

# ProbeLibrary

## “One Transcriptome - One Kit”

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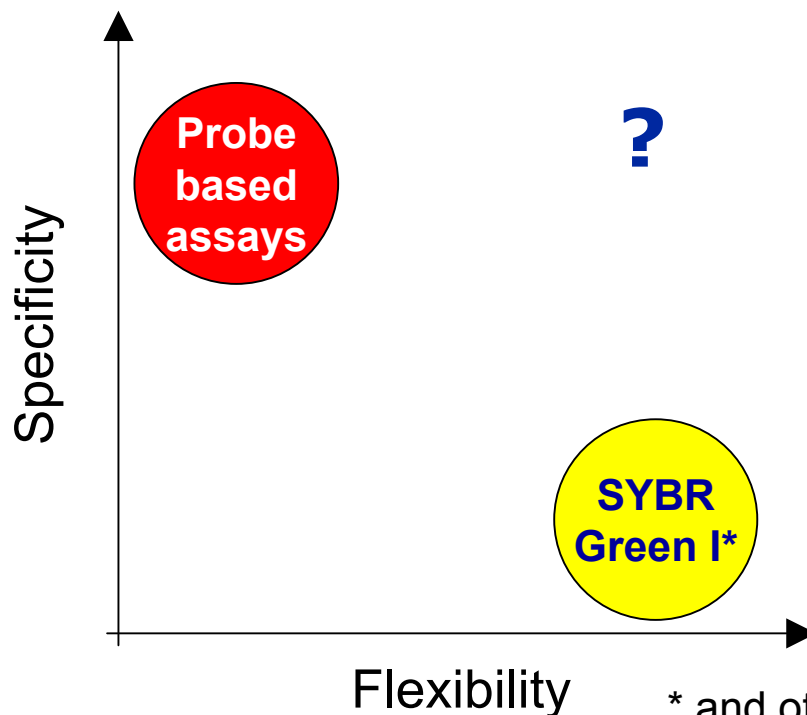
*qPCR-Symposium (Leipzig)*

*10.03.2005*

# Problem in real-time PCR: Specificity or flexibility ?

Today you have to decide whether using a specific probe-based assay or a flexible SYBR Green I assay !

**What about having both characteristics together in one assay ?**



\* and other intercalating dye based assays

# How to solve the problem of combining flexibility and specificity ?

## Use the ProbeLibrary\*

- Kit with 90 probes and 250 rxns (50 µl) each
- Very short dual-labeled probes (8 and 9 mers)
- Probes are modified with LNA to ensure compatibility with standard real-time PCR methods
- Each transcript matches with ~ 16 probes, each probe matches with ~ 7.000 transcripts
- Specificity of assays through specific primers and assisted by probes

\* ProbeLibrary is a trademark of Exiqon A/S

# Advantages of the ProbeLibrary against other detection formats

- **Advantage against SYBR Green I:**
  - No primer dimers or side products detectable due to detection with specific probes
  
- **Advantage against probe based assays:**
  - Time to result within 2 days (instead of 1 week)
  - Less expensive than regular probe assays
  
- **Customer workflow:**
  - Order kit
  - Design assay with ProbeFinder design software via web
  - Order primers at oligohouse
  - Run experiment

