# **PCR DIG Probe Synthesis Kit**

For generation of highly-sensitive probes labeled with DIG-dUTP (alkali-labile) in the polymerase chain reaction (PCR)

# Cat. No. 1 636 090

1. Kit contents 2. Product overview

Kit for 25 polymerase chain reactions (50 µl)

## Version 4, August 2000 Store at -15 to -25° C

## 2. Product overview

<ol> <li>Proc</li> <li>PCR r</li> <li>DNA</li> <li>Hybrid</li> <li>Chem</li> </ol>	eaction electrophores dization with t niluminescent ping and repro	<b>d requ</b> i sis, trans the DIG detectio			Kit description	required for t DNA fragmer reaction (PCI reaction is id ciently labele synthesis of f DIG-dUTP in centration in single-copy of	Probe Synthesis Kit contains all reagents the direct digoxigenin (DIG)-labeling of nts generated by the polymerase chain R) process (1, 2). The polymerase chain eally suited to prepare specific and effi- d hybridization probes. The kit enables the nighly sensitive probes by incorporation of to the PCR product. The nucleotide con- the PCR DIG Mix ensures the detection of genes in genomic blots after hybridization
1. Kit c Caution	Vi Th ind dL re Th Vi. cc	nis mix f cluding UTP pro agent, t nese nu al 2 (PC oncentra corpora	CR DIG probe synthesis mix) rom this kit contains a mixture of the DIG-dUTP. The concentration vided here differs from a separate he PCR DIG labeling mix (Cat. No cleotide mixes should not be into R DIG probe synthesis mix) conta titon of DIG-dUTP to achieve max tion, and maximal probe sensitivity used with this PCR DIG Probe S	of DIG- yet similar 1 585 550). erchanged. ins a higher imal DIG y. Only vial 2	Synthesis principle	The kit incluc Fidelity (e.g. yield and hig solution is su tration of lab cDNA for a h tive primers s PCR labeling PCR product from low amo	s can directly be amplified and labeled ounts of plasmid or genomic DNA and
Bottle/ Cap	Label		Content including function	Cat. No. (if avai- lable se-	Basic steps	further purifie	y be used as hybridization probes without cation.
2	Enzyme mix, Expand <sup>1</sup> High Fidelity PCR DIG probe	<ul> <li>3.5</li> <li>Stop</li> <li>pH</li> <li>1 n</li> <li>ED</li> <li>0.5</li> <li>509</li> <li>Enz</li> <li>proc</li> <li>125</li> </ul>	μl enzyme mix (105 units) units/μl rage buffer; 20 mM Tris-HCl, 7.5 (25°C), 100 mM KCl, M dithiothreitol (DDT), 0.1 mM TA, 0.5% Tween <sup>2)</sup> 20 (v/v)*, % Nonidet <sup>3)</sup> P40 (v/v)*, % glycerol (v/v). cyme mix for the labeling of PCR duct 5 μl ture containing dATP, dCTP,	parately) 1 732 641		Stage PCR labeling Hybridiza- tion	Description DIG-labeled DNA probes are generated according to the PCR labeling technique. DIG-labeled probes are used for hybri- dization to membrane blotted nucleic acids according to standard methods. The use of the alkali-labile form of DIG- 11-dUTP enables easier and more effi- cient stripping of blots for rehybridization with a second DIG-labeled probe.
	synthesis mix, 10 × conc.	dG 0.7 pH • Nu PC	TP (2 mM each); 1.3 mM dTTP; mM DIG-11-dUTP, alkali-labile; 7.0. cleotide mix for the labeling of R product			Immunolog- ical detec- tion	The hybridized probes are immunode- tected with anti-digoxigenin-AP, Fab fragments and are then visualized with a chemiluminescent substrate (CDP- <i>Star</i> or CSPD*) or by colorimetric detection with e.g. NBT/ BCIP.
3	PCR buffer with MgCl <sub>2</sub> , 10 × conc dNTP stock solution, 10 × conc.	cor • Bui • 125 • Miz dT	band High Fidelity buffer 10 × nc. with 15 mM MgCl <sub>2</sub> ffer for the PCR labeling reaction 5 µl cture contains dATP, dCTP, dGTP, IP (2 mM each), pH 7.0.	1 581 295	Application	sensitive hyb low (single) o alkali-labile f which is easi	ecially designed for generation of highly ridization probes suitable for detection of copy target sequences. Besides this the orm of DIG-dUTP generated a probe ly used for rehybridizations of (genomic) ely stripping of.
<ul> <li>Solution for the possible dilut the PCR labeling reaction</li> <li>Control template</li> <li>50 μl</li> <li>1 ng plasmid DNA [20 pg/ml Tris/EDTA buffer; pH 8.0.</li> <li>The 5 kb plasmid contains th cDNA for human tissue type plasminogen activator (tPA).</li> </ul>		μl g plasmid DNA [20 pg/ml] in /EDTA buffer; pH 8.0. e 5 kb plasmid contains the NA for human tissue type		Labeling efficiency	Optimal react DNA and print temperatures also concent	tion conditions are dependent on template mer. In particular incubation times and s, concentration of Mg <sup>2+</sup> and enzyme but ration of template DNA and primer should for optimal results for each new primer/	
6	Control PCR primer mix	1 a	µl pmol] of control PCR primer nd 2 (2 mM each). ner for the control reaction				



