MPCR Kit for Human Apoptosis Genes Set-6
Cat No. MP-70018: 50 reactions
Cat No. MP-70010: 100 reactions

*These products are designed and sold for use in the Multiplex PCR (MPCR) covered by patent # 5,822,989. Use of the MPCR process requires a license. A limited, non-automated research field license under the patent to use only this amount of the product to practice the MPCR process is conveyed to the purchaser by the purchase of this product.

The Polymerase Chain Reaction (PCR) process is covered by patents owned by Hoffman-LaRoche. Use of the PCR process requires a license. A license for diagnostic purposes may be obtained from Roche Molecular Systems. A license for research may be obtained by the purchase and the use of authorized reagents and DNA thermocyclers from the Perkin-Elmer Corporation or by negotiating a license with Perkin-Elmer.

This product is intended for research use only and not for diagnostic purposes.
Mammalian cells can self-destruct via intrinsic programmed cell death (1). Apoptosis, a form of programmed cell death, is characterized by specific morphologic and biochemical properties. Apoptosis plays a major role in many biological processes, including embryogenesis, development of the immune system and tissue regeneration. Like growth and differentiation, apoptosis requires active and coordinated regulation of specific genes. Some molecular components of the apoptotic program have been conserved through evolution. Genetic studies of C. elegans have led to the identification of mutations, on 14 genes, that affect programmed cell death in this organism (1). Three of these genes, CED-3, CED-4, and CED-9, are essential for apoptosis. Each of these has at least one functional equivalent in mammals. CED-9 is a member of the Bcl-2 family, CED-3 is the prototypical worm caspase, and CED-4 is homologous to Apaf-1.

Ecotropic expression of bcl-2 has shown to block apoptosis in many experimental systems (2). Although the mechanism of bcl-2 action is unknown, biochemical studies have implicated this protein in the regulation of cell redox potential. Genetic evidence indicates that bcl-2 belongs to an emerging family. Some members of the bcl-2 family, like bcl-xl (3), suppress apoptosis, while other members, such as bax and bcl-xS (3,4), increase the susceptibility of cells to apoptotic stimuli. In addition, bcl-2 family members form hetero- and homo-dimers, suggesting that cellular factors control susceptibility to apoptosis is partly influenced by the level of gene expression.

ICE (Caspase-1) encodes a cystein protease that cleaves peptide bonds after Asp residues (5). The ICE (Interleukin-1 beta Converting Enzyme) family -recently named the caspases for cysteine aspartate-specific proteases- plays a central role in apoptosis and may well constitute part of the conserved core mechanism of the process. A family of caspases (Caspase #1 to #13) have been identified, which cleave key targets in the cell (6). Cell death is the outcome of a programmed intracellular cascade of genetically steps involving caspase activation and their subsequently catalytic reactions. Caspases normally exist in cells as inactive proenzymes; Proteolytic processing at a few specific sites unleashes their latent enzymatic activity and may triggers cell destruction.

Caspases can be activated by two distinct mechanisms. Because all caspases have similar cleavage specificity, the simplest way to activate a caspase is to expose it to a previously activated caspase molecule. This "caspase cascade" is used extensively by cells for the activation of the downstream effector caspases: caspase-3, caspase-6 (6).

The second strategy, "induced proximity", was first observed in caspase-8, an initiator caspase that acts downstream of Fas/Fasl(7). Upon ligand binding, Fas receptor aggregate into a membrane-bound complex. This signaling complex recruits, via the receptor-bound adapter protein FADD, several procaspase-8 molecules, resulting in a high local concentration of procaspase-8. Under these conditions, the low protease activity inherent to procaspases is sufficient to drive intermolecular proteolytic activation of the receptor-associated procaspase-8 molecules (6).

