

Order information

REF	CONTENT		Analyzer(s) on which kit(s) can be used
07005806 190	LDL-Cholesterol Gen.3 (2 × 50 tests)		cobas c 111
12172623 122	Calibrator f.a.s. Lipids (3 × 1 mL)	Code 424	
10781827 122	Precinorm L (4 × 3 mL)	Code 304	
11778552 122	Precipath HDL/LDL-C ($4 \times 3 \text{ mL}$)	Code 319	
05117003 190	PreciControl ClinChem Multi 1 (20 × 5 mL)	Code 391	
05947626 190	PreciControl ClinChem Multi 1 (4 × 5 mL)	Code 391	
05117216 190	PreciControl ClinChem Multi 2 (20 × 5 mL)	Code 392	
05947774 190	PreciControl ClinChem Multi 2 (4 × 5 mL)	Code 392	
04774230 190	Diluent NaCl 9 %	Code 951	

English

System information

LDLC3: ACN 552

Intended use

In vitro test for the quantitative determination of LDL-cholesterol in human serum and plasma on the cobas c 111 system.

Summarv

Low Density Lipoproteins (LDL) play a key role in causing and influencing the progression of atherosclerosis and, in particular, coronary sclerosis.^{1,2} The LDLs are derived from VLDLs (Very Low Density Lipoproteins) rich in triglycerides by the action of various lipolytic enzymes and are synthesized in the liver. The elimination of LDL from plasma takes place mainly by liver parenchymal cells via specific LDL receptors. Elevated LDL concentrations in blood and an increase in their residence time coupled with an increase in the biological modification rate results in the destruction of the endothelial function and a higher LDL-cholesterol uptake in the monocyte/macrophage system as well as by smooth muscle cells in vessel walls. The majority of cholesterol stored in atherosclerotic plaques originates from LDL. The LDLcholesterol value is the most powerful clinical predictor among all of the single parameters with respect to coronary atherosclerosis. Therefore, therapies focusing on lipid reduction primarily target the reduction of LDLcholesterol which is then expressed in an improvement of the endothelial function, prevention of atherosclerosis and reducing its progression as well as preventing plaque rupture.

Various methods are available for the determination of LDL-cholesterol such as ultracentrifugation as the reference method, lipoprotein electrophoresis, HPLC and precipitation methods.^{3,4} In the precipitation methods apolipoprotein B-containing LDL-cholesterol is, for example, precipitated using either polyvinyl sulfate, dextran sulfate or polycyclic anions. The LDL-cholesterol content is usually calculated from the difference between total cholesterol and cholesterol in the remainder (VLDL and HDL cholesterol) in the superpote ofter preprinticing with polycing and HDL-cholesterol) in the supernate after precipitation with polyvinyl sulfate and dextran sulfate.5 Lipid Research Clinics recommend a combination of ultracentrifugation and precipitation methods using polyanions in the presence of divalent cations. The precipitation methods are, however, time-consuming, cannot be automated and are susceptible to interference by hyperlipidemic serum, particularly at high concentrations of free fatty acids. A more recent method is based on the determination of LDL-cholesterol after the sample is subjected to immunoadsorption and centrifugation.6

The calculation of the LDL-cholesterol concentration according to Friedewald's formula is based on two cholesterol determinations (total cholesterol and HDL-cholesterol) and one triglyceride determination.

Friedewald formula for calculation of LDL-C presumes that a direct relationship exists between VLDL-cholesterol and triglycerides in fasting blood samples (VLDL-cholesterol = Trigl./5 mg/dL, VLDL-cholesterol = Trigl./2.2 mmol/L). The bias in calculating LDL-C using this assumption is only acceptable in samples with a triglyceride concentration < 2.0 mmol/L (177 mg/dL).^{8,9} Even in the presence of small amounts of chylomicrons or abnormal lipoproteins, the formula gives rise to artificially low LDLcholesterol values. Non-fasting samples cannot be used for the calculation of LDL-C because they contain a high concentration of chylomicrons and in many cases the limit of acceptable triglyceride concentration is exceeded.

For these reasons, a simple and reliable method for routine measurement of LDL-cholesterol without any preparatory steps was developed. This automated method for the direct determination of LDL-cholesterol takes

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advantage of the selective micellary solubilization of LDL-cholesterol by a nonionic detergent and the interaction of a sugar compound and lipoproteins (VLDL and chylomicrons). When a detergent is included in the enzymatic method for cholesterol determination (cholesterol esterase cholesterol oxidase coupling reaction), the relative reactivities of cholesterol in the lipoprotein fractions increase in this order: HDL < chylomicrons < VLDL < LDL.

The combination of a sugar compound with detergent enables the selective determination of LDL-cholesterol in serum and plasma samples.

