

The Ups & Downs of Gene Expression:

Using Lipid-Based Transfection and RT-qPCR to Deliver Perfect Knockdown and Achieve Optimal Expression Results

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BIO-RAD

Topics

- What is RNAi?
- Methods of Delivery and Detection
- RNA Preparation
- Reverse Transcription
- qPCR Detection
- Case Study: ODC Pathway



What is RNAi?

RNA interference (RNAi) is a phenomenon where dsRNA specifically blocks the expression of its homologous gene. Also known as post-transcriptional gene silencing (PTGS) and quelling.

1990 RNAi was discovered as an endogenous property in petunias

1998 Fire & Mello at the Carnegie in Washington showed gene silencing pathway in c.elegans

2000 Tuschl and Elbashir at the Max Planck Institute showed that short interfering RNAs could be introduced into mouse cells.



The Power of RNA Interference

Why is RNAi so powerful?

- Allows fast characterization of gene / protein function
- Enables study of pathways
- Facilitates rapid identification and validation of targets
- Therapeutic potential

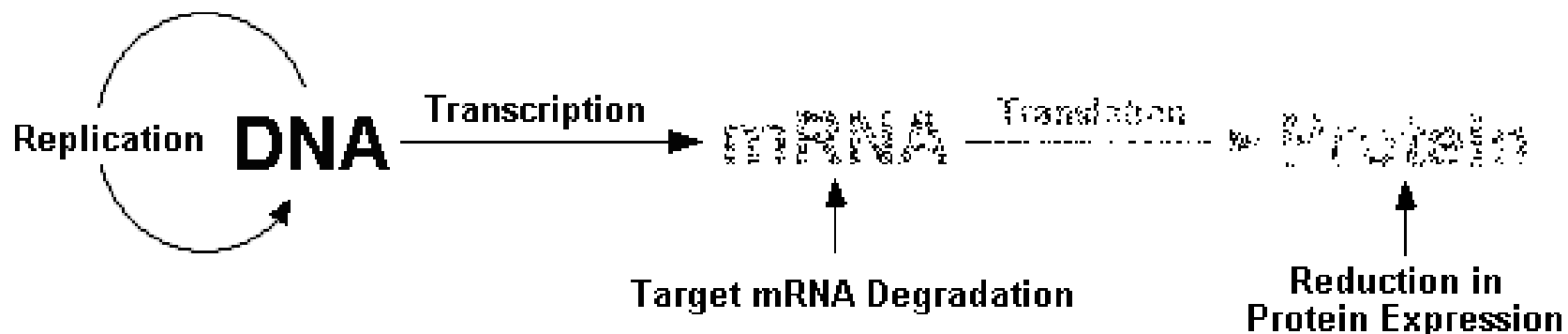


Molecular Biology and RNAi

Central Dogma of Molecular Biology:



Basic RNA interference Mechanism:



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RNAi: Challenge of Delivery

What delivery method is best?

- Electroporation – good for suspension & difficult cells
- Biolistics – good for neural & primary cells
- MicroInjection – offers greatest specificity
- Viral – very high efficiency
- Lipid Mediated – low cost, simple protocol, consistent results, good for high throughput applications



Lipid Mediated Delivery

Three Major Lipid Characteristics to Consider:

- Design / Development
- Efficiency
- Toxicity

Silencing (siRNA Activity)

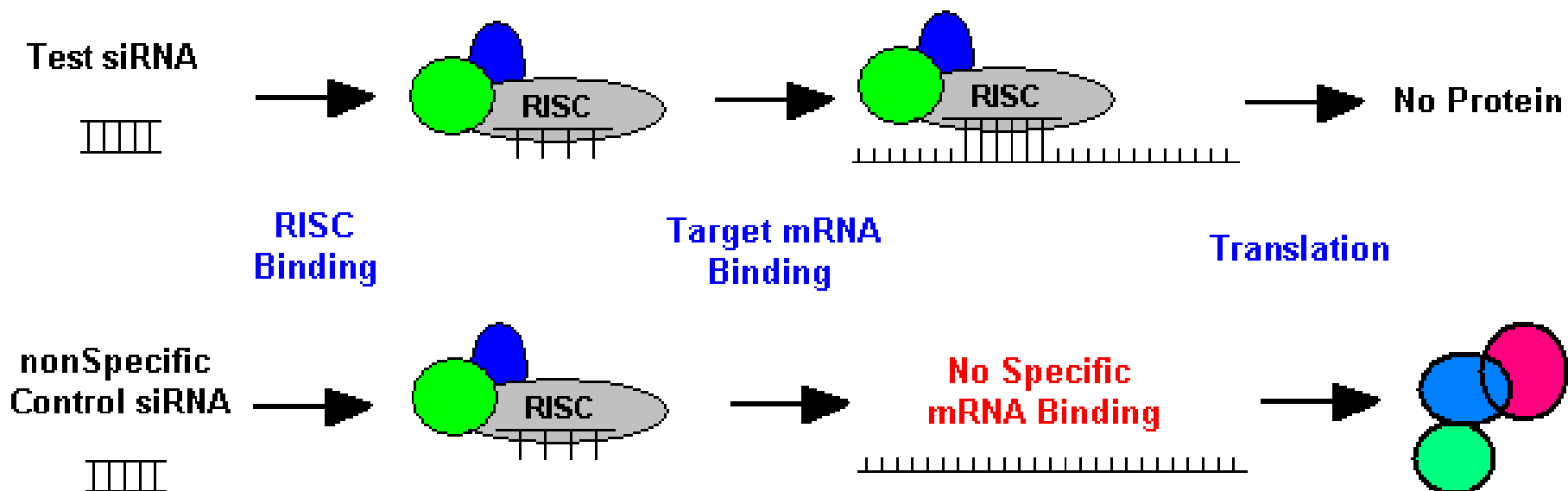


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graph TD; A[Design / Development] --> D[Silencing (siRNA Activity)]; B[Efficiency] --> D; C[Toxicity] --> D;
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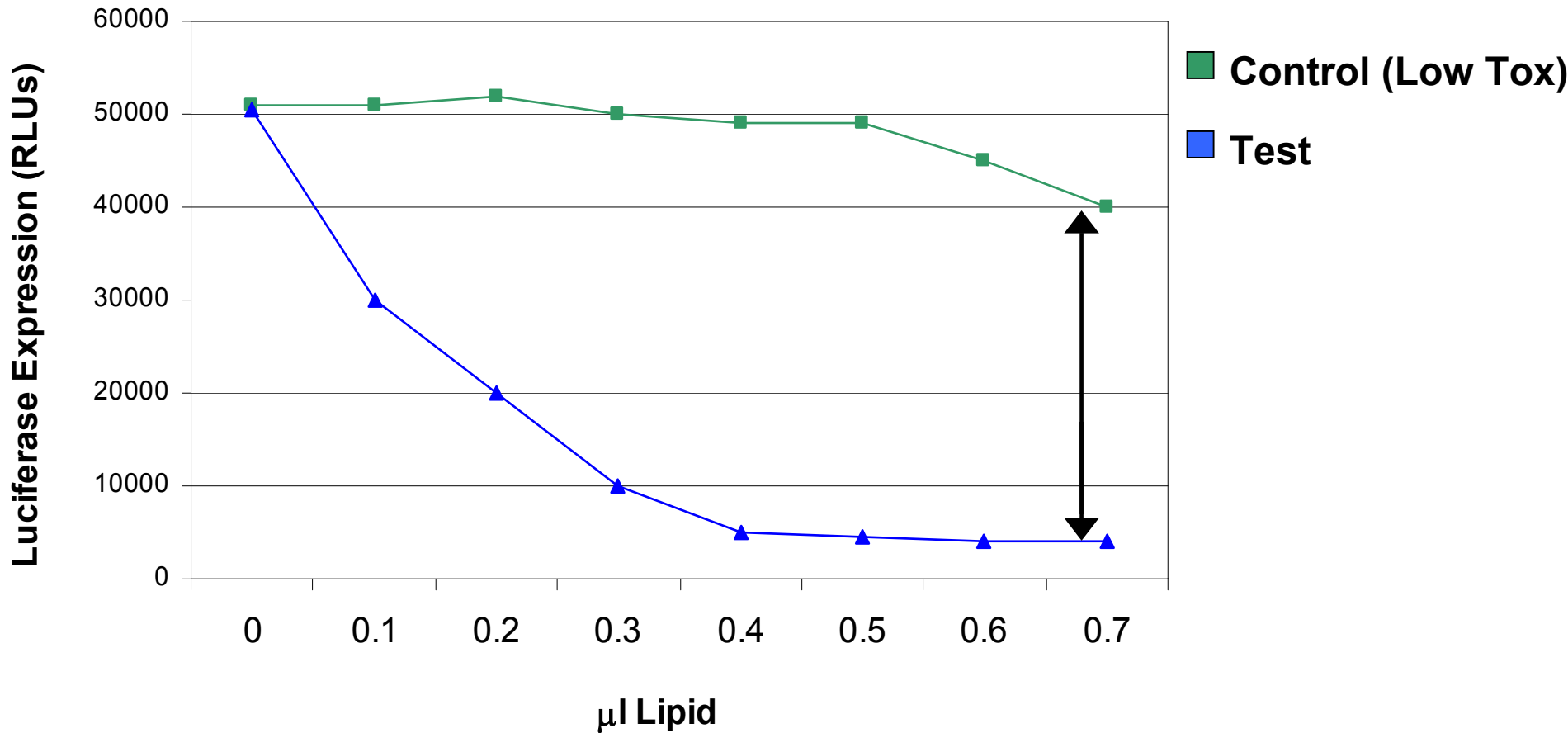
Experimental Design: Controls