Sample material	Partially purified DNA containing the sequence to be labeled. Use either:
	<ul> <li>Plasmid DNA, 10–100 pg (optimal amount, 10 pg)</li> </ul>
	<ul> <li>Genomic DNA, 1–50 ng (optimal amount, 10 ng)</li> </ul>
	<i><u>Note</u>:</i> Purity of template is not as critical for PCR labeling as for other types of labeling.
Number of tests	1 kit is sufficient for 25 polymerase chain reactions.
Quality Control	The PCR DIG Probe Synthesis Kit is function-tested in PCR. Amplification products are assayed in genomic Southern blots. Under PCR conditions described the control reaction generates an amplification product of 442 bp. Due to multiple incorporation of DIG-dUTP during the PCR process the molecular weight of the PCR product is increased significantly compared to the unlabeled PCR product. After hybridization of the PCR product to 5 µg human genomic DNA and chemiluminescent detection a specific fragment pattern is detected.
Kit storage/ stability	The unopened kit is stable at -15 to -25°C until the expiration date printed on the label. Shipping conditions on dry ice.
Sensitivity	The nucleotide concentration in the PCR DIG probe synthesis mix ensures the identification of single-copy genes in genomic blots after hybridization to DIG-labeled PCR products. Human single-copy genes typically are detectable in 5 $\mu$ g of genomic DNA.

## Advantages

Benefits	Feature
Small amounts of template DNA	Because incorporation of DIG-dUTP into the PCR product produce highly sensitive probes.
Purity of template DNA is not as critical for PCR labeling	As for other types of labeling the purity of template is not critical, espe- cially random primed labeling. For example, suitable templates include partially purified plasmid prepared by any of a variety of "quick preparation" methods, even simple boiling of cells.
The labeling reaction requires less optimization than other methods	<ul> <li>Can efficiently use GC-rich regions as template</li> <li>For most templates, no optimiza- tion of MgCl<sub>2</sub> concentration is required; most labeling reactions will work with the standard con- centrations of 1.5 mM MgCl<sub>2</sub>.</li> </ul>
Large quantities of DIG-labeled probes.	Yield of labeled probe is very high.
dNTP stock solution	Because of their unique sequences, some large fragments (larger than 1 kb) amplify poorly when high concen- trations of DIG-dUTP are used. If a template produces yields which are insufficient, we supply dNTP stock solution to dilute the DIG-dUTP. Please see "Procedure" and "Trouble Shooting" for additional instructions.
Evaluation of labeling efficiency	Very quickly by gel electrophoresis
No purification of labeled PCR prod- uct necessary	PCR-generated probes are very pure and can be used directly in the hybridization reaction.
The PCR DIG Probe Synthesis Mix contains the alkali-labile form of DIG- dUTP.	Subsequent rehybridizations of (genomic) blots by merely stripping off the DIG molecule under alkaline conditions.

## 3. Procedures and required materials

## Before you begin

	20.0.0 jou 20				
Critical Hints about PCR Labeling		Please refer to the following table:			
	PCR amplifi- cation parameters	Optimize PCR amplification parameters (cycling conditions, template concentration, primer sequence, and primer con- centration) for each template and primer set in the <b>absence</b> of DIG-dUTP before attempting incorporation of DIG.			
	DIG-dUTP concentration	According to the length of the probe being labeled the con- centration of DIG-dUTP has to be adapted:			

ncentration	centration of DIG-dUTP has to be adapted:
	<ul> <li>&lt; 1 kb, use a 1:3 ratio of DIG-dUTP : dTTP (standard</li> </ul>
	labeling mix vial 2).
	Note: DNA containing a high GC content may require a 1:6
	ratio of DIG-dUTP : dTTP.
	<ul> <li>&gt; 1 kb, use a 1:6 ratio of DIG-dUTP : dTTP. Mix equal</li> </ul>
	parts of PCR DIG probe synthesis mix (vial 2) and dNTP
	stock solution (vial 4)
	<ul> <li>&gt; 3 kb, use a 1:6 ratio of DIG-dUTP : dTTP and substitute</li> </ul>
	the Expand Long Template enzyme mix for the Expand
	High Fidelity enzyme mix included in the PCR DIG Probe
	Synthesis Kit.
	Note: You may eventually need to reduce the
	DIG-dUTP : dTTP ratio as low as 1:10.

