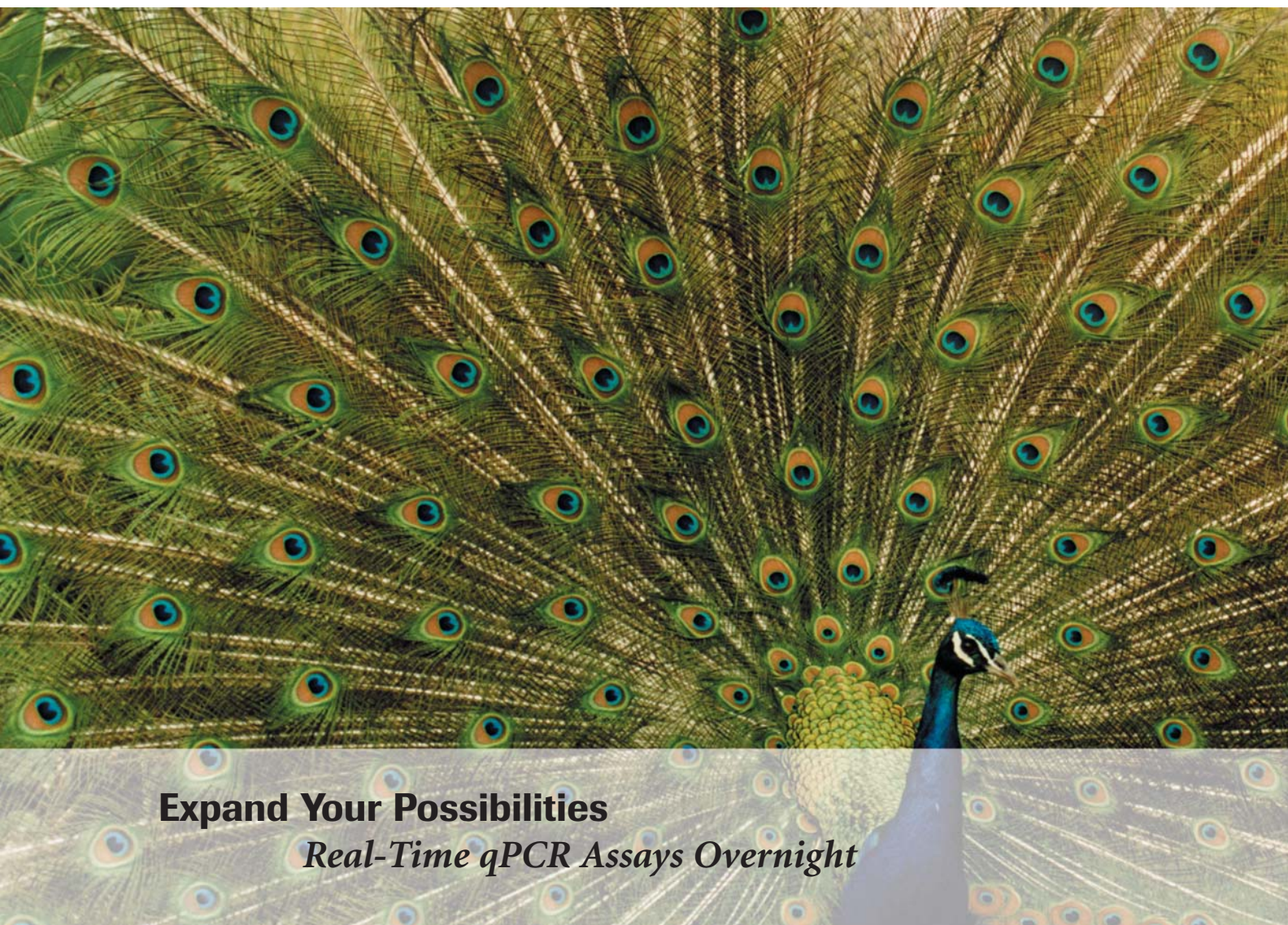


*Roche Applied Science*  
**Universal ProbeLibrary**



**Expand Your Possibilities**  
*Real-Time qPCR Assays Overnight*

## The Universal ProbeLibrary

*Design a gene-expression assay and quantify virtually any transcript in a genome in only two days by using the unique combination of rapid, online assay design and prevalidated probes. Choose from seven transcriptome-specific sets of 90 probes; individual probes are also available separately.*

