

UK BIOBANK

# Biospecimens Manual

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Collection of biological samples, processing and storage

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This manual details the collection of biological samples at UK Biobank Assessment Centres and the subsequent processing of those samples at the UK Biobank Central Processing Centre.

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## **1 Introduction**

1.1 This manual details the process for collection, processing and analysis of biological samples. The samples are taken at the 7<sup>th</sup> station of the assessment centre visit, shipped and then analysed at UK Biobank central processing centre.

The following samples were collected at the assessment centres before being processed at the UK Biobank central processing site:

1. Ethylenediaminetetraacetic acid, EDTA (10ml), Purple cap
2. Ethylenediaminetetraacetic acid, EDTA (10ml), Clear cap
3. Plasma Separator, PST (10ml), Green cap
4. Serum Separator, SST (10ml), Orange cap
5. Urine (10ml), Grey cap
6. Acid Citrate Dextrose, ACD (6ml), Pale yellow cap
7. Ethylenediaminetetraacetic acid, EDTA (4ml), Purple cap
8. Applied Biosystems Tempus Tube, RNA Stabilised Blood (3ml), Blue cap
9. 50ml Falcon Tube, Saliva

**Note:** Tubes 8 and 9 were an enhancement to the study and added for the last 100,000 participants.

Please see below for the sequence of the assessment centre visit.

Table 1: sequence of assessment visit

	<b>Visit Station</b>	<b>Assessments Undertaken</b>
<b>1</b>	Reception	<ul style="list-style-type: none"><li>• Welcome &amp; registration</li><li>• Generating a USB key for Participants</li></ul>
<b>2</b>	Touch-screen Section	<ul style="list-style-type: none"><li>• Consent</li><li>• Touch-screen questionnaire</li><li>• Hearing Test</li><li>• Cognitive function tests (Shape, Pairs, Fluid Intelligence, Snap)</li></ul>
<b>3</b>	Interview & blood pressure	<ul style="list-style-type: none"><li>• Interviewer questionnaire</li><li>• Blood pressure measurement</li><li>• Measurement of arterial stiffness</li></ul>
<b>4</b>	Eye measurements	<ul style="list-style-type: none"><li>• Visual acuity</li><li>• Auto-refraction</li><li>• Intraocular pressure</li><li>• Retinal image (OCT Scan)</li></ul>
<b>5</b>	Physical measurements	<ul style="list-style-type: none"><li>• Height (Standing and Sitting)</li><li>• Hip &amp; waist measurement</li><li>• Weight and Bio-impedance measurement</li><li>• Hand-grip strength</li><li>• Heel-bone ultrasound</li><li>• Spirometry (Lung function Test)</li></ul>

<b>6</b>	Cardio (Physical fitness)	• Exercise ECG (Cycling)
<b>7</b>	Sample collection & exit	• Blood samples collected • Urine sample sought • Saliva sample sought • Consent & result summary printed • Travel expense claim provided
<b>8</b>	Web-based diet questionnaire	• Dietary assessment

**1.2** Throughout this document, the term “Participant” signifies a study participant who is taking part in the Assessment Centre process, regardless of whether they eventually give or withhold consent to take part in the UK Biobank study.

**1.3** The collection of data from assessment visits uses the direct data entry system of the Assessment Centre Environment (ACE).

**1.4** At the start of their visit, each participant is issued with a USB Key at the Reception station. This USB Key acts as a participant identifier (it contains Participant ID, name, date of birth and gender) and as a temporary storage device for the recorded data. As the participant progresses between stations, the USB key acts as an identifying token and also as a data transfer mechanism. At the Reception & Exit module, all data on the USB key is removed, after it has been backed up to the Assessment Centre head PC.

## **2 Staff**

Blood draws are carried out by registered phlebotomists, trained and certified to conduct blood draws. The Assessment Centre Manager oversees that all staff work in accordance with the procedure.

## **3 Collection of blood samples into Vacutainers®**

Below is the protocol for blood sample collection in all Becton Dickinson Vacutainer® formats.

**3.1** Hold Vacutainer® barrel steady with one hand and select, with other hand, first Vacutainer® tube in sequence (purple cap 10ml EDTA tube) from pre-prepared rack. Push Vacutainer® tube into barrel until sharp end of needle within barrel pierces rubber bung of tube;

**3.2** Release tourniquet as blood begins to flow into Vacutainer® tube (most participants will not need tourniquet after access to vein has been established), but keep some pressure/tightness on tourniquet if flow is slow;

**3.3** When Vacutainer® 1 is full, blood flow will stop. Remove the Vacutainer® tube from the barrel

And insert Vactutainer® 2. Gently invert Vacutainer® 1 ten times while the second bottle is

filling then return it to rack.

**Note:** If you are unable to invert the Vacutainer® while the other bottle is filling then all bottles should be inverted immediately upon completion of blood sampling.

**3.4** Repeat this step to fill Vacutainers® 2-8 according to order in rack: pay particular attention to the gentle inversion of the tubes (1 to 6) after each is filled with blood.

**Note:** It is crucial that the Blue RNA tube must be shaken vigorously for 20 seconds immediately after sample is obtained.

1. Purple, EDTA (10ml)
2. Clear, EDTA (10ml)
3. Green, Plasma Separator (10ml)
4. Orange, Serum Separator (10ml)
5. Grey, Urine (10ml)
6. Pale yellow, ACD (6ml)
7. Purple, EDTA (4ml)
8. Blue, RNA Stabilised Blood (3ml)
9. 50ml Falcon Tube, Saliva

**3.5** If blood flow slows during collection, re-apply tourniquet and ask participant to clench and unclench their fist while keeping arm still. When blood flow starts again, release tourniquet and continue collection;

**Note:** If blood flow does not re-start, explain difficulties to participant and ask permission to repeat venepuncture in other forearm or, failing that, to collect sample from veins in back of hand using a Safety Lok butterfly needle (as described in Section 8.4 above);

**3.6** After blood collection is complete (with either all required Vacutainers® filled or as many as possible), apply clean cotton wool dressing over skin at insertion of needle. Remove needle and apply pressure on puncture site (applying pressure before needle removal will cause discomfort);

**3.7** Immediately scan each filled Vacutainer® using barcode reader. It is very important to check on the computer screen below that each filled tube has been properly scanned as this links these samples to the participant. If the scanner fails to recognise a barcode, the tube ID should be manually entered using the 13-digit tube ID;

**3.8** If one or more tubes are not collected, record the reason why not using the drop-down menus for each specific tube (see below); empty blood tubes do not need to be scanned.

**3.9** When all of the filled tubes have been scanned a message is sent to the sample processing station and activates a timer that ensures that tube 3 [orange cap] is allowed to stand for 30 minutes prior to centrifugation;

**3.10** A check is made to ensure that all tubes have been collected (or reason(s) for not being collected recorded).

**3.11** Sample collection is complete.

## **4 Sample storage and shipping**

All samples are stored immediately in a refrigerator at the clinic. At the end of the day, all samples are shipped at +4°C to the centralised processing centre in Stockport, Cheshire. The one exception is the ACD sample which is transported at 18°C.

## **5 Sample processing**

The samples were processed at the UK Biobank central processing site as follows:

### **5.1 EDTA (Sample #1 and Sample #2)**

EDTA samples were processed on an automated system following centrifugation. The automated system retrieved up to four 850ul aliquots of plasma, one aliquot of buffy coat and one aliquot of red blood cells. Samples were processed on a purpose made automated system, chilled to +4°C.

### **5.2 Plasma Separator (Sample #3)**

Plasma samples were processed on an automated system designed to retrieve the plasma layer from a gel separation tube. Up to four 850ul aliquots of plasma were recovered per sample. Samples were processed on a purpose made automated system, chilled to +4°C.

### **5.3 Serum Separator (Sample #4)**

Serum samples were processed on an automated system designed to retrieve the serum layer from a gel separation tube. Up to four 850ul aliquots of serum were recovered per sample. Samples were processed on a purpose made automated system, chilled to +4°C.

### **5.4 Urine (Sample #5)**

Urine samples were mixed before being aliquoted into six 850ul aliquots of urine using an automated liquid handling system. Processed samples were immediately moved to a +4°C incubator. Samples were processed at +4°C.

### **5.5 ACD (Sample #6)**

ACD samples were processed by aliquoting with the subsequent addition of RPMI (Roswell Park Memorial Institute media) media and DMSO (Dimethyl Sulphoxide). Two 700ul aliquots of ACD blood were aliquoted into 1.2ml Abgene tubes before the addition of 150ul of DMSO (30ul) and RPMI (120ul) mixture, resulting in two 850ul final volume aliquots. Both aliquots were frozen at a controlled rate down to -196°C. ACD samples were processed at room temperature and the DMSO-RPMI mixture was cooled to +4°C. The environment was HEPA filtered.

### **5.6 EDTA (Sample #7, Haematology Sample)**

Once received at the sample processing centre in Stockport, the samples are analysed on a Beckman Coulter® LH700 System. Samples are analysed within 24 hours of blood draw. The system determines the following hematologic parameters of whole-blood specimens:

WBC	White Blood Cell or leukocyte count
RBC	Red Blood Cell or erythrocyte count
Hgb	Hemoglobin concentration
Hct	Hematocrit (relative volume of erythrocytes)
MCV	Mean Corpuscular (erythrocyte) Volume
MCH	Mean Corpuscular (erythrocyte) Hemoglobin
MCHC	Mean Corpuscular (erythrocyte) Hemoglobin Concentration
RDW	Red Cell (erythrocyte volume) Distribution Width
Plt	Platelet or thrombocyte count
MPV	Mean Platelet (thrombocyte) Volume
LY%	Lymphocyte percent
MO%	Monocyte percent
NE%	Neutrophil percent
EO%	Eosinophil percent
BA%	Basophil percent
LY#	Lymphocyte number
MO#	Monocyte number
NE#	Neutrophil number
EO#	Eosinophil number
BA#	Basophil number
NRBC%	Nucleated Red Blood Cell percent
NRBC#	Nucleated Red Blood Cell number
RET%	Reticulocyte percent
RET#	Reticulocyte number
*HLR%	High Light scatter Reticulocytes %
*HLR#	High Light scatter Reticulocytes #
IRF	Immature Reticulocyte Fraction
MRV	Mean Reticulocyte Volume
*MSCV	Mean Sphered Cell Volume
*Pct	Plateletcrit
*PDW	Platelet Distribution Width

For more information on the calculation of the above haematological parameters, please see the Section 3.7 of the Beckman Coulter® LH700 Series System Manual attached in Appendix 1.

## **5.6.1 Quality Control**

### **5.6.1.1 Daily Controls**

The following controls were run on a daily basis:

#### **Latron controls**

LATRON primer prepares the tubing and instrument components for the LATRON control. LATRON control monitors the performance of the volume, conductivity and light scatter measurements.

#### **5C Cell Control**

5C®-ES control monitors the CBC (complete blood count) and differential (Diff) parameters.

## **Retic C control**

Retic-C cell control monitors the reticulocyte (Retic) parameters.

### **5.6.1.2 Calibration (every six months)**

The haematology systems were calibrated every six months according to the manufacturer's recommendations.

## **5.7 RNA stabilised blood (Sample #8)**

RNA stabilised blood was initially mixed before being aliquoted into 6 aliquots of 850ul. The liquid handling system was HEPA filtered and cooled to +4°C.

## **5.8 Saliva sample (sample #9)**

Saliva samples were mixed and aliquoted into 2 aliquots of 850ul. Saliva samples were processed by an automated liquid handling system in a HEPA controlled environment at +4°C.

## **6 Sample Storage**

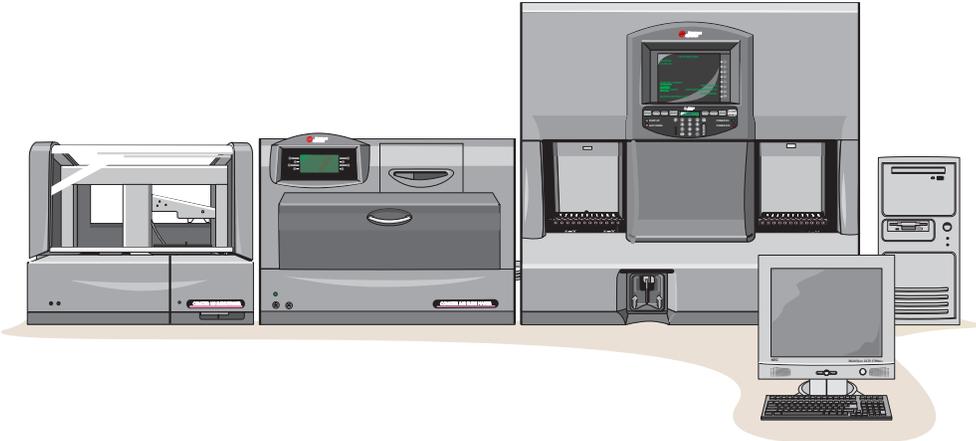
Samples are all stored at -80°C or in liquid nitrogen. All samples were taken, processed and put into storage within 24 hours. The various sample aliquots are stored as follows:

Sample Type	Aliquot Type	Total aliquots	No. of aliquots	
			-80°C	LN2
EDTA 1	Plasma	4	3	1
	Buffy Coat	1	1	
	Red Blood Cells	1	0	1
EDTA 2	Plasma	4	3	1
	Buffy Coat	1	1	
	Red Blood Cells	1	0	1
SST	Serum	4	3	1
PST	Plasma	4	3	1
Urine	Urine	6	4	2
ACD	ACD Blood + DMSO+ RPMI	2	0	2
Tempus (RNA)	RNA stabilised blood	6	0	6
Saliva	Saliva	2	0	2

The -80°C automated freezer is capable of storing up to 10 million samples and can pick in the region of 1000 samples per day.

## **Appendix 1 – Beckman Coulter® LH700 Series System Manual**

Reference



# WARNINGS AND PRECAUTIONS

READ ALL PRODUCT MANUALS AND CONSULT WITH BECKMAN COULTER-TRAINED PERSONNEL BEFORE ATTEMPTING TO OPERATE INSTRUMENT. DO NOT ATTEMPT TO PERFORM ANY PROCEDURE BEFORE CAREFULLY READING ALL INSTRUCTIONS. ALWAYS FOLLOW PRODUCT LABELING AND MANUFACTURER'S RECOMMENDATIONS. IF IN DOUBT AS TO HOW TO PROCEED IN ANY SITUATION, CONTACT YOUR BECKMAN COULTER REPRESENTATIVE.

## **HAZARDS AND OPERATIONAL PRECAUTIONS AND LIMITATIONS**

WARNINGS, CAUTIONS, and IMPORTANTS alert you as follows:

**WARNING** - Can cause injury.

**CAUTION** - Can cause damage to the instrument.

**IMPORTANT** - Can cause misleading results.

BECKMAN COULTER, INC. URGES ITS CUSTOMERS TO COMPLY WITH ALL NATIONAL HEALTH AND SAFETY STANDARDS SUCH AS THE USE OF BARRIER PROTECTION. THIS MAY INCLUDE, BUT IT IS NOT LIMITED TO, PROTECTIVE EYEWEAR, GLOVES, AND SUITABLE LABORATORY ATTIRE WHEN OPERATING OR MAINTAINING THIS OR ANY OTHER AUTOMATED LABORATORY ANALYZER.

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**WARNING** Risk of operator injury if:

- All doors, covers and panels are not closed and secured in place prior to and during instrument operation.
- The integrity of safety interlocks and sensors is compromised.
- Instrument alarms and error messages are not acknowledged and acted upon.
- You contact moving parts.
- You mishandle broken parts.
- Doors, covers and panels are not opened, closed, removed and/or replaced with care.
- Improper tools are used for troubleshooting.

To avoid injury:

- Keep doors, covers and panels closed and secured in place while the instrument is in use.
- Take full advantage of the safety features of the instrument. Do not defeat safety interlocks and sensors.
- Acknowledge and act upon instrument alarms and error messages.
- Keep away from moving parts.
- Report any broken parts to your Beckman Coulter Representative.
- Open/remove and close/replace doors, covers and panels with care.
- Use the proper tools when troubleshooting.

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**CAUTION** System integrity might be compromised and operational failures might occur if:

- This equipment is used in a manner other than specified. Operate the instrument as instructed in the Product Manuals.
- You introduce software that is not authorized by Beckman Coulter into your computer. Only operate your system's computer with software authorized by Beckman Coulter.
- You install software that is not an original copyrighted version. Only use software that is an original copyrighted version to prevent virus contamination.

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**IMPORTANT** If you purchased this product from anyone other than Beckman Coulter or an authorized Beckman Coulter distributor, and, if it is not presently under a Beckman Coulter service maintenance agreement, Beckman Coulter cannot guarantee that the product is fitted with the most current mandatory engineering revisions or that you will receive the most current information bulletins concerning the product. If you purchased this product from a third party and would like further information concerning this topic, call your Beckman Coulter Representative.

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## Issue A, 10/01

Software version 1A. Converted from Help Version 1A.012431.

## Issue B, Complete Revision, 5/02

Software version 2A. Converted from Help Version 2A.021501.

## Issue C, 10/03

Software version 2B. Converted from Help Version 2B.031971.

Changes were made to:

- Change the company name from Coulter Corporation to Beckman Coulter Inc.
- Change LH 750 to LH 700 Series.
- Change all variations of bar-code to be consistent.
- Change 5C-ES to 5C Series.

**Note:** Changes that are part of the most recent revision are indicated in text by a bar in the margin of the amended page.

*This document applies to the latest software listed and higher versions. When a subsequent software version changes the information in this document, a new issue will be released.*

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This introductory section contains the following topics:

- How to use your COULTER® LH 700 SERIES System hard-copy manuals
- About this manual
- Online Help System
- Conventions

## HOW TO USE YOUR COULTER® LH 700 SERIES SYSTEM HARD-COPY MANUALS

Use the **Getting Started** booklet to see an overview of the system hardware and software. This document comes with your LH 700 SERIES System.

Use the **Reference** manual for in-depth information about what the instrument does, the methods it uses, its specifications, and information on installation, safety and software options. The Reference manual for the LH 700 SERIES System is included in the online Help system; it is available in hard copy by request.

Use the **Special Procedures and Troubleshooting** manual to run calibration; to clean, replace or adjust a component on the instrument; and for troubleshooting the instrument. This document is made up of procedures from the online Help system; it is available in hard copy by request.

Use the **Operator's Guide** for the day-to-day operation of your instrument. This document is made up of procedures from the online Help system; it includes Startup, running controls and samples, reviewing data, Shutdown, and the software on the Analyzer and the Workstation. This document is available in hard copy by request.

Use the **SlideMaker Operator's Guide** for in-depth information about what the SlideMaker does, the methods it uses, its specifications, and information on installation, safety and software, as well as day-to-day operating and troubleshooting your SlideMaker. This document is made up of procedures from the online Help system; it is available in hard copy by request.

Use the **SlideStainer Operator's Guide** for the day-to-day operating and troubleshooting of your SlideStainer. This document is made up of procedures from the online Help system; it includes in-depth information about what the SlideStainer does, the methods it uses, its specifications, and information on installation, safety and software. This document is available in hard copy by request.

Use the **Master Index** to easily locate a subject in your hard-copy Reference manual, Operator's Guide or Special Procedures and Troubleshooting manual. The Master Index comes with the hard copy of both the Operator's Guide and the Special Procedures and Troubleshooting manual.

Use the **Host Transmission Specification** to find the information needed to program the transmission interface between the LH 700 SERIES System and your laboratory's host computer. This document comes with your LH 700 SERIES System.

See the Documentation page on the back cover of this manual for the contents of each manual. It can help you to determine quickly in which manual the information you need is located.

## ABOUT THIS MANUAL

Your LH 700 SERIES System Reference Guide is a source of information for the day-to-day operation of your instrument. This information is organized as follows:

- Chapter 1, Use and Function  
Contains the intended use of the instrument, a brief history of the methods used, the reagents, calibrators and controls used, and a short description of the major components.
- Chapter 2, Installation  
Contains the instrument requirements, and the diagrams of the reagent/pneumatic tubing connections and the interunit cable connections.
- Chapter 3, Operation Principles  
Contains descriptions of the Coulter Method, the normal sample flow through the instrument, how counting and sizing are accomplished, and what the DataPlots show.
- Chapter 4, Specifications/Characteristics  
Details the instrument and performance specifications, the performance characteristics, the interfering substances, and the bar-code label specifications.
- Chapter 5, Hazards  
Describes laser safety precautions and the location of the laser-related labels.
- References
- Glossary
- Index, hard copy only

## ONLINE HELP SYSTEM

The LH 700 SERIES Workstation has a comprehensive Online Help System, which includes reference information, all operating, maintenance and troubleshooting procedures. On the LH

700 SERIES Workstation, select  to access Help. Select  to access the tutorials.

## CONVENTIONS

This document uses the following conventions:



indicates a key on the Numeric keypad.



indicates a key on the LH 700 SERIES Workstation keyboard.



is the icon for Patient results on the LH 700 SERIES Workstation.



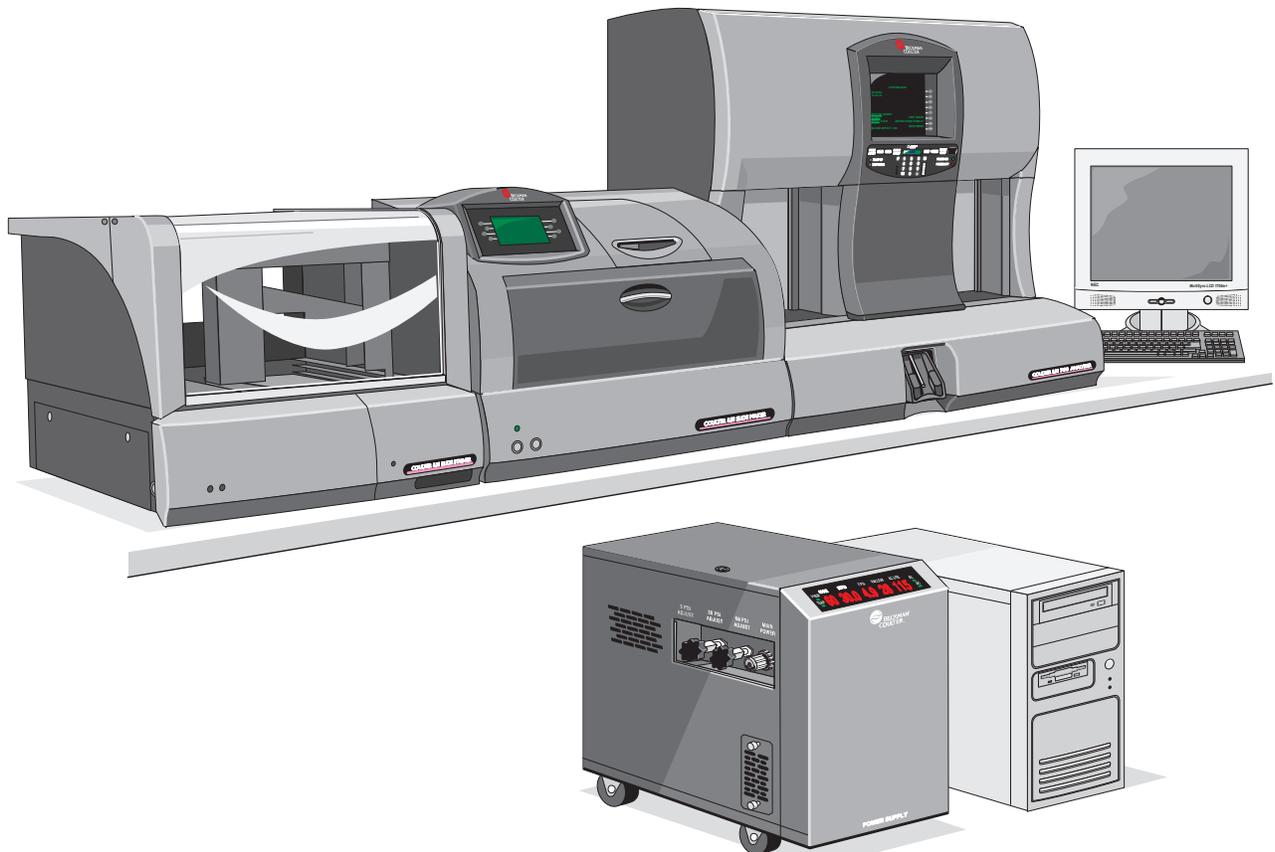
is the icon for the Printer on the LH 700 SERIES Workstation.

## 1.1 INTENDED USE

The COULTER® LH 700 Series, Figure 1.1, is a quantitative, automated hematology analyzer For In Vitro Diagnostic Use in clinical laboratories. The LH 700 Series provides automated complete blood count, leukocyte differential and Reticulocyte analysis and nucleated red blood cell (NRBC) enumeration.

The purpose of the LH 700 Series is to separate the normal patient, with all normal system-generated parameters, from the patient who needs additional studies of any of these parameters. These studies might include further measurements of cell size and platelet distribution, biochemical investigations, manual WBC differential or any other definitive test that helps diagnose the patient's condition.