Analysis of the temporal and spatial distribution of RNA expression provides researchers with important clues about the function of apoptosis regulating genes in their own systems. Northern Blot and RNase Protection Assay are the most widely used procedures for determining the abundance of a specific mRNA in a total or poly(A) RNA sample. RT-MPCR provides an alternate and accurate method to detect multiple gene expression by amplifying all the genes under the same conditions (9, 10, 11). Variations in RNA isolation, initial quantitation errors or tube-to-tube variations in RT and PCR can be compensated by including a house-keeping gene, such as GAPDH, in MPCR. Alternatively, a parallel RT-PCR using the same cDNA, PCR conditions and primers for one of house-keeping genes may be run to offset any variations. Differences in gene expression can be determined by normalizing its expression against GAPDH expression.

Maxim’s hAP06G-MPCR kits have been designed to detect the expression of human ICH1 (Caspase-2), ICH2 (Caspase 4), MCH2 (Caspase 6), MCH3 (Caspase 7), MCH4 (Caspase 10) and GAPDH genes. The PCR primers have similar Tm and no obvious 3'end overlap to enhance multiple and equal amplification. The 500 bp (GAPDH), 419 bp (Caspase-6), 360 bp (Caspase-4), 297 bp (Caspase-7), 230 bp (Caspase-10), and 187 bp (Caspase-2 & S3) PCR products can be generated from human RNA or the positive control, which is included in this kit. Therefore, the hAP06G-MPCR kit provides a quick and simple method to analyze human Caspase-2, #4, #6, #7 and #10 gene expression during apoptosis.
I: Radioactive Quantitation

In our experience, visual inspection of an EtBr-stained agarose gel is sensitive and precise enough to detect changes as low as two-fold. If greater discrimination is necessary, several methods are available. The simplest procedure is to add a radioactively labeled dNTP to the PCR reaction. After gel analysis, the band may be excised and counted in a scintillation counter. Alternatively, the gel may be dried and an autoradiogram may be generated which can be scanned in a densitometer. Another method is to label the 5’ end of one or both of the primers with 32P, which is incorporated into the PCR products and then assayed for radioactivity (13).

Southern blot hybridization with synthetic DNA probes may also be performed to verify and quantitate PCR generated products, either by densitometry of an autoradiogram or by excising and counting the signal from a hybridization membrane. This method also quantitates only the target product without interference from nontarget products or primer-generated artifacts.

II: Non-Radioactive Quantitation

Nonradioactive quantitation methods include the use of biotinylated or digoxigenin-labeled primers in conjunction with the appropriate detection methods (14), use of a bioanalyzer or WAVE. For an in-depth discussion of the various methods of PCR product quantitation, refer to the review article by Bloch (15).

In addition to the above methods, several companies now offer gel video systems which can scan and quantitate EtBr-stained gel bands in much the same way a densitometer does. Lab-on-a-chip (BioAnalyzer), CE, HPLC, and WAVE may also be used to analyze MPCR products and quantitate simultaneously.

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MPCR (Multiplex Polymerase Chain Reaction)

√ Non-isotope method with high sensitivity
0.1-1μg total RNA per MPCR

√ Whole process takes only a few hours

√ Detect Multiple Genes Simultaneously &
Quantitatively

√ Signal can be quantified directly from gel
if isotope is included in MPCR. Additional
techniques can be used to quantify MPCR
product (using Bioanalyzer, HPLC, and WAVE.)

√ Non-specific products can be eliminated by using
probes and southern hybridization.

√ Ready-to-use

RPA (RNase Protection Assay)

√ Isotope or Non-Isotope methods
1-20 μg total RNA per RPA assay

√ Whole process takes two days

√ Detect Multiple Genes Simultaneously &
Quantitatively

√ Signal can be quantified directly from gel

√ Non-specific signal can be generated
by either low stringent conditions or
high-secondary-structure template.

√ Make own "hot" RNA probes
MPCR KIT DESCRIPTION

MPCR Amplification Kits include all necessary MPCR amplification reagents with the exception of Taq Polymerase. These kits have been designed to direct the simultaneous amplification of specific regions of human DNA.

MPCR Kits come in two quantities:
- 50X 50µL reaction kits
- 100X 50µL reaction kits

Each kit offers Maxim's optimal primer/buffer system which will enhance amplification specificity.

Figure 1 shows quality control MPCR results obtained by following MPCR kit manual using different concentrations of positive control.