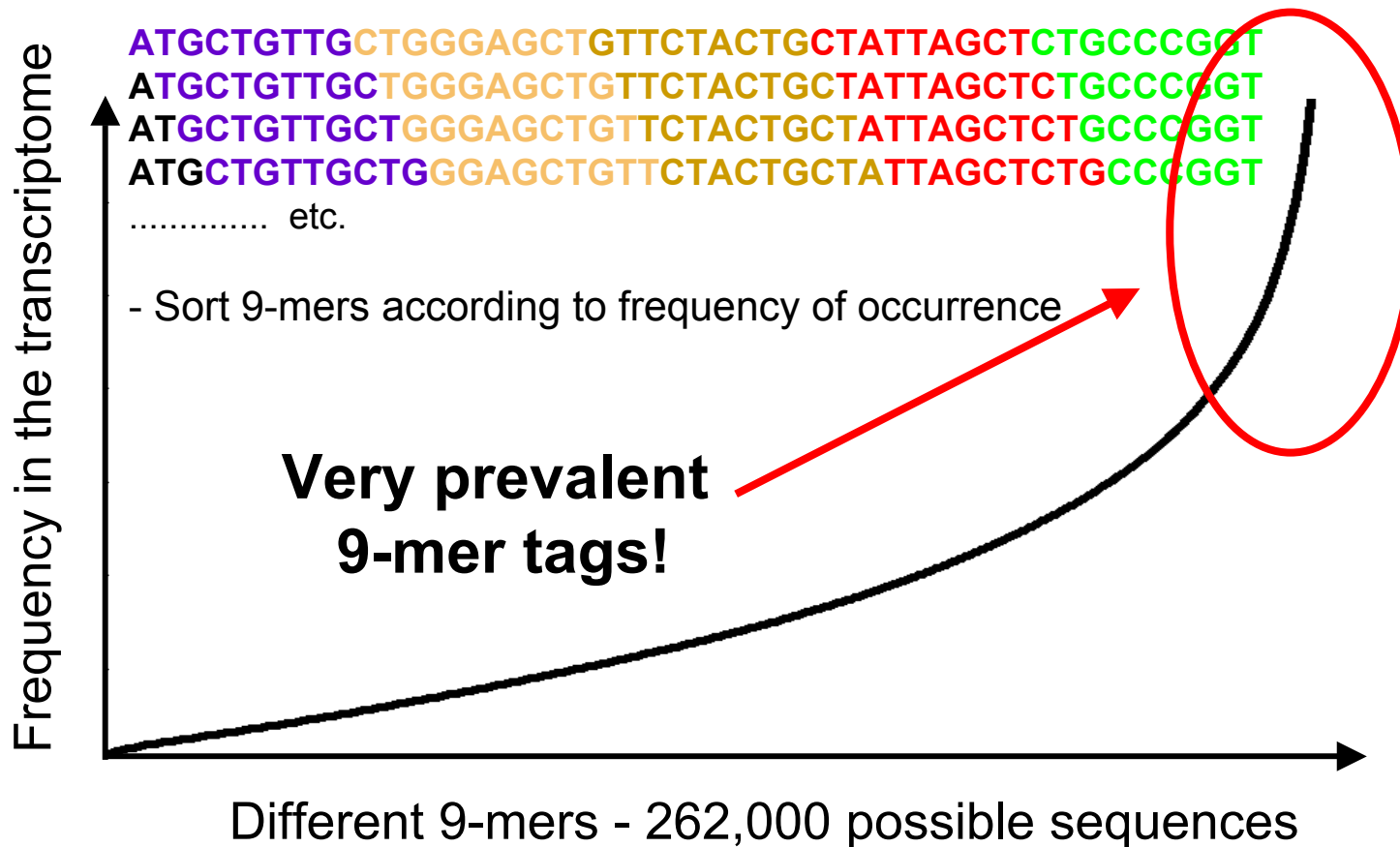
# What are the properties of the ProbeLibrary ?



- **Real-time PCR assays for Expression Profiling**
- **Complete transcriptome of one organism is covered by 98%**
- **Assay design is very fast and easy with web-based ProbeFinder design software**
- **Organisms available so far are human, mouse, rat, arabidopsis, drosophila and C. elegans**
- **The product is developed and already launched by Exiqon since May 2004**

# Why using 9-mers as probes ?

## Occurrence of 9-mers in human transcripts



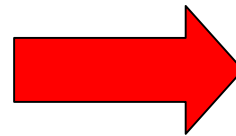
# 90 probes cover entire transcriptome

Bioinformatic analysis of the genome reveals:

- Some 8 & 9 mer motifs are very prevalent in the transcriptome
- The same probes can be used for the same motifs in several genes

