



PROFICIENCY TESTING REPORT
ISHTM-AIIMS EXTERNAL QUALITY ASSURANCE PROGRAMME
 NABL accredited program as per ISO/IEC 17043:2010 standard
 Organized By Department of Hematology, AIIMS, New Delhi-110029



Duration of stability testing - minimum upto 8 days at ambient temp. after dispatch of specimens

EQAP CODE No. : 4355

Distribution No.: 156-L

Month/Year: July/2022

Instrument ID: YUMIZEN H550

Name & Contact No. of PT Co-ordinator: Dr. Seema Tyagi (Prof.), Hematology, AIIMS, Delhi,
 Tel: 9013085730 , E-Mail : accuracy2000@gmail.com

Date of issue & status of the report: 27-09-2022[Final].

CBC and Retic Assessment

Test Parameters	S.No.	Among Lab (Accuracy Testing)						Within Lab (Precision Testing)			
		Your Result 1	Your Result 2	Your Results Sum of 2 Value	Consensus result sum of 2 values (Assigned Value)	Uncertainty of Assigned Values	Z Score	Yours Results Diff. of 2 Values	Consensus Result Diff. of 2 values (Assigned Value)	Uncertainty of Assigned Values	Z Score
WBC x10 ³ /µl	1	6.1	4.75	10.85	12.18	0.0990	-0.45	1.35	0.15	0.0120	7.36
RBC x10 ⁶ /µl	1	4.4	4.36	8.76	8.69	0.0140	0.18	0.04	0.05	0.0040	-0.15
Hb g/dl	1	12.5	12.5	25	24.8	0.0290	0.25	0	0.1	0.0100	-0.45
HCT%	1	39.5	39.3	78.8	79.6	0.2150	-0.14	0.2	0.4	0.0290	-0.34
MCV-fl	1	90.1	89.7	179.8	183.3	0.3570	-0.35	0.4	0.4	0.0270	0.00
MCH-Pg	1	28.7	28.5	57.2	57.3	0.0800	-0.05	0.2	0.3	0.0200	-0.34
MCHC-g/dl	1	31.9	31.7	63.6	62.4	0.1610	0.28	0.2	0.3	0.0230	-0.27
Plt. x10 ³ /µl	1	190	187	377	348	1.70	0.65	3	7	0.47	-0.45
Retic %	2										

P.S . Assesment

YOUR REPORT		CONSENSUS REPORT
DLC%	3	Nrbcs=5 , Poly=44 L=1, E=3, Mono/Promono=0 , B1=0 P.M.=1, Mye=33, Meta=18, Other=
RBC Morphology	3	Predominantly: Normocytic/Normochromic; Moderate: Anisocytosis, hypochromia, Microcytosis; Mild: Macrocytosis, Poikilocytosis
Diagnosis	3	Chronic Myeloid Leukemia (Chronic Phase)

COMBINED DATA VALUES OF TOTAL PARTICIPANTS

Test parameters	S.No.	Total participants covered in the current dist. 156--L	Total No. responded	% of Labs with Z Score 0-2		% of Labs with Z Score 2-3		% of Labs with Z Score >3	
				Among labs	Within lab	Among labs	Within lab	Among labs	Within lab
WBC x10³/µl	1	318	306	92.16	82.35	5.56	5.88	2.28	11.77
RBC x10⁶/µl	1	318	318	87.11	83.65	5.35	4.4	7.54	11.95
Hb g/dl	1	318	318	87.74	84.91	3.14	5.35	9.12	9.74
HCT%	1	318	307	92.18	89.25	4.56	5.21	3.26	5.54
MCV-fl	1	318	307	91.21	91.86	5.54	3.58	3.25	4.56
MCH-Pg	1	318	307	89.25	85.67	4.89	3.58	5.86	10.75
MCHC-g/dl	1	318	307	90.88	86.97	5.86	5.21	3.26	7.82
Plt. x10³/µl	1	318	307	91.86	92.18	5.86	2.61	2.28	5.21
ReticCount%	2	318	218	91.28	83.94	3.67	13.3	5.05	2.76
PS Assessment	3	318	216	Satisfactory :93.4%, Borderline Sat. :2.83%, Unsatisfactory :3.77%					

***Comments:**

1). **Among Lab (EQA) : Results acceptable.**

2). **Within Lab (IQA) : Difference in the CBC measurement values for WBC unacceptable, may be due to random/human error.**

Note-1: EQA (External Quality Assurance) : Your Performance among various of participating labs in PT, to determine the accuracy of your results.