**Figure 1.1 COULTER LH 700 Series**



## Parameters

The system determines these hematologic parameters of whole-blood specimens:

WBC	White Blood Cell or leukocyte count
RBC	Red Blood Cell or erythrocyte count
Hgb	Hemoglobin concentration
Hct	Hematocrit (relative volume of erythrocytes)
MCV	Mean Corpuscular (erythrocyte) Volume
MCH	Mean Corpuscular (erythrocyte) Hemoglobin
MCHC	Mean Corpuscular (erythrocyte) Hemoglobin Concentration
RDW	Red Cell (erythrocyte volume) Distribution Width
Plt	Platelet or thrombocyte count
MPV	Mean Platelet (thrombocyte) Volume
LY%	Lymphocyte percent
MO%	Monocyte percent
NE%	Neutrophil percent
EO%	Eosinophil percent
BA%	Basophil percent
LY#	Lymphocyte number
MO#	Monocyte number
NE#	Neutrophil number
EO#	Eosinophil number
BA#	Basophil number
NRBC%	Nucleated Red Blood Cell percent
NRBC#	Nucleated Red Blood Cell number
RET%	Reticulocyte percent
RET#	Reticulocyte number
*HLR%	High Light scatter Reticulocytes %
*HLR#	High Light scatter Reticulocytes #
IRF	Immature Reticulocyte Fraction
MRV	Mean Reticulocyte Volume
*MSCV	Mean Sphered Cell Volume
*Pct	Plateletcrit
*PDW	Platelet Distribution Width

\*For Research Use Only. Not For Use In Diagnostic Procedures.

Unless otherwise stated, all parameter results are shown in a US unit format throughout the manuals.

## 1.2 QUALITY CONTROL (QC)

Your laboratory can use these QC techniques with the LH 700 Series:

- Daily instrument checks
- Commercial controls
- Delta checks
- Patient controls
- XB Analysis
- Interlaboratory Quality Assurance Program (IQAP)

Quality Assurance includes routine maintenance and service in conjunction with the use of controls and calibrators. The combination of these methods provides the assurance of complete quality control and should be applied separately or in combination, in accordance with your laboratory, state and federal protocols.

## 1.3 METHOD HISTORY

### Development

W.H. Coulter (1956) describes the Coulter Principle:<sup>1</sup>

A suspension of blood cells is passed thru [sic] a small orifice simultaneously with an electric current. The individual blood cells passing thru the orifice introduce an impedance change in the orifice determined by the size of the cell. The system counts the individual cells and provides cell size distribution. The number of cells counted per sample is approximately 100 times greater than the usual microscope count to reduce the statistical error by a factor of approximately 10 times.

This substantial improvement in precision over previous methods helped to establish the erythrocyte count as a sensitive index of erythropoietic dyscrasia, particularly when considered together with Hct and Hgb measurements.<sup>2</sup>

The COULTER COUNTER® Model S analyzer was the first instrument that automated simultaneous multiparameter measurements on blood. Brittin et al., Gottmann, and Hamilton and Davidson, reviewed the performance and clinical value of the Model S.<sup>3, 4, 5</sup>

Refinements of the COULTER COUNTER analyzer to provide accurate size (volume) distribution data led to a reawakening of interest in pathological erythrocyte size distribution, first sparked by Price-Jones.<sup>6, 7</sup>

Among the advantages offered by the Coulter method of counting and sizing was the ability to derive an accurate Hct measurement by summing the electronic volume of erythrocytes. England et al. speculated that electronic Hct measurements did not contain the trapped plasma error of centrifugal Hct measurements.<sup>8</sup>

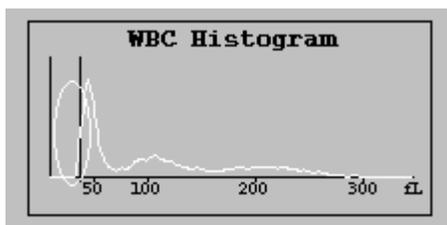
Bull et al. described the use of a COULTER COUNTER analyzer for counting thrombocytes.<sup>9</sup> This method, useful as it was, depended on preparing thrombocyte-rich plasma to avoid counting erythrocytes as thrombocytes. Mundschenk et al. and Schulz and Thom discussed the possibility of counting thrombocytes in the presence of erythrocytes and classifying them by size.<sup>10, 11</sup> Electronic refinements in the Model S-PLUS enhanced the accuracy of the

hydrodynamic method. Von Behrens and Paulus have also cited the feasibility of counting thrombocytes by the Coulter method.<sup>12, 13</sup>

### Corrected WBC Counts

White Blood Cell count results from the CBC analysis. The WBC count is adjusted for interfering substances when appropriate. If there is a population of cells in the far left of the WBC histogram, the number of cells is derived and the WBC count is corrected. No further correction of WBC is required.

**Figure 1.2 Corrected WBC**



When WBC correction has occurred, the uncorrected WBC will appear in the printout Comments field as UWBC = "value". The "Cellular Interference" suspect message is displayed and the corrected WBC count is reported. The uncorrected WBC can be found on the CBC data tab. When the separation between the WBC populations is poorly defined on the histogram, WBC correction will be performed and the corrected WBC will have an R flag.

### Hemoglobinometry

The lytic reagent used for the complete blood count (CBC) parameters prepares the blood so the system can count leukocytes and measure the amount of hemoglobin. The lytic reagent rapidly and simultaneously destroys the erythrocytes and converts a substantial proportion of the hemoglobin to a stable pigment while it leaves leukocyte nuclei intact. The absorbance of the pigment is directly proportional to the hemoglobin concentration of the sample.

The accuracy of this method equals that of the hemoglobincyanide method, the reference method of choice for hemoglobinometry recommended by the International Committee for Standardization in Hematology.<sup>14</sup>

### Differential Measurement

The COULTER VCS established WBC differential technology using three measurements: individual cell volume, high-frequency conductivity and laser-light scatter.

The combination of low-frequency current, high-frequency current and light-scattering technology provides abundant cell-by-cell information that is translated by the instrument into conventional stained-film leukocyte categories.

### Volume Analysis

Electronic leukocyte volume analysis, using low-frequency current, has been used since 1967.<sup>15</sup> It has been evaluated as a possible adjunct to the differential white cell count.<sup>16,17,18,19</sup>

### Conductivity Analysis

Cell walls act as conductors to high-frequency current. The current, while passing through the cell walls and through each cell interior, detects differences in the insulating properties of cell components. The current characterizes the nuclear and granular constituents and the chemical composition of the cell interior.<sup>20,21,22</sup>

### Light Scatter Analysis

Coulter's experience in flow cytometry dates back decades to Fulwyler's pioneering use of light scatter for cell analysis.<sup>23</sup> Loken et al. and Jovin et al. discuss the relationship of particle size and refractivity to the angle of light scattered from a laser beam.<sup>24,25</sup>

### Reticulocyte (Retic) Analysis

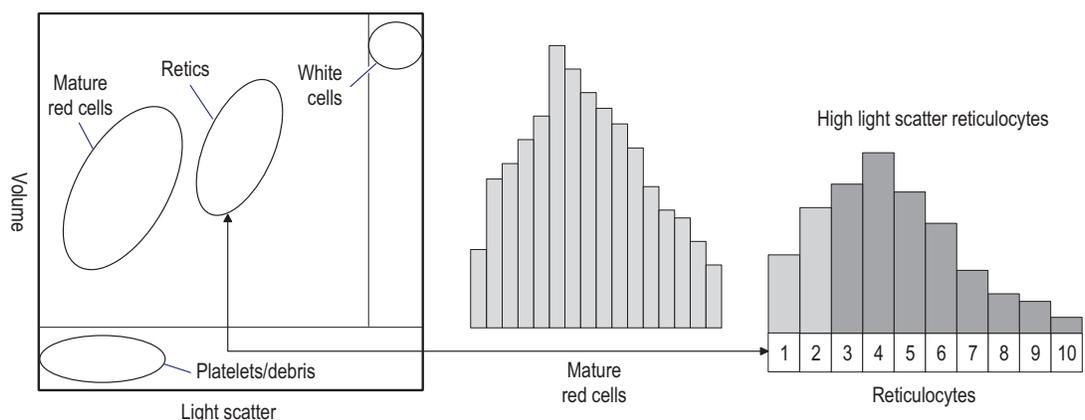
Reticulocytes are immature, nonnucleated erythrocytes retaining a small network of basophilic organelles, consisting of RNA and protoporphyrin. The enumeration of reticulocytes provides a simple, effective means to determine red cell production and regeneration.<sup>26,27,28,29</sup>

The most common means of measuring reticulocytes is to use supravital dyes, such as New Methylene Blue or Brilliant Cresyl Blue. These dyes precipitate and aggregate the basophilic substances within the reticulocyte, resulting in a granular, staining pattern easily seen with light microscopy.<sup>30</sup>

Reticulocyte immaturity is related to cell volume and light scatter. Since more immature reticulocytes are larger, contain more RNA and cause increased light scatter, the cell volume and light scatter will increase with immaturity of the cell.

Figure 1.3 illustrates the IRF and MRV algorithms. This figure is a representation of the VCS data that is shown on the three-dimensional and/or two dimensional analyzer displays.

**Figure 1.3 Illustration of the ten light scatter regions**



The detected light scatter intensity of the reticulocyte population is divided into the 11 equal regions as shown above. After the reticulocyte population is measured and identified as RET% (regions 0 to 10), the spectrum of light scatter intensity for the identified reticulocyte population is analyzed algorithmically. The RET% is calculated as the ratio of reticulocytes to

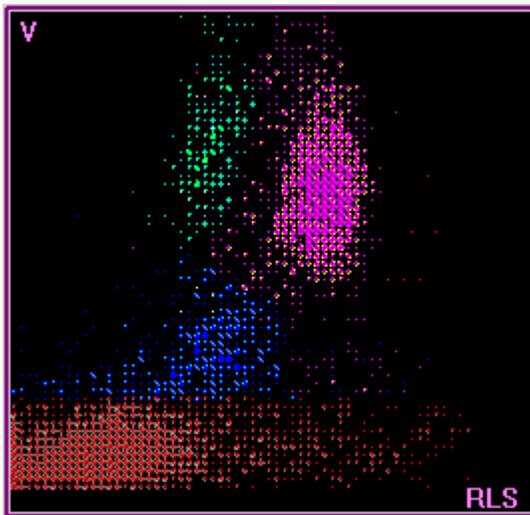
the total number of red cells. The IRF parameter is calculated as the ratio of the total number of reticulocyte events in the outermost 8 regions (regions 3 to 10) to the total number of reticulocytes (defined as regions 0 to 10). The MRV parameter is calculated as the average volume of all reticulocytes; or the mean volume of all retic events.

### NRBC Enumeration

The NRBC Enumeration is achieved through the combined use of impedance and VCS technology and a proprietary algorithm.

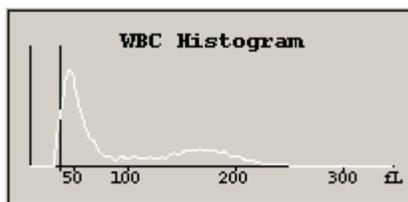
The first step in NRBC enumeration is the identification of particles in the NRBC signature position in the differential data plot. This information is generated from VCS analysis of the cells.

**Figure 1.4 NRBC signature position on Differential Dataplot**



Once cells have been identified in this region, the LH 700 Series examines the far left region of the WBC histogram for the presence of cells.

**Figure 1.5 NRBC location on WBC histogram**



If the VCS dataplot and the WBC histogram both indicate the presence of NRBCs, then the combined information is further evaluated for special data patterns -- such as small lymphocytes, giant platelets, and aging blood. If the combined information from the VCS dataplot and the WBC histogram are consistent with NRBCs, the NRBC count is derived from the WBC histogram.

## COULTER IntelliKinetics™ Application

The LH 700 Series utilizes the COULTER IntelliKinetics application. Control of reaction kinetics is extremely important to ensure the best performance of the automated white cell differential and reticulocyte analysis. The IntelliKinetics application is a management tool for the key step of system optimization when fluctuations in external variables in the laboratory, such as temperature, occur.

The IntelliKinetics application management ensures consistent reaction kinetics. This application intelligently manages variations in ambient laboratory temperature through automatic adjustments to reagent reaction temperature, exposure time and delivery volumes. Enhancements in instrument electronics, such as improved signal-to-noise ratio, work with the IntelliKinetics application to provide better data signals for the system algorithms to analyze. Reagent temperature control helps to increase the speed of dye uptake, thereby improving instrument throughput. Analysis occurs under controlled conditions.

The LH 700 Series with the IntelliKinetics application shows improved separation of populations, both for the white cell differential and reticulocytes. Cell populations made available for analysis by the algorithms are in a more consistent location in three-dimensional space. The IntelliKinetics application, working in concert with new algorithms, provides the instrument with the best signals for analysis, even when the laboratory environment varies throughout the day.

## XB Analysis

Dennis B. Dorsey, MD, proposed in 1963 that the relatively constant blood cell indices could be used to follow the performance of hematology instrumentation.<sup>32</sup> Brian Bull, MD, improved the technique and it is termed XB Analysis.<sup>32</sup>

XB Analysis uses a "weighted moving average" of patient sample results because Koepke and Protector said that QC materials "ideally should be similar in structure and in reactivity to the patient constituent being measured. Therefore freshly drawn patient blood samples seem to be the most appropriate [QC material]."<sup>33</sup> Bull explains, "The analyser [sic] is considered to be 'in control' when mean MCV, MCH, and MCHC determined on a batch of 20 patients by use of the algorithm XB are within 3% of the expected mean indices of the population."<sup>34</sup>

## 1.4 SYSTEM COMPONENTS

The LH 700 Series is a modular system that consists of the following units.

### Power Supply

This unit consists of two assemblies. The Electronic Power Supply assembly provides the regulated and unregulated voltages required by the circuitry of the system. The Pneumatic Power Supply assembly is the source of air pressure and vacuum.

### Diluter

This unit is the primary operating unit of the system. It performs the mixing, transporting, pipetting, diluting, lysing, and sensing functions.

## **Analyzer**

This unit controls the electronic sequence of each operating cycle and calculates the results. It receives count and size information directly from the Diluter while the sample is being cycled; then it counts, measures, and computes the parameters. The Analyzer then sends this information to the Workstation. Many of the controls and indicators needed for normal daily operation are on the front of the Analyzer.

## **LH 700 Series Workstation**

The LH 700 Series Workstation holds the data algorithms used to process the List Mode Data supplied by the Analyzer. From the List Mode Data, the Workstation computes Diff and Retic results, develops the histograms and DataPlots, and displays the results. The Workstation stores the data and transmits it to the Printer and Host computer.

The LH 700 Series Workstation is equipped with a mouse that allows operator interaction with the software.

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**CAUTION** System integrity can be compromised and operational failures can occur if:

- This equipment is used in a manner other than specified.
  - You introduce software that is not authorized by Beckman Coulter into your computer.
  - You install software that is not an original copyrighted version.
- 

Operate the instrument as instructed in your product documentation. Only operate your system's computer with software authorized by Beckman Coulter. Only use software that is an original copyrighted version to prevent virus contamination.

## **Handheld Scanner**

Use the handheld scanner to manually read bar-code labels.

# **1.5 HARDWARE OPTIONS**

## **Graphic/Laser Printer**

You can use any printer that is supported by the Microsoft® Windows® 2000 operating system. The Printer prints the data displayed on the LH 700 Series Workstation screen, including parameter data and graphics.

## **LH 700 Series SlideMaker**

The LH 700 Series SM SlideMaker makes blood smears from samples as they are being analyzed, according to user-defined criteria.

## **LH 700 Series SlideStainer**

The SlideStainer stains blood smears generated by the LH 700 Series SlideMaker or by manually prepared blood smears introduced into the SlideStainer.

## 1.6 CONTROLS AND CALIBRATOR

### Controls

Use stable reference controls to monitor the instrument performance as part of your quality control and to verify calibration. Refer to the package insert for detailed information before using a control.

5C®-ES control monitors the CBC and differential (Diff) parameters.

LATRON primer prepares the tubing and instrument components for the LATRON control.

LATRON control monitors the performance of the volume, conductivity and light scatter measurements.

Retic-C cell control monitors the reticulocyte (Retic) parameters.

### Calibrator

The S-CAL® calibrator kit is an acceptable alternative to the whole-blood reference method of calibration. S-CAL calibrator is traceable to reference methods and materials. Use S-CAL calibrator to ensure accurate instrument measurements. Refer to the package insert for detailed information before use.

The differential and reticulocyte measurement devices are set for optimum performance at the factory.

## 1.7 REAGENTS

Beckman Coulter recommends these reagents or their equivalents. All stated performance characteristics in this manual are based on the use of the LH 700 Series with these reagents. Refer to the container's label for detailed information before using the reagent.

### Diluent

LH 700 series diluent is an isotonic electrolyte solution that:

- Dilutes the whole-blood samples.
- Stabilizes cell membranes for accurate counting and sizing.
- Conducts aperture current.
- Rinses instrument components between analyses.
- Carries and focuses the sample stream in the flow cell to direct the blood cells through the aperture.

Since cell size (volume) is measured, the effect of diluent on osmosis or other phenomena must be tightly controlled. The diluent must not contain particles and must not support growth of bacteria or molds.

### CBC Lytic Reagent

LYSE S® III Diff lytic reagent:

- Rapidly lyses erythrocytes (RBCs), freeing hemoglobin (Hgb) and reducing the size of cellular debris to a level that does not interfere with leukocyte (WBC) count.
- Causes a substantial conversion of the Hgb to a stable cyanide-containing pigment, the absorbance of which is directly proportional to the Hgb concentration over the clinical range.

## LH 700 Series PAK Reagent System

The LH 700 Series PAK reagent system contains Erythrolyse II (PAK LYSE) and StabiLyse (PAK PRESERVE).

### Erythrolyse II (PAK LYSE), Diff Lytic Reagent

Erythrolyse II erythrocyte lytic reagent:

- Dilutes the blood samples.
- Rapidly lyses erythrocytes (RBCs).
- Reduces cellular debris to an insignificant level.

### StabiLyse (PAK PRESERVE), Diff Preservative

StabiLyse leukocyte preservative:

- Maintains leukocytes (WBCs) in their near-natural state.
- Allows the leukocytes to be differentiated into their subpopulations through the volume, conductivity and light-scatter measurements.

## LH 700 Series RETIC PAK Reagent System

The LH 700 Series RETIC PAK contains Reagent A and Reagent B.

### Reagent A, Retic Stain

Reticulocyte staining solution is a specially formulated, New Methylene Blue (NMB) dye that stains the reticulum.

### Reagent B, Retic Clearing Solution

Reticulocyte clearing solution is a clearing reagent that removes hemoglobin from the erythrocytes (RBCs) without removing the precipitated dye-RNA complex, keeping the cell and its membranes intact.

## Cleaning Agent

COULTER CLENZ® cleaning agent cleans and rinses the internal surfaces of the instrument components. Daily use prevents protein buildup and eliminates the need for routine aperture bleaching.

## 1.8 MATERIAL SAFETY DATA SHEETS (MSDS)

To obtain an MSDS for Beckman Coulter reagents used on the LH 700 SeriesLH 700 Series:

1. On the internet, go to [www.beckmancoulter.com](http://www.beckmancoulter.com) and select MSDS from the Customer Support dropdown menu.
2. If you do not have internet access:
  - In the USA, either call Beckman Coulter Customer Operations (800-526-7694) or write to:  
Beckman Coulter Inc.  
Attn: MSDS Requests

P.O. Box 169015  
Miami, FL 33116-9015

- Outside the USA, contact your Beckman Coulter Representative.

**USE AND FUNCTION**  
*MATERIAL SAFETY DATA SHEETS (MSDS)*

## 2.1 GENERAL

---

**CAUTION** Possible system damage can occur if you uncrate the instrument, install it or set it up. Keep the instrument in its packaging until your Beckman Coulter Representative uncrates it for installation and setup.

---

Your instrument is tested before it is shipped from the factory. International symbols and special handling instructions printed on the shipping cartons tell the carrier how to handle this electronic instrument.

Carefully inspect all cartons when they arrive. If you see any sign of mishandling or damage, file a claim with the carrier immediately. If the shipment is separately insured, file a claim with the insurance company.

## 2.2 SPECIAL REQUIREMENTS: HARDWARE

Install and operate this instrument in a conventional clinical laboratory environment. Since the individual units are all interrelated, you must determine the system location and layout before your Beckman Coulter Representative arrives to install the instrument. Consider the following special requirements.

### Space and Accessibility

In addition to the space required for the individual components, consider:

- Comfortable working height.
- Access to perform service procedures. Allow at least 46 cm (18 in.) for the rear doors plus sufficient room for work space. Units can be moved to obtain additional work space.

### Electrical Input

---

**CAUTION** Introduction of electrical interference causing the instrument to lock up or reset frequently can occur if you do not plug the primary power cables directly into an electrical outlet. Overheating, melting and burning of the power lines can occur if you use an extension cord with the primary power cables. Plug the primary power cables directly into an electrical outlet. Place the instrument close enough to an electrical outlet that an extension cord is not needed.

---

This instrument requires:

- An independent protected circuit
- A three-wire outlet furnishing the applicable line voltage, single-phase input power
- A ground path capable of carrying the full current of the circuit (confirmed thirdwire earth ground)
- That the 3-m (10-ft) primary power cord on the rear of the Power Supply be plugged directly into the electrical outlet. Do not use an extension cord.

Current-carrying capacity of 20 A is recommended, although the actual power consumption is only 2080 W as shown in the table below:

<b>Instrument Components</b>	<b>Watts</b>
Analyzer, Diluter, Power Supply	860
SlideMaker	430
SlideStainer	430
Computer	300
Monitor	60
	2080

Building outlets must be properly grounded and transients protected.

**Ambient Temperature and Humidity**

Operate the system in a room with a temperature of 15.5° to 32°C (60° to 90°F) and humidity up to 95% without condensation.

If the average room ambient temperature changes more than 5.5°C (10°F) from the calibrating temperature, verify calibration and recalibrate if necessary to ensure conformance to specifications.

**Air Conditioning**

In air-conditioned environments, an additional 5,500 Btu is required to compensate for the heat the system generates.

**Ventilation**

All ventilation fans must be at least 25 cm (10 in.) away from walls or obstructions that could interfere with the flow of air.

**Drainage**

---

**CAUTION** Incomplete waste chamber drainage and eventual waste chamber overflow into the vacuum system can occur if the waste line is too long. Contact your Beckman Coulter Representative if you need to increase the length of the waste line supplied with the instrument.

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**WARNING** Biohazardous contamination could occur from contact with the waste container and its associated tubing if not handled with care. Avoid skin contact. Clean up spills immediately. Dispose of the contents of the waste container in accordance with your local regulations and acceptable laboratory procedures.

---

The maximum waste line length is 3.7 m (12 ft). The waste drain tubing (rear panel of the Diluter) supplied with the system can be connected to either:

- An open drain, suitable for biohazardous waste, less than 76 cm (30 in.) above the floor
- A waste container with a minimum capacity of 20 L (5 gal.).

When using an open drain instead of a waste container:

- Mechanically secure the waste tube into the drain so the tube cannot accidentally come out of the drain. This prevents spillage.

