

# QIAGEN Supplementary Protocol

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## Pre-amplification of gDNA and cDNA using the QIAGEN® Multiplex PCR Plus Kit

This protocol has been developed for use with the QIAGEN Multiplex PCR Plus Kit (cat. nos. 206151 and 206152) to enable simultaneous amplification of up to 80 targets, up to 300 bp in length (ideally 80–150 bp) using cDNA or gDNA as a template. This protocol describes pre-amplification of cDNA for applications such as two-step RT-PCR for gene expression analysis of multiple targets. It also describes gDNA pre-amplification for SNP/mutation detection from small amounts of gDNA for subsequent qPCR, end-point PCR, and Pyrosequencing®.

**IMPORTANT:** Please read the *QIAGEN Multiplex PCR Plus Handbook*, paying careful attention to the “Safety Information” and “Important Notes” sections, before beginning this procedure. The QIAGEN Multiplex PCR Plus Kit is intended for molecular biology applications. This product is not intended for the diagnosis, prevention, or treatment of a disease.

### Equipment and reagents

When working with chemicals, always wear a suitable lab coat, disposable gloves, and protective goggles. For more information, consult the appropriate material safety data sheets (MSDSs), available from the product supplier.

- PCR tubes/strips/plates
- Pipets and pipet tips (aerosol resistant)
- Thermal cycler
- Primers
- Primers should be purchased from an established oligonucleotide manufacturer. Lyophilized primers should be dissolved in TE buffer to provide a stock solution of 100  $\mu\text{M}$ ; concentration should be checked by spectrophotometry. Primer stock solutions should be stored in aliquots at  $-20^{\circ}\text{C}$ .



## Important points before starting

- **Always use the cycling conditions specified in this protocol.**
- **Use equal concentrations (0.1  $\mu\text{M}$ ) of all primers.**
- For optimal results, we recommend using primer pairs with a  $T_m$  of  $\geq 68^\circ\text{C}$ . For multiplex PCR primer design, see Appendix A, page 38 of the *QIAGEN Multiplex PCR Plus Handbook*.
- Prepare a 10x primer mix as described in page 13 of the *QIAGEN Multiplex PCR Plus Handbook*.
- **Do not use Q-Solution<sup>®</sup>** reagent that is provided with the kit.
- **Do not use CoralLoad<sup>®</sup> Dye** provided with the kits when performing qPCR or capillary sequencing subsequently.
- **PCR must start with an activation step of 5 minutes at 95°C** to activate HotStarTaq<sup>®</sup> Plus DNA Polymerase (see step 5 of this protocol).

## Procedure

1. **Thaw the 2x Multiplex PCR Master Mix (if stored at  $-20^\circ\text{C}$ ), template cDNA or gDNA, RNase-free water, and primer mix. Mix the solutions completely before use.**

It is important to mix the solutions completely before use to avoid localized concentrations of salts. Preparing a mixture of all primers avoids pipetting of individual primers for each experiment, reducing pipetting time and increasing reproducibility of results.

2. **Prepare a reaction mix according to Table 1.**

The reaction mix typically contains all the components required for multiplex PCR, except the template DNA. Prepare a volume of reaction mix 10% greater than that required for the total number of reactions to be performed. Primer concentration should be 0.1  $\mu\text{M}$  per primer (up to 80 primer pairs/reaction).

**Table 1. Reaction composition using 2x Multiplex PCR Master Mix**

Component	Volume/reaction	Final concentration
<b>Reaction mix</b>		
2x Multiplex PCR Master Mix*	12.5 $\mu$ l	1x
10x primer mix, 1 $\mu$ M each primer	2.5 $\mu$ l	0.1 $\mu$ M <sup>†</sup>
RNase-free water	Variable	–
<b>Template DNA</b>		
Template cDNA or gDNA, added at step 4	Variable	1–50 ng DNA
<b>Total volume</b>	25 $\mu$ l	–

\* Provides a final concentration of 3 mM MgCl<sub>2</sub>.

<sup>†</sup> A final primer concentration of 0.1  $\mu$ M is optimal for most primer–template systems. However, in some cases, using other primer concentrations (i.e., 0.02–0.1  $\mu$ M) may further improve amplification performance.

**3. Mix the reaction mixture thoroughly and dispense appropriate volumes into PCR tubes or wells.**

Mix gently, for example, by pipetting the reaction mix up and down a few times. Due to the hot start, it is not necessary to keep samples on ice during reaction setup.

**4. Add template DNA (1–50 ng/reaction) to the individual PCR tubes or wells containing the reaction mix. See Table 1.**

**5. a) Program the thermal cycler according to the manufacturer’s instructions.**

**b) Place the PCR tubes/plates/strips in the thermal cycler and start the cycling program as outlined in Table 2 or 3.**

Each PCR program must start with an initial heat-activation step at 95°C for 5 min to activate HotStarTaq *Plus* DNA Polymerase.

**6. After amplification, samples can be stored overnight at 2–8°C or at –20°C for long-term storage or subjected to a second round of PCR or qPCR amplification.**

**Note:** Use 0.2–2  $\mu$ l for subsequent PCR amplification (25  $\mu$ l final reaction volume). Additional purification of samples prior to subsequent amplification is not required.

**Table 2. Thermal cycler conditions — using cDNA as template**

<b>Step</b>	<b>Time</b>	<b>Temperature</b>	<b>Additional comments</b>
<b>Initial activation step</b>	5 min	95°C	HotStarTaq <i>Plus</i> DNA Polymerase is activated by this step.
<b>2-step cycling:</b>			
Denaturation:	15 s	95°C	
Annealing/extension:	120 s	60°C	Do not change annealing/extension conditions.
<b>Number of cycles:</b>	8–14		Do not use more than 14 cycles as the PCR must still be in a linear range to avoid uneven amplification of different amplicons.

**Table 3. Thermal cycler conditions — using gDNA as template**

<b>Step</b>	<b>Time</b>	<b>Temperature</b>	<b>Additional comments</b>
<b>Initial activation step</b>	5 min	95°C	HotStarTaq <i>Plus</i> DNA Polymerase is activated by this step.
<b>3-step cycling:</b>			
Denaturation:	30 s	95°C	
Annealing:	30 s	60°C	The annealing temperature of 60°C is suitable for most PCR systems when using gDNA as a template. For bisulfite-converted DNA, use 55°C.
Extension:	120 s	72°C	
<b>Number of cycles:</b>	8–14		Do not use more than 14 cycles as the PCR must still be in a linear range to avoid uneven amplification of different amplicons.

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