperature; keep the plate in a humid chamber to avoid drying.
- 4. Pour a 0.01N (100-fold direction of a 1N stock) iodine solution onto the gel; clear areas indicate regions of amylase activity. **Do not over stain the plate.** Do this by monitoring the plate as it is staining; pour off the iodine solution when a sufficient amount of staining has occurred so that all the standards are clearly visible.
- 5. Photograph the results via the Mideo System. Ensure there is a scale in the photograph.
- 6. Measure the diameter of the clear areas and record on the worksheet.

Entire items (blood spatter patterns, etc.) can be tested for amylase. Prepare a large plate on a bordered glass plate (scale up reagents) and allow to solidify; bring item (or area of item) into contact with the gel for 5 minutes. Follow steps 3-6 above to visualize any amylase pattern.

AMYLASE DI	FFUSION PRESUMPTIVE TEST	F FOR SALIVA
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# **Interpretation of Results**

The values of diffusion for the 0.02 and 0.002 unit standards should fall in the ranges of 7-15 and 4-10 mm, respectively. In addition, the amount of diffusion of the 0.02 unit standard must be greater than that of the 0.002 unit standard.

The interpretation of amylase results depends on the source of the sample:

- 1. Body cavity swabs (e.g., vaginal and anal) are positive if the diameter is equal to or greater than the diameter of the 0.02 U standard. Designate as or in e ("O") on amylase worksheet.
- 2. Samples not from a body cavity (e.g., penile swabs, creater butts, cups, etc.) are positive if the diameter is equal to or greater than the 4002U standard. Designate as external ("E") on amylase worksheet.
- 3. The location from which a "dried secretion" swab is taken affects the interpretation. Swabs taken essentially from a body cavity or similar place (e.g., introitus, etc.) are interpreted as if the sample is from a body cavity. Other locations (e.g., breast, thigh, penis, etc.) may need to be interpreted differently.

Below is a general list of compon sample types and designation as **body cavity**/<u>orifice</u> samples:

OUTSIDE/OUTER ANUS **ANUS EXTERNAL GE** IA (female) OUTSIDE/OUTER VAGINA/L PERIANAL PERINEAL INTROITUS **PERINEUM** LABIA MAJORA PERIORAL LABIA MINORA **PERIVAGINAL** LIPS VESTIBULE **MOUTH** VULVA

#### AMYLASE DIFFUSION PRESUMPTIVE TEST FOR SALIVA APPROVED BY DATE EFFECTIVE PAGE 03-24-2010 **EUGENE LIEN** 5 OF 5

Below is a general list of common sample types and designation as **external** samples:

**BUTTOCKS INGUINAL CHEEK INNER THIGH** 

**CHIN** MONS VENEVIS (mons pubis)

**GROIN PENIS** 

**SCROTUM** 

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ACID PHOSI	PHATASE PRESUMPTIVE TEST	FOR SEMEN
DATE EFFECTIVE	APPROVED BY	PAGE
03-24-2010	EUGENE LIEN	1 OF 1

#### **GENERAL**

An Acid Phosphatase test is a presumptive test for semen. It may be performed directly on a cut out portion of a stain, an extract of a stain, or a "wipe" of the stained material. A wipe may be made using a piece of filter paper, thread, or swab. Wet the wipe with water, then rub over the stained area while still wet.

# **CONTROLS**

Analysts using Acid Phosphatase test reagents must test each lot/aliquor of reagent at least once per day, using positive and negative controls, before any evidence terms are tested. The results of this test shall be recorded in the case notes. Semen must be used as a positive control. A drop of deionized water may be used for the negative control. Resultiols do not pass, inform the Quality Assurance Team immediately.

#### **PROCEDURE**

- Apply a drop of the Alpha-Naphthyl Rhosphate reagent; wait 60 seconds.
   If a purple color occurs at this point, the testing results should indicate "inconclusive."
- 2. Apply a drop of the Fast Brue B reagent. An immediate purple color is a positive reaction.

BLOODS	TAIN PREPARATION FROM WHOL	E BLOOD
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# Staincards are prepared from all vouchered blood samples and from post-mortem blood samples:

- 1. Take custody of the blood vials awaiting bloodstain preparation.
- 2. Prepare the UltraSTAIN<sup>TM</sup> cards by affixing a pre-printed FB case number sticker (if available) and writing in the following:
  - Initials of person preparing the stain
  - FB number, if no sticker is available

Wear latex gloves when handling these cards.

- 3. Preparation of the bloodstain **must** be witnessed by mather laboratory staff member. The witness must confirm that the processor is mandling the correct blood vial and stain card BEFORE the stain is made. After each stair is made, the witness must initial the stain card and the evidence packaging worksheet.
- 4. Prepare stains one at a time. Staining of the cards and the opening of liquid blood samples MUST be performed under abiological safety cabinet with the exhaust fan operating. It is advisable that a new KimWipe<sup>TM</sup> be used to open each vial stopper. Make sure a blood tube is closed before preparing the next stain.
- 5. Fold back the paper "flat" and make four stains on the card, placing the blood in the outlined areas. Use four drops of blood per area; apply the drops slowly, allowing them to soak in. This with prevent appreciable transfer to the paper "flap".
- 6. Bring down the paper "flap", turn the entire card over, and allow it to air-dry upside down. The stain cards must be allowed to dry overnight before storage.
- 7. Package the air-dried stains into a 4x6" KAPAK<sup>TM</sup> bag. Heat seal the KAPAK<sup>TM</sup>. The person sealing the bag must date and initial the bag. Store at room temperature, and record the storage location for the chain of custody.

	BLOODS	TAIN PREPARATION FROM WHOL	E BLOOD
ĺ	DATE EFFECTIVE	APPROVING AUTHORITY	PAGE
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- 8. CLEAN THE BIOLOGICAL SAFETY CABINET (refer to QC Procedure #QC125 of the Quality Assurance/Quality Control Manual).
- 9. Place all case files that contain **any** sexual assault evidence in the designated area so that they may be processed. Place all cases files that contained any evidence from the NYPD or DA's office back from where they were retrieved (either "cases to be called on," "cases to be assigned," or the assigned analyst). Place all remaining case folders in the Forensic Biology office so that they may be filed.
- 10. Disposal of blood and blood vials:

For non-vouchered blood, the remainder of the liquid blood and the blood vial will be discarded immediately. Purple-topped vials **must** be tis a ded in a plastic BIOHAZARD "sharps" container.

For vouchered blood, the remainder of the liquid flood is discarded into bleach immediately after making the bloodstain and. The empty vial rinsed with 10% bleach. The empty vial is packaged for return to the Evidence Unit.



March 24, 2010 – Initial version of procedure.

July 16, 2012 - Specific terminology was removed and replaced with generic terminology to accommodate LIMS.

BLOODSTA	IN PREPARATION FROM WHO	OLE BLOOD
DATE EFFECTIVE	APPROVED BY	PAGE
03-24-2010	EUGENE LIEN	1 OF 2

# Staincards are prepared from all vouchered blood samples and from post-mortem blood samples:

- 1. Find the case files and obtain the blood vials; sign the chain of custody forms.
- 2. Prepare the UltraSTAIN<sup>TM</sup> cards by affixing a pre-printed FB case number sticker (if available) and writing in the following:
  - Subject's name
  - Date
  - Initials of person preparing the stain
  - FB number, if no sticker is available

Wear latex gloves when handling these cards.

- 3. Preparation of the bloodstain **must** be witnessed by another laboratory staff member. After each stain is made, the witness must initial the stain card and the evidence packaging worksheet. The witness must comfirm that the processor is handling the correct blood vial and stain card BEFOR the stain is made.
- 4. Prepare stains one at a time. Staining of the cards and the opening of liquid blood samples MUST be performed under a biological safety cabinet with the exhaust fan operating. It is advisable that a new KimWipe<sup>TM</sup> be used to open each vial stopper. Make sure a blood tube is closed before preparing the next stain.
- 5. Fold back the paper "flap" and make four stains on the card, placing the blood in the outlined areas. Use four drops of blood per area; apply the drops slowly, allowing them to soak in. This will prevent appreciable transfer to the paper "flap".
- 6. Bring down the paper "flap", turn the entire card over, and allow it to air-dry upside down. The stain cards must be allowed to dry overnight before storage.
- 7. Package the air-dried stains into a 4x6" KAPAK<sup>TM</sup> bag. Heat seal the KAPAK<sup>TM</sup>, the person sealing the bag must date and initial the bag. Store at room temperature. Continue the chain of custody to reflect their final location.

BLOODSTA	AIN PREPARATION FROM WHO	OLE BLOOD
DATE EFFECTIVE	APPROVED BY	PAGE
03-24-2010	EUGENE LIEN	2 OF 2

- 8. CLEAN THE BIOLOGICAL SAFETY CABINET (refer to QC Procedure #QC125 of the Quality Assurance/Quality Control Manual).
- 9. Place all case files that contain **any** sexual assault evidence in the designated area so that they may be processed. Place all cases files that contained any evidence from the NYPD or DA's office back from where they were retrieved (either "cases to be called on," "cases to be assigned," or the assigned analyst). Place all remaining case folders in the Forensic Biology office so that they may be filed.
- 10. Disposal of blood and blood vials:

For non-vouchered blood, the remainder of the liquid blood and the blood vial will be discarded immediately. Purple-topped vials **must** be tis added in a plastic BIOHAZARD "sharps" container.

For vouchered blood, the remainder of the liquid flood is discarded into bleach immediately after making the bloodstain (and). The empty vial rinsed with 10% bleach. The empty vial is packaged for return (a) he Evidence Unit.

Revision History:

CHRIST	MAS TREE STAIN FOR SPERM	ATOZOA
DATE EFFECTIVE	APPROVED BY	PAGE
03-24-2010	EUGENE LIEN	1 OF 1

The nuclear material within the cell is stained red by the Nuclear Fast Red stain. Sperm heads are usually well differentiated with the acrosome staining significantly less dense than the distal region of the head. Epithelial membranes and sperm tails are stained green by the Picric Indigo Carmine (PIC) stain; nuclei inside epithelial cells appear purple. Yeast cells also stain red, however the stain is uniform throughout the cell and extends into polyp-like structures that are occasionally seen in yeast.

# Reagents: Nuclear Fast Red and Picric Indigo Carmine

- 1. Fix cells to the slide by heating (approximately 5 to 10 seconds).
- 2. Cover cell debris with Nuclear Fast Red stain and allow to sit for at least 10 minutes.
- 3. Wash away the nuclear fast red with deionized water.
- 4. Add PIC stain to the still-wet slide; allow to it for no more than 30 seconds.
- 5. Wash away the PIC stain with ethanol
- 6. Place slide over a heat source to conslete drying.
- 7. Examine the slide at 100X of 400X (don't use immersion oil).

	GENERAL GUIDELINES	
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- 1. The procedures within this Serology Procedures Manual are intended to support the processes outlined in the Evidence Examination Procedure in the Evidence and Case Management Manual.
- 2. In general, screening tests and/or confirmatory tests are used to identify physiological fluids such as blood, semen, and saliva prior to further analysis.
- 3. All reagents are available pre-made and are quality control checked, where possible. Do not make your own or use supplies that have not been quality control checked. If reagents are needed, contact the Quality Assurance Unit for assistance.

Revision History:

March 24, 2010 – Initial version of procedure.

KASTLE-MEYE	R (KM) PRESUMPTIVE TESTI	NG FOR BLOOD
DATE EFFECTIVE	APPROVED BY	PAGE
03-24-2010	EUGENE LIEN	1 OF 1

#### **GENERAL**

A Kastle-Meyer test may be performed directly on a cut out portion of a stain, an extract of a stain, or a "wipe" of the stained material. A wipe may be made using a piece of filter paper, thread, or swab. Wet the wipe with water, then rub over the stained area while still wet.

#### **CONTROLS**

Positive and negative controls must be used to test each lot/aliquot of reagent at least once per day and before any evidence items are tested. Blood must be used as a positive control. A drop of deionized water may be used for the negative control. If controls do not pass, inform the Quality Assurance Team immediately.

#### **PROCEDURE**

1. Apply a drop of KM reagent if using aw pe. If performing directly on a cut out portion of a stain, use enough until sample is covered. Observe any color change.

A normal color reaction is a green sh/gray tint with the presence of possible blood.

A PINK COLOR HERE TO THE PRESENCE OF AN OXIDIZING AGENT (e.g., a chemical oxidant), NOT BLOOD. If a pink color occurs at this point, the testing results should indicate "inconclusive."

2. Add a drop of % hydrogen peroxide. An immediate pink color is a positive result.

#### P30 ENZYME LINKED IMMUNOSORBENT ASSAY (ELISA) DATE EFFECTIVE APPROVING AUTHORITY PAGE 07-16-2012 QUALITY ASSURANCE MANAGER 1 OF 14

**Solutions:** Phosphate buffered saline (PBS)

PBS-Casein (PBS with 0.02% w/v casein)

P30 antigen [prostate specific antigen (PSA)] and phosphate buffered saline **Standards:** 

**Plates:** Immulon II microELISA plates (microtiter plate)

**Antibodies:** Mouse monoclonal anti-human PSA (prostate specific antigen, P30)

Rabbit polyclonal anti-human PSA

Goat anti-rabbit IgG alkaline phosphatase conjugate Mouse IgG1, Kappa chain (MOPC 21, mouse myelona) 2012/1/8

Note: Store all antibodies at 4°C.

