

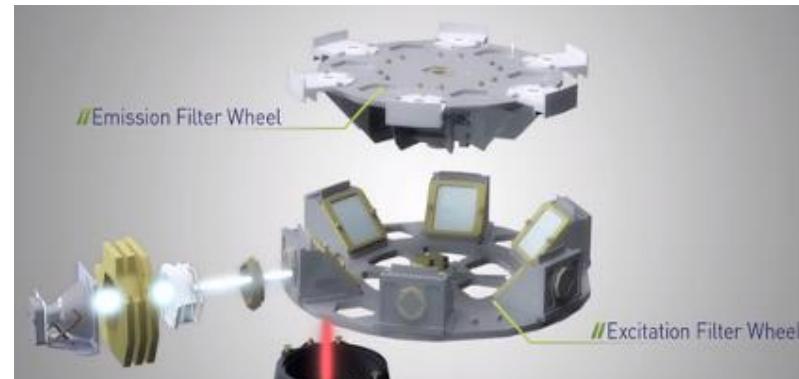


QuantStudio qPCR System Overview

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Field Applications Scientist

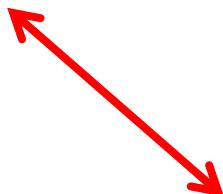
The Optics: Enhanced OptiFlex™ System

- light source
- CCD Camera captures data of all filters at every cycle



| | m1(520±15) | m2(558±11) | m3(586±10) | m4(623±14) | m5(682±14) | m6(711±12) |
|------------|-------------------------------------|--------------------------|--------------------------|--------------------------|--------------------------|--------------------------|
| x1(470±15) | <input checked="" type="checkbox"/> | <input type="checkbox"/> |
| x2(520±10) | | <input type="checkbox"/> |
| x3(550±11) | | | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> |
| x4(580±10) | | | | <input type="checkbox"/> | <input type="checkbox"/> | <input type="checkbox"/> |
| x5(640±10) | | | | | <input type="checkbox"/> | <input type="checkbox"/> |
| x6(662±10) | | | | | | <input type="checkbox"/> |

QuantStudio Connectivity



USB

Thermo Fisher Connect™



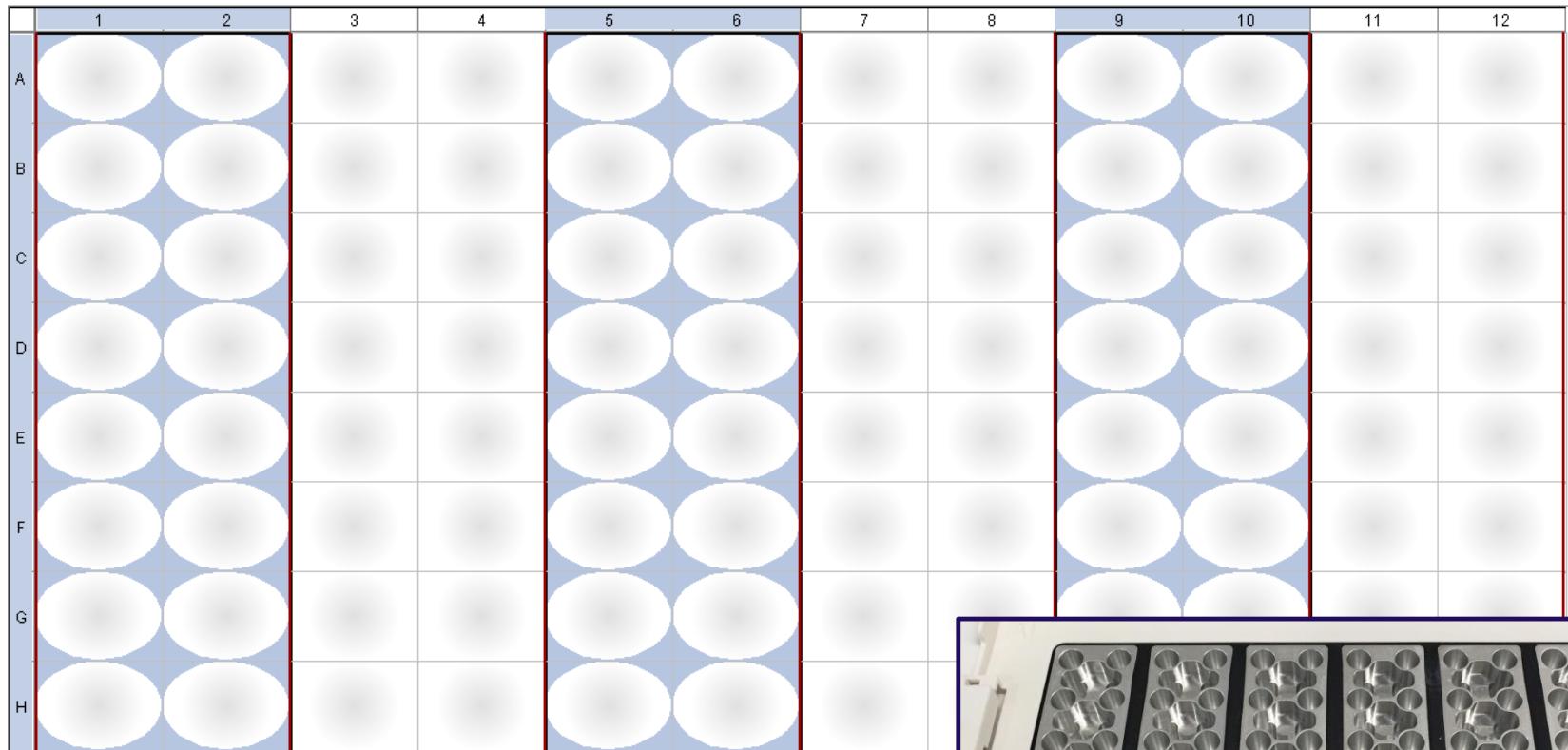
PC or Mac®
Connect with a web
browser

Connect to your data anytime, anywhere

ThermoFisher
SCIENTIFIC

VeriFlex™ Block technology

6 independently controlled programmable zones ($\leq 5^{\circ}\text{C}$ of adjacent zone)

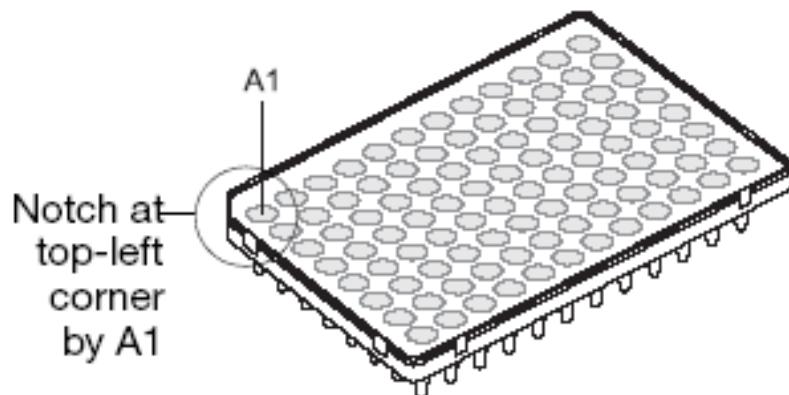


| Zone 1 | Zone 2 | Zone 3 | Zone 4 |
|--------|--------|--------|--------|
| 55.0 | 57.0 | 59.0 | 61.0 |

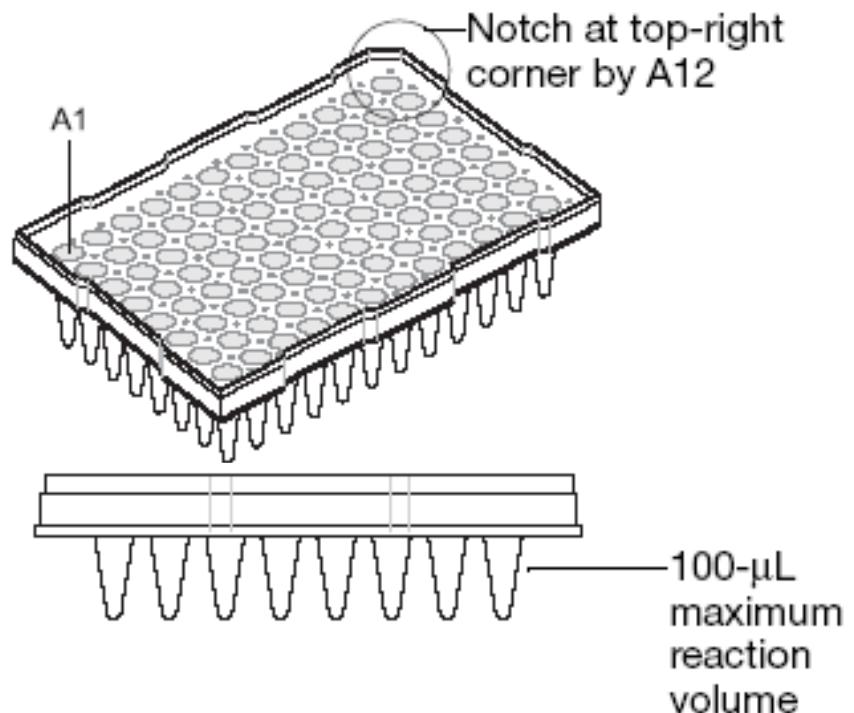


96 well

Fast 0.1 mL plates



Standard 0.2 mL plates

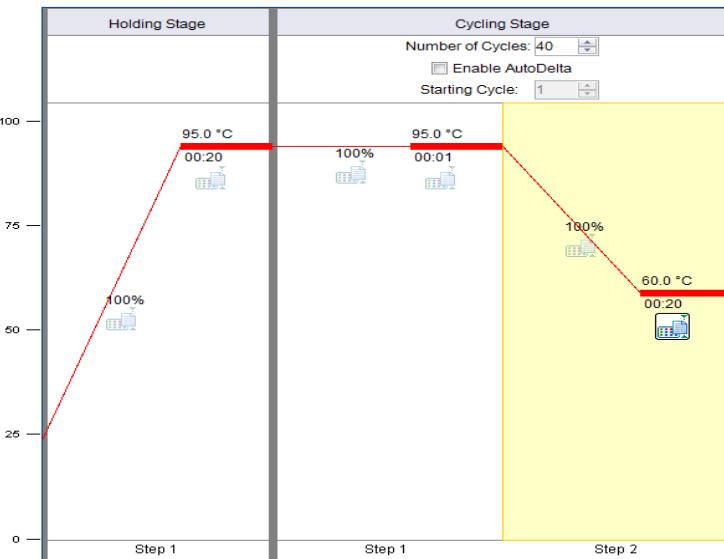


| | <u>0.1mL</u> | <u>both</u> | <u>0.2mL</u> |
|----------------------|--------------|-------------|--------------|
| 96 well plates | 4346906 | | 4306737 |
| 96 well plate holder | | 4312063 | |
| adhesive covers | | 4360954 | |
| tray | 4379983 | | 4381850 |
| 8strip tubes | 4358293 | | 4316567 |
| 8strip tube caps | | 4323032 | |
| Tube Capping tool | | 4330015 | |
| single tubes | 4358297 | | N8010540 |
| Validated RXN volume | 10-30uL | | 10-100uL |

Instrument Default Cycling Options

Fast: 40 cycles in ~40min.

- faster ramping & shorter hold times



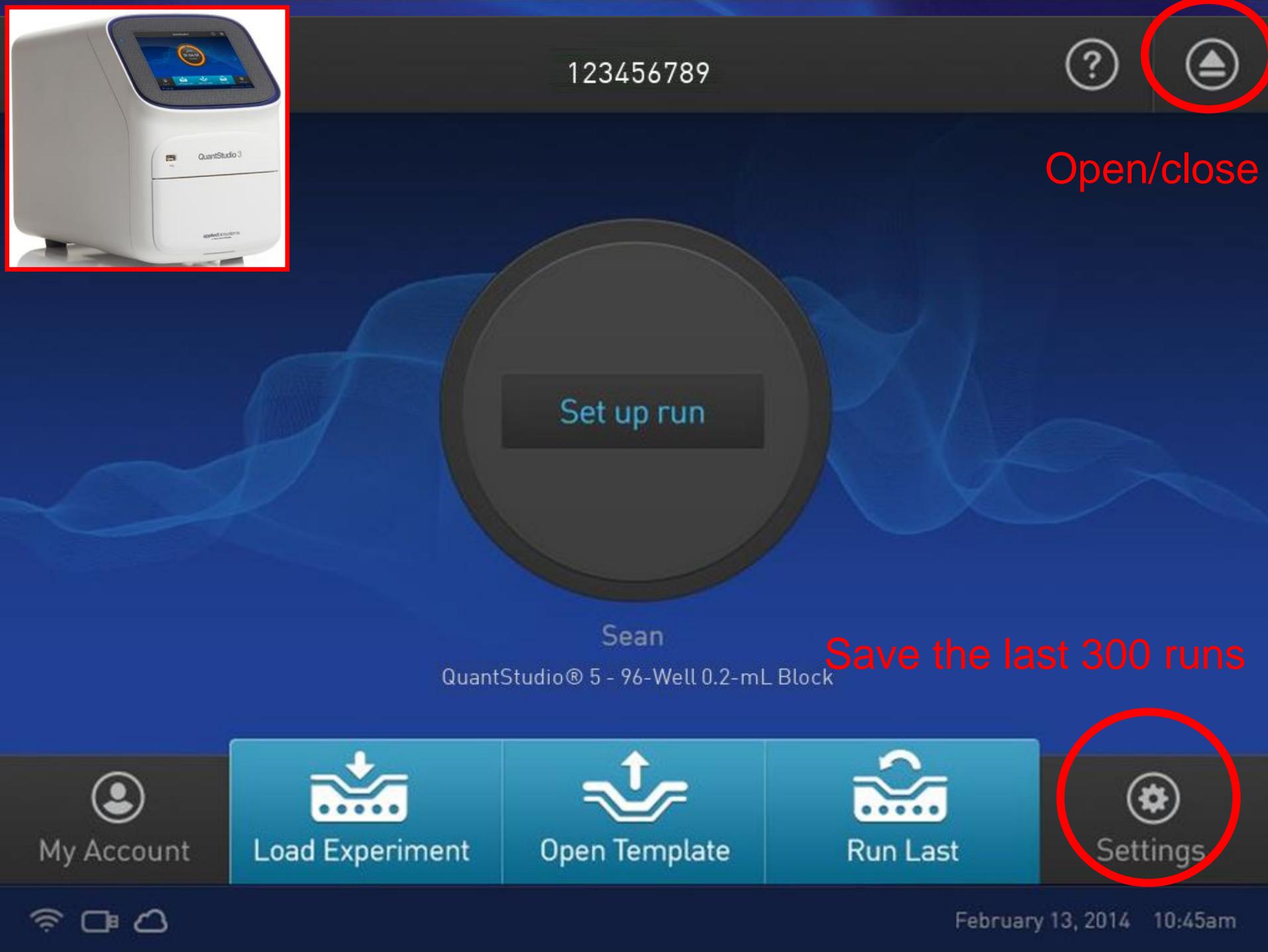
Standard: 40 cycles in ~1hr 45min



TaqMan Fast Advanced Master Mix (TaqPath ProAmp for SNP)

PowerUp SYBR Green Master Mix

There are many master mixes, read the protocol for the master mix!



Recommended Calibration and Verification

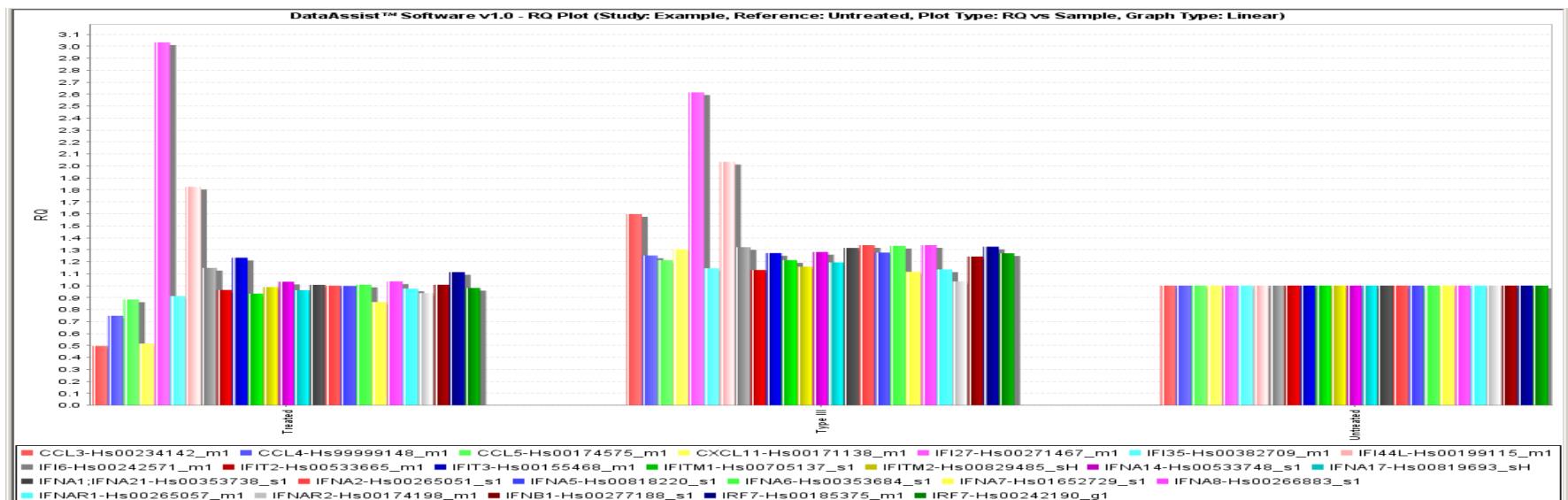
| Calibration | Recommended frequency |
|----------------------|---|
| ROI/uniformity | Every two years <ul style="list-style-type: none">Always perform new Background and Dye calibrations after an ROI/Uniformity calibration. |
| Dye | Every two years <ul style="list-style-type: none">During a Dye calibration, only the dyes on the given spectral calibration plate are calibrated. |
| Background | Every month <ul style="list-style-type: none">As needed: To check for contaminationAs needed: To obtain the most accurate data for the removal of background fluorescence |
| RNase P verification | After installing or moving the instrument. After performing instrument or block calibrations. As needed to confirm instrument performance. |