IQA (Internal Quality Assurance) : Your Performance of comparison of two consecutive measurement values within your lab to test the precision of your autoanalyzer.

Note-2: Z score among & within lab were calculated, as per to ISO/IEC 13528:2015 standard. Z score among lab (EQA)= (Your Result Sum of two values - Consensus Result sum of two values)/(Normalised IQR)

Z score within lab (IQA)= (Your Result Difference of two values - Consensus Result difference of two values)/(Normalised IQR)

IQR = Quartile 3 - Quartile 1 of participant data, Normalised IQR = 0.7413 x IQR

Note-3: Z score 0 to ±2: Acceptable, Z score ±2 to ±3 :Warning Signal, Z score > ±3 : Unacceptable [As per ISO/IEC 13528:2015 standard]

Note-4: Z score value between "0 to ±2" are texted in green colour. Z score value between "±2 to ±3" are texted in orange colour. Z score value > ±3 are texted in red colour.

Note-5: Homogeneity and stability testing of PT sample were done as per ISO 13528:2015 standard. To pass homogeneity test, between sample SD (Ss) should be smaller than the check value (0.3*SDPA). To pass the stability test, average difference in measurement values of first and last day sample ($\bar{x}-\bar{y}$) should be smaller than the check value (0.3*SDPA).

Note-6: ISHTM-AIIMS-EQAP does not subcontract any task of its scheme

Note-7: Participants are free to use methods/analyzer of their own choice.

Note-8: Proficiency testing (PT) samples are sent quarterly to each participant.

Note-9: All the necessary details regarding design and implementation of PT, are provided in the instruction sheet as well as on programme's website www.ishtmaiimseqap.com.

Note 10: Reports are kept confidential.

Report authorized by,



Dr. Seema Tyagi (Prof.)

PT Co-ordinator: ISHTM-AIIMS-EQAP

Department of Hematology, AIIMS, New Delhi

-----End Of Report-----

Title	PT/ EQAS EVALUATION RECORD
Document Number	FRM.QCM.03
Version	01
Amendment No	01
Effective Date	01.09.2022

Date of Investigation: 29/09/2022

PT/EQAS Set Identification:	ISHTM - AIIMS (156-L)
Date of PT/EQAS:	July-22
Acceptable/ Unacceptable Results	- 7.36
Acceptable Result Range:	
Previous Trends/ Unacceptable Results from this Analyte/ Test:	- No
<p>Classification of Problems: (Please tick)</p> <p>Clerical:</p> <p><input type="checkbox"/> Transcription error (may be pre- or post-analytical factors)</p> <p><input type="checkbox"/> Wrong method has been registered for analysis or method change not updated.</p>	
<p>Details of Investigation:</p> <p>Nil</p>	
<p>Methodological</p> <p><input type="checkbox"/> Instrument function checks (e.g., temperatures, blank readings, pressures) not performed as necessary, or results not within acceptable range.</p> <p><input type="checkbox"/> Scheduled instrument maintenance not performed appropriately.</p> <p><input type="checkbox"/> Incorrect instrument calibration.</p> <p><input type="checkbox"/> Standards or reagents improperly reconstituted and stored, or inadvertently used beyond expiration date.</p> <p><input type="checkbox"/> Instrument probes misaligned.</p> <p><input type="checkbox"/> Problem with instrument data processing functions. The laboratory may need to contact the manufacturer to evaluate such problems.</p> <p><input type="checkbox"/> Problem in manufacture of reagents / standards, or with instrument settings specified by manufacturer</p> <p><input type="checkbox"/> Carry-over from previous specimen.</p>	

Title	PT/ EQAS EVALUATION RECORD
Document Number	FRM.QCM.03
Version	01
Amendment No	01
Effective Date	01.09.2022

- Automatic pipettor not calibrated to acceptable precision and accuracy.
- Imprecision from result being close to detection limit of method.
- QC material not run within expiration date, or improperly stored.
- QC material not run at relevant analyte concentration.
- Result not within reportable range (linearity) for instrument / reagent system.
- Obstruction of instrument tubing / orifice by clot or protein.
- Incorrect incubation times.