#### SOLUTION PREPARATION

#### **Phosphate buffered saline (PBS):**

- tablets in 1 liter of deionized water. This can be To prepare 1 liter, dissor 1. stored at 4°C for up
- On label write init 2. preparer, the date of make (DOM) and the date of discard (DOI

pax of plates you will need 1 bottle (1 liter). For each plate or

# PBS-Casein (PBS with 0.02% w/v Hammerstein casein):

#### If preparing from Frozen Casein Aliquots:

- 1. Thaw casein at room temperature.
- 2. Dilute 20mL of casein stock solution to 1 liter of PBS.

#### If preparing from a Bottle of Liquid Stock Casein (Refrigerated):

- 1. Shake the bottle well.
- 2. Using a graduated cylinder, take 20mL of casein stock solution and dilute it into 1 liter of PBS.

One plate uses approximately 500mL.

P30 ENZYN	ME LINKED IMMUNOSORBENT AS	SAY (ELISA)
DATE EFFECTIVE	APPROVING AUTHORITY	PAGE
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#### PLATE PREPARATION:

#### COATING THE PLATES

- 1. Prepare an appropriate dilution of mouse monoclonal anti-human PSA by adding the antiserum to PBS as determined by the QC of the current lot of antiserum. Each plate requires about 5mL of diluted antiserum. **Always make dilutions in glass, not in plastic.**
- 2. Prepare a 1/8000 dilution of MOPC by adding 10uL MOPC to 80 mL RB3. Always make dilutions in glass, not in plastic. This is enough for 16 plates. For collates add 2.5uL MOPC to 20mL PBS.
- 3. Coat the plate as shown in the diagram. Use 100uL of the appropriate solution per well.

PBS = phosphate buffered saline αPSA = mouse monoclonal anti-human PSA

M = MOPC

	1	2	3	4	5	6	7	8	9	10	11	12
A	PBS	M	M	M	C <sub>W</sub> X	M	M	M	M	M	M	M
В	PBS	M	M	М	M	M	M	M	M	M	M	M
C	PBS	αPSA	αPSA		αPSA	αPSA	αPSA	αPSA	αPSA	αPSA	αPSA	αPSA
D	PBS	αPSA	αPSA	αPSA	αPSA	αPSA	αPSA	αPSA	αPSA	αPSA	αPSA	αPSA
E	PBS	M	M	M	M	M	M	M	M	M	M	M
F	PBS	M	М	M	M	M	M	M	M	M	M	M
G	PBS	αPSA	αPSA	αPSA	αPSA	αPSA	αPSA	αPSA	αPSA	αPSA	αPSA	αPSA
Н	PBS	αPSA	αPSA	αΡSΑ	αPSA	αΡSΑ	αΡSΑ	αPSA	αPSA	αΡSΑ	αΡSΑ	αPSA

4. Cover plates with Parafilm® and label "Coating" with initials and date. Incubate overnight (approximately 17-20 hours) at 4°C.

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# **BLOCKING THE PLATES:**

- 1. Prime/Rinse washer (See Rinsing and Priming procedure below.)
- 2. Use the "Block 1" program to aspirate contents of well. (This will wash each plate twice and fill the wells with wash buffer) Let the filled plate sit at room temperature for 15-20 minutes (see Block 1 procedure).
- 3. Use the "Block 2" program to aspirate the final wash (See Block 2 procedure).
- 4. Tap plate upside-down on paper towels to remove access liquid
- 5. Plates that are not used immediately should be wrapped in the film® and stored at 4°C; they can be stored for up to 2 weeks.

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# **Instructions for Bio-Rad 1575 washers**

The Bio-Rad 1575 plate washer has a cover that can be swiveled to an opened or closed position.

# **Rinsing and priming**

- 1. The on/off switch is located on the right-hand side of the lower back of the washer. Turn plate washer on.
- 2. After initialization (concurrent with a quick flash of a Bio-Rad version#), the **main menu** appears and looks like this:



The circles with the triangles above represent the soft keys that you will be pressing to access various functions on the machine. In this menu, soft keys corresponding to "In" and "Out", when pressed instruct the plate washer to move the plate holder in and out of the washer, respectively. Soon after turning on the plate washer, the plate holder automatically sets itself in the out position.

- 3. Fill the **Rinse** bottle with deionized water, close the lid, and attach the tube with the blue colored line to the top of the container. Also, make sure that the stopper connected to the yellow and red colored lines is snugly connected to the **Waste** bottle (if not, the plate washer will not aspirate which can result in flooding of the plate holder).
- 4. Prime the washer with deionized water (from the Rinse bottle). Do this by pressing the very first soft key corresponding to the upward arrow in the main menu. Now you will see the Prime/Rinse menu. Press the soft key corresponding to "YES" and the washer will proceed to prime itself. This step is important to ensure proper vacuum pressure and for the washer to dispense correct volumes. Always make sure to re-prime the washer whenever changing the blue colored tube from the Wash bottle to the Rinse bottle and vice versa.
- 5. Fill the **Wash** bottle with PBS-casein solution. Remove the blue-striped tube from the **Rinse** bottle and attach it to the **Wash** bottle. Repeat the priming procedure 1 time using the PBS-casein solution from the **Wash** bottle. You are now ready to run the plate washer.

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#### **Running plate washer programs**

#### BLOCK 1

- 1. To start the blocking procedure, insert a coated plate into the plate washer lengthwise, so that plate well #A1 is located on the upper right portion of the plate
- 2. Proceed with the first portion of the blocking procedure in which the contents of the wells are aspirated, washed 2 times, and then filled with PBS-casein. The program hat does this is called **BLOCK1**.
- 3. To access the **BLOCK1** program, press the soft key that corresponds to "**YES**" while in the main menu. You will now be in the **run menu**. While in the **un** menu, scroll to the **BLOCK1** program by pressing the soft keys that correspond to the upward and downward arrows.
  - Before proceeding, you should see "FON: BLOCK1" on the first line of the display. Press the "YES" key.
  - You will then see "LAST STKP 12" on the first line of the display. Press the "YES" key again.
  - Usually you will then be one more prompt ("CONNECT THE WASH R9") on the first line of the display. Press the "YES" key once again.

The program will then can on its own, leaving the wells filled at the end of the program.

4. Incubate the plactor 15-20 min. at room temperature as described in the P30 protocol.

#### **BLOCK 2**

5. If you have removed your plate for the incubation step, place plate back into the plate washer by using the "IN" and "OUT" keys from the main menu. The last step of the blocking procedure requires aspiration of the wells. This is achieved by the BLOCK2 program. Access the BLOCK2 program from the run menu as described in step 3 for the BLOCK1 program. Before proceeding, you should see "RUN: BLOCK2" on the first line of the display. Run the BLOCK2 program by pressing the "YES" key.

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#### **WASH**

- 6. The washing program required for the P30 run has been named "WASH". The WASH program can be accessed from the **run menu** as described for the **BLOCK1** program above. Before running the **WASH** program, make sure that "RUN: WASH" appears on the first line of the display. Press the "YES" key to run the WASH program. This program will wash/aspirate the plate three times with PBS-casein.
- 7. When finished using the plate washer, remove plate from the washer, replace it with the "test" plate, a dummy wash plate that is provided at each plate washing station.
  - Remove the blue colored tubing from the **WASH** bottle and attach it to the **RINSE** bottle.
  - Prime the machine with deionized water; access in priming command from the main menu as described above.
  - Run the **WASH** program so that the lines are moroughly flushed with deionized water.
- 8. Turn off plate washer. Discard liquids from the **WASH** and **WASTE** bottles and rinse thoroughly with water. Loosely replace covers onto these bottles and leave them near the plate washer.
- 9. When the plate washer sits ide for about 10 min., it will display a "PLEASE RINSE" message. When you are ready to use the washer again, press the "YES" key to return to the main menu.

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#### SAMPLE AND STANDARDS PREPARATION:

Prepare samples and standards on the day of use.

1. Stains/swabs Using the pipette tip/test tube method, extract 2.5 x 2.5 mm samples in 100uL of PBS for 30 minutes at room temperature. Centrifuge. Prepare a 1/25 dilution by adding 20uL sample extract to 480uL wash buffer (PBS-casein).

Cover and refrigerate the remainder of the sample stract until the ELISA is complete and/or ready for amylase analysis.

**2. Standards** Prepare a 2ng/ml standard by first preparite a 10ng/mL solution followed by a 5-fold dilution of this solution as follows:

a. 10 ng/mL: 50uL P30 (1 kg/m**L**) + 5.0mL PBS-casein

b. 2 ng/mL: 1000uL of Nng/mL P30 standard + 4.0mL of PBS-

casein

# SAMPLE AND STANDARD APPLICATION:

1. Have a witness verify the documentation and tube order. Apply PBS, PBS-casein, standards and samples as shown in the per am. Use 100 uL of the appropriate solution per well.

	1	2	3	4	5	6	7	8	9	10	11	12
A	PBS	W	2ng	S2	S4	<b>S</b> 6	S8	S10	S12	S14	S16	S18
В	PBS	W	2ng	S2	S4	<b>S</b> 6	<b>S</b> 8	S10	S12	S14	S16	S18
C	PBS	W	2ng	S2	S4	<b>S</b> 6	<b>S</b> 8	S10	S12	S14	S16	S18
D	PBS	W	2ng	S2	S4	S6	<b>S</b> 8	S10	S12	S14	S16	S18
E	PBS	W	<b>S</b> 1	<b>S</b> 3	S5	<b>S</b> 7	<b>S</b> 9	S11	S13	S15	S17	S19
F	PBS	W	<b>S</b> 1	<b>S</b> 3	S5	<b>S</b> 7	<b>S</b> 9	S11	S13	S15	S17	S19
G	PBS	W	S1	S3	S5	S7	<b>S</b> 9	S11	S13	S15	S17	S19
Н	PBS	W	S1	<b>S</b> 3	S5	S7	<b>S</b> 9	S11	S13	S15	S17	S19

PBS = phosphate buffered saline

W = PBS-casein

S = samples

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- 2. Cover the plates with Parafilm® and incubate at room temperature for 1 hour. Prime/Rinse washer.
- 3. Use the "Wash" program to aspirate contents of wells. Make sure that there is no excess buffer remaining in the wells by gently tapping the plate upside-down on papertowels.
- 4. For each plate, prepare an appropriate dilution of rabbit polyclonal anti-PSA by adding the antiserum to PBS as determined by the QA of the current lot of anti-serum.
- 5. Apply 100 uL of the diluted rabbit anti-human polyclonal anti-PSA to each well of columns 2-12. Apply 100 uL of PBS to each well of column 100 uL of PBS to
- 6. Cover the plates with Parafilm® and incubate the plates from temperature for 1 hour.
- 7. Use the "Wash" program to aspirate contents of yells. Make sure that there is no excess buffer remaining in the wells by gently tapping the plate upside-down on papertowels.
- 8. For each plate, prepare an appropriate di union of the goat anti-rabbit IgG alkaline phosphatase conjugate by adding the anti-serum to PBS as determined by the QA of the current lot of antiserum.
- 9. Apply 100 uL diluted goat anti-rabbit alkaline phosphatase conjugate to each well of columns 2-12. Apply 100 pl. of PBS to each well of column 1.
- 10. Cover the plates with Parafilm® and incubate at room temperature for 1 hour. During the incubation, aliquot 20mL of alkaline substrate buffer (ASB) and allow to come to room temperature, about 30 minutes. For each two plates, dissolve one 20mg p-nitrophenyl phosphate tablet (PNPP) in the 20mL of ASB. Store in a dark place until needed.
- 11. Use the "Wash" program to aspirate contents of wells. Make sure that there is no excess buffer remaining in the wells.
- 12. Apply 100 uL of PNPP substrate solution to each well of columns 1-12.
- 13. Cover the plates with Parafilm and incubate for 1 hour at 37°C. Uncover and read at 405nm.

