GoTaq® Probe qPCR: A Real-Time PCR Master Mix Optimized for Hydrolysis Probe Assay

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Abstratct

The **GoTaq**® **Probe qPCR Master Mix** is a ready-to-use, 2X master mix for qPCR using hydrolysis probe detection, and is designed for sensitive detection and quantification over a broad range of DNA or RNA targets in the presence of a wide range of PCR inhibitors. Rapid hot-start activation and processive enzymes makes these mixes compatible with any real-time PCR.

The **GoTaq**[®] **Probe 2-Step RT-qPCR** facilitates detection and relative quantification of RNA expression levels via a two-step RT-qPCR method. The GoScript[™] Reverse Transcription System enables efficient synthesis of first-strand cDNA which may be added directly to downstream qPCR reactions.

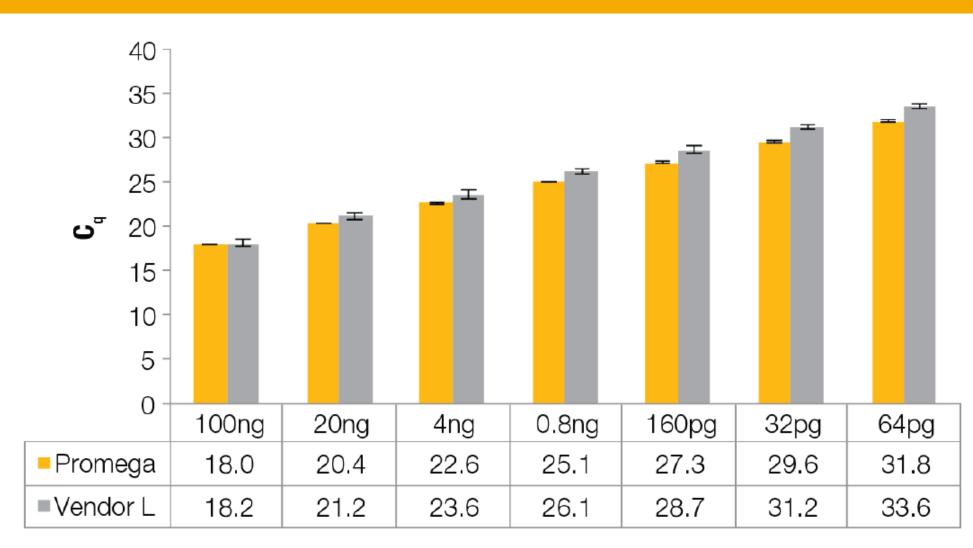
The **GoTaq**[®] **Probe 1-Step RT-qPCR with dUTP** is optimized for detection and relative quantification of RNA expression levels using a one-step RT-qPCR method, combining GoScript™ Reverse Transcriptase and GoTaq[®] Probe qPCR Master Mix in single-step real-time amplification reactions.

The master mix does not contain a reference dye; a separate tube of carboxy-X-rhodamine (CXR) reference dye is included.

Features of the GoTag® Probe qPCR Master Mix

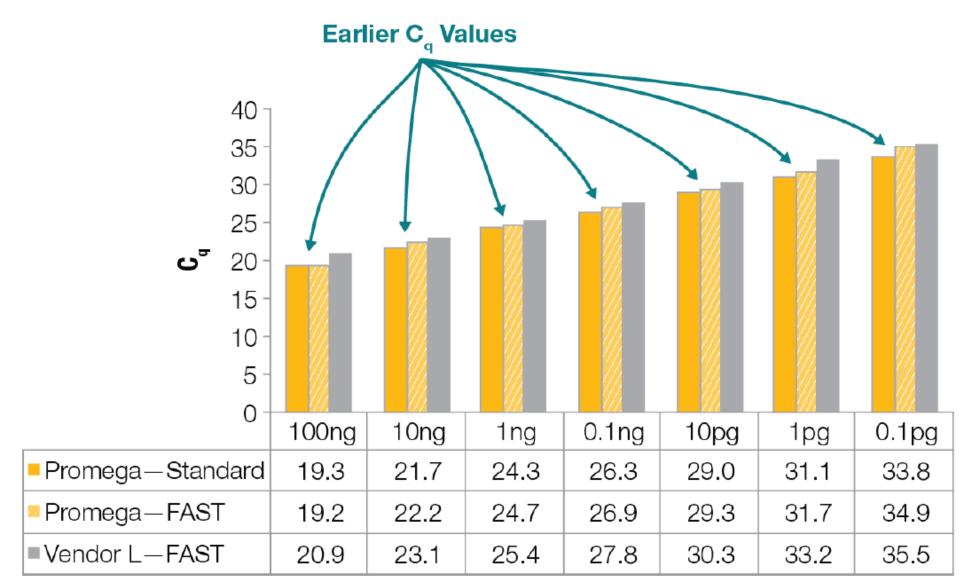
- Designed to provide resistance to a wide range of PCR inhibitors.
- Uses antibody-mediated hot-start chemistry, allowing reaction setup to be performed at room temperature.
- Employs rapid hot-start activation and processive enzymes, making it compatible with both standard and fast instrument cycling programs.