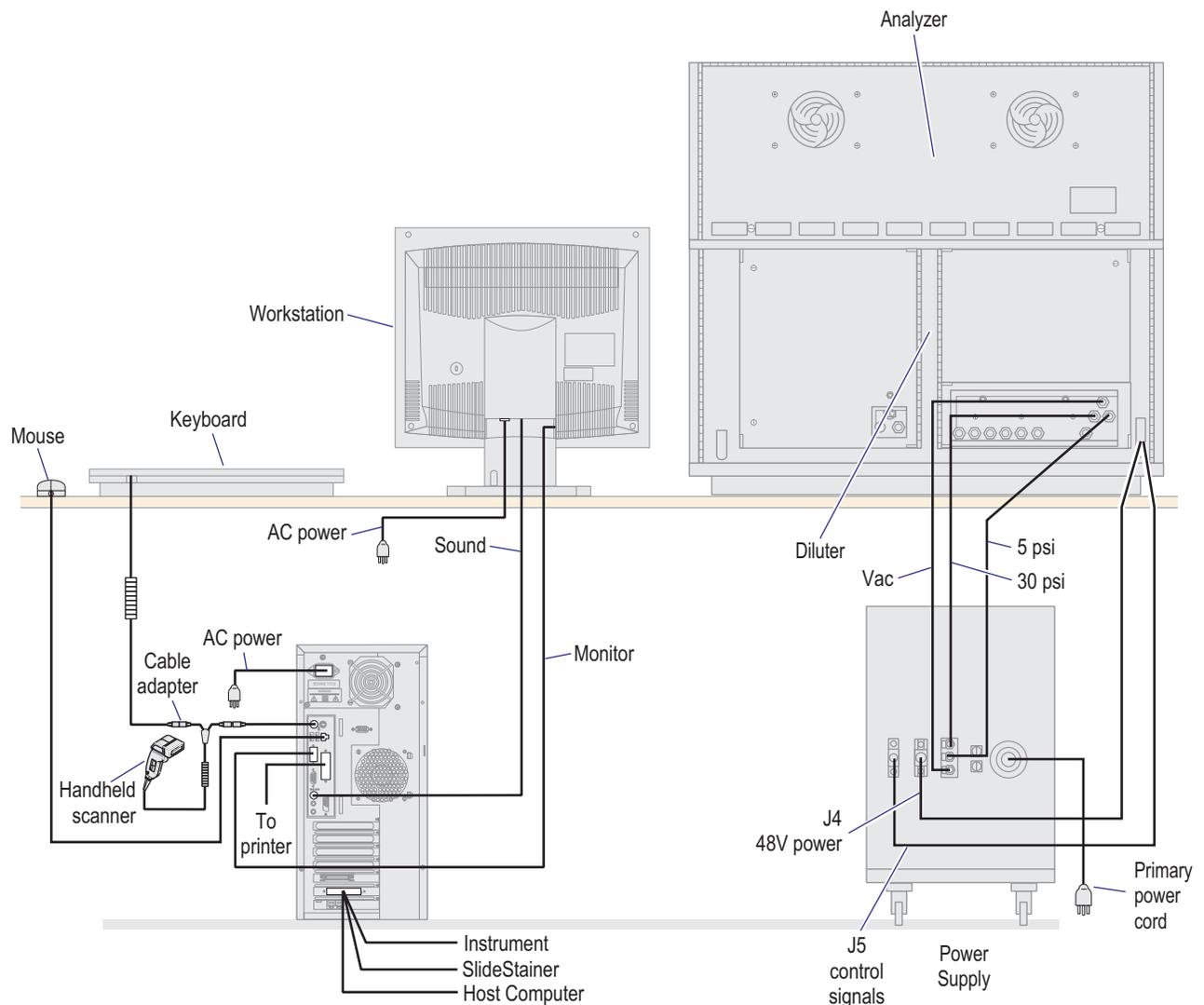
Be sure to dispose of waste in accordance with environmental protection regulations.

## 2.3 INTERUNIT CONNECTIONS

### Power and Signal Cables

Figure 2.1 shows the interunit connections of the power and signal cables that are supplied with the instrument. Your Beckman Coulter Representative makes these connections when installing the instrument.

**Figure 2.1 Interunit Power and Signal Cable Connections**



## Pneumatic/Hydraulic Tubing Connections

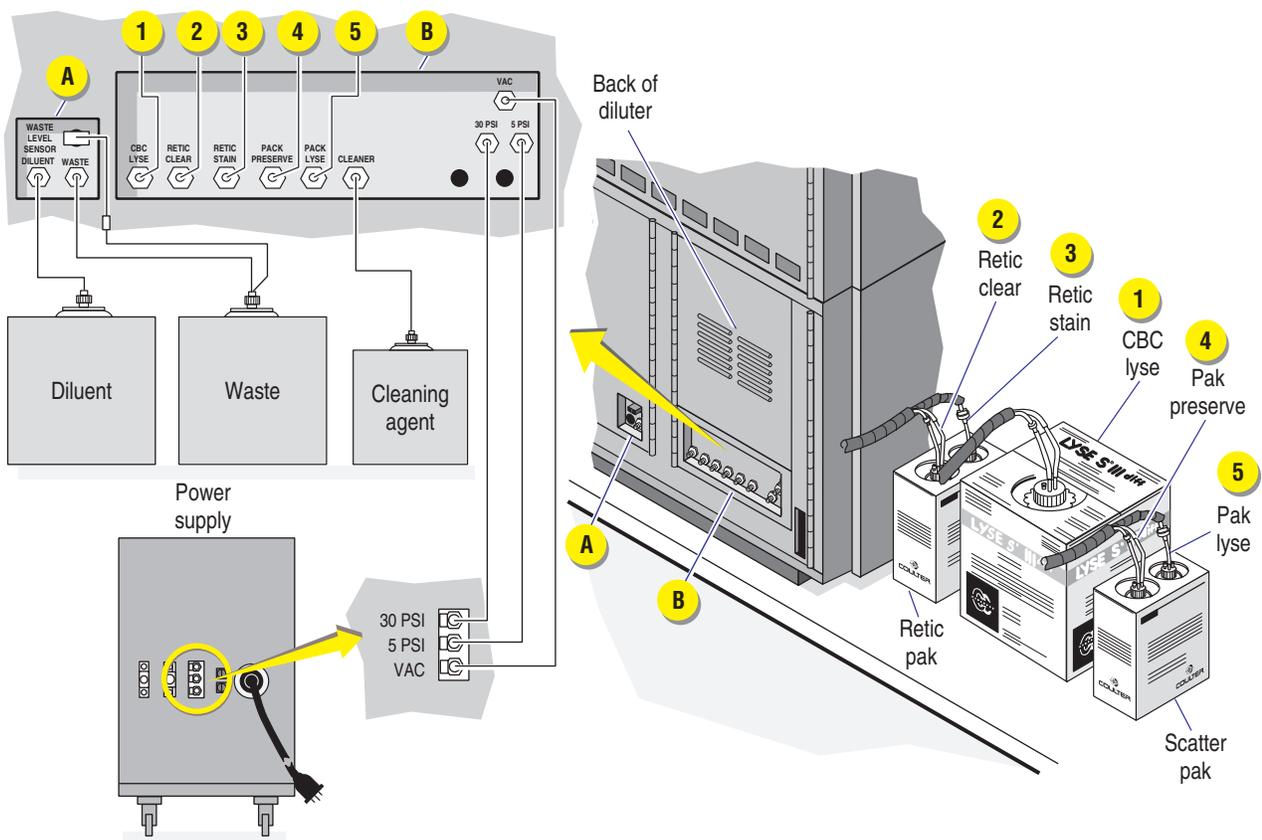
**CAUTION** Possible reagent siphoning effect and priming problems can occur if a reagent container is placed above the level of the Analyzer. Do not place reagent containers above the level of the Analyzer.

**IMPORTANT** Placing Reagent Paks in any location other than on the counter next to the instrument may cause erroneous results.

Figure 2.2 shows the tubing connections between the Diluter and the:

- Reagent containers
- Waste container
- Power Supply pressure and vacuum supplies.

**Figure 2.2 Pneumatic/Hydraulic Connections**

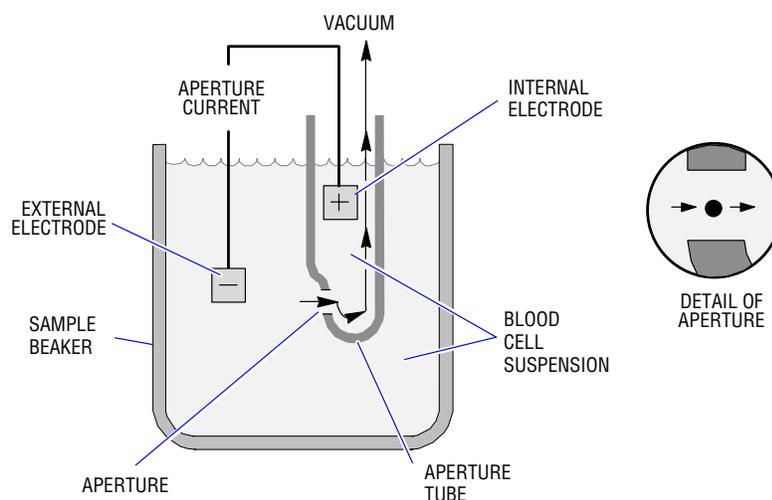


### 3.1 COULTER METHOD

#### CBC Analysis

The Coulter method counts and sizes cells by detecting and measuring changes in electrical resistance when a particle (such as a cell) in a conductive liquid goes through a small aperture. See Figure 3.1.

**Figure 3.1 Coulter Method of Counting and Sizing**



Each cell suspended in a conductive liquid (diluent) acts as an insulator. As each cell goes through the aperture, it momentarily increases the resistance of the electrical path between two submerged electrodes, one located on each side of the aperture. This causes an electrical pulse that can be counted and sized.

While the number of pulses indicates particle count, the size of the electrical pulse is proportional to the cell volume.<sup>35,36,37,38</sup>

#### Differential Analysis

WBC differential analysis and classification occurs in the flow cell, where:

- Low-frequency current measures volume,
- High-frequency current senses cellular internal content through measuring changes in conductivity,
- Light from the laser bouncing off the individual WBC cells characterizes cellular surface, shape and reflectivity.

### **Effect of Reagents**

The conductive diluent must affect cells minimally, if at all.

Both lytic reagents must destroy erythrocytes without significantly affecting leukocytes. They must work rapidly to satisfy the speed with which the system works.

The leukocyte preservative must

- Provide clear separation of the white blood cell populations, and
- Preserve leukocytes in their near-natural state for accurate cytometric measurement.

### **Reticulocyte Analysis**

A supravital dye, New Methylene Blue, is incubated with whole-blood samples. The dye precipitates the basophilic RNA network found in reticulocytes. Hemoglobin and unbound stain are removed by adding a clearing reagent, leaving clear spherical mature RBCs and darkly stained reticulocytes.

Stained reticulocytes are differentiated from mature red cells and other cell populations by light scatter, direct current measurements, and opacity characteristics.

## **3.2 AUTOMATIC ASPIRATION MODE**

The system automatically transports, mixes, aspirates and processes specimens.

### **Loading Specimens**

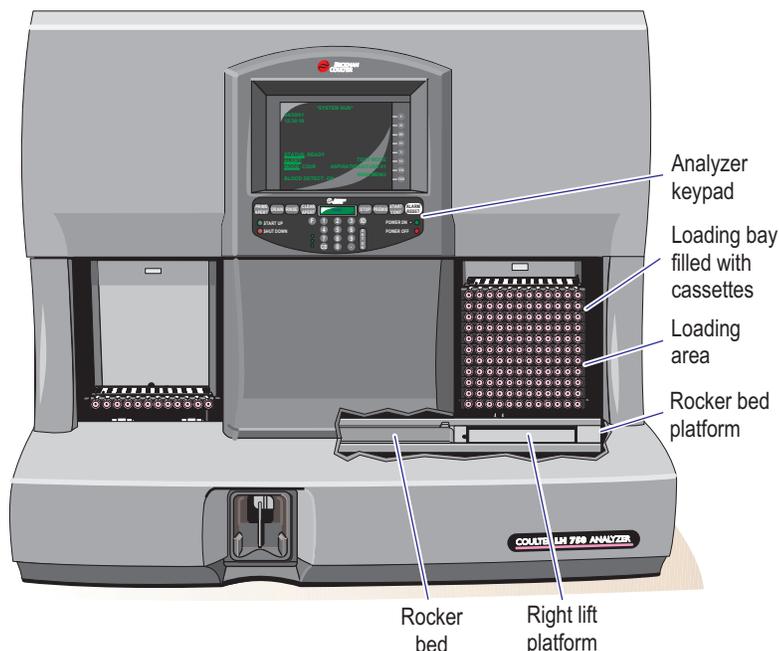
1. The operator places specimen tubes, which can be identified by bar-code labels, into cassettes. Each cassette and the tube positions in the cassette are identified by bar-code labels.
2. You can load up to 12 standard or Hemogard cassettes with 144 samples into the loading bay at one time. Figure 3.2 shows the loading bay filled with cassettes.

### **Transporting Cassettes**

The system transports each cassette from the loading bay to the sampling station.

1. The right lift platform (Figure 3.2) rises beneath the stacked cassettes in the loading bay.
2. The bottom cassette is deposited on the platform.
3. The platform lowers the cassette to the level of the rocker bed.

Figure 3.2 Transport System



4. The cassette moves onto the rocker bed where it rocks back and forth, mixing the specimens.
5. The cassette moves toward the sensing station until it reaches the tube sensor.
6. When the first tube is sensed, the stripper plate locks onto the tube.
7. After at least 14 rocks from the time the cassette was loaded, the rocker bed locks in a 45-degree forward position.
8. At the sampling station, the tube is locked in position.
9. The tube ram pushes the tube out from the cassette, causing the needle to pierce the tube stopper.
10. The bar-code reader scans the cassette and tube labels on both its forward and return passes; an audible indicator can be enabled to indicate each correctly-read bar-code.
  - If the two bar-code scans are identical, the bed continues to rock, and the cassette moves forward until the next available tube reaches the sampling station. If the bar-code reader detects a discrepancy between the forward and return readings, it makes an additional pass.
  - If the bar-code reader cannot read the cassette bar-code label, the Diluter stops, posts a message on the Analyzer, Diluter and Workstation displays, backwashes the system, then returns to the Ready state. If Cass/Pos is the Positive Identifier, and the bar-code reader cannot read the tube bar-code label, dashes (-----) appear in the Sample ID field on the Workstation with the sample results.
11. After the last tube in the cassette is aspirated, the cassette continues along the rocker bed and is deposited onto the unloading bay platform. The left platform lifts the cassette into the exit bay, where up to 12 cassettes can be stacked.

## **Aspiration**

At the sampling station, after the cap is pierced:

1. A pump draws a maximum of 300  $\mu\text{L}$  (LH 700 Series only) or 550  $\mu\text{L}$  (LH 700 Series SM SlideMaker) of sample through the needle and through the Blood Sampling Valve (BSV).
2. The blood detectors monitor the passage of sample through the BSV and aspiration lines.
3. The needle is withdrawn and the sample tube is resealed in the cassette.

## **Delivery**

### **CBC**

After the sample is aspirated:

- The center section of the BSV rotates and segments the sample into two separate volumes.
- Beginning a few seconds before the delivery of the dilutions to the appropriate baths, 5 psi of pressure is sent to the WBC bath. This pressure allows drainage of any residual liquid in the WBC bath, thus preventing carryover.
- The pressure continues during delivery and forms bubbles that mix each cell suspension before sensing begins.
- At the beginning of the delivery, any residual rinse in the Hgb cuvette drains into the waste chamber and the waste chamber drains.
- Diluent from the diluent dispensers drives the separated volumes of sample from the BSV to the baths.
- One volume of sample, 1.6  $\mu\text{L}$ , is delivered with 10 mL of diluent to the RBC bath. This dilution is used for RBC/Plt counting and MCV/Plt sizing.
- The other volume, 28  $\mu\text{L}$ , is delivered with 6 mL of diluent to the WBC bath. This dilution is used to count WBC and develop Hgb.
- During delivery to the WBC bath, 1 mL of lytic reagent is added to the dilution to lyse the red cells and convert Hgb.
- At the same time the lytic reagent is dispensed, 5 mL of diluent from the backwash tank is transferred into the Hgb cuvette for the Hgb blank reading.
- The final dilution in the WBC bath is 1 part whole blood in a total volume of 251 parts. The final dilution in the RBC bath is 1 part whole blood in a total volume of 6250 parts.
- The vent section of the piercing needle is rinsed, then dried by high vacuum.
- The center section of the BSV returns to the aspirate position.

### **Differential (Diff) and Retic**

At the same time as the segmented parts of the sample are being delivered to the baths, the Diff and Retic segmenting modules segment additional 31  $\mu\text{L}$  samples of the blood for analysis.

- For the WBC differential, the sample and approximately 0.506 mL of heated Diff lytic reagent are delivered to the mixing chamber, which agitates to mix them thoroughly.

During the mixing process, approximately 0.2 mL of Diff preservative enters the mixing chamber to preserve the leukocyte populations, and the instrument initiates the sheath stream of diluent in the triple-transducer flow cell.

- For the Retic analysis, the sample and approximately 166 µL of Retic stain are delivered to the Stain chamber, where they are mixed, heated and incubated for 31 seconds.

A 2-µL segment of the stain/blood mixture and 2.00 mL of heated or cooled Retic clearing solution are delivered to the Retic chamber, where they are mixed and incubated for 25 seconds.

The sample line between the Retic chamber and the flow cell is primed with the stain/blood/clearing solution before cell counting begins. After the sample line is primed, the flow cell is cleaned and primed with diluent, and the sheath stream of diluent begins.

For both the WBC differential and the Retic analysis, the instrument injects the sample into the center of the sheath stream and activates the flow cell aperture current. The laminar flow guides the sample through the center of the flow cell aperture; the sheath stream on the exit side of the flow cell aperture prevents the sample's cells from re-entering the aperture.

### **CBC Sensing System**

Vacuum, equal to 6 in. of mercury, draws a precise volume of suspension from each bath through the three apertures. At the same time, sweep flow is drawn behind the RBC apertures to prevent cells from re-entering the sensing zone.

When the vacuum starts to draw the suspension, current is supplied to the electrode. The electrical path allows sensing of the number and volume of each cell pulled through the apertures.

While the sample in each bath is sensed, the photometer reads the Hgb-blank and the Analyzer retains this reference voltage.

### **CBC Analysis in the Baths**

The RBC and Plt data is generated by the RBC bath. The WBC and Hgb data is generated by the WBC bath and Hgb cuvette.

In the analyzer the R/W Proc counts and sizes the RBC and WBC data. The PLT Proc sizes the PLT data. The Comm Intrfc measures the HGB blank and sample. The RBC and WBC raw data consists of counts, waittime counts, count time, and channelyze time, as well as histograms for each of the three apertures. The PLT raw data consists of the histogram and channelyze time for each aperture. The HGB raw data consists of two voltage measurements for the blank and two measurements for the sample. The Analyzer then sends the raw data to the workstation.

The workstation then:

- Wait time and Coincidence corrects the RBC and WBC raw counts
- Fit the PLT histogram and Coincidence correct the PLT histogram count.
- Calculate HGB from the blank and sample readings
- Scale for calibration and dilution

- Perform Voting on the three apertures for RBC, WBC, PLT.
- Derives the MCV and RDW parameters from the RBC histogram;
- Derives the MPV and PDW parameters from the Plt histogram.
- Displays results and histograms.

### Differential and Retic Multiparameter Sensing System

For the WBC differential and for the Retic analysis, the multiparameter sensing system produces the three measurement signals. Figure 3.3 shows the Triple Transducer Module and its protective housing as it resides in the Diluter.

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**WARNING** The laser beam can cause damage to your eyes and to the instrument. Do not attempt to remove the laser from the Diluter module. If removal is required, it must be done only by a Beckman Coulter Representative.

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Tamper-proof screws secure the protective housing; they can only be removed with a special tool.

The laser is a helium-neon laser that complies with the United States' performance standard for laser products, Title 21 Code of Federal Regulations 1040.10 and 1040.11. Figure 3.4 shows the laser module without its protective housing to display the flow cell and label locations.

**Figure 3.3 Triple Transducer Module with Protective Housing**

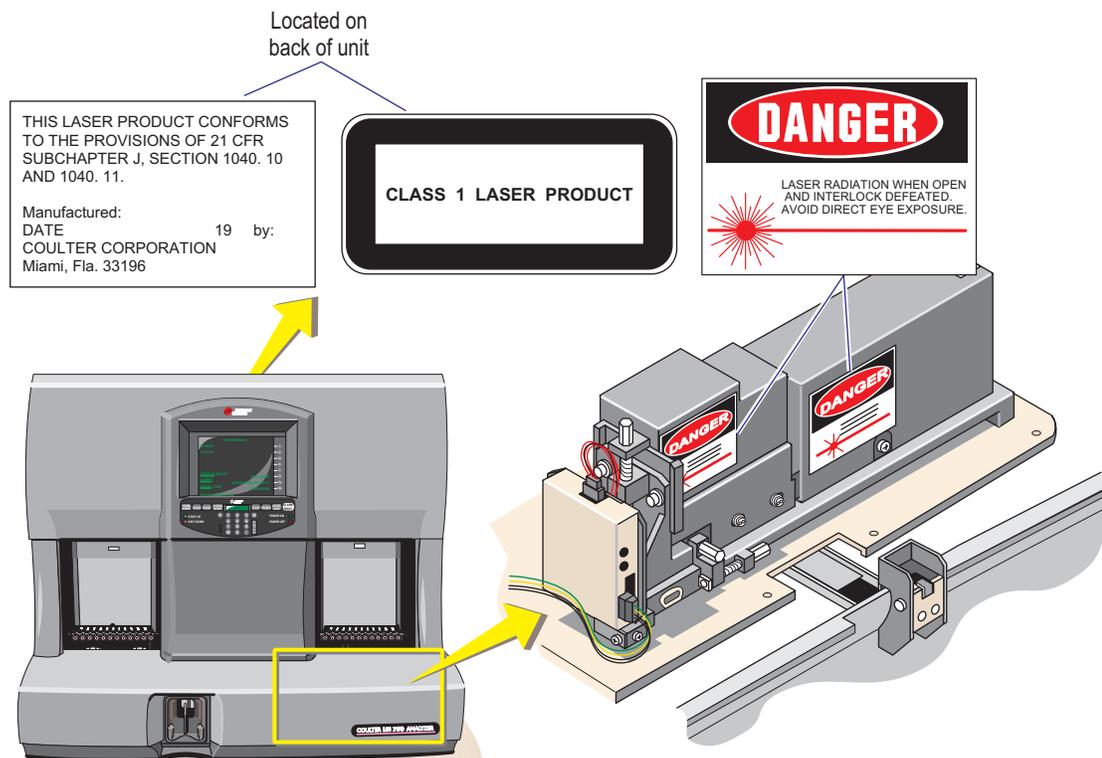
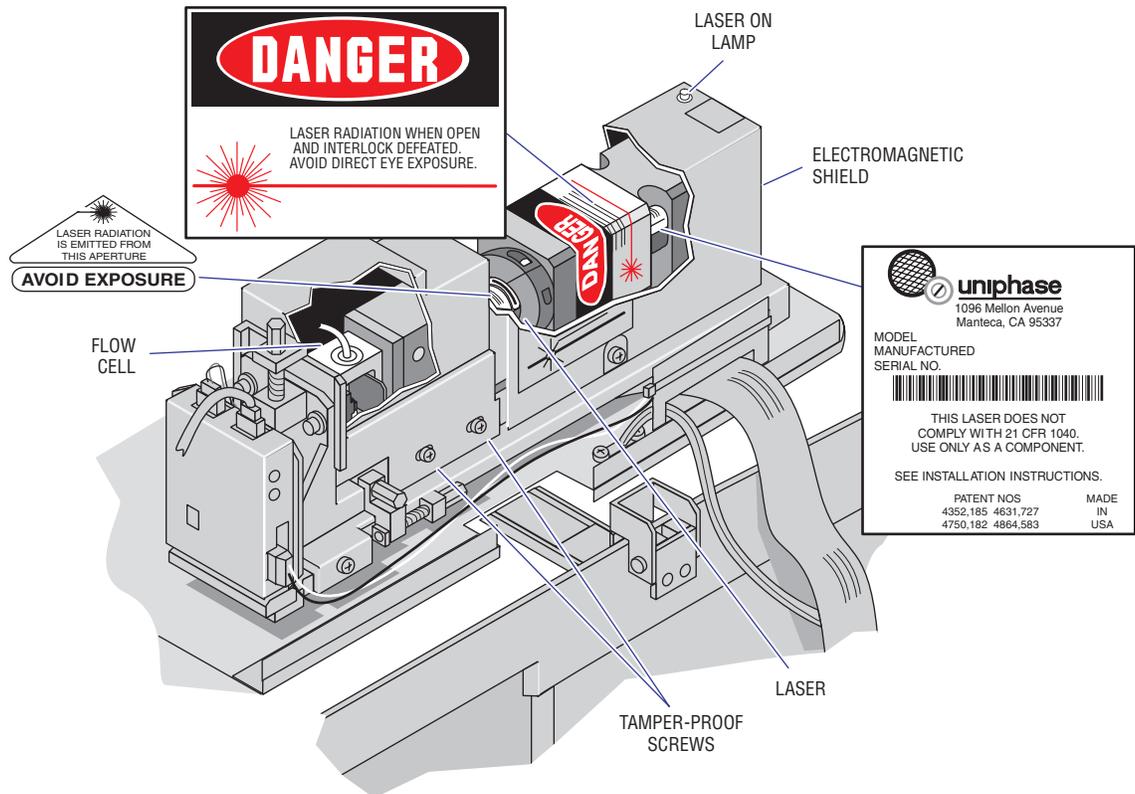


Figure 3.4 Triple Transducer Module with Protective Housing Cut Away



**Note:** As installed in the Triple Transducer Module (TTM) safety fixture, the laser presents no radiation hazard to users and complies with 21 CFR 1040.

### WBC Differential Analysis

The LH 700 Series makes three measurements (volume, conductivity, and scatter) as each cell passes through the flow cell.

The low-frequency impedance measurement defines cell volume. The high-frequency conductivity measurement indicates the internal conductivity. The light-scatter measurement indicates the structure and shape.

Three raw analog signals are sent to the Analyzer for amplification and signal processing. The Analyzer sends this raw data to the Workstation for computation and Data Plot generation of the five differential parameters. The Analyzer sends the CBC results to the Workstation. All results are displayed at the same time.