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#### READING THE PLATE

#### **BioRad Benchmark and 680 XR Plate Reader**

1. Turn on the computer and allow the *Windows* software to boot up. Turn on the Benchmark or 680 XR plate reader and let it warm up. For the 680 XR plate reader, type in the password "00000" and hit the **Enter** soft key to start the program. This will take approximately 3 minutes following an initial self-diagnosis that requires about 1 minute. When the plate reader has warmed up, its screen should read as follows:

Benchmark	680 XR
PLATE READING:	01: Forensic <b>1</b> 30 M405 (2 <b>R</b> 655 (4)
M=2:405, R=4:655	M405 (2005) (4)
Mixing = ON (05s)	Shake. 5 Low
Incu.= OFF	$\mathbf{O}_{\mathbf{A}}^{\mathbf{A}}$

- 2. From this point on, you will be controlling the plate reader from the computer it is attached to.
  - Double click on the Foreign-p30 Microplate Manager 5.2.1 shortcut icon.
  - The screen for the plate readers should now read **Remote Mode**
- 3. A protocol window will appear. The default settings should read as follows:

Reader:	Benchmark	Model 680 XR
Read speed:	N/A	Fast Read
Reading Parymeters:	Dual	Dual
Measurement Filter:	405	405
Reference Filter:	655	655
Dual Wavelength Operation:	Subtract	Subtract
Incubator On:	deselected	deselected
Mix Speed:	N/A	Low
Initial Wait:	0 sec	0 sec
Mix Time:	5 sec	5 sec

4. Place your microtiter plate into the plate reader.

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- 5. Click **Run** located at the upper right corner of the protocol window. The plate analysis takes about 20 seconds.
- 6. When the analysis is finished the **Raw Data** window appears. Save the Raw Data onto the network by doing the following:
  - Go to **File** from the main menu and select **Export**.
  - Change the following:
    - i. Save In: box to read M:\FBIOLOGY\_MAIN\P30FLISA\RAW\_DATA folder.
    - ii. Save As Type: box to Tab delimited (.txt)
    - iii. **File Name** to your plate name.

      Name plates using the date (*MMDDYY*) and plate letter (i.e. *021210A*).
      - The plate letter is written on the top letter of the plate.
  - Click Save.

## **Printing Raw Data:**

- 1. On the **Benchmark** plate reader, press the **START/STOP** soft key once and then the **PAGE**(+) key three times, and then press **ENTER**. The raw data will print on thermal paper using the built in thermal printer. (Note: The raw data can also be printed using an external network printer by going to **File** and selecting **Print**.)Write on the printout your initials, date and plate name
- 2. The **680 XR** plate render does not have a built in thermal printer so the raw data has to be printed using an external network printer by going to **File** and selecting **Print**. Write on the printout your initials, date and plate name.
- 3. When you are finished with the Microplate Manager software, go to **File** in the main menu and select **Exit**. Click **NO** when prompted to save changes to your plate.
- 4. When done, shut down the computer and turn off the plate reader.

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#### P30 result calculations

Refer to the LIMS manual for Forensic Biology for specific procedures within the LIMS system.

- 1. Open the appropriate P30 batch in the LIMS system.
- 2. Select all of the Test Batch samples and navigate to Data Entry for the batch.
- 3. Import the Instrument Data from the appropriate P30 run. Save the Data Entry to trigger the P30 calculations.
- 4. Fill in the Orifice/External values for each sample, and size. The final Interpretation values should populate.
- 5. Select all of the data values and release the data threcord the data into the LIMS system.
- 6. Confirm the QC Batch Parameter data entry for the P30 batch. Save and Release the data to update LIMS system.
- 7. Save the P30 batch to update the entire batch data into the LIMS system.

8.

# **CALCULATION OF P30 ELISA RESULTS**

#### **Manual calculations**

The calculations are done automatically by transferring the data from the microtiter plate reader to the LIMS system to perform them. If necessary, the calculations can be done manually as follows:

- 1. Subtract the mean value for column 1 from each value in all remaining columns.
- 2. Calculate the plate threshold value (PT) of the plate:

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Determine the average (AVE) and standard deviation (standard deviation =  $s = s^{1/2}$ ) (SD) of the values in column 2.

PT value = 2(AVE) + 3(SD)

- 3. Subtract the PT value from the values in columns 3-12.
- 4. For all standards and samples, calculate the average of the duplicate camples for both the MOPC and monoclonal PSA coated wells.
- 5. Subtract the MOPC averages from their corresponding monocloral PSA averages. The remaining value is the P30 ELISA result.

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# P30 ENZYME LINKED IMMUNOSORBENT ASSAY (ELISA) DATE EFFECTIVE APPROVING AUTHORITY PAGE 07-16-2012 QUALITY ASSURANCE MANAGER 13 OF 14

## **Example:**

The 2 ng standard gave a final ELISA value of 0.055.

Sample 1 is in wells A3-D3; sample 2 is in wells E3-H3

<b>Value</b>	<u>Value</u>	
A2 0.114	A3 0.081 } MOPC	. Ca
B2 0.091	B3 0.063 } MOPC	
C2 0.123	C3 0.356 } monoclonal PSA	
D2 0.063	D3 0.325 } monoclonal PSA	
E2 0.081	E3 0.266 } MOPC	
F2 0.085	F3 0.272 } MOPC	NO.
G2 0.085	G3 0.711 } monoclonal PSA	
H2 0.070	H3 0.847 } monoclonal PSA	
		こレ

Average of column 2 values: 0.089

Standard deviation of column 2 values 1.02

**Plate threshold** = 2(0.089) + 3(0.020) = 0.238

Subtract the PT value from each value in column 3; if less than zero, enter zero.

```
A3 0.000
B3 0.000
C3 0.118
D3 0.087
E3 0.028
F3 0.034
average of C3 and D3 = 0.103
G3 0.473
H3 0.609
average of G3 and H3 = 0.541
```

#### **P30 ELISA results:**

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#### INTERPRETATION OF RESULTS

The interpretation of P30 results depends on the source of the sample:

- 1. Body cavity swabs (e.g., oral, vaginal, and anal) are positive if the result is greater than the 2ng standard.
- 2. Samples not from a body cavity (e.g., panties, etc.) are positive if the result is greater than 0.05 absorbance units.
- 3. The location from which a "dried secretion" swab is taken will affect the interpretation. Swabs taken essentially from a body cavity or similar place (e.g., introitus, etc.) are interpreted as if the sample is from a body cavity. Other (ocations (e.g., breast, thigh, etc.) may need to be interpreted differently.
- 4. If P30 results are close to the 2ng level (for blody avity swabs) or the 0.05 level (for other samples), a slide should be prepared from the sample and a sperm search done. A general guideline is that sperm searches should be performed when the P30 values is >70% of the 2ng cutoff.
- 5. Off scale P30 values are indicated by "HIGH" in the results table of the P30 ELISA spreadsheet. All "HIGH" values are interpreted as positive results.



March 24, 2010 – Initial version of procedure.

July 16, 2012 – Specific names of worksheets were removed and replaced with generic terminology to accommodate LIMS. Additionally, procedures concerning "sample and standard application" were revised to allow the use of different dilutions of the antibodies as determined by the quality control tests.

#### P30 ENZYME LINKED IMMUNOSORBENT ASSAY (ELISA) DATE EFFECTIVE APPROVED BY PAGE 03-24-2010 **EUGENE LIEN** 1 OF 15

**Solutions:** Phosphate buffered saline (PBS)

PBS-Casein (PBS with 0.02% w/v casein)

P30 antigen [prostate specific antigen (PSA)] and phosphate buffered saline **Standards:** 

**Plates:** Immulon II microELISA plates (microtiter plate)

**Antibodies:** Mouse monoclonal anti-human PSA (prostate specific antigen, P30)

Rabbit polyclonal anti-human PSA

Goat anti-rabbit IgG alkaline phosphatase conjugate Mouse IgG1, Kappa chain (MOPC 21, mouse myelona) 2012/1/8

Note: Store all antibodies at 4°C.

#### SOLUTION PREPARATION

#### **Phosphate buffered saline (PBS):**

tablets in 1 liter of deionized water. This can be To prepare 1 liter, dissor 1. stored at 4°C for up

On label write init 2. of preparer, the date of make (DOM) and the date of discard (DOI

For each plate or par of plates you will need 1 bottle (1 liter).

# PBS-Casein (PBS with 0.02% w/v Hammerstein casein):

#### If preparing from Frozen Casein Aliquots:

- 1. Thaw casein at room temperature.
- 2. Dilute 20mL of casein stock solution to 1 liter of PBS.

#### If preparing from a Bottle of Liquid Stock Casein (Refrigerated):

- 1. Shake the bottle well.
- 2. Using a graduated cylinder, take 20mL of casein stock solution and dilute it into 1 liter of PBS.

One plate uses approximately 500mL.

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#### PLATE PREPARATION:

#### COATING THE PLATES

- 1. Prepare an appropriate dilution of mouse monoclonal anti-human PSA by adding the antiserum to PBS as determined by the QC of the current lot of antiserum. Each plate requires about 5mL of diluted antiserum. **Always make dilutions in glass, not in plastic.**
- 2. Prepare a 1/8000 dilution of MOPC by adding 10uL MOPC to 80 mL RB9. Always make dilutions in glass, not in plastic. This is enough for 16 plates. For the lates add 2.5uL MOPC to 20mL PBS.
- 3. Coat the plate as shown in the diagram. Use 100uL of the appropriate solution per well.

  PRS = phosphate buffered soling.

PBS = phosphate buffered saline

 $\alpha PSA = mouse monoclonal anti-human PSA$ 

M = MOPC

	1	2	3	4	5	_ 6	7	8	9	10	11	12
A	PBS	M	M	M	<b>C</b> <sub>W</sub> <b>X</b>	M	M	M	M	M	M	M
В	PBS	M	M	М	M	M	M	M	M	M	M	M
C	PBS	αPSA	αPSA	NI NI	αPSA	αPSA	αPSA	αPSA	αPSA	αPSA	αPSA	αΡSΑ
D	PBS	αPSA	αPSA	αPSA	αPSA	αPSA	αPSA	αPSA	αPSA	αPSA	αPSA	αΡSΑ
E	PBS	M	M	M	M	M	M	M	M	M	M	M
F	PBS	M	М	M	M	M	M	M	M	M	M	M
G	PBS	αPSA	αPSA	αPSA	αPSA	αPSA	αPSA	αPSA	αPSA	αPSA	αPSA	αΡSΑ
Н	PBS	αPSA	αPSA	αPSA	αPSA	αPSA	αPSA	αPSA	αPSA	αPSA	αPSA	αΡSΑ

4. Cover plates with Parafilm® and label "Coating" with initials and date. Incubate overnight (approximately 17-20 hours) at 4°C.

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# **BLOCKING THE PLATES:**

- 1. Prime/Rinse washer (See Rinsing and Priming procedure below.)
- 2. Use the "Block 1" program to aspirate contents of well. (This will wash each plate twice and fill the wells with wash buffer) Let the filled plate sit at room temperature for 15-20 minutes (see Block 1 procedure).
- 3. Use the "Block 2" program to aspirate the final wash (See Block 2 procedure).
- 4. Tap plate upside-down on paper towels to remove access liquid
- rapped it 5. Plates that are not used immediately should be wrapped in Frifilm® and stored at 4°C; they can be stored for up to 2 weeks.

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# **Instructions for Bio-Rad 1575 washers**

The Bio-Rad 1575 plate washer has a cover that can be swiveled to an opened or closed position.

# **Rinsing and priming**

- 1. The on/off switch is located on the right-hand side of the lower back of the washer. Turn plate washer on.
- 2. After initialization (concurrent with a quick flash of a Bio-Rad version#), the **main menu** appears and looks like this:



The circles with the triangles above represent the soft keys that you will be pressing to access various functions on the machine. In this menu, soft keys corresponding to "In" and "Out", when pressed instruct the plate washer to move the plate holder in and out of the washer, respectively. Soon after turning on the plate washer, the plate holder automatically sets itself in the out position.

- 3. Fill the **Rinse** bottle with deionized water, close the lid, and attach the tube with the blue colored line to the top of the container. Also, make sure that the stopper connected to the yellow and red colored lines is snugly connected to the **Waste** bottle (if not, the plate washer will not aspirate which can result in flooding of the plate holder).
- 4. Prime the washer with deionized water (from the Rinse bottle). Do this by pressing the very first soft key corresponding to the upward arrow in the main menu. Now you will see the Prime/Rinse menu. Press the soft key corresponding to "YES" and the washer will proceed to prime itself. This step is important to ensure proper vacuum pressure and for the washer to dispense correct volumes. Always make sure to re-prime the washer whenever changing the blue colored tube from the Wash bottle to the Rinse bottle and vice versa.
- 5. Fill the **Wash** bottle with PBS-casein solution. Remove the blue-striped tube from the **Rinse** bottle and attach it to the **Wash** bottle. Repeat the priming procedure 1 time using the PBS-casein solution from the **Wash** bottle. You are now ready to run the plate washer.

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#### **Running plate washer programs**

#### BLOCK 1

- 1. To start the blocking procedure, insert a coated plate into the plate washer lengthwise, so that plate well #A1 is located on the upper right portion of the plate
- 2. Proceed with the first portion of the blocking procedure in which the contents of the wells are aspirated, washed 2 times, and then filled with PBS-casein. The program hat does this is called **BLOCK1**.
- 3. To access the **BLOCK1** program, press the soft key that corresponds to "**YES**" while in the main menu. You will now be in the **run menu**. While in the **un** menu, scroll to the **BLOCK1** program by pressing the soft keys that correspond to the upward and downward arrows.
  - Before proceeding, you should see "FON: BLOCK1" on the first line of the display. Press the "YES" key.
  - You will then see "LAST STKP 12" on the first line of the display. Press the "YES" key again.
  - Usually you will then get one more prompt ("CONNECT THE WASH R9") on the first line of the display. Press the "YES" key once again.