Thermo Fisher Connect

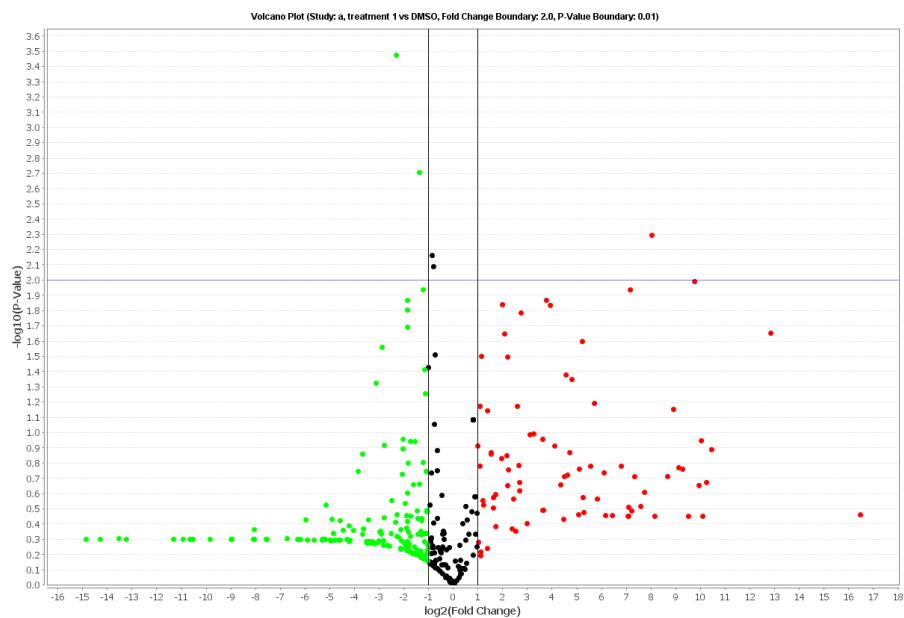
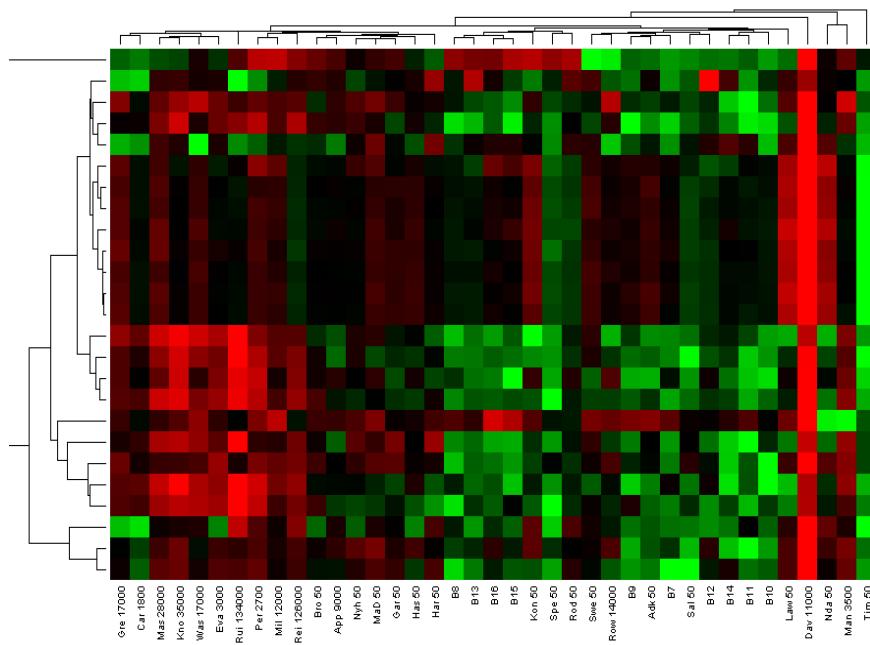
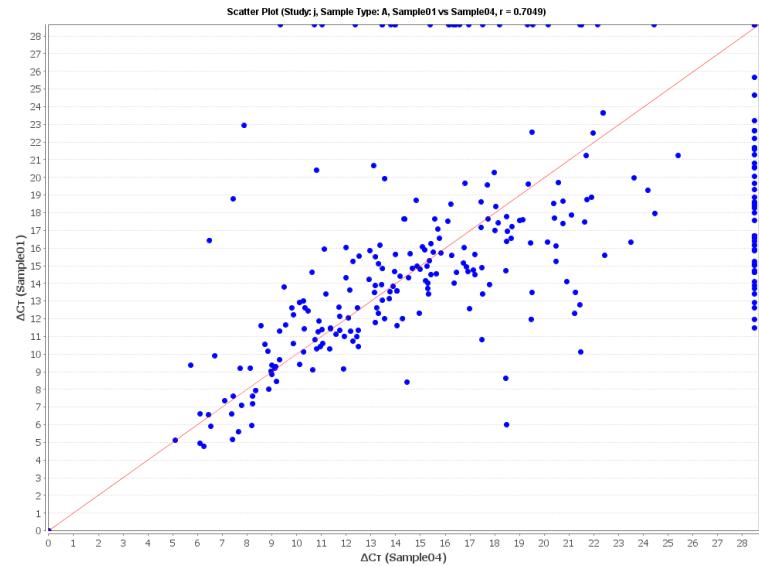
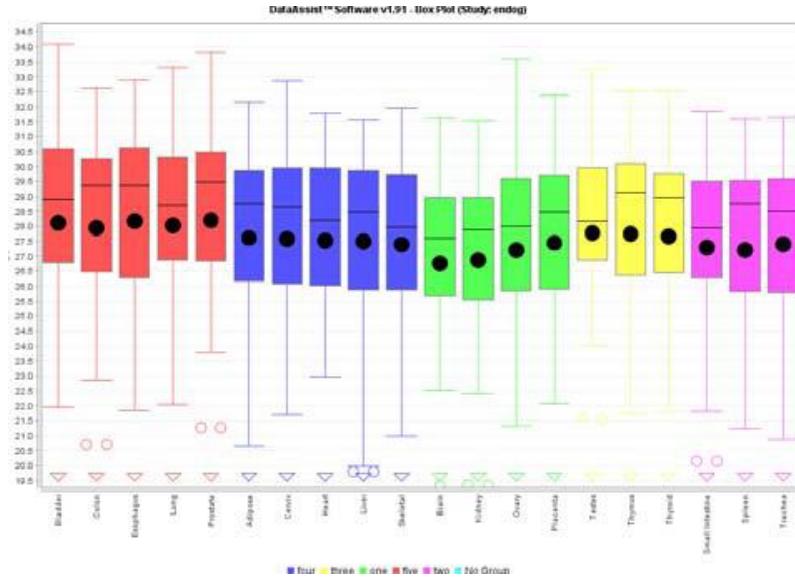
FREE web-based Gene Expression Analysis Software

How it works

- Import .eds data files from system software
- Analyze **>100 plates** in one study
- Graphical representation of the RQ data
- Export Analyzed Results, Data & Plots



Thermo Fisher Connect



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Get your free product, rewards, benefits, and more

Rediscover your
scientific aspirations

aspire
member program



Join the free member program that supports your lifelong love of science.

Enroll today and receive a free, full-size product, 500 points towards rewards and discounts, and access to science career benefits and more...

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qPCR Applications & Information

Gene Expression

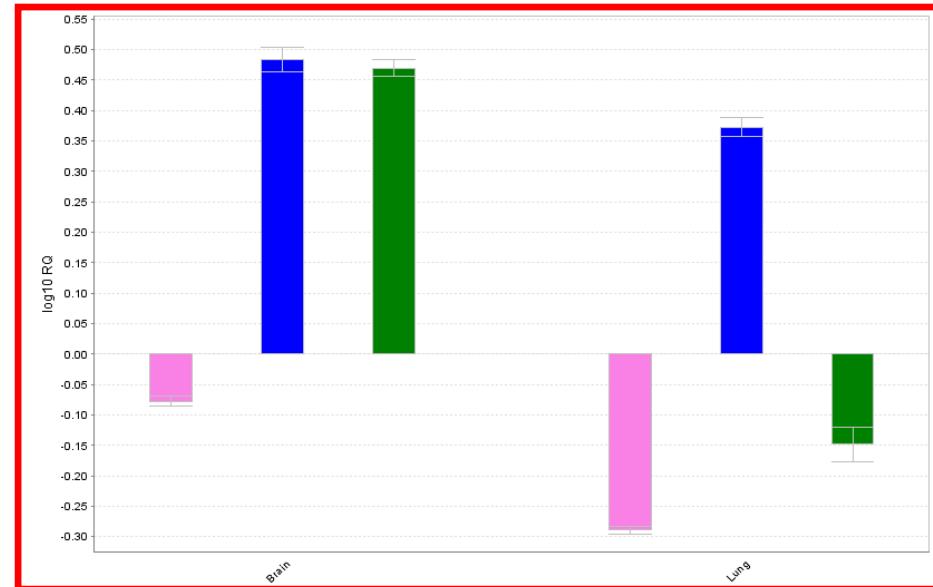
miRNA expression

SNP Genotyping

Copy Number Variation

CAST PCR for Mutation Analysis

Protein Thermal Shift



Digital PCR (QuantStudio 3D)
Service Plans on qPCR systems



Follow-up email with today's slides & survey

qPCR is the Technology to detect amplification of PCR products in real-time

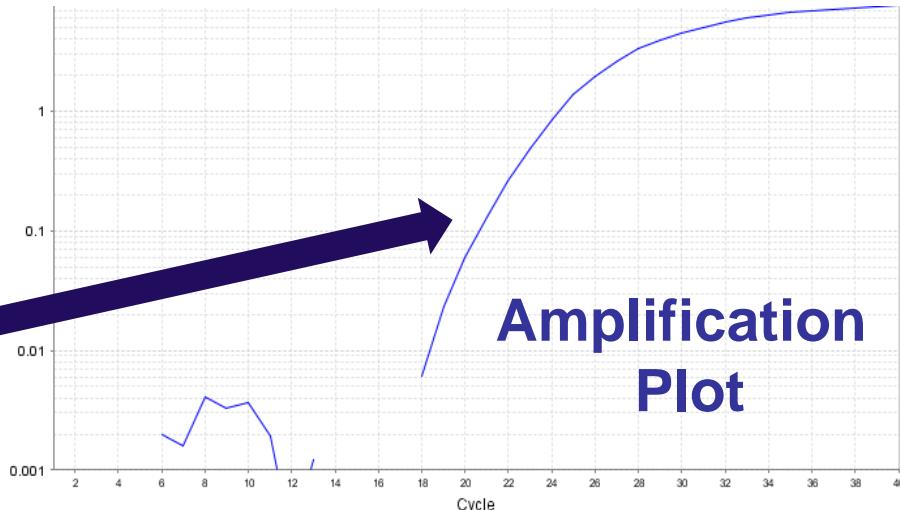
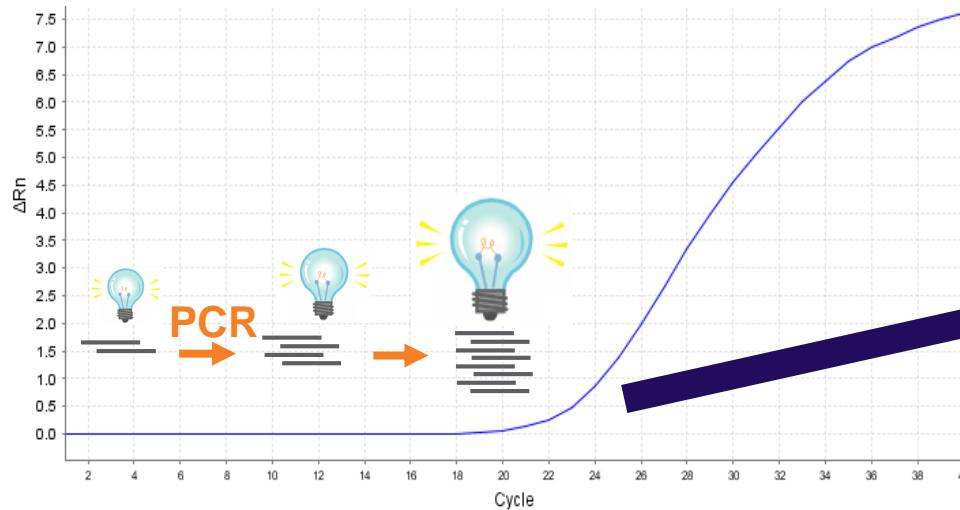
Similar to traditional PCR...



Denature template
Anneal primers
Extension of primers → amplicons
40 cycles

A **fluorescent** dye is in the reaction

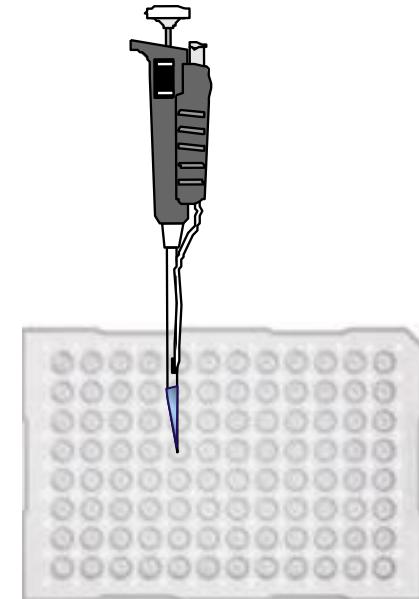
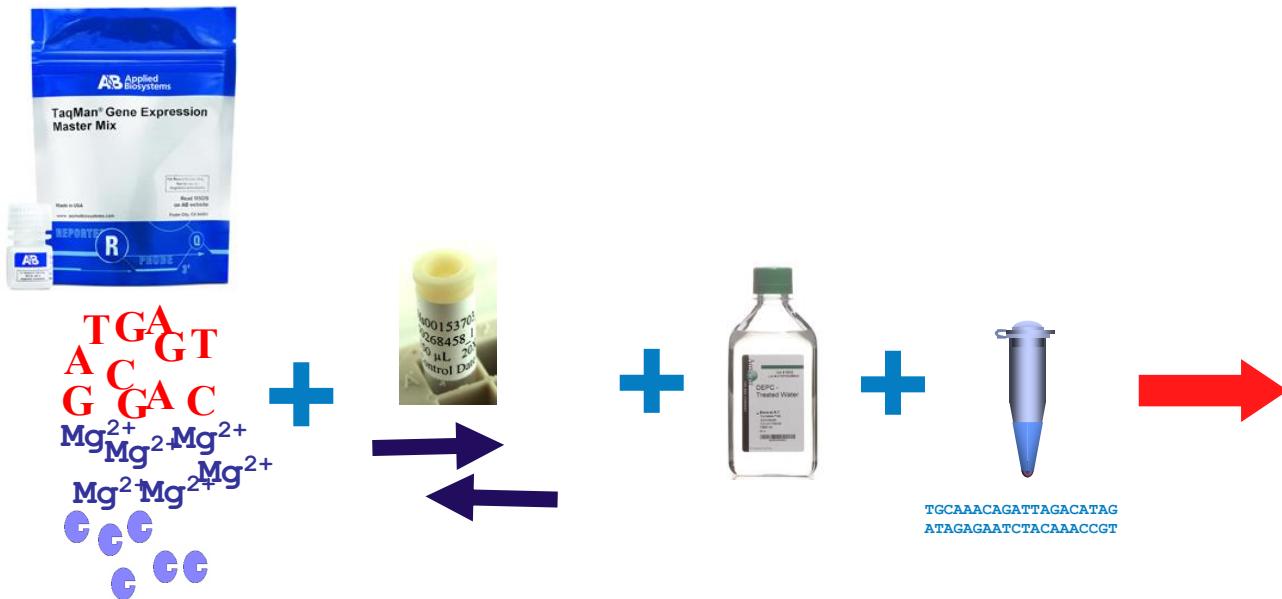
As Amplification proceeds, fluorescence increases with PCR product formation



How to set up Real-Time PCR

Combine:

- | | |
|---|----------------------|
| 1) TaqMan® Master Mix or SYBR® Green Master Mix | <u>20uL reaction</u> |
| 2) TaqMan® Gene Expression Assay or SYBR® primers | (2X) 10uL |
| 3) water (#AM9935) | (20X) 1uL |
| 4) Sample (template) | 7uL |
| | (10 ng) 2uL |



Nucleic Acid Isolation Kits

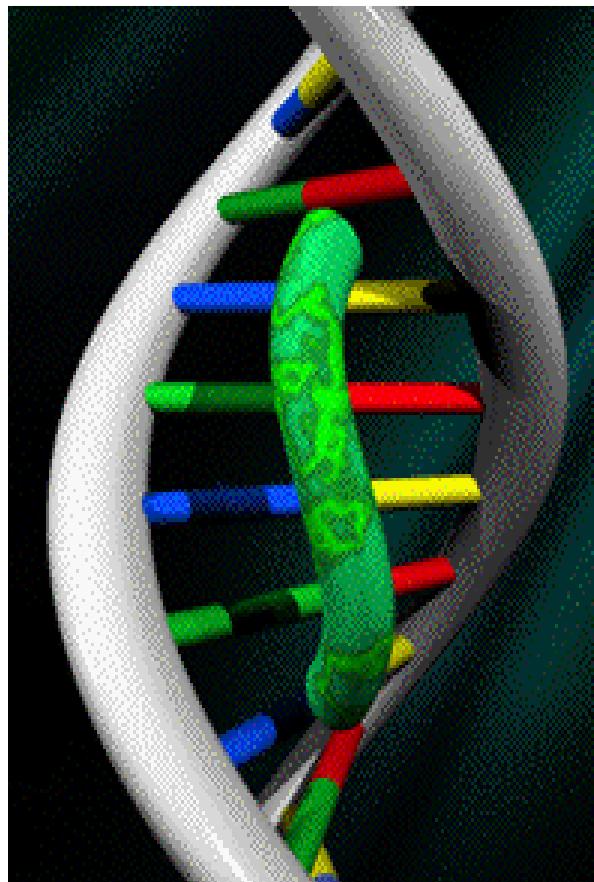
| DNA & RNA Selection Guide | | | | | | |
|---------------------------|---------------------|------------------|---------------|----------------------|--------|--|
| Displaying 7 results. | | | | | | |
| | | | DNA | RNA | | |
| Sample Type | Starting Amount | Throughput | Application | | Format | |
| Cells | less than 1000 | 1-50 | Real-Time | Silica matrix column | | |
| Tissue | $10^3\text{-}10^4$ | 50-100 | RT-PCR | Purification-free | | |
| Blood | $10^4\text{-}10^5$ | greater than 100 | microRNA | Organic Solution | | |
| Virus | greater than 10^5 | | Cloning | Magnetic beads | | |
| Bacteria/Yeast | | | Microarrays | | | |
| Plant | | | Northern Blot | | | |
| FFPE | | | Sequencing | | | |
| LCM | | | messenger RNA | | | |