Unlabeled positive control	Identical to experimental sample except the reaction mix contains no DIG-dUTP. <u><i>Note:</i></u> Always include this control reaction in every experiment. It is required for evaluating probe labeling efficiency.
Labeled positive control	Substitutes the tissue plasminogen activator (tPA) tem- plate and primers (included in the PCR Probe Synthesis Kit) for your experimental template and primers. Pro- duces a labeled probe that recognizes human tPA sequences.
	The tPA probe generated in the control PCR recognizes a restriction length polymorphism (RFLP) for an <i>Eco</i> RI site in human DNA (5). This results in the detection of variable fragment patterns depending on the human DNA used. One of the following fragment patterns of human single-copy tPA gene should be detected: • 2.9 kb + 1.7 kb • 2.5 kb + 1.7 kb • 2.9 kb + 2.5 kb + 1.7 kb
	<i><u>Mote</u>:</i> As you gain experience with the labeling kit, you may choose not to run this control reaction.
3.1 PCR reaction	 ו

	all equipment needed for PCR reactions
ment and reagents . required	sterile double dist. water
	mineral oil (optional, depending on your PCR equiment)
	Primer:
	<ul> <li>1–10 µM solution of upstream primer</li> </ul>
	<ul> <li>1–10 µM solution of downstream primer</li> </ul>
•	Template DNA

Procedure

Cycling conditions depend on the combination of template, primers, and thermal cycler. The conditions given below may not be optimal for your template/primer combination, but are a good starting point for initial experiments.