The program will then run on its own, leaving the wells filled at the end of the program.

4. Incubate the plactor 15-20 min. at room temperature as described in the P30 protocol.

#### **BLOCK 2**

5. If you have removed your plate for the incubation step, place plate back into the plate washer by using the "IN" and "OUT" keys from the main menu. The last step of the blocking procedure requires aspiration of the wells. This is achieved by the BLOCK2 program. Access the BLOCK2 program from the run menu as described in step 3 for the BLOCK1 program. Before proceeding, you should see "RUN: BLOCK2" on the first line of the display. Run the BLOCK2 program by pressing the "YES" key.

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#### **WASH**

- 6. The washing program required for the P30 run has been named "WASH". The WASH program can be accessed from the **run menu** as described for the **BLOCK1** program above. Before running the **WASH** program, make sure that "RUN: WASH" appears on the first line of the display. Press the "YES" key to run the WASH program. This program will wash/aspirate the plate three times with PBS-casein.
- 7. When finished using the plate washer, remove plate from the washer, replace it with the "test" plate, a dummy wash plate that is provided at each plate washing station.
  - Remove the blue colored tubing from the **WASH** bottle and attach it to the **RINSE** bottle.
  - Prime the machine with deionized water; access in priming command from the main menu as described above.
  - Run the **WASH** program so that the lines are moroughly flushed with deionized water.
- 8. Turn off plate washer. Discard liquids from the **WASH** and **WASTE** bottles and rinse thoroughly with water. Loosely replace covers onto these bottles and leave them near the plate washer.
- 9. When the plate washer sits ide for about 10 min., it will display a "PLEASE RINSE" message. When you are ready to use the washer again, press the "YES" key to return to the main menu.

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#### SAMPLE AND STANDARDS PREPARATION:

Prepare samples and standards on the day of use.

1. Stains/swabs Using the pipette tip/test tube method, extract 2.5 x 2.5 mm samples in 100uL of PBS for 30 minutes at room temperature. Centrifuge. Prepare a 1/25 dilution by adding 20uL sample extract to 480uL wash buffer (PBS-casein).

Cover and refrigerate the remainder of the sample stract until the ELISA is complete and/or ready for amylase analysis.

**2. Standards** Prepare a 2ng/ml standard by first preparity a 10ng/mL solution followed by a 5-fold dilution of this solution as follows:

a. 10 ng/mL: 50uL P30 (1 kg/m**L**) + 5.0mL PBS-casein

b. 2 ng/mL: 1000uL of Nng/mL P30 standard + 4.0mL of PBS-

casein

# SAMPLE AND STANDARD APPLICATION:

1. Have a witness verify the worksheet and tube order. Apply PBS, PBS-casein, standards and samples as shown in the diagran. Use 100 uL of the appropriate solution per well.

	1	2	y	4	5	6	7	8	9	10	11	12
A	PBS	W	2ng	S2	S4	<b>S</b> 6	S8	S10	S12	S14	S16	S18
В	PBS	W	2ng	S2	S4	S6	S8	S10	S12	S14	S16	S18
C	PBS	W	2ng	S2	S4	S6	S8	S10	S12	S14	S16	S18
D	PBS	W	2ng	S2	S4	S6	S8	S10	S12	S14	S16	S18
E	PBS	W	<b>S</b> 1	<b>S</b> 3	S5	S7	<b>S</b> 9	S11	S13	S15	S17	S19
F	PBS	W	<b>S</b> 1	<b>S</b> 3	S5	S7	<b>S</b> 9	S11	S13	S15	S17	S19
G	PBS	W	<b>S</b> 1	<b>S</b> 3	S5	S7	<b>S</b> 9	S11	S13	S15	S17	S19
Н	PBS	W	S1	<b>S</b> 3	S5	S7	<b>S</b> 9	S11	S13	S15	S17	S19

PBS = phosphate buffered saline

W = PBS-casein

S = samples

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- 2. Cover the plates with Parafilm® and incubate at room temperature for 1 hour. Prime/Rinse washer.
- 3. Use the "Wash" program to aspirate contents of wells. Make sure that there is no excess buffer remaining in the wells by gently tapping the plate upside-down on papertowels.
- 4. For each plate, prepare a 1/1000 dilution of rabbit polyclonal anti-PSA by adding 10 uL antiserum to 10 mL PBS. For 2 plates add 20 uL antiserum to 20 mL PBS.
- 5. Apply 100 uL of the 1/1000 dilution of rabbit anti-human polycronal anti-PSA to each well of columns 2-12. Apply 100 uL of PBS to each well of column 1.
- 6. Cover the plates with Parafilm® and incubate the plates from temperature for 1 hour.
- 7. Use the "Wash" program to aspirate contents of yells. Make sure that there is no excess buffer remaining in the wells by gently tapping the plate upside-down on papertowels.
- 8. For each plate, prepare a 1/3000 dilution of goat anti-rabbit IgG alkaline phosphatase conjugate by adding 4uL antiserum to 12 nL PBS. For 2 plates add 8uL antiserum to 24mL PBS.
- 9. Apply 100 uL of the 1/3000 chlution of goat anti-rabbit alkaline phosphatase conjugate to each well of columns 2-12. Apply 100 uL of PBS to each well of column 1.
- 10. Cover the plates with Parafilm® and incubate at room temperature for 1 hour. During the incubation, aliquot 20mL of alkaline substrate buffer (ASB) and allow to come to room temperature, about 30 minutes. For each two plates, dissolve one 20mg p-nitrophenyl phosphate tablet (PNPP) in the 20mL of ASB. Store in a dark place until needed.
- 11. Use the "Wash" program to aspirate contents of wells. Make sure that there is no excess buffer remaining in the wells.
- 12. Apply 100 uL of PNPP substrate solution to each well of columns 1-12.
- 13. Cover the plates with Parafilm and incubate for 1 hour at 37°C. Uncover and read at 405nm.

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#### READING THE PLATE

### BioRad Benchmark and 680 XR Plate Reader

1. Turn on the computer and allow the *Windows* software to boot up. Turn on the Benchmark or 680 XR plate reader and let it warm up. For the 680 XR plate reader, type in the password "00000" and hit the **Enter** soft key to start the program. This will take approximately 3 minutes following an initial self-diagnosis that requires about 1 minute. When the plate reader has warmed up, its screen should read as follows:

Benchmark	680 XR
PLATE READING:	01: Forensic <b>F</b> 30 M405 (2 <b>R</b> 655 (4)
M=2:405, R=4:655	M405 (2005 (4)
Mixing = ON (05s)	Shake. Low
Incu.= OFF	(0,1)

- 2. From this point on, you will be controlling the plate reader from the computer it is attached to.
  - Double click on the Foreign-p30 Microplate Manager 5.2.1 shortcut icon.
  - The screen for the plate readers should now read **Remote Mode**
- 3. A protocol window will appear. The default settings should read as follows:

Reader:	Benchmark	Model 680 XR
Read speed:	N/A	<b>Fast Read</b>
Reading Parameters:	Dual	Dual
Measurement Filter:	405	405
Reference Filter:	655	655
Dual Wavelength Operation:	Subtract	Subtract
Incubator On:	deselected	deselected
Mix Speed:	N/A	Low
Initial Wait:	0 sec	0 sec
Mix Time:	5 sec	5 sec

4. Place your microtiter plate into the plate reader.

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- 5. Click **Run** located at the upper right corner of the protocol window. The plate analysis takes about 20 seconds.
- 6. When the analysis is finished the **Raw Data** window appears. Save the Raw Data onto the network by doing the following:
  - Go to **File** from the main menu and select **Export**.
  - Change the following:
    - i. Save In: box to read M:\FBIOLOGY\_MAIN\P30FLISA\RAW\_DATA folder.
    - ii. Save As Type: box to Tab delimited (.txt)
    - iii. **File Name** to your plate name.

      Name plates using the date (*MMDDYY*) and plate letter (i.e. *021210A*).

      The plate letter is written on the top part corner of the plate.
  - Click Save.

# **Printing Raw Data:**

- 1. On the **Benchmark** plate reader, press the **START/STOP** soft key once and then the **PAGE**(+) key three times, and then press **ENTER**. The raw data will print on thermal paper using the built in thermal printer. (Note: The raw data can also be printed using an external network printer by going to **File** and selecting **Print**.)Write on the printout your initials, date and plate name
- 2. The **680 XR** plate render does not have a built in thermal printer so the raw data has to be printed using an external network printer by going to **File** and selecting **Print**. Write on the printout your initials, date and plate name.
- 3. When you are finished with the Microplate Manager software, go to **File** in the main menu and select **Exit**. Click **NO** when prompted to save changes to your plate.
- 4. When done, shut down the computer and turn off the plate reader.

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# **Spreadsheet calculations**

- 1. Open the *Microsoft Excel* program by double clicking on the "P30XL" template worksheet in the **M:\FBIOLOGY\_MAIN\P30ELISA\WRKSHEET** folder.
- 2. The P30 spreadsheet will appear (this a **READ-ONLY** file; it cannot be altered). Two lettered A and B will be at the bottom of the page. Using the mouse, click once on the **B** tab.
- 3. The cursor should be at the A1 box position. If not, position the cursor in the A1 block using the mouse (simply click once on the A1 box; if it is not visible, use the scroll bars/buttons on the right and bottom sides of the screen to bring that box into view).
- 4. Pull down the <u>Data</u> menu and move the arrow to the **loport External <u>Data</u>** menu. Click once on **Import <u>Data...</u>**. The **Select Data Sou** ce window will pop up. Click once on the box directly to the right of **Look in:** and then rick once on **M:\FBIOLOGY\_MAIN\P30ELISA\WRNSHEET**. (NOTE: This may be out of view. Simply scroll up using the small scroll bar in the right of the window.)
- 5. You should now be able to see icoms representing the raw data files that were generated by the plate reader. Double click on the file that needs to be examined (or click once to highlight the file and press **FINTER**). This window will now close and the spreadsheet window will remain be open. Press **Finish**.
- 6. The **Import Data** window will pop up. The **Existing Worksheet** option should have a black dot to the **left** and **=\$A\$1** should appear in the window below it. Press **OK**. The saved data from the plate reader should appear on the page in the upper left corner.
- 7. Below the data are a series of bold headings for the following: name, date, plate (number), QC # (for QC purposes only), and the sample series (3E-H, 4A-D, 5E-H, and so on). Using the mouse, click on the box to the immediate right of each heading and enter with the keyboard the appropriate information.
  - **NOTE:** a) The date **MUST** be entered in the following format: **June 21, 20YY** (Y is the last digits of the year)
    - **b)** Plate: refers to the plate number (*i.e.*, A, B, *etc.*)
- 8. Enter the sample series in the following manner: *FBYY-XXXXX vaginal*

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- 9. When all the information has been entered, click once on the **A** tab at the bottom of the page with the mouse. All the information should now be in the pre-made template. Print the page by pulling down the **File** menu and clicking once on **Print**.... When the **PRINT** window appears, click once on the **Print**... button (or press **ENTER**).
- 10. The file should now be saved. Pull down the **File** menu once more. Click once on **Save As...**.
- 11. Click on the **File name:** window and enter the correct filename. The flename should be the date entered as such: 06210XA (where A designates the **first flate**, if more than one and X designates the last digit of the year). Also check that the file is being saved in the proper location; "**Wrksheet**" should be present in the **Save in** window. If not, click on the arrow directly to the right of the **Save in** window. Save file in the "**Wrksheet**" folder via this path:

# M:\FBIOLOGY\_MAIN\P30ELISA\WPASHEET

- 12. To do calculations for a second plate, close the file by pulling down the **File** menu and clicking once on **Close**. Repeat step 2 through 12.
- When you are finished, exit in program by pulling down the **File** menu and click once on **Exit**.

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#### **CALCULATION OF P30 ELISA RESULTS**

# **Manual calculations**

The calculations are done automatically by transferring the data from the microtiter plate reader to an *Microsoft Excel* spreadsheet set up to perform them. If necessary, the calculations can be done manually as follows:

- 1. Subtract the mean value for column 1 from each value in all remaining columns
- 2. Calculate the plate threshold value (PT) of the plate:

Determine the average (AVE) and standard deviation (SD) of the values in column 2.

$$PT \text{ value} = 2(AVE) + 3(SD)$$

- 3. Subtract the PT value from the values in columns 3-12.
- 4. For all standards and samples, calculate the average of the duplicate samples for both the MOPC and monoclonal PSA coated wells.
- 5. Subtract the MOPC averages from their corresponding monoclonal PSA averages. The remaining value is the P30 ELISA result.

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# **Example:**

The 2 ng standard gave a final ELISA value of 0.055.