For optimal results, please read and follow the instructions in this manual carefully. If you have any questions, please contact Maxim Biotech Customer Service at (650) 871-1919.

Lane N: PCR using hAPO6G Primers without positive (Negative)
Lane 1: PCR using hAPO6G Primers with 1X Positive without Enhancer
Lane 2: PCR using hAPO6G Primers with 1X Positive & Enhancer
Lane 3: PCR using Human GAPDH Primers
Lane 4: PCR using Human Caspase 6 Primers
Lane 5: PCR using Human Caspase 4 Primers
Lane 6: PCR using Human Caspase 7 Primers
Lane 7: PCR using Human Caspase 10 Primers
Lane 8: PCR using Human Caspase 2 Primers
Lane M: DNA M.W. Marker

MPCR PRIMER INFORMATION

<table>
<thead>
<tr>
<th>Product Code</th>
<th>Gene</th>
<th>5'/'3' Tm</th>
<th>Amplicon Size</th>
<th>Accession No.</th>
<th>Intron Span</th>
<th>Genomic Size</th>
</tr>
</thead>
<tbody>
<tr>
<td>hAPO6G-ICE2</td>
<td>Human Caspase-2 gene</td>
<td>71°C/70°C</td>
<td>187bp both S &amp; L</td>
<td>U13021</td>
<td>yes</td>
<td>1989bp</td>
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<tr>
<td>hAPO6G-ICE4</td>
<td>Human Caspase-4 gene</td>
<td>70°C/71.5°C</td>
<td>360bp</td>
<td>U25804</td>
<td>yes</td>
<td>29424bp</td>
</tr>
<tr>
<td>hAPO6G-ICE6</td>
<td>Human Caspase-6 gene</td>
<td>67°C/67°C</td>
<td>419bp</td>
<td>U20536</td>
<td>yes</td>
<td>1700bp</td>
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<tr>
<td>hAPO6G-ICE7</td>
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<td>hAPO6G-ICE10</td>
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<td>U60519</td>
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<td>230bp</td>
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<tr>
<td>hAPO6G-GAP</td>
<td>Human GAPDH</td>
<td>62°C/63°C</td>
<td>500bp</td>
<td>M33197</td>
<td>yes</td>
<td>2533bp</td>
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## KIT COMPONENTS

**MP-70018**

5X50μL MPCR reaction kit  
*Store all reagents at -20°C*

<table>
<thead>
<tr>
<th>Product Code</th>
<th>Kit Component</th>
<th>Amount</th>
</tr>
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<tbody>
<tr>
<td>hAPO6G-B001</td>
<td>2X hAPO6G MPCR Buffer (containing chemicals, enhancer, stabilizer and dNTPs)</td>
<td>1250 μl</td>
</tr>
<tr>
<td>hAPO6G-CD01</td>
<td>10X hAPO6G MPCR Pos. Control</td>
<td>50μl</td>
</tr>
<tr>
<td>hAPO6G-P001</td>
<td>10X hAPO6G MPCR Primers</td>
<td>250μl</td>
</tr>
<tr>
<td>MRB-0014</td>
<td>DNA M.W. Marker (100bp Ladder)</td>
<td>100 μl</td>
</tr>
<tr>
<td></td>
<td>ddH₂O (DNase free)</td>
<td>2.0 ml</td>
</tr>
<tr>
<td></td>
<td>Instruction Manual</td>
<td></td>
</tr>
</tbody>
</table>

**MP-70010**

100X50μL MPCR reaction kit  
*Store all reagents at -20°C*

<table>
<thead>
<tr>
<th>Product Code</th>
<th>Kit Component</th>
<th>Amount</th>
</tr>
</thead>
<tbody>
<tr>
<td>hAPO6G-B001</td>
<td>2X hAPO6G MPCR Buffer (containing chemicals, enhancer, stabilizer and dNTPs)</td>
<td>1250 μl X2</td>
</tr>
<tr>
<td>hAPO6G-CD01</td>
<td>10X hAPO6G MPCR Pos. Control</td>
<td>50μl X2</td>
</tr>
<tr>
<td>hAPO6G-P001</td>
<td>10X hAPO6G MPCR Primers</td>
<td>250μl X2</td>
</tr>
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<td>MRB-0014</td>
<td>DNA M.W. Marker (100bp Ladder)</td>
<td>100 μl X2</td>
</tr>
<tr>
<td></td>
<td>ddH₂O (DNase free)</td>
<td>2.0 ml</td>
</tr>
<tr>
<td></td>
<td>Instruction Manual</td>
<td></td>
</tr>
</tbody>
</table>