Test siRNA vs. nonSpecific Control siRNA



Experimental Design: Controls

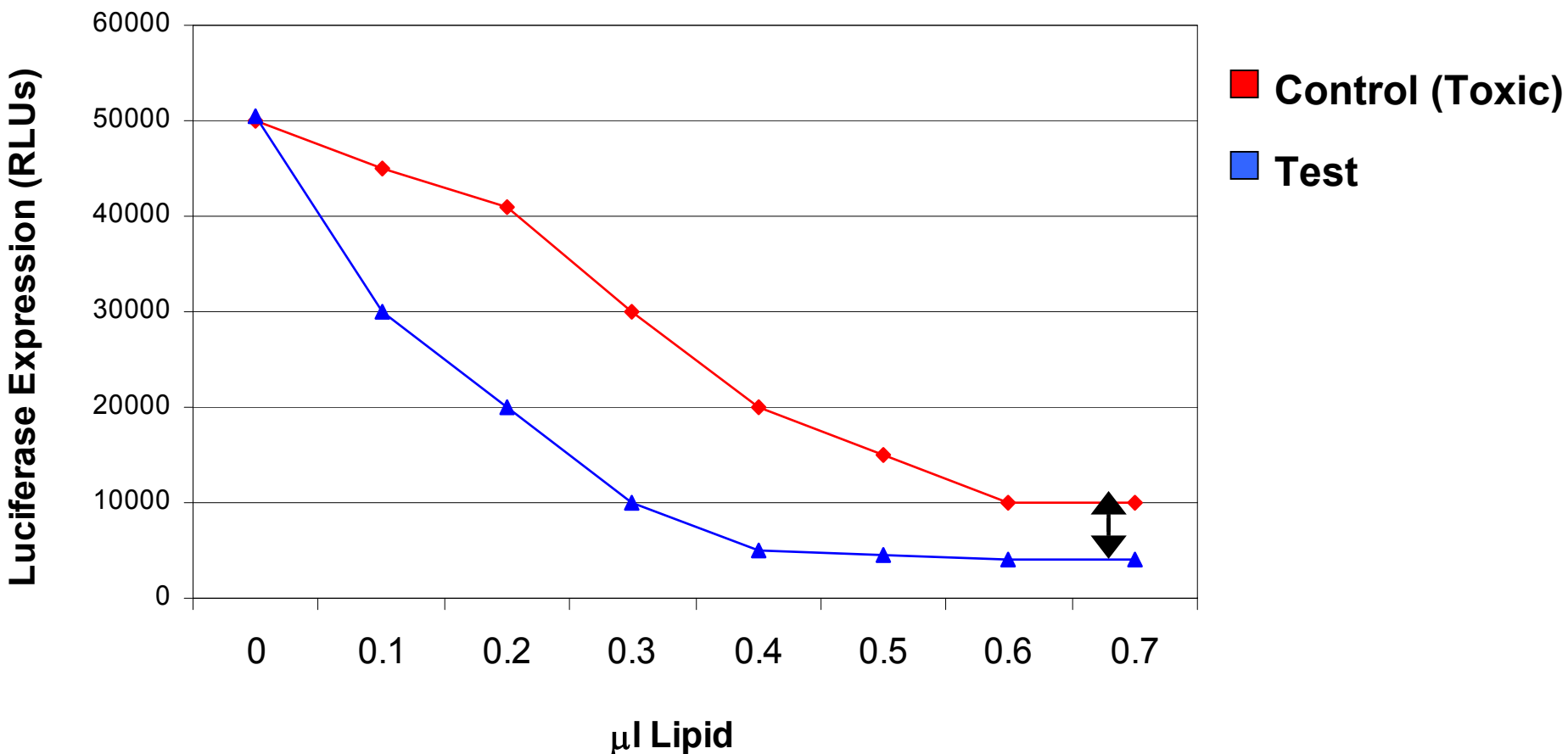
How this will look as Data.....



Experimental Design: Controls

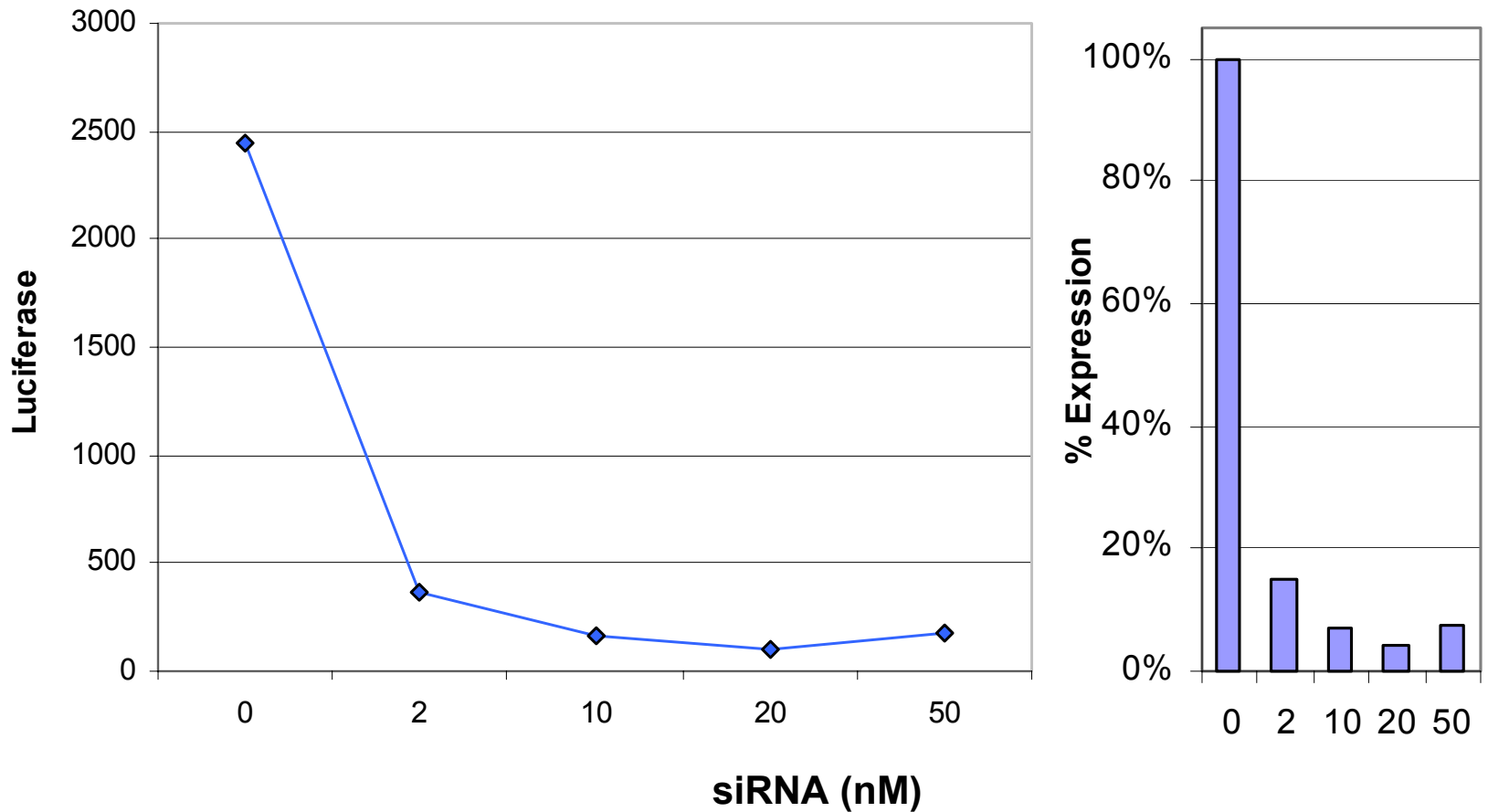


How this will look as Data.....