Sample 1 is in wells A3-D3; sample 2 is in wells E3-H3

<b>Value</b>	<u>Value</u>	
A2 0.114	A3 0.081 } MOPC	. Ca
B2 0.091	B3 0.063 } MOPC	
C2 0.123	C3 0.356 } monoclonal PSA	
D2 0.063	D3 0.325 } monoclonal PSA	
E2 0.081	E3 0.266 } MOPC	
F2 0.085	F3 0.272 } MOPC	<b>N'O</b> *
G2 0.085	G3 0.711 } monoclonal PSA	
H2 0.070	H3 0.847 } monoclonal PSA	
		ンレ

Average of column 2 values: 0.089

Standard deviation of column 2 values 1.52

**Plate threshold** = 2(0.089) + 3(0.020) = 0.238

Subtract the PT value from each value in column 3; if less than zero, enter zero.

```
A3 0.000
B3 0.000
C3 0.118
D3 0.087
E3 0.028
F3 0.034
average of C3 and D3 = 0.103
G3 0.473
H3 0.609
average of G3 and H3 = 0.541
```

# **P30 ELISA results:**

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## INTERPRETATION OF RESULTS

The interpretation of P30 results depends on the source of the sample:

- 1. Body cavity swabs (e.g., oral, vaginal, and anal) are positive if the result is greater than the 2ng standard.
- 2. Samples not from a body cavity (e.g., panties, etc.) are positive if the result is greater than 0.05 absorbance units.
- 3. The location from which a "dried secretion" swab is taken will affect the interpretation. Swabs taken essentially from a body cavity or similar place (e.g., introitus, etc.) are interpreted as if the sample is from a body cavity. Other ocations (e.g., breast, thigh, etc.) may need to be interpreted differently.
- 4. If P30 results are close to the 2ng level (for body avity swabs) or the 0.05 level (for other samples), a slide should be prepared from the sample and a sperm search done. A general guideline is that sperm searches should be performed when the P30 value is >70% of the 2ng cutoff.
- 5. Off scale P30 values are indicated by "HIGH" in the results table of the P30 ELISA spreadsheet. All "HIGH" values are interpreted as positive results.

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# A. Receipt of postmortem specimens

This task should be performed reasonably soon after a batch of samples arrives in the laboratory. The assigned Criminalist I will report to the postmortem (PM) processing supervisor, and perform any and all tasks related to PM processing. Criminalist I's assigned to the Exemplar rotation will be responsible for PM exemplar processing, and witnessing of the PM bloods.

1. Specimens from all five boroughs are delivered to the laboratory in Goled red plastic containers. The LIMS system will automatically update the PM bin's chain of custody once the PM bin's custody has transferred from the Evidence Unit to the Forensic Biology Personnel..

Note: if samples arrive late in the day, inventory readins (Step 2) and store samples in a refrigerator. Samples will be processed the next day.

- 2. To inventory the contents of the red plastic containers proceed with the following:
  - Inventory each container separately (Check for completeness and record any discrepancies. Report any discrepancies to the PM supervisor.)
  - Compare the plastic tags with serial numbers to the serial numbers written on the chain of custody.
  - The person on the intation must record the chain of custody.
  - Scan the included than of custody to a PDF document, and incorporate into the LIMS system. The original is given back to the Evidence Unit.
  - Scan the manifest to a PDF document, and incorporate into the LIMS system. Discard the original in a red biohazard waste container.
  - Sort the manifests by borough and set aside.

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- 3. For discrepancies or problems with the inventory, refer to "Section E: Troubleshooting" and proceed as specified.
- 4. Fill out the PM documentation for each bin. The LIMS system will automatically create the chain of custody for each sample, and record the packaging and processing as the analyst unpacks the postmortem evidence and exemplar samples.
- 5. Ensure that the PM items all have barcode labels and are stored in an appropriate container (See Table 1).

If items are not packaged properly, repackage according to the table below. Seal the package with Evidence Tape or using a heat-sealer for the 4x6" KAPAK<sup>TM</sup> bag, except where indicated. Initial and date all seals. Note: the evidence tape should not obscure the ME # on the barcode label.

# Table 1

Sample	Packaging Packaging
Bloodstain cards	4x6" KAPAK <sup>T</sup> bag (seal KAPAK bag)
Hair, Nails, Trace Evidence*	Coin envelopes placed into 4x6" KAPAK <sup>TM</sup> bag (do not seal KAPAK bag)
Oral, vaginal, anal, penile, and bladdor swabs*	Coin envelopes placed into 4x6" KAPAK <sup>TM</sup> bag (do not seal KAPAK bag)
Bone	Plastic specimen container
Muscle or soft tissue	Plastic specimen container or 15 ml Falcon tube

<sup>\*</sup> Store samples from the same ME # in the same KAPAK bag. Do not seal the bag.

6. Once inventoried and processed, store samples in the appropriate storage area (See Table 2).

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#### Table 2

Room Temperature	Refrigerator (4°C)	Freezer (-20°C)
(20°C)		
- Bloodstain cards	- Oral, vaginal, anal, penile,	- Bone
- Fingernails	and bladder swabs	- Muscle or Soft Tissue
- Hair	- SAK	- Product of conception
- Other Trace Evidence	- Samples in RNAlater®	(PQC)

- 7. Spray the inside of the red bins with disinfectant and let air dry. Set the red containers aside in the designated area for pick up.
- B. Postmortem bloodstain processing (non-vouchered bloods)
- 1. Make the ME barcode labels for the bloodstan cards using the LIMS system. Wear gloves when handling the bloodstain cards. Bandwrite the ME # if unable to generate labels. Initial each bloodstain card prepared.
  - The preparer of the bloodstain taids must initial and date each card.
- 2. The setup of the bloods and bloodstain cards must be witnessed by another laboratory staff member. That person nuss confirm that the order of the blood vials in the rack match the order of the prevated bloodstain cards. The witness will record the witnessing setup in the documentation.
- 3. The bloodstain ords should have the following information prior to processing:
  - a) ME case number (on affixed label or handwritten)
  - b) Initials of the person preparing the stain
  - c) Date the stain card was prepared
  - d) LIMS' stain card ID
- 4. Prepare stains one at a time. Staining of the cards and the opening of liquid blood samples MUST be performed under a biological safety hood with the exhaust fan operating. A new KimWipe<sup>TM</sup> should be used to open each vial stopper. Make sure the blood vial is closed before preparing the next bloodstain card.

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- 5. Use a transfer pipet to make four stains for each bloodstain card, filling in the four circles on each card with blood.
- 6. Re-cap non-vouchered PM blood vials and discard in the plastic biohazard "sharps" container.
- 7. Allow the bloodstain cards to dry overnight in the hood with the exhaust fan running. Document that the stain cards are being stored in the hood.
- 8. Package the air-dried stains into a 4x6" KAPAK<sup>TM</sup> bag. Seal the log with evidence tape or using a heat sealer. Initial and date the seal.
- 9. Organize the bloodstain cards by borough and in ME toder. Add the cards to the appropriate yellow borough bin located on the benefit where they are temporarily stored until a supervisor has had a chance to review the cards. Document the cards' new storage location.
- 10. Bloodstain cards of ME cases that have been assigned FB #'s by a supervisor will be labeled with the FB # and transferred to the red bin on the bench. Cards of ME cases that will not be assigned an FB # are transferred to the blue borough bins. The transfer of cards reviewed by the supervisor are placed to their appropriate long-term storage locations by the assigned Criminalist III on PM Processing:
  - Cards with FB # vare stored numerically by FB # in the designated bloodstain card box.
  - Cards without FB # are stored numerically by borough and ME # in the designated bloodstain card box.

The electronic chain of custody will document the transfer between storage locations and Criminalists.

11. CLEAN THE BIOLOGICAL SAFETY CABINET (refer to Quality Control Procedure #QC125 in the Quality Assurance/Quality Control Manual).

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# C. Assignment of case numbers

This task should be performed by the PM supervisor or trained supervisor.

- 1. Gather all appropriate documentation The daily case census sheets are available electronically through the MEANS system (see Appendix I). The autopsy case worksheets are available electronically through the Document Archiving system (see Appendix II).
- 2. Compare each autopsy case documentation with the manifest and the specimens received to ensure that all of the specimens designated for Forensic Biology have been received. See **Section E. Troubleshooting** if there are discrepancies.
- 3. Screen all the documentation for potential Forensia Bology cases. The following types of cases should be assigned an FB case number.
  - Homicides
  - Any case in which sexual assault evidence (SAK or orifice/penile swabs) has been collected
  - Any case in which a Forensic Biology test is requested via email, phone, or noted on the manifest. Note: Homoglobin, thrombophilia, and sickle cell cases are assigned an MG # and not an FB #. Contact the Molecular Genetics group.
  - Any unknown body with PM samples requiring DNA identification (must verify the victim is still unknown by checking MEANS or the ID Unit)
  - Any case in which evidence from the NYPD or DA's office has been submitted
  - POC/fetus (only if criminal activity is involved)

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- 4. **For cases that will be assigned an FB case number:** Check the database to determine if FB case numbers have been assigned to the ME numbers.
  - a. If the database has an FB # for the ME #, the PM samples will be signed into the pre-existing case numbers.
  - b. If the database does not have an FB # for the ME #, review and assign the PM samples an FB case number. Enter the appropriate information into the database. Create a new case folder by obtaining a manila folder with he FB case number.

Upon electronically assigning a FB # to the ME #, LIMS will create a unique PM number for each specimen.

Exception: For Missing Persons cases (unknown victor) the PM sexual assault evidence (PM SAK or PM orifice/penile swabs) should be pixed on a separate chain of custody from the other PM samples.

- 5. PM SAK and PM orifice/penile swabs must be signed over to the Evidence Unit so that they may be processed. All other specimens must be placed in retained storage. Continue to document the chain of custody for these items to reflect their final location.
- 6. Give the FB cases to the evidence sign-in supervisor.
- 7. All other cases are not as a pared an FB case number. These would include cases where the Manner of Death is:
  - Pending Studies (possible homicides, i.e.- CUPPI, case unknown pending police investigation)
  - Natural
  - Therapeutic Complication
  - Accident/Motor vehicle accidents (MVA's) which are under investigation (i.e.-hit and run)
  - Suicide
  - Undetermined
  - Or any case which involves child abuse or suspected child abuse
- 8. **For cases that will NOT be assigned an FB case number:** File the daily case census sheets and respective autopsy worksheets in chronological order for archival purposes. After 30 days, discard the paperwork. Electronic copies are available through MEANS and DMS.

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# **D.** Discarding postmortem items

Refer to the table below regarding storage and discarding of blood and non-blood items:

**Table 3:** 

	Bloodstain?	Non-Blood?	Discard?
FB cases	Y	Y	Retain all indefinitely.
Non-FB cases	Y	Y	Discard non blood after 6 months, discard bloodstain after 5 years.
	N	Y	Discard non-blood after 5 years.
	Y	N	Discard bloodstain after 5 years.
POC/Fetus (criminal activity)	n/a	h	Retain a small piece and discard the remainder.

A copy of the manifest will be filed in a binder for discarded postmortem samples.

# E. Troubleshooting

Problem	<b>5</b>	Recommended Action

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Problem	Recommended Action
Unlabeled specimen; unscanable label	Criminalist I: For an unlabeled specimen, do not process; record the deviation and notify supervisor. Store questionable samples in designated refrigerated area.  For an unscanable label, process as long as the ME number is legible.  Criminalist III/IV: Nation down possible ME by process of elemenation. Contact ME who performed the autopsy to request an additional sample. If not available, retrieve sample from Department of Toxicology.
Unreadable but scannable barcode label	Criminalist I: Scan barcode and generate new label. Use new label to confirm ME# with manifest and place label on staincard. Continue with processing.
Specimen collected but not listed on manifest	Criminalist I: Record the deviation and continue with processing. Criminalist III/IV: Confirm what samples were collected by the ME who performed the autopsy.
Specimen not collected but listed on manifest	Criminalist I: Record the deviation and notify the supervisor. Criminalist III/IV: Contact ME who performed the autopsy to request an additional sample. If not available retrieve sample from Department of Toxicology.

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Problem	Recommended Action
Blood vial labeled "Hospital Blood" and/or has the ME # written on the hospital label	Criminalist I: Record the deviation continue with processing, and notify supervisor. Criminalist III/IV: Verify on the autopsy worksheet that ME submitted hospital blood. If so, do nothing. If not contact ME who performed the autopsy to inform them of the situation and attempt to retrieve sample in a purple top tube.
Missing manifest	Criminalist I: Record the deviation and continue with processing, and notify supervisor. Criminalist III/IV: Contact the respective borough Deputy ME.
Container not sealed with black ties	Criminalist I: Record the deviation ,continue with processing, and notify supervisor. Criminalist III/IV: Notify Dan Stevelman.
Broken blood vials Blood vial with a detached rubber stopper	Criminalist I: Record the deviationand notify supervisor. Criminalist III/IV: Contact ME who performed the autopsy to request an additional sample. If not available, retrieve sample from Department of Toxicology.
Blood vial with a non-purple stopper	Criminalist I: Record the deviation and continue with processing. Criminalist III/IV: Contact ME who performed the autopsy to inform them of the situation and attempt to retrieve sample in a purple top tube.