**Only 90 pre-validated probes are sufficient to cover the entire human transcriptome !**

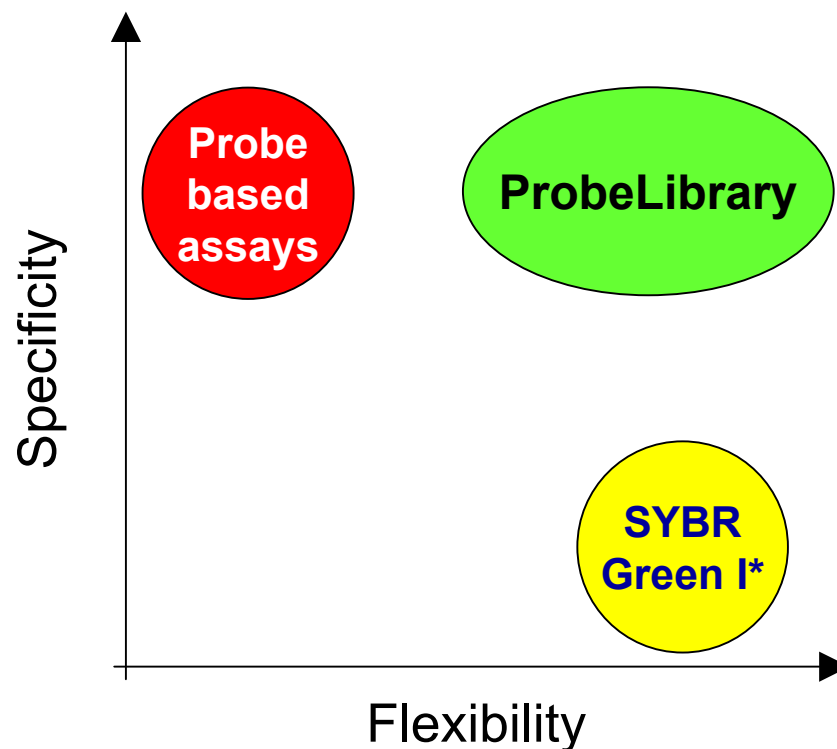
## ProbeLibrary



# ProbeLibrary: High flexibility and specificity

## Just 90 Probes but ...

- 99 % of mRNA transcripts in RefSeqHum are targeted
- 90 % of Ensembl mRNA transcripts are targeted with intron spanning assays
- Targeting 999 of the 1000 most cited human genes in RefSeq db with at least one assay



\* and other intercalating dye based assays



# Workflow of the ProbeLibrary: From assay design to results within 2 days



Day 1

Identify gene or target sequence of interest

Identify probes and primers at [www.probelibrary.com](http://www.probelibrary.com)

Order primers from your preferred oligo supplier for overnight delivery

Day 2

Select the relevant probe from the ProbeLibrary

Perform real-time PCR

Evaluate results

# Assay design, step 1: Choose an organism



## Real-Time PCR Assay Design Center

The Assay Design Center allows you to fully benefit from the power of the ProbeLibrary, by quickly and easily designing your real-time PCR assays.

The core component of the Assay Design Center is the ProbeFinder software: a free-access, web-based tool that provides you with clear and detailed real-time PCR assay designs for your genes of choice.

- **Save Time**
  - Design your assays in seconds
- **Simplicity**
  - Just enter your target gene and you will get optimal design of your real-time PCR primers and a ProbeLibrary probe selection
- **Design Assays You Can Trust**
  - *In silico* PCR feature
  - Intron-spanning options
  - 96% assay design success rate

Start the Assay Design Software by choosing an organism:



# Assay design, step 2: Enter your sequence / gene of interest



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Human Mouse Rat Arabidopsis Drosophila C.elegans

**This is ProbeFinder for the Human ProbeLibrary**

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

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

**Option 1: Enter sequence ID**

By sequence: e.g. Human sequence in FASTA format or plain sequence format

```
CACGGCATCGTCACCAACTGGGACGACATGGAGAAAATCTGGCACCACACCTTCTACAAT
GAGCTGCGTGTGGCTCCCGAGGAGCACCCCGTGCTGCTGACCGAGGCCCCCTGAACCCC
AAGGCCAACCGCGAGAAGATGACCCAGATCATGTTTGAGACCTTCAACACCCAGCCATG
TACGTTGCTATCCAGGCTGTGCTATCCCTGTACGCCTCTGGCCGTACCACTGGCATCGTG
ATGGACTCGGGTACCGGGTACCCACACATGAGCCATCTACGAGGGGTATGCCCTCCCC
```

**Option 2: Paste sequence**

Automatically select an [intron spanning assay](#).

Next >>

# Assay design, step 3: Output information of optimal assay



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Human Mouse Rat Arabidopsis Drosophila C.elegans

This is ProbeFinder for the Human ProbeLibrary

ProbeFinder has designed the optimal real-time PCR assay for:

NM\_001101.2 Homo sapiens actin, beta (ACTB), mRNA.