Non-fasting sample results are slightly lower than fasting results. Comparable non-fasting results were observed with the beta quantification bias \leq 4 % versus reference method, and \leq 12 % total analytical error.^{11,12,13,14} method.¹⁰ This direct assay meets the 1995 NCEP goals of < 4 % total CV,

Test principle

Homogeneous enzymatic colorimetric assay

Cholesterol esters and free cholesterol in LDL are measured on the basis of a cholesterol enzymatic method using cholesterol esterase and cholesterol oxidase in the presence of surfactants which selectively solubilize only LDL. The enzyme reactions to the lipoproteins other than LDL are inhibited by surfactants and a sugar compound. Cholesterol in HDL, VLDL and chylomicron is not determined.

LDL-cholesterol esters + H₂O

LDL-cholesterol + O_2

cholesterol esterase

detergent

cholesterol + free fatty acids (selective micellary solubilization)

>

Cholesterol esters are broken down quantitatively into free cholesterol and fatty acids by cholesterol esterase.

cholesterol oxidase

 Δ^4 -cholestenone + H₂O₂

In the presence of oxygen, cholesterol is oxidized by cholesterol oxidase to Δ^4 -cholestenone and hydrogen peroxide.

peroxidase >

red purple pigment + 5 H₂O

I

a) N-ethyl-N-(3-methylphenyl)-N-succinylethylenediamine

In the presence of peroxidase, the hydrogen peroxide generated reacts with 4-aminoantipyrine and EMSE to form a red purple dye. The color intensity of this dye is directly proportional to the cholesterol concentration and is measured photometrically.

Reagents - working solutions

R1 Bis-tris^{b)} buffer: 20.1 mmol/L, pH 7.0; 4-aminoantipyrine: 0.98 mmol/L; ascorbate oxidase (AOD, Acremonium spec.): \geq 66.7 µkat/L; peroxidase (recombinant from Basidiomycetes): \geq 166.7 µkat/L; BSA: 4.0 g/L; preservative



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SR MOPS^{c)} buffer: 20.1 mmol/L, pH 7.0; EMSE: 2.16 mmol/L; cholesterol esterase (Pseudomonas spec.): ≥ 33.3 µkat/L; cholesterol oxidase (recombinant from E. coli): ≥ 31.7 µkat/L; peroxidase (recombinant from Basidiomycetes): ≥ 333.3 µkat/L; BSA: 4.0 g/L; detergents; preservative

b) bis(2-hydroxyethyl)-amino-tris-(hydroxymethyl)methane

c) 3-morpholinopropane-1-sulfonic acid

Precautions and warnings

For in vitro diagnostic use.

Exercise the normal precautions required for handling all laboratory reagents.

Disposal of all waste material should be in accordance with local guidelines. Safety data sheet available for professional user on request.

This kit contains components classified as follows in accordance with the Regulation (EC) No. 1272/2008:



Warning

H317 May cause an allergic skin reaction.			
Prevention:			
P261	Avoid breathing dust/fume/gas/mist/v	/apours/spray.	
P272	Contaminated work clothing should r the workplace.	not be allowed out of	
P280	Wear protective gloves.		
Response:			
P333 + P313	If skin irritation or rash occurs: Get m advice/attention.	nedical	
P362 + P364	Take off contaminated clothing and v	vash it before reuse.	
Disposal:			
P501	Dispose of contents/container to an a disposal plant.	approved waste	
	/ labeling follows EU GHS guidance.		
Contact phone	e: all countries: +49-621-7590		
Reagent hand Ready for use			
Storage and	stability		
LDLC3			
Shelf life at 2-8 °C:		See expiration date on reagent.	
On-board in use and refrigerated on the analyzer:		3 weeks	
Diluent NaCl S	9 %		
Shelf life at 2-8 °C:		See expiration date on reagent.	
On-board in u	4 weeks		
Specimen collection and preparation For specimen collection and preparation only use suitable tubes or collection containers.			
Only the specimens listed below were tested and found acceptable. Serum			
Plasma: Li-heparin, K_2 - and K_3 -EDTA plasma.			
Fasting and non-fasting samples can be used.6			

The sample types listed were tested with a selection of sample collection tubes that were commercially available at the time of testing, i.e. not all available tubes of all manufacturers were tested. Sample collection systems from various manufacturers may contain differing materials which could affect the test results in some cases. When processing samples in primary tubes (sample collection systems), follow the instructions of the tube manufacturer.

Centrifuge samples containing precipitates before performing the assay.

Stability:15,16

7 days at 2-8 °C

12 months at -20 °C

12 months at -70 °C

It is reported that EDTA stabilizes lipoproteins.13

Materials provided

See "Reagents - working solutions" section for reagents.