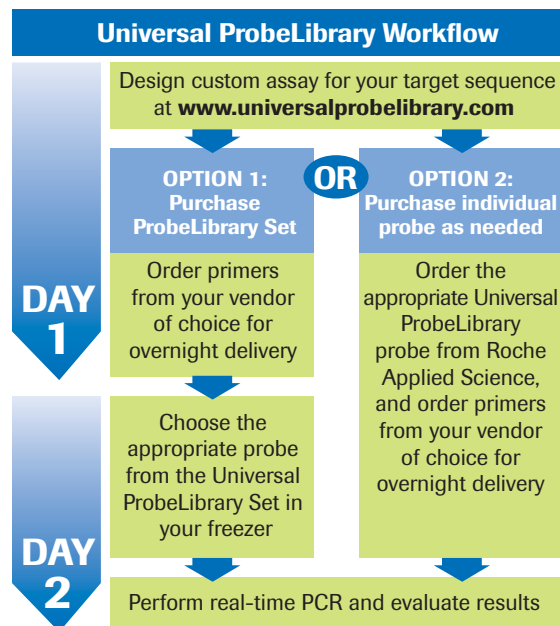
**Significantly reduce assay design time** – in just seconds, design specific, intron-spanning assays for multiple organisms: human, mouse, rat, *Drosophila*, *Arabidopsis*, *C. elegans*, and primates.

**Store a complete set of prevalidated, specific probes in your freezer** – there is no need to wait weeks for an assay or custom probe.

**Attain hydrolysis probe specificity at prices comparable to SYBR Green I** – without sacrificing flexibility.

**Utilize standard protocols on any real-time PCR instrument** – no special hardware or unique reaction conditions are required.

**Obtain consistent results in gene-expression applications** – probes can be used in microarray validation, gene-knockdown quantification assays, and more.



### Generate over Two Million Real-Time qPCR Assays with the Universal ProbeLibrary

The unique combination of software and 165 prevalidated probes enables the design of over two million real-time PCR assays (Table 1). Over one-half million of these are intron-spanning assays. Achieve 96% or greater assay design success, and save time by minimizing assay redesign.

	Intron-Spanning Assays Available	Total Assays Available	Universal ProbeLibrary Coverage
Human	>155,000	>639,500	99%
Mouse	>140,400	>509,500	99%
Rat	>128,000	>364,000	98%
Primates	>174,500	>519,500	96%
<i>Drosophila</i>	>65,500	>253,500	99%
<i>Arabidopsis</i>	>59,500	>199,000	98%
<i>C. elegans</i>	>38,500	>134,000	95%
<b>TOTAL</b>	<b>&gt;761,400</b>	<b>&gt;2,619,000</b>	

▲ Table 1: Number of real-time qPCR assays available using the Universal ProbeLibrary, and total transcriptome coverage for each organism.



## Experimental Data using Various Quant

### Redefine and Revolutionize your Real-Time qPCR Assays with the Universal ProbeLibrary

Sets of 90 prevalidated hydrolysis probes allow you to quantify virtually any transcript of the human, mouse, rat, primates, *Drosophila*, *C. elegans*, or *Arabidopsis* transcriptomes. Transcripts from many other organisms can also be detected with high probability.

The Universal ProbeLibrary utilizes short hydrolysis probes of only 8-9 bases. Specificity and high melting temperature are retained by using Locked Nucleic Acid (LNA) nucleotide chemistry.

Each probe can hybridize to over 7,000 transcripts; on average, each transcript contains binding sites for up to 19 different probes. As a result, several intron-spanning assays can be designed for the same gene.

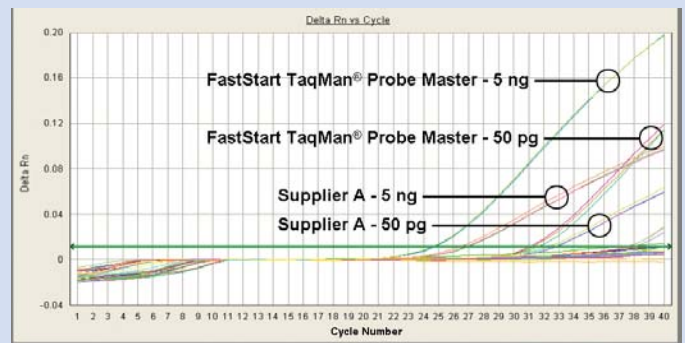
Assay specificity is attained through the combination of specific primers and the Universal ProbeLibrary probes.



### Perform Universal ProbeLibrary Assays on any Real-Time PCR Platform

For optimal results when performing Universal ProbeLibrary assays on thermal block-based real-time PCR instruments, use the **FastStart TaqMan® Probe Master** (Figure 1).

This 2x-concentrated master mix enables very sensitive and specific detection and quantification in combination with Universal ProbeLibrary or other hydrolysis probes.