Assay details:

Use ProbeLibrary probe: Human#64

Primer	Length	Position	Tm	%GC	Sequence
Left Primer	18	414 - 431	60	56	ccaaccgcgagaagatga
Right Primer	20	491 - 510	59	60	ccagaggcgtacagggatag
Amplicon (97 nt)					
ccaaccgcgagaagatgaccagatcatgtttgagaccttcaacaccccagccatgtacg ttgctatccaggctgtgctatccctgtacgacctctgg					

Primer information  
(sequence, Tm, %GC)

Amplicon information

Download report Order Probes or Kit

# Assay design, step 4 (optional): Choose another assay from the list

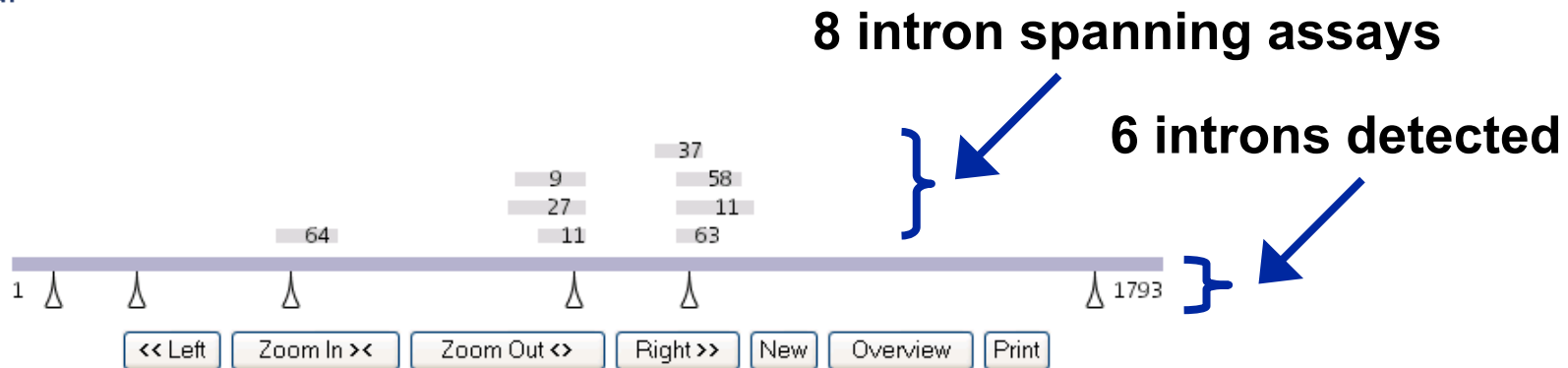


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**Human** | Mouse | Rat | Arabidopsis | Drosophila | C.elegans

Additional good real-time PCR assays designed for:

[NM\\_001101.2](#) Homo sapiens actin, beta (ACTB), mRNA.

Transcript overview:



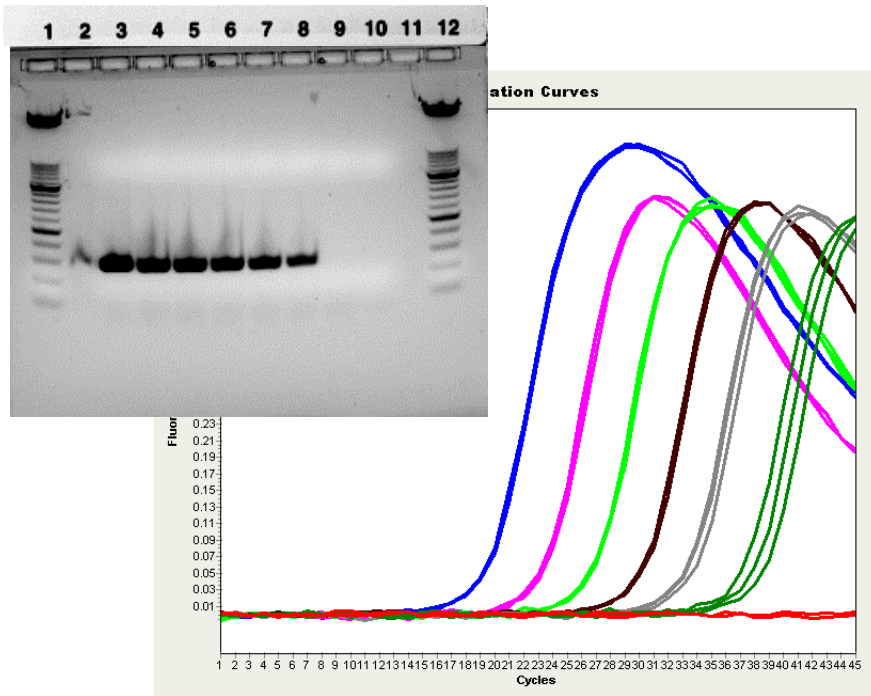
[Figure legend ?](#)