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Problem	Recommended Action
Blood that appears to be decomp fluid, grayish in color, or clotted	Criminalist I: Record the deviation and continue with processing, and notify supervisor. For blood clots, smear clot onto the stain card. Discard leftover blood clot properly.  Criminalist III/IV: Contact ME who performed the autopsy and ask for a bone sample.
Blood labeled "decomp" on blood vial or autopsy case worksheet	Criminalist I: Record the deviation, continue with processing, and notify supervisor. Criminalist III/IV: Contact ME who performed the autopsy and ask for a bone sample.
Blood vial labeled for HIV testing (or paperwork for HIV testing included)	Cominalist I: Do not process; Record the deviation and notify supervisor. Store questionable samples in designated refrigerated area.  Criminalist III/IV: Return items to the Manhattan morgue.
RNAlater® samples: liver, spleen, and heart and/or requisition forms	Criminalist I: Do not process; record the deviation and notify supervisor. Place samples in designated refrigerated area. Criminalist III/IV: Notify the Molecular Genetics group to pick up samples and sign Batch Chain.
Incorrect or no sample submitted for decomposed victim or a case for FB	Criminalist III/IV: Contact ME who performed the autopsy and ask for an appropriate sample (long bone, rib, etc.) Retrieve sample from Toxicology as a last resort.

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# F. Civil paternity requests

Do not accept any phone calls from family members. Direct all phone calls to the OCME Legal Department.

- 1. A paternity request is initiated with an email from the Legal Department indicating the family plans to have DNA paternity testing done and to place any specimens on hold.
- 2. Check the PM database to determine the following:
  - A. Was a sample collected?
  - B. What type of PM sample is available (blood, hair, etc.)
  - C. Is this an FB or non-FB case?
  - D. Verify subject's name with autopsy sheet (See Appendix II, Section A for viewing autopsy sheet in DMS).

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- 3. Locate the appropriate PM sample and verify that you have the correct PM sample and subject name. Place PM sample into paternity bin for FB case # assignment.
- 4. Send a "reply to all" email answering all of the questions listed above in #2. List all samples in FB custody. Indicate if there is an inconsistency between the subject's name listed in the email from the Legal Department and what is listed in the autopsy sheet.
- 5. If no sample is available in FB, contact the Toxicology Department a potential sample.

If a sample is available, retrieve it from EU, and process the sample. Store the stain card in the appropriate retained storage to alion. Update all appropriate databases. Retain the email requesting a specimen from the Toxicology Department and your reply. Place PM Sample into the Paternity Bin for FB Case Number assignment.

- 6. FB will be contacted by the Legal Department when a paternity kit has arrived for the subject. Retrieve the kit.
- 7. Locate the appropriate FB case file sample.
- 8. Open kit and discard any glass containers for liquid blood in the sharps container.
- 9. Submit a quarter of the AM sample for testing. If PM sample appears to be decomposed, submit half of the sample. (Example- If four circles are stained, submit one circle. If the bloodstain is decomp fluid, submit two circles.) Do not send the entire sample; a minimum of 50% of the sample should be retained. If the testing laboratory or family is requesting the entire item, verify this with the Legal Department and proceed as advised.
- 10. Submit the portion of stain card in a coin envelope labeled with the subject name, ME #, and any other relevant information. Submit a portion of the tissue or bone sample in a plastic, puncture- and leak-proof container labeled as described previously. Seal, initial, and date packaging. Return unused sample to their original storage location.
- 11. Fill out an OCME autopsy specimen chain of custody documentation and shipping paperwork. Refer to the autopsy sheet for information regarding the subject's age, race, time of death, and medical examiner who performed the autopsy.

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- 12. If requested, have autopsy specimen chain of custody notarized. Consult with the PM Blood Processing Supervisor for a list of Public Notaries within the agency.
- 13. Make copies of the paperwork and save the sender's receipt from the shipping envelope. File the relevant paperwork in the FB file. Update the paternity database.
- 14. Place sample, court order, and other appropriate paperwork in the kit.
- Seal and place kit in appropriate area to be sent. Call the appropriate shipping company to 15. arrange pick-up, as needed. Record the confirmation number in FB file.
- 16.

Email the original contact and inform them that the kit will be picked up. Include the confirmation number. File the email with the relevant paperwork in the FB file.

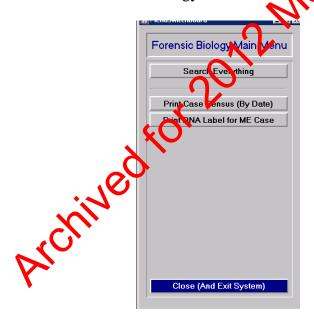
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# **APPENDIX I: DataEase MEANS (Forensic Biology Version)**

<u>Note:</u> A user must obtain access rights from DoITT in order to use MEANS. DoITT will issue the username and password.

# A. Printing Barcode Labels

- 1. Double-click on the MEANS icon on desktop
  - a. Enter login name.
  - b. Enter password.
  - c. Make sure that "CSC" is selected for the field "Ug on to:"
- 2. The MEANS "Forensic Biology Main Menu" (pictured below) will appear:



3. Select "Print DNA label for ME Case". The "Print Barcode Label for ME Case" will appear (pictured below):



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- 4. To print a label, either type in the ME# without spaces or hyphens or scan the barcode from the labels on the manifest sheet.
- 5. Click on "Print Label."
- 6. Click on "OK" to print.
- 7. To print a different label, select "Clear" and repeat steps #4-#6. You must clear the ME# otherwise the previous label will be reprinted.
- 8. Log out as soon as you are done. Failure to log out probables other users from accessing the program (See Section B for logging out).

# **B.** Logging Out of MEANS:

- 1. Select "Close" to exit from each mean open.
- 2. Select "Close (and exit system)" (sictured below) to quit out of MEANS.

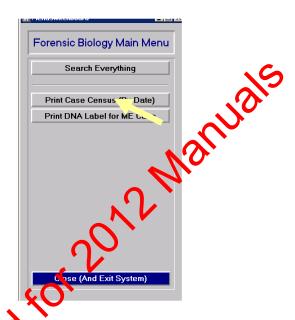
**Note:** Do not use the "x" on the upper-right corner to close out of menus in MEANS.



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# **C.** Printing Daily Case Census Sheets

1. Select "Print Case Census (By Date)" from the Forensic Biology Main Menu (pictured below):



a. The "Print Dill Case Census" screen will appear (pictured below). Enter a "From" of and a "To" date. For the current, select the "Click here for today of the" button.

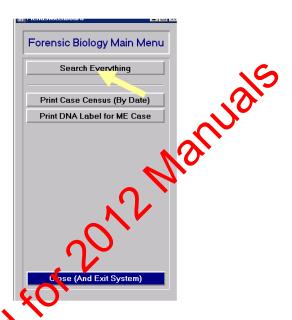


- 2. Select the borough (Brooklyn, Queens, Manhattan, Bronx, or Richmond) by clicking on the desired borough button.
- 3. Select "Print" to run the report.

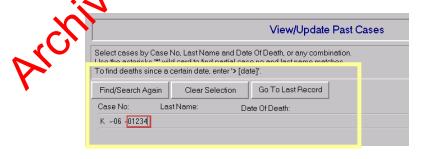
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# D. Resolving Issues Using MEANS

1. Select "Search Everything" from the MEANS Forensic Biology Main Menu (pictured below):



2. On the "View/Update Past Cases" (pictured below), type in the ME # using borough, year, an odigit ME #.



- 3. If the ME # is not available, type in last name and/or date of death in the proper fields.
- 4. Click on "Find/Search again."
- 5. To review more cases, click on "Clear Selection" and repeat steps 2-4.

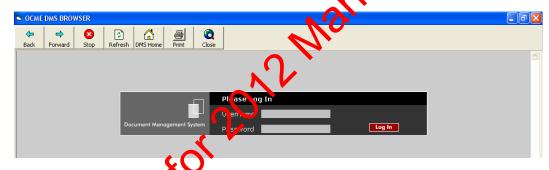
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# APPENDIX II: DMS (Document Imaging and Management System) Browser

<u>Note:</u> A user must obtain access rights from DoITT in order to use MEANS. DoITT will issue the username and password.

# A. Printing out Autopsy Worksheets

- 1. Double click on OCME DMS Browser icon on desktop
  - a. Enter username.
  - b. Enter password.
  - c. Click "Log In" button or hit "Enter."



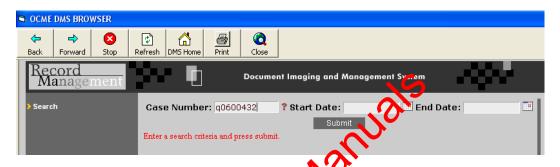
2. The "Document Imaging and Management System" main screen (pictured below) will appear:



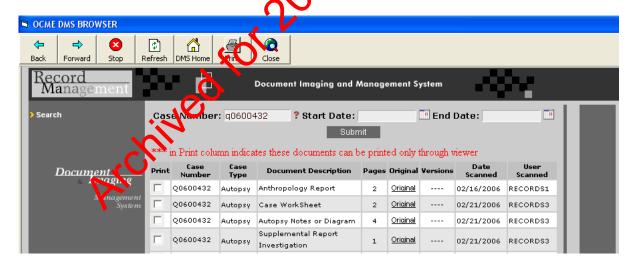
3. Double click on "Record Management".

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4. Enter Medical Examiner case number in the field called "Case Number" in the following format: if the ME # is Q06-00432, enter q0600432 (no hyphens or spaces). See example below:

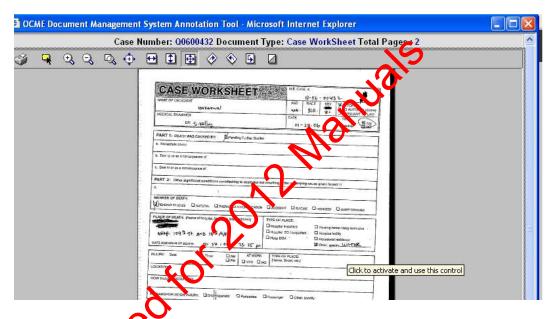


- 5. Click on the "Submit" button (hitting "arter" will not work.)
- 6. The following screen will appear:



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- 7. Look for "Case Worksheet"; this is a scanned .pdf image of the autopsy worksheet.
  - a. To obtain a copy, click on "Original."
  - b. The autopsy worksheet will open up in an Internet Explorer window (pictured below):



- c. Double click on the printer icon to obtain a printed copy of the autopsy worksheer. Click "X" to close the window.
- d. Go tack to the main login screen and double click the red "Log Out" button. Then click the "Close" icon at the top of the screen.

**Revision History:** 

March 24, 2010 – Initial version of procedure.

July 16, 2012 - Specific terminology was removed and replaced with generic terminology to accommodate LIMS.

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# A. Receipt of postmortem specimens

This task should be performed reasonably soon after a batch of samples arrives in the laboratory. The assigned Criminalist I will report to the postmortem (PM) processing supervisor, and perform any and all tasks related to PM processing. Criminalist I's assigned to the Exemplar rotation will be responsible for PM exemplar processing, and witnessing of the PM bloods.

1. Specimens from all five boroughs are delivered to the laboratory in Goled red plastic containers.

Note: if samples arrive late in the day, inventory red bits (Step 2) and store samples in a refrigerator. Samples will be processed the next (Day).

- 2. To inventory the contents of the red plastic containers proceed with the following:
  - Inventory each container separately. (Check for completeness and make a note of any discrepancies. Report any discrepancies to the PM supervisor.)
  - Compare the plastic tags with serial numbers to the serial numbers written on the chain of custody.
  - The person on the rotation must sign the chain of custody.
  - Copy the chain of custody. The copy is kept by the X-team and the original is given back to the Exidence Unit.
  - Make a copy of the manifest for our records and discard the original in a red biohazard waste container.
  - Sort the manifests by borough and set aside.
  - Write down the container number, initial and date the manifest (see below).

	<u>cc</u>	ONTAINER#: 12	-
Container received in Forensic Biology and opened by	<u>MB</u> (Forensi: Biologist Initial:	on <u>3/4/06</u> s) (Date)	-

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- 3. For discrepancies or problems with the inventory, refer to "Section E: Troubleshooting" and proceed as specified.
- 4. Fill out a PM Sample Batch, Blood Processing and Chain of Custody Sheet (referred to as the Batch Chain from hereon) for each borough and list the date received.

Record the ME numbers in numerical order, indicate if blood was received (circle Y or N), use the abbreviations found on the form to list all the other items received in the Additional Samples column, and tally up the total number of samples received (see below).