Details of Investigation:

Nil

Technical

- EQA material improperly reconstituted.
- Testing delayed after reconstitution of EQA material (with problem from evaporation or deterioration).
- Sample not placed in proper order on instrument.
- Result released despite unacceptable QC data.
- QC data within acceptable limits but showed trend suggestive of problem with the assay.
- Inappropriate quality control limits / rules. If the acceptable QC range is too wide, the probability increases that a result will fall within the acceptable QC range yet exceed acceptable limits for EQA.
- Manual pipetting / diluting performed inaccurately, at an incorrect temperature or with incorrect diluent.
- Calculation error or result reported using too few significant digits.
- Secondary specimen tubes incorrectly labeled.
- In addition to above discipline specific errors may also occur

Details of Investigation:

Nil

Problem with PT/EQAS Material

- Matrix effects: The performance of some instrument / method combinations may be affected by the matrix of the PT/EQAS sample. This can be overcome to some extent by assessing participants in peer groups – to be done by the PT/EQAS provider.

Title	PT/ EQAS EVALUATION RECORD
Document Number	FRM.QCM.03
Version	01
Amendment No	01
Effective Date	01.09.2022

- Non-homogenous test material due to variability in fill volumes, inadequate mixing, or inconsistent heating of lyophilized specimens.
- Non-viable samples for microbiology PT/EQAS program.
- Haemolysis on an immune-haematology program samples.

Details of Investigation:

Nil

Problem with PT/EQAS Evaluation

- Peer group not appropriate.
- Inappropriate target value: Target values developed from participant consensus can be inappropriate from non-homogeneous testing material or lingering ("masked") outliers. However, occasional inappropriate target values occur in every PT program. Inappropriate evaluation interval: An evaluation interval may be inappropriately narrow e.g. if ± 2 standard deviation units are used with an extremely precise method; the acceptable range may be much narrower than needed for clinical usefulness.
- Incorrect data entry by PT provider.

Details of Investigation:

Nil

No Explanation: Attributed to Random Error

Any Others (explain)

Summary of Investigation:

NO any specific deviations noted in IQC

Title	PT/ EQAS EVALUATION RECORD
Document Number	FRM.QCM.03
Version	01
Amendment No	01
Effective Date	01.09.2022

Was patient data affected? & Corrective action taken if Patient data was affected.	
NO	
Corrective/ Preventive action taken to prevent Reoccurrence	
Performance montly monitored closely in next sample	
Conclusions	
Based on inter Laboratory Comparison Study and IOL performance Suspected random error.	
<p>Sudhanshu Saini</p> <p>Quality Manager/ Team Leader</p>	<p>Date: 24.12.2022</p>
<p>Lab Head</p> <p><i>[Signature]</i></p>	<p>Date: 21.12.2022</p>

Controlled Copy




NON AVAILABILITY OF EQAP MATERIAL



me

dr.biplabkumarbiswas@yahoo.in

[Show less](#)

- To
-  EQAP accuracy2000@gmail.com
 -  Md Ehtesham mdehtesham@lupindia...
 -  Kanchan Amar Mundhe kanchanmund...
 -  Manisha Khanna manishakhanna@lup...
 -  HLM JeevanSuraksha hlmjsh@lupindi...

23 Dec 2022 at 7:33 pm

RESPECTED SIR/ MADAM,

I AM DR BIPLAB BISWAS, LAB HEAD, LUPIN DIAGNOSTICS & HLM JEEBAN SURAKSHA HOSPITAL, BANKURA, WEST BENGAL. WE HAVEN'T RECEIVED THE EQAP MATERIAL OF FOURTH QUARTER (OCT - DEC '22 TILL NOW.

REGARDS
DR BIPLAB BISWAS
9775689990



Delete



Move to



Forward



Reply



More

Objective:

Inter Laboratory comparison study conducted because AIMMS EQAS sample of period October-December-2022 is not received. As a part of preventive action ILC study performed with two patient samples.