LibraryProbe	Rank	Position	Intron spanning	InSilicoPCR test	Score	Class
Human#64	1	481	+	+		Good
Human#11	2	856	+	+		Good
Human#27	3	796	+	+		Good
Human#09	4	814	+	+		Good
Human#63	5	1065	+	+		Good

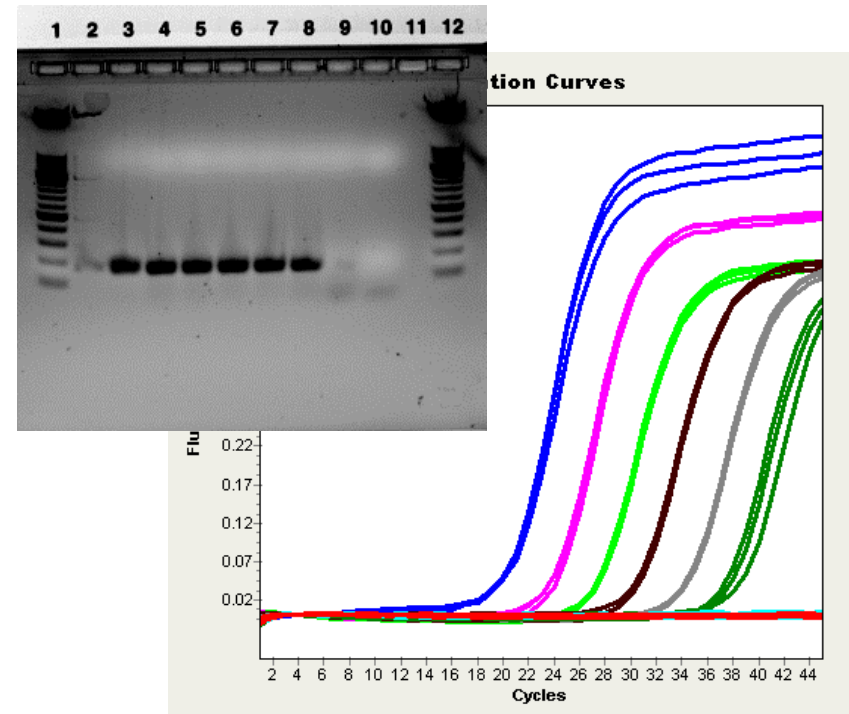
# Comparison HybProbe\* vs ProbeLibrary assays