### Reticulocyte Analysis

Reticulocyte samples are run in one of the three RETIC modes. The modes, which are set at the Workstation, are:

- RETIC
- CBC/RETIC
- CBC/DIFF/RETIC

The LH 700 Series uses the Triple Transducer Module to measure these parameters:

- Reticulocyte percent (RET%) - the number of reticulocytes per 100 RBCs, directly measured and reported as a percentage of RBCs.
- Reticulocyte number (RET#) - the absolute number of reticulocytes, calculated from RET% and RBC number. Expressed and reported as  $10^6$  cells/ $\mu$ L or  $10^9$  cells/L (US units).
- Mean reticulocyte volume (MRV) – the average cell volume of the reticulocytes, determined by the VCS technology algorithm and reported in femtoliters (fL).
- Immature reticulocyte fraction (IRF) - a calculated ratio using the count of the highest light scatter reticulocytes to the total count of reticulocytes. The higher the light scatter the more immature the reticulocytes are. The degree of stained reticulum (using New Methylene Blue) is proportional to the degree of the RNA present in the cell and thus the amount of light scatter produced. It is reported as a decimal.

### Backwash and Rinse

The Diluter performs backwash of aspiration pathways and rinses its components.

Approximately 0.5 mL of Erythrolyse II erythrocyte lytic reagent is delivered to the mixing chamber to remove residual material from the previous cycle.

10 mL of diluent rinse for the WBC bath comes from the RBC diluent dispenser, and 6 mL of diluent rinse for the RBC bath comes from the WBC diluent dispenser. The WBC bath needs a larger rinse volume to

- Remove RBC cell stroma after lysing
- Remove remaining lytic reagent
- Rinse above the 7 mL fill line
- Rinse the hemoglobin cuvette.

## 3.3 MANUAL ASPIRATION MODE

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**IMPORTANT** Clots in the specimen can cause misleading results. The blood detectors are not active in the Manual mode. Be sure to inspect the specimen for clots, and use good laboratory practices to verify results.

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The Manual mode of operation is like the Automatic mode except:

- Before you run the sample, you enter the sample identification number by performing any of the following--
  - ▶ Scanning the bar-code label on the tube
  - ▶ Entering the ID number on the Numeric Keypad
  - ▶ Entering the ID number in the bar-code field on the LH 700 Series Workstation from its keyboard.
- You use an open vial and introduce the sample at the aspirator tip.
- You begin the cycle by either
  - ▶ Obstructing the optical sensor's beam

- ▶ Pressing the panel behind the tip.
- The system aspirates a maximum of 200  $\mu\text{L}$  of sample.
- The blood detectors are not active in the Manual mode.

## 3.4 COUNTING AND SIZING

### Red and White Blood Cell Counting

The RBC and WBC baths each have three discrete apertures that function as independent systems. The six aperture currents are individually adjusted during calibration. When the Aperture Current/Signal Generator (API-SIG GEN) card applies current to the apertures, there is a delay. During this delay, the system conditions the electronics to perform the counting and sizing of the sample.

Regulated vacuum draws a precise volume of sample dilution through the apertures. At each aperture, the system gathers pulses for 4 seconds. The RED/WHITE PRE-AMP cards amplify these pulses and the Analyzer screen displays them.

The system sends these pulses to the RED/WHITE PROCESSOR card. The RED/WHITE PROCESSOR card counts and sizes the RBC and WBC data. The RBC and WBC raw data consists of counts, waittime counts, count time, and channelize time, as well as histograms for each of the three apertures. The Analyzer then sends the raw data to the workstation.

The workstation then:

- Wait time and Coincidence corrects the RBC and WBC raw counts
- Fit the PLT histogram and Coincidence correct the PLT histogram count.
- Calculate HGB from the blank and sample readings
- Scale for calibration and dilution
- Perform Voting on the three apertures for RBC, WBC, PLT.
- Derives the MCV and RDW parameters from the RBC histogram;
- Derives the MPV and PDW parameters from the Plt histogram.
- Displays results and histograms.

### Coincidence Correction

Occasionally, more than one cell passes through the aperture at the same time. When cells coincide, the Analyzer counts only one pulse. As the frequency of coincidence is proportional to the actual count, the system automatically corrects results for coincidence.

### Voting

To prevent data errors due to statistical outliers or obstructions that may block an aperture, the Analyzer votes on the data from all of the apertures, and rejects any questionable data. For the WBC count, RBC count, MCV, RDW, Plt count, and MPV, the Analyzer computer compares the data from the three apertures. It verifies that at least two apertures have produced data within an established statistical range of each other.

If the data from one aperture is outside the established statistical range, the computer votes out the data and histograms from that aperture. The computer derives the affected parameter by averaging the data from the two remaining apertures.

The data from at least two of the three apertures must be within an established statistical range of each other, or the system totally votes out the parameter and histograms. When a parameter totally votes out, the system does not give any results for the affected parameter or for any parameters that are derived from it. See your Online Help System or Operator's Guide for codes and messages that appear in these circumstances.

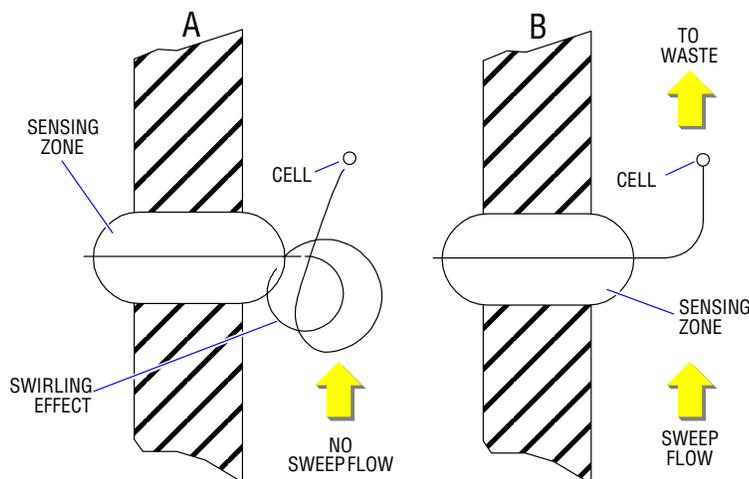
### Pulse Editing

When cells pass through the aperture near the edge or at an angle rather than at the center, they create atypical pulses. The RED/WHITE PROCESSOR card edits RBC and WBC pulses to exclude these atypical pulses from analysis because they distort the true size of the cell. This prevents the atypical pulses from influencing size measurement. Each of the six apertures has an editor.

### Sweep Flow

The sweep flow is a steady stream of diluent that flows behind the RBC aperture during the sensing period. This prevents cells from re-entering the sensing zone and being counted as platelets. See Figure 3.5.

**Figure 3.5 Sweep Flow**



### RBC Size Distribution

After editing, the RED/WHITE PROCESSOR converts the three sets of both the RBC and WBC pulses. It converts each pulse to a number that corresponds to the size of the cell. The Analyzer Microcontroller card (AMC) reads this data and generates histograms. The digital information from each aperture is stored according to volume in 256-channel, size-distribution histograms.

To ensure that the size-distribution curve accurately reflects the true cell population, the system extends RBC sensing for up to eight additional 2-second sensing periods whenever

the RBC data accumulations are below a predetermined value. The RBC size distribution curve reflects the total data accumulated in all of the sensing periods.

After the sensing periods are completed, the system sends these histograms through the Communication Interface (COMM INTFC) card to the Workstation for display.

### Plt Count and Size Distribution

Pulses representing cells from 2 to 20 fL are classified as platelets in the PLATELET PROCESSOR card.

The system digitizes the Plt pulses. The AMC card reads the digital information and channelizes it into three 64-channel size-distribution histograms. The PLT channelizing time is 2 seconds minimum, up to 20 seconds maximum, in 1 second increments. PLT channelizing stops when there are at least 1500 cells channelized in each of the 3 histograms, or the 20 second maximum time is reached.

After the sensing periods are completed, the system sends these histograms and channelizing time through the COMM INTFC card to the Workstation. In the Workstation, an initial platelet count is derived from the fitted or raw histograms and the channelizing time. Then, coincidence correction is applied. MPV and PDW are derived from the histograms. Pct is calculated. The computer votes on the parameter results from each histogram (fitted or not fitted). Results are scaled for reported unit and calibration, and the parameters displayed on the Workstation.

### Plt Fitting Process

The Platelet Algorithm smooths the platelet histogram from each aperture. The algorithm then finds a maximum (mode) point and two minimum points (valleys) in each smoothed histogram. The minimum points are located to the left and the right of the maximum point. The algorithm uses all the information in the raw histogram within the two minimum points to fit an extrapolated curve to the log-normal smoothed curve, using a least-squares fit method.

The Algorithm verifies that each of the fitted curves:

1. Is positive and unimodal
2. Has a mode between 3 and 15 fL
3. Has a PDW less than 20. (NOTE: This is an approximate value since the measurement is made before results are scaled for calibration.)
4. Has a ratio between the raw and fitted histograms mode amplitudes of less than 1.5
5. Has a ratio between channels 0 and 1 and the mode amplitudes of less than 1.5
6. Has a plt count approximately greater than 20.

If any of these conditions are not met, the algorithm derives the platelet count from the portion of the raw histogram between the two minimum points.

Condition 5 above causes generation of a Plt no-fit condition where the Plt is flagged with a Review (R) flag, and the 'Low Plt Interference' message is displayed on the Research screen. Additionally, Plt results are Review (R) flagged when the 'High Plt Interference' flag occurs (the text message is displayed on the Research screen).

**Retic Parameters**

The system makes three measurements as each cell passes through the flow-cell aperture:

- The low-frequency impedance measurement, which defines the cell's volume
- The high-frequency impedance measurement, which indicates the cell's internal conductivity
- The light-scatter measurement, which indicates the cell's structure and shape.

**3.5 MEASUREMENT OF HEMOGLOBIN CONCENTRATION**

After the WBC count, the lysed WBC dilution drains into the hemoglobin cuvette for Hgb measurement.

A beam of white light from an incandescent lamp goes through the cuvette and then through an optical filter that has a center transmission wavelength of 525 nm. Light passing through the filter falls on a photocell. The photocurrent thus generated is proportional to the transmittance of the contents of the cuvette at the chosen wavelength. It is sent to the COMM INTFC card where it is digitized. The digital information is sent to the Analyzer computer, then to the Workstation.

A significant refinement of the Beckman Coulter systems is the introduction of a reagent blank into the cuvette during each operating cycle. After the percent transmittance is converted to absorbance, the reagent-blank signal level provides a reference to which the sample signal is compared.

**3.6 DATAPLOT DEVELOPMENT**

The Workstation performs a series of operations on the stored digital raw values received from the flow cell to identify memberships (subpopulations) and calculate the frequency of cells within each membership group. It also produces the DataPlot displays for visual representation of the WBC Diff and Reticulocyte/RBC membership and density.

In the LH 700 Series, the Beckman Coulter AccuGate™ algorithm uses adaptive contouring methods designed for finding optimal separation between overlapping clusters of data. The AccuGate algorithm provides a statistical analysis tool that discovers the overlapping populations using nonlinear separation techniques. The newly developed adaptive gating techniques use multidimensional data to distinguish the presence of even the faintest subpopulation. The AccuGate algorithm can adapt to unusual population shifts and overlaps. It can define highly irregular separation, and signal the need for further analysis. It can then make a subsequent analysis of the identified regions and correct deficiencies in separation.

In the DataPlots, different colors represent different memberships (types of cells). Shades of colors represent density (concentration): dark colors for low density, bright colors for high density.

<b>DIFF:</b>	Lymphocytes	Blue
	Neutrophils	Purple
	Eosinophils	Orange
	Monocytes	Green
	Basophils	White
	Non-white debris	Red
<b>RETIC:</b>	RBCs	Red
	Reticulocytes	Blue
<b>Platelets</b>		Green
<b>WBCs</b>		Purple

### Two-Dimensional (2D) DataPlots

The 2D WBC Diff DataPlot shows the five memberships: lymphocytes (LY), monocytes (MO), neutrophils (NE), eosinophils (EO) and basophils (BA), plus the non-white cell populations. Cell volume (VOL), determined by the low-frequency impedance measurement, is plotted on the Y-axis; rotated light scatter (RLS) is plotted on the X-axis.

The Reticulocyte DataPlot shows mature red cells and Reticulocytes. Cell volume is plotted on the Y-axis, and linear light scatter (LLS) is plotted on the X-axis.

### Three-Dimensional (3D) DataPlots

The 3D DataPlot view classifies by density, light scatter and opacity. The axes are color coded.

On the WBC differential DataPlot the axes are:

Volume = green  
RLS (rotated light scatter) = red  
OP (opacity) = blue

On the Retic DataPlot the axes are:

Volume = green  
LLS (linear light scatter) = red  
OP (opacity) = blue

## 3.7 PARAMETERS AND THEIR DERIVATION

Mathematical expressions in this section are in U.S. units of measurement. Parameter units can be changed to other International System of Units (SI) through Setup on the LH 700 Series Workstation.

### White Blood Cell (WBC) Count

WBC is the number of leukocytes measured directly, multiplied by the calibration factor. This number is expressed as

$$\text{WBC} = n \times 10^3 \text{ cells} / \mu\text{L}$$

### Red Blood Cell (RBC)

RBC is the number of erythrocytes measured directly, multiplied by the calibration factor. This number is expressed as

$$\text{RBC} = n \times 10^6 \text{ cells} / \mu\text{L}$$

### Hemoglobin (Hgb) Concentration

The transmittance of light (525 nm wavelength) through the lysed WBC solution in the hemoglobin cuvette is compared to the transmittance of the same light through a reagent blank. The system converts this ratio to an Hgb value in g/dL using a calibration factor.

Weight (mass) of hemoglobin determined from the degree of absorbance found through photocurrent transmittance is:

$$\text{HGB (g/dL)} = \text{Constant} \times \log_{10} \frac{\text{Reference \%T}}{\text{Sample \%T}}$$

### **Mean Corpuscular Volume (MCV)**

MCV is the average volume of individual erythrocytes derived from the RBC histogram. The system multiplies the number of RBCs in each channel by the size of the RBCs in that channel. The products of each channel between 36 fL and 360 fL are added. This sum is divided by the total number of RBCs between 36 fL and 360 fL. The analyzer then multiplies by a calibration constant and expresses MCV in femtoliters.

### **Hematocrit (Hct)**

Hct is the relative volume of packed erythrocytes to whole blood, computed by the formula:

$$\text{HCT (\%)} = (\text{RBC} \times \text{MCV}) / 10$$

### **Mean Corpuscular Hemoglobin (MCH)**

MCH is the weight of hemoglobin in the average erythrocyte, computed by the formula:

$$\text{MCH (pg)} = (\text{HGB/RBC}) \times 10$$

### **Mean Corpuscular Hemoglobin Concentration (MCHC)**

MCHC is the average weight of hemoglobin in a measured dilution, computed by the formula:

$$\text{MCHC (g/dL)} = (\text{HGB/HCT}) \times 100$$

### **Red Distribution Width (RDW)**

RDW is the size distribution spread of the erythrocyte population derived from the RBC histogram. It is the coefficient of variation (CV) expressed in percent of the RBC size distribution.

### **Platelet (Plt) Count**

Plt is the number of thrombocytes derived from the Plt histogram and multiplied by a calibration factor. This number is expressed as:

$$\text{PLT} = n \times 10^3 \text{ cells} / \mu\text{L}$$

### **Mean Platelet Volume (MPV)**

MPV is the average volume of individual platelets derived from the Plt histogram. It represents the mean volume of the Plt population under the fitted Plt curve multiplied by a calibration factor. This number is expressed in femtoliters.

### **NRBC % (Nucleated Red Blood Cells)**

NRBC enumeration is achieved through the combined use of VCS technology along with an advanced algorithm applied to the WBC count. NRBC% is defined as the number of nucleated red blood cells per 100 WBC.

### NRBC # (Nucleated Red Blood Cells)

A parameter that is calculated from the NRBC % and the total WBC count. NRBC # represents the total number of nucleated Red Blood Cells.

$$\text{NRBC (10}^3 \text{ cells } \mu\text{L)} = \text{NRBC\%} \times \text{WBC count}$$

### Differential (Diff)

#### Diff Percentages (DIFF%)

The percentages of leukocytes from each category are derived from the DataPlot.

$$\text{NE\%} = \frac{\text{no. of cells inside NE area}}{\text{no. of cells inside NE + LY + MO + EO + BA}} \times 100$$

$$\text{LY\%} = \frac{\text{no. of cells inside LY area}}{\text{no. of cells inside NE + LY + MO + EO + BA}} \times 100$$

$$\text{MO\%} = \frac{\text{no. of cells inside MO area}}{\text{no. of cells inside NE + LY + MO + EO + BA}} \times 100$$

$$\text{EO\%} = \frac{\text{no. of cells inside EO area}}{\text{no. of cells inside NE + LY + MO + EO + BA}} \times 100$$

$$\text{BA\%} = \frac{\text{no. of cells inside BA area}}{\text{no. of cells inside NE + LY + MO + EO + BA}} \times 100$$

#### Diff Absolute Numbers (DIFF#)

The absolute numbers of leukocytes in each category are computed from the WBC count and differential percentage parameters.

$$\text{NE (10}^3 \text{ cells / } \mu\text{L)} = \frac{\text{NE\%}}{100} \times \text{WBC count}$$

$$\text{LY (10}^3 \text{ cells / } \mu\text{L)} = \frac{\text{LY\%}}{100} \times \text{WBC count}$$

$$\text{MO (10}^3 \text{ cells / } \mu\text{L)} = \frac{\text{MO\%}}{100} \times \text{WBC count}$$

$$\text{EO (10}^3 \text{ cells / } \mu\text{L)} = \frac{\text{EO\%}}{100} \times \text{WBC count}$$

$$\text{BA (10}^3 \text{ cells / } \mu\text{L)} = \frac{\text{BA\%}}{100} \times \text{WBC count}$$

## Reticulocyte (Retic) Parameters

### Retic Percent (RET%)

Number of reticulocytes per 100 RBCs. This parameter is directly measured and reported as %, a percentage of RBCs. The RET% is calculated as the ratio of reticulocytes to the total number of red cells; this calculation is shown below:

$$\text{Retic\%} = \frac{\sum \text{Retics}}{\sum \text{Red\_Cells}} \times 100$$

### Retic Absolute Number (RET#)

Absolute number of reticulocytes computed from the reticulocyte percent (RET%) multiplied by the RBC count.

$$\text{RET\#} = \frac{\text{RET\%} \times \text{RBC Count}}{100}$$

### Immature reticulocyte fraction (IRF)

The percentage of the count of the highest light scatter reticulocytes relative to the total count of the reticulocytes is calculated and is reported optionally as a decimal ratio.

The IRF parameter is an indication of new reticulocyte synthesis. It is calculated from the RET% as the total number of reticulocyte events in the outermost light scattering region, corresponding to immature reticulocytes, relative to the total number of reticulocytes and is reported as this ratio.

The algorithm for calculating the IRF parameter is shown below:

$$\text{IRF} = \frac{\sum_{\text{Region - 10}} \text{Retics} - \sum_{\text{Region - 3}} \text{Retics}}{\sum_{\text{Region - 10}} \text{Retics} - \sum_{\text{Region - 0}} \text{Retics}}$$

### Mean reticulocyte volume (MRV)

The average volume of individual reticulocytes derived using VCS reticulocyte algorithm. It is calculated and is reported in femoliters.

The MRV parameter is the average volume of all reticulocytes (or the mean volume of all retic events). It is calculated from the RET% and reported in femoliters (fL).

The MRV algorithm is shown below:

$$\text{MRV} = \frac{\sum_{\text{Region - 10}} \text{Retics\_Volume}}{\sum_{\text{Region - 10}} \text{Retic} - \sum_{\text{Region - 0}} \text{Retic}}$$

## Summary of Parameter Derivations

Parameters are either:

- Measured directly.
- Derived from the RBC or Plt histogram.
- Computed.

### Measured Directly

Parameter	Method
RBC	Coulter Principle
WBC	Coulter Principle
Hgb	Photometric Measurement
DIFF% parameters	VCS Technology
RET%	VCS Technology

### Derived From RBC or Plt Histogram

Parameter	Derived From
MCV	RBC Histogram
RDW	RBC Histogram
Plt	Plt Histogram
MPV	Plt Histogram

### Derived From WBC Histogram

Parameter	Derived From
NRBC%	WBC Histogram and VCS Technology

### Computed

Hct  
MCH  
MCHC  
DIFF# parameters  
RET#  
IRF  
MRV  
NRBC#

### 3.8 XB ANALYSIS

Studies (Bull 1974, Koepke 1981) indicate that the red cell indices (MCV, MCH, and MCHC) of patient populations are stable over time.<sup>39,40</sup> This stability characteristic of the indices is the basis of a quality-control technique called XB Analysis. In a manually implemented system, population means (target values) are established by analyzing as large a sample as possible, at least 250, but ideally 1,000 blood samples. (The XB Analysis used in the Workstation does all the calculating automatically.) Once the target values have been established, the XB Analysis can be applied using quite small batches from the patient population. A 20-patient sample batch is a typical size, and can be used in the LH 700 Series Workstation.

Here is the XB formula.

$$X(B,i) = X(B,i - 1) + \text{SGN} \left\{ \sum_{j=1}^N \text{SGN}[X(j,i) - X(B,i - 1)] \times \sqrt{|X(j,i) - X(B,i - 1)|} \right\} \times F$$

$$F = \frac{\left\{ \sum_{j=1}^N \text{SGN} \left[ X(j,i) - X(B,i - 1) \times \sqrt{|X(j,i) - (B,i - 1)|} \right] \right\}^2}{N^2}$$

Where:

- X(B,i) = ith XB value
- X(B, i-1) = (i-1)th XB value
- X(j,i) = the jth X value in the ith batch
- SGN = the arithmetic sign of number in parentheses
- N = number of samples in the batch
- x = symbol used to represent multiplication

The formula is easily implemented with a computer. Its function is to enable reliable estimates of the values for these parameters to be made for a population from small samples of that population. It is superior to the traditional moving average because it reacts quickly to changes. Small batch sizes allow for more frequent, therefore tighter quality control. The formula both trims the data by giving less weight to outliers, and smoothes it by incorporating information from the previous patient batch in the analysis of the current batch. As each sample is processed, the mean of the previous set of samples is subtracted from each of the red cell indices. The square root of this deviation (difference between the means) is stored. After 20 samples have been processed, the sum of the square roots is divided by 20. The result is squared to recover the mean (average) deviation. The individual deviations carry a positive or negative sign, so then it can be added to or subtracted from the corresponding previous means. The resulting new mean is then used for the succeeding batch of 20 samples.

The hematology system is considered "in control" when the batch means are within established limits of the target values. Using the XB Analysis, the direction and amount of change due to the instrument, the reagent, flagged samples or sample handling can be detected. Because of the characteristic appearance of the graphs of the XB results, it is also often possible to identify changes.

The LH 700 Series Workstation calculates and displays the percent difference between each batch mean and its corresponding preset target value. The percent difference is derived as follows:

#### MCV

$$\text{percent diff} = \left( \frac{\text{MCV Batch Mean}}{\text{MCV Target Value}} - 1 \right) \times 100$$

#### MCH

$$\text{percent diff} = \left( \frac{\text{MCH Batch Mean}}{\text{MCH Target Value}} - 1 \right) \times 100$$

#### MCHC

$$\text{percent diff} = \left( \frac{\text{MCHC Batch Mean}}{\text{MCHC Target Value}} - 1 \right) \times 100$$

### Adjusting Initial XB Target Values

The recommended target values for initial entry are:

MCV	89.5
MCH	30.5
MCHC	34.0

As samples are run and laboratory values established, the recommended target values can be adjusted to fit your laboratory's population. After 20 XB batches have been analyzed, calculate the mean and CV% for each of the XB indices. The mean values should not differ from the target values by more than 3%, and the CV should be less than 1.5%. If the CVs are less than 1.5% and the means are less than 3% different from the target values, use the calculated means as new target values.

If the CVs are greater than 1.5%, or the mean values are greater than 3% different from the recommended target values, there may be an instrument or population problem. In this case, repeat this procedure using the next 20 XB batches. If the indices themselves are stable in a hospital population, then any deviation from the Target Values and Action Limits may point to an instrument or reagent problem. These problems would involve the parameters directly measured by the instrument and used to calculate the red cell indices. Table 3.1 lists the directly measured parameters that would be involved with out-of-limits XB batch values for each of the red cell indices.

If the XB indices are still out-of-limits, you should investigate the instrument and reagent systems associated with the directly-measured parameter(s) as indicated by Table 3.1 and call your Beckman Coulter Service Representative.

**Table 3.1 Effect of Directly-Measured Parameters on the Red Cell Indices**

Index	Directly-Measured Parameter					
	MCV		RBC		HGB	
	Increased	Decreased	Increased	Decreased	Increased	Decreased
MCV	HIGH	LOW	NORMAL	NORMAL	NORMAL	NORMAL
MCH	NORMAL	NORMAL	LOW	HIGH	HIGH	LOW
MCHC	LOW	HIGH	LOW	HIGH	HIGH	LOW

See the Glossary for terms used with the XB Analysis.

