

CelGREEN Universal qPCR mix (2x)

| | |
|-------------|-----------|
| CM3111-0200 | 200 rxns |
| CM3111-0500 | 500 rxns |
| CM3111-1000 | 1000 rxns |

Store the kit at -20°C



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What's in the box

| Cat no. | Pack size | 2x CelGREEN Universal qPCR mix | ROX Additive (50µM) |
|-------------|-----------|--------------------------------|---------------------|
| CM3111-0200 | 200 rxns | 2 x 1ml | 1 x 0.2ml |
| CM3111-0500 | 500 rxns | 5 x 1ml | 1 x 0.2ml |
| CM3111-1000 | 1000 rxns | 10 x 1ml | 2 x 0.2ml |

Product description

CelGREEN Universal qPCR mix combines an antibody-mediated hot-start polymerase with an advanced buffer system to achieve superior PCR sensitivity and greater PCR specificity by limiting the formation of primer dimers. The proprietary intercalating dye that enables amplicon quantification does not affect the efficiency of the PCR (as reported for other commonly used dyes). This mix performs well, even with AT- and GC- rich templates and requires minimal optimization.

Notes

For research use only.

Reaction set-up

Briefly vortex 2x CelGREEN Universal qPCR mix to ensure homogeneity.

Prepare a master mix using the guidelines below.

| Reagent | 20µl reaction | Final concentration |
|--------------------------------|------------------------------|-----------------------------|
| 2x CelGREEN Universal qPCR mix | 10µl | 1x |
| Forward primer (10µM) | 0.8µl | 400nM |
| Reverse primer (10µM) | 0.8µl | 400nM |
| Template DNA | Variable | See template considerations |
| PCR grade water | Up to a final volume of 20µl | |

Shipping and storage

Store the kit at -20°C upon arrival. This product is shipped on ice blocks and can be kept at 4°C for 4 weeks without affecting the product performance. Full enzyme activity is retained for 12 months when stored at the recommended conditions and can endure up to 30 freeze/thaw cycles. Limit direct light exposure for extended periods.

Recommended cycling conditions

| Description | Temperature | Time | Cycles |
|--|-------------|-------------|--------|
| Enzyme activation and initial denaturation. Use 2min for cDNA and 3min for genomic DNA | 95°C | 2 to 3min | 1 |
| Denaturation | 95°C | 5sec | 40 |
| Annealing and extension, maximum 30sec, minimum 60°C* | 60 to 65°C | 20 to 30sec | |
| Melt analysis (optional) – refer to the instrument manual | | | |

*Data acquisition in the FAM channel

Instrument compatibility

CelGREEN Universal qPCR mix is suitable for use on all real time PCR cyclers. The Universal qPCR mix is supplied with a separate tube of 50µM ROX dye that may be used with instruments requiring ROX as a passive reference dye. Please refer to the instrument manufacturer's manual to confirm the required ROX concentration.

ROX protocol:

Add the 50µM ROX additive directly to the 1ml tube of 2x CelGREEN Universal qPCR mix and vortex thoroughly. Use immediately or store as per normal storage instructions. Refer to the directions below for guidelines to obtain the desired concentration.

Hi-ROX instruments:

Add 20µl of ROX additive (50µM) to 1ml of 2x CelGREEN Universal qPCR mix to obtain a final concentration of 1µM. This will result in a concentration of 500nM per 20µl reaction.

Lo-ROX instruments:

Add 2µl of ROX additive (50µM) to 1ml of 2x CelGREEN Universal qPCR mix to obtain a final concentration of 100nM. This will result in a concentration of 50nM per 20µl reaction.

General Considerations

Primer design: Design primers with a predicted melting temperature of around 60°C using software such as Primer 3 (<http://frodo.wi.mit.edu/primer3/>). The amplicon should be between 80 and 200bp in length to allow efficient amplification under fast cycling conditions, and no longer than 400bp for standard cycling conditions. Bear in mind that shorter amplicons will result in quicker cycling times.

Template: Add 1µg or less high-quality genomic DNA per 20µl reaction or up to 100ng cDNA per 20µl reaction. Perform PCR with a template dilution series to determine the optimal amount of template to use with new primer pairs for maximum amplification efficiency.

Troubleshooting

Non-specific amplification – (presence of more than one peak during melt analysis) – possible actions to resolve the problem:

1. Increase the reaction annealing temperature to enhance the specificity of priming.
2. Systematically eliminate potential contamination in reaction components by replacing single components in separate PCR reactions, until the source of contamination is identified and isolated.
3. Reduce the primer concentration to increase the specificity of priming.
4. Perform PCR set-up and PCR product analysis in separate areas

No PCR product visible – possible actions to resolve the problem:

1. Confirm all required components were added (in the correct volumes) during PCR reaction set-up. Ensure the FAM channel is selected in the instrument software for successful signal acquisition.
2. Ensure components were stored at the recommended storage conditions listed above. Test each component in separate reactions to eliminate potential defective reagent.
3. Decrease the amount of DNA added to the reaction to reduce PCR inhibitors that may be present in samples.
4. Decrease the annealing temperature to reduce primer specificity.
5. Increase the number of cycles to allow for additional exponential amplification.

Technical support

For technical support please e-mail info@celticmolecular.com