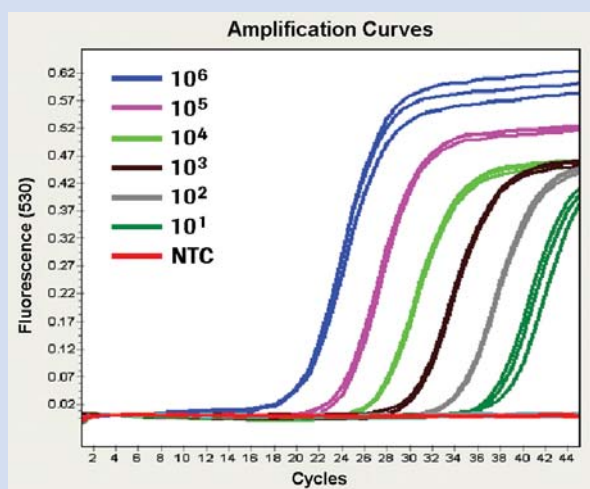


▲ **Figure 1: The Universal ProbeLibrary Software was used to design an assay to detect the GC-rich human Apolipoprotein E gene.** A 90 bp fragment with a GC-content of 60% was selected and used in combination with Probe 72 of the Universal ProbeLibrary Set, Human. A real-time qPCR was performed on an Applied Biosystems 7500 Real-Time PCR System using the FastStart TaqMan® Probe Master and a kit from another supplier. All reactions were performed in duplicate; the ROX reference dye was applied in every reaction. **Result:** The reactions with the **FastStart TaqMan® Probe Master** and Probe 72 from the **Universal ProbeLibrary Set, Human** produced the best sensitivity.

# titative PCR Platforms

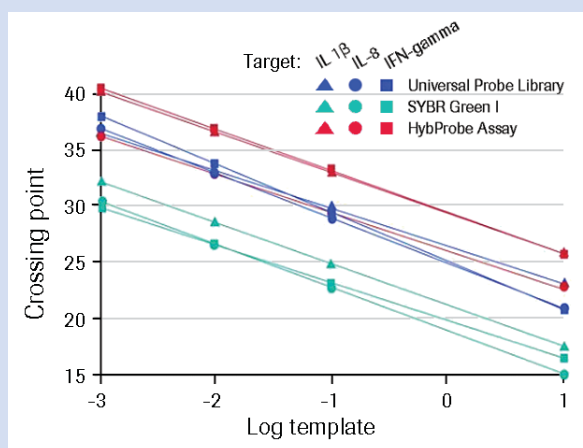
## Perform Universal ProbeLibrary Assays on all Roche LightCycler® Instruments

For Universal ProbeLibrary assays on the LightCycler® 2.0 and 1.5 Instruments, use the **LightCycler® TaqMan® Master** to achieve highest sensitivity combined with the convenience of a ready-to-use master mix (Figure 2).



▲ **Figure 2: Sensitivity down to 10 copies is shown for a Universal ProbeLibrary assay targeting the  $\beta$ -2-microglobulin reference gene.** All samples were run in triplicate. The graph shows 10-fold serial dilutions of *in vitro*-transcribed RNA, ranging from 10<sup>6</sup> to 10<sup>1</sup> copies (NTC = no template control). The assay was performed using the LightCycler® TaqMan® Master qPCR master mix on the LightCycler® 2.0 Instrument.

Comparison of the performance of Universal ProbeLibrary assays with other detection formats on the LightCycler® 2.0 Instrument shows that a high degree of sensitivity, linearity, and fluorescent signal is reproducibly achieved (Figure 3).



▲ **Figure 3: Comparison of linearity of different detection formats.** Assays were run on the LightCycler® 2.0 Instrument. The Universal ProbeLibrary assays were performed using the LightCycler® TaqMan® Master.

**LightCycler® TaqMan® Master** — the choice for laboratories that want to maximize the sensitivity of their LightCycler® 2.0 or 1.5 Instrument.

- Minimize primer-dimer formation with this hot start master mix.
- Save time by eliminating the need for MgCl<sub>2</sub> optimization.
- Achieve high sensitivity — detect as little as 3 pg genomic DNA (~one copy).
- Ensure success with a reagent specifically optimized for LightCycler® 2.0 or 1.5 Instruments.

# Design a Customized Gene-Expression Assay in Three Simple Steps

The starting point for design of a real-time qPCR assay resides in the **Universal ProbeLibrary** Assay Design Center. The core component is the free-access, web-based **ProbeFinder** Software.

Use this software to create customized real-time PCR assays for your gene of interest in seconds.

Product Search

Go

^ Universal ProbeLibrary

Assay Design Center

Home

Human

Mouse

Rat

C.elegans

Drosophila

Arabidopsis

Primates

Probe No. Conversion

Need Help ?