Step		Action						
1	Add the following components to a sterile microcentrifuge tube o ice:							
	10 <del>0</del> .							
		Reagent		DIG labeled probe	Unlabeled DNA con- trol	kit (	eled con- ol	Final conc.
	Sterile dou dist. water		le	variable	variable	29.2	25 µl	-
		PCR buffer with MgCl <sub>2</sub> , 10 x conc., (vial 3)		5 µl	5 mµl	5	μI	1x
		PCR DIG Labeling Mi (vial 2)	х	5 µl	-	5	μΙ	200 μM dNTP
		dNTP stock solution (vial 4)		-	5 mµl		-	200 µM dNTP
		Upstream an downstream primer		variable	variable		µl al 6)	0.1 -1 µl each
		Enzyme mix (vial 1)		0.75 µl	0.75 mµl	0.7	5 µl	2.6 units
		Template DI	٨V	variable	variable		µl al 5)	
		Total		50 µl	50 µl	50	)µl	
2		x the reagent			je brieny to c	Uncer	110 30	ampie at ti
2	bot Ove dur	ttom of the tu erlay with 10 ring amplifica	ibe. 0 µl atior	mineral oil	to reduce ev	apora	ition o	f the mix
	bot Ove dur <u>No</u>	ttom of the tu erlay with 10 ring amplifica	ibe. 0 µl atior	mineral oil		apora	ition o	f the mix
	bot dur <u>No</u> neo	ttom of the tu erlay with 100 ring amplifica <u>ote:</u> If your th cessary.	ibe. 0 µl atior erm	mineral oil n. al cycler ha	to reduce ev	apora er, the	tion o	f the mix
3	bot dur <u>No</u> neo	ttom of the tu erlay with 100 ring amplifica <u>ote:</u> If your th cessary.	ibe. 0 µl atior erm n th	mineral oil n. al cycler ha	to reduce ev as a top heate	apora er, the irt PC	tion o oil ov	f the mix
3	bot dur <u>No</u> neo	ttom of the tu erlay with 100 ring amplifica <u>offer</u> If your th cessary. ce samples i Cycle	ibe. 0 µl atior erm n th	mineral oil n. aal cycler ha	to reduce ev as a top heate cycler and sta Tempera	apora er, the irt PC	tion o oil ov R. <b>Tim</b>	f the mix verlay is no
3	bot dur <u>Na</u> nec	ttom of the tu erlay with 100 ring amplifica <u>offer</u> If your th cessary. ce samples i Cycle	ibe. 0 µl ation erm n th Ini ati De Ar	mineral oil n. al cycler ha te thermal c <b>Reaction</b> tial denatur	to reduce ev as a top heate cycler and sta Tempera	apora er, the irt PC	tion o oil ov R. <b>Tim</b>	f the mix rerlay is no e period
3	bot Ove dur <u>No</u> nec Pla	ttom of the tu erlay with 100 ring amplifica <u>te</u> : If your th cessary. ce samples i Cycle number	ibe. 0 μl ation erm n th Ini ati Δe Ar Elc Ar Elc	mineral oil n. lal cycler ha le thermal c <b>Reaction</b> tial denatur on enaturation mealing ongation enaturation mealing ongation	to reduce ev as a top heate cycler and sta <b>Tempera</b> 95°C 95°C 60°C 72°C 95°C 60°C 72°C	apora rt PC ture	40 s tiona	f the mix verlay is no e period 2 min 30 s 30 s 30 s 30 s 30 s 30 s 30 s 30 s
3	bot Ove dur <u>No</u> nec Pla	ttom of the tu erlay with 100 ring amplifica <u>vte:</u> if your th cessary. ce samples i Cycle number Cycles 1 - 10 Cycles	Ibe. D µl ation erm n th Ini ati De Ar Elc Ar Elc Mar rec ca	mineral oil n. al cycler ha te thermal o <b>Reaction</b> tial denatur on enaturation ongation enaturation nealing ongation	to reduce ev as a top heater cycler and state Tempera - 95°C 60°C 72°C 95°C 60°C 72°C	apora rt PC ture	40 s tiona tiona each time tis. Fo nue to	f the mix rerlay is no e period 2 min 30 s 30 s 30 s 30 s 30 s 30 s 30 s 30 s
3	bot Ove dur <u>No</u> nec Pla	ttom of the tu erlay with 100 ring amplifica <u>vte:</u> if your th cessary. ce samples i Cycle number Cycles 1 - 10 Cycles	Ibe. D µl ation erm n th Ini ati De Ar Elc De Ar Elc Mar rec ca 40 Fir tion	mineral oil n. aal cycler ha ae thermal o <b>Reaction</b> tial denatur on enaturation nnealing ongation enaturation nealing ongation <u>ofte:</u> The inc quired for k tin of short s elongation nat leonga- n step	to reduce ev as a top heater cycler and state Tempera - 95°C 60°C 72°C 95°C 60°C 72°C 95°C 60°C 72°C 95°C 60°C 72°C 72°C	apora rt PC ture	40 s tiona R. Tim 40 s tiona each time ts. Fo nue to ycles.	f the mix rerlay is no e period 2 min 30 s 30 s 30 s 30 s 30 s 30 s 30 s 30 s
3	bot Ove dur <u>No</u> nec Pla	ttom of the tu erlay with 100 ring amplifica <u>vte:</u> if your th cessary. ce samples i Cycle number Cycles 1 - 10 Cycles	Ibe. 0 µl ation erm Initiati De Ar Elc Mar Ca 40 Fir	mineral oil n. aal cycler ha ae thermal o <b>Reaction</b> tial denatur on enaturation nnealing ongation enaturation nealing ongation <u>ofte:</u> The inc quired for k tin of short s elongation nat leonga- n step	to reduce ev as a top heater cycler and state Tempera - 95°C 60°C 72°C 95°C 60°C 72°C 60°C 72°C 60°C 72°C 95°C 60°C 72°C	apora rt PC ture	40 s tiona R. Tim 40 s tiona each time ts. Fo nue to ycles.	f the mix rerlay is not e period 2 min 30 s 30 s 30 s 30 s 30 s 30 s 30 s 30 s
3	boti Ovu dur <u>Na</u> nec Pla	ttom of the tu erlay with 100 ring amplifica <u>vte:</u> if your th cessary. ce samples i Cycle number Cycles 1 - 10 Cycles	Ibe. 0 µl ation erm n th Ini ati De Ar Elc De Ar Elc Fir tion Ho	mineral oil n. nal cycler ha ne thermal c Reaction tial denatur on maturation mealing ongation maturation mealing ongation <u>ote:</u> The inc quired for k tin of short 's elongation natep old	to reduce ev as a top heater cycler and state <b>Tempera</b> <b>7-</b> 95°C 60°C 72°C 95°C 60°C 72°C 95°C 60°C 72°C 95°C 60°C 72°C 20°C 72°C 4°C	apora rt PC ture	40 s tiona R. Tim 40 s tiona each time ts. Fo nue to ycles.	f the mix rerlay is no e period 2 min 30 s 30 s 30 s 30 s 30 s 30 s 30 s 30 s
3	boti Ovu dur <u>Na</u> nec Pla	ttom of the tu erlay with 100 ring amplifica te: If your th cessary. ce samples i Cycle number Cycles 1 - 10 Cycles 11 - 30	Ibe. Ibe. Ibe. Ibe. Ibe. Ibe. Ibe. Ibe.	mineral oil h. hal cycler ha he thermal c <b>Reaction</b> tial denatur on maturation mealing ongation enaturation nealing ongation enaturation nealing ongation enaturation nealing ongation enaturation nealing ongation enaturation nealing ongation enaturation nealing ongation enaturation nealing ongation enaturation nealing ongation enaturation nealing ongation enaturation nealing ongation enaturation nealing ongation enaturation nealing ongation enaturation nealing ongation enaturation nealing ongation enaturation nealing ongation enaturation nealing ongation enaturation nealing ongation enaturation nealing ongation enaturation nealing ongation enaturation nealing ongation enaturation nealing ongation enaturation nealing ongation enaturation nealing ongation enaturation nealing ongation enaturation nealing ongation enaturation notep old	to reduce ev as a top heater cycler and state <b>Tempera</b> <b>7</b> 95°C 60°C 72°C 95°C 60°C 72°C 95°C 60°C 72°C 00°C 72°C 40°C 4°C 4°C 4 be stored:	apora er, the rt PC ture gmer conti 30 cy	40 s tiona each time ts. Fo nue to ycles.	f the mix rerlay is not e period 2 min 30 s 30 s 30 s 30 s 30 s 30 s 30 s 30 s