Materials required (but not provided) See "Order information" section

General laboratory equipment

Assay

For optimum performance of the assay follow the directions given in this document for the analyzer concerned. Refer to the appropriate operator's manual for analyzer-specific assay instructions.

The performance of applications not validated by Roche is not warranted and must be defined by the user.

Application for serum and plasma

cobas c 111 test definition

Measuring mode		Absorbance	
Abs. calculation mode		Endpoint	
Reaction direction		Increase	
Wavelength A/B		583/659 nm	
Calc. first/last		16/37	
Unit		mmol/L	
Reaction mode		R1-S-SR	
Pipetting parameters			
			Diluent (H ₂ O)
R1		150 μL	
Sample		2 µL	7 μL
SR		50 µL	
Total volume		209 µL	
Calibration			
Calibrators	C.f	f.a.s. Lipids	
		ionized water is use instrument as the i	ed automatically by zero calibrator.
Calibration mode	Lin	ear regression	
Calibration frequency	2-p	point calibration	
	•	after reagent lot ch	ange
	•	as required following procedures	ng quality control
Calibration interval may be extend calibration by the laboratory.	led	based on acceptab	le verification of

Traceability: This method has been standardized against the beta quantification method as defined in the recommendations in the LDL Cholesterol Method Certification Protocol for Manufacturers.¹⁷

Quality control

For quality control, use control materials as listed in the "Order information" section.



In addition, other suitable control material can be used.

The control intervals and limits should be adapted to each laboratory's individual requirements. Values obtained should fall within the defined limits. Each laboratory should establish corrective measures to be taken if values fall outside the defined limits.

Follow the applicable government regulations and local guidelines for quality control.

Calculation

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LDL-Cholesterol Gen

The **cobas c** 111 analyzer automatically calculates the analyte concentration of each sample.

Conversion factor: mmol/L × 38.66 = mg/dL

Limitations – interference

Criterion: Recovery within \pm 0.40 mmol/L of initial values of samples \leq 4.0 mmol/L and within \pm 10 % for samples > 4.0 mmol/L.

Icterus:¹⁸ No significant interference up to an I index of 60 for conjugated and unconjugated bilirubin (approximate conjugated and unconjugated bilirubin concentration: 1026 μ mol/L or 60 mg/dL).

Hemolysis:¹⁸ No significant interference up to an H index of 1000 (approximate hemoglobin concentration: 621μ mol/L or 1000 mg/dL).

Lipemia (Intralipid):¹⁸ No significant interference up to an L index of 1000. There is poor correlation between the L index (corresponds to turbidity) and triglycerides concentration.

No significant interference from HDL-C (\leq 3.03 mmol/L or \leq 117 mg/dL), VLDL-C (\leq 3.63 mmol/L or \leq 140 mg/dL), or chylomicrons (\leq 22.6 mmol/L or \leq 2000 mg/dL triglycerides).

Drugs: No interference was found at the rapeutic concentrations using common drug panels. $^{19,20}\,$

Nicotinic acid (Niacin), statins (Simvastatin) and fibrates (Clofibrate) tested at therapeutic concentration ranges did not interfere.

Acetaminophen intoxications are frequently treated with N-acetylcysteine. N-acetylcysteine at the therapeutic concentration when used as an antidote and the acetaminophen metabolite N-acetyl-p-benzoquinone imine (NAPQI) independently may cause falsely low LDL-C results. Venipuncture should be performed prior to the administration of metamizole. Venipuncture immediately after or during the administration of metamizole may lead to falsely low results.

Ascorbic acid up to 28.4 mmol/L (500 mg/dL) does not interfere.

Abnormal liver function affects lipid metabolism; consequently HDL and LDL results are of limited diagnostic value. In some patients with abnormal liver function, the LDL-cholesterol result is significantly negatively biased versus beta quantification results.

EDTA plasma may cause decreased values compared to serum.²¹

In very rare cases, gammopathy, in particular type IgM (Waldenström's macroglobulinemia), may cause unreliable results.²²

For diagnostic purposes, the results should always be assessed in conjunction with the patient's medical history, clinical examination and other findings.

ACTION REQUIRED

Special Wash Programming: The use of special wash steps is mandatory when certain test combinations are run together on the **cobas c** 111 analyzer. For information about test combinations requiring special wash steps, please refer to the latest version of the carry over evasion list found with the CLEAN Method Sheet and the operator's manual for further instructions.

Where required, special wash/carry-over evasion programming must be implemented prior to reporting results with this test.

Limits and ranges

Measuring range

0.10-14.2 mmol/L (3.87-549 mg/dL)

Determine samples having higher concentrations via the rerun function. Dilution of samples via the rerun function is a 1:2 dilution. Results from samples diluted using the rerun function are automatically multiplied by a factor of 2.

