Adeuriev Vic	Indicated as a correlation factor with the reference data when at least 100 samples			
differential blood: (ount)	(at least 20 samples for NRBC and IG) of peripheral blood (samples with nucleated RBC for NRBC, samples with immature granulocyte for IG) are analyzed.			
Milaja islojeki mede:	The reference data is obtained by a standard analysis method that uses the flow			
HPCImodes keeps and	cytometry method, based on the standard instrument, standard 5-category white blood cell analysis method, standard NRBC analysis method, or standard immature.			
		e analysis method.	So unaryolo mouros, or olaridara ii	
		- r = 0.80 or more		
		r = 0.90 or more	re, Armania de Carlos de C	
		r = 0.90 or more	i again da agus na galaigh agus agus agus agus agus agus agus agus	
	Allege eta mate Malanti e Apolit da eta	r ≡ 0.75 or more	Standard Color Col	
	100 100 100 100 100 100 100 100 100 100	r = 0.80 or more	e digitari sa sa sa sa giliku kulondari. Manan di inggan digitari sa	
	BASO%	r = 0.50 or more		
	the public of the specific of the specific of the second o	r = 0.80 or more	A STATE OF THE STA	
	Indicated as the average value of the difference between the measured values of a least 100 samples (at least 20 samples for NRBC and IG) of peripheral blood (samples with nucleated RBC for NRBC, samples with immature granulocyte for IG			
	and values	measured on a standard instr	ument.	
	NEUT%	within ±3.0 NEUT%		
	LYMPH%	within ±3.0 LYMPH%		
	MONO%	within ±2.0 MONO%		
	EO%	within ±1.0 EO%	natural de la companya del companya del companya de la companya de	
	BASO%	within ±1.0 BASO%	A serialist	
	IG%	within ±1.5 IG%		

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3/2	Accuracy (differential blood count) (Pre-Dillution) mode	Indicated as a correlation factor with the reference data when at least 100 samples (at least 20 samples for NRBC and IG) of diluted peripheral blood (samples with nucleated RBC for NRBC, samples with immature granulocyte for IG) are analyzed. The reference data is obtained by a standard analysis method that uses the flow cytometry method, based on the standard instrument, standard 5-category white
		blood cell analysis method, standard NRBC analysis method, or standard immature
(8		granulocyte analysis method. NRBC% r = 0.70 or more
		_NEUT%
4		MONO% r = 0.60 or more
		"EO% r = 0.60 or more BASO% r = 0.50 or more
		Indicated as the average value of the difference between the measured values of at
		least 100 samples (at least 20 samples for NRBC and IG) of diluted peripheral blood (samples with nucleated RBC for NRBC, samples with immature granulocyte for IG) and values measured on a standard instrument. NEUT% within ±3.0 NEUT% LYMPH% within ±3.0 LYMPH% MONO% within ±2.0 MONO% EO% within ±1.0 EO% BASO% within ±1.0 BASO%
	Accuracy (differential blood count) (Body Fluid) mode ²	Indicates the correlation with the reference method and the slope of the regression line when 50 or more body fluid samples are analyzed. The reference data are obtained by a method in which slides created by sight spin method are visually classified. MN# $r = 0.9$ or more, and within slope=1 ± 0.5 PMN# $r = 0.9$ or more, and within slope=1 ± 0.5 MN% $r = 0.7$ or more, and within slope=1 ± 0.5 PMN% $r = 0.7$ or more, and within slope=1 ± 0.5

*1 The HPC analysis can only be performed if the instrument offers the HPC analysis mode.

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*2 The body fluid analysis can only be performed if the instrument offers the body fluid analysis mode.