## HybProbe\* assay



## ProbeLibrary assay

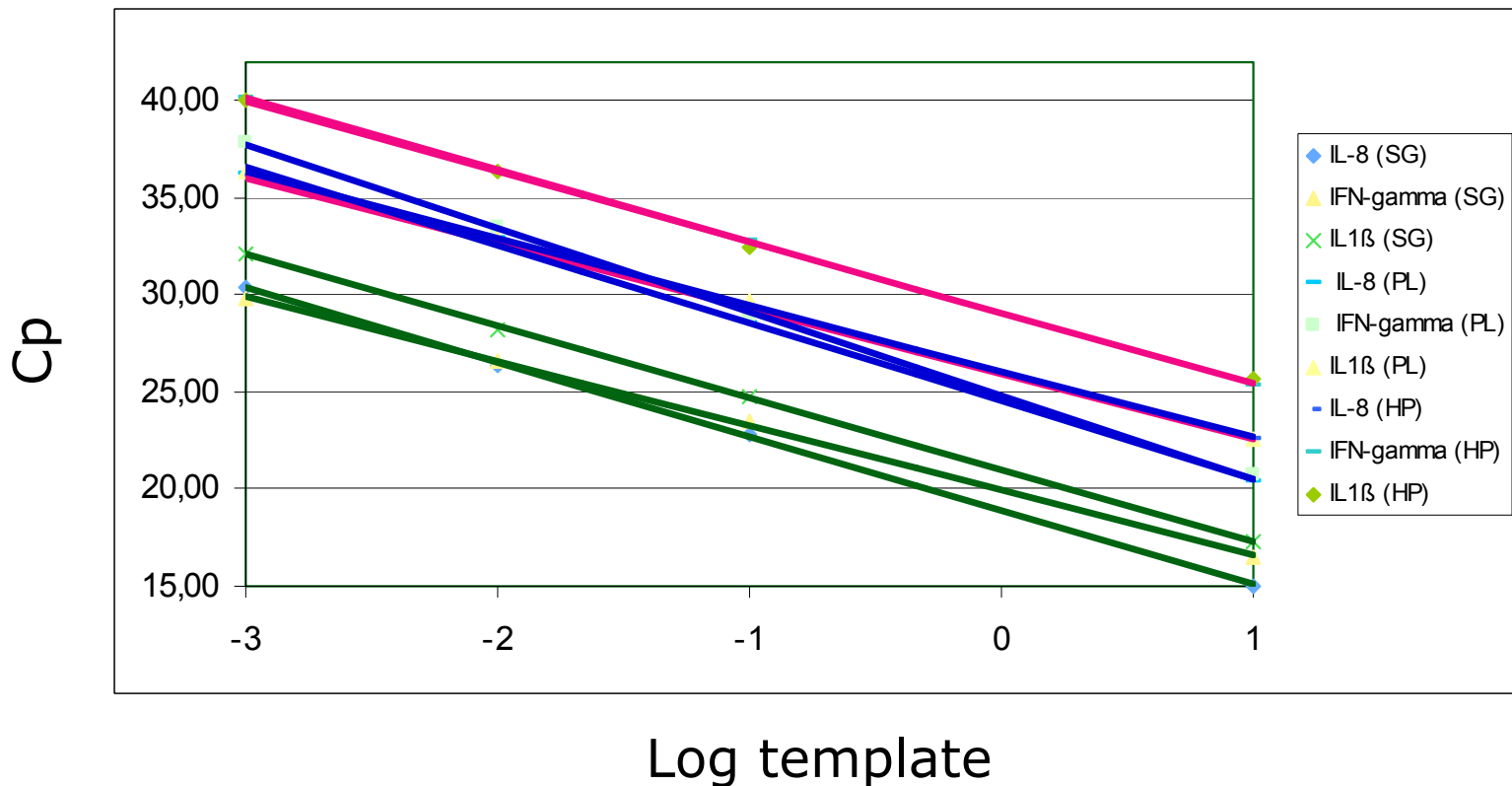


Target:  $\beta$ 2M; Template: In-vitro RNA-Transcript; Dilution series of  $10^6$  (blue line, slot3),  $10^5$  (pink line, slot4), ...,  $10^0$  (red line, slot 9) copies of transcript, negativ control (NTC, turquoise), all samples in triplicates; Instrument: LightCycler 2.0; PCR Kit: LightCycler TaqMan Master