KingFisher Flex
Up to 96 samples in a run

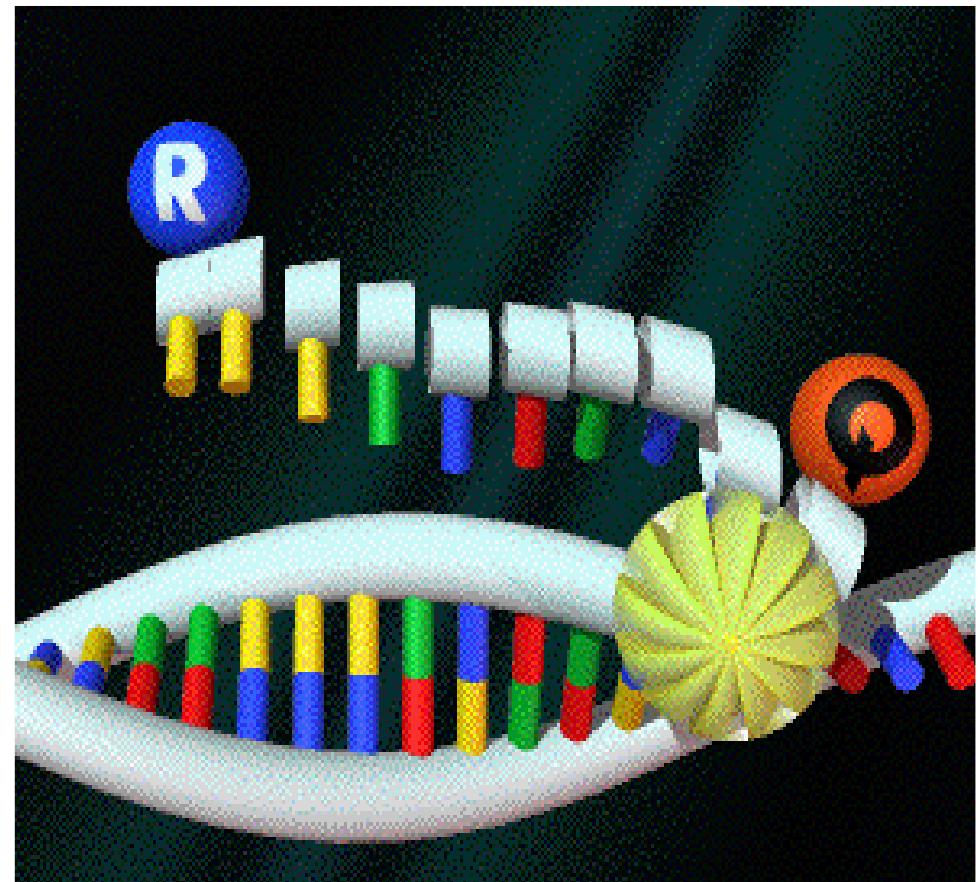


Supported Fluorescent Chemistries

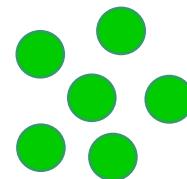
SYBR® Green



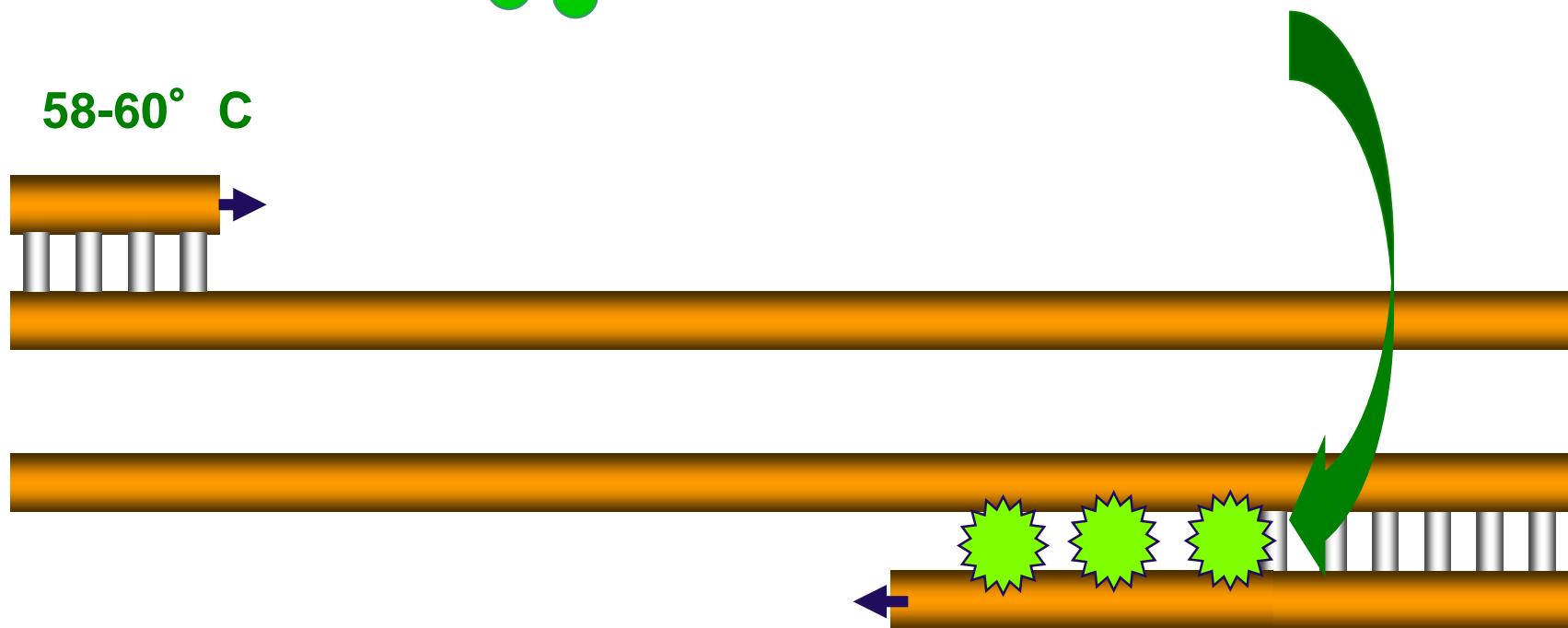
TaqMan®



SYBR® Green Reaction



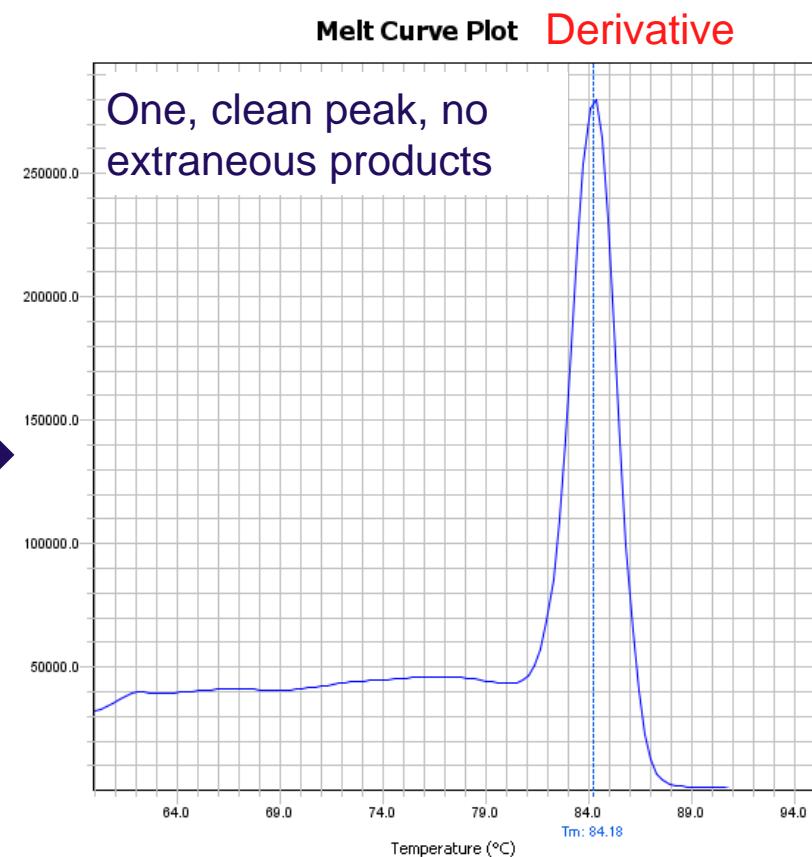
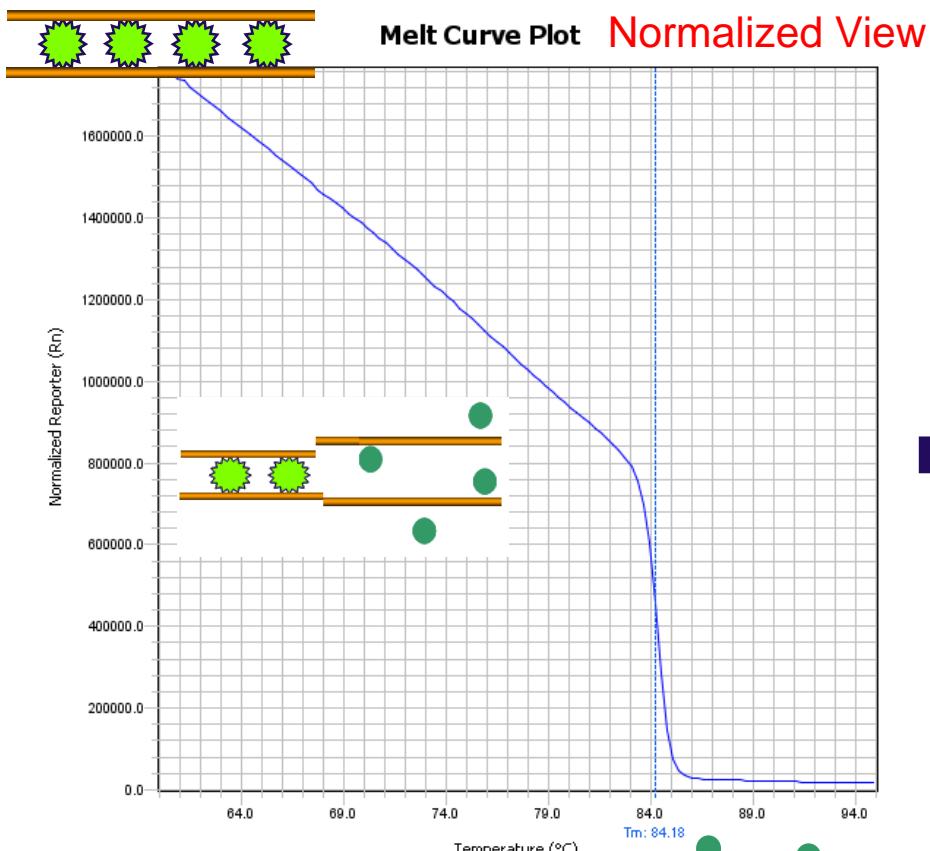
SYBR® Green I Dye



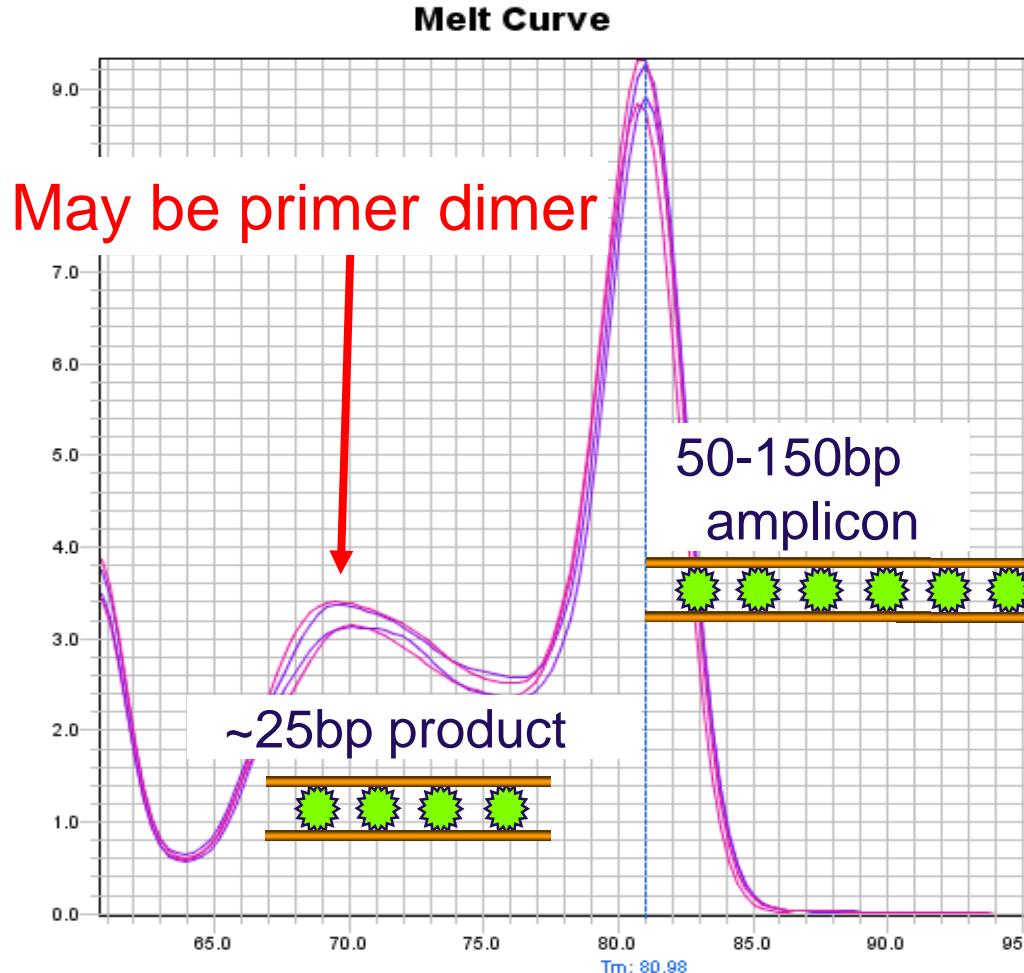
Binds to *any* double-stranded product in PCR reaction

ThermoFisher
SCIENTIFIC

Check specificity of reactions by Melt Curve protocol



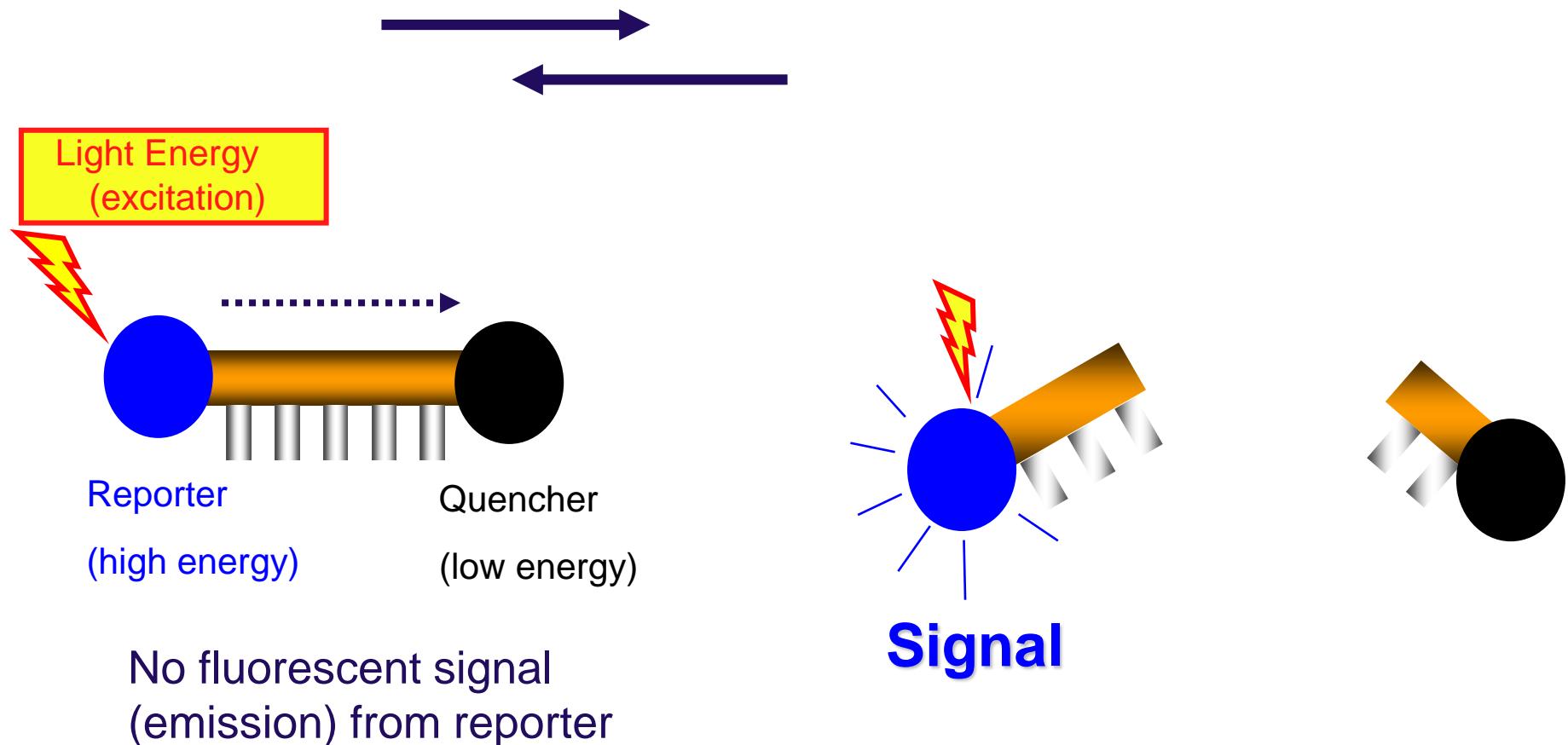
Extra peaks in melt curve



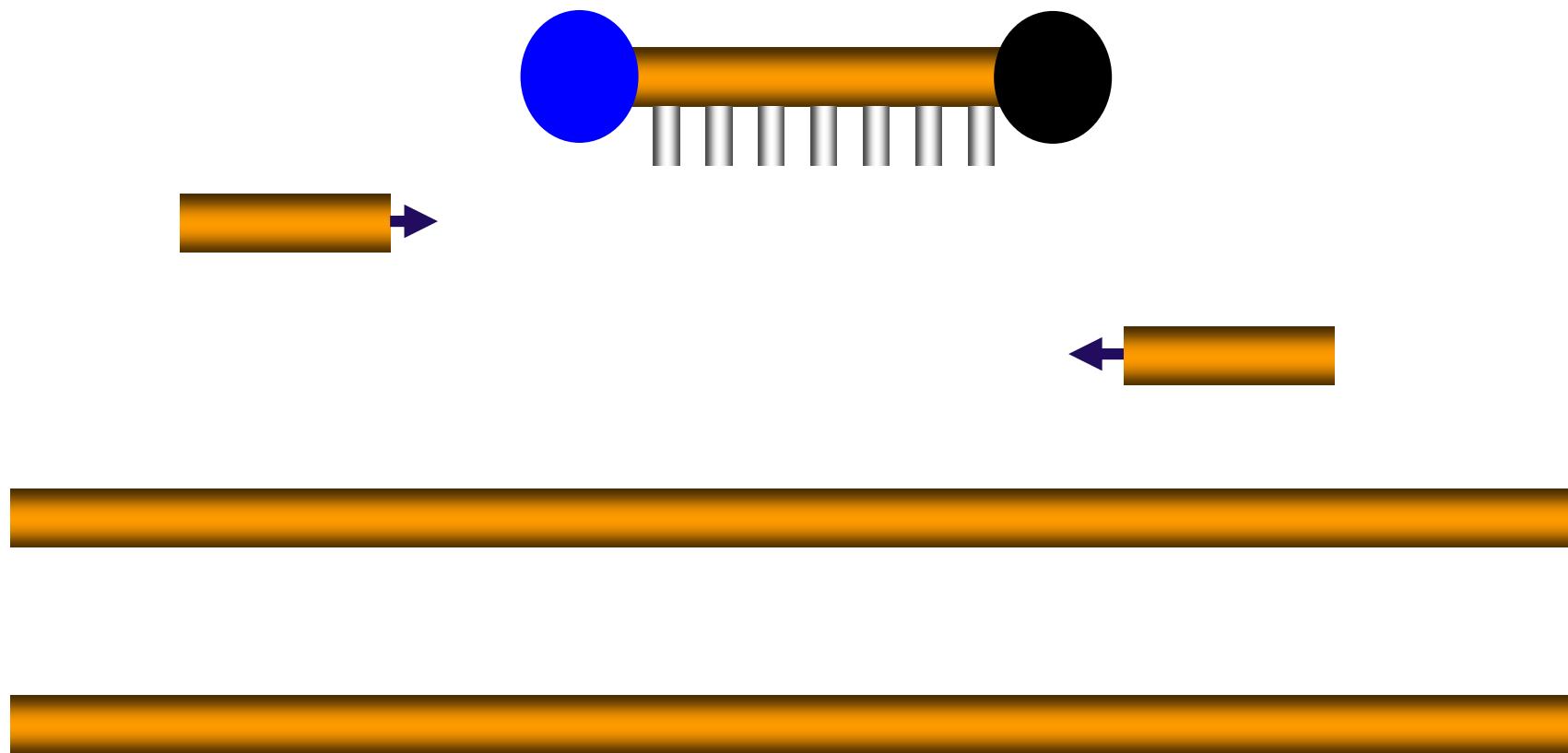
- Primer dimers can contribute towards fluorescent signal
- Primers are too complementary for each other
- Primer : template ratio is too high
- Resolved by testing out various primer concentrations

TaqMan Probe

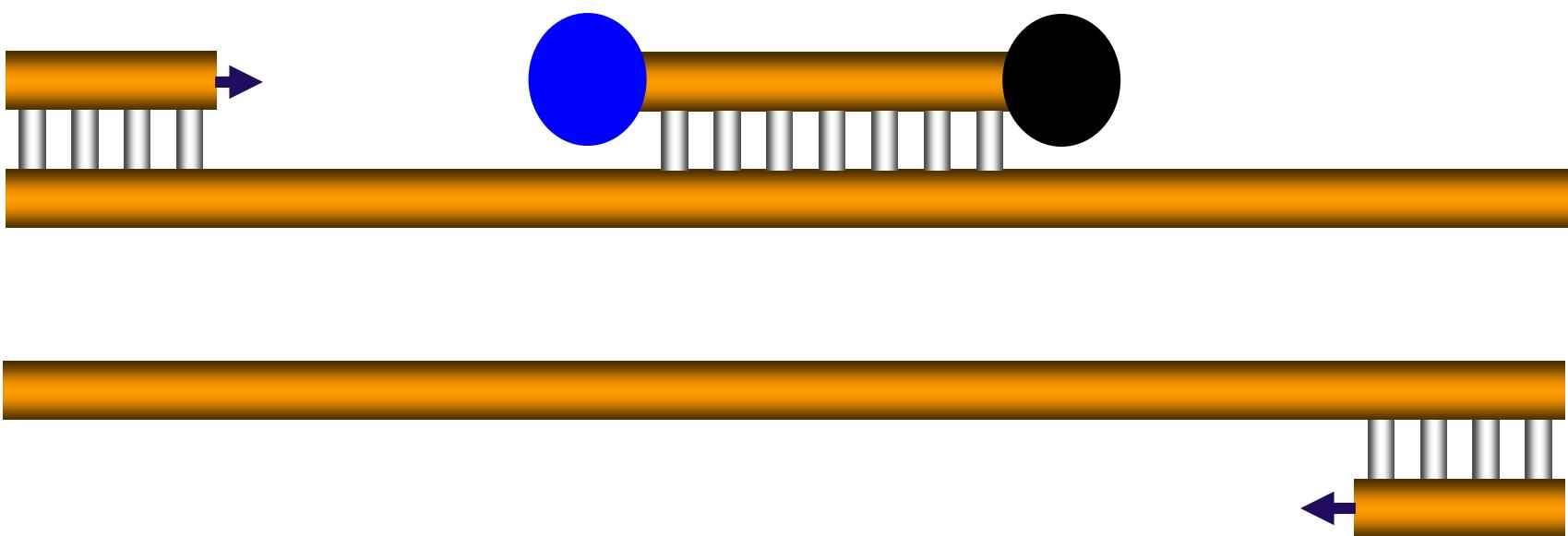
- FRET; phenomenon that describes an energy transfer mechanism between 2 fluorescent molecules when in close proximity.