Reference Laboratory- Lupin Diagnostics, NRL

Date of study conducted- 24.12.2022

Sr No	Parameters	NRL	JSH	%Diff	Reference range	%Clinical Correlation
1	RBC	5.68	5.68	0.00	3.8-6	100
2	HB	15.8	16	-1.25	11.5-17	100
3	PCV	50.2	48.5	3.51	35-52	100
4	MCV	88.3	85.3	3.52	76-100	100
5	HCH	27.7	28.2	-1.77	27-34	100
6	HCHC	31.4	33	-4.85	32-35	0
7	RDWCV	15.3	14.1	8.51	11.0-17	100
8	RDWSD	50.4	47.9	5.22	37-49	75
9	PLT	316	281	12.46	150-400	100
10	PCT	0.38	0.34	11.76	0.15-0.40	100
11	MPV	11.9	12.1	-1.65	8.0-11	75
12	PDW	23.2	22	5.45	11.0-22	75
13	PLCC	162	143	13.29	44-140	100
14	PLCR	51.4	50.8	1.18	18-50	100
15	WBC	7.11	7.23	-1.66	3.5-10	100
16	NEUT	3.54	3.58	-1.12	1.6-7	100
17	LYMP	2.76	2.63	4.94	1.0-3	100
18	MONO	0.49	0.56	-12.50	0.2-0.8	100
19	EOS	0.24	0.3	-20.00	0.0-0.50	100
20	BASO	0.04	0.07	-42.86	0.0-0.15	100
21	LIC	0.04	0.09	-55.56	0.0-0.10	75
22	NEUT%	49.8	50.3	-0.99	40-73	75
23	LYM%	39.1	36.8	6.25	15-45	75
24	MONO%	7	7.8	-10.26	4.0-12	100
25	EOS%	3.5	4.2	-16.67	0.5-7	100
26	BASO%	0.6	0.9	-33.33	0.0-2.0	100
27	LIC%	0.6	1.2	-50.00	0.0-1.0	75
Average						89.81

Sr No	Parameters	NRL	JSH	%Diff	Reference range	%Clinical Correlation
1	RBC	5.15	5.24	-1.72	3.8-6	100
2	HB	15.3	15.3	0.00	11.5-17	100
3	PCV	46.7	46	1.52	35-52	100
4	MCV	90.6	87.8	3.19	76-100	100
5	HCH	29.8	29.2	2.05	27-34	100
6	HCHC	32.9	33.2	-0.90	32-35	75
7	RDWCV	15.7	15.2	3.29	11.0-17	100
8	RDWSD	51.2	52.1	-1.73	37-49	100
9	PLT	165	170	-2.94	150-400	100
10	PCT	0.13	0.13	0.00	0.15-0.40	100
11	MPV	8	7.9	1.27	8.0-11	100
12	PDW	11.5	10.6	8.49	11.0-22	25
13	PLCC	31	28	10.71	44-140	100

14	PLCR	18.9	16.7	13.17	18-50	50
15	WBC	6.43	6.32	1.74	3.5-10	100
16	NEUT	4.39	4.18	5.02	1.6-7	100
17	LYMP	1.36	1.38	-1.45	1.0-3	100
18	MONO	0.49	0.57	-14.04	0.2-0.8	100
19	EOS	0.15	0.13	15.38	0.0-0.50	100
20	BASO	0.03	0.05	-40.00	0.0-0.15	100
21	LIC	0.01	0.01	0.00	0.0-0.10	100
22	NEUT%	68.4	66.5	2.86	40-73	100
23	LYM%	21.2	21.8	-2.75	15-45	100
24	MONO%	7.7	9	-14.44	4.0-12	100
25	EOS%	2.3	2	15.00	0.5-7	100
26	BASO%	0.4	0.7	-42.86	0.0-2.0	100
27	LIC%	0.2	0.2	0.00	0.01.0	100
						94.44

Observations-

- ✓ >80% Clinical correlation noted in both samples.
- ✓ High % Difference noted due to statistical limitations.

Conclusion:

Based on obtained result recovery Inter laboratory comparison study successfully passed for CBC test parameter.

Swarup Saini

Documented By
(Mr Swarup Saini)



Approved by
(Dr Biplab K.)