Efficiency: siRNA Amount

CHO-Luc / siLentFect – 0.3 μ l (96-well)

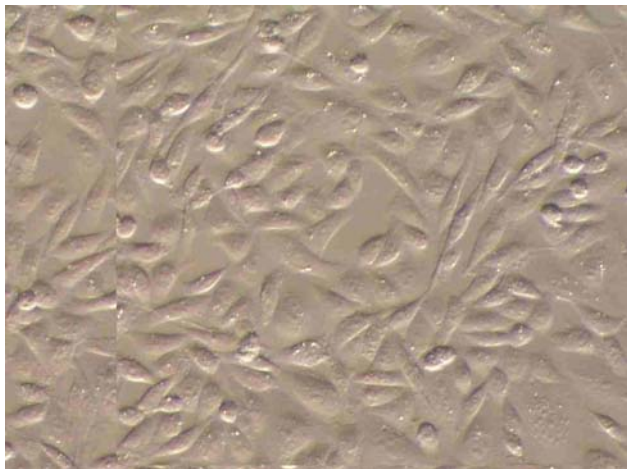


Toxicity Evaluations

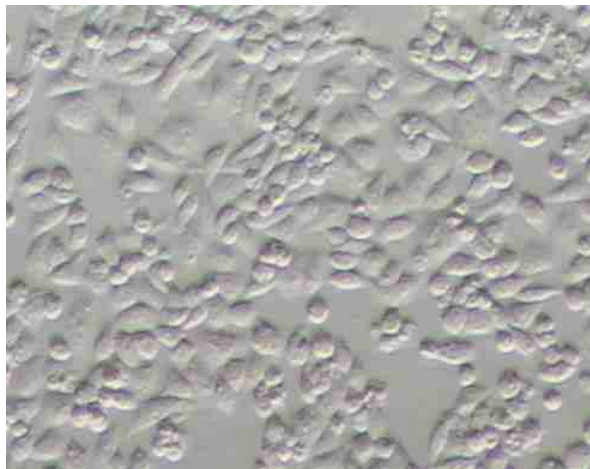
Visual Analysis

- Morphology changes
- Detachment
- Lysis

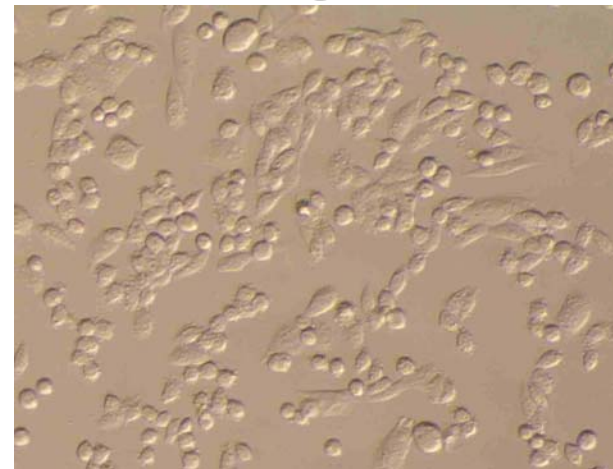
Low



Moderate

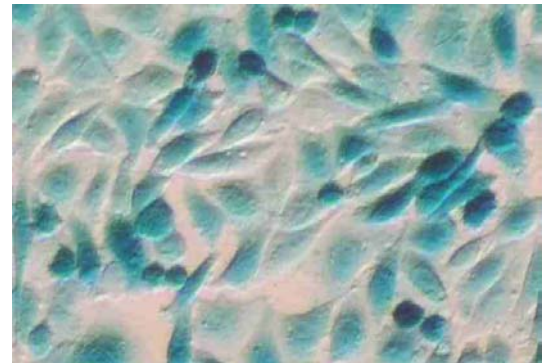
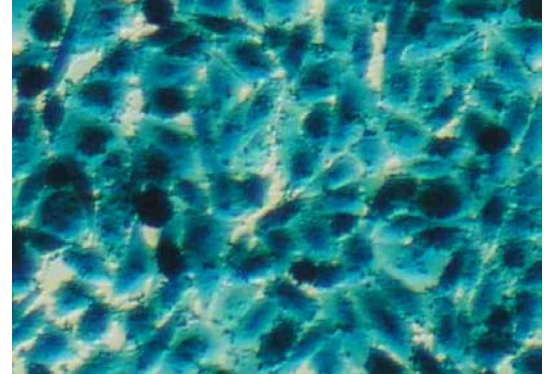


High



RNAi Detection Strategies

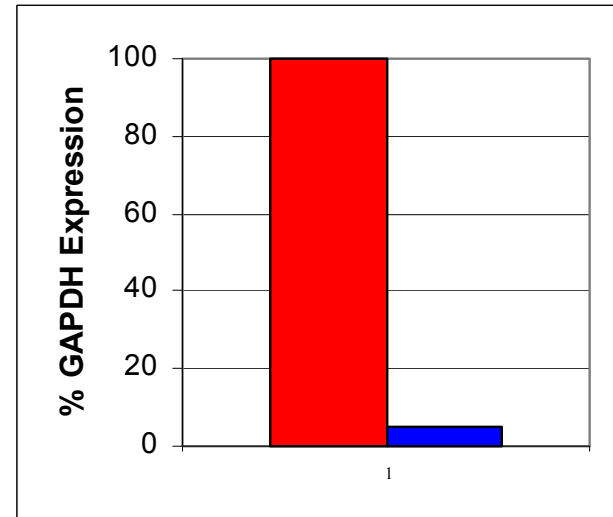
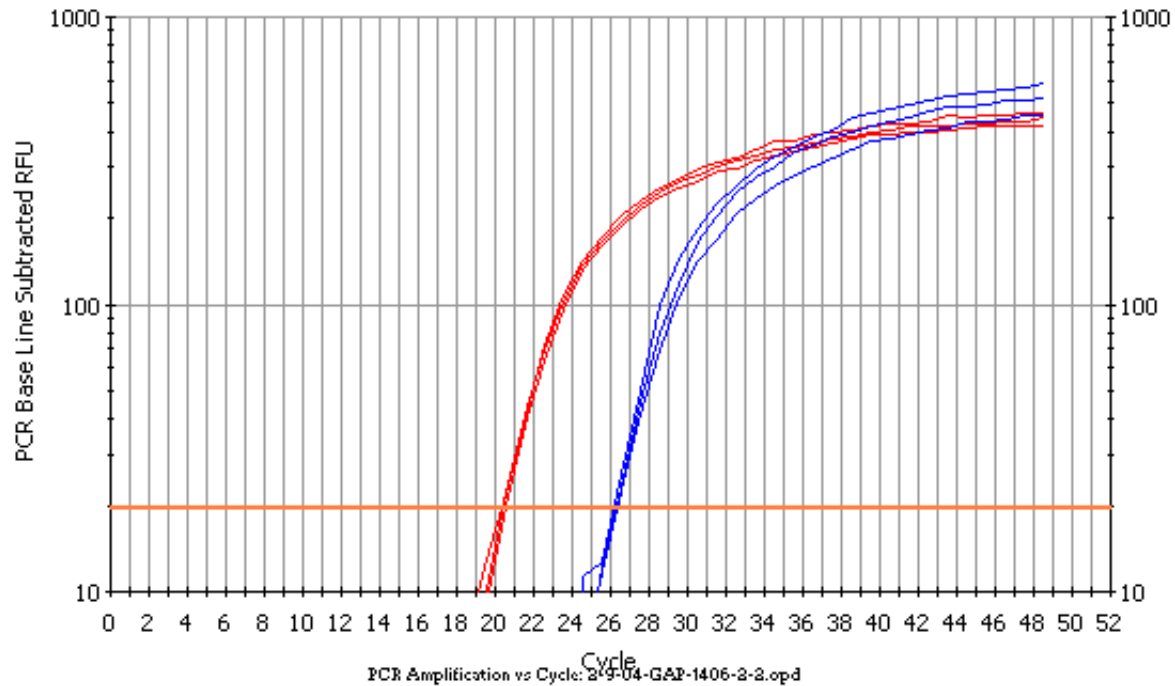
- Western Blots
- Northern Blots
- MicroArrays
- qPCR
 - 1.0 Cycle Threshold = 50% silencing
 - 3.3 Cycle Threshold = 90% silencing
 - 6.6 Cycle Threshold = 99% silencing