Universal ProbeLibrary for Human

Enter your sequence(s) either by sequence ID or by pasting the sequence:

By sequence ID: e.g. ENST00000158302, NM\_001101 or AB062273

By sequence: e.g. >Human sequence in FASTA format or plain sequence form  
CACGGCAGTCGTCACCAACTGGGACGACATGGAGAAATCTGGCACCACACCTTC  
GAGCTGCTGTGTGGCTCCGAGGAGCACCCTGCTGCTGACCGAGGCCCCCTG  
AAGGCCAACCGGAGAGATGACCCAGATCATGTTTGAGACCTTCAACACCCCA  
TACGTTGCTATCCAGGCTGTGCTATCCCTGTACGGCTCTGGCGGTACCACTGG  
ATCGACTCCGGTACCGGGTCAACCCACACTGTGCCCATCTACGAGGGGTATGGC

☐ Automatic

Next

1

Go to the free, web-based Assay Design Center at [www.universalprobelibrary.com](http://www.universalprobelibrary.com)

Select the organism of interest and enter the target-gene name, accession number, or nucleotide sequence; click “Next”.

2

Within seconds, the ProbeFinder Software will display a custom qPCR assay for your target gene. The output includes the primer sequences and the number of the appropriate Universal ProbeLibrary probe. Complete sequence information is provided for both primers and probe.

Use Universal ProbeLibrary probe: #6, cat.no. 04685032001  
(Formerly Exon ProbeLibrary probe: Human#06)

Primer	Length	Position	Tm	%GC	Sequence
Left Primer	20	126 - 145	60	55	gaaccaatctcaccgacagg
Right Primer	20	173 - 192	60	55	gccacccgagtgtaaccata

Amplicon (67 nt)

gaaccaatctcaccgacaggagctggcgagaggaaataacctgtaccgctatgggttacactcgggtggc

Download pack insert

Download report

Order probes or set

gaaccaatctcaccgacagg

gcaggaga

ataccaaatgtgagccacccg

ggagacc

tgagaaacaaatctcaccgacaggcagc

tgccagaggaatacc

gtaccgc

tatgggttacac

tcgggtggcagagatgcgt

116

798

202

3

Select “More Assays” to view all potential assays designed for your target gene. This function allows the rapid and simple creation of multiple real-time PCR assays for the same gene.

87  
53  
27  
53  
6

79 61 43 24 20 20 77 43

1 2334

<< Left

Zoom In <

Zoom Out >

Right >>

New

Overview

Print

LibraryProbe	Rank	Position	Intron spanning	InSilicoPCR test	Score	Class
<a href="#">#6(Human#06)</a>	1	153	+	+		Perfect
<a href="#">#53(Human#53)</a>	2	150	+	+		Perfect
<a href="#">#29(Human#29)</a>	3	826	+	+		Perfect
<a href="#">#43(Human#43)</a>	4	829	+	+		Perfect
<a href="#">#27(Human#27)</a>	5	142	+	+		Perfect
<a href="#">#53(Human#53)</a>	6	189	+	+		Perfect
<a href="#">#87(Human#87)</a>	7	187	+	+		Good
<a href="#">#80(Human#80)</a>	8	113	+	+		Good
<a href="#">#43(Human#43)</a>	9	1420	+	+		Good
<a href="#">#77(Human#77)</a>	10	1380	+	+		Good
<a href="#">#37(Human#37)</a>	11	777	+	+		Good





## Assay Design with the Universal ProbeLibrary Product Family

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

For more information about the Universal ProbeLibrary, visit [www.universalprobelibrary.com](http://www.universalprobelibrary.com) or contact your local Roche representative.

Product	Cat. No.	Pack Size
LightCycler® TaqMan® Master	04 535 286 001	1 kit for 96 reactions

Product	Cat. No.	Pack Size
FastStart TaqMan® Probe Master (Rox) <sup>†§</sup>	04 673 450 001	2.5 ml (2 x 1.25 ml) for 100 (50 µl) reactions
	04 673 468 001	12.5 ml (10 x 1.25 ml) for 500 (50 µl) reactions
	04 673 476 001	50 ml (10 x 5 ml) for 2,000 (50 µl) reactions
FastStart TaqMan® Probe Master <sup>†§</sup>	04 673 409 001	2.5 ml (2 x 1.25 ml) for 100 (50 µl) reactions
	04 673 417 001	12.5 ml (10 x 1.25 ml) for 500 (50 µl) reactions
	04 673 433 001	50 ml (10 x 5 ml) for 2,000 (50 µl) reactions

Other brands or product names are trademarks of their respective holders.

§ No rights for any other application, including any in vitro diagnostic application, are conveyed expressly, by implication or by estoppel under patents owned by Roche Molecular Systems, Inc. and F. Hoffmann-La Roche Ltd covering the 5' nuclease detection technology.



**www.roche-applied-science.com**