Results

Run Date 24/12/2022 08:09:38 PM

Operator JSH

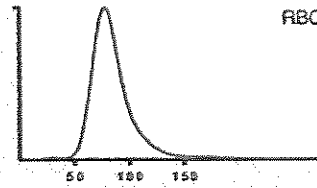
Last Name
First Name
Gender
Patient ID
Birth Date
Sample comments

Sample ID JITENDRA VALID
Rack/Pos
Department
Physician
Type Standard

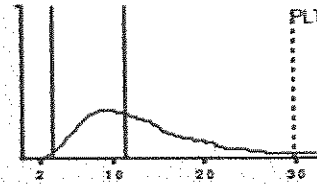
			Range
RBC	5.68	10 ⁹ /μL	3.80 - 6.00
HGB	16.0	g/dL	11.5 - 17.0
HCT	48.5	%	35.0 - 52.0
MCV	85.3	μm ³	76.0 - 100.0
MCH	28.2	pg	27.0 - 34.0
MCHC	33.0	g/dL	32.0 - 35.0
RDW-CV	14.1	%	11.0 - 17.0
RDW-SD	47.9	μm ³	37.0 - 49.0

Recommended actions

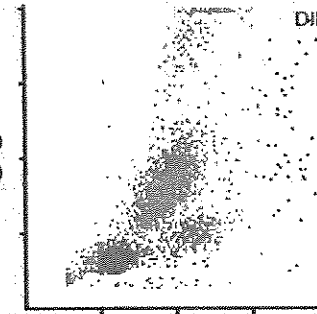
- Slide review
- Abnms**
- Control failed
- Control solution expired
- WBC**
- LYM Interference
- Susp. Pathologies**
- Macroplatelets
- PLT aggregate or NRBC ?



		*		Range
PLT	281	*	10 ⁹ /μL	150 - 400
PCT	0.34	*	%	0.15 - 0.40
MPV	12.1	H*	μm ³	8.0 - 11.0
PDW	22.0	*	μm ³	11.0 - 22.0
P-LCC	143	h	10 ³ /μL	44 - 140
P-LCR	50.8	h	%	18.0 - 50.0



		*		Range	
WBC	7.23	*	10 ³ /μL	3.50 - 10.00	
	#		Range	%	Range
NEU	3.58	*	1.60 - 7.00	50.3	* 40.0 - 73.0
LYM	2.63	*	1.00 - 3.00	36.8	* 15.0 - 45.0
MON	0.56	*	0.20 - 0.80	7.8	* 4.0 - 12.0
EOS	0.30	*	0.00 - 0.50	4.2	* 0.5 - 7.0
BAS	0.07	*	0.00 - 0.15	0.9	* 0.0 - 2.0
LIC	0.09	*	0.00 - 0.10	1.2	h* 0.0 - 1.0



Slide Review

Neutrophil	Myeloblast	Anisocytosis
Lymphocyte	Promyelocyte	Hypochromia
Monocyte	Myelocyte	Polychromasia
Eosinophil	Metamyelocyte	Poikilocytosis
Basophil	Blast	Microcytosis
Atypical Lymphocyte	Target Cell	Macrocytosis
Other	Sickle Cell	Platelet Clumps

Reviewed on _____ by _____ Signature :

Results

Run Date 24/12/2022 07:55:44 PM

Operator LUPIN

Last Name

Sample ID AUTO_SID0005

First Name

Rack/Pos 041731/1

Gender

Age

Department

Patient ID

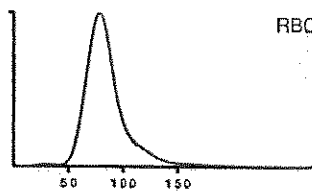
Physician

Birth Date

Type Standard

Sample comments

				Range
RBC	5.68	*	10 ⁶ /μL	3.80 - 6.00
HGB	15.8		g/dL	11.5 - 17.0
HCT	50.2	*	%	35.0 - 52.0
MCV	88.3	*	μm ³	76.0 - 100.0
MCH	27.7		pg	27.0 - 34.0
MCHC	31.4	L	g/dL	32.0 - 35.0
RDW-CV	15.3		%	11.0 - 17.0
RDW-SD	50.4	h	μm ³	37.0 - 49.0