Sensitivity - Detection of picograms levels of RNA input



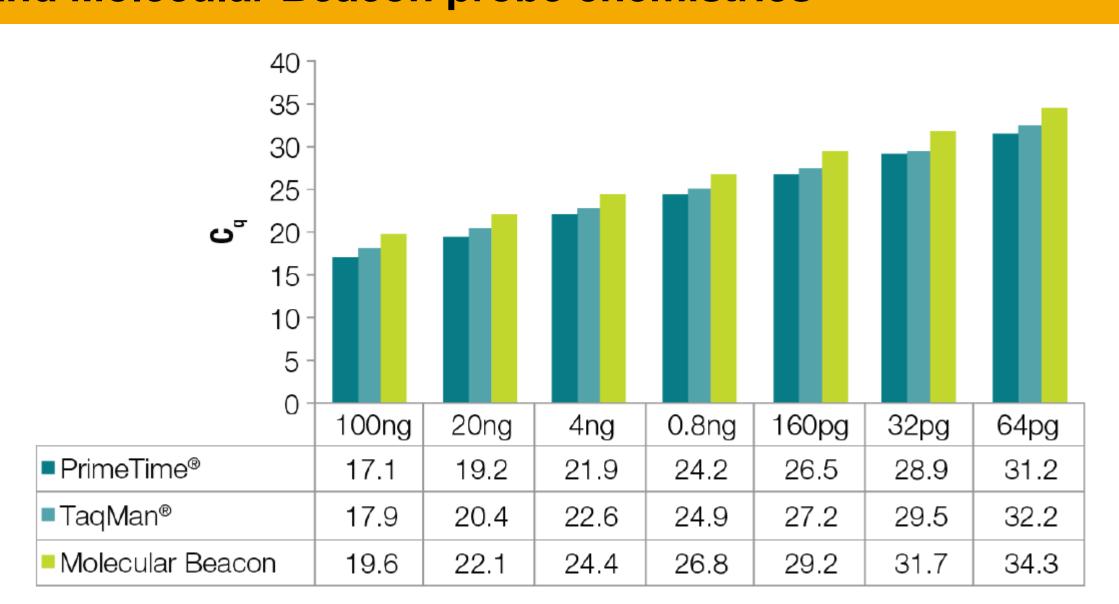
Amplification curve for GAPDH detection from human pancreas RNA using the GoTaq® Probe 2-Step RT-qPCR System. A series of fivefold serial dilutions of human pancreas cDNA was used to measure GAPDH gene expression using the GoTaq® Probe 2-Step RT-qPCR System and a competing product (Vendor L). The GoTaq® Probe 2-Step RT-qPCR System shows earlier Cq values (>0.2) for all samples. Two microliters of human pancreas RNA was reverse transcribed using the GoScript™ Reverse Transcriptase supplied in the kit followed by qPCR with GoTaq® Probe qPCR Master Mix.

Versatility - Works with both standard and fast thermal-cycling conditions



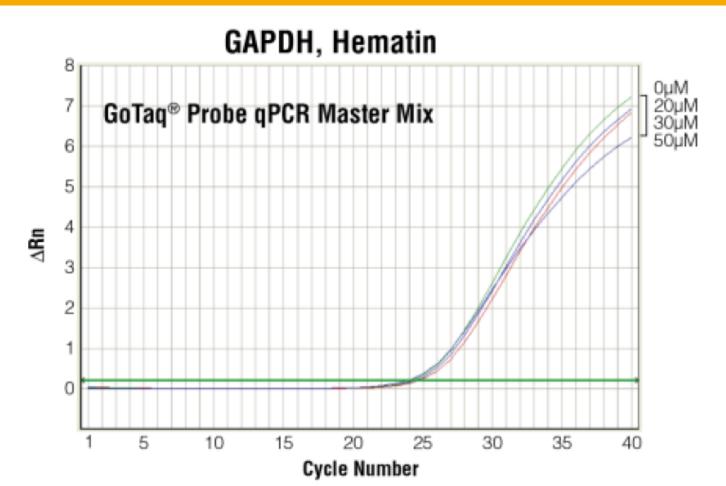
GoTaq® Probe qPCR Master Mix shows comparable results when GAPDH is amplified from human pancreas cDNA using standard and fast thermal-cycling conditions. Earlier Cq values were observed for both standard and fast thermal-cycling conditions when GoTaq® Probe qPCR Master Mix was compared to a competing product (Vendor L).