The best way to check the success of the reaction is to run a portion (5  $\mu$ I) of each reaction on an agarose mini gel and then stain the gel with ethidium bromide (EtBr).

The tPA control probe will have an apparent size of 500–550 bp.

**Note:** The actual size of the amplicon is 442 bp. The presence of DIG in DNA makes it run slower in the gel than unlabeled DNA. The agarose concentration of the gel determines the extent to which the migration rate deviates from its "true" value.

- Your unlabeled control probe should be the predicted size.
- Both the labeled experimental probe and the unlabeled control probe should be clearly visible on the gel.
- Your labeled experimental probe should migrate slower than your unlabeled control probe (due to the presence of DIG).
- The EtBr staining of the labeled DNA will be somewhat less than the unlabeled control DNA.
   <u>Note</u>: When you are using a high ratio of DIG-dUTP: dTTP (1:3), the reaction will make less labeled probe than unlabeled probe. The polymerase is slowed by the presence of the DIG hapten.

## 3.2 DNA electrophoresis, transfer, and fixation

General

Analysis of

labeled probe

Standard protocols for gel electrophoresis, denaturation and neutralization of the gel are described in Sambrook et al. (6). Gels lacking ethidium bromide are preferred, because ethidium can cause uneven background problems. All common types of DNA transfer methods are suitable for subsequent DIG hybridization (7,8).

In our experience, best results are obtained when gels are blotted by capillary transfer with 20x SSC on nylon membranes, positively charged\* (Cat. No. 1 209 272).

<u>Note</u>: Alkali transfer (e.g., in 0.4 M NaOH) is not suitable for the transfer of DIG-labeled molecular weight markers\*.

#### Overview

Stage	Description
1	Gel electrophoresis.
2	Denaturation and neutralization of DNA in the gel.
3	Fixation of DNA on the membrane.

#### Electrophoresis We recommend the following:

- Prepare a suitable electrophoresis gel as thin as possible.
- · Load small amounts of the target DNA samples:

genomic DNA1–5 μg per laneplasmid DNA< 1 ng per lane
plasmid DNA < 1 ng per lane
pidoinia Briti
DIG-labeled Molecular Weight Marker* <u>Note:</u> The needed will depend on the expected size of your hybridization product. Make sure to load enough marker to produce promi- nent bands that are about the same size as your hybrid.

· Run the gel until the DNA bands are well separated.

- To assess the quality of the target DNA, stain the gel briefly in 0.25–0.50 µg/ml ethidium bromide, then destain with water.
- · Examine the gel under UV light.

Fixation procedure After the transfer, while the blot is still damp, fix the DNA to the blot by either of the following methods:

IF you want to	THEN
UV-crosslinking (nylon mem- brane)	<ul> <li>place the membrane on Whatman 3MM-paper soaked with 2x SSC.</li> <li>UV-crosslink the wet membrane without prior washing.</li> <li>after the UV-crosslinking, rinse the membrane briefly in double distilled water and allow to airdry.</li> </ul>
bake at 120°C (nylon mem- brane)	<ul> <li>wash the membrane briefly in 2x SSC.</li> <li>bake the nylon membrane at 120°C for 30 min or according to the manufacturer's instructions.</li> </ul>

Storage of membrane

Please refer to the following table.

IF	THEN
you want to go ahead.	use the membrane imme- diately for prehybridization.
you want to work later on	store the membrane dry at 2 to 8°C.

