## ratiogen

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# ApoE ToolSet<sup>TM</sup> for LightCycler<sup>TM</sup> (ApoE E2/E3/E4, codons 112 Cys/Arg & 158 Cys/Arg)

**Application on LightCycler 480 II** 

Lyophilized ToolSet for PCR using the LightCycler™ Instrument. Licensed by Roche Diagnostics GmbH

Order#: ApoE - 16

1 ToolSet for 16 reactions

Store at 4°C, protected from light. Exposure to light may especially damage the Oligotool ™ tube (vial with red cap).

For use with LightCycler FastStart DNA Master HybProbe, 10 x conc. (Roche Cat.No.: 03003248001)

#### 1. ToolSet contents

Vial	Label	Content	Quantity
			ApoE - 16
1, Red cap	OligoTool	- lyophilized oligos for PCR - contains mutation detection	For 16 tests
		and anchor probe, primers	Dissolved:
			50 μL
2, Green cap	Control	- lyophilized ApoE E2/E4 HET DNA	Dissolved:
			20 μL
3, Blue cap	Solvent	- to dissolve OligoTool / Control	1000 μL of Solvent
4, Yellow Cap	DMSO	- PCR additive for GC rich regions	100 μL of DMSO
	<b>Caution Irritant!</b>		·
	May be absorbed		Note : Fluid above 18 °C
	through skin!		Solid below 18 °C

Additional equipment and reagents required but not supplied:

LightCycler FastStart DNA Master HybProbe, 10 x conc. (Roche Cat.No.: 03003248001),including 25mM MgCl<sub>2</sub>; LightCycler instrument, LightCycler capillaries, DNA extraction materials

#### 2. Introduction

#### 2.1. Product overview

**ToolSet description** This ToolSet is specifically designed for multiplex genotyping of the codon 112 Cys / Arg

and codon 158 Cys / Arg polymorphisms in the Apolipoprotein E (ApoE) gene, encoding the **E2/E3/E4 isoforms of Apolipoprotein E**, by LightCycler PCR with Melting Curve Analysis. Primers and fluorescent probes have been optimized for specific amplification of a 462 bp segment containing the relevant site and for optimal genotype discrimination.

**Control material** E2 / E4 Heterozygous Control DNA, lyophilized

(= Cys112Arg / Cys158Arg compound heterozygote).

Storage of ToolSet and Solutions

Store at +4°C when lyophilized, protected from light.

The unopened lyophilized ToolSet is stable at +4°C for 12 months from date of manufacture if protected from light. When dissolved store at +4°C for a maximum of 4 weeks, or at -20°C for longer periods (months),

protected from light. Avoid freezing and thawing > 3 times.

#### 3. Preparation for LightCycler PCR

#### **Toolset** preparation

Dissolve the content of the OligoTool tube (Red Cap) with 50 µl of Solvent. Dissolve the content of the Control tube (Green Cap) with 20 µl of Solvent.

- 1. Before opening tubes, centrifuge them guickly.
- 2. Add Solvent into OligoTool tube and Control tube as above.
- 3. Recap tubes, vortex gently.
- 4. Before opening tubes, centrifuge them quickly.
- 5. Proceed to Reaction Mix preparation.

#### **Primers?** Probes?

You don't have to add primers. You don't have to add probes.

#### **Reaction Mix Preparation**

For 1 (One) reaction, prepare the Reaction Mix as shown in the following table:

Reagent	μL	
OligoTool ApoE -16, dissolved	2.8	
Solvent ApoE -16	8.0	
MgCl <sub>2</sub> 25 mM	1.2 (final 2.5 mM)	
DMSO (yellow cap)	2.0	
FastStart DNA Master HybProbe, 10x	2.0	
Total Reaction Mix	16.0	
+ Your DNA or Control ApoE E2/E4 HET –16	4.0	
Grand Total	20.0	

Use FastStart DNA Master HybProbe 10x and MgCl, 25 mM from Roche LightCycler FastStart DNA Master HybProbe, 10 x conc. (Roche Cat.No.: 03003248001, including 25mM MgCl<sub>2</sub>). For multiple reactions, multiply the indicated volumes appropriately.

#### **Positive Control**

Always run a positive control with the samples. Use the dissolved heterozygous Control ApoE E2/E4 HET - 16 DNA (Green Cap).

**Negative control** Always run a negative control with the samples. To prepare a negative control, replace the template DNA with Solvent (Blue Cap).

#### **Extraction of** genomic DNA

You can use different Kits for DNA isolation, either with a manual method or with an automated system. The elution buffers should be salt-free. Example: Roche High Pure PCR Template Preparation Kit (Cat.No. 1 796 828)

#### **Application**

The **ApoE** ToolSet<sup>™</sup> for LightCycler<sup>™</sup> allows detection of the **codon 112** Cys/Arg and codon 158 Cys/Arg variants in the ApoE gene, encoding the E2/E3/E4 isoforms of Apolipoprotein E.

Note: This ToolSet was developed for use in life science research only.

### 4. LightCycler 480 Settings and Experimental Protocol

For use with LC 480 Program Version 1.5 series.