Borough: Manhattan Date Received: 3/4/06

Item	Medical Examiner #	PM blood received	ADDITIONAL SAMPLES see abbreviations below)	Total # of Samples
1	M06-00163	(Y) N	HH, OS, VS, AS, O - debris	6
2	M06-00172	Y (N	SB	1
3	M06-00184	(Y) N		1

HH:head hair, PH: pubic hair, LF:left hand fingernails, RF:right hand fingernails, SAK: sexual assault kit, OS:oral swabs, VS:vaginal swabs, AS:anal swabs, PS:penile swabs, BS:bladder swabs, ST:soft tissue/muscle, SB:skeletal remains/bar. Sother – specify

4a. Complete the Chain of Custody section of the Batch Chain. If the PM samples are processed on the same day the samples are received and put away the following day, the chain of custody will reflect this (see below):

## **CHAIN OF CUSTODY**

ITEM(S)	RECEIVED FROM	RECEIVED BY	now FB #:	DATE
1-3	sealed red container	M. Shernaw	n/a	3/4/06
1-3	M. Shernaw	Fbiology Storage	n/a	3/4/06
1-3	Fbiology Storage	B. Donner		3/5/06
1-3	B. Donner	Fbiology Storage		3/5/06

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4b. If PM samples are **not processed on the same day the samples are received**, the chain of custody will reflect this (see below):

#### **CHAIN OF CUSTODY**

ITEM(S)	RECEIVED FROM	RECEIVED BY	now FB #:	DATE
1-3	sealed red container	M. Shernaw	n/a	3/4/06
1-3	M. Shernaw	Fbiology Storage	n-a	3/4/06
1-3	Fbiology Storage	B. Donner	\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\	3/5/06
1-3	B. Donner	Fbiology Storage		3/5/06
1-3	Fbiology Storage	B. Domer		3/6/06
1-3	B. Donner	Fbiology Storage		3/6/06

Use the Comments section to document damaged packaging or other unusual observations. (Ex: If a sample was not received in a purple top tube, list affected ME # and the color top tube that had been used.)

5. Enter the inventoried PM samples from the Batch Chains into the Post-Mortem Inventory spreadsheet. This step can be performed before or after PM samples are processed. Go to M:\FBIOLOGY\_MAIN\_\(\sqrt{PERSONS\PM'S\PM'S\DB}\). Select the appropriate worksheet tab based on the year. Important: enter data after the last entry (start from the bottom), do not sow the database, and always remember to save.

Enter the MC# date samples received, "blood" or "no blood", and additional samples (using abbre liations listed on the Batch Chain). Do not fill out the other categories unless instructed (see below).

	3.577//	Date Received			
FB#	ME#	by FB	Blood	Addtl. PM Sample (use codes)	Location
	B06-00001	1/3/2006	blood	SB (bone)	
	B06-00003	1/3/2006	blood	PH, HH	

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- 6. Staple the Batch Chains to the respective manifest and put aside until processing.
- 7. Ensure that the PM items all have barcode labels and are stored in an appropriate container (See Table 1).

If items are not packaged properly, repackage according to the table below. Seal the package with Evidence Tape or using a heat-sealer for the 4x6" KAPAK<sup>TM</sup> bag, except where indicated. Initial and date all seals. Note: the evidence tape should not obscure the ME # on the barcode label.

## Table 1

Sample	Packaging
Bloodstain cards	4x6" KAPAK™ bag (sear KAPAK bag)
Hair, Nails, Trace	Coin envelopes placed into 4x6" KAPAK <sup>TM</sup> bag (do not
Evidence*	seal KAPAK bag)
Oral, vaginal, anal,	Coin envelopes placed into 4x6" KAPAK <sup>TM</sup> bag (do not
penile, and bladder	seal KA(AK bag)
swabs*	-87
Bone	Pastic specimen container
Muscle or soft tissue	Plastic specimen container or 15 ml Falcon tube

<sup>\*</sup> Store samples from the same ME # in the same KAPAK bag. Do not seal the bag.

8. Once inventoried and processed, store samples in the appropriate storage area (See Table 2).

### Table 2

Room Temperature (20°C)	Refrigerator (4°C)	Freezer (-20°C)
- Bloodstain cards - Fingernails	- Oral, vaginal, anal, penile, and bladder swabs	- Bone - Muscle or Soft Tissue
<ul><li> Hair</li><li> Other Trace Evidence</li></ul>	- SAK - Samples in RNAlater®	- Product of conception (POC)

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9. Spray the inside of the red bins with disinfectant and let air dry. Place a blank manifest into each container. Set the red containers aside in the designated area for pick up.

## **B.** Postmortem bloodstain processing (non-vouchered bloods)

- 1. Make the ME barcode labels for the bloodstain cards using the MEANS program (see Appendix I-A). Wear gloves when handling the bloodstain cards. Handwrite the ME # if unable to generate labels. Initial each bloodstain card prepared.
- 2. Fill out the Blood Processing section of the Batch Chain (see below). Make sure that only those item numbers for which blood was received are listed here. The preparer of the bloodstain cards must initial and date each card. Leave the Witnessed by field blank.

Blood processing:	
Stain cards from the above blo	ods # 1 were made on Date: 3/4/06
By analyst: M. Shernaw	Witnested by:
•	

- 3. The setup of the bloods and proodstain cards must be witnessed by another laboratory staff member. That person pust confirm that the order of the blood vials in the rack match the order of the prepared bloodstain cards. The witness will then initial the bloodstain cards, and initial the Witnessed By field of the Batch Chain (see Figure 6).
- 4. The bloodstain cards should have the following information prior to processing:
  - a) ME ase number (on affixed label or handwritten)
  - b) Initials of the person preparing the stain
  - c) Initial of the witness
  - d) Date
- 5. Prepare stains one at a time. Staining of the cards and the opening of liquid blood samples MUST be performed under a biological safety hood with the exhaust fan operating. A new KimWipe<sup>TM</sup> should be used to open each vial stopper. Make sure the blood vial is closed before preparing the next bloodstain card.

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- 6. Use a transfer pipet to make four stains for each bloodstain card, filling in the four circles on each card with blood.
- 7. Re-cap non-vouchered PM blood vials and discard in the plastic biohazard "sharps" container.
- 8. Allow the bloodstain cards to dry overnight in the hood with the exhaust fan running.
- 9. Package the air-dried stains into a 4x6" KAPAK<sup>TM</sup> bag. Seal the bag with evidence tape or using a heat sealer. Initial and date the seal.
- 10. Organize the bloodstain cards by borough and in ME # order. Add the cards to the appropriate yellow borough bin located on the bench there they are temporarily stored until a supervisor has had a chance to review the cards.
- 11. Bloodstain cards of ME cases that have been assigned FB #'s by a supervisor will be labeled with the FB # and transferred to the red bin on the bench. Cards of ME cases that will not be assigned an FB # are transferred to the blue borough bins. The transfer of cards reviewed by the supervisor are placed to their appropriate long-term storage locations by the assigned Criminalist III on PM Processing:
  - Cards with FB #'s are tored numerically by FB # in the designated bloodstain card box.
  - Cards without NB # are stored numerically by borough and ME # in the designated bloodstain card box.
- 12. CLEAN THE BIOLOGICAL SAFETY CABINET (refer to Quality Control Procedure #QC125 in the Quality Assurance/Quality Control Manual).

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## C. Assignment of case numbers

This task should be performed by the PM supervisor or trained supervisor.

- 1. Retrieve the daily case census sheets and autopsy case worksheets for all five boroughs. The paperwork is typically sent in envelopes addressed to the "Serology Department" except for loose paperwork retrieved from the ID Unit. The daily case census sheets are available electronically through the MEANS system (see Appendix 1). The autopsy case worksheets are available electronically through the Document Archiving system (see Appendix II).
- 2. Organize the daily case census sheets in chronological order. For each borough the daily case census sheet is a list of the autopsies performed by the ME for that day. Attach the respective autopsy worksheets to the daily case centre sheets. Compare each autopsy case worksheet with the manifest and the specimens received to ensure that all of the specimens designated for Forensic Biology have been received. See Section E. Troubleshooting if there are discrepancie.
- 3. Screen all the paperwork for potential Forensic Biology cases. The following types of cases should be assigned an FB case number:
  - Homicides
  - Any case in which saval assault evidence (SAK or orifice/penile swabs) has been collected
  - Any case in which a Forensic Biology test is requested via email, phone, or noted on the manifest. Note: Hemoglobin, thrombophilia, and sickle cell cases are assigned an MG # and not an FB #. Contact the Molecular Genetics group.
  - Any variown body with PM samples requiring DNA identification (must verify the victim is still unknown by checking MEANS or the ID Unit)
  - Any case in which evidence from the NYPD or DA's office has been submitted
  - POC/fetus (only if criminal activity is involved)

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- 4. **For cases that will be assigned an FB case number:** Separate the autopsy worksheets from the others. Check the Access database to determine if FB case numbers have been assigned to the ME numbers.
  - a. If Access has an FB # for the ME #, the PM samples will be signed into the preexisting case numbers. Retrieve any existing case folders.
  - b. If Access does not have an FB # for the ME #, assign the PM samples the next available FB case number. Enter the appropriate information into the FB logbook and Access. Create a new case folder by obtaining a manife folder with the FB case number.
- 5. Fill out a chain of custody for the PM sample(s). Numerically list each specimen in the evidence submitted section. Give each specimen a bridge PM number (i.e.- blood is usually designated "PM 1" and sexual assault line are usually designated "PM 2"). In the chain of custody section, enter the date the specimens were received from Forensic Biology storage.

Exception: For Missing Persons cases (unknown victim), the PM sexual assault evidence (PM SAK or PM orifice/penile swats) should be placed on a separate chain of custody from the other PM samples.

- 6. On the corresponding Batch Chain, indicate which ME # was signed over to an FB file. Make a copy of the Batch Chain and place it behind the chain of custody in the case file. Both should be affired to the left side of the folder.
- 7. Autopsy case worksheets are affixed to the right side of the folder.
- 8. PM SAK and PM orifice/penile swabs must be signed over to the Evidence Unit so that they may be processed. All other specimens must be placed in retained storage. Continue the chain of custody for these items to reflect their final location.
- 9. Give the FB cases to the evidence sign-in supervisor.

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- 10. All other cases are not assigned an FB case number. These would include cases where the Manner of Death is:
  - Pending Studies (possible homicides, i.e.- CUPPI, case unknown pending police investigation)
  - Natural
  - Therapeutic Complication
  - Accident/Motor vehicle accidents (MVA's) which are under investigation (i.e.-hit and run)
  - Suicide
  - Undetermined
  - Or any case which involves child abuse or suspected child abuse
- 11. **For cases that will NOT be assigned an FB case rember:** File the daily case census sheets and respective autopsy worksheets in chronological order for archival purposes. After 30 days, discard the paperwork. Electronic copies are available through MEANS and DMS.
- D. Discarding postmortem items

Refer to the table below regarding storage and discarding of blood and non-blood items:

**Table 3:** 

.()	Bloodstain?	Non-Blood?	Discard?
FB cases	Y	Y	Retain all indefinitely.
Non-FB cases	Y	Y	Discard non-blood after 6 months; discard bloodstain after 5 years.
	N	Y	Discard non-blood after 5 years.
	Y	N	Discard bloodstain after 5 years.
POC/Fetus (criminal activity)	n/a	Y	Retain a small piece and discard the remainder.

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Prepare a manifest for discarding postmortem samples. The manifest must reflect:

- ME #
- Samples that are discarded
- Date of discard
- The individual discarding the samples.

A copy of the manifest will be filed with Batch Chain for the sample being discarded. The original manifest will be filed in a binder for discarded postmorem samples.

# E. Troubleshooting

Problem	Recommended Action
Unlabeled specimen; unscanable label	Crimina is I: For an unlabeled specimen, do not process; make a note on the Batch Chain and notify supervisor. Store questionable samples in designated relargerated area.
Archivedio	For an unscanable label, process as long as the ME number is legible.  Criminalist III/IV: Narrow down possible ME by process of elimination. Contact ME who performed the autopsy to request an additional sample. If not available, retrieve sample from Department of Toxicology.
Unreadable but scannable barcode label	Criminalist I: Scan barcode and generate new label in MEANS. Use new label to confirm ME# with manifest and place label on staincard. Continue with processing.

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Problem	Recommended Action
Specimen collected but not listed on manifest	Criminalist I: Make a note on the Batch Chain and continue with processing. Criminalist III/IV: Confirm what samples were collected by the ME who performed the autopsy.
Specimen not collected but listed on manifest	Criminalist I: Make a note on the Batch Chain and notify the supervisor. Criminalist III/IV: Contact ME who performed the autopsy to request an additional sample. If not available retrieve sample from Repartment of Toxicology.
Blood vial labeled "Hospital Blood" and/or has the ME # written on the hospital label	Criminalist I: Make a note on the Batch Chain continue with processing, and notify subervisor. Criminalist III/IV: Verify on the autopsy worksheet that ME submitted hospital blood. If so, do nothing. If not, contact ME who performed the autopsy to inform them of the situation and attempt to retrieve sample in a purple top tube.
Missing manifest	Criminalist I: Handwrite manifest, make a note on the batch chain of custody, continue with processing, and notify supervisor. Criminalist III/IV: Contact the respective borough Deputy ME.
Container not sealed with black ties	Criminalist I: Make a note on the Batch Chain, continue with processing, and notify supervisor. Criminalist III/IV: Notify Dan Stevelman.