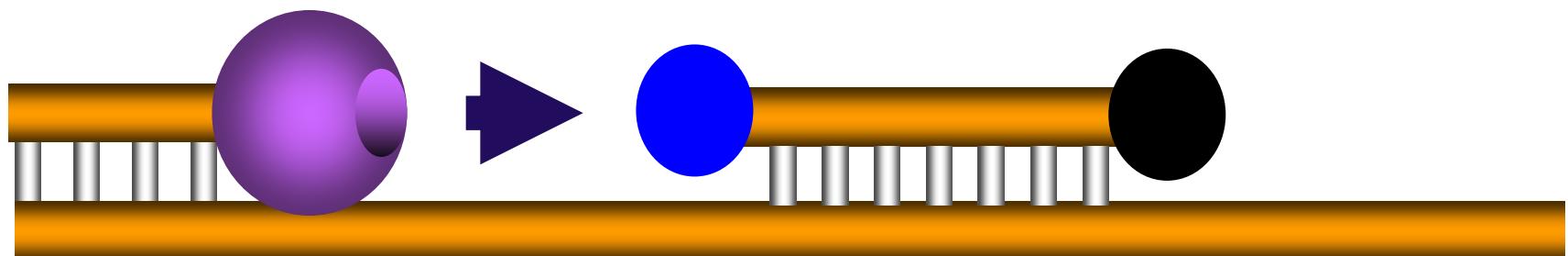
Denaturation (95°)



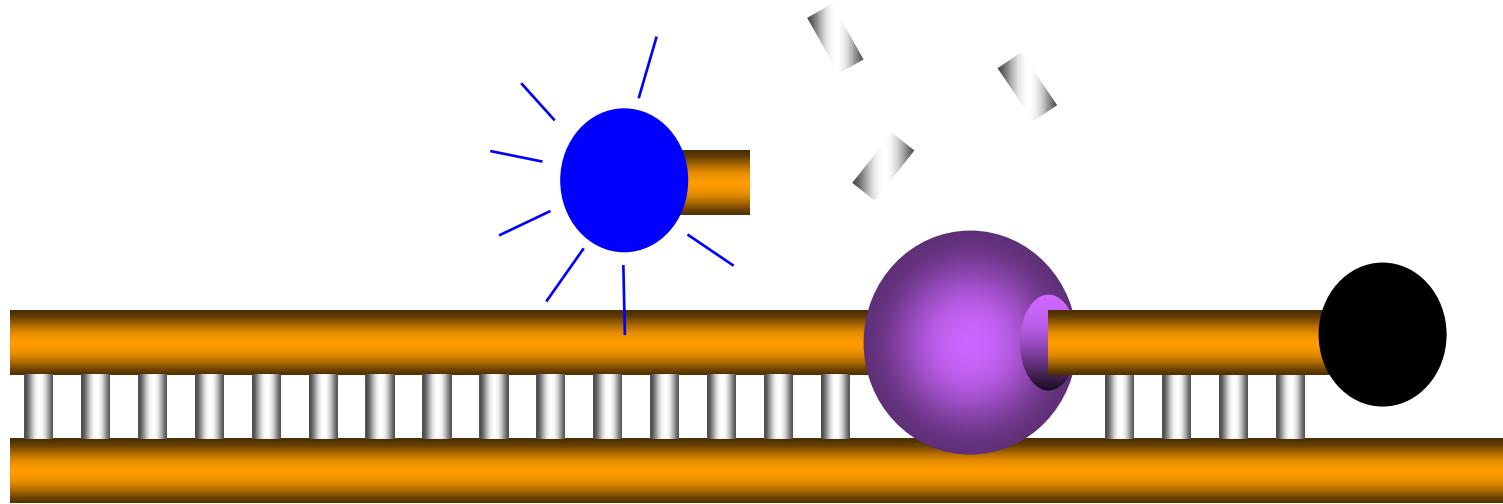
Annealing (60°)



Taq polymerase binds, then extends from upstream primer

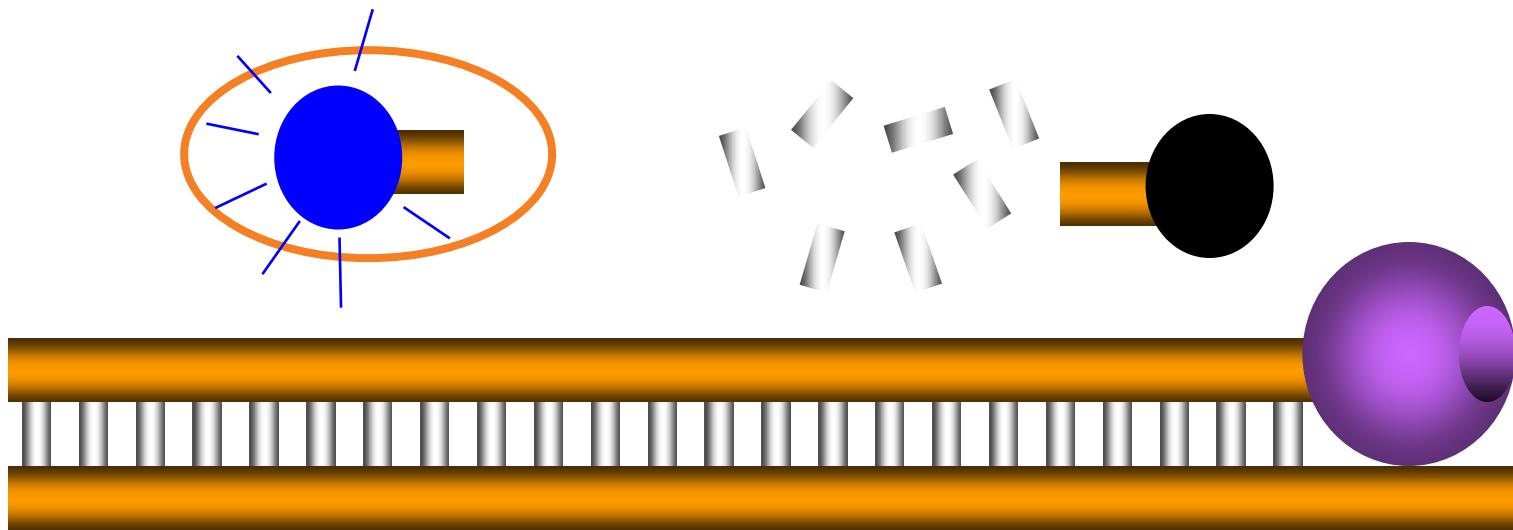


**Taq cleaves probe →
releases reporter dye**



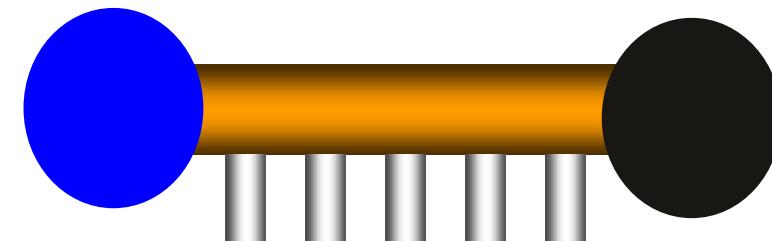
5'-Nuclease activity digests probe

Probe digested; *Taq* completes product



Reporter signal generated in tube/well

TaqMan® probe dye Choices



Reporter
(high energy)

Quencher
(low energy)

FAM™

VIC™

ABY™

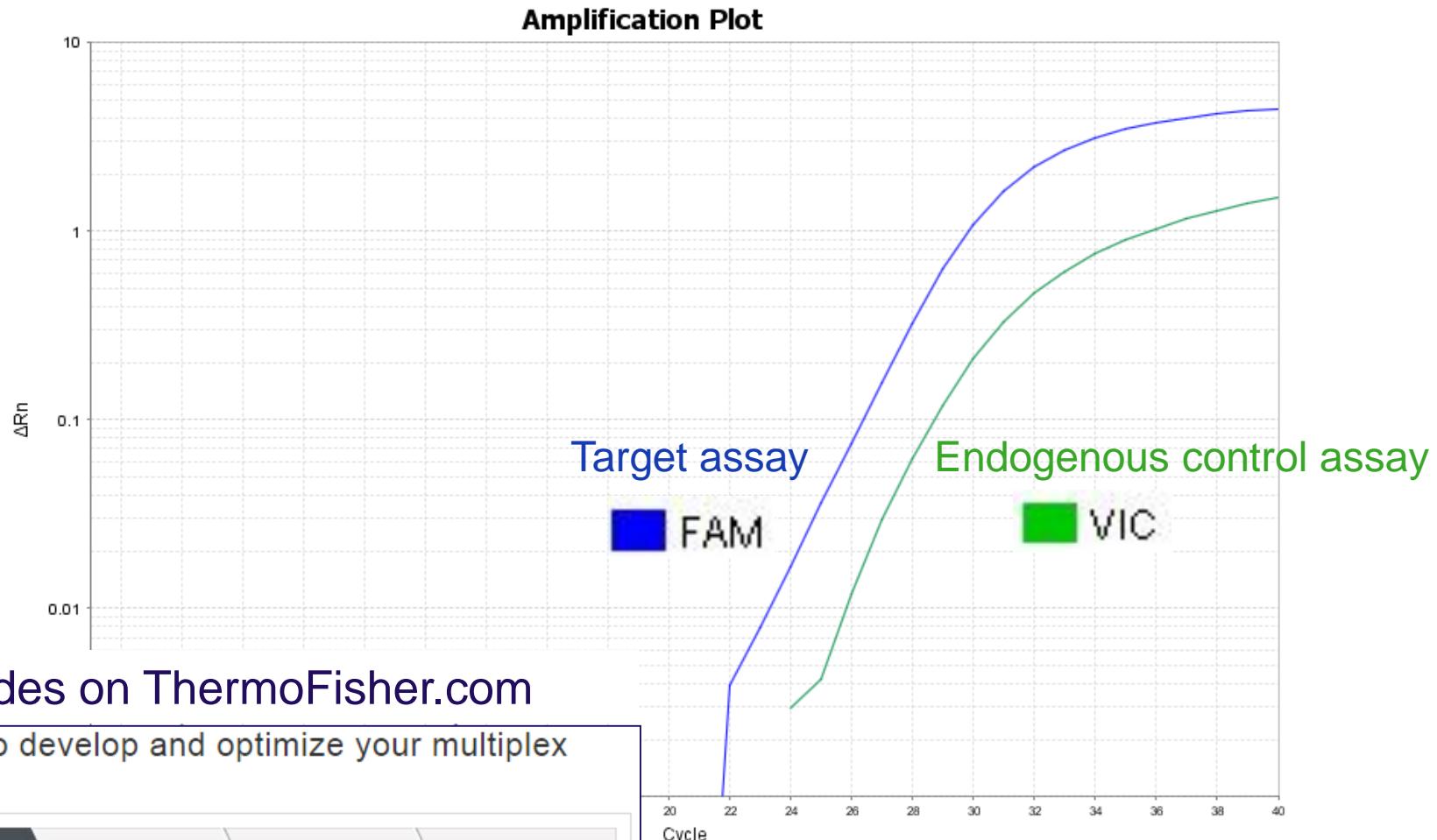
JUN™

MGB (non-fluorescent)

TAMRA™

QSY - for multiplexing >2 targets
- easily convert from BHQ

Pure Dye Spectra of System Enables Multiplexing (TaqMan chemistry)



Assay Set-up

SYBR

- primer optimization experiment... 300nM, 200nM, 100nM
- Use the concentration that provides the strongest amplification fluorescent signal and a single product in the melt curve

TaqMan

- Primers @ **900nM**, Probe @ **250nM** (final concentration in reaction).
- Probe Tm (**68-70c**) is ~10c higher than primers (**58-60c**)
- Pre-designed assays (**20X no need to worry about this!**)



TaqMan® Gene Expression Assays



Search All

Search

Popular Applications & Techniques Shop All Pro



TaqMan Real-Time PCR
Assays

| | | |
|-------------|---------------|-----------------|
| Human | Soybean | Wine grape |
| Mouse | Rhesus Monkey | Clawed frog |
| Rat | Rabbit | Chicken |
| Arabidopsis | Rice | Wheat |
| Cattle | Pathogen | Frog |
| Nematode | Baker's yeast | Chinese hamster |
| Dog | Fission yeast | Maize |
| Guinea Pig | Pig | Fruit fly |
| Zebrafish | Bread wheat | Corn |
| Horse | | |

**2,300,000
Pre-developed Assays**

1 tube contains: Primers & probe (Optimized at 20x concentration)

“Probe spans Exons”: no gDNA detected



Popular Applications



TaqMan Real-Time PCR Assays

Hs02786624_g1 GAPDH 4 RefSeq (NM) Both primers 157 User: FAM-MGB

Catalog n Target sp Importan Gene Transcripts Gene Symbol GAPDH Entrez Gene ID 2597 Gene Name glyceraldehyde-3-phosphate dehydrogenase Gene Aliases CDABP0047, G3PD, GAPD Location Chromosome Chr.12: 6643657 - 6647536 on Build 37 UniGene Hs.544577

[View Details](#) [Re](#)

99 / 100 Bloz Stars

| Interrogated Sequence | | Translated Protein | Exon Boundary | Assay Location | Amplicon Length |
|----------------------------|--------------------------------|--------------------------------|---------------|----------------|-----------------|
| RefSeq | NM_001256799.1 | NP_001243728.1 | 6 - 7 | 728 | 93 |
| | NM_002046.4 | NP_002037.2 | 7 - 8 | 704 | 93 |
| GenBank mRNA | AB062273.1 | - | 7 - 8 | 582 | 93 |
| | AF261085.1 | - | 7 - 8 | 633 | 93 |
| | AK026525.1 | - | 7 - 8 | 655 | 93 |
| | AK308198.1 | - | 7 - 8 | 606 | 93 |
| | AY007133.1 | - | 7 - 8 | 582 | 93 |
| | AY633612.1 | - | 6 - 7 | 554 | 93 |
| | BC001601.1 | - | 7 - 8 | 584 | 93 |
| | BC004109.2 | - | 7 - 8 | 582 | 93 |
| | BC009081.1 | - | 7 - 8 | 582 | 93 |
| | BC013310.2 | - | 7 - 8 | 582 | 93 |
| BC020308.1 | - | 7 - 8 | 582 | 93 | |
| BC023632.2 | - | 7 - 8 | 582 | 93 | |
| BC025925.1 | - | 7 - 8 | 582 | 93 | |
| BC026907.1 | - | 7 - 8 | 582 | 93 | |

Build 37.1 Human Chr.12: Hs02758991_g1

Availability: **Inventoried**
Price (USD): **173.00**
[Check your price](#)

[View Details](#) [Related Reagents](#) [Related Controls](#) [View Assay Map](#)

[Add to cart](#)

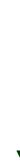


Cell Culture Plas

Custom TaqMan® Assay Design Tool

| Status | Assay Name ? | Primer Sequence ? | Probe Sequence ? |
|--------|------------------------------|--|--|
| | | Forward Primer: <input type="text"/> | Probe 1 Dye: 6-FAM <input type="text"/> |
| | | Reverse Primer: <input type="text"/> | Do you wish to permanently hide your assay primer/probe sequences in all documents? More Info ? <input checked="" type="radio"/> No <input type="radio"/> Yes |

| Status | Name ? | Sequence ? | Target Position & Name ? |
|--------|------------------------|--|--|
| Valid | TEST | GTCAGAGAGAAAGTAGGAGGCCATTGAAGTCGTGGACTGAGGAACAA AATAACACAAGTGTAGAGAGAAAGTAGGAGGCCATTGAAGTCGTGG ACTGAGGAACAAAATACACAAGTGTAGAGAGAAAGTAGGAGGCCATT GAGCTCTCCGCTCTCGACGACGATGAGCTGAGCTGAGCTGAGCTGAG Do you wish to permanently hide your assay primer/probe sequences in all documents? More Info ? <input checked="" type="radio"/> No <input type="radio"/> Yes | <input type="radio"/> Manual <input checked="" type="radio"/> Automatic <input type="text"/> ANY <input type="text"/> ANY |



Design Job Details

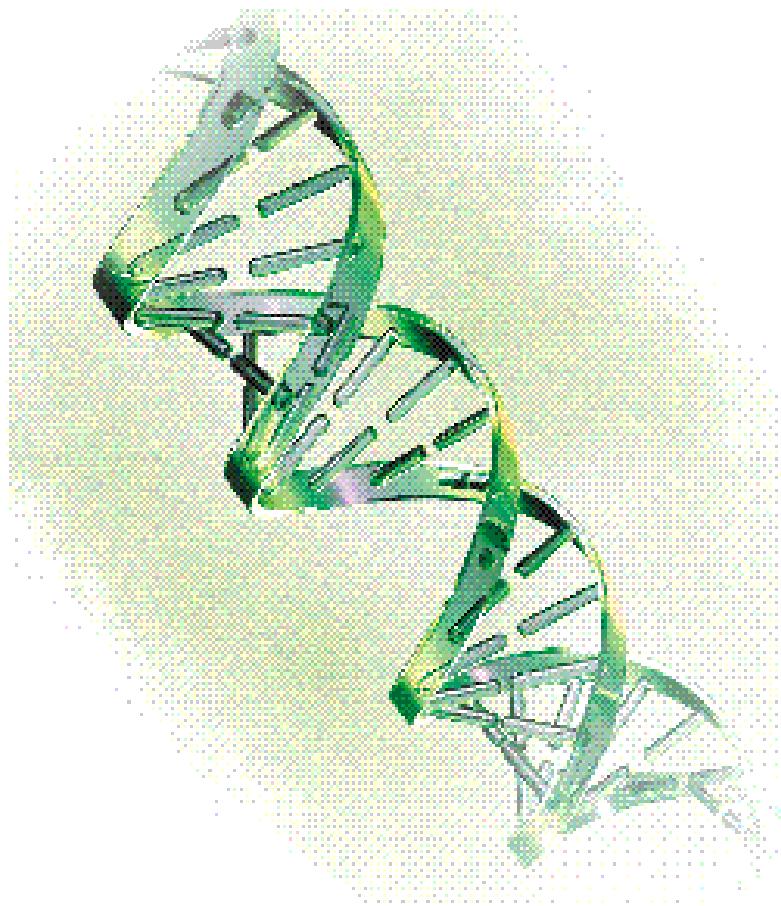
Refresh Batch List

| Batch ID | Submitted | Status | Details ? |
|----------------|---------------------|-----------|---------------------------------|
| w0906118406000 | 2009-06-13 16:59:41 | COMPLETED | 1 Passed, 0 Failed, 0 No_Design |

Design Results

| ID / Name | Type | Design Status | Size | Quantity | Add All |
|-----------------|--------|---------------|-----------------|------------------------|------------------------------------|
| AILIWHU TEST | Custom | Passed | 4331348 : small | <input type="text"/> 1 | <input type="button" value="Add"/> |