## 3.3 Hybridization with the DIG-labeled Probe

Hybridization	The procedures for using the DIG-labeled probes for the detection of human genomic DNA on a Southern blot can also be used for the detection on Dot blots.				
	We recommend to use				
	<ul> <li>DIG Easy H</li> </ul>	yb buffer* (Cat. No. 1 603 558) and			
	<ul> <li>Hybridization</li> </ul>	on bags* (Cat. No. 1 666 649)			
		s. For detailed information about the protocol please see the DIG Easy Hyb e insert.			
Probe concentration	The standard probe concentration is 2 µl probe per ml hybridization buffer.				
	<ul> <li>If the labeled PCR product band on the evaluation gel was very faint, use up to 4 µl probe per ml hybridization buffer.</li> </ul>				
	- If the labeled band was very strong, use only 1 $\mu I$ (or even as little as 0.5 $\mu I$ ) probe per mI hybridization buffer.				
Overview	Do not allow the membrane to dry at any time from the beginning of prehybridization through probe-hybrid visualization. If the membrane dries or sticks to a second membrane (e.g., during simultaneous processing of blots), the assay will have a high background.				
	Stage	Description			
	1	Prehybridization of blot.			
	2	Hybridization with DIG-labeled probe.			
	3	Stringency washes.			

Storage of membrane After finishing the last high stringency wash, you may air dry the membranes and store it in a sealed bag at 2 to 8°C for later detection.

#### 3.4 Chemiluminescent detection

Chemilumines- cent detection	For the chemiluminescent detection of the labeled DIG- probe we recommend to use		
	<ul> <li>CDP-Star*(Cat. No. 2 041 677) or CSPD* (Cat. No. 1 755 633)</li> </ul>		
	DIG Block and Wash Buffer Set* (Cat. No. 1 585 762) and		
	<ul> <li>Anti-digoxigenin-AP, Fab fragments * (Cat. No. 1 093 274)</li> </ul>		
Overview	This table lists the single steps of the chemilumines- cent detection.		

Stage	Description	
1	Washing and blocking of membrane.	
2	Antibody binding.	
3	Washing and equilibration of membrane.	
4	Chemiluminescent reaction.	
5	Film exposure.	

For detailed information about the detection protocol please see the package insert of CDP-*Star* or CSPD.

## 3.5 Stripping and reprobing of DNA blots

General	The alkali-labile form of DIG-11-dUTP enables easier and more efficient stripping of blots for rehybridization experiment.		
Additional	<ul> <li>0,2 N NaOH, 0,1% SDS (w/v)</li> </ul>		
reagents required	• 2x SSC		
Protocol	Please refer to the following table.		
	<u>Note</u> : When stripping and rehybridization of blots is planned, the membrane should not dry off at any time.		
	Step Action		
	1	Rinse membrane briefly in <b>double distilled</b> water.	

I	water.
	Wash for 2x 15 min in <b>0.2 N NaOH</b> , <b>SDS</b> , <b>0.1%</b> (w/v) at 37°C under constant agitation.
3	Equilibrate briefly in <b>2x SSC</b> .
4	Prehybridize and hybridize with the next probe according to the protocol.

#### 4. Appendix

Trouble shooting table

This table describes various trouble shooting parameters for DIG-labeling and detection

Problem	Possible cause	Recommendation	
Low yield of DIG-labeled PCR product	PCR is not opti- mized	Always optimize the PCR parameters (cycling conditions, template concen- tration, primer sequence, and primer concentration) for each template and primer set in the <b>absence</b> of DIG- dUTP before attempting incorporation of DIG.	
	Too much DIG- dUTP in reaction <sup>1</sup>	Reduce the concentration of DIG-dUTP in the reaction. This is especially impor- tant for long templates, for further information see chapter 3.	
Cloudy hybri- dization back- ground	Probe concen- tration too high in the hybridization solution.	Reduce probe concentration to 1 µl probe per ml DIG Easy Hyb buffer. <u>Note:</u> Evaluate the amount of labeled probe in the PCR product. If the amount of labeled PCR pro-duct band is very strong on the gel, use as little as 0.5 µl probe per ml hybridization buffer.	
Hybridization smear	Template concen- tration too high during PCR	For best results, use only small amounts of template. Ideal amounts: 10 pg plasmid DNA or 10 ng genomic DNA. <u>Note:</u> Our experience indicates that higher concentrations of template lead to large amounts of primary extension products in the labeled probe.	
	Target concentra- tion too high on the blot	We recommend to use 1–5 µg per lane genomic DNA or < 1 ng plasmid DNA per lane	

#### References

- Saiki, R. et al. (1985) *Science* 230, 1350–1354.
   Lion, T. & Haas, O.A. (1990) *Anal. Biochem.* 188, 335–337.
   Rolfs, A. et al. (1992) PCR: Clinical Diagnostic and Research, Springer Verlag, Berlin.