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Problem	Recommended Action
Broken blood vials/ Blood vial with a detached rubber stopper	Criminalist I: Make a note on the Batch Chain and notify supervisor. Criminalist III/IV: Contact ME who performed the autopsy to request an additional sample. If not available, retrieve sample from Department of Toxicology.
Blood vial with a non-purple stopper	Criminalist I: Make a note on the Batch Chain and continue with processing. Criminalist III/IV: Contact ME who performed the rutorsy to inform them of the situation and attempt to retrieve sample in a purple top tube.
Blood that appears to be decomp fluid, grayish in color, or clotted	Criminalist I: Make a note on the Batch Cham, continue with processing, and notify supervisor. For blood clots, smear clot onto the stain card. Discard leftover blood clot properly.  Criminalist III/IV: Contact ME who performed the autopsy and ask for a bone sample.
Blood labeled "decomp" on blood vial or autops case worksheet	Criminalist I: Make a note on the Batch Chain, continue with processing, and notify supervisor. Criminalist III/IV: Contact ME who performed the autopsy and ask for a bone sample.
Blood vial labeled for HIV testing (or paperwork for HIV testing included)	Criminalist I: Do not process; make a note on the Batch Chain and notify supervisor. Store questionable samples in designated refrigerated area. Criminalist III/IV: Return items to the Manhattan morgue.

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Problem	Recommended Action
RNAlater® samples: liver, spleen, and heart and/or requisition forms	Criminalist I: Do not process; make a note on the Batch Chain and notify supervisor. Place samples in designated refrigerated area. Criminalist III/IV: Notify the Molecular Genetics group to pick up samples and sign Batch Chain.
Incorrect or no sample submitted for decomposed victim or a case for FB	Criminalist III/IV: Contact ME who performed the autopsy and ask for an appropriate sample from Bone, rib, etc.) Retrieve sample from Toxicology as a last resort.

## F. Civil paternity requests

Do not accept any phone calls from family members. Direct all phone calls to the Legal Department.

- 1. A paternity request is init and with an email from the Legal Department indicating the family plans to have DNA paternity testing done and to place any specimens on hold.
- 2. Check the PM database to determine the following:
  - A. Was a sample collected?
  - B. What you of PM sample is available (blood, hair, etc.)?
  - C. Is this an FB or non-FB case?
  - D. Verify subject's name with autopsy sheet (See Appendix II, Section A for viewing autopsy sheet in DMS).
- 3. Locate the appropriate PM sample and verify that you have the correct PM sample and subject name. Place PM sample into paternity bin for FB case # assignment.
- 4. Send a "reply to all" email answering all of the questions listed above in #2. List all samples in FB custody. Indicate if there is an inconsistency between the subject's name listed in the email from the Legal Department and what is listed in the autopsy sheet.

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5. If no sample is available in FB, contact the Toxicology Department for a potential sample.

If a sample is available, retrieve it, make a chain of custody, and process the sample.

Store the stain card in the appropriate retained storage location. Update the PM \_DB and the paternity database. Retain the email requesting a specimen from the Toxicology Department and your reply. Place PM Sample into the Paternity Bin for FB Case Number assignment.

- 6. FB will be contacted by the Legal Department when a paternity kit has arrived for the subject. Retrieve the kit.
- 7. Locate the appropriate FB case file & sample.
- 8. Open kit and discard any glass containers for Niquid blood in the sharps container.
- 9. Sign the Chain of Custody found in the Belle when retrieving samples from FB Storage.
- 10. Submit a quarter of the PM sample for testing. If PM sample appears to be decomposed, submit half of the sample. (Example- If four circles are stained, submit one circle. If the bloodstain is decomp fluid, adamit two circles.) Do not send the entire sample; a minimum of 50% of the sample should be retained. If the testing laboratory or family is requesting the entire item, verify this with the Legal Department and proceed as advised.
- 11. Submit the portion of stain card in a coin envelope labeled with the subject name, ME #, and any other relevant information. Submit a portion of the tissue or bone sample in a plastic, puncture- and leak-proof container labeled as described previously. Seal, initial, and date packaging. Return unused sample to their original storage location and sign the Forensic Biology Chain of Custody.
- 12. Fill out an OCME autopsy specimen chain of custody documentation (found at M:\FBIOLOGY\_MAIN\FORMS\PM\PrivateLabCOC.doc) and shipping paperwork. Refer to the autopsy sheet for information regarding the subject's age, race, time of death, and medical examiner who performed the autopsy (Refer to DMS for a scanned image of the autopsy sheet).

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- 13. If requested, have autopsy specimen chain of custody notarized. Consult with the PM Blood Processing Supervisor for a list of Public Notaries within the agency.
- 14. Make copies of the paperwork and save the sender's receipt from the shipping envelope. File the relevant paperwork in the FB file. Update the paternity database.
- 15. Place sample, court order, and other appropriate paperwork in the kit.
- 16. Seal and place kit in appropriate area to be sent. Call the appropriate shipping company to arrange pick-up, as needed. Record the confirmation number in FB file.
- 17.

Email the original contact and inform them that the kit will be picked up. Include the confirmation number. File the email with the relevant paperwork in the FB file.

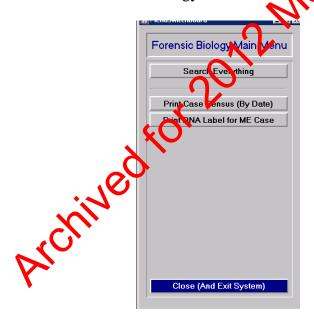
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### **APPENDIX I: DataEase MEANS (Forensic Biology Version)**

**Note:** A user must obtain access rights from DoITT in order to use MEANS. DoITT will issue the username and password.

## A. Printing Barcode Labels

- 1. Double-click on the MEANS icon on desktop
  - a. Enter login name.
  - b. Enter password.
  - c. Make sure that "CSC" is selected for the field "Ug on to:"
- 2. The MEANS "Forensic Biology Main Menu" (pictured below) will appear:



3. Select "Print DNA label for ME Case". The "Print Barcode Label for ME Case" will appear (pictured below):



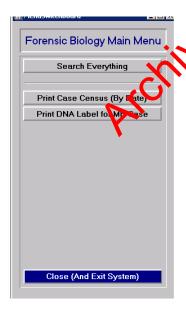
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- 4. To print a label, either type in the ME# without spaces or hyphens or scan the barcode from the labels on the manifest sheet.
- 5. Click on "Print Label."
- 6. Click on "OK" to print.
- 7. To print a different label, select "Clear" and repeat steps #4-#6. You must clear the ME# otherwise the previous label will be reprinted.
- 8. Log out as soon as you are done. Failure to log out probables other users from accessing the program (See Section B for logging out).

## **B.** Logging Out of MEANS:

- 1. Select "Close" to exit from each mean open.
- 2. Select "Close (and exit system)" victured below) to quit out of MEANS.

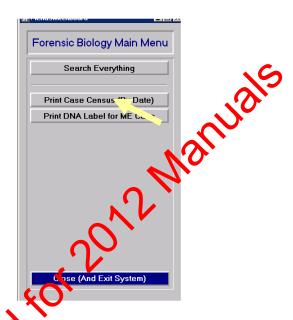
Note: Do not use the "x" on the upper-right corner to close out of menus in MEANS.



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## C. Printing Daily Case Census Sheets

1. Select "Print Case Census (By Date)" from the Forensic Biology Main Menu (pictured below):



a. The "Print Dill Case Census" screen will appear (pictured below). Enter a "From" of and a "To" date. For the current, select the "Click here for today's the" button.

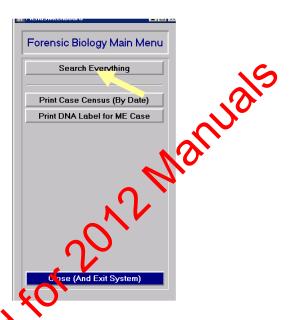


- 2. Select the borough (Brooklyn, Queens, Manhattan, Bronx, or Richmond) by clicking on the desired borough button.
- 3. Select "Print" to run the report.

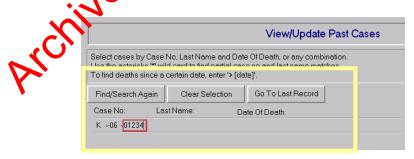
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## D. Resolving Issues Using MEANS

1. Select "Search Everything" from the MEANS Forensic Biology Main Menu (pictured below):



2. On the "View/Update Past Cases" (pictured below), type in the ME # using borough, year, an odigit ME #.



- 3. If the ME # is not available, type in last name and/or date of death in the proper fields.
- 4. Click on "Find/Search again."
- 5. To review more cases, click on "Clear Selection" and repeat steps 2-4.

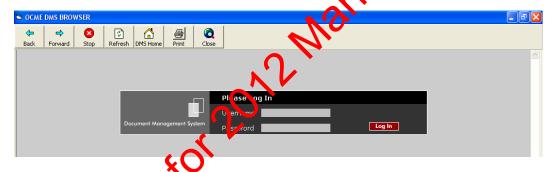
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## APPENDIX II: DMS (Document Imaging and Management System) Browser

<u>Note:</u> A user must obtain access rights from DoITT in order to use MEANS. DoITT will issue the username and password.

## A. Printing out Autopsy Worksheets

- 1. Double click on OCME DMS Browser icon on desktop
  - a. Enter username.
  - b. Enter password.
  - c. Click "Log In" button or hit "Enter."



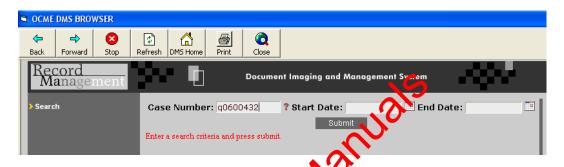
2. The "Document Imaging and Management System" main screen (pictured below) will appear:



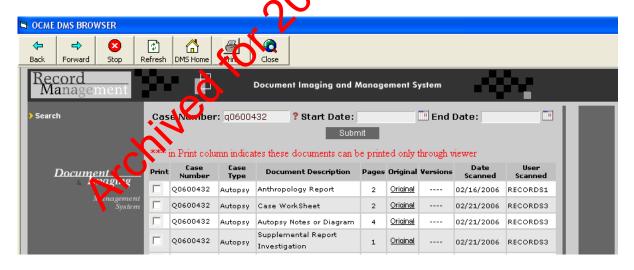
3. Double click on "Record Management".

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4. Enter Medical Examiner case number in the field called "Case Number" in the following format: if the ME # is Q06-00432, enter q0600432 (no hyphens or spaces). See example below:

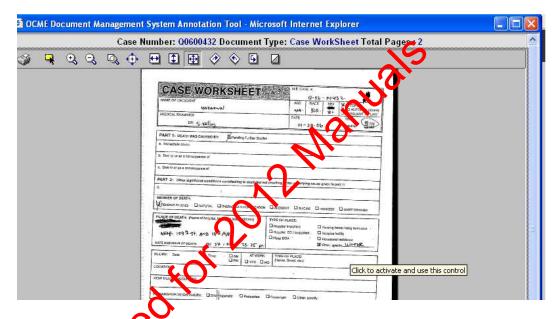


- 5. Click on the "Submit" button (hitting "mer" will not work.)
- 6. The following screen will appear:



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- 7. Look for "Case Worksheet"; this is a scanned .pdf image of the autopsy worksheet.
  - a. To obtain a copy, click on "Original."
  - b. The autopsy worksheet will open up in an Internet Explorer window (pictured below):



- c. Double click on the printer icon to obtain a printed copy of the autopsy worksheer. Click "X" to close the window.
- d. Go tack to the main login screen and double click the red "Log Out" button. Then click the "Close" icon at the top of the screen.

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SLIDE PREPARATION FOR SPERMATOZOA SEARCHES				
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There are two methods to prepare slides for spermatozoa searches. Either may be used:

## 1. Mashing

- A. Cut 1.0 x 1.0 mm of the sample and place it on a clean microscope slide.
- B. Add a drop of distilled water.
- C. Tweeze apart sample until fibers are in a thin even layer across the slide
- D. Fix sample to the slide by heating on a hot-plate (approximately 5 to 10 seconds).
- E. Stain slide using the Christmas Tree Staining volcure.

## 2. Pipette Tip/Test Tube Extraction

- A. Using the pipette tip/test tube me hod, extract 1.5 x 1.5 mm samples in 50uL of distilled water for 30 minutes at room temperature.
- B. Centrifuge sample for 2 whates.
- C. Pipette pellet onto pricroscope slide.
- D. Fix sample to the slide by heating on a hot-plate (approximately 5 to 10 seconds).
- E. Stain slide using the <u>Christmas Tree Staining procedure</u>.

Revision History:

September 17, 2012 – Initial version of procedure.