Primer Express® Software



AB applied
biosystems

Primer Express® Software for Real-Time PCR

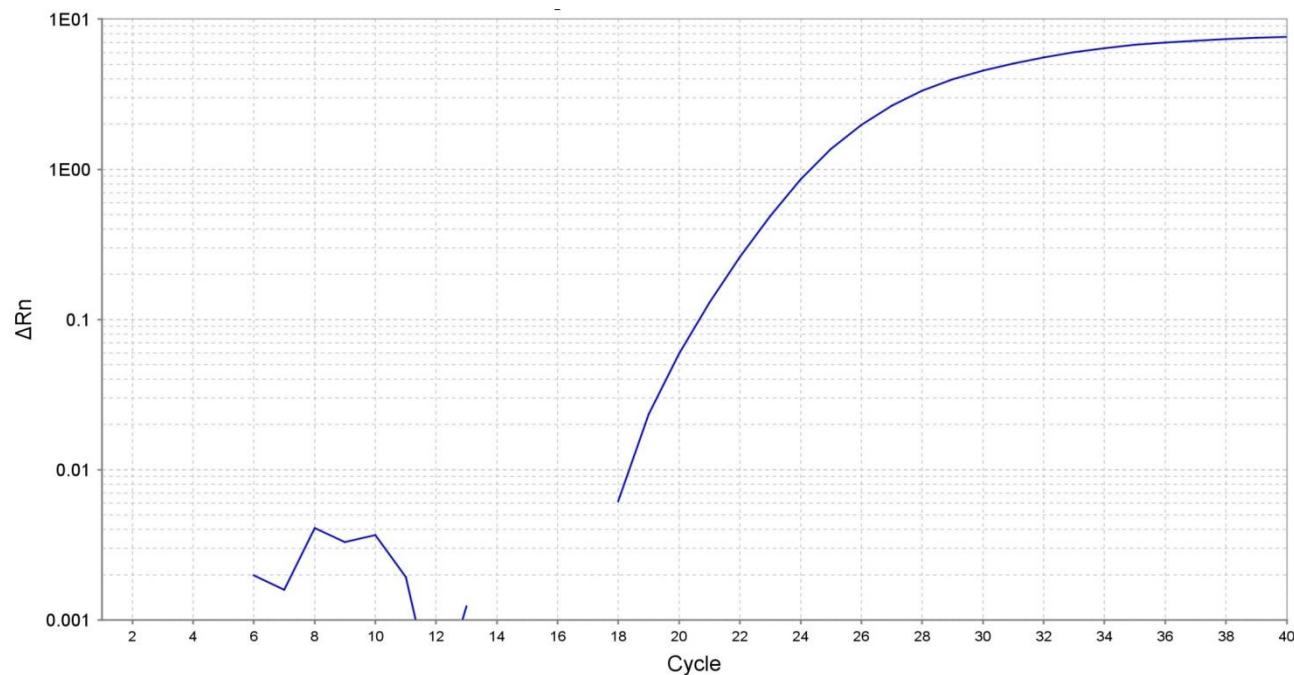
Version 3.0.1

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<https://www.thermofisher.com/order/catalog/product/4363991>

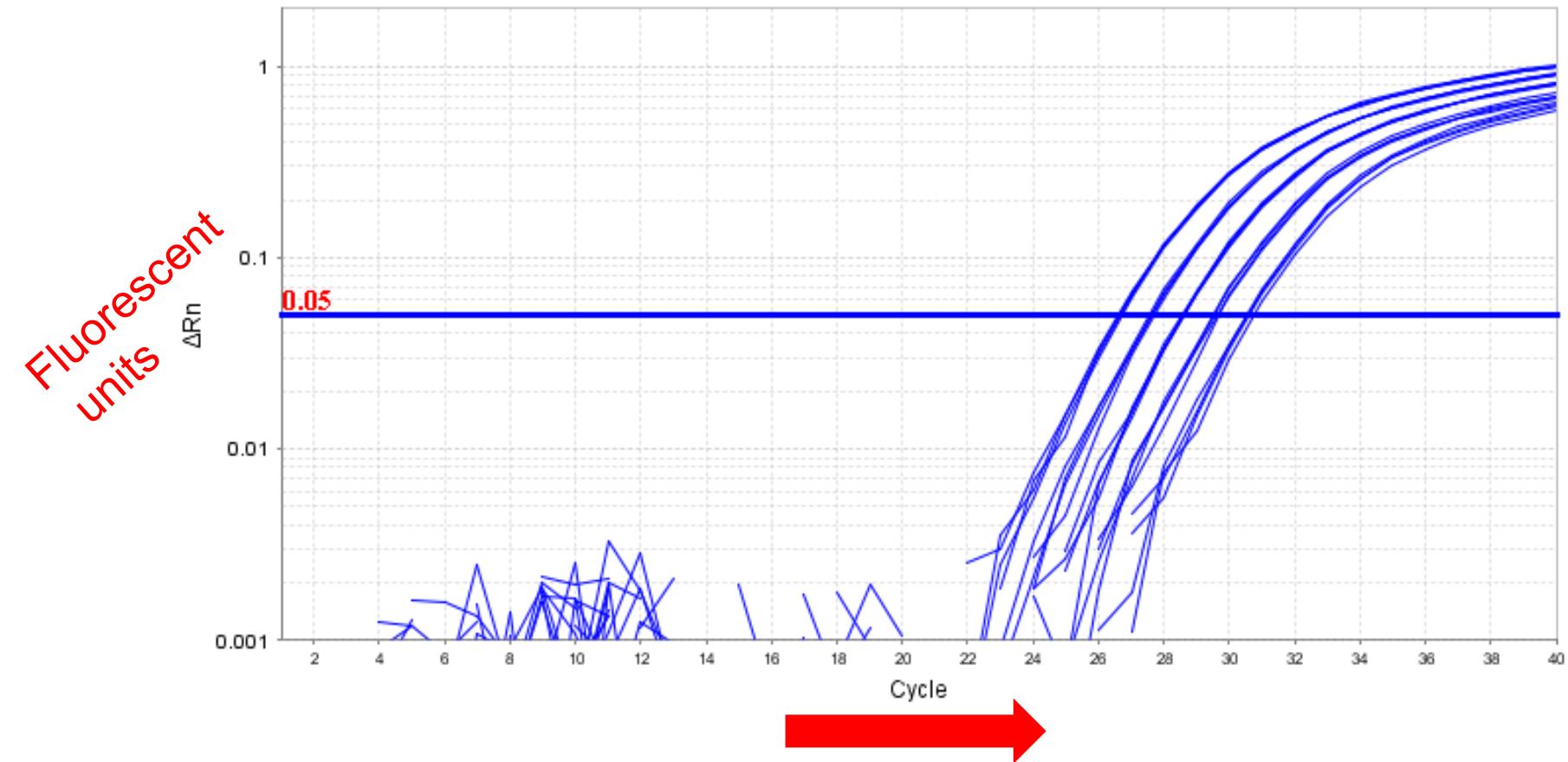
ThermoFisher
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Examining qPCR amplification plots



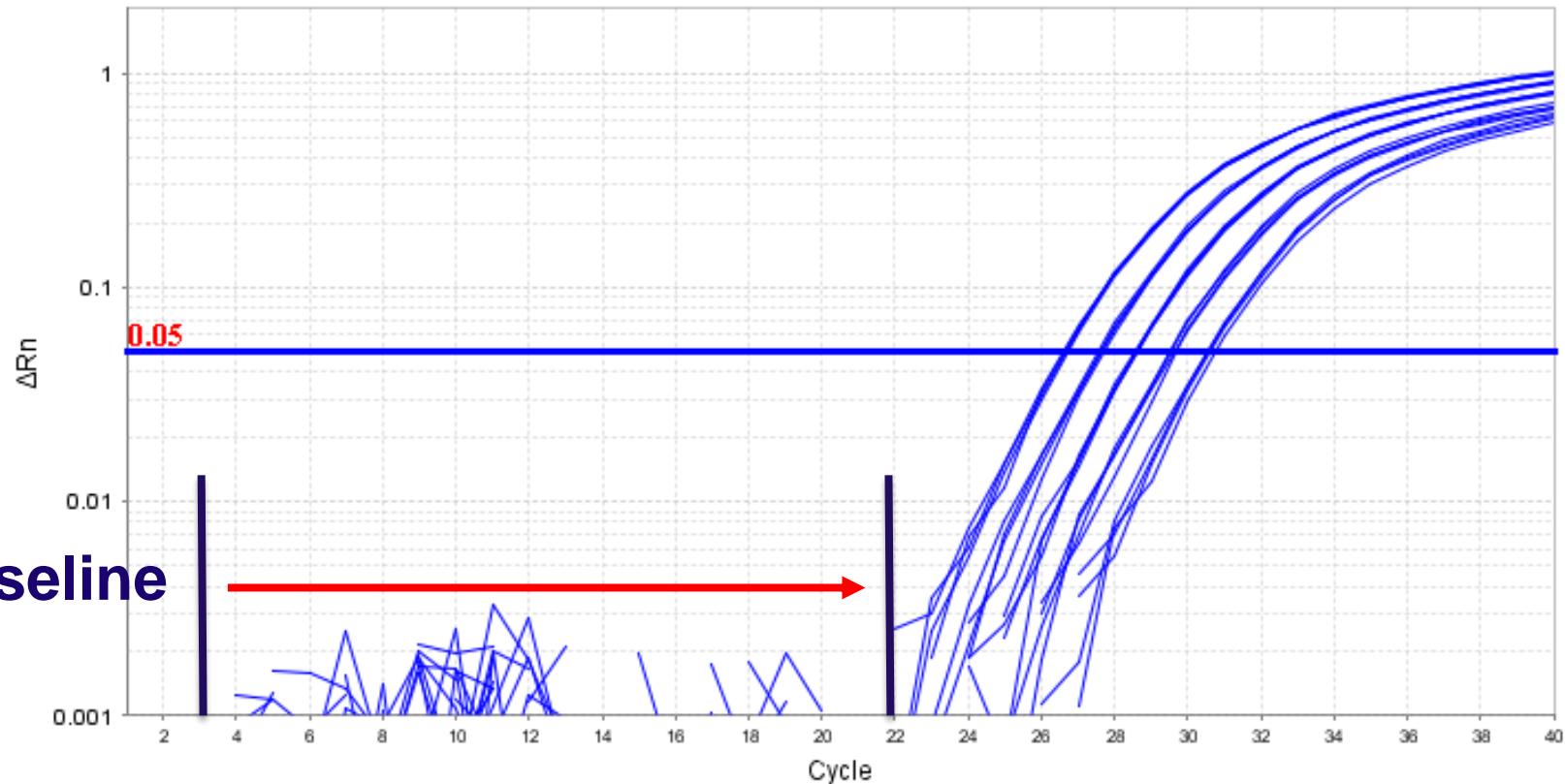
Real-time PCR Terminology

**Amplification plots
graph cycle vs. Δ in fluorescence**



Baseline

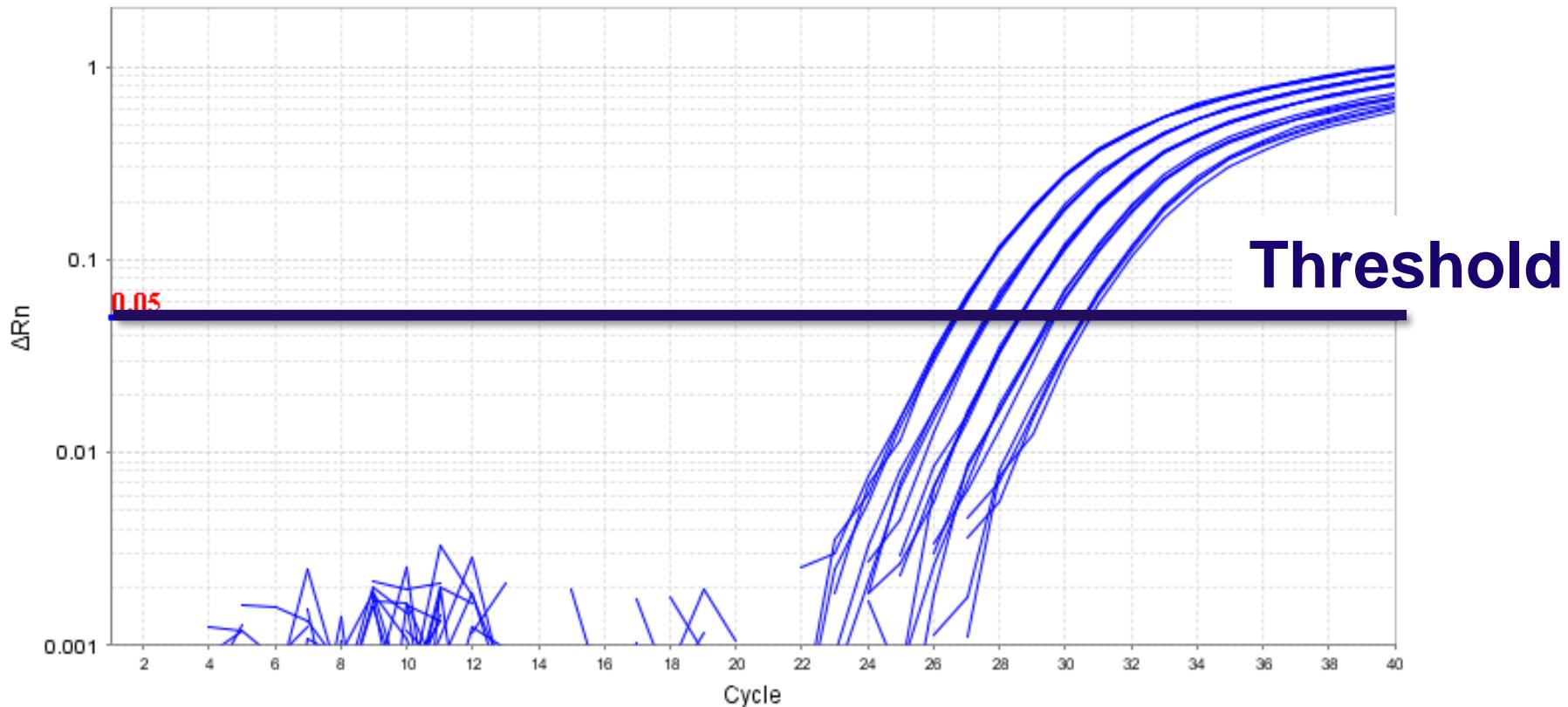
Defines region of background noise by a cycle-to-cycle range ex. cycles 3-22



Automatic/manual baseline

Threshold

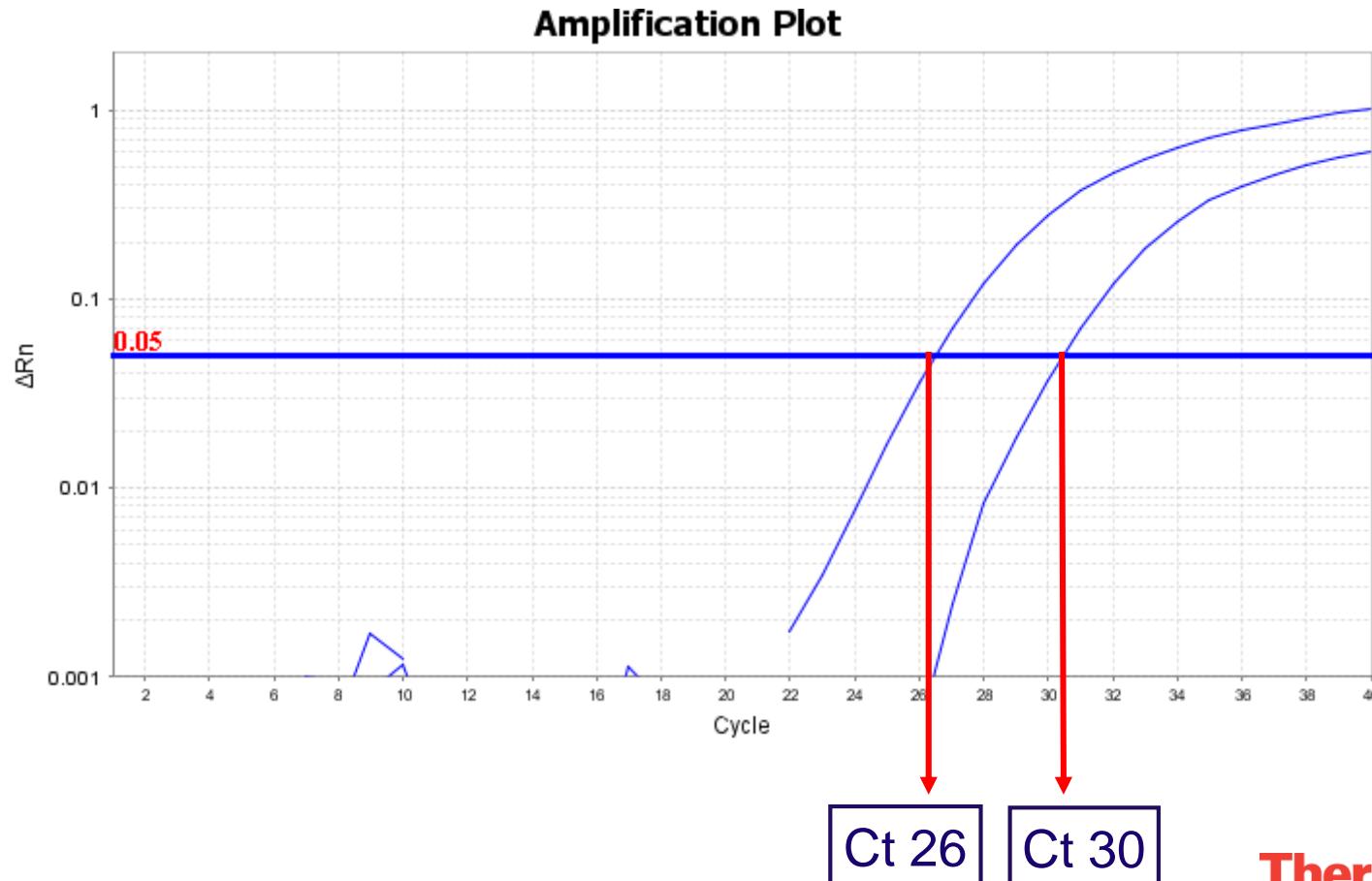
An adjustable horizontal line in the amplification plot.
Tells the software where to capture the data.



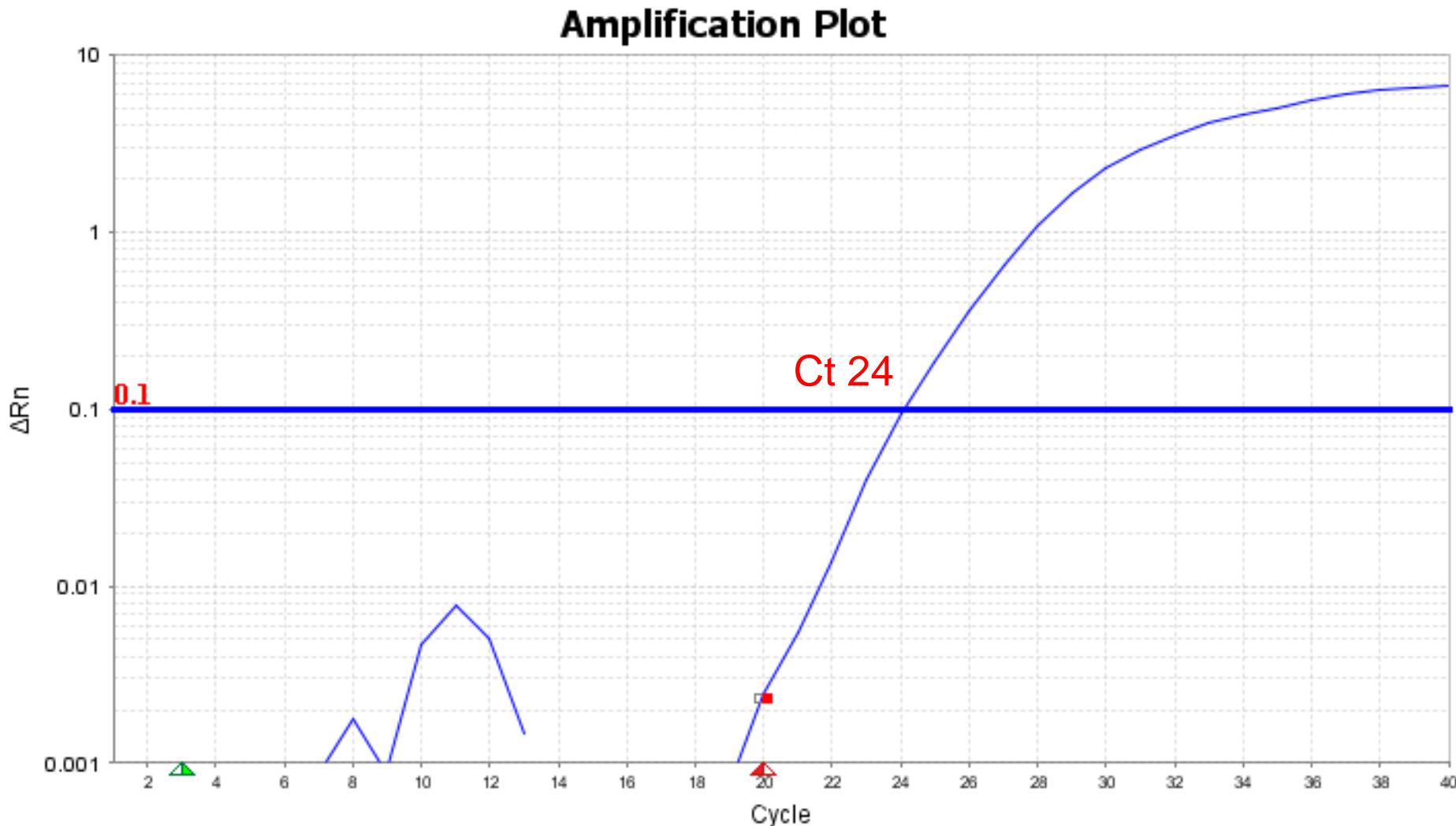
AutoThreshold (default); Software sets the threshold for each assay.
Different assays may have different exponential phases

Cycle threshold (Ct)

The fractional cycle number at which each amplification curve crosses the threshold.