Detection: Dynamic Red 640 (498-640 nm) and Dynamic Red 705 (498-660 nm) with color compensation (obligatory!)

#### **Fast Start Enzyme Activation and DNA Denaturation**

Cycle Program Data	Value		
Cycles	1		
Analysis Mode	None		
Temperature Targets	Segment 1		
Target Temperature (°C)	95		
Incubation time (s) <b>Hold</b>	600		
Temperature Transition Rate (°C/s) Ramp Rate	4.4		
Acquisitions (per °C)			
Secondary Target Temperature (°C)	0		
Step Size (°C)	0		
Step Delay (Cycles)	0		
Acquisition Mode	None		

#### **Amplification**

Cycle Program Data	Value			
Cycles	45	45		
Analysis Mode	Quantification	Quantification		
Temperature Targets	Segment 1	Segment 2	Segment 3	
Target Temperature (°C)	95	60	72	
Incubation time (s) <b>Hold</b>	5	10	15	
Temperature Transition Rate (°/s) Ramp Rate	4.4	2.2	4.4	
Acquisitions (per °C)				
Secondary Target Temperature (°C)	0	0	0	
Step Size (°C)	0	0	0	
Step Delay (Cycles)	0	0	0	
Acquisition Mode	None	Single	None	

#### **Melting Curve Analysis**

Cycle Program Data	Value		
Cycles	1		
Analysis Mode	Melting Curves		
Temperature Targets	Segment 1	Segment 2	Segment 3
Target Temperature (°C)	95	40	85
Incubation time (s) <b>Hold</b>	30	120	
Temperature Transition Rate (°/s) Ramp Rate	4.4	1.5	0.1
Acquisitions (per °C)			3
Secondary Target Temperature (°C)	0	0	0
Step Size (°C)	0	0	0
Step Delay (Cycles)	0	0	0
Acquisition Mode	None	None	Continuous

### Cooling

Cycle Program Data	Value		
Cycles	1		
Analysis Mode	None		
Temperature Targets	Segment 1		
Target Temperature (°C)	40		
Incubation time (s) <b>Hold</b>	30		
Temperature Transition Rate (°/s) Ramp Rate	1.5		
Secondary Target Temperature (°C)	0		
Step Size (°C)	0		
Step Delay (Cycles)	0		
Acquisition Mode	None		

#### 5. Typical results

#### Introduction

Use the Melting Curve program to genotype the human genomic DNA research samples. The melting peaks allow discrimination between the possible genotypes of the codon 112 and 158 polymorphisms in the ApoE gene. Figure 1 shows a typical result obtained with the ApoE ToolSet™ for LightCycler™:

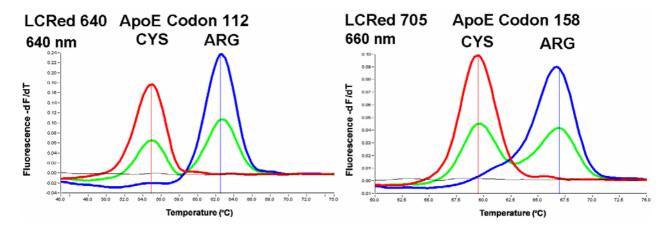


Figure 1: Melting curve analysis of genotypes of codons 112 and 158 in the ApoE gene.

BLUE: Homozygotes Arg112Arg (left) and Arg158Arg (right)

**GREEN**: Compound Heterozygote **Cys112Arg** / **Cys158Arg** = ApoE E2 / E4 HET Control contained in the ToolSet.

**RED**: Homozygotes **Cys112Cys** (left) and **Cys158Cys** (right)

**BLACK**: No DNA Control.

Conditions: LC 480 program version 1.5, With Color compensation,

#### For conversion of codon 112 / 158 melting curves into ApoE genotypes, use the following Table:

Table 1 : Conversion of Codon 112 / Codon 158 melting curve results into ApoE genotype YES : peak observed at indicated Tm no : No peak observed at indicated Tm

	<b>Codon 112</b> (LCRed 640, 640nm)		<b>Codon 158</b> (LCRed 705, 660 nm)		
Genotype	Tm 54 - 56 C	Tm 62 - 64 C	Tm 59 - 61 C	Tm 66 – 68 C	Genotype
E2/E2 (Cys/Cys)	YES	no	YES	no	E2/E2 (Cys/Cys)
E3/E3 (Cys/Arg)	YES	no	no	YES	E3/E3 (Cys/Arg)
E4/E4 (Arg/Arg)	no	YES	no	YES	E4/E4 (Arg/Arg)
E2/E3	YES	no	YES	YES	E2/E3
E2/E4	YES	YES	YES	YES	E2/E4
E3/E4	YES	YES	no	YES	E3/E4

**Note**: The values for the respective melting temperatures may vary for +/- 2.5 °C between different experiments. The Delta T between the melting peaks for different genotypes may vary +/- 1.0 °C. The ApoE ToolSet™ has been developed for and validated with the LightCycler™ and its original accessory materials and reagents. Performance of the ToolSet with other instruments, accessories and reagents has not been validated by ratiogen.

#### 7. Notices to Purchaser: Licenses and Trademarks, Prohibition of Resale

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