CHO-lacZ cells transfected with scrambled siRNA control (top) and beta-gal siRNA (bottom)

Detection: qPCR Analysis

GAPDH, Primary Fibroblasts, 48 hr, 6-well

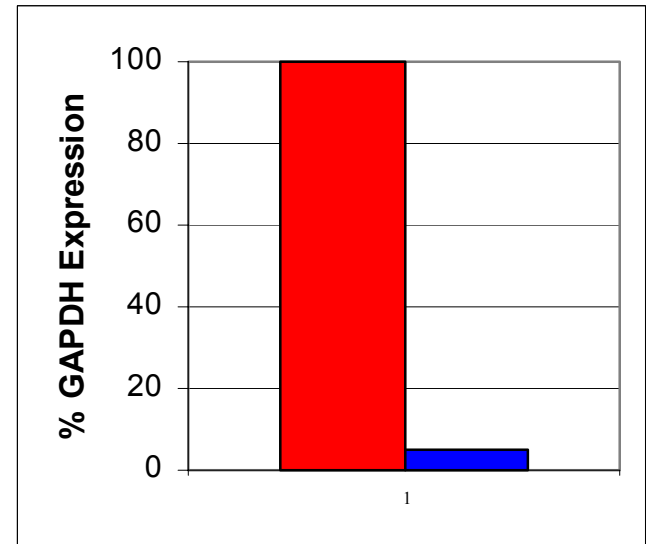
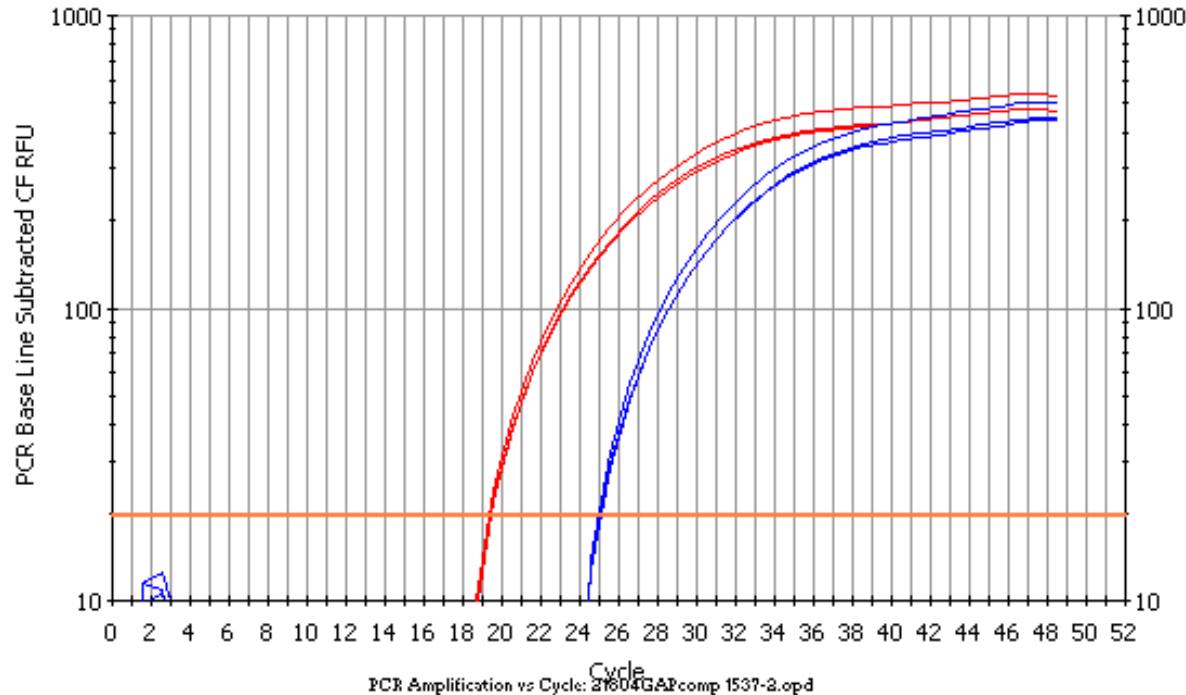


- 5.9 C_t Difference
- Over 95% knockdown
- 1.25 μ l siLentFect
- 10 nM siRNA



Detection: qPCR Analysis

GAPDH, HeLa Cells, 48 hr, 6-well



- 5.6 C_t Difference
- Over 95% knockdown
- 1.25 μ l siLentFect
- 10nM siRNA



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RNA Preparation

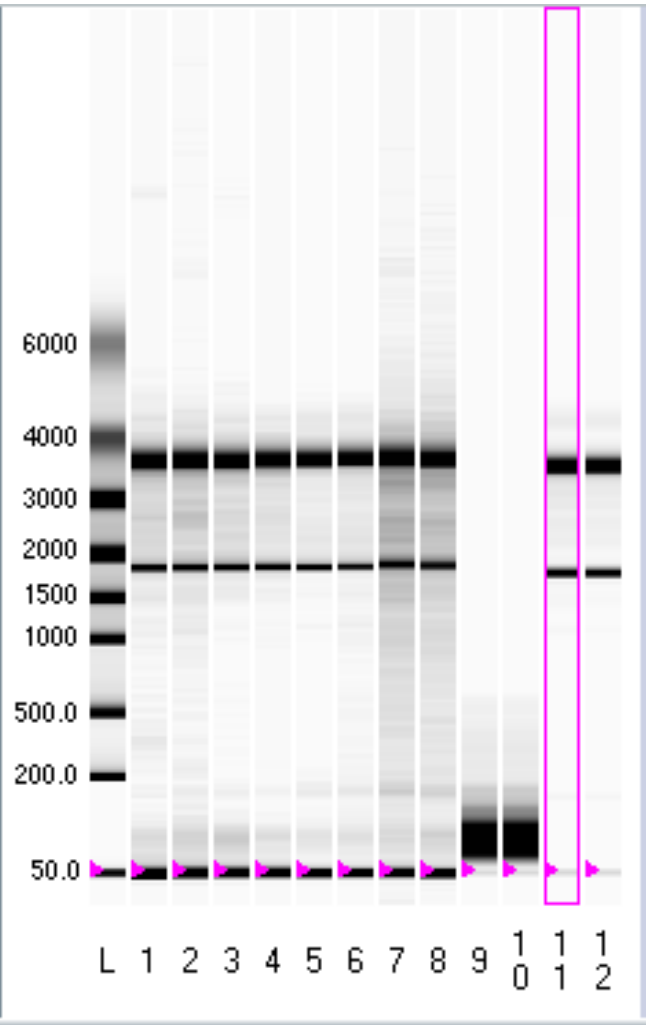
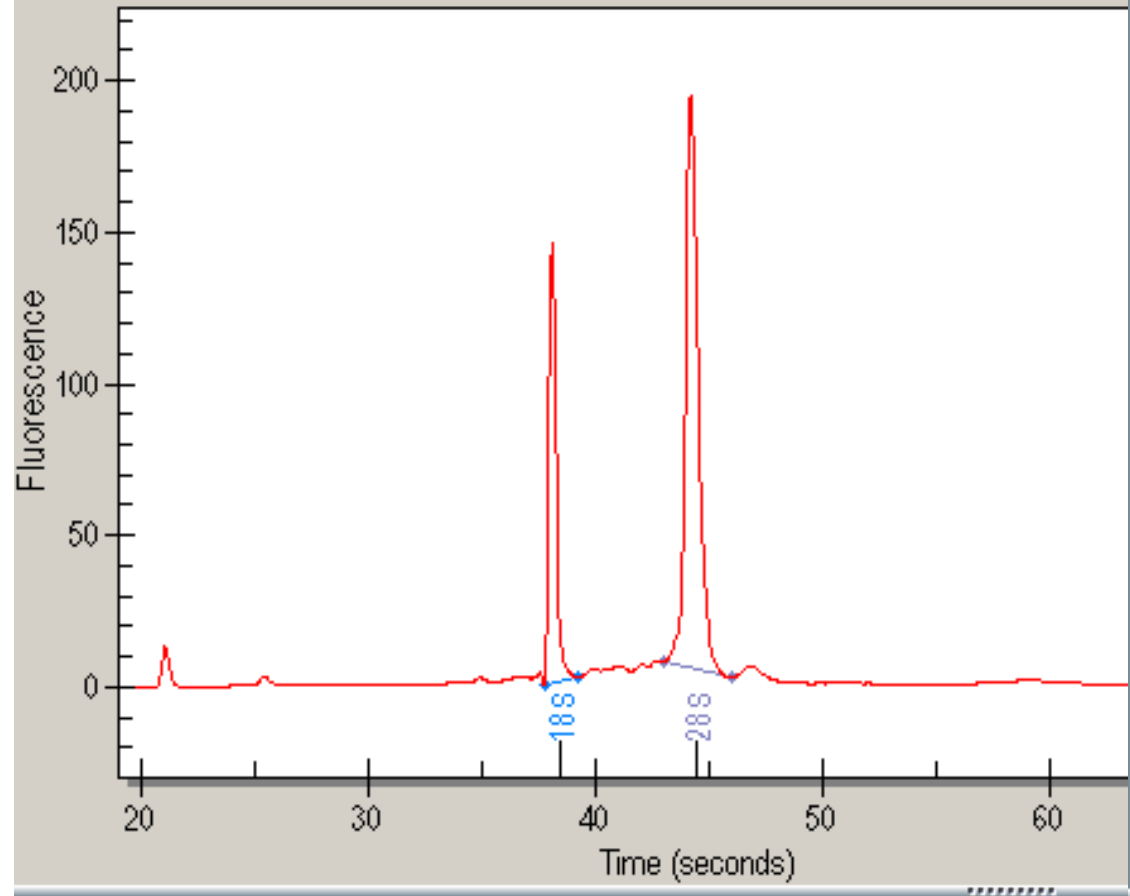
- Extract RNA (DNase treatment optional)
- Analyze RNA, careful quantification is necessary:
 - RiboGreen assay
 - Experion™ System