**NOTE:** SPIN ALL TUBES BEFORE USING!!
**RT Protocol:**

The isolation of undegraded, intact RNA is an essential prerequisite for successful first strand synthesis and PCR amplification. Care should be taken to avoid RNase contamination of buffers and containers used for RNA work by pretreating with DEPC, autoclaving, and baking. Always wear sterile gloves when handling reagents. Use cDNA derived from 10^6 cells (1 mg cDNA) and apply them to each MPCR reaction.

1. Prepare total RNA, mRNA or use the control GAPDH RNA which is provided in Maxim's MPCR kit. **NOTE:** It is best to use cDNA derived from 0.5-1 x 10^6 cells (0.5-1 mg cDNA derived from RNA) for each MPCR reaction.

2. Equilibrate 3 water baths: 37°C, 70°C and 95°C.

3. **On ice,** pipet 1-2 mg mRNA or 10 µg total RNA (from 10^6 cells) dissolved in pure water or 2 ml control GAPDH RNA into a RNase free reaction vial. We strongly recommend including a positive control reaction when setting up an RT-PCR reaction for the first time.

4. Add sterile water to a final volume of 14.5 µl.

5. Add 4 µl random hexamer (50 mM) or Oligo(dT) (50 mM).
   **NOTE:** The hexamer and Oligo(dT) RT reactions may be run simultaneously.

6. Incubate tube(s) at 70°C for 5 minutes and quickly chill on ice.

7. Begin your RT reaction by adding the following reagents to your hexamer or Oligo mixture:

<table>
<thead>
<tr>
<th>Reagent</th>
<th>Description</th>
<th>Volume per Reaction</th>
</tr>
</thead>
<tbody>
<tr>
<td>RNase Inhibitor</td>
<td>130U/µl</td>
<td>0.5µl</td>
</tr>
<tr>
<td>5 X RT buffer</td>
<td>250mM Tris-HCl (pH8.3)</td>
<td>10µl</td>
</tr>
<tr>
<td></td>
<td>375mM KCl, 15mM MgCl2, 50mMDTT</td>
<td></td>
</tr>
<tr>
<td>dNTPs</td>
<td>1mM each</td>
<td>20µl</td>
</tr>
<tr>
<td>MMLV RT</td>
<td>250U/µl</td>
<td>1µl</td>
</tr>
</tbody>
</table>

8. Incubate the RT mixture at 37°C for 60 minutes.

9. Then, heat RT mixture at 95°C for 10 minutes and quickly chill on ice.

10. Add another 50 µl water or 0.1X TE buffer.

11. 2-5 µl of above cDNA is sufficient for most genes in a standard MPCR reaction. However, more or less DNA may be needed in PCR depending on the copy number of the specific gene.

**PCR Protocol:**

1. Taq DNA polymerase from Perkin-Elmer or its derivatives are highly recommended for MPCR. AmpliTaq Gold, however, is not recommended because its own optimal buffer system is required.

2. **Reaction Mixture Preparation:**
   
   A. Set up MPCR reactions with the test samples and MPCR buffers provided in the MPCR kit according to the table below:
### Volume (Per assay) | Reagent (Add in order)
---|---
25.0 µl | 2X MPCR BufferMixture
5.0 µl | 10X MPCR Primers
0.5 µl | Taq DNA Polymerase(5U/ml)
5.0 µl | Specimen cDNA or
14.5 µl | 10X Control cDNA from kit
50.0 µl | H₂O

*: ³²P dNTPs may be used here to achieve higher sensitivity and better quantitation. 5-10 µCi [α-³²P]dCTP (3000 Ci/mmmole) should be used here per MPCR. Keep final dNTPs concentration same as without ³²P-dNTPs.