## 4.1 PHYSICAL SPECIFICATIONS

### Dimensions

Unit	Height*	Width*	Depth*	Weight
Analyzer/Diluter	88.9 cm (35 in.)	101.6 cm (40 in.)	61 cm (24 in.)	93.2 k (205 lb)
Power Supply	59 cm (23.3 in.)	35.5 cm (14 in.)	60 cm (24 in.)	56.7 k (125 lb)

\* ±5.1 cm (2 in.)

### Power

#### Input

Power Supply:	90 - 264 Vac, 47 - 63 Hz
Workstation:	90-135 Vac, 47-63 Hz or 180-265 Vac, 47-63 Hz

#### Consumption

2080 W (5500 BTU/h) maximum

Installation Category: per IEC 1010-1, Category II

#### Temperature

Ambient operating range: 15.5 to 32°C (60 to 90°F)

#### Humidity

0 to 95% without condensation

### Sample Stability

Refer to [Heading 4.3, PERFORMANCE CHARACTERISTICS on page 8](#).

#### CBC/Diff Parameters

- Stored at room temperature (23.9°C or 75°F), up to 24 hours after collection
- Stored between 2 and 8°C (35.6 and 46.4°F), up to 48 hours after collection

#### Reticulocyte Parameters

- Stored at room temperature (23.9°C or 75°F), up to 24 hours after collection
- Stored between 2 and 8°C (35.6 and 46.4°F), up to 72 hours after collection

#### NRBC Parameters

- Stored at room temperature (23.9°C or 75°F), up to 24 hours after collection
- Stored between 2 and 8°C (35.6 and 46.4°F), up to 24 hours after collection

### **Sample Storage**

Room temperature (23.9°C, 75°F) up to 24 hours

### **Sample Type**

Anticoagulated human whole blood

### **Recommended Anticoagulant**

K<sub>2</sub>EDTA or K<sub>3</sub>EDTA

## **SAMPLING MODES**

### **Aspiration**

Automatic cap-piercing, whole blood

Manual, open vial, whole blood and diluted specimens.

### **Test/Cycle**

CBC

CBC/DIFF

CBC/RETIC

CBC/DIFF/RETIC

RETIC

## **THROUGHPUT, AUTOMATIC MODE**

Typical throughput performance is described as "average" for samples exhibiting parameter levels within the normal range, and "maximum" for samples with elevated parameter levels. Approximate throughput performance data, not including sample preparation, is:

<b>LH 700 Series</b>	<b>Maximum Sample/hour</b>
CBC	110
CBC/Diff	110
CBC/Diff/Retic	45

<b>LH 755 (LH 700 Series with SlideMaker and SlideStainer)</b>	<b>Maximum Sample/hour</b>
CBC	105
CBC/Diff	100
CBC/Diff/Retic	45

Throughput requirements are based on the sample criteria in Table 4.1.

**Table 4.1 Throughput Sample Criteria**

Parameter	RETIC modes	CBC	CBC/DIFF
WBC	N/A	> 7.0 x 10 <sup>3</sup> /μL	> 7.0 x 10 <sup>3</sup> /μL
RBC	> 5.0 x 10 <sup>6</sup> /μL	> 5.0 x 10 <sup>6</sup> /μL	> 5.0 x 10 <sup>6</sup> /μL
Plt	N/A	> 300.0 x 10 <sup>3</sup> /μL	> 300.0 x 10 <sup>3</sup> /μL

**Sample Volume Aspirated**

Automatic, cap-piercing mode: maximum 300 μL  
 Manual, open-vial mode: maximum 200 μL  
 With optional LH 700 Series SlideMaker 550 uL

**Waste**

20-liter waste container

**Pneumatic Supplies (Internally Regulated)**

Pressure = 60 psi (pounds per square inch)

Vacuum = 22 in. Hg (inches of mercury) minimum at sea level

**Calibration Stability**

Electronic measurement system: < 1% per month

Variation with temperature: If ambient room temperature changes by less than 5.5°C (10°F) from the calibrating temperature, and the temperature is within the temperature specifications, then the LH 700 Series does not require calibration. Under these conditions, the expected variation in calibration factors is:

WBC	<1.25%	MCV	<1.18%
RBC	<0.70%	Plt	<2.70%
Hgb	<0.78%	MPV	<5.00%

---

**IMPORTANT** The operating temperature influences the rate of kinetic reactions. The online help states that the LH 700 SERIES should be recalibrated whenever the ambient temperature changes by 10 degrees Fahrenheit. If you have to recalibrate due to a large change in laboratory ambient temperature, you should re-evaluate the differential flagging sensitivity settings for your typical patient population.

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**LH 700 Series Workstation Storage**

Patient results,

- up to 20,000 with numeric, graphic and list mode data.

**4.2 PERFORMANCE SPECIFICATIONS--LH 700 Series**

The LH 700 Series consists of three subsystems, designated as "CBC" (Complete Blood Count), "WBC Differential" and "Retics." The CBC subsystem is based on the established Coulter principles of automated cell counting. The WBC differential subsystem is based on

the Coulter principles of leukocyte differential counting as embodied in the COULTER VCS. The Retics subsystem is based on the Coulter volume, conductivity and light scatter technology.

Performance specifications stated apply only to an instrument that has been properly maintained as indicated in the COULTER LH 700 Series documentation, using the recommended reagents.

If the average room temperature should change more than 5.5°C (10°F) from the calibrating temperature, verify calibration and recalibrate if necessary to ensure conformance to specifications.

**Precision**

**Within-Run Precision**

Within-run Precision is based on at least 31 determinations of the same sample. See Table 4.2.

**Table 4.2 Within-Run Precision (n = 31)**

Parameter	@ Approximate Level	Limit
WBC	9 to 11 x 10 <sup>3</sup> cells/μL	≤1.7% CV
RBC	4.5 to 5.5 x 10 <sup>6</sup> cells/μL	≤0.8% CV
Hgb	14 to 16 g/dL	≤0.8% CV
MCV	80 to 90 fL	≤0.8% CV
RDW	12 to 14%	≤2.2% CV
Pit	280 to 320 x 10 <sup>3</sup> cells/μL	≤3.3% CV
Pit	90 to 110 x 10 <sup>3</sup> cells/μL	≤6.6% CV
Pit	10.0 to 15.0 x 10 <sup>3</sup> cells/μL	≤14.0% CV
MPV	8 to 10 fL	≤2.2% CV
NE%	50 to 60%	2SD ≤3.0
LY%	25 to 35%	2SD ≤3.0
MO%	5 to 10%	2SD ≤2.0
EO%	2 to 5%	2SD ≤1.0
BA%	0.5 to 1.5%	2SD ≤1.0
RET%	0.00 to 0.49%	1SD ≤0.23 or 16.5% CV
RET%	0.50 to 1.49%	1SD ≤0.23 or 14.5% CV
RET%	1.50 to 4.00%	1SD ≤0.68 or 11.0% CV
RET%	4.01 to 15.00%	1SD ≤0.68 or 5.5% CV

**Accuracy**

**Accuracy Qualification**

Accuracy is tested on a minimum of 150 non-flagged, morphological normal samples for CBC and Diff, and 50 non-flagged samples for Retics. Distributional abnormal samples should be

included. NCCLS H20-A<sup>42</sup> criteria should be used in the sample selection and manual differential examination.

**Accuracy, CBC**

For the CBC parameters, the LH 700 Series can be adjusted within the resolution of the readout to agree with a predetermined reference value at any point in the operating range. Accuracy was tested against a predicate hematology analyzer. Table 4.3 lists the specifications for the mean difference and mean percent difference.

**Table 4.3 Accuracy, CBC**

Parameter		Mean Difference	Mean Percent Difference
WBC	0.00 to 100.0	±0.2	3.5%
	100.1 to 400.0	N/A	9.0%
RBC		±0.05	2.0%
Hgb		±0.2	3.0%
MCV		N/A	2.0%
RDW		±0.5	5.0%
Plt		±10	7.0%
MPV		N/A	5.0%

**Accuracy, WBC Differential**

Accuracy of the WBC differential should be determined by comparison against either or both of two comparator methods. Accuracy can be tested against the manual differential reference method (NCCLS H20-A [n = 400]). Accuracy was tested against a predicate hematology analyzer. When the LH 700 Series is compared to either method, using non-flagged, morphological normal samples, expect results within tolerance limits listed in Table 4.3.

**Table 4.4 Accuracy Tolerance Limits, WBC Differential**

Cell Type	Mean Difference % using NCCLS H20-A	Mean Difference % using predicate analyzer
Lymphocyte	±3.0	±1.5
Monocyte	±3.0	±1.0
Neutrophil	±2.0	±2.0
Eosinophil	±1.0	±0.5
Basophil	±1.0	±0.5

**Accuracy, Reticulocyte**

Reticulocyte parameter (RET%) accuracy can be determined against any of three comparator methods. Accuracy can be tested against a clinical flow cytometer following the guidelines of NCCLS H44-P. Accuracy was tested against the manual method using NCCLS H16-P. See Table 4.4. Accuracy was tested against a predicate hematology analyzer. See Table 4.5

**Table 4.5 Accuracy, Reticulocyte, using NCCLS H16-P or clinical flow cytometer**

Parameter/Population	Mean Difference	SD of Difference
RET% Range 0.00 to 15.00% <30% have a RET% >4.00	±1.00	≤1.50
RET% Range 0.00 to 30.00% <50% have a RET% >4.00 <10% have a RET% >20.00	±1.50	≤3.00

**Table 4.6 Accuracy, Reticulocyte, using a predicate hematology analyzer**

Parameter/Population	Mean Difference
RET%	±0.5

### Linearity

When tested using a stable material having no interfering substances, the LH 700 Series value should be equal to the expected value within the limits given in Table 4.7. To minimize the effects of imprecision, take multiple readings at each dilution point. Subtract the background count from the values obtained.

**Table 4.7 Linearity Limits**

Parameter	Linearity Range	Limits (mean difference or % difference, whichever is greater)
WBC x 10 <sup>3</sup> cells/μL	0.00 to 100.0 100.1 to 400.0	±0.2 or 3.0% ±9%
RBC x 10 <sup>6</sup> cells/μL	0.00 to 8.00	±0.05 or 2.0%
Hgb g/dL	0.0 to 25.0	±0.2 or 3.0%
Plt x 10 <sup>3</sup> cells/μL	0 to 3000	±10 or 7.0%

### Background

Background counts for CBC parameters are performed during the Startup cycle. The background limits are

WBC	≤0.20
RBC	≤0.01
Hgb	≤0.20
Plt	≤3.00
DIFF	≤100 events
RETIC	≤600 events

### Carryover

High-to-Low Carryover on the LH 700 Series should meet these limits:

WBC	≤2.0%
RBC	≤1.0%
Hgb	≤2.0%
Plt	≤2.0%
DIFF	≤200 events
RETIC	≤600 events

### Operating and Reportable Ranges

The operating ranges reflect the range of values over which the instrument displays, prints and transmits results. Values that are between the linear range and the operating range, and values outside the reportable range, are displayed, printed and transmitted with an over linear range flag (+). Values that are above the operating range are inhibited, and the value is replaced by pluses (+++++). The reportable range identifies the values where the instrument is accurate, and reflects the range studied in accuracy testing. See Table 4.8.

**Table 4.8 Operating and Reportable Ranges**

Parameter	Operating Range	Reportable Range
WBC	0.00 to 900.00 x 10 <sup>3</sup> cells/μL	0.00 to 400.00 x 10 <sup>3</sup> cells/μL
RBC	0.00 to 20.00 x 10 <sup>6</sup> cells/μL	0.00 to 8.00 x 10 <sup>6</sup> cells/μL
Hgb	0.0 to 99.9 g/dL	0.0 to 25.0 g/dL
Hct	0.0 to 99.9%	N/A
MCV	0.0 to 300.00 fL	0.0 to 150.0 fL
MCH	0.0 to 99.9 pg	N/A
MCHC	0.0 to 99.9 g/dL	N/A
RDW	0.0 to 99.9%	N/A
Plt	0.00 to 5000 x 10 <sup>3</sup> cells/μL	0.00 to 3000 x 10 <sup>3</sup> cells/μL
Pct	0.0 to 9.999%	N/A
MPV	0.0 to 99.9 fL	N/A
PDW	0.0 to 99.9%	N/A
NE%, LY%, MO%, EO%, BA%	0 to 100%	0 to 100%
NE#, LY#, MO#, EO#, BA#	0.00 to 900.00 x 10 <sup>3</sup> cells/μL	0.00 to 400.00 x 10 <sup>3</sup> cells/μL
RET%	0.00 to 100%	0.00 to 30.0%
RET#	0.00 to 999.9 x 10 <sup>6</sup> cells/μL	0.0000 to 0.7500 x 10 <sup>6</sup> cells/μL
NRBC%	0.0 or 2.0 to 600 %	0.0 or 2.0 to 600 %

### Mode-to-Mode Comparison

Automatic (closed vial) and Manual (open vial) mode-to-mode differences of the means of 10 hematologically "normal" blood specimens, measured in triplicate, or 50 samples from accuracy testing should be less than or equal to the following:

WBC     ± 0.4 x 10<sup>3</sup> cells/µL or 5%, whichever is greater  
RBC     ± 0.2 x 10<sup>6</sup> cells/µL or 2%, whichever is greater  
Hgb     ± 0.3 g/dL or 2%, whichever is greater  
Plt     ± 20 x 10<sup>3</sup> cells/µL or 7%, whichever is greater

In addition, 95% of the individual differences should be within the stated limits.

### 4.3 PERFORMANCE CHARACTERISTICS

The information in this section describes *typical performance* for an instrument that has been properly maintained as indicated in the COULTER LH 700 Series documentation, using the recommended reagents.

#### Sample Stability

The following tables show the average results for specimens from five normal donors collected in K<sub>3</sub>EDTA†. The specimens were stored at room temperature and cold temperature. For this study, room temperature was 72 - 73° (22 - 23°C) and cold temperature was 37 - 39°F (3 - 4°C). Room temperature specimens were mixed for 22 inversions and then immediately analyzed. Upon removal from the refrigerator, the cold specimens were mixed for 22 complete inversions and analyzed within five minutes

**Table 4.9 CBC Sample Stability, Room Temperature**

Hours	WBC	RBC	Hgb	MCV	RDW	PLT	MPV
1	8.45	4.416	13.49	86.4	13.84	248.7	8.59
4	8.75	4.482	13.65	86.8	13.87	249.2	8.76
8	8.76	4.485	13.69	87.9	14.17	247.7	8.68
16	8.94	4.468	13.65	88.5	14.38	256.0	8.78
24	8.81	4.462	13.61	89.5	14.59	261.0	8.96

**Table 4.10 CBC Sample Stability, Cold Temperature**

Hours	WBC	RBC	Hgb	MCV	RDW	PLT	MPV
1*	8.45	4.416	13.49	86.4	13.84	248.7	8.59
4	8.54	4.438	13.49	86.3	13.90	252.8	8.64
8	8.59	4.437	13.51	87.0	13.77	244.9	8.65
16	8.65	4.446	13.54	87.1	13.84	240.4	8.80
24	8.71	4.419	13.51	87.5	13.80	235.6	8.96
36	8.68	4.451	13.52	87.5	13.67	232.1	9.07
48	8.48	4.437	13.62	87.8	13.76	227.0	9.21

\*Results analyzed @ Room Temperature

**Table 4.11 DIFF% Sample Stability, Room Temperature**

Hours	NE	LY	MO	EO	BA
1	56.02	34.36	7.45	1.82	0.35
4	55.60	34.76	7.35	1.90	0.39
8	55.15	35.58	7.25	1.63	0.40
16	55.92	35.45	6.68	1.63	0.32
24	56.56	35.25	6.30	1.56	0.33

**Table 4.12 DIFF% Sample Stability, Cold Temperature**

Hours	NE	LY	MO	EO	BA
1*	56.02	34.36	7.45	1.82	0.35
4	55.96	35.24	6.73	1.73	0.34
8	56.63	34.92	6.32	1.76	0.37
16	58.08	34.96	5.05	1.65	0.27
24	58.71	33.57	5.80	1.63	0.29
36	60.58	31.41	6.33	1.47	0.22
48	61.00	31.14	6.43	1.11	0.32

\*Results analyzed @ Room Temperature

**Table 4.13 DIFF# Sample Stability, Room Temperature**

Hours	NE	LY	MO	EO	BA
1	5.45	2.32	0.54	0.11	0.03
4	5.60	2.45	0.53	0.12	0.03
8	5.58	2.51	0.53	0.10	0.03
16	5.79	2.54	0.47	0.10	0.02
24	5.71	2.48	0.46	0.10	0.03

**Table 4.14 DIFF# Sample Stability, Cold Temperature**

Hours	NE	LY	MO	EO	BA
1*	5.45	2.32	0.54	0.11	0.02
4	5.50	2.40	0.50	0.11	0.03
8	5.59	2.43	0.44	0.11	0.03
16	5.72	2.45	0.36	0.10	0.02
24	5.81	2.35	0.42	0.11	0.02
36	5.94	2.20	0.43	0.09	0.01
48	5.77	2.16	0.46	0.06	0.02

\*Results analyzed @ Room Temperature

**Table 4.15 RETIC Sample Stability, Room Temperature**

Hours	RETIC%	RETIC#	MRV	IRF
1	1.771	0.078	103.87	0.356
4	1.711	0.076	105.09	0.353
8	1.683	0.075	105.21	0.336
16	1.743	0.077	105.99	0.341
24	1.595	0.070	107.69	0.352

**Table 4.16 RETIC Sample Stability, Cold Temperature**

Hours	RETIC%	RETIC#	MRV	IRV
1*	1.771	0.078	103.87	0.356
4	1.900	0.084	103.92	0.354
8	1.874	0.083	103.94	0.347
16	1.957	0.086	105.48	0.357
24	1.907	0.084	104.63	0.360
36	1.670	0.074	106.86	0.360
48	1.644	0.073	104.70	0.345
60	1.674	0.074	106.08	0.374
72	1.786	0.079	105.84	0.368

\*Results analyzed @ Room Temperature

**Table 4.17 NRBC Sample Stability, Room Temperature**

Hours	NRBC%	NRBC#
1	0.00	0.00
4	0.00	0.00
8	0.00	0.00
16	0.00	0.00
24	0.00	0.00

**Table 4.18 NRBC Sample Stability, Cold Temperature**

Hours	NRBC%	NRBC#
1*	0.00	0.00
4	0.00	0.00
8	0.00	0.00
16	0.00	0.00
24	0.00	0.00

Results analyzed @ Room Temperature

### WBC Differential Flagging Stability

The tables below indicate flagging at timed intervals using mid-level flagging sensitivity with NRBC enumeration enabled. For additional information regarding flagging sensitivity, refer to Flagging Preferences section of the LH 700 SERIES System Operator's Guide.

**Table 4.19 Differential Suspect Flagging at Room Temperature**

Sample #	1 hour	4 hour	8 hour	16 hour	24 hour
1	Imm Ne 1	Imm Ne 1	Imm Ne 1	Imm Ne 1	Mo Blast, Imm Ne 1, Imm Ne 2
2	*	*	*	*	Imm Ne 2
3	*	*	*	*	*
4	*	*	*	*	*
5	*	*	*	*	*
*Indicates that there were no flags present.					

**Table 4.20 Differential Suspect Flagging at Cold Temperature**

Sample #	4 hour	8 hour	16 hour	24 hour	36 hour	48 hour
1	Imm Ne 1	*	Imm Ne 1	Imm Ne 1	Imm Ne 1	Imm Ne 1
2	*	*	*	*	*	*
3	*	*	*	PLT Clumps	*	*
4	*	*	*	*	*	*
5	*	*	*	*	*	*
*Indicates that there were no flags present.						

### NRBC Paired Sample Imprecision

Table 4.21 shows the NRBC results for 60 replicate determinations of whole blood in K<sub>3</sub>EDTA, Automatic Mode. The individual differences for each specimen were calculated (run1 - run 2) and the Mean and Standard Deviation of the Differences were determined.

**Table 4.21 NRBC Imprecision Characteristics, Paired Sample Analysis - Replicate 1 vs. Replicate 2**

Parameter	Units	N	Population Minimum	Population Maximum	Mean Difference	SD of Difference
NRBC	%	488	0.00	184.44	-0.14	1.69

### NRBC Accuracy Characteristics

NRBC Accuracy Characteristics was defined as the agreement between the LH 700 Series and the results given by the reference method (NCCLS H20A) and a predicate hematology analyzer.

All NRBC % results greater than or equal to two and less than or equal to fifteen will be flagged with an R symbol. Beckman Coulter, Inc. recommends a slide review as per your

individual laboratory protocol. The presence of this flag does not affect the correction of the white blood cell count in the presence of NRBC's.

Table 4.22 shows the NRBC results for whole blood specimens with and without NRBC collected in a salt of EDTA.

**Table 4.22 NRBC Accuracy Characteristics, Compared Samples**

Parameter	Units	N	Population Minimum	Population Maximum	Mean Difference
NRBC*	%	496	0.0	344.00	-0.84
NRBC**	%	56	0.0	178.00	4.00

\* NCCLS

\*\* predicate hematology analyzer

### Platelet Accuracy Characteristics

Table 4.23 shows the Plt results for whole blood specimens collected in a salt of EDTA.

**Table 4.23 Plt Accuracy Characteristics vs ICSH/ISLH Platelet Method**

Parameter	Units	N	Population Minimum	Population Maximum	Mean Difference	Mean % Difference
PLT	$\times 10^3$ cells/L	104	0.37	1138.00	-4.42	19.16

### Reference Ranges

A Normal Range study was conducted to assess the Reference Ranges for the LH 700 Series. Whole-blood samples were collected from 131 donors (males and females). The selection of donors was consistent with guidelines stated in NCCLS, C28-A.

**Table 4.24 Normal Population Study**

Parameter	Units	Gender	Mean	95% Confidence Low Limit	95% Confidence High Limit
WBC	$\times 10^3$ cells/ $\mu$ L	M/F	6.03	3.53	9.57
RBC	$\times 10^6$ cells/ $\mu$ L	M/F	4.64	3.96	5.50
Hgb	g/dL	M/F	13.83	12.14	16.27
Hct	Ratio	M/F	40.22	35.39	47.19
MCV	fL	M/F	87.0	76.9	94.7
MCH	pg	M/F	29.98	25.69	32.95
MCHC	g/dL	M/F	34.42	33.34	35.47
RDW	%	M/F	13.30	12.09	15.19
Plt	$\times 10^3$ cells/ $\mu$ L	M/F	248.65	169.06	397.10
MPV	fL	M/F	9.31	7.54	11.24
NE	%	M/F	59.78	41.71	73.74

**Table 4.24 Normal Population Study (Continued)**

LY	%	M/F	28.67	18.39	44.42
MO	%	M/F	7.89	4.69	12.66
EO	%	M/F	2.96	0.75	8.06
BA	%	M/F	0.62	0.17	1.37
NE	x 10 <sup>3</sup> cells/ $\mu$ L	M/F	3.74	1.47	7.06
LY	x 10 <sup>3</sup> cells/ $\mu$ L	M/F	1.84	0.65	4.25
MO	x 10 <sup>3</sup> cells/ $\mu$ L	M/F	0.50	0.17	1.21
EO	x 10 <sup>3</sup> cells/ $\mu$ L	M/F	0.20	0.03	0.77
BA	x 10 <sup>3</sup> cells/ $\mu$ L	M/F	0.04	0.01	0.13
RET	%	M/F	1.18	0.45	2.28
RET	x 10 <sup>6</sup> cells/ $\mu$ L	M/F	0.05	0.02	0.11
MRV	fl	M/F	114.15	102.73	124.89
IRF	Ratio	M/F	0.27	0.163	0.362

## Known Interfering Substances

### CBC

All	Misleading results can occur if the specimen is not properly collected, stored or transported. Beckman Coulter recommends that you follow NCCLS or equivalent procedures to ensure proper specimen collection, storage and transport. Always follow manufacturer's recommendations when using microcollection devices for capillary specimen collection.
All	Misleading results can occur if specimens contain clots. Always use good laboratory practices for inspecting specimens for clots and verifying results.
All	Misleading results can occur if the specimen is not properly mixed. Always use good laboratory practices to ensure specimens are appropriately mixed. Do not bypass or circumvent the automated mixing process used on the LH 700 Series.
WBC	NRBCs, giant platelets, platelet clumps, malarial parasites, precipitated elevated proteins, microlymphoblasts, very small lymphocytes, fragmented white cells, agglutinated white cells, lyse resistant red cells, unlysed particles > 35 fL in size.
RBC	Very high WBC count, high concentration of very large platelets, auto-agglutination .
Hgb	Very high WBC count, severe lipemia, heparin, certain unusual RBC abnormalities that resist lysing.
MCV	Very high WBC count, high concentration of very large platelets, auto-agglutination.
RDW	Very high WBC count, high concentration of very large platelets, auto-agglutination.