Accuracy	Indicated as a correlation factor with the reference data when at least 100 samples of peripheral blood are analyzed. The reference data are obtained by the standard
(refletjoeyte) parametere	instrument method or the visual observation method.
[HPC] mode ²⁰	RET% r = 0.90 or more RET# r = 0.90 or more RETHE r = 0.90 or more
	(More than the half of the samples are RET# $0.020 \times 10^6 / \mu L$ or more) RBC-He r = 0.9 or more
	Delta-He RET-He r = 0.9 or more, RBC-He r = 0.9 or more Indicated as the average value of the difference between the measured values of at least 100 samples of peripheral blood and values measured on a standard
	instrument. RET% within ±20% or ±0.30 RET% RET# within ±20% or ±0.0150 x 10 ⁶ /µL IRF within ±30% or ±10.0 IRF (within 40.0 IRF*) LFR within ±30% or ±10.0 LFR (within 35.0 LFR*) MFR within ±30% or ±10.0 MFR (within 30.0 MFR*) HFR within ±30% or ±5.0 HFR (within 15.0 HFR*) * Control blood or calibrator
Accuracy (reticulocyte parameters*1) [Pre-Dilution] mode	Indicated as a correlation factor with the reference data when at least 100 samples of diluted peripheral blood are analyzed. The reference data are obtained by the standard instrument method or the visual observation method. RET% r = 0.80 or more RET# r = 0.80 or more RET-He r = 0.7 or more RBC-He r = 0.7 or more Delta-He RET-He r = 0.7 or more, RBC-He r = 0.7 or more Indicated as the average value of the difference between the measured values of at least 100 complex of diluted paripheral blood and values measured on a standard
	least 100 samples of diluted peripheral blood and values measured on a standard instrument. RET% within ±30% or ±0.50 RET% RET# within ±30% or ±0.020 x 10 ⁶ /µL IRF within ±50% or ±10.0 IRF LER within ±50% or ±10.0 LER MFR within ±50% or ±10.0 MFR
	HFR within ±50% or ±5.0 HFR

^{*1} These items do not appear with all analyzer types.

^{*2} The HPC analysis can only be performed if the instrument offers the HPC analysis mode.

Linearity [Mitole Shood) mode [FIRG] mode **The state of the state of	Indicated as a logical value or a residual or residual rate with respect to the value measured on a standard instrument. WBC within ±3% or ±0.20 x 10³/μL (0.00 to 100.00 x 10³/μL) within ±6% (100.01 to 310.00 x 10³/μL) within ±11% (310.01 to 440.00 x 10³/μL) RBC within ±2% or ±0.03 x 10⁵/μL (0.00 to 8.00 x 10⁶/μL) within ±4% or ±0.06 x 10⁶/μL (8.01 to 8.60 x ⁴πο⁶/μL) HGB within ±2% or ±0.2 g/dL (0.0 to 25.0 g/dL, 0.00 to 15.52 mmol/L) within ±5% or ±0.5 g/dL (25.1 to 26.0 g/dL, 15.53 to 16.14 mmol/L) HCT within ±3% or ±1.0 HCT (0.0 to 75.0%) PLT*1 within ±5% or ±10 x 10³/μL (0 to 1000 x 10³/μL) within ±6% (1001 to 5000 x 10³/μL) PLT*2,4 within ±7% or ±10 x 10³/μL (0 to 5000 x 10³/μL) PLT*3,4 within ±5% or ±10 x 10³/μL (0 to 1000 x 10³/μL) within ±6% (1001 to 5000 x 10³/μL) NRBC# within ±10% or ±0.20 x 10³/μL (0.00 to 20.00 x 10³/μL) NRBC# within ±20% or ±2.0 NRBC% (0.0 to 600.0/100WBC) RET** Within ±20% or ±0.30 RET% (0.00 to 30.00%) RET#* within ±20% or ±0.0150 x 10⁶/μL (0.0000 to 0.7200 x 10⁶/μL) *1 PLT counted in the RBC/PLT channels (PLT particle size distribution). *2 PLT counted in the PLT-F channels. *3 PLT counted in the PLT-F channels.
Einearity	Indicated as a logical value or a residual or residual rate with respect to the value measured on a standard instrument. This specification is based on the verification using control blood.
[Body:Fluid] mode*2	WBC-BF within $\pm 0.010 \times 10^3 / \mu L$ (0.000 to $0.050 \times 10^3 / \mu L$, RBC < $1.000 \times 10^6 / \mu L$) within $\pm 20\%$ (0.051 to $10.000 \times 10^3 / \mu L$, RBC < $1.000 \times 10^6 / \mu L$) within $\pm 2\%$ or $\pm 0.010 \times 10^6 / \mu L$ (0.000 to $5.000 \times 10^6 / \mu L$) within $\pm 0.010 \times 10^3 / \mu L$ (0.000 to $0.050 \times 10^3 / \mu L$, RBC < $1.000 \times 10^6 / \mu L$) within $\pm 20\%$ (0.051 to $10.000 \times 10^3 / \mu L$, RBC < $1.000 \times 10^6 / \mu L$)

The availability of these functions depends on your system configuration.