\* HybProbe is a trademark of a Member of the Roche Group

# Comparison of different detection formates due to linearity - 1

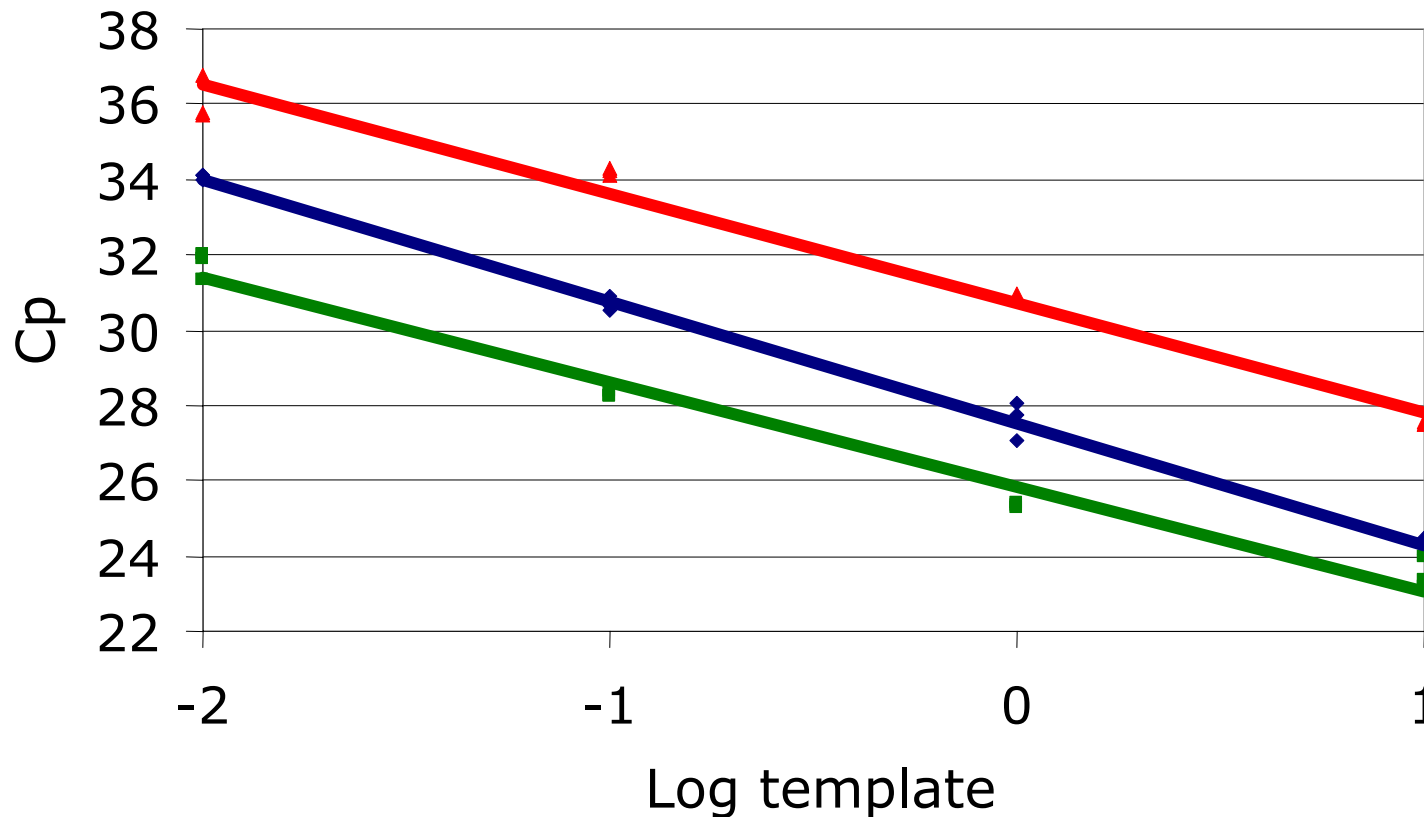
◆ ProbeLibrary      ■ SYBR Green I      ▲ HybProbe assay



3 assays (IL-8, IL1β, IFN-gamma) for each detection formate

# Comparison of different detection formats due to linearity - 2

◆ ProbeLibrary    ■ SYBR Green I    ▲ Pre-validated assay





# Conclusion 1



**What is the benefit by using the ProbeLibrary:**

- **Custom real-time PCR assays**
- **Flexible and specific assays**
- **Fast and easy assay design by web-based ProbeFinder software**
- **Faster time to result (Design one day, run PCR the next day)**
- **Probes are compatible with any real-time PCR instrument**
- **Cost sensitive assays in the range of SYBR Green I assays**

# Conclusion 2



## Performance:

- **Equal sensitivity of ProbeLibrary against SYBR Green I and other probe-based assays**
- **Advantage against SYBR Green I: No primer dimers or side products detectable due to detection with specific probes**
- **Advantage against probe based assays: Time to result within 2 days (instead of 1 week), less expensive than regular probe assays**
- **Reproducible and reliable results**

# Take home message



- You can quantify just about any human gene using only 90 pre-validated real-time PCR probes
- You can actually design assays one day and run them the next
- You can save money as costs for ProbeLibrary assays are similar to SYBR Green I assays

*This is what we call:  
One Transcriptome – One Kit  
or just:  
The ProbeLibrary*