Plt	Giant platelets, platelet clumps, white cell fragments, electronic noise, very small red cells, red cell fragments
Hct	Known interferences related to RBC and MCV.
MCH	Known interferences related to Hgb and RBC.
MCHC	Known interferences related to Hgb, RBC and MCV.
<b>NRBC</b>	Known interferences related to lyse resistant red cells, platelet clumps, giant platelets, malarial parasites, very small or multi-population lymphocytes, precipitated elevated proteins. NRBC > 35 fL may not be counted and may result in a false negative condition. If an NRBC% is reported as zero and there are any suspect messages or parameter codes, Beckman Coulter, Inc. recommends a slide review per your laboratory protocol.  Cord blood is a recognized source of high numbers of NRBC. However, due to the possible matrix effect of the sample constituents that may be introduced during sample, cord blood is not recommended for NRBC studies.
<b>Differential</b>	Hypogranular granulocytes, agranular granulocytes, lyse-resistant red cells, very small or multi-population lymphocytes, elevated triglycerides, precipitated elevated proteins.
<b>Reticulocytes</b>	Erythrocyte inclusions stained by New Methylene Blue, if sufficiently numerous within a sample, and some hemoglobinopathies (SS, SC) might affect the accuracy of the reticulocyte enumeration. <sup>42</sup>

## 4.4 BAR-CODE SYMBOLOGY OVERVIEW

A bar-code symbol is a series of adjacent bars and spaces. The widths of the bars and spaces represent the actual data encoded within the symbol. A device, usually called a scanner, or reader is used to decode this data into human readable information. A bar-code symbology, such as Interleaved 2-of-5, describes the unambiguous conventions or rules used for the way in which data is encoded into the arrangement of the adjacent bars and spaces.

A symbology is defined by the following characteristics:

### Character Set

This refers to the range of data characters that can be encoded for a specific symbology. There are essentially three types:

- Numeric—The symbology can encode only numeric characters.
- Alphanumeric—Can encode the full alpha and numeric characters of the ASCII character set.
- 128—Can encode all the 128 available ASCII characters.

### Symbology Type

There are essentially two types:

- Discrete—Each character is separated from the next by an [intercharacter gap](#). This allows each character to be decoded independently. Every character has a bar on each end.
- Continuous—Has no intercharacter gaps. Every character begins with a bar and ends with a space.

### Fixed or Variable Length

Some symbologies can only encode messages of a fixed length. Some can be used to encode variable length data. Some should only be used as a fixed length to increase the security of data decoding.

### Self-Checking

A self-checking symbology has the ability to prevent character transposition due to a single printing defect.

### Start Code, Stop Code

Start and Stop Codes are bar and space patterns that are placed at the beginning and end of the encoded data. A start code indicates the beginning of the symbol, and indicates the start of the encoded information. The stop code indicates the end of the symbol and marks the end of the encoded information.

### Check Character, Checksum Algorithm

Virtually all bar-code symbologies use a check character. The check character is derived by a mathematical calculation using the characters in the encoded information. It is used by the scanning device to verify that the correct information has been decoded. It is highly recommended that a check character, also called a checksum algorithm, is used.

### Quiet Zone, Quiet Area

The Quiet Zone is a clear area on either side of the start and stop codes. It helps the scanning device establish the reflectivity characteristics of the label.

## 4.5 BAR-CODES AND THE LH 700 Series

The LH 700 Series supports a wide variety of [bar-code symbologies](#). When choosing a symbology to use in your laboratory, it is important to understand the different bar-codes available. Some bar-codes are more stable, and therefore less prone to misreads than others.

The following information is intended as a guide to help you in your selection process.

---

**IMPORTANT** Risk of misidentification. Use of poor quality, dirty, improperly placed or damaged bar-code labels could keep the instrument from reading the bar-code labels. Ensure the bar-code labels are undamaged. Ensure the bar-code labels conform to the specifications provided in Chapter 4 of the Reference manual.

---

### Checksum Algorithm

Beckman Coulter strongly recommends the use of bar-code [checksums](#) to provide automatic checks for read accuracy.

---

**IMPORTANT** Use of bar-codes is an extremely accurate and effective method of positive patient identification. Certain features, such as checksum digits, maximize accuracy in reading Codabar, Code 39 and Interleaved 2-of-5 labels. In one study, the use of checksum digits detected 97% of misread errors. Use checksums to provide protection against occasional misread errors caused by problems such as damaged or misapplied labels. If you must use bar-codes without checksums, Beckman Coulter recommends that you verify each bar-code reading to assure correct patient identification.

---

## Bar-code Symbologies Supported by the LH 700 Series

### Interleaved 2-of-5

Interleaved 2-of-5 is a high density, continuous numeric symbology. It is [self-checking](#). Every character in the symbology encodes two digits, one in the bars and one in the spaces.

This symbology is susceptible to an incorrect read due to a partial scan (a scanning path that does not include both leading and trailing [quiet zones](#)). The most common incorrect read is a shorter, but valid decoding of the information. The presence of a checksum does not eliminate this risk. It is recommended that any Interleaved 2-of-5 label contain [Bearer Bars](#). Alternatively, this label should be used with a fixed length only, with the scanning devices set to recognize labels of a specific length (for example 12 digits).

### Code 39 (Also called 3-of-9 Code)

Code 39 was the first alphanumeric symbology developed. It is a self-checking, discrete and variable length symbology. The Health Industry Bar-code Council (HIBCC) has adopted the use of a check character for health care applications. This is encouraged, particularly where print quality is less than optimum.

### Codabar

Codabar has a character set of 16 characters. It is a discrete, self-checking symbology and is most commonly used in libraries and blood banks.

### NW-7

NW-7 is very similar to Codabar. It uses the same character set and check character. The difference is in the bar and space structure.

### Code 128/USS 128

Code 128 is a continuous, variable length symbology. This symbology has 106 different printed characters.

Code 128 is character dependent. See AIM® Uniform Symbol Specification (USS) Rev. 1986 for additional required dimensional tolerances.

You must use and print a checksum character, and it must conform to the AIM USS 128 checksum generation procedure.

---

**IMPORTANT** If a laboratory uses Code 128 bar-code format, it must have checksum enabled. If the checksum is disabled, the LH software produces a bar-code that does not comply with the Code 128 standard and cannot be read by bar-code scanners.

---

Do not use these values:

Code set A - 0, 64 through 102

Code set B - 0, 95 through 102

Code set C - 100 through 102

Do not use leading or trailing spaces in the ID.

## Bar-code Tips

No bar-code symbology is perfect. You will occasionally get a read error. Below are some tips to help you get the best performance from the symbology you choose:

- Use high quality labels.
- Always use a check character.
- Use fixed length bar-code information.
- Use Bearer Bars with Interleaved 2-of-5.
- When using the handheld scanner, check that the label was decoded correctly.

## 4.6 BAR-CODE LABEL SPECIFICATIONS

### Handheld Scanner

Information about configuring the handheld scanner is located elsewhere.

### General

A bar-code consists of black lines (bars) and white lines (spaces), which are called elements.

There are narrow elements (NE) and wide elements (WE); their arrangement is determined by the code.

---

**IMPORTANT** Possible misidentified results. For accurate reading by the scanner, it is important that bar-code labels for specimen tubes adhere strictly to the specifications given in this section. Labels that meet these specifications are available from Beckman Coulter.

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### Optical Characteristics at 880 nm $\pm 10\%$ and 633 nm $\pm 10\%$

- Print Contrast Signal (PCS): 80% min.
- Reflectivity of Media (RW): 80% min.
- Reflectivity of Ink (Rb): 16% max.
- No spots or voids; no ink smearing.
- Edge roughness is included in the bar and space tolerances.

Measurement method is according to American National Standards Institute's MH10-8M-1983.

### Printing Method

Photographic, or thermal transfer.

### Label Thickness

Maximum label thickness must be such that:

- The tube's outer diameter including the label is not greater than 13.3 mm.
- The label including adhesive = 0.006  $\pm$  0.003 in.

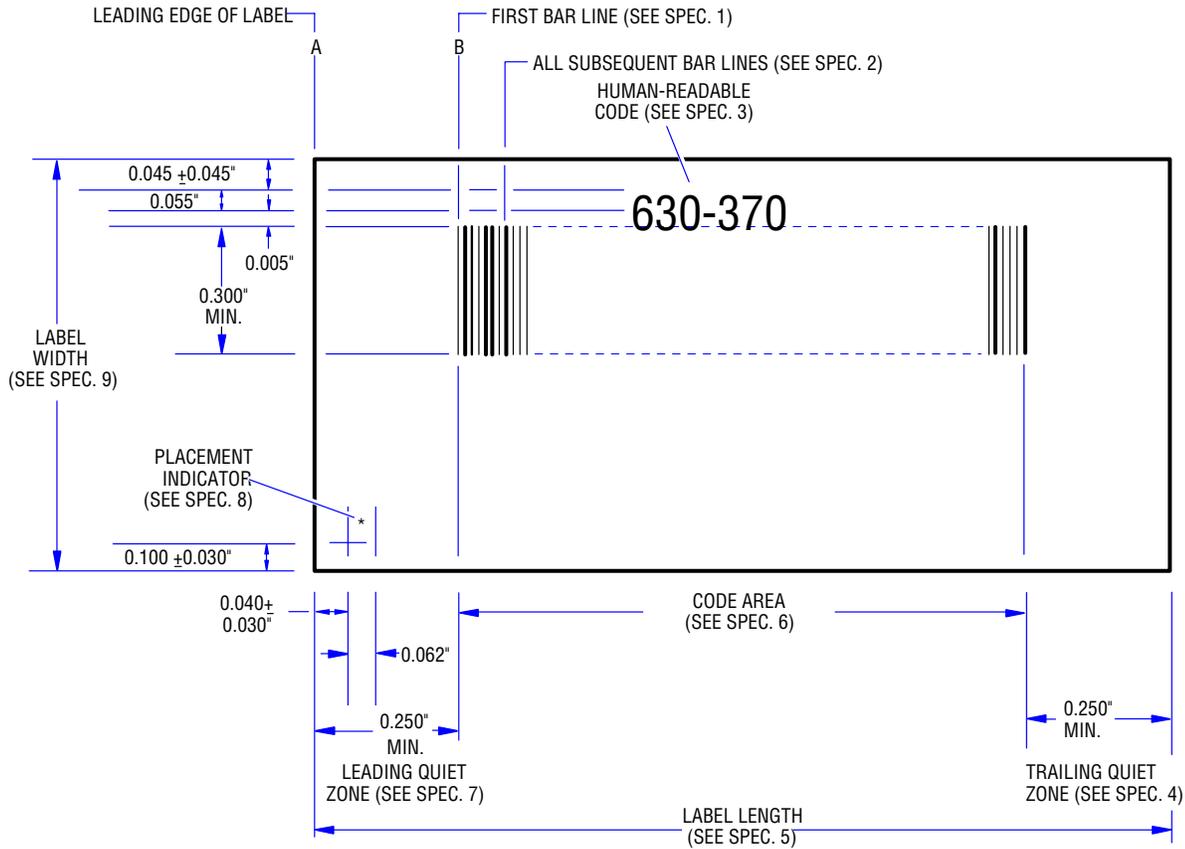
### NE/WE Ratio

Must remain constant over code length.

**Label Dimensions and Data**

The dimensional and data specifications are illustrated in Figure 4.1. Table 4.13 explains the specifications called out in Figure 4.1

**Figure 4.1 Bar-Code Label Specifications**



**Table 4.25 Bar-Code Label Specifications**

Specification Called Out in Figure 4.1	Explanation
1	The first bar of the code (B) must be parallel to the label edge (A) within 0.002".
2	All subsequent bar lines must be parallel to (B) within 0.001".
3	The human-readable code (HRC) does not include the checksum; the dash in the HRC is not encoded in the bar-code.
4	The trailing quiet zone must be 0.250" minimum.
5	The maximum label length is determined by the tube length. The scanner can accommodate labels up to 2.35". With HEMOGARD™ tubes, the maximum label length is 2.04".
6	The bar-code area contains the start character, data digits, checksum, and stop character.
7	The leading quiet zone must be 0.250" minimum.

**Table 4.25 Bar-Code Label Specifications**

8	The placement indicator shows you which end of the label goes next to the tube stopper. This is an optional feature, not a mandatory one.
9	The width of the label must leave at least a 1/8" window for viewing the contents of the tube. The maximum label width for a 10-mm diameter tube is 1.1". The minimum label width is 0.400".

**Acceptable Bar-codes**

Within the given specifications, the scanner automatically distinguishes the following bar-codes.

- Interleaved 2-of-5
- Code 39
- Codabar
- NW7
- Code 128/USS 128

The following table summarizes the code-related specifications.

**Table 4.26 Code-Related Specifications**

Code	Interleaved 2-of-5	Codabar	Code 39****	NW7	Code 128**** USS 128
Narrow element (NE) width	0.0105" ±0.001"	0.010"* Scaling Factor = 1.538*	0.010" ±0.001"	0.0105" ±0.001"	0.010" ±0.001"
Wide element/narrow element ratio (WE/NE)	(2.2 to 3): 1	N/A	(2.21 to 3): 1	(2.2 to 3): 1	(2 to 4): 1**
Intercharacter gap	No	0.010" Min.	>=NE	0.010" Min.	No
Data digits	3 to 11	3 to 9	3 to 9 (3 to 8 with HEMOGARD tubes)	3 to 9	LH 700 Series, 3 to 11 data digits SlideMaker, 3 to 9 data digits Checksum always printed***

\* According to American National Standard for bar-code specifications that yield 10 characters per inch at NE = 0.0065".

\*\* Code 128 is character dependent. See AIM® Uniform Symbol Specification Rev. 1986 for additional required dimensional tolerances.

\*\*\* You must use and print a checksum character, and it must conform to the AIM USS 128 checksum generation procedure.

**IMPORTANT** If a laboratory uses Code 128 bar-code format, it must have checksum enabled. If the checksum is disabled, the LH software produces a bar-code that does not comply with the Code 128 standard and cannot be read by bar-code scanners.

Do not use these values:

- Code set A - 0, 64 through 102
- Code set B - 0, 95 through 102
- Code set C - 100 through 102

\*\*\*\* Do not use leading or trailing spaces in the ID

## Checksum Algorithm

Beckman Coulter strongly recommends the use of bar-code checksums to provide automatic checks for read accuracy.

---

**IMPORTANT** Use of bar-codes is an extremely accurate and effective method of positive patient identification. Certain features, such as checksum digits, maximize accuracy in reading Codabar, Code 39 and Interleaved 2-of-5 labels. In one study, the use of checksum digits detected 97% of misread errors. Use checksums to provide protection against occasional misread errors caused by problems such as damaged or misapplied labels. If you must use bar-codes without checksums, Beckman Coulter recommends that you verify each bar-code reading to assure correct patient identification.

---

The algorithm for determining the checksum for each code is given below.

### Interleaved 2-of-5

This code requires 3 to 11 data digits plus a checksum.

To determine the value of the checksum character:

1. Identify even- and odd-positioned characters in the message with the right-hand message character **always** defined as an even-positioned character.
2. Sum the numeric values of the odd-positioned characters.
3. Sum the numeric values of the even-positioned characters and multiply the total by 3. Sum the odd and even totals from steps 2 and 3.
4. Determine the smallest number which, when added to the sum in step 4, results in a multiple of 10.

This number is the value of the checksum character.

5. Determine whether total number of characters (message plus checksum) is odd or even. If odd, add a leading nonsignificant zero to the message to produce an even number of characters as required by the symbology.

Example:

<b>MESSAGE</b>	1	2	5	6	7	8
<b>PARITY</b>	0	E	0	E	0	E

STEP 2:  $1+5+7=13$

STEP 3:  $(2+6+8) \times 3=48$

STEP 4:  $13+48=61$

STEP 5:  $61+9=70$

Therefore, the checksum is 9, and the final decoded message is 01256789.

### Codabar and NW7

These codes use three to eight data digits.

Note: Codabar and NW7 codes have the same character set and the same checksum algorithm. The difference between these two codes is that Codabar has 18 different bar and space dimensions, and NW7 has only NE and WE structure.

The value assigned to each of the characters is presented in the following table.

<b>CHARACTER</b>	<b>VALUE</b>	<b>CHARACTER</b>	<b>VALUE</b>
0	0	-	10
1	1	\$	11
2	2	:	12
3	3	/	13
4	4	.	14
5	5	+	15
6	6	A	16
7	7	B	17
8	8	C	18
9	9	D	19

The checksum technique is:

- The character value of a message is obtained from the above table and added together.
- This sum is divided by 16, and the remainder corresponds to the value of the checksum character.

Examples:

1.

<b>MESSAGE</b>	2	3	4	7	1	3
<b>VALUE</b>	2	3	4	7	1	3

$$2+3+4+7+1+3 = 20$$

$$20 \div 16 = 1, \text{ REMAINDER } 4$$

The value 4 corresponds to character 4; therefore, the checksum is 4 and the final decoded message is 2347134.

2.

<b>MESSAGE</b>	\$	\$	/	/	+	+	+	+
<b>VALUE</b>	11	11	13	13	15	15	15	15

$$11+11+13+13+15+15+15+15 = 108$$

$$108 \div 16 = 6, \text{ REMAINDER } 12$$

The value 12 corresponds to character :, therefore, checksum is :, and the final decoded message is: \$\$//++++:

### Japan Red Cross NW7 Decoding

This code uses three to nine data digits.

Japan Red Cross Hospitals use the following NW7 values:

CHARACTER	VALUE
0	0
1	1
2	2
3	3
4	4
5	5
6	6
7	7
8	8
9	9

The checksum technique is:

- The data digit value that is the difference between 11 and the Mod 11 sum of the weighted values of the data digits is used as the check digit. The start and stop digits are not used as part of the checksum calculation.
- NW7 is made up of 1 start digit, 9 data digits and 1 stop digit.
- The checksum digit immediately precedes the stop digit.

WEIGHTED MODULUS 11:

DIGIT POSITION (Right Justified)	12	11	10	9	8	7	6	5	4	3	2	1
WEIGHT (1)	6	3	5	9	10	7	8	4	5	3	6	2
WEIGHT (2)	5	8	6	2	10	4	3	7	6	8	5	9

The first 9 digits from the right are used for the calculation of the check digit.

To determine the value of the checksum character:

1. Determine the Modulus 11 value corresponding to each character in the message.
2. Multiply each character in the message by the corresponding Mod 11 value.
3. Add the resulting values and divide by 11.
4. Subtract the remainder from 11.
5. Determine the character that corresponds to the result from step 4. This is the checksum character.

Examples:

1. MESSAGE 011529007  
 USE WEIGHT (1): 6 3 5 9 10 7 8 4 5 3 6 2

<b>DIGIT POSITION (Right Justified)</b>	<b>0</b>	<b>0</b>	<b>0</b>	<b>0</b>	<b>1</b>	<b>1</b>	<b>5</b>	<b>2</b>	<b>9</b>	<b>0</b>	<b>0</b>	<b>7</b>
WEIGHT (1)	6	3	5	9	10	7	8	4	5	3	6	2
Result	0	0	0	0	10	7	40	8	45	0	0	14

$$0 + 10 + 7 + 40 + 8 + 45 + 0 + 0 + 14 = 124$$

$$124 \div 11 = 11, \text{ REMAINDER } 3$$

When the REMAINDER IS 0, 0 is the check digit.  
 $11 - 3 = 8$

The value 8 corresponds to character 8, therefore the checksum is 8 and the final decoded message is 0115290078

2. MESSAGE 023229006  
 USE WEIGHT (1): 6 3 5 9 10 7 8 4 5 3 6 2

<b>DIGIT POSITION (Right Justified)</b>	<b>0</b>	<b>0</b>	<b>0</b>	<b>0</b>	<b>2</b>	<b>3</b>	<b>2</b>	<b>2</b>	<b>9</b>	<b>0</b>	<b>0</b>	<b>6</b>
WEIGHT (1)	6	3	5	9	10	7	8	4	5	3	6	2
Result	0	0	0	0	20	21	16	8	45	0	0	12

$$0 + 20 + 21 + 16 + 8 + 45 + 0 + 0 + 12 = 122$$

$$122 \div 11 = 11, \text{ REMAINDER } 1$$

When the REMAINDER is 1, the calculation must be repeated using weight (2): 5 8 6 2  
 10 4 3 7 6 8 5 9

<b>DIGIT POSITION (Right Justified)</b>	<b>0</b>	<b>0</b>	<b>0</b>	<b>0</b>	<b>2</b>	<b>3</b>	<b>2</b>	<b>2</b>	<b>9</b>	<b>0</b>	<b>0</b>	<b>6</b>
WEIGHT (2)	5	8	6	2	10	4	3	7	6	8	5	9
Result	0	0	0	0	20	21	6	14	54	0	0	54

$$0 + 20 + 12 + 6 + 14 + 54 + 0 + 0 + 54 = 160$$

$$160 \div 11 = 14, \text{ REMAINDER } 6$$

When the REMAINDER is 0, 0 is the check digit.  
 $11 - 6 = 5$

The value 5 corresponds to character 5, therefore the checksum is 5 and the final decoded message is 0232290065.

**Code 39® Bar-code**

This code uses three to nine data digits.

The value assigned to each of the characters is:

CHARACTER	VALUE	CHARACTER	VALUE	CHARACTER	VALUE
0	0	F	15	U	30
1	1	G	16	V	31
2	2	H	17	W	32
3	3	I	18	X	33
4	4	J	19	Y	34
5	5	K	20	Z	35
6	6	L	21	-	36
7	7	M	22	.	37
8	8	N	23	SPACE	38
9	9	O	24	\$	39
A	10	P	25	/	40
B	11	Q	26	+	41
C	12	R	27	%	42
D	13	S	28		
E	14	T	29		

The checksum technique is:

- The character values of the message are obtained from the above table and added together.
- This sum is divided by 43, and the remainder corresponds to the value of the checksum character.

Example:

CHARACTER	S	T	U	V	W	X	Y	F
VALUE	28	29	30	31	32	33	34	15

$$28+29+30+31+32+33+34+15 = 232$$

$$232 \div 43 = 5, \text{ REMAINDER } 17; 17 = H = \text{CHECKCHARACTER}$$

The value 17 corresponds to character H; therefore, checksum is H, and the final decoded message is: STUVWXYFH.

**Code 128**

This code uses three to eleven data digits on the LH 700 Series and three to nine data digits for the SlideMaker bar-code label printing.

The checksum character immediately precedes the stop character. The checksum character used with Code 128 must conform to the AIM USS 128 checksum generation procedure. Do not use these values:

Code set A - 0, 64 through 102

Code set B - 0, 95 through 102

Code set C - 100 through 102

The checksum value (see the following table) is equal to the modula 103 sum of the value of the start character and the weighted values of the data/special characters. The weights are one for the first data/special character and continuing with two, three, four and so forth for the following data/special characters.

For example, a label contains a START character (Code C), Data (25), a Check character and a STOP character. The value of the Start character C is 105, and the data character for 25 is 25. The weight of the first data character is one, so the check character value is calculated as follows:

$$105 + (25 \times 1) = 130$$

where 105 and 25 are the values and 1 is the weight.

The checksum is equal to 130 modula 103 (the remainder of 130 divided by 103):

$$130 \div 103 = 1, \text{ REMAINDER } 27$$

Therefore the check character equals character value 27, which is ; in Code Set A.

For additional information on this procedure, refer to AIM USS-128 Rev. 1986, published by AIM, Inc., 1326 Freeport Road, Pittsburgh, PA 15238.

VALUE	CODE A	CODE B	CODE C
0	SP	SP	00
1	!	!	01
2	"	"	02
3	#	#	03
4	\$	\$	04
5	%	%	05
6	&	&	06
7	'	'	07
8	(	(	08
9	)	)	09
10	*	*	10
11	+	+	11
12	,	,	12
13	-	-	13

**SPECIFICATIONS/CHARACTERISTICS**  
*BAR-CODE LABEL SPECIFICATIONS*

14	.	.	14
15	/	/	15
16	0	0	16
17	1	1	17
18	2	2	18
19	3	3	19
20	4	4	20
21	5	5	21
22	6	6	22
23	7	7	23
24	8	8	24
25	9	9	25
26	:	:	26
27	;	;	27
28	<	<	28
29	=	=	29
30	>	>	30
31	?	?	31
32	@	@	32
33	A	A	33
34	B	B	34
35	C	C	35
36	D	D	36
37	E	E	37
38	F	F	38
39	G	G	39
40	H	H	40
41	I	I	41
42	J	J	42
43	K	K	43
44	L	L	44
45	M	M	45
46	N	N	46
47	O	O	47
48	P	P	48
49	Q	Q	49
50	R	R	50
51	S	S	51

52	T	T	52
53	U	U	53
54	V	V	54
55	W	W	55
56	X	X	56
57	Y	Y	57
58	Z	Z	58
59	[	[	59
60	\	\	60
61	]	]	61
62			62
63	—	—	63
64	NUL	`	64
65	SOH	a	65
66	STX	b	66
67	ETX	c	67
68	EOT	d	68
69	ENQ	e	69
70	ACK	f	70
71	BEL	g	71
72	BS	h	72
73	HT	i	73
74	LF	j	74
75	VT	k	75
76	FF	l	76
77	CR	m	77
78	SO	n	78
79	SI	o	79
80	DLE	p	80
81	DC1	q	81
82	DC2	r	82
83	DC3	s	83
84	DC4	t	84
85	NAK	u	85
86	SYN	v	86
87	ETB	w	87
88	CAN	x	88
89	EM	y	89

**SPECIFICATIONS/CHARACTERISTICS**  
*BAR-CODE LABEL SPECIFICATIONS*

90	SUB	z	90
91	ESC	{	91
92	FS		92
93	GS	}	93
94	RS	~	94
95	US	DEL	95
96	FNC 3	FNC 3	96
97	FNC 2	FNC 2	97
98	SHIFT	SHIFT	98
99	CODE C	CODE C	99
100	CODE B	FNC 4	CODE B
101	FNC 4	CODE A	CODE A
102	FNC 1	FNC 1	FNC 1
103	START (CODE A)		
104	START (CODE B)		
105	START (CODE C)		

## 5.1 LASER SAFETY

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**WARNING** Possible harm to operator. Do not use any controls, make any adjustments, or perform any procedures other than those specified herein. To do so may result in hazardous radiation exposure.

