PCRBIOSYSTEMS
simplifying research

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Clara® Inhibitor-Tolerant Probe Mix No-ROX

Product description

Clara® Inhibitor-Tolerant Probe Mix offers reliable probe-based qPCR detection of DNA target sequences in the presense of qPCR inhibitors. This 4x mix gives superior target amplification, in single or multiplex assays, even from highly dilute samples.

Clara® Inhibitor-Tolerant Probe Mix contains hot start Taq polymerase, dNTPs, and $MgCl_2$. It is developed to work well with the full range of probe types, including TaqMan, Scorpions and molecular beacons.

The mix contains excipients which ensure reliable performance in crude saliva samples and in the presence of PCR-inhibitory compounds that include but are not limited to: standard laboratory chemicals (SDS, guanidine, and ethanol), and biological sample inhibitors, such as those found in blood samples (hemin, hematin, haemoglobin, heparin, IgG immunoglobulins , lactoferrin, sodium citrate), urine (urea), plant, and environmental samples (humic acid, catechin, quercetin, tannic acid, cellulose, and chlorophyl).

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	4x Clara® Inhibitor-Tolerant Probe Mix No-ROX
200 reactions	1 x 1 mL
600 reactions	3 x 1 mL
1000 reactions	5 x 1 mL
10000 reactions	1 x 50 mL

Shipping and storage

On arrival the kit should be stored between -30 °C and -20 °C. Avoid prolonged exposure to light. If stored correctly, the kit will retain full activity until the indicated expiry date. The kit can be stored at 4 °C for 1 month.

Technical support

Scan or click the QR code for troubleshooting help and answers to frequently asked technical questions. For further technical support, please email technical@pcrbio.com with the following information:

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

TROUBLESHOOT

FAQS

Important considerations

Instrument compatibility: Different qPCR instruments may require different levels of ROX passive reference for normalisation. Use our qPCR Selection Tool to determine which ROX concentration your instrument requires (https://pcrbio.com/resources/qpcr-selection-tool/).

Template: The mix can be used with cDNA or DNA synthesised or extracted by most commercial kits or standard extraction methods and is also suitable for crude sample input. Addition of 2 to 5 µL volumes of sample will improve assay precision. For genomic DNA, 1 µg or less is recommended. For cDNA, 100 ng or less is recommended. However, users are encouraged to attempt a dilution series for new template/primer pairs to ensure that the PCR is efficient at that template concentration.

Primer design: For efficient amplification we recommend amplicon lengths between 80-200 bp and not exceeding 400 bp. Shorter amplicons allow for faster cycling. Primers should have an approximate Tm of ~60 ˚C using default Primer 3 settings (https://bioinfo.ut.ee/primer3/). To verify the best annealing temperature for your primers in our products, please visit https://pcrbio.com/resources/tm-calculator/. For TaqMan probes choose a probe close to the 5' primer, avoiding terminal guanosine residues.

Reaction setup

1. Before starting, thaw and briefly vortex the 4x Clara® Inhibitor-Tolerant Probe Mix.
2. Prepare a master mix based on the following table:

Reagent	20 µL reaction	Final concentration	Notes
4x Clara® Inihbitor-Tolerant Probe Mix	5 µL	1x	
Forward primer (10 µM)	0.8-2 µL	400 nM - 1 µM	See above for optimal primer design
Reverse primer (10 µM)	0.8-2 µL	400 nM - 1 µM	
Probe (10 µM)	0.25-1 µL	125 - 500 nM	
DNA Template	2-5 µL	Variable	<100 ng cDNA, <1 µg genomic DNA
PCR grade dH_2O	Up to 20 µL final volume		

3. Program the instrument using the following conditions, acquiring data on the appropriate channel(s) for your chosen probe(s):

Cycles	Temperature	Time	Notes
1	95 ˚C	2 minutes	Polymerase activation
50	95 ˚C 55 ˚C to 65 ˚C	15 seconds 30 seconds	Denaturation Anneal/Extension
Melt analysis	Refer to instrument instructions		Optional melt profile analysis, available for hybridisation probes only