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Carryover (Whole blood) mode	WBC 1.0% or less
[Pre-Billinton; mode	HGB 1.0% or less
[HRC] mode:	HCT: 1.0% or less
	NRBC# 2.0% or 0.02 x 10 ³ /µL or less NEUT# 2.0% or 0.05 x 10 ³ /µL or less LYMPH# 2.0% or 0.05 x 10 ³ /µL or less
	MONO# 2.0% or 0.03 x 10 ³ /μL or less EO# 2.0% or 0.03 x 10 ³ /μL or less BASO# 2:0% or 0.03 x 10 ³ /μL or less
Carrydvels [Body/Fluid) mode ¹²	WBC-BF 0.3 % or 0.001 x 10 ³ /µL or less RBC-BF 0.3 % or 0.003 x 10 ⁶ /µL or less TC-BF# 0.3 % or 0.001 x 10 ³ /µL or less

- *1 The HPC analysis can only be performed if the instrument offers the HPC analysis mode.
- *2 The body fluid analysis can only be performed if the instrument offers the body fluid analysis mode.

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Sample Stability with Time after Blood	<u> </u>	r blood is taken are shown below.
Collection	T WARRANT R	MMART CONTRACTOR AND ARABAN AND A
3 hours	нст	within +5.0% 3
	MCV	within +5.0%
		Anna and the state of the state
24960188	HCT	within +8.0% (in a refrigerator), within +15.0% (stored at 18 to 26°C)
	MCV	within +8.0% (in a refrigerator), within +15.0% (stored at 18 to 26°C)
Carra National Paragraphs Control	NRBC%	within ±10.0% or ± 3.0 / 100 WBC
	IG%	Within ±2.0 IG%
	RET%*1	within ± 20.0% or ±0.3 RET%
	RET#*1	within ± 20.0% or ±0.015 x 10 ⁶ /μL
	IRF* ¹	within ± 30.0% or ±10.0 IRF
	LFR* ¹ MFR* ¹	within ± 30.0% or ±10.0 LFR
	HFR*1	within ± 30.0% or ±10.0 MFR within ± 30.0% or ±5.0 HFR
	RET-He* ¹	within ± 30.0% or ±5.0 AFA within ± 8.0% (RET# 0.0100 x 10 ⁶ /µL or more)
	RBC-He*1	within ± 8.0% (ΝΕΤ# 0.0100 Χ 10 /μΕ οι Πίοιθ)
	Delta-He*1	RET-He, RBC-He within ± 8.0% (RET# 0.0100 x 10 ⁶ /μL or more)
	NEUT-RI*2	within ±8.0%
	NEUT-GI*2	within ±8.0%
	IPF	within ± 30.0% or ±2.0 IPF% (PLT 100 x 10 ³ /µL or more, IPF 2.0% or more)
	engye	
36 hours	NEUT%	within ±8.0 NEUT%
	LYMPH%	within ±7.0 LYMPH%
	MONO%	within ±3.0 MONO%
	EO%	within ±3.0 EO%
	BASO%	within ±1.0 BASO%
48 hours	PLT* ³	within ±10% or ± 30 x 10 ³ /μL
	PLT*1,4	within ±15.0%
	PLT*1,5	within $\pm 10.0\%$ or $\pm 30 \times 10^3 / \mu L$
	NEUT%	within ±8.0 NEUT%
	LYMPH%	within ±7.0 LYMPH%
	MONO%	within ±4.0 MONO%
	EO%	within ±3.0 EO%
	BASO%	within ±1.0 BASO%
70.1	wsa	
7/2 hours	WBC	within ±10.0% within ±5.0%
	RBC HGB	within ±5.0%
	HGD	With HT ±5.0 %
	*1 These iter	ns do not appear with all analyzer types.
	W	bility of these functions depends on your system configuration.
		ed in the RBC/PLT channels (PLT particle size distribution).
	*4 PLT count	ed in the RET channels.
	*5 PLT count	ed in the PLT-F channels.
	@	
	No 🏈	te:
	The data are	the values when analyzing the samples stored at 18 to 26°C or
	in a refrigera	tor (2 to 8°C). If the samples were refrigerated, they were
		oom temperature before analyzing. Depending on how the
	samples wer	e stored, the values may not fall within the above range.

PAM Storage Capacity	Samples stored: 100,000 samples
	Patient information: 10,000 records
	Wards registered: 200 wards
	Doctor names registered: 200 names
	Analysis registration function: 2,000 records
	QC files: 99 files per analyzer (300 plots per file)
	Reagent replacement history: 5,000 records %2
	Maintenance history: 5,000 records
Diality Control	X-bar control (L-J control): 300 plots x 94 files
	X-barM control: 300 plots x 5 files

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