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The Triple Transducer Module and Bar-Code Reader contain lasers. A laser is a unique light source that exhibits characteristics different from conventional light sources. The safe use of the laser depends upon familiarity with the instrument and the properties of coherent, intense beams of light. The beam can cause eye damage and instrument damage. There is enough power from the laser to ignite substances placed in the beam path, even at some distance. The beam might also cause damage if contacted indirectly from reflective surfaces (specular reflection). The lasers on the LH 700 Series are covered by protective housings that are held in place by tamper-proof screws.

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**WARNING** Possible harm to operator. Do not attempt to remove the laser or to open it. Failure to comply can result in hazardous radiation exposure. If removal is required, it must be done only by a Beckman Coulter Representative.

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All service and maintenance of the laser must be done at the Beckman Coulter factory by trained personnel. If removal is required, it must be done by a Beckman Coulter Representative.

## 5.2 RADIATION HAZARDS

In the design and manufacture of the LH 700 Series, Beckman Coulter Inc. has complied with the requirements governing the use and application of a laser as stipulated in regulatory documents issued by the

- U.S. Department of Health and Human Services, and
- Center for Devices and Radiological Health (CDRH).

In compliance with these regulatory documents, every measure has been taken to ensure the health and safety of users and laboratory personnel from the possible dangers of laser use.

## 5.3 LASER WARNING LABELS

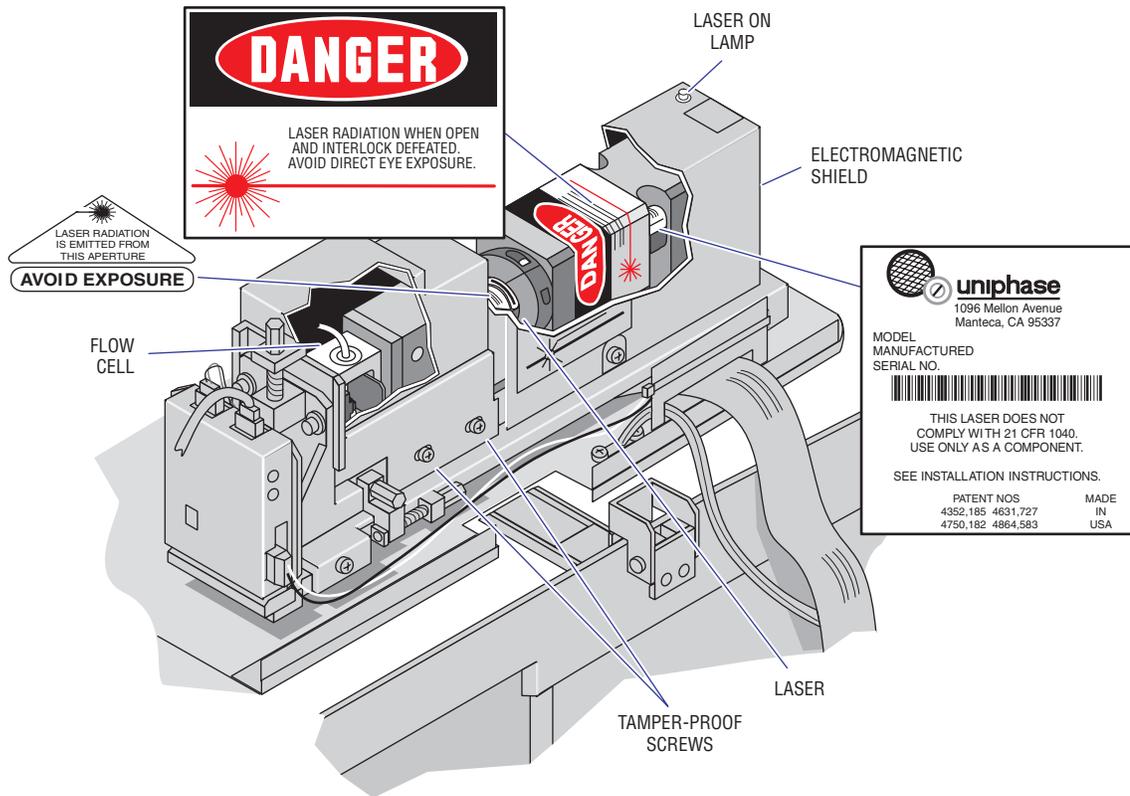
---

**WARNING** WARNING Possible harm to operator. This instrument contains components dangerous to the operator. If any attempt has been made to defeat a safety feature, or if this instrument fails to perform as listed in this manual, disconnect power and call your Beckman Coulter Representative.

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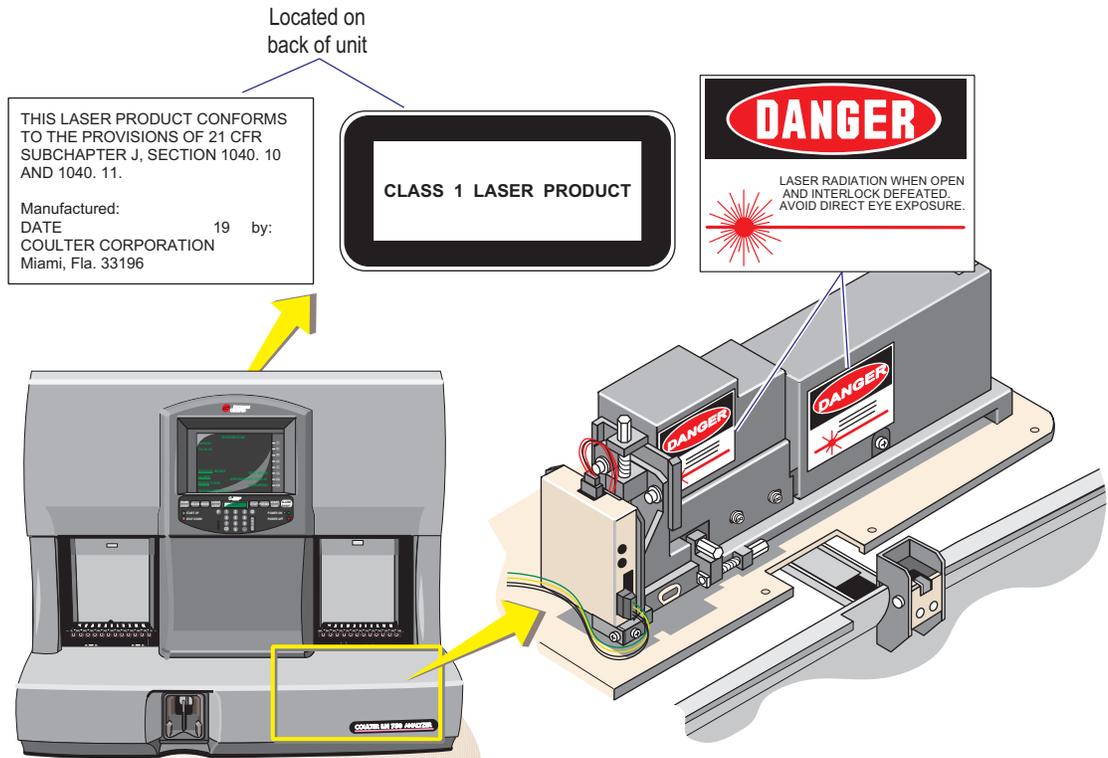
CDRH-approved labels are placed near or on those covers that, when removed, might expose laser radiation. Figure 5.1 shows the laser cover and the protective housing cut away. This illustration is intended only to show you what the system looks like, in compliance with CDRH. See Figure 5.1 for the labels and their locations on the laser head. See Figure 5.2 for the label location on the beam cover between the laser head and the sampling compartment. Figure 5.1 and Figure 5.2 show certification labels.

**Figure 5.1 Laser Warning Label, Protective Housing Cut Away**



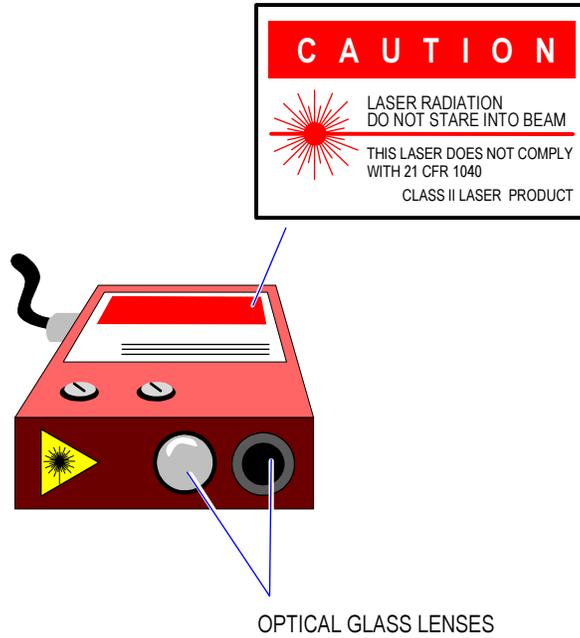
Note: As installed in the Triple Transducer Module (TTM) safety fixture, the laser presents no radiation hazard to users and complies with 21 CFR 1040.

Figure 5.2 Laser Warning Label Locations, Protective Housing On



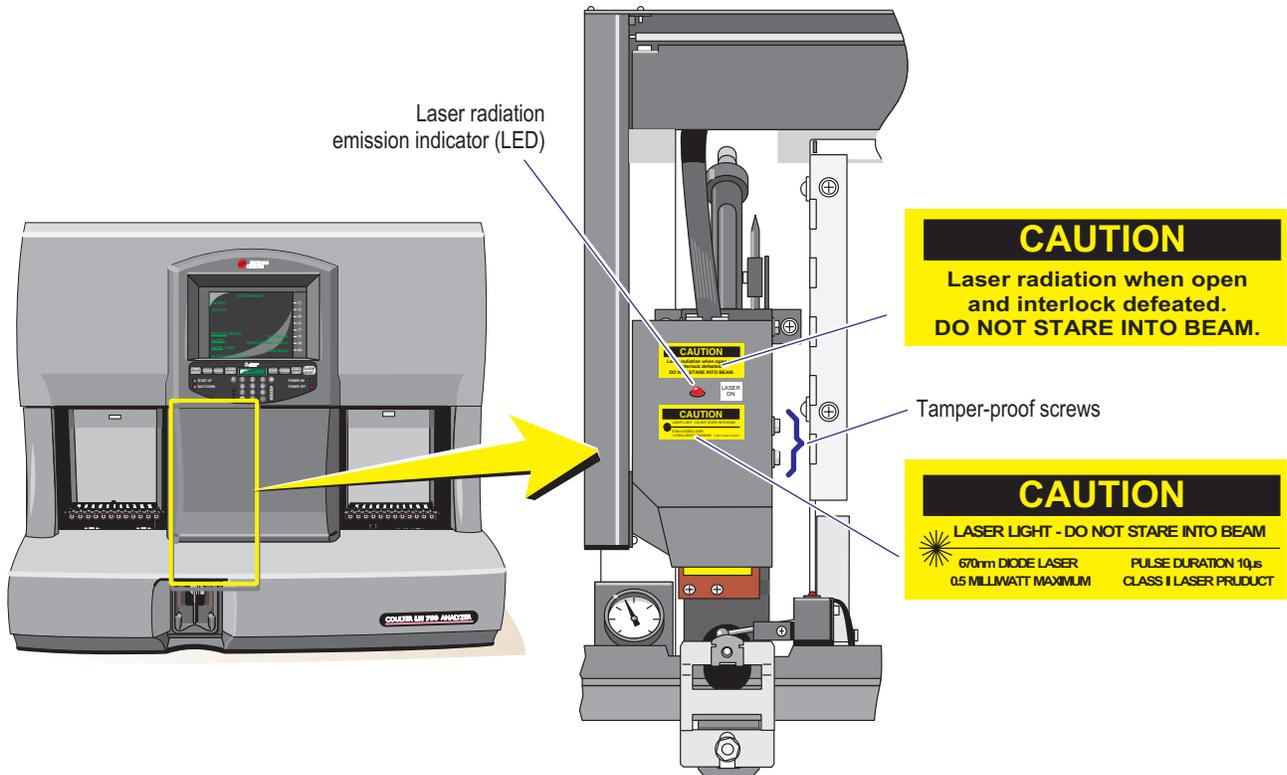
## 5.4 BAR-CODE READER

Figure 5.3 Bar-Code Reader Laser Warning Label Location, Protective Housing Cut Away



Note: As installed in the Bar-Code Reader Assembly, the laser presents no radiation hazard to users and complies with 21 CFR 1040.

Figure 5.4 Bar-Code Reader Laser Warning Label Location, Protective Housing On

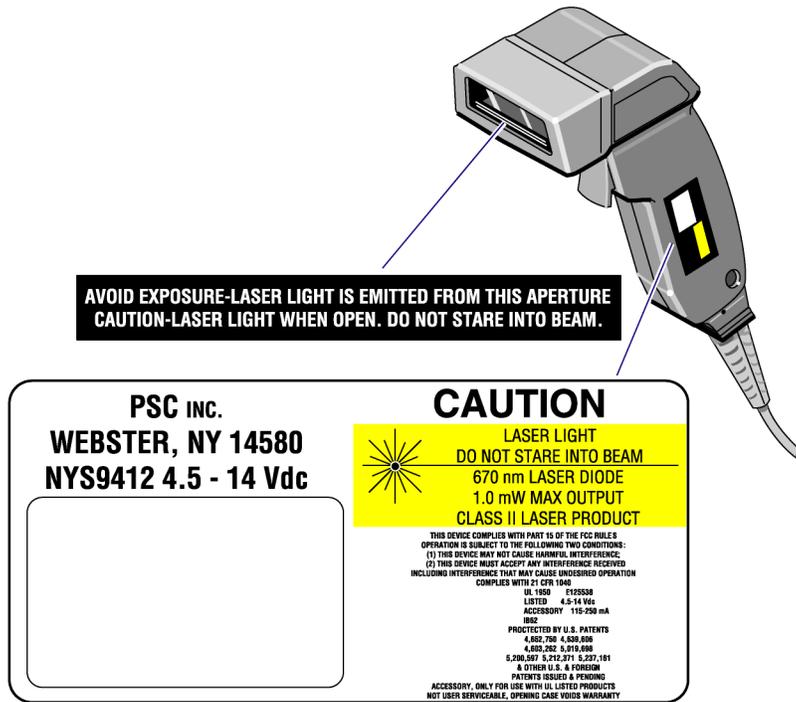


Note: As installed in the Bar-Code Reader Assembly, the laser presents no radiation hazard to users and complies with 21 CFR 1040.

## 5.5 HANDHELD SCANNER

Figure 5.5 illustrates the label on the PSC® handheld scanner.

Figure 5.5 Handheld Scanner



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<b>µL</b>	Microliter, a unit of volumetric measurement equal to one millionth of a liter.
<b>µm</b>	Micrometer, a unit of linear measurement equal to one millionth of a meter.
<b>A</b>	Ampere, a unit of electric current.
<b>Accuracy</b>	Ability of the instrument to agree with a predetermined reference value.
<b>Action Limits</b>	Values established by your laboratory to flag results requiring action.
<b>Active Set</b>	<p>For Flagging Limits, Decision Rules and Auto Validation logic, the Active Set of parameters are defined as those parameters that,</p> <ul style="list-style-type: none"> <li>• are enabled for the System (System Setup – Patient – Parameters window),</li> <li>• are enabled for the Report Profile you are using (System Setup – Patient –Reporting Options window),</li> <li>• are part of the Test Mode you are running (C, CD, CDR, CR, or R on the Command Center) or have been edited into the sample,</li> <li>• are included in the parameter block of the validation code you are looking for.</li> </ul>
<b>ADMS</b>	<p>The computer hardware and software that controls instrument operation; displays, stores, and recalls sample data; automates QC and calibration procedures; and assists you in troubleshooting.</p> <p>Also called: ADMS, DMS, LH 700 Series Workstation.</p>
<b>Algorithm</b>	A particular procedure for performing an analysis.
<b>Analytical Station</b>	The Analyzer, Diluter, and Power Supply of the LH 700 Series.
<b>Analyzer Cover</b>	Several circuit cards appear behind the Analyzer cover. You should only open the Analyzer cover at the instruction of your Beckman Beckman Coulter Representative.
<b>Analyzer Pulse</b>	A graphical display on the Analyzer screen that indicates the condition of all the apertures. You can detect noise, bubbles, or a clogged flow cell by monitoring the pulses.
<b>ANSI</b>	American National Standards Institute.
<b>Application</b>	A computer program, generally started from a button or icon. For example, starts the Patient Tests application.
<b>ASCII</b>	American Standard Code for Information Interchange.
<b>Aspiration Mode</b>	The method of running a sample, closed vial (Automatic aspiration mode) or open vial (Manual aspiration mode).
<b>Assay Values</b>	<p>Values of all parameters for a control established by extensive assay of that control. These values are provided by the manufacturer of the control.</p> <p>Also called: assigned values.</p>
<b>Assigned Value</b>	The Workstation adds and subtracts the expected range to each value to develop an acceptable limit for the control. When you cycle a control, the Workstation checks the control results against the acceptable limits. If the control results are outside these limits, flags appear.

## GLOSSARY

<b>ASTM</b>	American Society for Testing and Materials.
<b>AutoCollation</b>	A Workstation feature that automatically combines results of different test modes (CBC/Diff and Retic) analyzed from a sample with an identical patient or sample ID. The different test modes must be performed within a predetermined time.
<b>Automatic Aspiration Mode</b>	The closed-vial method of running a sample. You place a closed tube sample in a cassette and place the cassette on the LH 700 Series loading bay. The LH 700 Series automatically reads the cassette label and tube label (if present) and aspirates the sample.  Also called: closed-vial mode, primary mode, automatic sampling mode.
<b>Automatic Startup Cycles</b>	When you press the Startup button, the system flushes the cleaning agent from all Diluter components and tubing if cleaning reagent is not already removed. It also performs electronic and fluidic checks to ensure the instrument is ready to analyze control or whole-blood samples.
<b>AutoNumbering</b>	A Workstation feature that assigns a number to each sample received from the instrument.
<b>Autopurge</b>	If 24 hours elapse with the power on, pneumatics off, and the instrument in shutdown, the system automatically: turns on the pneumatics; drains and fills the baths; purges the flow cell and associated sample lines with cleaning agent.
<b>AutoSequencing</b>	A Workstation feature used with the ToDo list to automatically increment the identifiers you specify (Cass/Pos, sample ID, and sequence number).  When you enter the sample ID, ensure you type the same number of digits as exist on your bar-code labels, including leading zeros.
<b>AutoStop</b>	A Workstation feature that instructs the instrument to stop processing based on specific criteria.
<b>Average Value</b>	A value that is computed by dividing the sum of a set of values by the number of values. Usually referred to as the simple arithmetic mean.
<b>BA # (Basophils)</b>	A WBC differential parameter result from the Diff sample analysis. The number (#) is computed from the WBC count and the BA %.
<b>BA % (Basophils)</b>	A WBC differential parameter result from the Diff sample analysis. The percent (%) is measured directly using VCS technology.
<b>Background Count</b>	A measure of the amount of electrical or particle interference using a diluent sample or no sample.
<b>Background Results</b>	Background results indicate reagent quality and the presence of electronic noise or bubbles. Background results above certain levels cause concern since they might falsely elevate (or interfere with) the results obtained for blood and control samples.

<b>Batch</b>	<p>A group or set of results.</p> <p>For Analysis, a batch equals 20 patient samples used for monitoring MCV, MCH and MCHC as an automated QC procedure.</p> <p>For printing or transmitting, a batch equals the samples you selected from the database.</p>
<b>Batch Mean</b>	The mean or average of a set of samples. For Analysis, the batch mean is a value based on a statistical averaging technique and is a type of "weighted moving average." It is used to estimate what a simple average result of a very large number of samples (population mean) might be by using a small number of samples.
<b>Batch Print Job</b>	Selecting multiple items from the database, then selecting the print icon to print all of them at the same time.
<b>Baud</b>	A rate defining how many data bits per second are transferred during communications between two pieces of equipment.
<b>Calibration</b>	A procedure to standardize the instrument for accuracy by determining its deviation from calibration references and applying any necessary correction factors.
<b>Calibration Disk</b>	A data diskette containing the Assigned Values for the lot number of the S-CAL® calibrator kit being used.
<b>Calibration Factors</b>	Values the system uses to fine-tune instrument accuracy.
<b>Carryover</b>	The percentage or particle count (for Diff and Retic) of blood cells that are retained from one blood sample to the next.
<b>Cass/Pos</b>	The Workstation considers this field an optional identifier for the cassette number and cassette position number. This identifier appears on bar-code labels so that cassettes and positions are automatically read using a bar-code reader.
<b>Cassette</b>	The cassette is the carrier for the sample tubes (patient, control, or special test) used in Automatic aspiration mode where automatic loading, mixing, and sampling occurs.
<b>Cassette Clips</b>	Clips that you permanently install in each opening of the special cassettes for tubes with HEMOGARD™ Closure. These clips allow the special cassette to accommodate additional types of tubes.
<b>CBC (Complete Blood Count)</b>	A measure of the cell number and the indices. CBC includes WBC, RBC, Hgb, Hct, MCH, MCHC, Plt, MPV, and RDW.
<b>CDC</b>	Centers for Disease Control and Prevention.
<b>CEE</b>	Commission for Electrical Equipment.
<b>Cell Control</b>	A preparation made of human blood with stabilized cells and surrogate material. It is used for daily instrument QC.
<b>Characters</b>	All letters A-Z and numbers 0-9.
<b>Check Valve</b>	<p>A one-way valve that routes liquid or air through the Diluter.</p> <p>Also called: mono-flow valve, one-way valve</p>

## GLOSSARY

<b>CLIA</b>	Clinical Laboratory Improvement Amendments.
<b>Closed-vial Mode</b>	The closed-vial method of running a sample. You place a closed tube sample in a cassette and place the cassette on the LH 700 Series loading bay. The LH 700 Series automatically reads the cassette label and tube label (if present) and aspirates the sample.  Also called: closed-vial mode, primary mode, automatic sampling mode.
<b>cm</b>	Centimeter, a unit of linear measurement.
<b>Codes</b>	On windows and printouts, symbols, such as +, -----, +++, ::::, appear with, or in place of sample or control results because of irregularities. A code can only be generated by the instrument.
<b>Coefficient of Variation</b>	An expression, in percent, of data spread as related to the mean.  $\%CV = (SD/Mean) \times 100$
<b>Coincidence</b>	More than one cell within aperture-sensing boundaries at the same time. The system counts only one pulse and automatically corrects results for coincidence.
<b>Collation</b>	The process of combining the results of different test modes(CBC/Diff and Retic) analyzed from the same sample.
<b>Command Center</b>	A blue bar that appears at the bottom of the screen.
<b>Control Disk</b>	A data diskette that contains the assay values for the lot number of the control being used.
<b>Coulter Principle</b>	A method of counting and sizing cells by detecting and measuring changes in electrical resistance when a particle in a conductive liquid goes through a small aperture.
<b>Critical Limits</b>	Values established by your laboratory to flag results requiring immediate action.
<b>CSA</b>	Canadian Standards Association.
<b>Cursor</b>	On the screen, a place shown by a little blinking indicator or by a highlighted area. The cursor shows where you can select an option or type information.
<b>CV (Coefficient of Variation)</b>	An expression, in percent, of data spread as related to the mean.  $\%CV = (SD/Mean) \times 100$
<b>Cycle Counter</b>	A number that appears on the Analyzer MAIN MENU screen that represents the actual number of instrument cycles that have occurred.
<b>DataPlot</b>	A graphic representation of results. DataPlots present a combined view of population density and membership. Colors represent different types of cells. Shades of colors represent the number of cells--bright colors are the most dense.
<b>Data Management System (DMS)</b>	The computer hardware and software that controls instrument operation; displays, stores, and recalls sample data; automates QC and calibration procedures; and assists you in troubleshooting.  Also called: ADMS, DMS, LH 700 Series Workstation.

