

Principles of Real Time PCR

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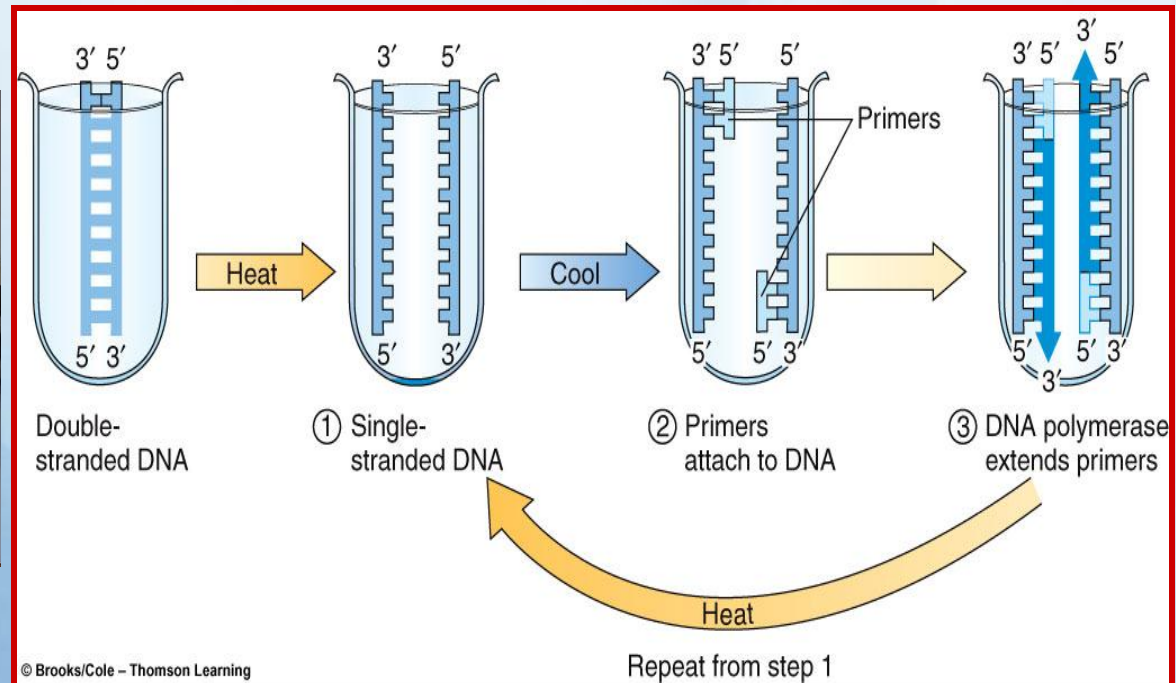
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Types of PCR