Experion System Data



Molgen2_TotalRNA_001072_2-23-2005_12-31-19 PM
Jurkat1:3-1

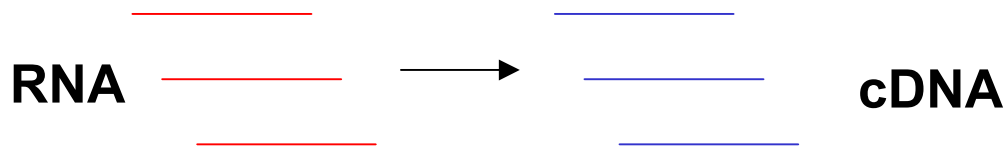


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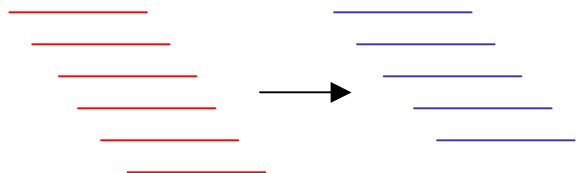
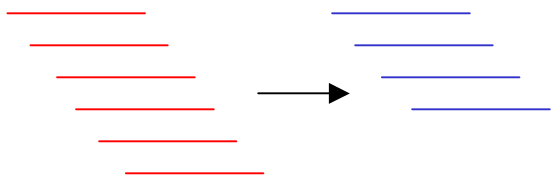
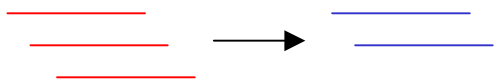
Testing the Reverse Transcriptase



RT Efficiency: Its Effect on the Assay

RNA  →  cDNA

Reliability ?

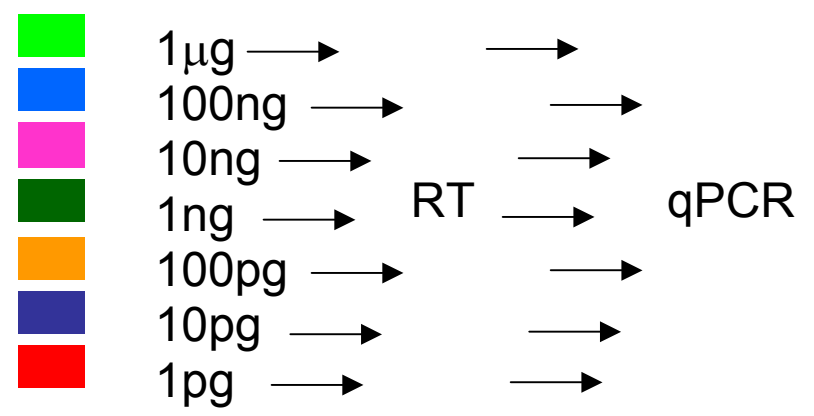
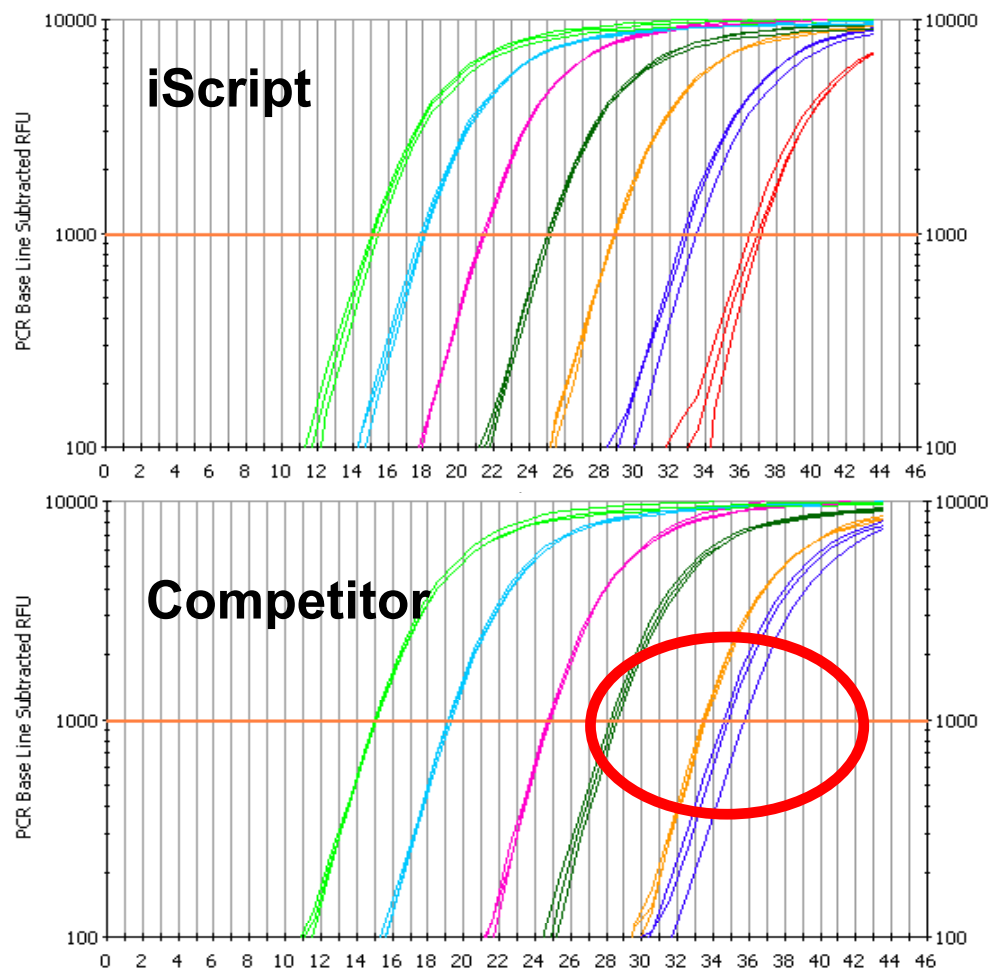