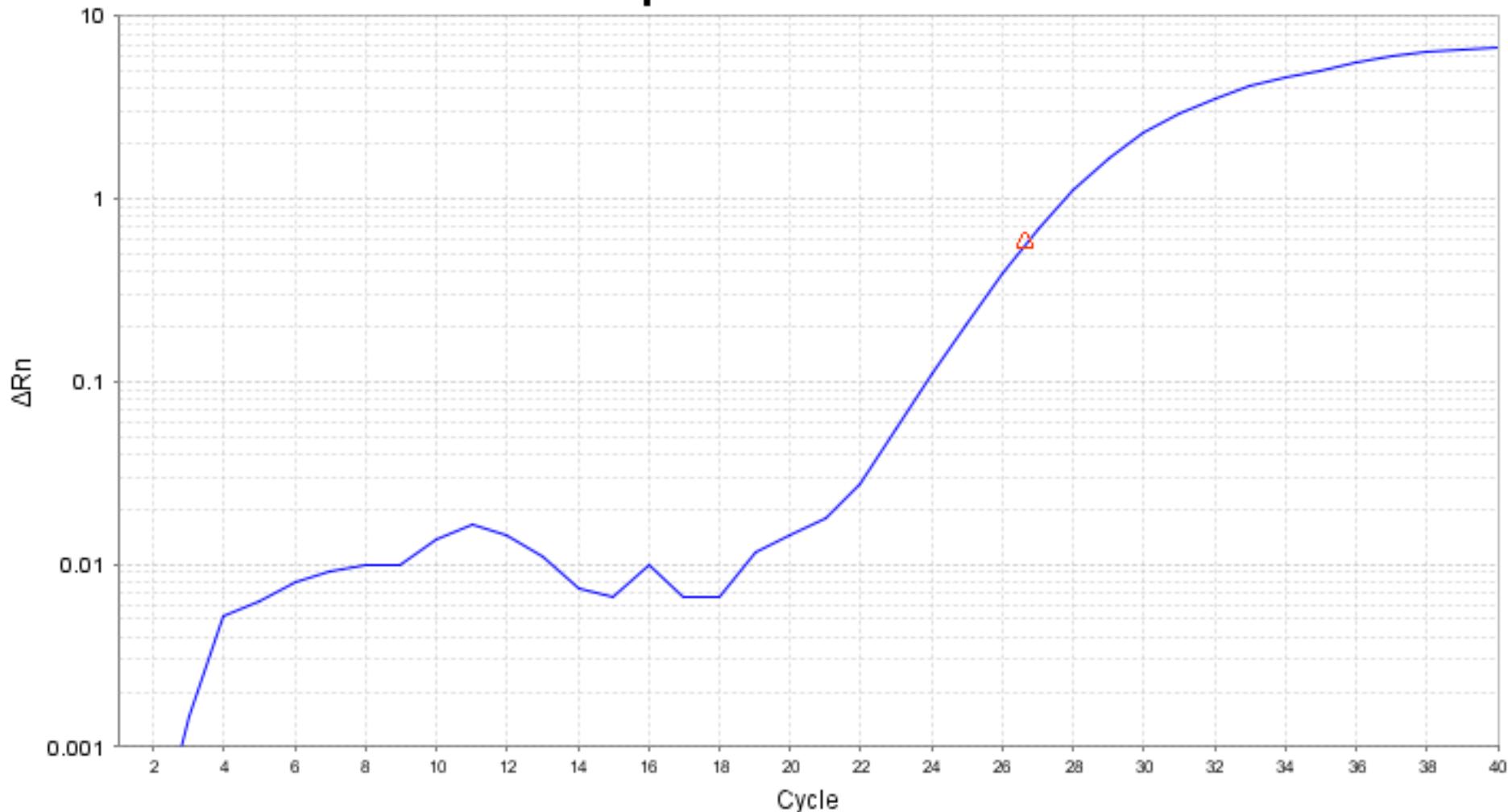


Summary of Ct algorithm (“baseline threshold”)



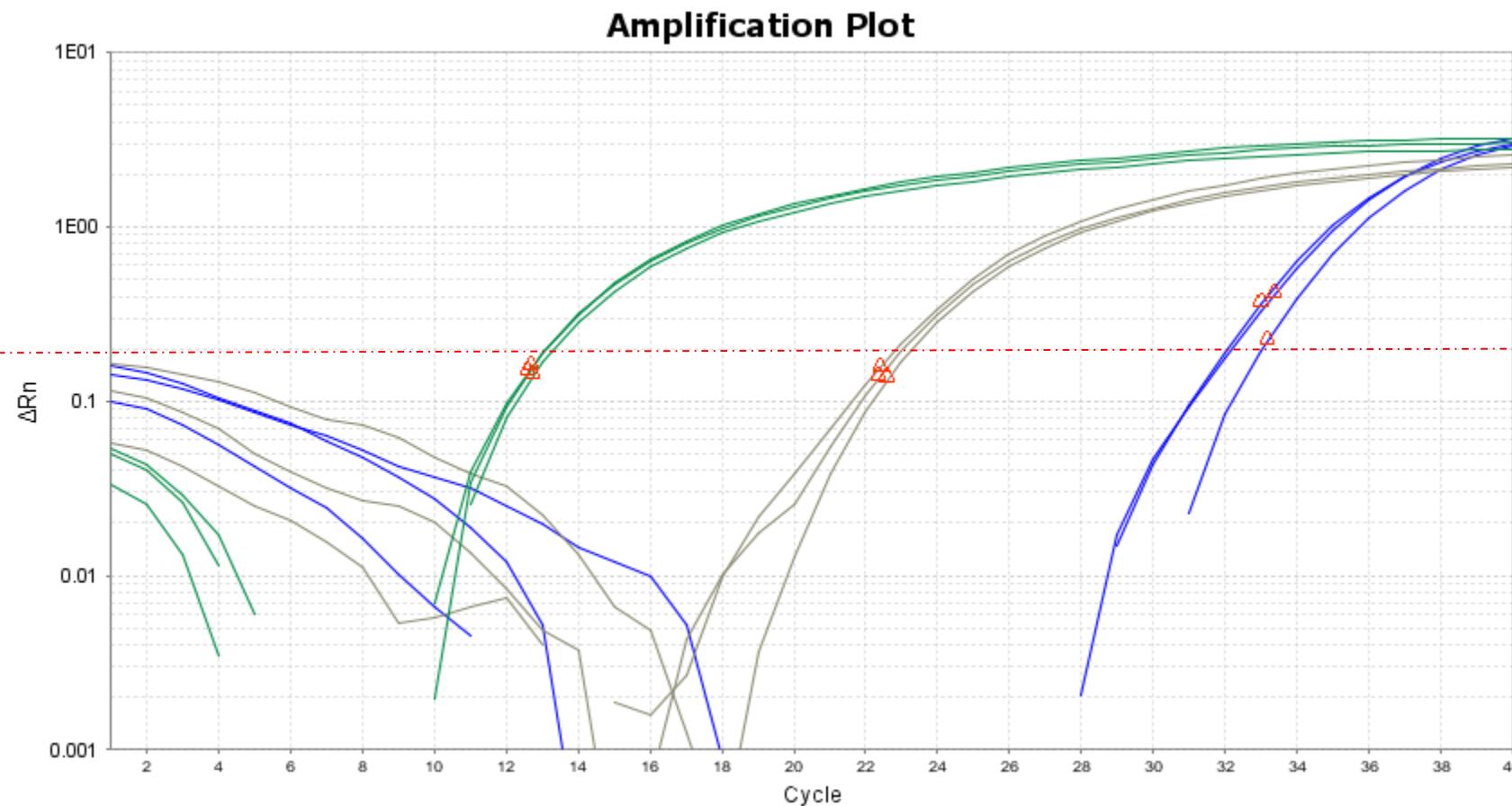
CRT algorithm (“relative threshold”)

Amplification Plot



App note: http://tools.thermofisher.com/content/sfs/brochures/CO28730-Crt-Tech-note_FLR.pdf

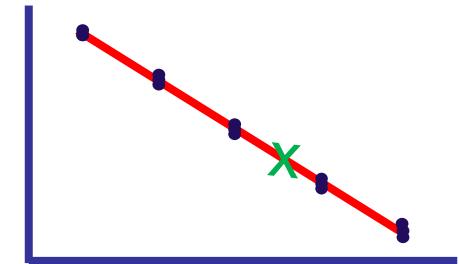
CRT may help out for samples that may need to be omitted with CT, to tighten up technical replicates.



Use Ct values to provide quantitative results

Absolute Quantitation calculates a copy number by using standard curve analysis

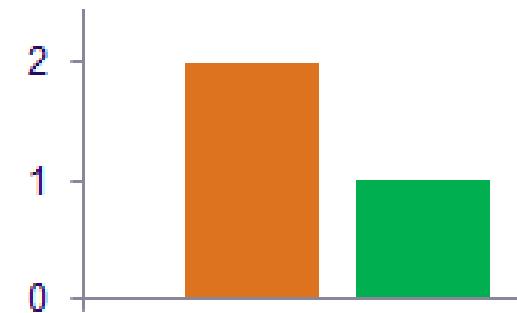
“I have 500 copies of TNF α in sample 1”



$\Delta\Delta Ct$ method calculates relative fold change

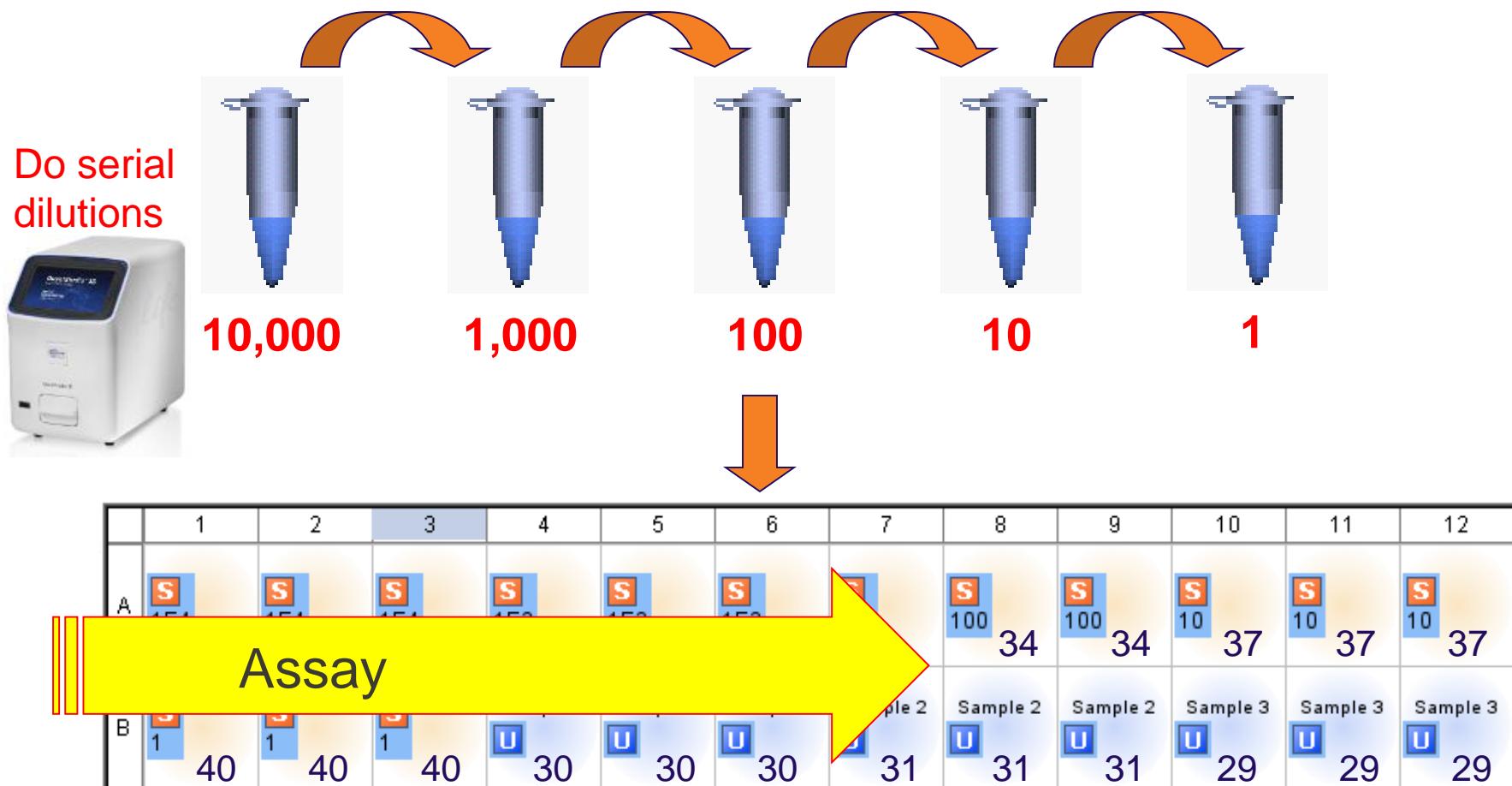
- no standard curves, assays have similar efficiencies

“TNF α is upregulated 2 fold in sample 1 compared to the control sample”

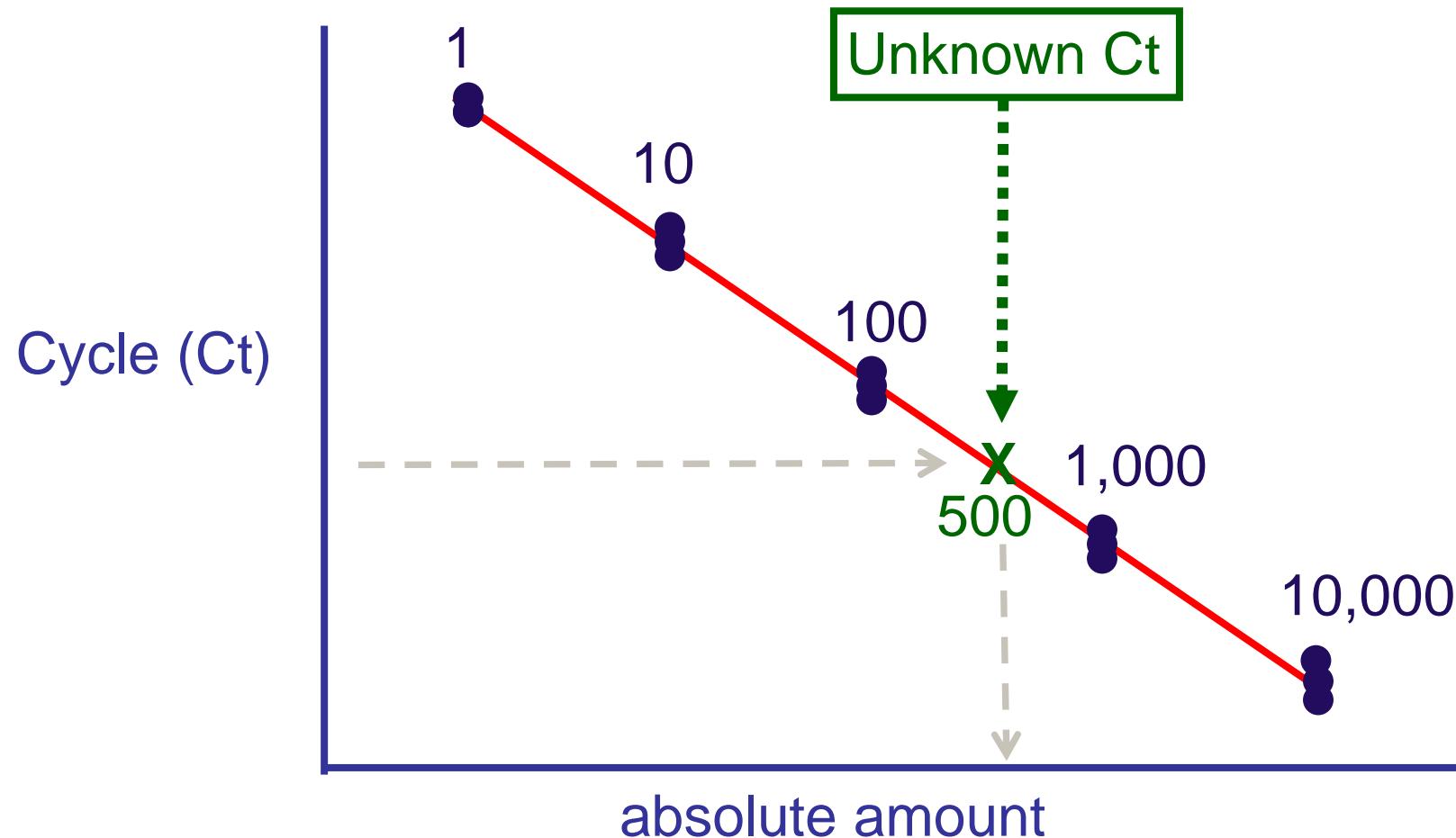


Generate an absolute standard curve

Recommend 5 or more, point standard curve of 10 fold dilutions



After a run, plot unknown Ct on curve



Another way to analyze expression data for relative comparisons (2^n)

ΔCt of 1 = 2^1 = 2 fold difference

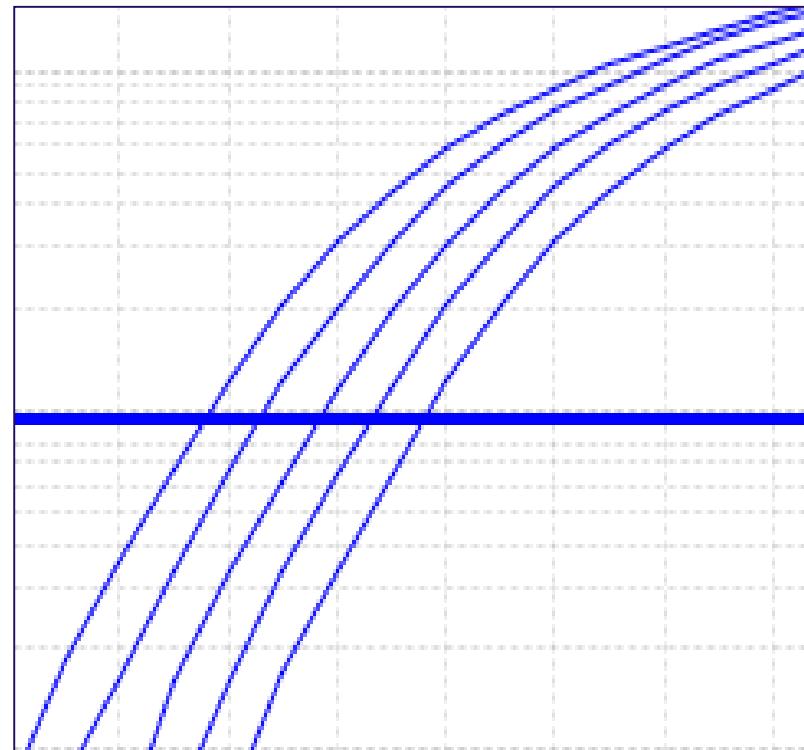
ΔCt of 2 = 2^2 = 4 fold

ΔCt of 3 = 2^3 = 8 fold

ΔCt of 3.3 = $2^{3.3}$ = 10 fold

ΔCt of 4 = 2^4 = 16 fold

This assumes that the amount of product is doubling with each cycle in the exponential phase, when it does, we call this 100% amplification efficiency.

