

Field Safety Notice

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cobas m 511: Potential for discrepancy in red cell parameters in patients with severe microcytic anemia and/or thalassemia

Product Name	cobas m 511 integrated hematology analyzer
GMMI / Part No Device Identifier	07261691190
Type of Action	Field Safety Corrective Action (FSCA)

Dear Valued Customer,

Roche Diagnostics regrets to inform you of reported cases affecting the **cobas m 511** integrated hematology analyzer.

Description of Situation

Discrepant results have been reported during the technical evaluation in patients with severe microcytic anemia (e.g. iron deficiency, thalassemia) and the following parameters are affected: RBC (red blood cell count), HGB (hemoglobin concentration), MCH (mean corpuscular hemoglobin), HCT (hematocrit), MCV (mean corpuscular volume), and RDW-SD (Red blood cell distribution width - standard deviation).

Internal investigations did show that the following parameters might be also affected: MCHC (Mean corpuscular hemoglobin concentration), RDW (Red blood cell distribution width - coefficient of variation), #RET (Reticulocyte count), %RET (Reticulocyte percent), and HGB-RET (Mean reticulocyte hemoglobin content).

This issue is preliminarily linked to the **cobas m 511** software version 1.0.

For global epidemiology of haemoglobin disorders and derived service indicators refer to the attachment.

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Potential medical impacts and risks

Of particular clinical concern are the HGB differences observed within the transfusion decision limits, which might lead to an incorrect transfusion decision.

Root cause analysis

In some cases, with extreme central pallor or hypochromia the RBC count may be low. When there is severe anisocytosis, there is a bias toward measuring smaller cells, thereby underestimating MCV and MCH. The calculated values HGB and HCT will also be lower. Further, derived parameters MCHC, RDW, RDW-SD, #RET, %RET, and HGB-RET might also show erroneous results.

Actions taken by Roche Diagnostics

In all reported cases associated with significant red blood cell differences, the **cobas m 511** integrated hematology analyzer, as designed, displayed messages, including "Anemia", "Anisocytosis", "Hypochromia", "Microcytosis", "RBC fragments", and "RBC interference". These messages prevent the results from being automatically released to the Laboratory Information System (LIS), thus triggering a laboratory review.

In addition, Roche has generated the new message "*RBC discrepancy?*" triggered by the rule " $\text{HGB} \leq 9.0 \text{ g/dL}$ and $\text{MCH} \leq 21.0 \text{ pg}$ ". The rule is finally validated and released and is now implemented by Roche's Application Specialist on your **cobas m 511** analyzer. The rule is marking all relevant eleven (11) parameters with an asterisk (*) which indicates that the results may be unreliable.

Roche will provide an update of the **cobas m 511** software and the corresponding user documentation which will be rolled-out in Q4, 2018.

Actions to be taken by the customer/user

When the message "*RBC discrepancy?*" is reported for a patient result, alternative methods shall be used to verify the relevant parameters as they may be unreliable. Review of the patient sample is recommended.

After implementing the rule, when the existing rule set needs to be changed, do NOT use the "Create" button (refer to button 1 in the image below) because "Create" will generate a new rule set without the "*RBC discrepancy?*" rule. ONLY use the "Duplicate" button (refer to button 2 in the image below) to duplicate the existing rule set including the "*RBC discrepancy?*" rule and modify the duplicated rule set as required.

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The screenshot shows the 'cobas m 511 CIR' software interface. The 'Configuration' tab is active, and the 'Rules' section is expanded. The 'Messages' table lists various parameters and their associated reasons. Red boxes highlight the 'Duplicate' and 'Create' buttons in the 'Rules' section.

Message	Type	Parameters	Hold	Reason	Repeat/Reflex
Microcytosis	RBC	MCV	✓	RBC	None
nRBCs?	RBC	%NRBC	✓	NRBC	None
RBC fragments?	RBC	RBC	✓	RBC	None
RBC inclusions?	RBC	%NEUT	✓	RBC with inclusion	None
RBC linearity	RBC	RBC	✓	None	None
Retic linearity	RBC	%RET	✓	None	None
Reticulocytosis	RBC	%RET	✓	None	None
Giant PLTs?	PLT	Parameters: 2	✓	Giant platelet	None
PLT abnormal first time	PLT	PLT	✓	None	None
PLT linearity	PLT	PLT	✓	None	None
Thrombocytopenia	PLT	PLT	✓	Platelet	None
Thrombocytosis	PLT	PLT	✓	Platelet	None
<600 cells counted for differential	SYS	Parameters: 5	✓	WBC differential	None

You are kindly advised to daily check the presence of the rule in your activated rule set. This can be done in the following way:

1. Select the “Factory settings” rule set and verify that an uneditable row appears with the message “RBC discrepancy?” and the condition “HGB ≤ 9 g/dL –And– MCH ≤ 21 pg”.