Reproducible
Data

Not
Reproducible



Reproducibility of RT

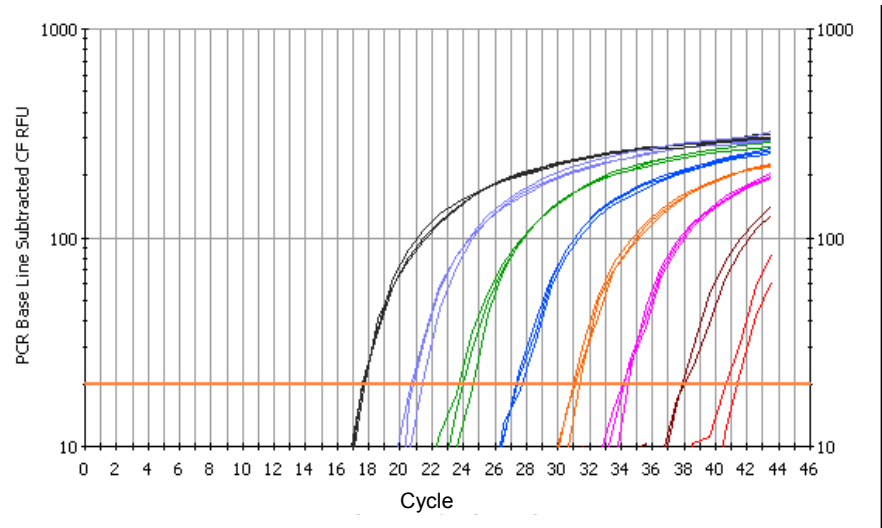


No discrimination at low Concentration
No detection at 1 pg



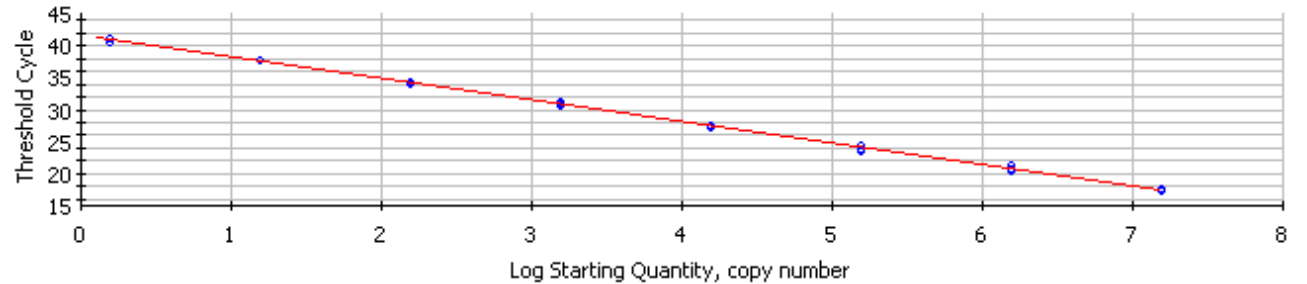
Dynamic Range of iScript

- 1.6×10^7
- 1.6×10^6
- 1.6×10^5
- 1.6×10^4
- 1.6×10^3
- 1.6×10^2
- 1.6×10^1
- 1.6×10^0



Correlation Coefficient: 0.999 Slope: -3.360 Intercept: 41.674 $Y = -3.360 X + 41.674$
PCR Efficiency: 98.4 %

□ Unknowns
○ Standards



PCR Standard Curve: kp120103.opd



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What makes for a good qPCR?

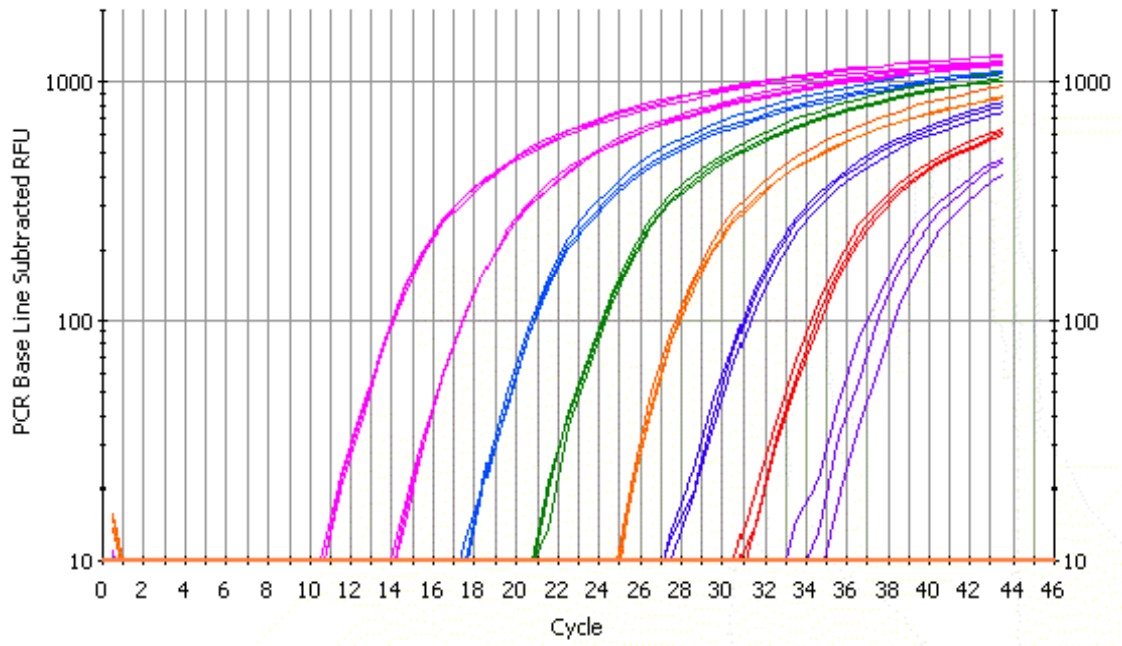
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- High Sensitivity
- Good Reproducibility
- Broad Dynamic Range



Dynamic Range of One-Step RT-qPCR

Beta-actin target, FAM-labeled probe, 1 μ g to 100fg input



$r = 1.000$, slope = -3.39 , efficiency = 97.2%

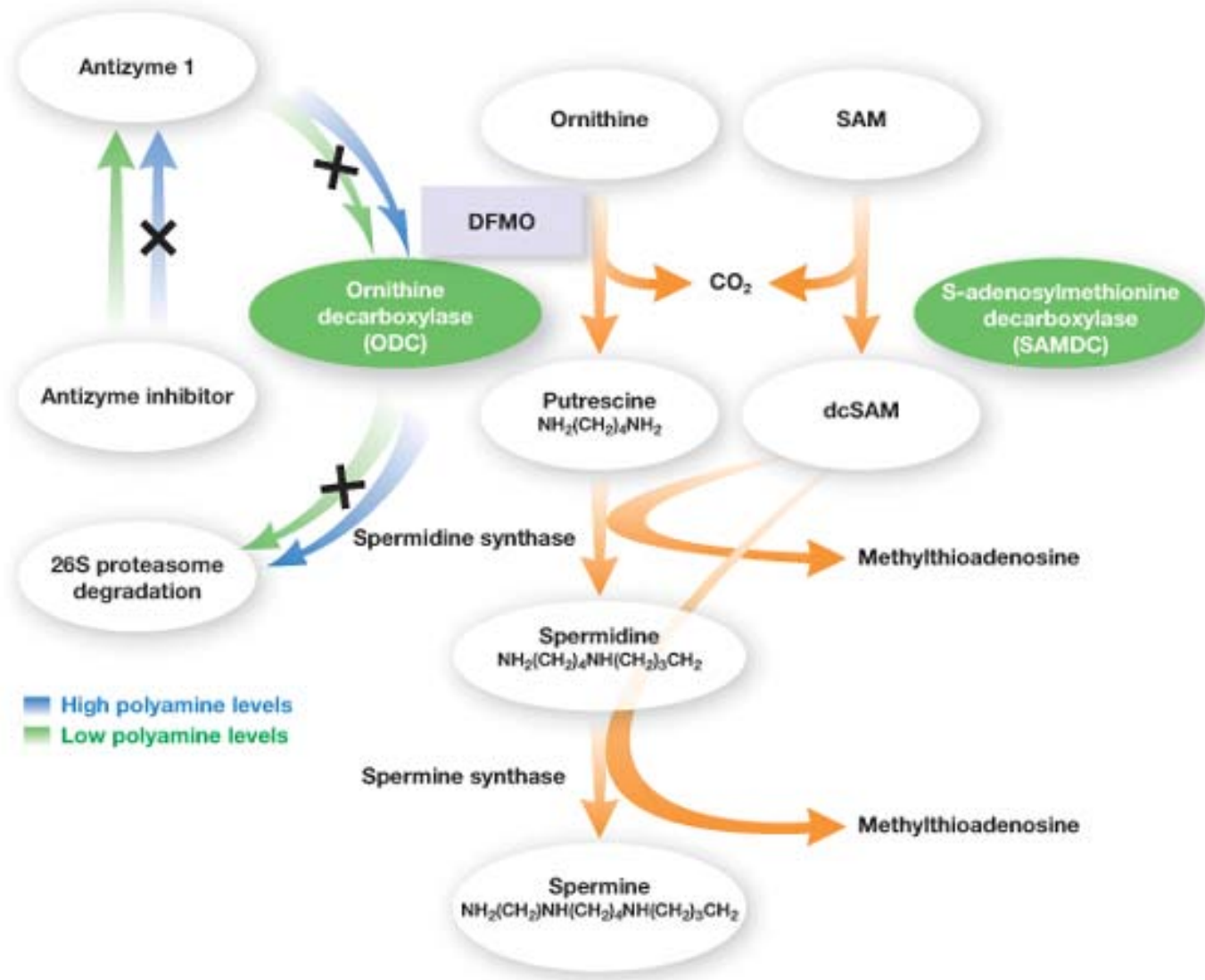


Topics

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Case Study: Polyamine Pathway