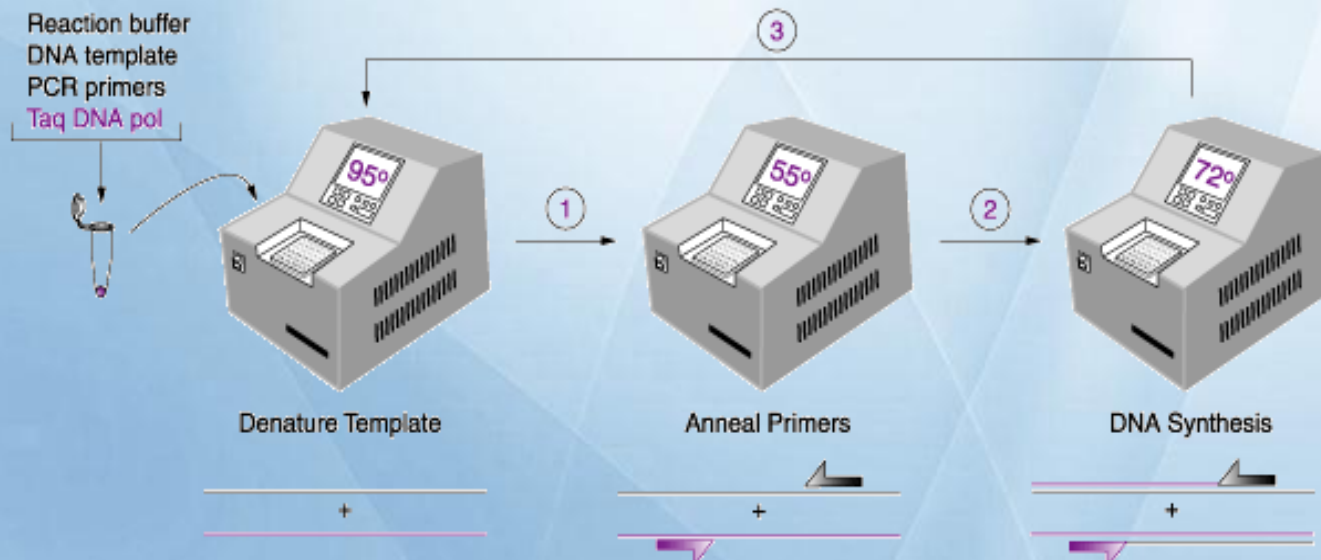
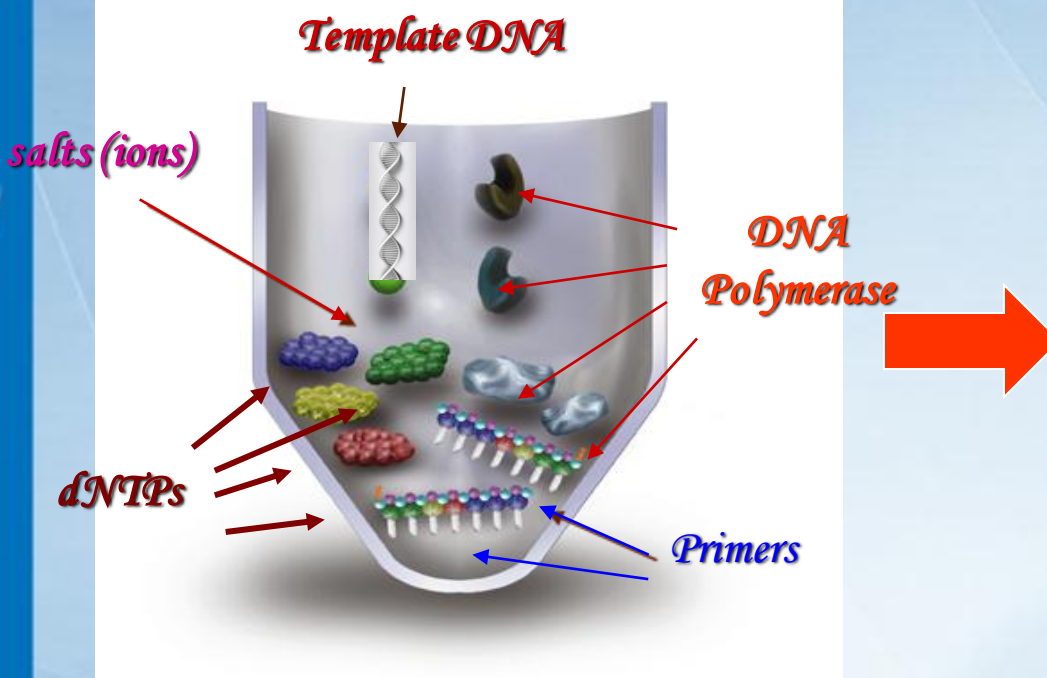
- **Standard PCR (conventional)**
- **RT-PCR (Reverse Transcriptase PCR)**
- **Real Time PCR (qRT-PCR)**

Conventional PCR

- **Each cycle (Round) of PCR contains 3 steps:**
 - 1- Denaturation**
 - 2- Primer annealing**
 - 3- Primer extension**
- **The cycle usually repeated for 25 – 40 times.**