The screenshot shows the 'cobas m 511 CIR' software interface. The 'Configuration' tab is active, and the 'Rules' section is expanded. The 'Messages' table lists various parameters and their associated reasons. Red boxes highlight the 'Duplicate' and 'Create' buttons in the 'Rules' section. Red arrows point to the 'RBC discrepancy?' message and the 'HGB ≤ 9 g/dL –And– MCH ≤ 21 pg' condition, with labels 'Not editable!' and 'Not active!'.

Message	Conditions
Neutrophilia	#NEUT > 20.00×10 ⁹ /L -And- History unavailable or previous #NEUT < 20.00×10 ⁹ /L
nRBCs?	%NRBC > 2.0/100 WBC -And- History unavailable or previous %NRBC ≤ 2.0/100 WBC
PLT abnormal first time	PLT < 100×10 ⁹ /L -Or- PLT > 1,000×10 ⁹ /L -And- History unavailable or previous PLT ≥ 100×10 ⁹ /L -And- History unavailable or previous PLT ≤ 1,000×10 ⁹ /L
PLT clumps?	clump_area > 30,000
PLT linearity	PLT > 5,000×10 ⁹ /L
RBC agglutination?	%agglutination > 50%
RBC discrepancy?	HGB ≤ 9.0 g/dL -And- MCH ≤ 21.0 pg
RBC fragments?	%RBCfrag > 1.00%
RBC inclusions?	%RBCi ≥ 0.04%
RBC interference?	MCHC < 31.0 g/dL -Or- MCHC > 37.0 g/dL
RBC linearity	RBC > 8.00×10 ⁹ /L
Retic linearity	#RET > 0.60×10 ⁹ /L
Reticulocytosis	#RET > 0.10×10 ⁹ /L -And- History unavailable or previous #RET ≤ 0.10×10 ⁹ /L

2. Select a user-created rule set (if available) and verify that the same rule appears.
3. Roll back that rule set to a previous version (if available) and verify that the same rule appears.
4. Navigate to the Configuration > Rules > Messages tab.
5. Verify that the “RBC discrepancy?” message exists and that it is uneditable, except for the “Reason” and “Repeat/Reflex” columns.

For customers evaluating the instrument Roche kindly advises to not report values for diagnostic use until your instrument is updated with a new **cobas m 511** software version including this rule.

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Labelling update

The User Assistance and, more specifically, the table in the section “List of default system messages about the sample” will be updated to include a description of the message “*RBC discrepancy?*” as presented in Table 1.

Message	Default Description	Recommended action
RBC discrepancy?	MCH ≤ 21.0 pg and HGB ≤ 9.0 g/dL	Use alternative methods to verify the following parameters, as they may be unreliable: RBC, HGB, HCT, MCV, MCH, MCHC, RDW, RDW-SD, %RET, #RET, and HGB-RET.

Table 1: New item that will be introduced in the User Assistance, in the section “List of default system messages about the sample”.

The User Assistance will also be updated to include the following message:

*For diagnostic purposes, **cobas m 511 integrated hematology analyzer** results should always be assessed in conjunction with the patients’ medical history, clinical examination, and other findings.*

The “System analytical performance characteristics” document will be updated to reflect the identified potentially interfering conditions. More specifically, the section currently titled as “System limitations and interfering substances” will be updated and the “Interfering conditions” sub-section will include the following applicable statements:

Red blood cells If any of the following are present, an erroneous low value may result for the RBC, HGB, HCT, MCV, MCH, RDW and/or RDW-SD parameters, and/or an erroneous low or high value may result for the MCHC parameter.

- Erythrocyte aggregation (e.g., cold agglutinin)
- Microerythrocytes
- Fragmented red blood cells

Reticulocytes If any of the following are present, an erroneous high value may result for the #RET parameter.

- Malaria
- Howell-Jolly body
- Thrombocytosis

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If any of the following are present, an erroneous low value may result for the #RET parameter, and/or an erroneous low or high value may result for the %RET and/or HGB-RET parameters.

- Erythrocyte aggregation (e.g., cold agglutinin)
- Microerythrocytes
- Fragmented red blood cells

Communication of this Field Safety Notice

This notice must be passed on to all those who need to be aware within your organization or to any other organization/individual where the potentially affected devices have been distributed/supplied. Please pass on this notice to the Chairman Medical Board and Head of Department as well, as required by HSA.

Please maintain awareness of this notice and resulting action for an appropriate period to ensure the effectiveness of the corrective action

We apologize for any inconvenience this may cause and hope for your understanding and your support.

Sincerely,

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