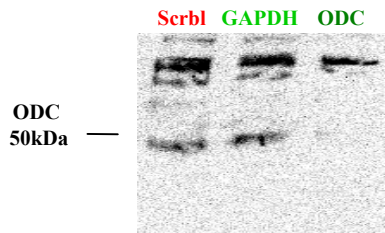
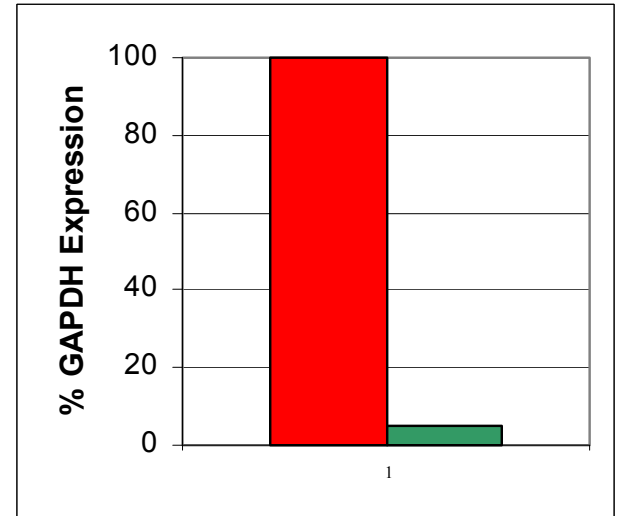
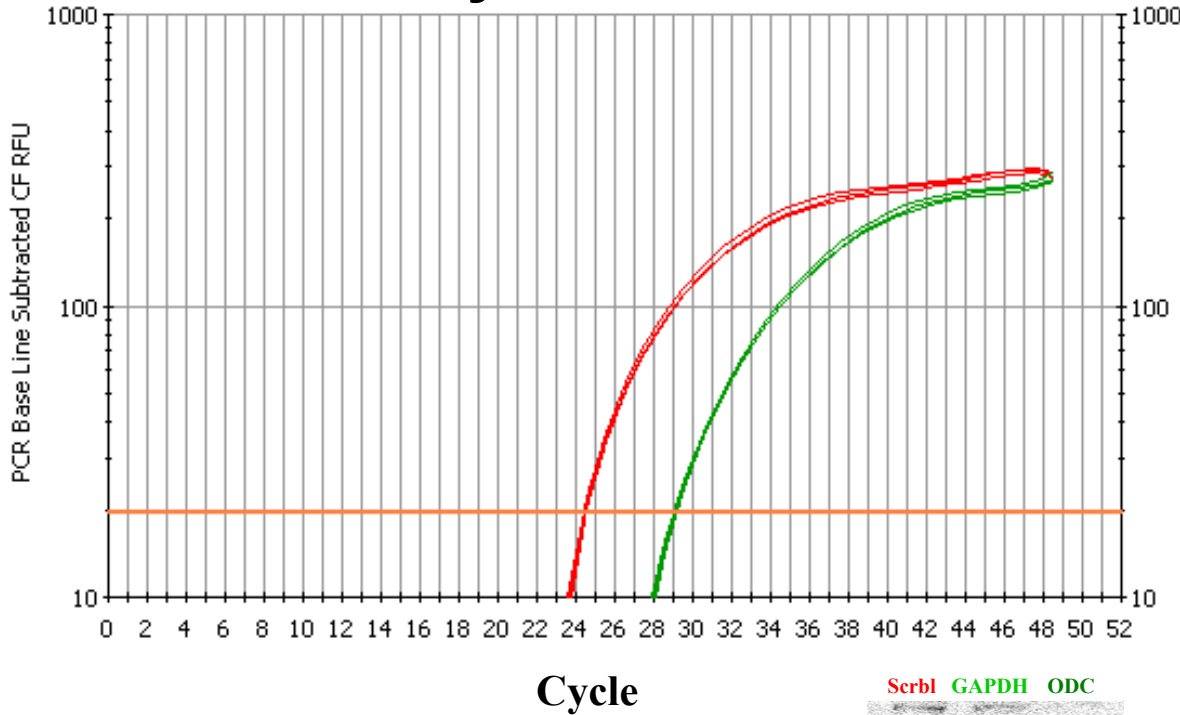
Points of Regulation

Central Dogma of Molecular Biology:



Down regulation of ODC

ODC, Primary Fibroblasts, 48 hr, 6-well

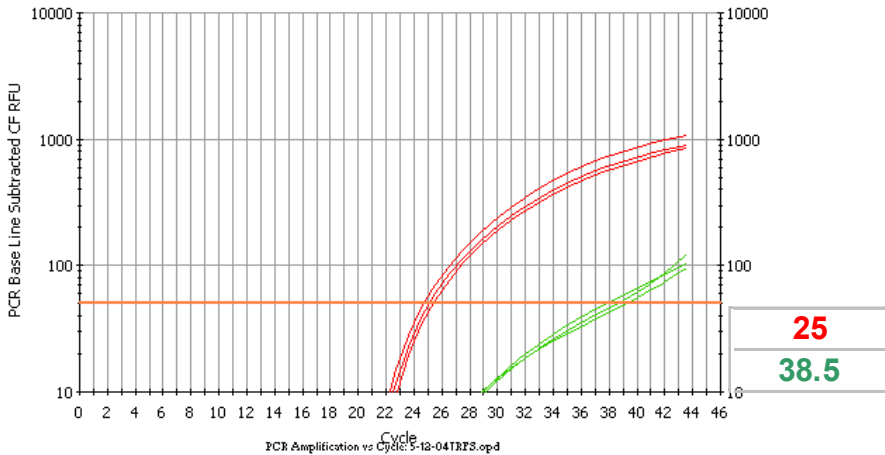


- 4.7 CT Difference
- Over 90% knockdown
- 2 μ l siLentFect
- 10 nM siRNA

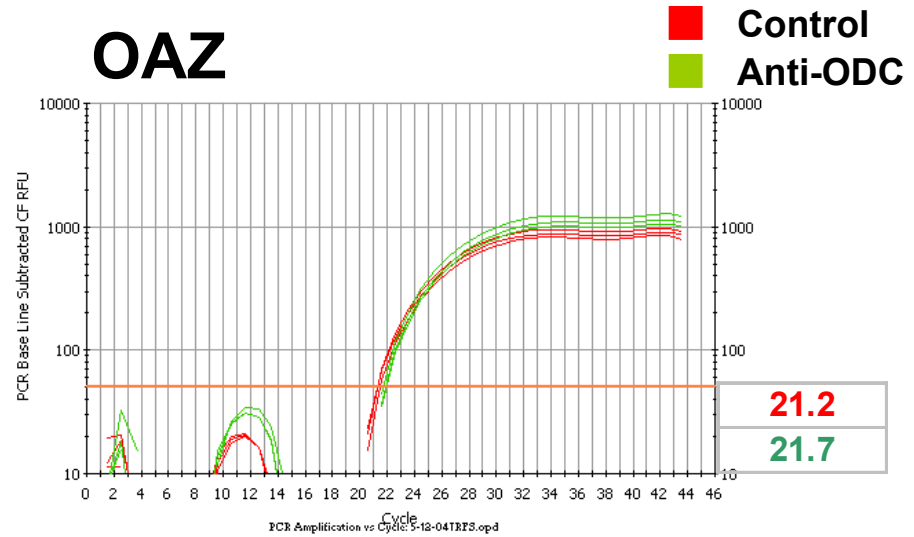


Effect of ODC Down Regulation

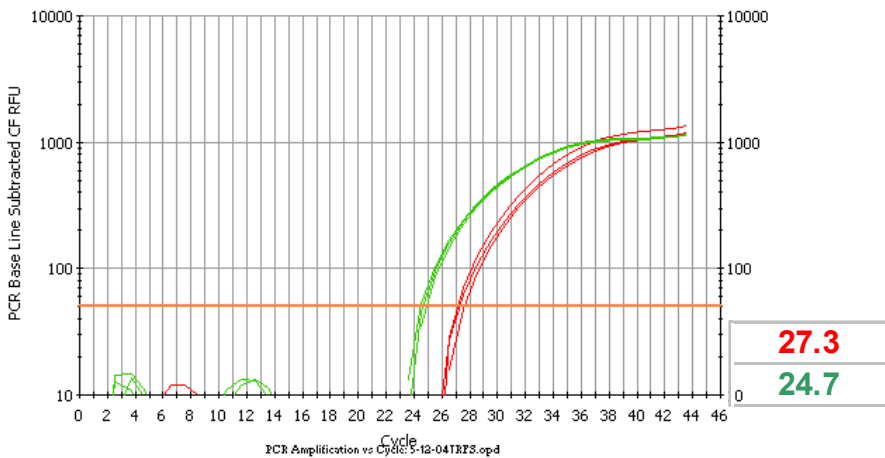
ODC



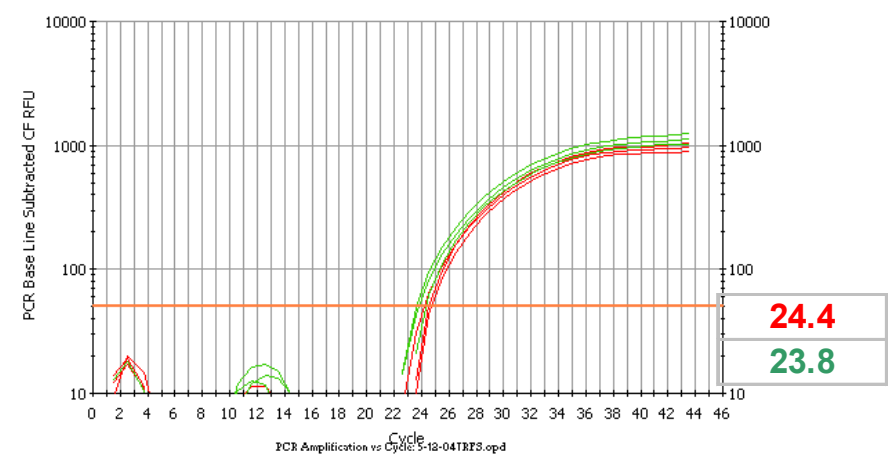
OAZ



SAMDC



AZI

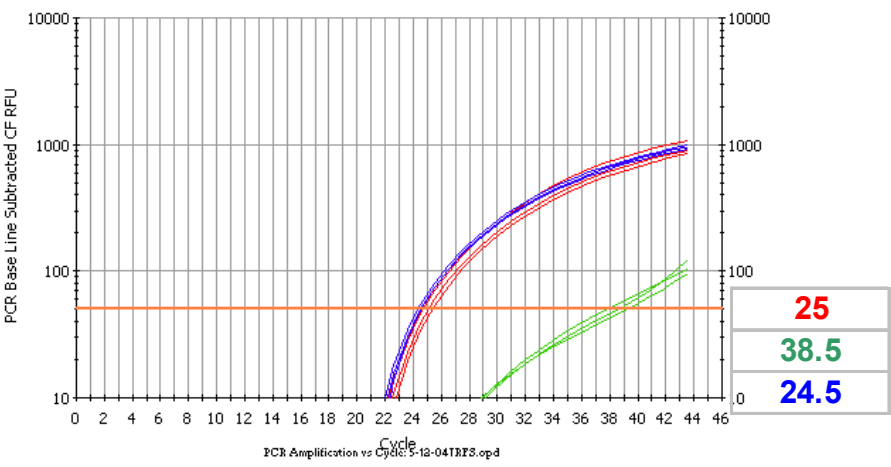


Effect of DFMO Treatment

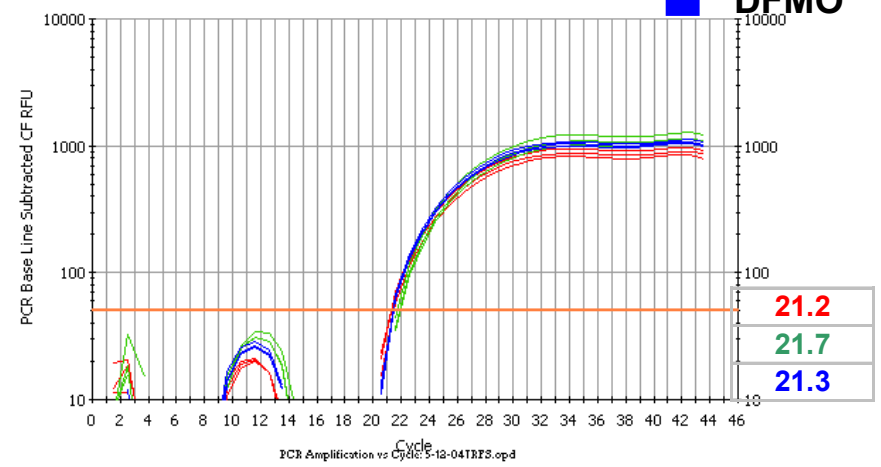


- Control
- Anti-ODC
- DFMO

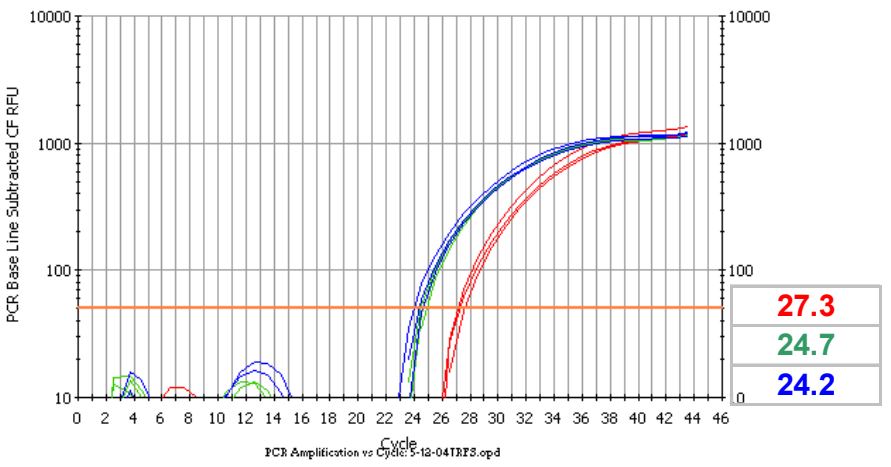
ODC



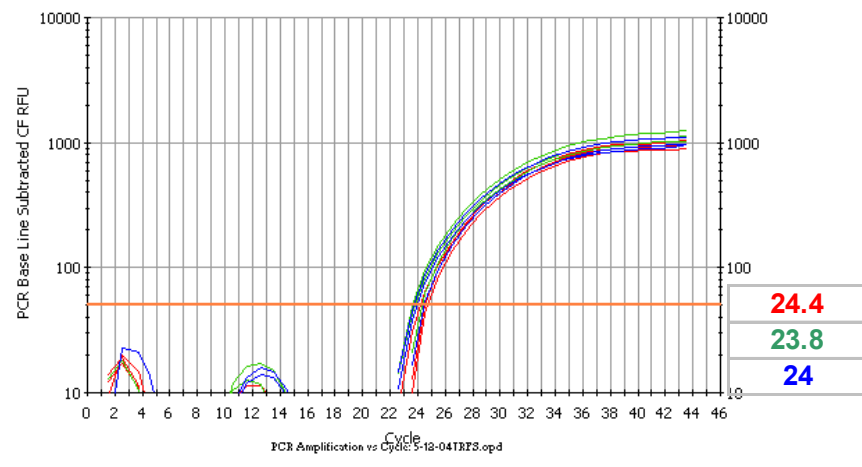
OAZ



SAMDC



AZI



Summary

- Transfection of primary fibroblasts with anti-ODC siRNA
 - results in a reduction of cellular ODC protein levels
 - results in up regulation of SAMDC transcript levels
 - regulatory enzymes OAZ and AZI were not affected (at the level of mRNA)
- Application of DFMO, which inactivates ODC protein
 - does not affect ODC transcript levels
 - results in the up regulation of SAMDC transcript levels



Summary continued

RNAi: Perfect Knockdown

- Choose a high quality RNA purification method (garbage in = garbage out)
- Good RT is critical to accurate transcript quantification
- Use a good, quantitative detection method: qPCR provides a fast, accurate, sensitive method for RNAi analysis