#### **Related products**

#### Kits

Product	Pack Size	Cat. No
DNA Isolation Kit for Cells and Tissue	10 isolations for 400 mg tissue or 5 x 10 <sup>7</sup> cells	1 814 770
High Pure Plasmid Isolation Kit	50 purifications 250 purifications	1 754 777 1 754 785
High Pure PCR Product Purification Kit	1 kit (50 purifications) 1 kit (250 purifications)	1 732 668 1 732 676
High Pure PCR Template Preparation Kit	1 kit (100 reactions)	1 796 828
Plant DNA Isolation Kit for convenient isolation of plant genomic DNA	up to 150 purifications	1 667 319

#### Single reagents

Product	Pack Size	Cat. No.
Agarose MP	20 g 100 g 500 g	1 444 964 1 388 983 1 388 991
Blocking reagent	50 g	1 096 176
CDP-Star	1 ml 2 ml	1 685 627 1 759 051
CDP-Star, ready-to-use	2x 50 ml	2 041 677
CSPD	1 ml 2 ml 4 ml	1 655 884 1 759 035 1 759 043
CSPD, ready-to-use	2x 50 ml	1 755 633
NBT/BCIP ready to use tablets	20 tablets	1 697 471
NBT/BCIP Stock solution	8 ml	1 681 451
DIG Easy Hyb (ready-to-use hybridization solution without formamide)	500 ml	1 603 558
DIG Easy Hyb Granules	1 set (6 x 100 ml)	1 796 895
DIG Quantification Test Strips	50 strips	1 669 958
DIG Wash and Block Buffer Set	30 blots (10 x 10 cm <sup>2</sup> )	1 585 762
DNA Molecular Weight Marker, Digoxigenin-labeled: DNA MWMII DNA MWMIII DNA MWM V DNA MWM VI DNA MWM VII DNA MWM VIII	5 µg (500 µl) 5 µg (500 µl)	1 218 590 1 218 603 1 669 931 1 218 611 1 669 940 1 449 451
Hybridization bags	50 bags	1 666 649
Lumi-Image F1	240 Volt 120 Volt	2 015 170 2 012 847
Nylon Membrane, positively charged (20 x 30 cm) (10 x 15 cm) (0.3 x 3 m roll)	10 sheets 20 sheets 1 roll	1 209 272 1 209 299 1 417 240

\* available from Roche Molecular Biochemicals

- <sup>1)</sup> Expand is a trademark of a Member of the Roche Group.
   <sup>2)</sup> Tween 20 is a trademark of ICI Americas Inc., Wilmington, USA.
- 3) Nonidet P40 is a trademark of Shell International Petroleum Company Limited, U.K.
- <sup>4)</sup> CSPD is a trademark of Tropix, Inc., Bedford, MA, USA and covered under US patent 5112960 <sup>5)</sup> CDP-*Star* is a trademark of Tropix, Inc., Bedford, MA, USA and
- covered under US patent 5,326,882.

Notice to purchaser:

#### limited license

The purchase price of this product includes a limited, non-transferable license under U.S. Patents 4.683.202, 4.683.195, 4.965.188 or their foreign counterparts owned by Hoffmann-La-Roche Inc. and F. Hoffmann-La Roche Ltd. ("Roche"), to use only this amount of the product to practice the Polymerase Chain Reaction ("PCR") and related processes described in said patents solely for the research and development activities of the purchaser when this product is used in conjunction with an authorized thermal cycler.

No right to perform or offer commercial services of any kind using PCR, including without limitation reporting the results of purchaser's activities for a fee other commercial consideration, is hereby granted by implication or estoppel. Further information on purchasing licenses to practice the PCR process may be obtained by contacting the Director of Licensing at The Perkin-Elmer Corporation, 850 Lincoln Center Drive, Foster City, California 94404 or at Roche Molecular Systems, Inc. 1145 Atlantic Avenue, Alameda, California 94501.

#### Sold through an arrangement with ENZO DIAGNOSTICS, INC. Purchase of this product does not include any right or license to exploit this product commercially.

This product or the use may be covered by one or more ENZO patents, including the following:

U.S. Patent Nos. 4,711,955; 5,328,824; 5,449,767; 5,241,060; 4,994,373; and 5,175,269; EP 0 063 897 BI; EP 0 117 440 BI; EP 0 122 614 BI; and EP 0 128 332 BI; and Canadian Patent Nos. 1,219,824; 1,223,831; 1,309,672; 1,254,525; and 1,228,811.