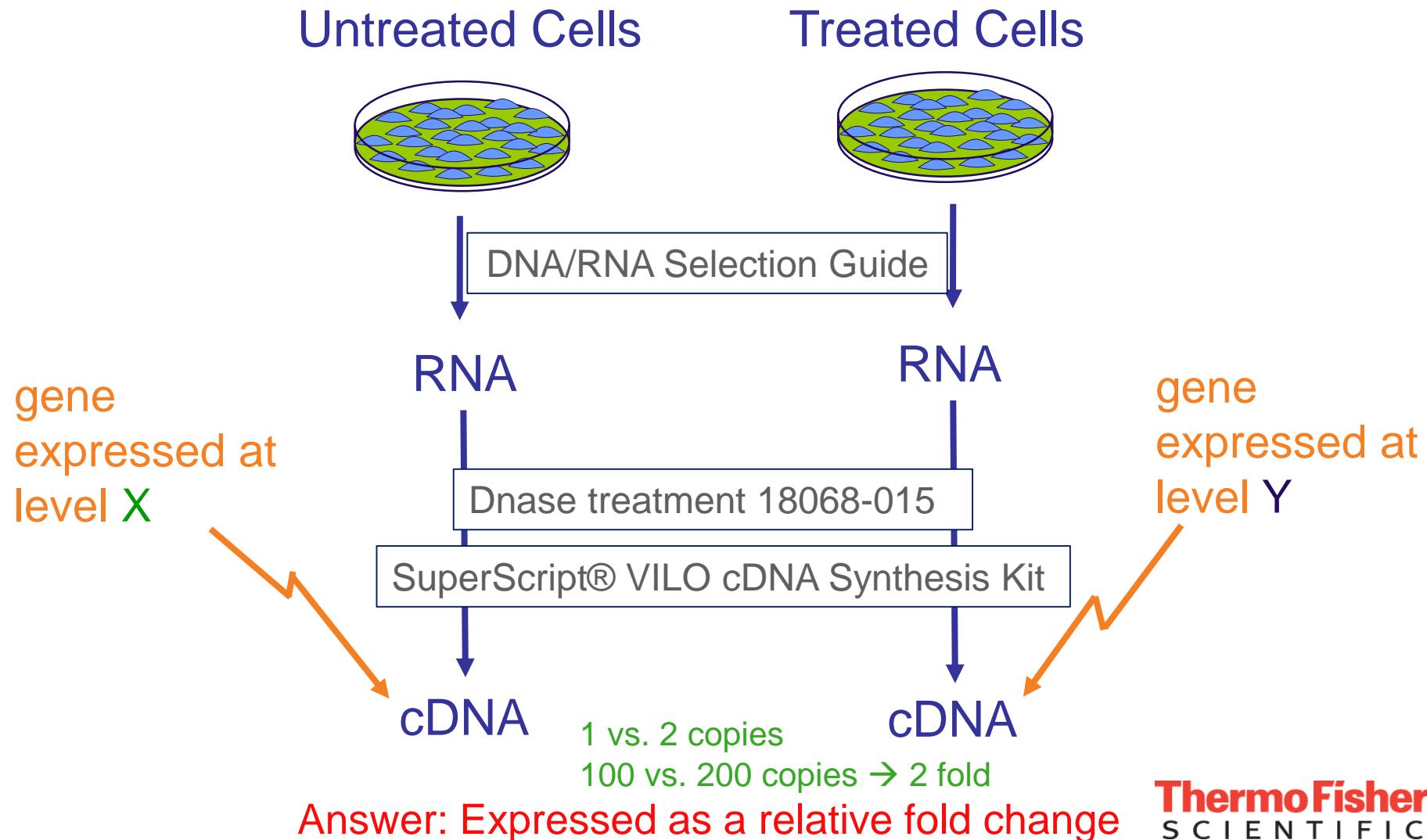


Standard curve statistics example:

Slope: -3.384 R²: 0.995 Eff%: 97.477

$\Delta\Delta Ct$

Question: What happens to the expression of a gene when I apply a treatment?



In relative gene expression experiments, we examine two types of genes. . .

Gene of interest (target)

Normalizing gene (control)

A gene that is **expressed consistently** in all samples in an experiment.
(a.k.a. “control gene,” “housekeeping gene,” “endogenous control”,
“reference gene”)

TaqMan® Endogenous Control Array

- Human & Rodent available
- Includes 32 commonly studied control genes
- Genes range in expression from:
high, medium, low (ex. 18s, GAPDH, HPRT1)

| Gene Symbols | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 | 10 | 11 | 12 |
|--------------|-------|--------|---------|---------|-------|--------|---------|---------|-------|--------|---------|---------|
| A | 18S | GAPDH | HPRT1 | GUSB | 18S | GAPDH | HPRT1 | GUSB | 18S | GAPDH | HPRT1 | GUSB |
| B | ACTB | B2M | HMBS | IP08 | ACTB | B2M | HMBS | IP08 | ACTB | B2M | HMBS | IP08 |
| C | PGK1 | RPLP0 | TBP | TFRC | PGK1 | RPLP0 | TBP | TFRC | PGK1 | RPLP0 | TBP | TFRC |
| D | UBC | YWHAZ | PPIA | POLR2A | UBC | YWHAZ | PPIA | POLR2A | UBC | YWHAZ | PPIA | POLR2A |
| E | CASC3 | CDKN1A | CDKN1B | GADD45A | CASC3 | CDKN1A | CDKN1B | GADD45A | CASC3 | CDKN1A | CDKN1B | GADD45A |
| F | PUM1 | PSMC4 | EIF2B1 | PES1 | PUM1 | PSMC4 | EIF2B1 | PES1 | PUM1 | PSMC4 | EIF2B1 | PES1 |
| G | ABLI | ELF1 | MISATP6 | MRPL19 | ABLI | ELF1 | MISATP6 | MRPL19 | ABLI | ELF1 | MISATP6 | MRPL19 |
| H | POP4 | RPL37A | RPL30 | RPS17 | POP4 | RPL37A | RPL30 | RPS17 | POP4 | RPL37A | RPL30 | RPS17 |

TaqMan® Endogenous Control Array

(Human & Rodent available)

| Gene Symbols | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 | 10 | 11 | 12 |
|--------------|-------|--------|---------|---------|-------|--------|---------|---------|-------|--------|---------|---------|
| A | 18S | GAPDH | HPRT1 | GUSB | 18S | GAPDH | HPRT1 | GUSB | 18S | GAPDH | HPRT1 | GUSB |
| B | ACTB | B2M | HMBS | IP08 | ACTB | B2M | HMBS | IP08 | ACTB | B2M | HMBS | IP08 |
| C | PGK1 | RPLP0 | TBP | TFRC | PGK1 | RPLP0 | TBP | TFRC | PGK1 | RPLP0 | TBP | TFRC |
| D | UBC | YWHAZ | PPIA | POLR2A | UBC | YWHAZ | PPIA | POLR2A | UBC | YWHAZ | PPIA | POLR2A |
| E | CASC3 | CDKN1A | CDKN1B | GADD45A | CASC3 | CDKN1A | CDKN1B | GADD45A | CASC3 | CDKN1A | CDKN1B | GADD45A |
| F | PUMI | PSMC4 | EIF2B1 | PES1 | PUMI | PSMC4 | EIF2B1 | PES1 | PUMI | PSMC4 | EIF2B1 | PES1 |
| G | ABL1 | ELF1 | MIR1826 | MRPL19 | ABL1 | ELF1 | MIR1826 | MRPL19 | ABL1 | ELF1 | MIR1826 | MRPL19 |
| H | POP4 | RPL37A | RPL30 | RPS17 | POP4 | RPL37A | RPL30 | RPS17 | POP4 | RPL37A | RPL30 | RPS17 |



Hs99999902_m1

Example of $\Delta\Delta Ct$ Math

| Sample | X | N |
|-----------|----|----|
| Treated 1 | 24 | 14 |
| Treated 2 | 20 | 11 |
| Treated 3 | 28 | 12 |
| Untreated | 24 | 13 |

X = Target N = Normalizing gene

Example of $\Delta\Delta Ct$ math

| Sample | X | N | ΔCt |
|-----------|---------|---|-------------|
| Treated 1 | 24 - 14 | = | 10 |
| Treated 2 | 20 - 11 | = | 9 |
| Treated 3 | 28 - 12 | = | 16 |
| Untreated | 24 - 13 | = | 11 |

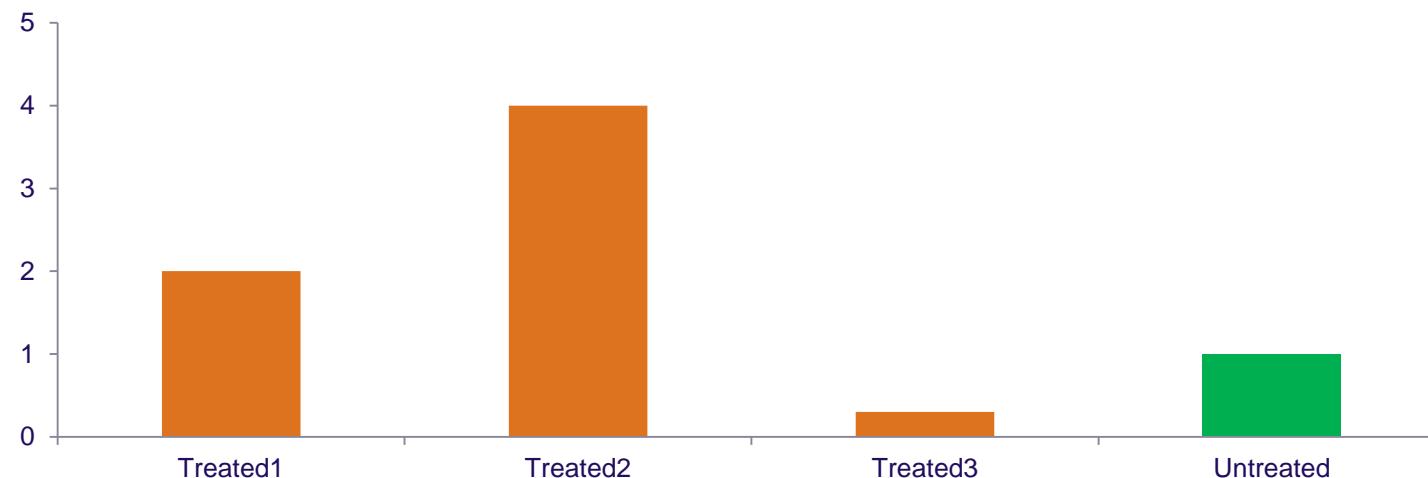
Choose a **calibrator**; all samples are relatively expressed to this sample

| Sample | X | N | ΔCt | $\Delta\Delta Ct$ |
|---------------|----------|----------|-------------------------------|-------------------------------------|
| Treated 1 | 24 | 14 | 10 – 11 = -1 | |
| Treated 2 | 20 | 11 | 9 – 11 = -2 | |
| Treated 3 | 28 | 12 | 16 – 11 = 5 | |
| Untreated | 24 | 13 | 11 – 11 = 0 | |

Example of $\Delta\Delta Ct$ math

RQ (fold change)

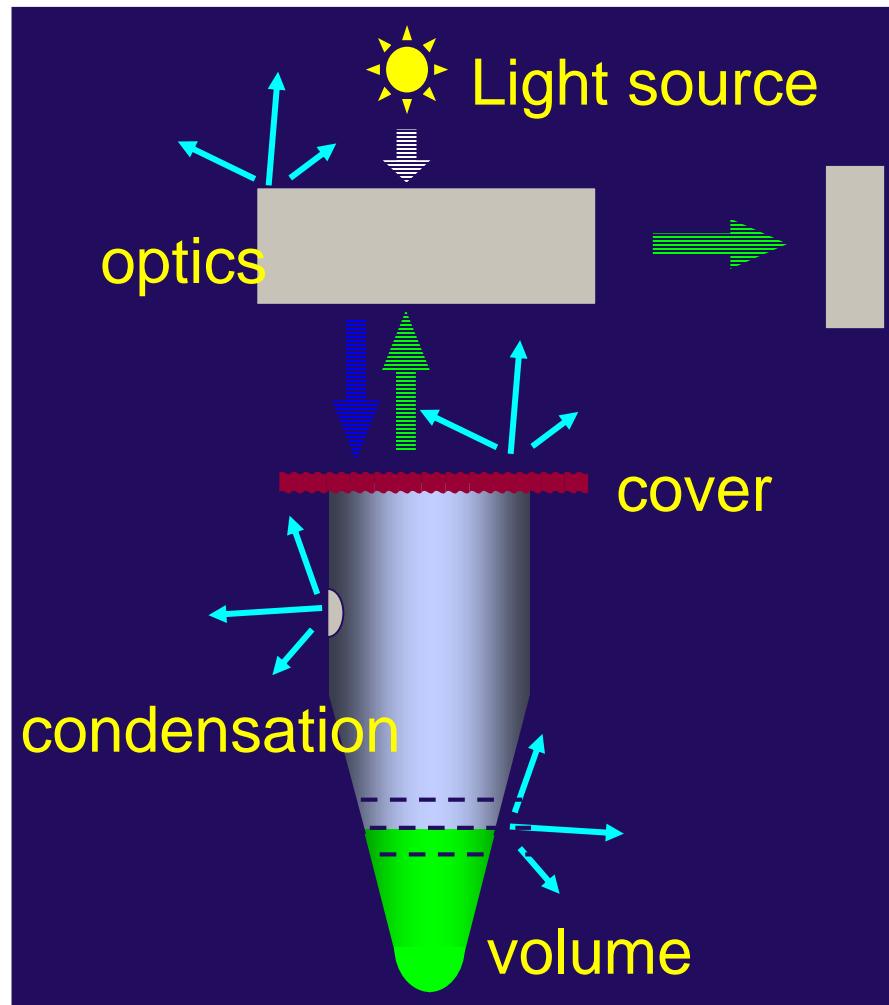
| Sample | X | N | ΔCt | $\Delta\Delta Ct$ | $2^{-\Delta\Delta Ct}$ | RQ (fold change) |
|-----------|----|----|-------------|-------------------|------------------------|------------------|
| Treated 1 | 24 | 14 | 10 | -1 | 2 | ↑ |
| Treated 2 | 20 | 11 | 9 | -2 | 4 | ↑ |
| Treated 3 | 28 | 12 | 16 | 5 | 0.3 | ↓ |
| Untreated | 24 | 13 | 11 | 0 | 1 | |



Software can perform the calculations!

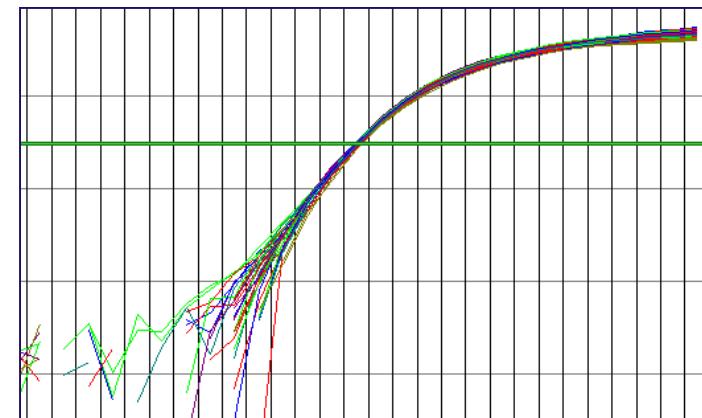
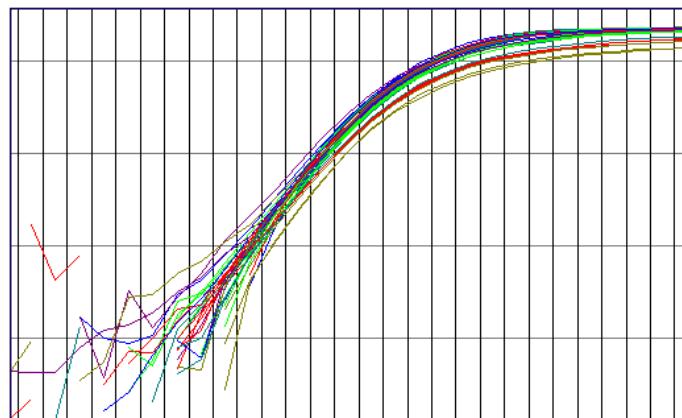
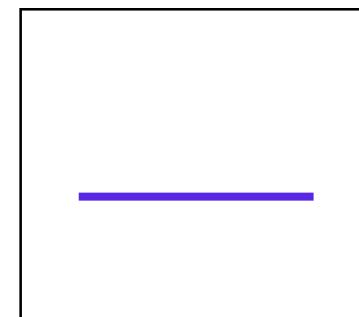
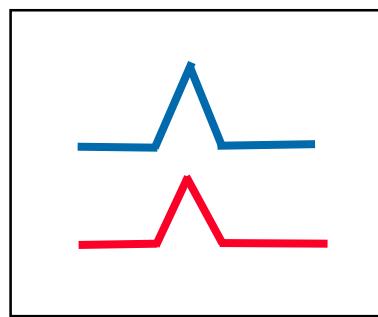
ThermoFisher
SCIENTIFIC

Other Tips & Information



ROX™ Passive Reference Dye

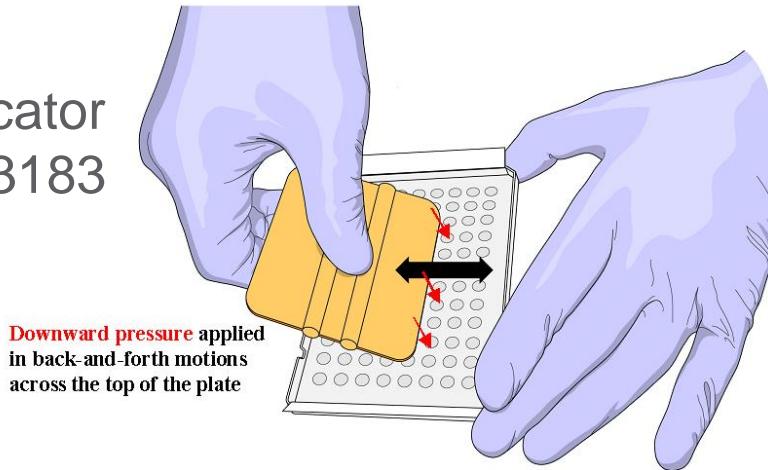
Improves precision of replicates by normalizing for non-PCR related variations.



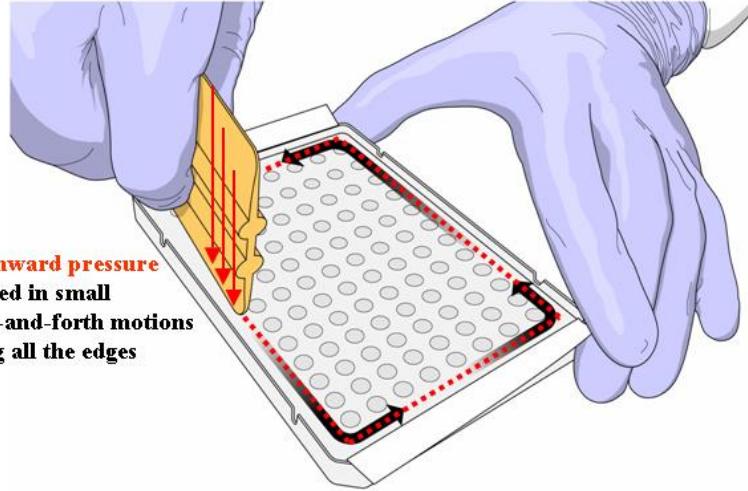
Prevent evaporation & bubbles during PCR

Use adhesive covers . . .

Applicator
#4333183



Downward pressure applied in back-and-forth motions across the top of the plate



Downward pressure applied in small back-and-forth motions along all the edges

SPIN PLATES before loading into instrument!!!



Guides to multiplexing

Real-time multiplexing

Part 1 - What is multiplexing? How does it work?

Multiplexing is the amplification of more than one target in one tube or well on a plate. Duplexing is specifically the amplification of two targets in one tube, triplexing is the amplification of three targets, and so on. Multiplexing is possible with TaqMan® probe-based assays, where each assay has a specific probe, and that probe is labeled with a different colored dye. The instruments can detect the different dyes and measure the signal from each one separately, and use that information to quantitate the amounts of different targets.

How many different dyes can the instruments detect?

The number of different dyes that can be detected varies by instrument. The Applied Biosystems 7500 Real-Time PCR System can detect up to 5 different dyes, and the 7000, 7300, 7700 and 7900 can detect up to 4. However, AB prefers to reserve one of these dyes for a passive reference dye such as ROX™ dye. See Figure 1 and Figure 2.