Compatibility - Suitable for PrimeTime®, TaqMan®, and Molecular Beacon probe chemistries



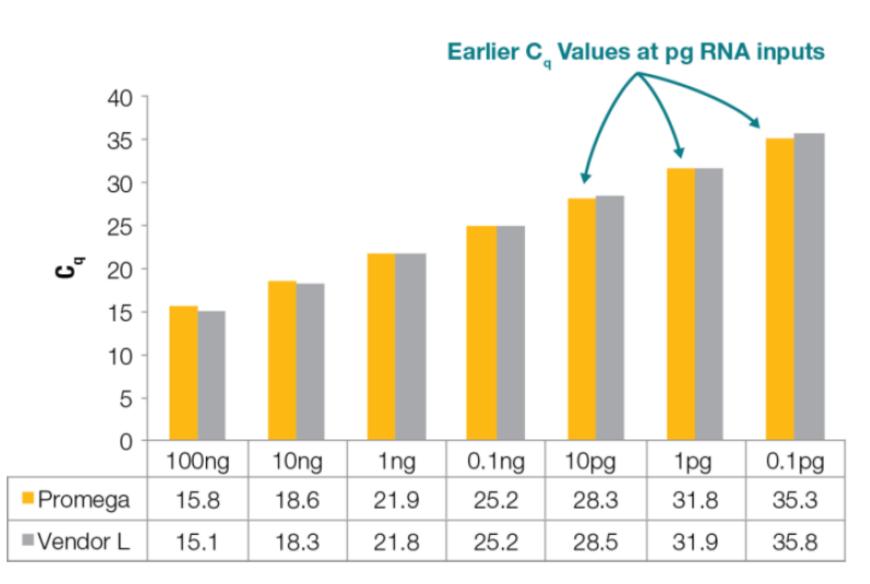
GoTaq® Probe 2-Step RT-qPCR System works with a variety of probe assay chemistries. A series of 7 fivefold dilutions of human heart RNA was analyzed for GAPDH gene expression using PrimeTime®, TaqMan®, and Molecular Beacon hydrolysis probe chemistries and the GoTaq® Probe 2-Step RT-qPCR System. All sample dilutions generated results with all chemistries tested using the GoTaq® Probe 2-Step RT-qPCR System.

Robustness - Resistant to a wide range of qPCR inhibitors

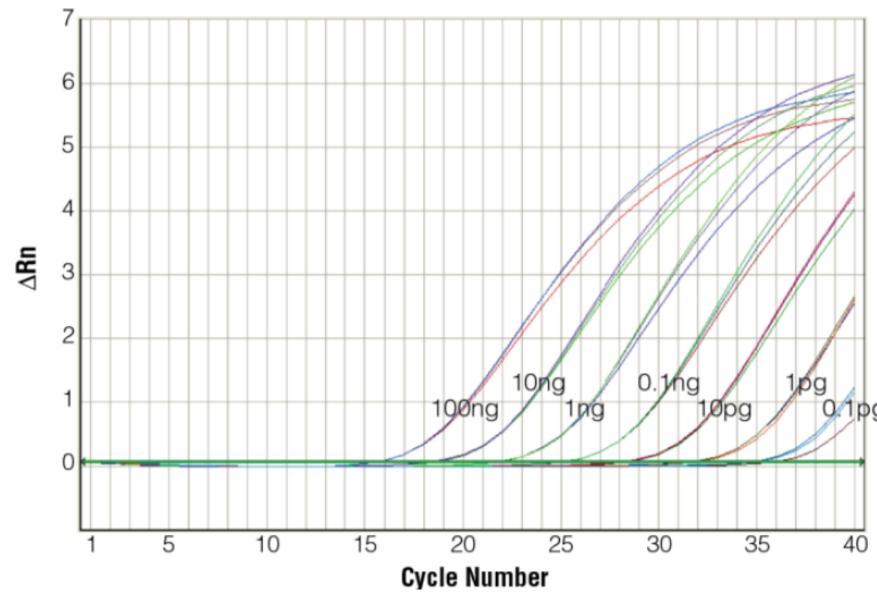


0–50µM of Hematin added to qPCR reactions amplifying GAPDH from human liver RNA. RNA was converted to cDNA with GoScript™ Reverse Transcription System. GoTaq® was robust enough to amplify all samples containing Hematin.

Dynamic Range – RNA detection from 100 ng down to 1pg



Cq values for 6 x tenfold dilution series of human control RNA detecting GAPDH in a 20µl 1-step RT-qPCR reaction comparing GoTaq® Probe 1-Step RT-qPCR System to a 1-Step RT-qPCR master mix from Vendor L. GoTaq® Probe 1-Step RT-qPCR System gave earlier Cq values with the pg of RNA samples showing greater sensitivity.



Amplification curve for the 6 x tenfold dilution series of human control RNA detecting GAPDH with the GoTaq® Probe 1-Step RT-qPCR System. The qPCR reaction efficiency was calculated to be 102.2% with an r2 of 0.998.

Summary

GoTaq® Probe qPCR and RT-qPCR Master Mixes are optimized for qPCR assays in the hydrolysis probe detection format and offers:

- Optimized formulation for a variety of PCR applications
- Easy room-temperature setup for automated and high-throughput detection
- Sensitive detection on most real-time instruments
- Versatility with both fast and standard cycling methods
- Resistant to a wide variety of PCR inhibitors
- Compatible with multiple probe chemistries
- Robust, highly efficient, full-length cDNA synthesis in the presence of inhibitors
- Dynamic range >6 orders of magnitude
- Backed by the Promega PCR Performance Guarantee



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