#### E-mail Adress

argentina.biochem@roche.com biochem.au@roche.com Gerhard.Muehlbauer@roche.com biochem.be@roche.com Valent@mbox.cit.bg africhem@camnet.cm biochem.ca@roche.com biochem.cn@roche.com Info@medisell.com.cy Bm-comp@bm-comp.cz dk.biochem@roche.com ou.melestrum@neti.ee nharsc et@telecom net et helsinki.biochem\_diagnostics@roche.com biochem.fr@roche.com mannheim.biochemInfo@roche.com Bm\_roche@hotmail.com h.hajian@tebtech.com tubanegin@istn.irost.com Dyn@netvision.net.il it biochem@roche.com biochemicals@rdj.co.jp pharmakp@net2000ke.com Bmskorea@chollian.net react@ncc.moc.kw Raitis@invitros.lv Sakkijha@rdleb.com Gintaras@eksma.lt diagnostics@prophac.lu Vccl@vol.net.mt Aiouche echo@dounia net ma biocheminfo.nl@roche.com biochem.nz@roche.com bofungwu@linkserve.com.ng biochem.se@roche.com biochem.pt@roche.com Topdiag@fx.ro biochem.sg@roche.com roche.diagnostics@siol.net south\_africa.bioboffin@roche.com biochem.es@roche.com biochem.se@roche.com BiochemInfo.CH@roche.com Jean-Marie.kindbeiter@roche.com bmuae@emirates.net.ae uk.biochem@roche.com biochemts.us@roche.com Mvalentiner@telcel.net.ve dusica@eunet.yu biochemts.row@roche.com

#### Country Argentina Australia Austria Belgium Bulgaria Cameroon Canada China Cyprus Czech Republic Denmark Estonia Ethiopia Finland France Germany India Iran Iran Israel Italy Japar Kenya Korea Kuwait Latvia Lebanon Lithuania Luxembourg Malta Morocco Netherlands New Zealand Nigeria Norway

Portugal

Romania

Slovenia South Africa

Sweden

Switzerland

Spain

Singapore

Tunisia United Arab Emirates United Kingdom USA Venezuela Yugoslavia All other countries http://biochem.roche.com/pack-insert/1636090a.pdf Argentina 541 954 5555; Australia (02) 9899 7999; Austria (01) 277 87; Belgium (02) 247 4930; Brazil +55 (11) 3666 3565; Bulgaria +35929625408; Cameroon 237-370269; Canada (450) 086 7050; (800) 361 2070; Chile 00 56 (2) 22 33 737 (central) 00 56 (2) 22 32 099 (Exec); China 86 21 6427 5586; Colombia 0057-1-3412797; Cyprus +357-2-311362; Czech Republic (0324) 45 54, 58 71-2; Denmark +45 363 999 58; Egypt 20-2-3619047; Estonia 372-7-447600; Ethiopia 251-1-552799; Finland +358 9 525 333 66; France 04 76 76 30 87; Germany (0621) 759 8568; Greece 3 (01) 67 40 238; Hong Kong (852) 2485 7596; India +91-22-8379906; Indonesia 62 (021) 252 3820 ext. 755; Iran +98-21-8072374 / +98-21-8797027; Israel 972-6- 6380569; Italy 039 247 4109-4181; Japan 03 3432 3155; Kenya +254-2-750112; Korea 82-2-3471-6500; Kuwait +965-4837859; Latvia 371-787828309; Lebanon Fax: 00961-1-399667; Lithuania 370-2-729715; Luxembourg +352-496098; Malta Fax: +356-341087; Morocco Fax: +212-2-944040; Malaysia 60 (03) 755 5039; Mexico (5) 227 8967; Netherlands (036) 539 4911; New Zealand (09) 276 4157; Nigeria +234-1-521767; Norway (47) 23 373300; Philippines (632) 810 7246; Poland +48 (22) 22 66 84 305; Portugal (01) 4171717; Republic of Ireland 1 800 40 90 41; Romania +40-1-2123763; Russia (49) 621 759 8636 Fax: (49) 621 759 8611; Saudia Arabia +966-1-4010364; Singapore 0065 272 9200; Slovenia +386 61 1363528; South Africa (011) 886 2400; South Korea 02 569 6902; Spain (93) 201 4411; Sweden (08) 404 8800; Switzerland +41 (41) 799 6161; Taiwan (02) 736 7125; Thailand 66 (2) 274 07 08 (12 line); Turkey 0090 212 216 32 80; United Arab Emirates +971-4-694351; United Kingdom (0800) 521578; USA (800) 428 5433. Venezuela Fax: +0058-4810697; Yugoslavia +381 11 137163.



Roche Diagnostics GmbH Roche Molecular Biochemicals Sandhofer Strasse 116 D-68305 Mannheim Germany