<b>Default</b>	A setting the instrument uses automatically. For example, you can set up a default printer. Every time you print sample results, the Workstation automatically prints the sample results to the default printer.
<b>Definitive Messages</b>	Definitive messages appear in a separate area of the screen display, printout, and host transmission. Definitive messages such as Anisocytosis, Leukopenia, etc, are laboratory-defined. Definitive messages are defined based on numeric limits generated by your laboratory. If results exceed the limits, the Workstation generates a message. Results that generate these messages may require review. Check your laboratory's protocol for handling the particular message.
<b>Delta Check</b>	A check on sample results that is made by clinical laboratories to determine if the current result on a particular patient is within certain limits of the last result obtained on that same patient.
<b>Density</b>	The number of cells in a particular region, regardless of the type of cell.  On DataPlots, as more cells appear in a region, the color of the region gets brighter.
<b>Diff (Differential)</b>	Leukocyte differential parameters (NE, LY, EO, BA, and MO) and processes that relate to them.  The differential count for all parameters, when totaled, equals 100% except when ALL differential parameters display as zeros. If the differential count displays all zeros, check for additional flagging and take appropriate action according to your laboratory's protocol.
<b>Digits</b>	All numbers 0-9.
<b>Diluter</b>	This is the fluidics portion of the LH 700 Series System. The Diluter is the subsystem that aspirates the sample, dilutes it and mixes it. Move the cursor over the illustration to see links to additional information.
<b>dL</b>	Deciliter, a unit of volumetric measurement equal to 0.1 liter.
<b>DMS</b>	The computer hardware and software that controls instrument operation; displays, stores, and recalls sample data; automates QC and calibration procedures; and assists you in troubleshooting.  Also called: ADMS, DMS, LH 700 Series Workstation.
<b>Double-click</b>	Using Means Light Pen Quickly pressing the light pen against the screen twice. Mouse Quickly pressing the left mouse button twice. Touch Screen Quickly touching the screen with your finger twice.
<b>Drag</b>	The process of selecting an item and maintaining the selection while you move the pointer to a different location.
<b>EDTA</b>	Ethylenediaminetetraacetic acid--the recommended anticoagulant for hematology analysis.
<b>EO # (Eosinophils)</b>	A WBC differential parameter result from the Diff sample analysis. The number (#) is computed from the WBC count and the EO %.
<b>EO % (Eosinophils)</b>	A WBC differential parameter result from the Diff sample analysis. The percent (%) is measured directly using VCS technology.

## GLOSSARY

<b>Expiration Date</b>	A manufacturer's recommended last day of use for a reagent, control, or calibrator.
<b>Field</b>	Area on a window or screen for entering or viewing data. When you move the cursor, you are moving it from field to field.
<b>First Name</b>	The Workstation considers this field an optional identifier you can use for a sample.  This field appears blank unless you provide the information when you add a sample request to the ToDo list or edit a sample result on the Edit Sample window.
<b>fL</b>	Femtoliter, a unit of volumetric measurement equal to $10^{-15}$ liter.
<b>Flags</b>	A flag is a single letter or symbol and will always appear to the right of a result. A flag can be instrument-generated (R, P), or laboratory-defined (H, L, c, a). On windows and printouts, the letters, such as H, L, and R appear next to parameter results to indicate specific conditions.
<b>Flow Cytometry</b>	A process for measuring the characteristics of cells or other biological particles as they pass through a measuring apparatus in a fluid stream.
<b>Function Key</b>	One of the keys labeled F1 to F12. To request a system or window-specific command, press the Function Key displayed on the screen next to the command. Sometimes these keys are used with other keys to access specific functions.
<b>g</b>	Gram, a unit of weight.
<b>HCT (Hematocrit)</b>	A computed value that represents the packed cell volume (PCV) relative to a given volume of whole blood.  $\text{HCT (\%)} = (\text{RBC} \times \text{MCV}) / 10$
<b>Hemoglobinometry</b>	Measurement of hemoglobin in the blood. In the LH 700 Series, this is done by comparing the amount of light that passes through a diluted lysed sample, in which the released Hgb has been chemically converted, with the amount of light that passes through a blank (LH 700 series diluent).
<b>HGB (Hemoglobin)</b>	Hemoglobin results from the CBC analysis. This parameter is measured directly using Photometric Measurement.  $\text{HGB (g / dL)} = \text{Constant} \times \log_{10} \frac{\text{Reference \%T}}{\text{Reference \%T}}$
<b>HGB Voltages</b>	If the HGB voltage readings do not meet the following criteria, you must repeat the background test.

<b>Histograms (RBC &amp; Plt)</b>	<p>Misleading results can occur if you estimate the number of cells from the distribution curves because curves show only the relative, not the actual, number of cells in each size range. Do not estimate the number of cells from the distribution curves.</p> <p>Graphic representations of cell frequency vs. size. Histograms provide information about erythrocyte and thrombocyte frequency and their distribution about the mean. They also might show the presence of subpopulations.</p>
<b>HLR (High Light Scatter Reticulocytes)</b>	<p>This field and parameter label disappear if the parameter was excluded as part of parameter selection in setup. These fields cannot be edited.</p> <p>These parameters are derived from VCS technology.</p> <p><math>HLR \% = (HLR / \text{Total Erythrocytes}) \times 100</math></p> <p><math>HLR \# = (HLR / \text{Total Erythrocytes}) \times RBC</math></p>
<b>Hz</b>	Hertz, a unit of frequency.
<b>i.d.</b>	Inside diameter.
<b>IEC</b>	International Electrical Commission.
<b>Information System</b>	Any host or laboratory computer system.
<b>Instrument</b>	The Analyzer and Diluter portion of the LH 700 Series.
<b>IQAP (Interlaboratory Quality Assurance Program)</b>	Beckman Coulter Inc. provides this program, which statistically compares your 5C®-ES and Retic-C cell control data to a group of other laboratories' control recovery data with the same control lot number.
<b>IRF (Immature Reticulocyte Fraction)</b>	<p>This field and parameter label disappear if the parameter was excluded as part of parameter selection in setup. This field cannot be edited.</p> <p>This parameter is derived from VCS technology.</p> <p><math>IRF = HLR / \text{Total reticulocytes}</math></p>
<b>L</b>	Liter, a unit of volumetric measurement.
<b>Lab Administrator</b>	A person at your location who can access the full set of Workstation functions. The lab administrator creates user names and assigns access levels. A specific user name and password is assigned to the lab administrator in your laboratory.
<b>Laser (Light Amplification by Stimulated Emission of Radiation)</b>	The instrument uses a laser for WBC Diff and Retic analysis.
<b>Last Name</b>	<p>The Workstation considers this field an optional identifier you can use for a sample.</p> <p>This field appears blank unless you provide the information when you add a sample request to the ToDo list or edit a sample result on the Edit Sample window.</p>

## GLOSSARY

<b>Levey-Jennings Control Chart</b>	A commonly used control procedure in which control measurements are plotted directly on a control chart with limit lines drawn as mean plus or minus expected ranges.
<b>Linearity</b>	<p>The ability of an instrument to recover expected results (reference values or calculated values) for such parameters as WBC, RBC, Hgb and Plt at varying levels of concentration of these parameters within specified limits.</p> <p>Beckman Coulter Inc. provides the LIN-C' linearity control for your convenience. Use it according to the instructions on its package insert.</p>
<b>Lot Number</b>	An identifier assigned by a manufacturer to identify a control, reagent or calibrator.
<b>LY #(Lymphocytes)</b>	A WBC differential parameter result from the Diff sample analysis. The number (#) is computed from the WBC count and the LY %.
<b>LY % (Lymphocytes)</b>	<p>A WBC differential parameter result from the Diff sample analysis. The percent (%) is measured directly using VCS technology.</p> $\text{LY \%} = \frac{\text{no. of cells inside LY area}}{\text{no. of cells inside NE+LY+MO+EO+BA}} \times 100$
<b>m</b>	Meter, a unit of linear measurement.
<b>Manual aspiration mode</b>	<p>The open-vial method of running a sample. You immerse the aspirator tip in the sample. The instrument automatically aspirates the sample. If the instrument fails to aspirate automatically, you can press and release the activator to aspirate the sample.</p> <p>Also called: open-vial mode, secondary mode, manual sampling mode.</p>
<b>Manual Diff Box</b>	A blank area included on a report where you can record your manual differential results for the sample.
<b>Manual Print Job</b>	Selecting the print icon to print a single item.
<b>MCH (Mean Corpuscular Hemoglobin)</b>	<p>This field and parameter label disappear if the parameter was excluded as part of parameter selection in setup.</p> <p>A computed value that represents the amount of hemoglobin by weight in the average red cell. The system automatically recalculates this value if HGB or RBC change.</p> $\text{MCH (pg)} = (\text{HGB/RBC}) \times 10$
<b>MCHC (Mean Corpuscular Hemoglobin Concentration)</b>	<p>This field and parameter label disappear if the parameter was excluded as part of parameter selection in setup.</p> <p>A computed value that represents the concentration of hemoglobin (weight/volume) in the average red cell. The system automatically recalculates this value if HGB or HCT change.</p> $\text{MCHC (g/dL)} = (\text{HGB/HCT}) \times 100$

<b>MCV (Mean Corpuscular Volume)</b>	<p>The volume of the average red cell derived from the RBC histogram. Manually calculated by dividing the hematocrit (HCT) by the red cell count and multiplying by 10.</p> $MCV = (HCT/RBC) \times 10$
<b>Mean</b>	Arithmetic average of a group of data.
<b>Membership</b>	<p>The different types of cells in a particular region, regardless of the number of cells.</p> <p>On DataPlots, membership is represented showing different types of cells in different colors.</p>
<b>mL</b>	Milliliter, a unit of volumetric measurement, equal to $10^{-3}$ liter.
<b>mm</b>	Millimeter, a unit of linear measurement, equal to one-thousandth of a meter.
<b>MO # (Monocytes)</b>	<p>A WBC differential parameter result from the Diff sample analysis. The number (#) is computed from the WBC count and the MO %.</p> $MO (10^3 \text{ cells } \mu\text{L}) = \frac{MO\%}{100} \times \text{WBC count}$
<b>MO % (Monocytes)</b>	<p>A WBC differential parameter result from the Diff sample analysis. The percent (%) is measured directly using VCS technology.</p> $MO \% = \frac{\text{no. of cells inside MO area}}{\text{no. of cells inside NE+LY+MO+EO+BA}} \times 100$
<b>Mode</b>	The method of running a sample, closed vial (Automatic aspiration mode) or open vial (Manual aspiration mode).
<b>Modes (Test)</b>	Use Default Type on the Command Center to specify the test mode. It determines how the LH 700 Series processes data.
<b>Mode-to-Mode Matching</b>	Agreement between patient results in Automatic aspiration mode and Manual aspiration mode.
<b>MPV (Mean Platelet Volume)</b>	<p>This field and parameter label disappear if the parameter was excluded as part of parameter selection in setup.</p> <p>MPV is the mean cell volume for platelets.</p> <p>The system derives this parameter from the Plt histogram.</p>
<b>MRV (Mean Reticulocyte Volume)</b>	<p>This field and parameter label disappear if the parameter was excluded as part of parameter selection in setup. This field cannot be edited.</p> <p>Mean volume of the Retic population. This parameter is derived from VCS technology.</p>

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MSCV (Mean Sphered Cell Volume)	<p>THIS PARAMETER IS FOR RESEARCH USE ONLY. NOT FOR USE IN DIAGNOSTIC PROCEDURES.</p> <p>This field and parameter label disappear if the parameter was excluded as part of parameter selection in setup. This field cannot be edited.</p> <p>Mean volume of red cell population of Reticulocyte analysis. This parameter is derived from VCS technology. The clearing solution causes a spherizing effect within the mature red cells.</p>
mW	Milliwatt, a unit of power equal to one-thousandth of a watt.
n	Number.
NCCLS	National Committee for Clinical Laboratory Standards.
NE # (Neutrophil)	<p>A WBC differential parameter result from the Diff sample analysis. The number is computed from the WBC count and the NE%.</p> $NE (10^3 \text{ cells } \mu\text{L}) = \frac{NE\%}{100} \times \text{WBC count}$
NE % (Neutrophil)	<p>A WBC differential parameter result from the Diff sample analysis. The percent is measured directly using VCS technology.</p> $NE \% = \frac{\text{no. of cells inside NE area}}{\text{no. of cells inside NE+LY+MO+EO+BA}} \times 100$
NEMA	National Electrical Manufacturers Association.
nm	Nanometer, a unit of linear measurement, equal to $10^{-9}$ meter.
NRBC # (Nucleated Red Blood Cells)	<p>A parameter that is calculated from the NRBC % and the total WBC count. NRBC # represents the total number of nucleated Red Blood Cells.</p> $NRBC (10^3 \text{ cells } \mu\text{L}) = NRBC\% \times \text{WBC count}$
NRBC % (Nucleated Red Blood Cells)	A parameter derived from both the WBC histogram and VCS information and represents the number of nucleated Red Blood Cells per 100 White Blood Cells.
o.d.	Outer diameter.
Operator ID	<p>A 16-character identifier that uniquely identifies the processor of the samples. If you forget your user name or password, contact your lab administrator. The user name works with your password to uniquely identify you to the system.</p> <p>Also called: operator ID.</p>
Open-vial Mode	<p>The open-vial method of running a sample. You immerse the aspirator tip in the sample. The instrument automatically aspirates the sample. If the instrument fails to aspirate automatically, you can press and release the activator to aspirate the sample.</p> <p>Also called: open-vial mode, secondary mode, manual sampling mode.</p>
Outlier	A data value far outside the range of the rest of the data.

<b>Output Device</b>	A physical device that is capable of receiving information from a computer and formatting it in a logical fashion. Examples: printer, fax machine or another computer.
<b>Parameters</b>	Characteristics of blood that the instrument measures and reports.
<b>Parity</b>	Method of detecting errors in data handling. The computer generates a parity bit such that the sum of the data bits for a data word are odd or even and stored in the parity bit for checking by the receiver of the data word.
<b>Patient ID</b>	The Workstation considers this field an optional sample identifier.  Your laboratory may use it as a specific identifier for the patient, such as the medical record or Social Security Number. It is intended for laboratories that want to track results of several different samples or tests for the same patient.
<b>Patient Population</b>	A large number of patient sample results for Analysis, used to give a fairly consistent average result for each of the three red blood cell indices: MCV, MCH and MCHC.
<b>PCT (Plateletcrit)</b>	THIS PARAMETER IS FOR RESEARCH USE ONLY. NOT FOR USE IN DIAGNOSTIC PROCEDURES.  This field and parameter label disappear if the parameter was excluded as part of parameter selection in setup.  A computed value that represents the platelet packed cell volume.
<b>PDW (Platelet Distribution Width)</b>	THIS PARAMETER IS FOR RESEARCH USE ONLY. NOT FOR USE IN DIAGNOSTIC PROCEDURES.  This field and parameter label disappear if the parameter was excluded as part of parameter selection in setup.  The coefficient of variation of Plt size.  The system derives this parameter from the Plt histogram. This parameter appears as a percent.
<b>pg</b>	Picogram, a unit of gravimetric measure equal to one trillionth of a gram.
<b>Photometric Measurement</b>	A process where a beam of white light from an incandescent lamp goes through an optical filter and is read by a photocell. This generates a current that can be measured.
<b>Plt Histogram</b>	Misleading results can occur if you estimate the number of cells from the distribution curves because curves show only the relative, not the actual, number of cells in each size range. Do not estimate the number of cells from the distribution curves.  A Plt distribution curve. It displays only the area between 0 fL and 36 fL.
<b>PLT (Platelet)</b>	Platelet count results from the CBC analysis. The system derives this parameter from the Plt histogram.  $PLT = n \times 10^3 \text{ cells}/\mu\text{L}$

<b>Positive Identifier</b>	An identifier that is linked irrevocably to the date and time of instrument analysis on a sample and the sample results. A positive identifier must be entered before a sample analysis can occur. You can use either sample ID or Cass/Pos as positive identifiers of the sample and its results. You can also choose to use both sample ID and Cass/Pos as positive identifiers.
<b>Precision</b>	A measure of the ability of the instrument to reproduce similar results when a sample is run repeatedly. Precision of the instrument is a CV (or an SD for differential results), based on at least 31 replicate determinations of the same sample. Precision shows the closeness of test results when repeated analyses of the same material are performed.  Also called: reproducibility.
<b>Precision Test</b>	The precision test is performed as part of the automatic startup cycles. If any test value exceeds the reference value by 1% or more, the test value appears in red on the Workstation screen and is flagged with an H (high) or L (low).
<b>Predilute</b>	
<b>Primary Mode</b>	
<b>Print Profile</b>	A set of characteristics that define what you want printed and transmitted for sample runs. You can set up print profiles as part of System Setup.  Also called: Reports, Reporting Options
<b>psi</b>	Pounds per square inch, a unit of pressure measurement.
<b>QC (Quality Control)</b>	A comprehensive set of procedures your laboratory uses to ensure that the instrument is working accurately and precisely.
<b>Ramp Test</b>	The ramp test is performed as part of the automatic startup cycles. If any test value exceeds the reference value by 1% or more, the test value appears in red on the Workstation screen and is flagged with an H (high) or L (low).
<b>RBC (Red Blood Cell)</b>	Red Blood Cell count results from the CBC analysis. This parameter is measured directly using the Coulter Principle.  $\text{RBC} = n \times 10^6 \text{ cells}/\mu\text{L}$
<b>RBC Histogram</b>	Misleading results can occur if you estimate the number of cells from the distribution curves because curves show only the relative, not the actual, number of cells in each size range. Do not estimate the number of cells from the distribution curves.  An RBC distribution curve. The normal curve ranges from 36 to 360 fL. The display starts at 24 fL.

<b>RDW (Red Distribution Width)</b>	<p>This field and parameter label disappear if the parameter was excluded as part of parameter selection in setup.</p> <p>The Coefficient of Variation of RBC size.</p> <p>The system derives this parameter from the RBC histogram.</p> <p>If you edit RBC, HGB, HCT, or MCV values, this field appears with the incomplete computation code (.....).</p>
<b>Record</b>	A collection of related information, for example, sample identification and results information, stored in the database and treated as a unit.
<b>Reflex Manager List</b>	A list automatically generated by the Workstation based on sample result criteria set up by your laboratory. You can use this list as a management tool to allow automation of post-analysis decisions. As part of setting up the Reflex Manager List, you define the decision criteria and the follow-up action required.
<b>Reportable range</b>	This range represents the clinical limits of values that have been tested and found to be accurate, precise, and linear. Reportable range can be the same as the linear range, but it is usually a subset of the linear range.
<b>Reproducibility</b>	A procedure to check that the system gives consistent results (within established limits) every time it measures the same sample. Also called: precision.
<b>RET # (Reticulocytes)</b>	<p>This field and parameter label disappear if the parameter was excluded as part of parameter selection in setup.</p> <p>A computed value that represents the absolute number of reticulocytes.</p> $\text{RET\#} = \frac{\text{RET\%} \times \text{RBC Count}}{100}$ <p>The sample's RBC must be known to calculate this value.</p>
<b>RET % (Reticulocytes)</b>	Reticulocyte count results from Retic analysis of samples or controls. The percent is measured directly using VCS technology.
<b>Sample ID (SID)</b>	<p>One of two possible positive identifier for sample analysis. The other possible positive identifier is Cass/Pos.</p> <p>For Automatic aspiration mode, this identifier is read from the bar-code label of the sample. For samples using Manual aspiration mode, this is the identifier specified at the Numeric Keypad or read from the bar-code label on the sample by using the wedge scanner.</p>
<b>Screen Saver</b>	<p>The Workstation has a screen saver that appears when you are not interacting with the Workstation. You can set up the screen saver to appear at specific intervals of non-use.</p> <p>The screen returns to its normal view when the Workstation receives sample data or when you interact with the Workstation by pressing a key on the keyboard.</p>

## GLOSSARY

<b>SD (Standard Deviation)</b>	<p>A measure of deviation from the mean.</p> $SD = \sqrt{\frac{\sum(x - \bar{x})^2}{n - 1}}$
<b>Select</b>	<p>When an item is selected, a visual cue is present. For example, appears in a field.</p>
<b>Sequence Number</b>	<p>The Workstation considers this field an optional identifier you can use for a sample.</p> <p>The Workstation assigns a sequence number to a sample when a sample request is added to the ToDo list. If AutoSequencing is ON for the Patient Level Seq # field the Workstation automatically assigns the next available sequence number to the sample request.</p>
<b>Shift</b>	<p>Consecutive values that abruptly move from one side of the mean to the other, then maintain a constant level. Contrast with trend.</p> <p>A scheduled period of work.</p>
<b>SID (Sample ID)</b>	<p>One of two possible positive identifier for sample analysis. The other possible positive identifier is Cass/Pos.</p>
<b>Standard Deviation (SD)</b>	<p>A measure of deviation from the mean.</p> $SD = \sqrt{\frac{\sum(x - \bar{x})^2}{n - 1}}$
<b>Status Bar</b>	<p>A horizontal bar that appears toward the bottom of the screen. It displays short instructions that can help you work with the LH 700 Series Workstation. It also displays information about various LH 700 Series options.</p>
<b>Stop Bit</b>	<p>A computer code that indicates the end of a character.</p>
<b>Suspect Messages</b>	<p>Suspect messages appear in a separate area of the screen display, printout, and host transmission. Suspect messages, such as Imm NE 1, Platelet Clumps, etc, are instrument-generated. Suspect messages appear for sample results based on an abnormal cell distribution or population. The system generates these messages according to an internal algorithm. Abnormalities should be confirmed by microscopic review.</p>
<b>Sweep Flow</b>	<p>A steady stream of diluent that flows behind the RBC aperture during sensing periods to keep RBCs from swirling back into the sensing zone.</p>
<b>Test (Modes)</b>	<p>Use Default Type on the Command Center to specify the test mode. It determines how the LH 700 Series processes data.</p> <p>The test mode appears as a field on the QA Results &amp; Graphics window.</p>

<b>ToDo List</b>	<p>A feature of the Workstation that lets you view a list of samples the instrument has not yet processed. You can view a list of all the unprocessed samples or a list of unprocessed samples for a particular test mode. You can sort these lists by any column you choose.</p> <p>Also called: Worklist, Pending List, Work Order.</p>
<b>Toolbar</b>	<p>A group of graphic buttons on a Workstation window. You can select a button on a Toolbar to quickly access commonly used functions.</p>
<b>Trend</b>	<p>Values that continue to increase or decrease gradually over a period of time. Contrast with shift.</p>
<b>Tube Adapters</b>	<p>Special holders that enable small tubes to fit in standard cassettes. Two sizes of gray sleeve adapters accommodate 2.0-mL tubes and 3.0-mL tubes.</p>
<b>UL</b>	<p>Underwriters Laboratory.</p>
<b>User Name</b>	<p>A 16-character identifier that uniquely identifies the processor of the samples. If you forget your user name or password, contact your lab administrator. The user name works with your password to uniquely identify you to the system.</p> <p>Also called: operator ID.</p>
<b>Vac</b>	<p>Volts of alternating current.</p>
<b>VCS (Volume, Conductivity and Scatter)</b>	<p>A flow cytometry technology applied in hematology to enhance WBC subpopulation classification and percent Reticulocyte measurement.</p>
<b>Voting</b>	<p>After the computer corrects for coincidence, it compares the three count periods. Voting may occur for WBC, RBC, Plt, MCV, RDW and MPV. Agreement among the three count periods causes them to be averaged to determine the parameter result.</p> <p>If the instrument finds disagreement among all three count periods, the Workstation displays the total voteout code, ----, instead of the parameter result.</p>
<b>W</b>	<p>Watt, a unit of power.</p>
<b>WBC (White Blood Cell)</b>	<p>White Blood Cell count results from the CBC analysis. The WBC count is adjusted for interfering substances when appropriate. No further correction of WBC is required. Interfering substances include, but are not limited to NRBC, giant platelets, platelet clumps, unlysed RBCs and RBC fragments.</p> <p>This parameter is measured directly using the Coulter Principle.</p> $\text{WBC} = n \times 10^3 \text{ cells}/\mu\text{L}$

<p><b>Wettability</b></p>	<p>The ability of any solid surface to be wetted when in contact with a liquid. Wettability affects the smear quality:</p> <p>Dirty slides concentrate the blood droplet in one location resulting in a poor quality smear.</p> <p>Clean slides allow for the blood droplet to disperse evenly across the slide providing for a good quality smear.</p>
<p><b>Window</b></p>	<p>A rectangular area on your computer screen that enables you to work with an application.</p>
<p><b>Windows® 2000 Workstation Operating System</b></p>	<p>The Microsoft® operating system for your computer.</p>
<p><b>XB Analysis</b></p>	<p>A method of quality control that automatically compares patient red blood cell indices (MCV, MCH and MCHC) with known target values. It is used to monitor for proper operation of automated instruments in hematology.</p>

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