Lower limits of measurement

Limit of Blank, Limit of Detection and Limit of Quantitation

Limit of Blank	= 0.10 mmol/L (3.87 mg/dL)
Limit of Detection	= 0.10 mmol/L (3.87 mg/dL)
Limit of Quantitation	= 0.10 mmol/L (3.87 mg/dL)

The Limit of Blank, Limit of Detection and Limit of Quantitation were determined in accordance with the CLSI (Clinical and Laboratory Standards Institute) EP17-A2 requirements.

C(n)ha

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The Limit of Blank is the 95^{th} percentile value from $n \ge 60$ measurements of analyte-free samples over several independent series. The Limit of Blank corresponds to the concentration below which analyte-free samples are found with a probability of 95 %.

The Limit of Detection is determined based on the Limit of Blank and the standard deviation of low concentration samples.

The Limit of Detection corresponds to the lowest analyte concentration which can be detected (value above the Limit of Blank with a probability of 95 %).

The Limit of Quantitation for LDL-C is 0.10 mmol/L determined in accordance with the guidelines in CLSI document EP17-A2, based on a minimum of 48 determinations, and a total error goal of 10 % calculated using RMS error model.

Expected values²³

Levels in terms of risk for coronary heart disease.

Adult levels:

Optimal	< 2.59 mmol/L (< 100 mg/dL)
Near optimal/above optimal	2.59-3.34 mmol/L (100-129 mg/dL)
Borderline high	3.37-4.12 mmol/L (130-159 mg/dL)
High	4.14-4.89 mmol/L (160-189 mg/dL)
Very high	≥ 4.92 mmol/L (≥ 190 mg/dL)

Risk classification of patients and treatment therapies are described in international guidelines. $^{\rm 24}$

Each laboratory should investigate the transferability of the expected values to its own patient population and if necessary determine its own reference ranges.

Specific performance data

Representative performance data on the **cobas c** 111 analyzer are given below. Results obtained in individual laboratories may differ.

Precision

Repeatability and intermediate precision were determined using human samples and controls in accordance with the CLSI (Clinical and Laboratory Standards Institute) EP5 requirements (4 aliquots per run, 1 run per day, 21 days). The following results were obtained:

Repeatability	Mean mmol/L (mg/dL)	SD mmol/L (mg/dL)	CV %
Precinorm L	2.63 (102)	0.04 (2)	1.6
Precipath HDL/LDL-C	4.98 (193)	0.04 (2)	0.9
Human serum 1	0.291 (11.3)	0.009 (0.4)	3.0
Human serum 2	2.86 (111)	0.03 (1)	0.9
Human serum 3	3.50 (135)	0.04 (2)	1.1
Human serum 4	8.10 (313)	0.12 (5)	1.4
Human serum 5	13.6 (526)	0.2 (8)	1.4
Intermediate precision	Mean mmol/L (mg/dL)	SD mmol/L (mg/dL)	CV %
Precinorm L	2.63 (102)	0.06 (2)	2.4
Precipath HDL/LDL-C	4.98 (193)	0.08 (3)	1.6
Human serum 1	0.291 (11.3)	0.009 (0.4)	3.2

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Intermediate precision	Mean mmol/L (mg/dL)	SD mmol/L (mg/dL)	CV %
Human serum 2	2.86 (111)	0.03 (1)	1.2
Human serum 3	3.50 (135)	0.04 (2)	1.2
Human serum 4	8.10 (313)	0.16 (6)	1.9
Human serum 5	13.6 (526)	0.2 (8)	1.6

Method comparison

LDL-cholesterol values for human serum samples obtained on a **cobas c** 111 analyzer (y) were compared to those determined using the corresponding reagent on a COBAS INTEGRA 400 plus analyzer (x). Sample size (n) = 167

Linear regression

Passi	na/Ra	ahlak ²	5

T assilly/Dablok	Linear regression
y = 0.993x - 0.022 mmol/L	y = 0.979x + 0.031 mmol/L
т = 0.932	r = 0.999

The sample concentrations were between 0.184 and 14.0 mmol/L (7.11 and 541 mg/dL).

References

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A point (period/stop) is always used in this Method Sheet as the decimal separator to mark the border between the integral and the fractional parts of a decimal numeral. Separators for thousands are not used.

Symbols

Roche Diagnostics uses the following symbols and signs in addition to those listed in the ISO 15223-1 standard (for USA: see https://usdiagnostics.roche.com for definition of symbols used):

CONTENT	Contents of kit
REAGENT	Reagent
\rightarrow	Volume after reconstitution or mixing
GTIN	Global Trade Item Number

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