



PCRBIO SYSTEMS

simplifying research

## Clara™ HRM Mix

[www.pcrbio.com](http://www.pcrbio.com)

### Product description:

Clara™ HRM Mix comprises our ultra-pure, Taq polymerase in a unique blend containing dNTPs and MgCl<sub>2</sub>. The mix is powered by our third-generation DNA-intercalating SyGreen 2 dye for greatly reduced PCR inhibition. This new generation qPCR mastermix offers superior performance for accurate SNP discrimination and quantification of methylation differences.

HRM analysis is a powerful technique for the analysis of mutations, polymorphisms and epigenetic differences in double-stranded DNA samples.

Clara™ HRM Mix relies on our ultra-pure Taq polymerase, purified using our 12-step purification method to totally eliminate host DNA contamination and improve reaction sensitivity and specificity.

Our high-throughput smart-screen technology screening has resulted in a buffer system that allows efficient amplification from GC-rich and AT-rich templates, under fast and standard cycling conditions and results in higher sensitivity in distinguishing every class of SNP.

### Quality control

PCR Biosystems operates under an ISO 13485 certified Quality Management System. Our products are extensively tested and undergo a comprehensive, multi-step quality control process according to ISO 13485 standards, to ensure optimum performance, consistency and traceability.

| Pack Size      | 2x Clara™ HRM Mix |
|----------------|-------------------|
| 100 reactions  | 1 x 1 mL          |
| 500 reactions  | 5 x 1 mL          |
| 2000 reactions | 20 x 1 mL         |

### Shipping and storage

On arrival the kit should be stored between -30°C and -15°C. Avoid prolonged exposure to light. If stored correctly the kit will retain full activity for 12 months. The kit can be stored at 4°C for 1 month.

### Limitations of product use

This product is for research use only.

### Technical support

Help and support is available on our website at <https://pcrbio.com/resources/> including answers to frequently asked technical questions and our qPCR technical guide. For technical support and troubleshooting you can submit a technical enquiry online, or alternatively email [technical@pcrbio.com](mailto:technical@pcrbio.com) with the following information:

- Amplicon size
- Reaction setup
- Cycling conditions
- Screen grabs of amplification traces and melting profile

## Instrument compatibility

| Manufacturer          | Instrument  |
|-----------------------|---|
| Applied Biosystems    | 7900, 7900HT, 7900HT FAST, StepOne™, StepOne™ Plus, 7500, 7500 FAST, Viia7™ |
| Bio-Rad®              | CFX96™, CFX384™   |
| Eppendorf             | Mastercycler® ep realplex, Mastercycler® realplex 25                        |
| Illumina®             | Eco™  |
| Qiagen/Corbett        | 6000, Q   |
| Roche Applied Science | Lightcycler®480, Lightcycler® 96, Lightcycler®Nano                          |
| Bio Molecular Systems | Mic qPCR Cycler   |

**Instrument Selection:** Not all real time thermocyclers are suitable for HRM analysis. We provide a chart above for quick reference. You can also use our selection tool (<https://pcrbio.com/resources/qpcr-selection-tool/>). If your instrument is not listed please refer to the manufacturer's manual or get in touch with us at [technical@pcrbio.com](mailto:technical@pcrbio.com) to find out whether or not it is suitable for HRM experiments.

## Important considerations

**Primer design:** For efficient amplification under fast cycling conditions we recommend amplicon lengths between 80bp and 200bp. With all manufacturers master mixes the shorter the amplicon length the faster the reaction can be cycled. Amplicon lengths should not exceed 400bp. Primers should have a predicted melting temperature of around 60°C, using default Primer 3 settings (<https://bioinfo.ut.ee/primer3/>).

## Reaction setup

1. Before starting, briefly vortex 2x Clara™ HRM Mix.
2. Prepare a master mix based on following table:

| Reagent                     | 20µL reaction            | Final concentration | Notes                               |
|-----------------------------|--------------------------|---------------------|-------------------------------------|
| 2x Clara™ HRM Mix           | 10 µL                    | 1 x                 |                                     |
| Forward primer (10 µM)      | 0.8 µL                   | 400 nM              | See above for optimal primer design |
| Reverse primer (10 µM)      | 0.8 µL                   | 400 nM              |                                     |
| Template DNA                | 0.5-50 ng genomic        | variable            |                                     |
| PCR grade dH <sub>2</sub> O | Up to 20 µL final volume |                     |                                     |

3. Program the instrument using following conditions, acquiring data on the FAM channel:

| Cycles       | Temperature                      | Time                       | Notes  |
|--------------|----------------------------------|----------------------------|--|
| 1            | 95 °C                            | 2 min                      | Polymerase activation, 2 minutes for cDNA and 3 minutes for genomic                                |
| 45           | 95 °C<br>60 °C to 65 °C          | 5 seconds<br>20-30 seconds | Denaturation<br>Annealing/Extension, do not exceed 30 seconds, do not use temperatures below 60 °C |
| HRM analysis | Refer to instrument instructions |                            |  |