RBC

Recommended actions

Slide review

Alert

Control failed

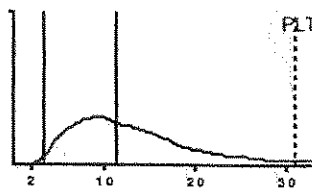
PLT

RBC PLT Interference

Susp. Pathologies

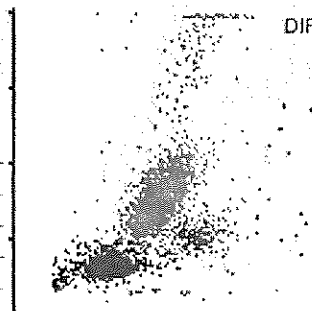
PLT aggregate ?

				Range
PLT	316	*	10 ³ /μL	150 - 400
PCT	0.38	*	%	0.15 - 0.40
MPV	11.9	h*	μm ³	8.0 - 11.0
PDW	23.2	h*	μm ³	11.0 - 22.0
P-LCC	162	h	10 ³ /μL	44 - 140
P-LCR	51.4	h	%	18.0 - 50.0



PLT

			Range
WBC	7.11		10 ³ /μL 3.50 - 10.00
	#	Range	% Range
NEU	3.54	1.60 - 7.00	49.8 40.0 - 73.0
LYM	2.76	1.00 - 3.00	39.1 15.0 - 45.0
MON	0.49	0.20 - 0.80	7.0 4.0 - 12.0
EOS	0.24	0.00 - 0.50	3.5 0.5 - 7.0
BAS	0.04	0.00 - 0.15	0.6 0.0 - 2.0
LIC	0.04	0.00 - 0.10	0.6 0.0 - 1.0



DIF

Slide Review

Neutrophil	Myeloblast	Anisocytosis
Lymphocyte	Promyelocyte	Hypochromia
Monocyte	Myelocyte	Polychromasia
Eosinophil	Metamyelocyte	Poikilocytosis
Basophil	Blast	Microcytosis
Atypical Lymphocyte	Target Cell	Macrocytosis
Other	Sickle Cell	Platelet Clumps

Reviewed on _____ by _____ Signature :

Results

Run Date 24/12/2022 08:13:01 PM

Operator JSH

Last Name

Sample ID RUSHIKESHVALID

First Name

Rack/Pos

Gender

Age

Department

Patient ID

Physician

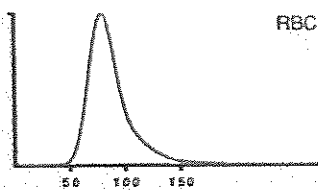
Birth Date

Type Standard

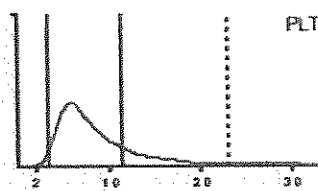
Sample comments

Alarms
Control failed
Control solution expired

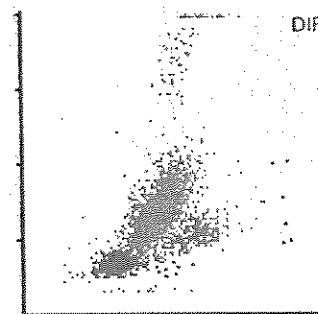
			Range
RBC	5.24	10 ⁶ /μL	3.80 - 6.00
HGB	15.3	g/dL	11.5 - 17.0
HCT	46.0	%	35.0 - 52.0
MCV	87.8	μm ³	76.0 - 100.0
MCH	29.2	pg	27.0 - 34.0
MCHC	33.2	g/dL	32.0 - 35.0
RDW-CV	15.2	%	11.0 - 17.0
RDW-SD	52.1	h μm ³	37.0 - 49.0



			Range
PLT	170	10 ³ /μL	150 - 400
PCT	0.13	%	0.15 - 0.40
MPV	7.9	μm ³	8.0 - 11.0
PDW	10.6	μm ³	11.0 - 22.0
P-LCC	28	10 ³ /μL	44 - 140
P-LCR	16.7	%	18.0 - 50.0