B. EDTA concentration in test sample must not exceed 0.5 mM because Mg²⁺ concentration in MPCR Buffers is limited to certain ranges. Additional Mg²⁺ may be added to the PCR mixture to compensate for EDTA. We strongly recommend running an MPCR reaction with the positive control provided in the kit. Since the MPCR DNA polymerase needed in each reaction is in a very small volume, it is recommended that all of the PCR components be premixed in a sufficient quantity for daily needs and then dispensed into individual reaction vials. This will help you to achieve more accurate measurements.

### 3. PCR thermocycle profile:

Reaction profiles will need to be optimized according to the machine type and needs of user. Please take note that temperature variations occur between different thermocyclers, therefore, the annealing temperature in the sample profile below is given as a range. It will be necessary to determine the optimal temperature for your individual thermocycler. An example of a time-temperature profile for the positive control PCR reaction optimized for Perkin Elmer machine types 480, 2400, and 9600 is provided below:

<table>
<thead>
<tr>
<th>Temperature</th>
<th>Time</th>
<th>Cycles</th>
</tr>
</thead>
<tbody>
<tr>
<td>96°C</td>
<td>1 min</td>
<td>2X</td>
</tr>
<tr>
<td>58-60°C</td>
<td>4 min</td>
<td>2X</td>
</tr>
<tr>
<td>94°C</td>
<td>1 min</td>
<td>28-35X</td>
</tr>
<tr>
<td>58-60°C</td>
<td>2 min</td>
<td>1X</td>
</tr>
<tr>
<td>70°C</td>
<td>10 min</td>
<td>1X</td>
</tr>
<tr>
<td>25°C</td>
<td>soak</td>
<td></td>
</tr>
</tbody>
</table>

**Note:** A 2-step PCR thermocycle profile was found to be more effective than a 3-step PCR thermocycle profile for MPCR amplification. For 2-step PCR, use 94-95°C for denaturation and 58-60°C for annealing and extension. The 72°C step is omitted.

### 4. Agarose Gel Electrophoresis:

To fractionate the MPCR DNA product electrophoretically, mix 10µl of the MPCR product with 2µl 6X loading buffer. Run the total 12µl alongside 10 µl of DNA marker from the MPCR kit on a 2 % agarose gel containing 0.5 mg/ml ethidium bromide. Electrophoresise and photograph. (Hint: Best results are obtained when the gels are run slowly at less than 100 volts).
# 1. MPCR AMPLIFICATION

<table>
<thead>
<tr>
<th>Observation</th>
<th>Possible Cause</th>
<th>Recommended Action</th>
</tr>
</thead>
</table>
| 1.1. No signal or missing some bands during amplification even using positive control provided in kit. | 1.1a. The annealing temperature in thermocycler is too high.  
1.1b. Dominant primer dimers. | 1.1a. Decrease PCR annealing temperature 3-5°C gradually.  
1.1b. Use any one of "Hot Start" PCR procedures. |
| 1.2. Too many nonspecific bands.                                             | 1.2a. The annealing temperature in the thermocycler is too low.  
1.2b. Pre-PCR mispriming.  
1.2c. cDNA is interfering with MPCR | 1.2a. Increase PCR annealing temperature 3-5°C gradually.  
1.2b. Use any one of "Hot Start" PCR procedures.  
1.2c. Clean cDNA with Phenol/Chloroform.  
1.2d. Use Maxim's 3M™-MPCR Kit. |
| 1.3. No difference in gene expression among treatments                      | 1.3a. PCR amplification of this specific gene has passed the exponential phase.  
1.3b. Variation in sample preparation, RT reaction and amounts of input cDNA. | 1.3a. Decrease PCR cycle number or decrease the input cDNA.  
1.3b. Run a parallel PCR with a house-keeping gene to eliminate variables. |
Storage

1. Store all MCPR Kit components at -20°C. Under these conditions components of the kit are stable for 1 year.
2. Isolate the kits from any sources of contaminating DNA, especially amplified PCR product.
3. Do not mix MCPR kit components that are from different lots. Each lot is optimized individually.

REFERENCES