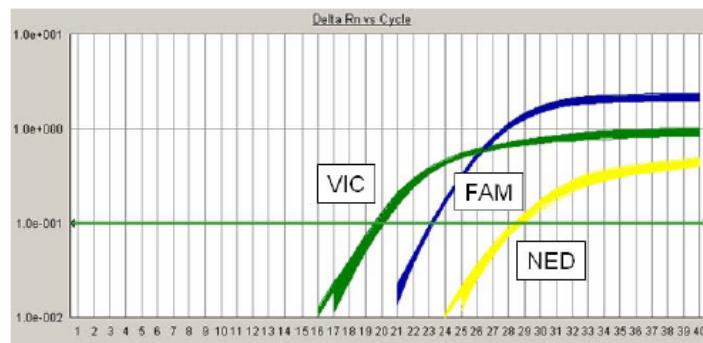


Figure 1 – Three color multiplexing on the AB 7300 instrument. Data is normalized to a fourth dye, ROX™ dye, as a passive reference.

User Bulletin #5

ABI PRISM® 7700 Sequence Detection System

August 10, 1998 (updated 01/2001)

SUBJECT: Multiplex PCR with TaqMan® VIC Probes

Overview Applied Biosystems now offers probes constructed with the new TaqMan® VIC reporter dye. The characteristics of the VIC dye make it an excellent candidate to replace existing TaqMan® JOE and HEX reporter dyes. The increased signal strength and improved spectral resolution also make VIC-labeled probes the ideal second probe for a multiplex PCR system.

This user bulletin describes the characteristics of VIC probes in relation to the existing JOE probes. It also contains guidelines for defining limiting primer concentrations in a one- or two-step multiplex reverse transcription-polymerase chain reaction (RT-PCR) system using VIC probes.

The following topics are covered in this user bulletin:

| Topic | See Page |
|--------------------------------------|----------|
| Characteristics of TaqMan VIC Probes | 2 |
| Multiplex RT-PCR | 5 |
| Technical Support | 13 |

IMPORTANT To use VIC probes on the ABI Prism® 7700 Sequence Detection System (SDS), you must first calibrate the instrument with the Sequence Detection Systems Spectral Calibration Kit (PN 4305822). This kit contains the new SYBR® Green and VIC fluorescent dye standards used to update the spectra components file in the SDS software. See User Bulletin #4: Generating New Spectra Components (PN 4306234).

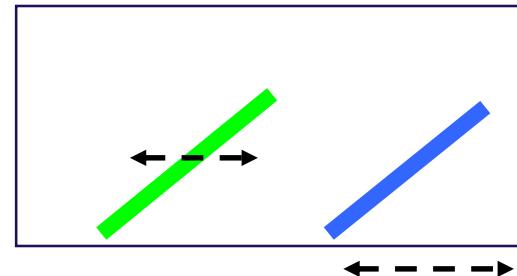
Note All documents referred to in this user bulletin are available through the Internet at the Applied Biosystems technical support documentation library or through Fax-on-Demand (see "To Obtain Documents on Demand" on page 16 for information).

The technical support documentation library is located at:

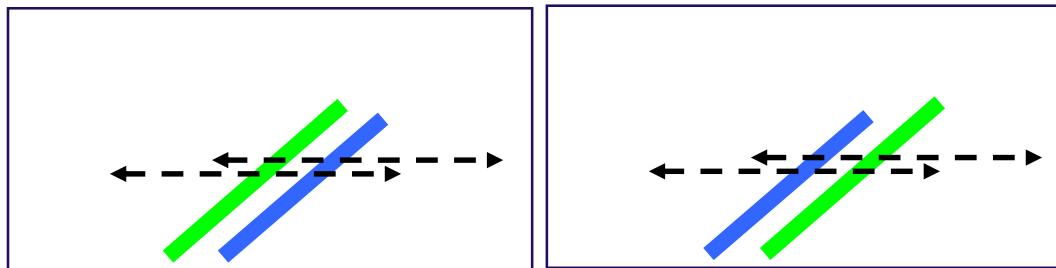
♦ www.appliedbiosystems.com/techsupport

Multiplex Scenarios

#1: One gene is more abundant.

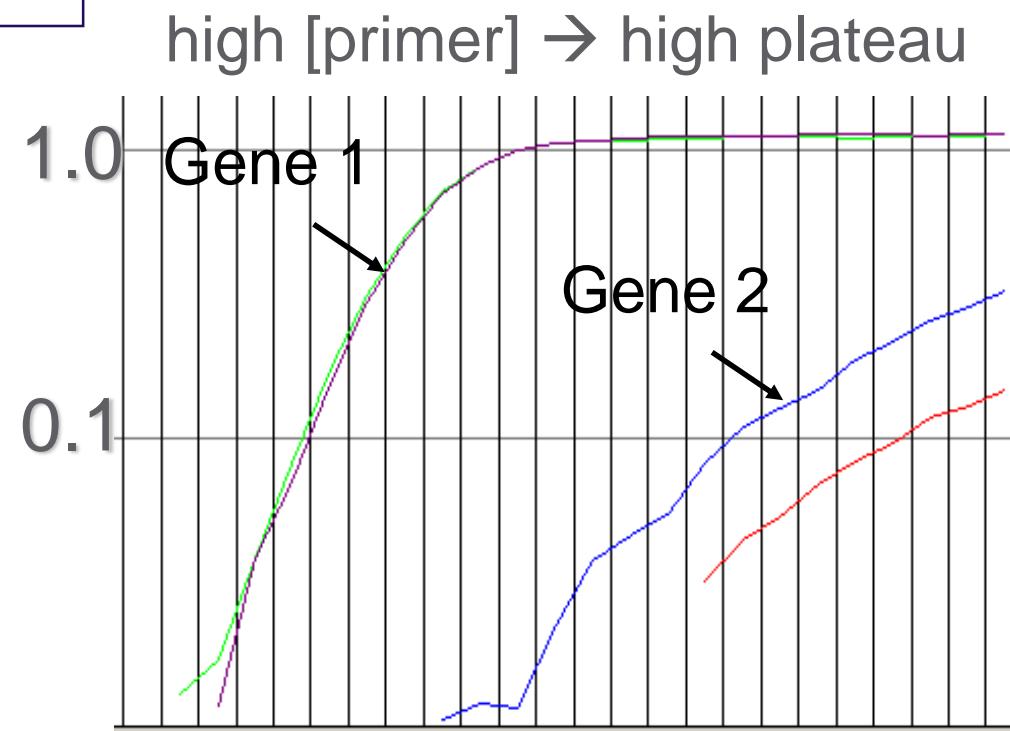
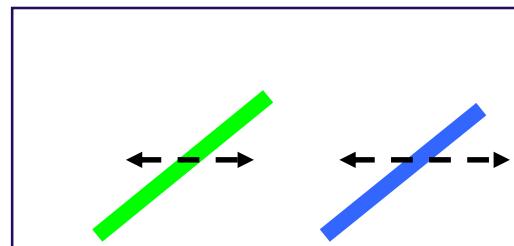


#2: Both genes are similar in expression

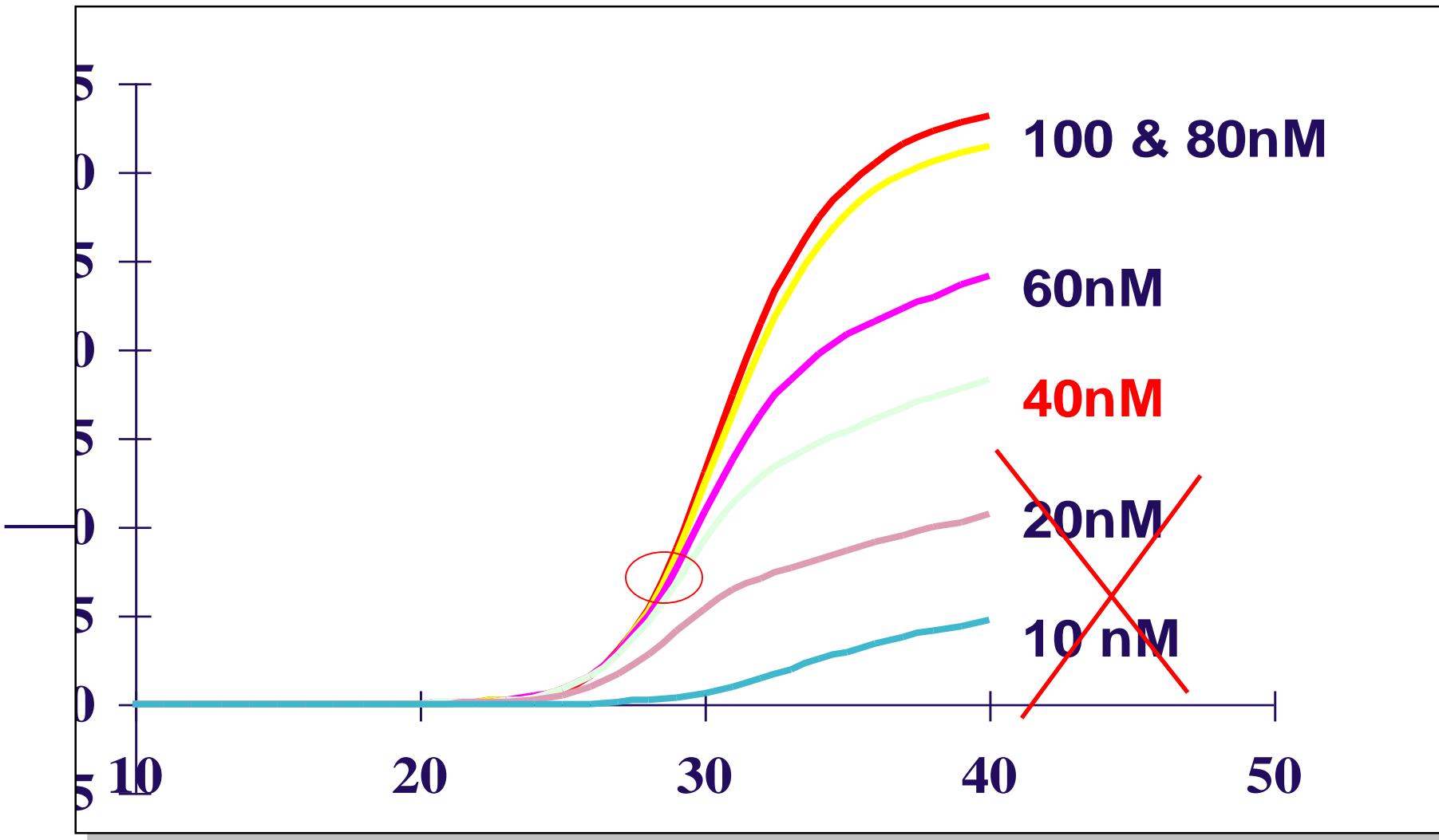


#1: One gene is more abundant

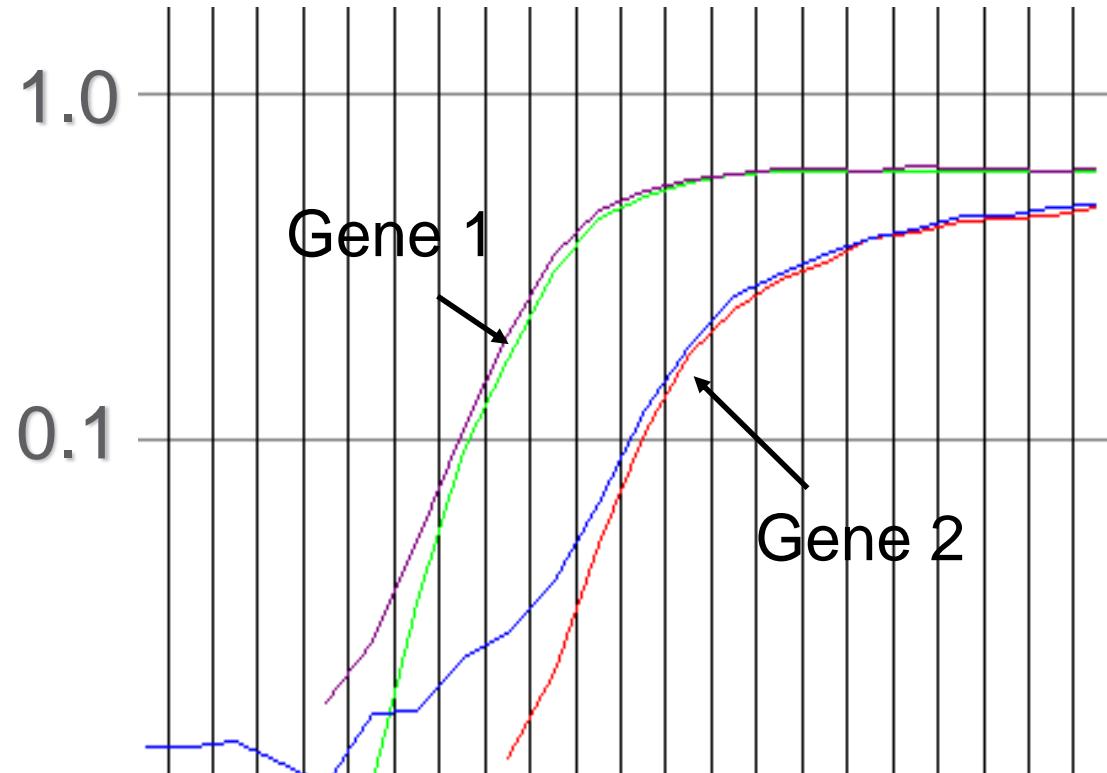
one gene uses all the reagents and there is nothing left for the other gene



Reduce primers of more abundant gene-Optimization



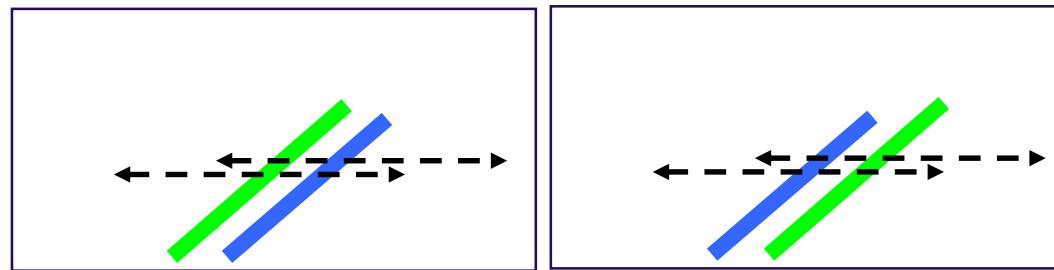
Solution:
lower the [primer] of the early expressed assay →
reaction plateau quickly



- Exponential phase intact for both assays
- “primer limited” assays
- Order primers & probe separately

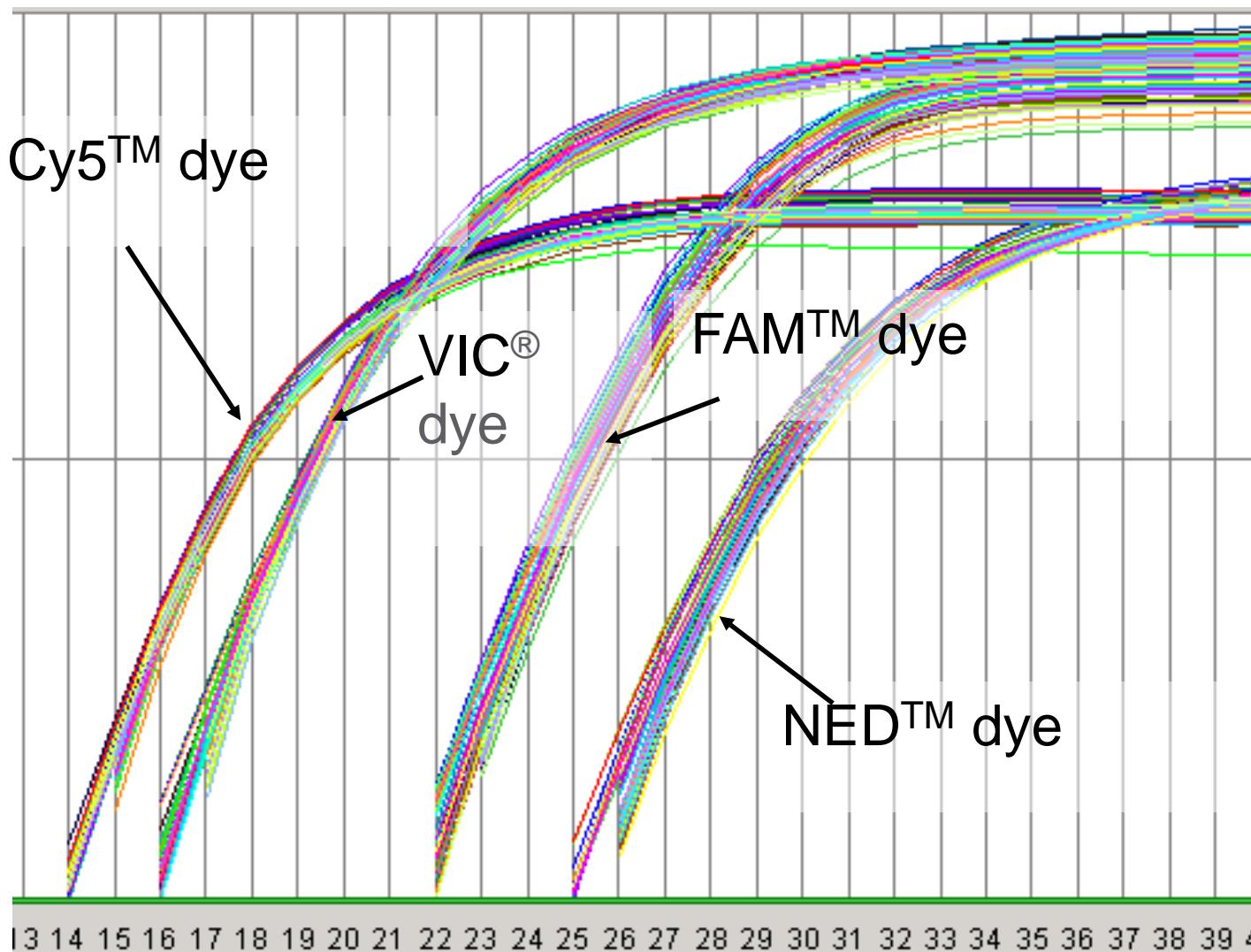
#2: Low stress multiplexing

both genes are similar in relative abundance



- Run in singleplex then duplex and ensure valid exponential phases and CTs are achieved
- If validation fails, primer limit one or both assays

Multiplexing becomes exponentially difficult with every additional assay (triplex, quadplex...)



Service Plans

Please contact Scott.Gardner@thermofisher.com for more details

Your qPCR system includes a 1 year manufacturers warranty. If a service plan is purchased, it will lock you into today's pricing and will start at the conclusion of the manufacturers warranty.

Discounts are available for multi-year coverage, locks you in at today's pricing.

Benefits of AB Assurance Service Plan

- Reduces downtime by providing proactive maintenance service
- Includes parts, labor, & travel at no additional cost
- Priority on-site guaranteed 2-day response time, & priority access to remote service engineer
- Scheduled on-site planned maintenance (PM) includes:
 - Calibration services (cost of the calibration kit is included)
 - Additional tests to ensure system performance
 - Computer repair and replacement



Support
(800) 955-6288
techsupport@thermofisher.com

Scott.gardner@thermofisher.com

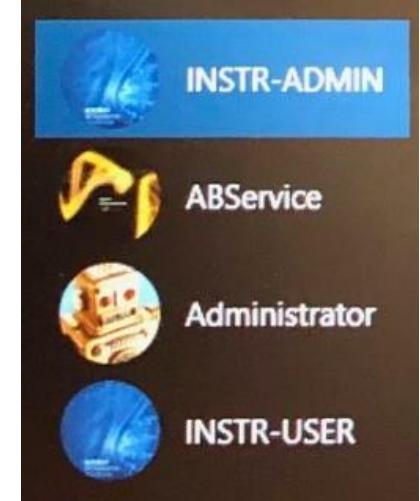
Instrument software

Computer login (default),

user name & password are the same

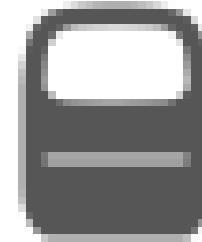
INSTR-ADMIN

INSTR-ADMIN



Free Software Download (link in follow-up email)

- .edt template
- .eds single data file



Merge multiple .eds files in ThermoFisher Connect or
TaqMan Genotyper (for SNP data)