			Range
WBC	6.32	10 ³ /μL	3.50 - 10.00
	#		
NEU	4.18	Range	1.60 - 7.00
		%	66.5
		Range	40.0 - 73.0
LYM	1.38	Range	1.00 - 3.00
		%	21.8
		Range	15.0 - 45.0
MON	0.57	Range	0.20 - 0.80
		%	9.0
		Range	4.0 - 12.0
EOS	0.13	Range	0.00 - 0.50
		%	2.0
		Range	0.5 - 7.0
BAS	0.05	Range	0.00 - 0.15
		%	0.7
		Range	0.0 - 2.0
LIC	0.01	Range	0.00 - 0.10
		%	0.2
		Range	0.0 - 1.0




Slide Review

Neutrophil	Myeloblast	Anisocytosis
Lymphocyte	Promyelocyte	Hypochromia
Monocyte	Myelocyte	Polychromasia
Eosinophil	Metamyelocyte	Poikilocytosis
Basophil	Blast	Microcytosis
Atypical Lymphocyte	Target Cell	Macrocytosis
Other	Sickle Cell	Platelet Clumps

Reviewed on _____ by _____ Signature :

Results

Run Date 24/12/2022 07:58:02 PM

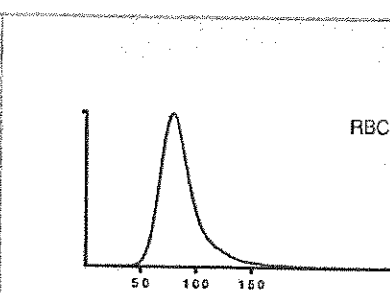
Operator LUPIN 

Last Name
First Name
Gender
Patient ID
Birth Date
Sample comments

Age

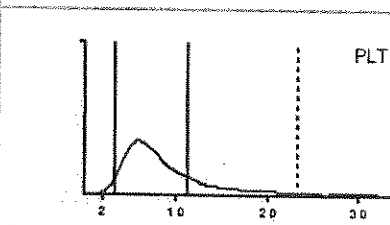
Sample ID AUTO_SID0006
Rack/Pos 041731/2
Department
Physician
Type Standard

			Range
RBC	5.15	10 ⁶ /μL	3.80 - 6.00
HGB	15.3	g/dL	11.5 - 17.0
HCT	46.7	%	35.0 - 52.0
MCV	90.6	μm ³	76.0 - 100.0
MCH	29.8	pg	27.0 - 34.0
MCHC	32.9	g/dL	32.0 - 35.0
RDW-CV	15.7	%	11.0 - 17.0
RDW-SD	51.2	h μm ³	37.0 - 49.0

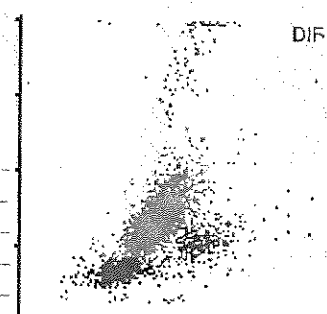


Alarms
Control failed

			Range
PLT	165	10 ³ /μL	150 - 400
PCT	0.13	%	0.15 - 0.40
MPV	8.0	μm ³	8.0 - 11.0
PDW	11.5	μm ³	11.0 - 22.0
P-LCC	31	10 ³ /μL	44 - 140
P-LCR	18.9	%	18.0 - 50.0



			Range	
WBC	6.43	10 ³ /μL	3.50 - 10.00	
	#	Range	%	Range
NEU	4.39	1.60 - 7.00	68.4	40.0 - 73.0
LYM	1.36	1.00 - 3.00	21.2	15.0 - 45.0
MON	0.49	0.20 - 0.80	7.7	4.0 - 12.0
EOS	0.15	0.00 - 0.50	2.3	0.5 - 7.0
BAS	0.03	0.00 - 0.15	0.4	0.0 - 2.0
LIC	0.01	0.00 - 0.10	0.2	0.0 - 1.0



Slide Review

Neutrophil	Myeloblast	Anisocytosis
Lymphocyte	Promyelocyte	Hypochromia
Monocyte	Myelocyte	Polychromasia
Eosinophil	Metamyelocyte	Poikilocytosis
Basophil	Blast	Microcytosis
Atypical Lymphocyte	Target Cell	Macrocytosis
Other	Sickle Cell	Platelet Clumps

Reviewed on _____ by _____ Signature :