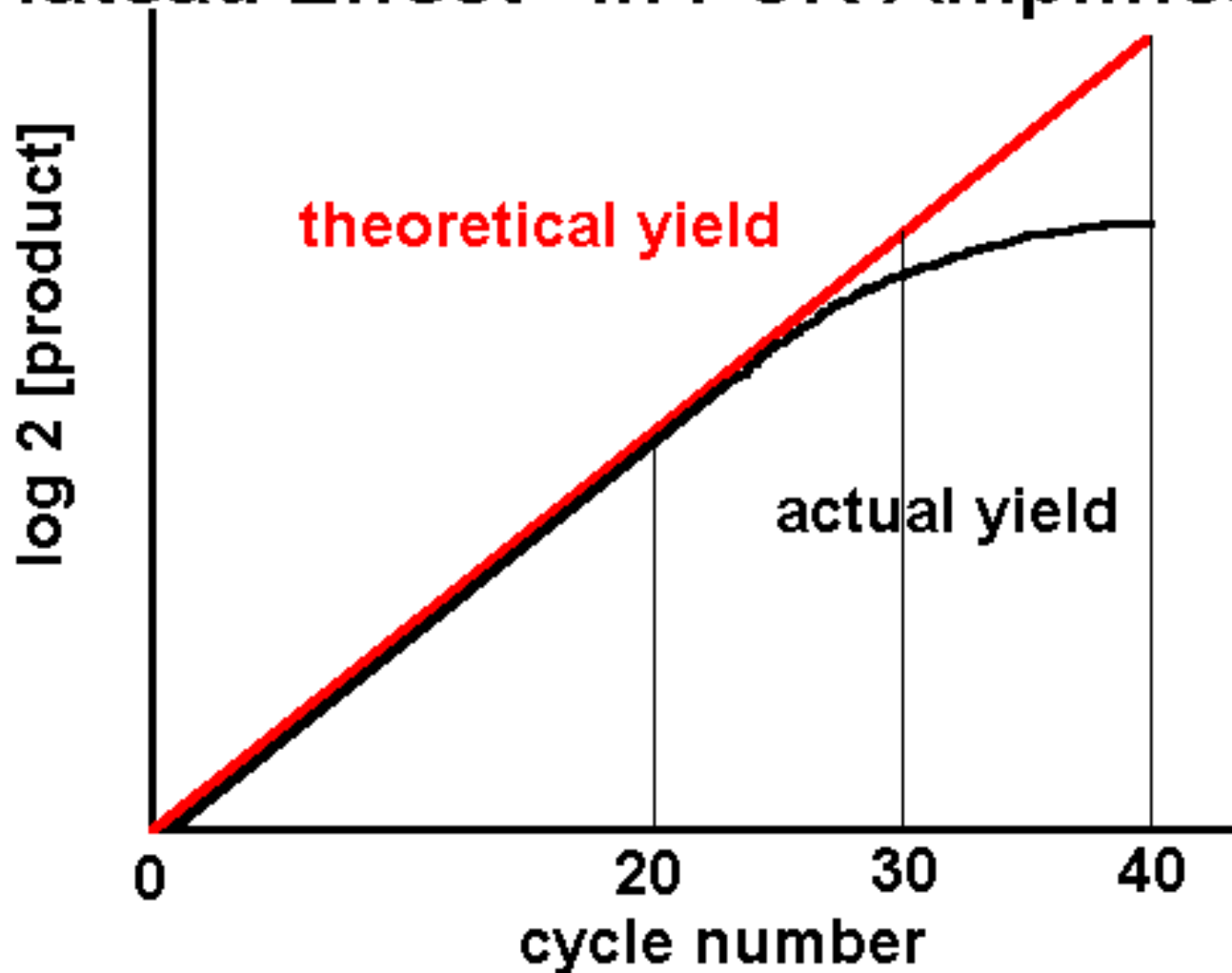


PCR Procedure

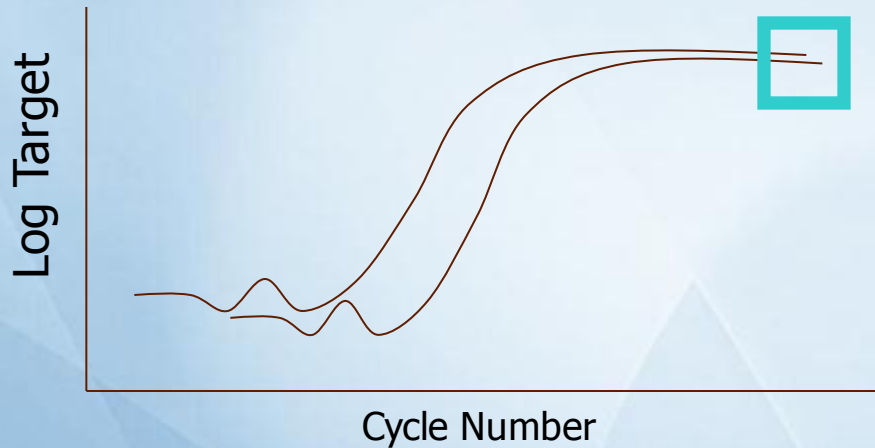


Cycle Number

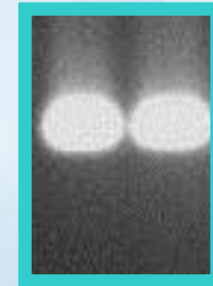
"Plateau Effect" in PCR Amplification



Conventional PCR problem



Agarose gel
electrophoresis
following PCR



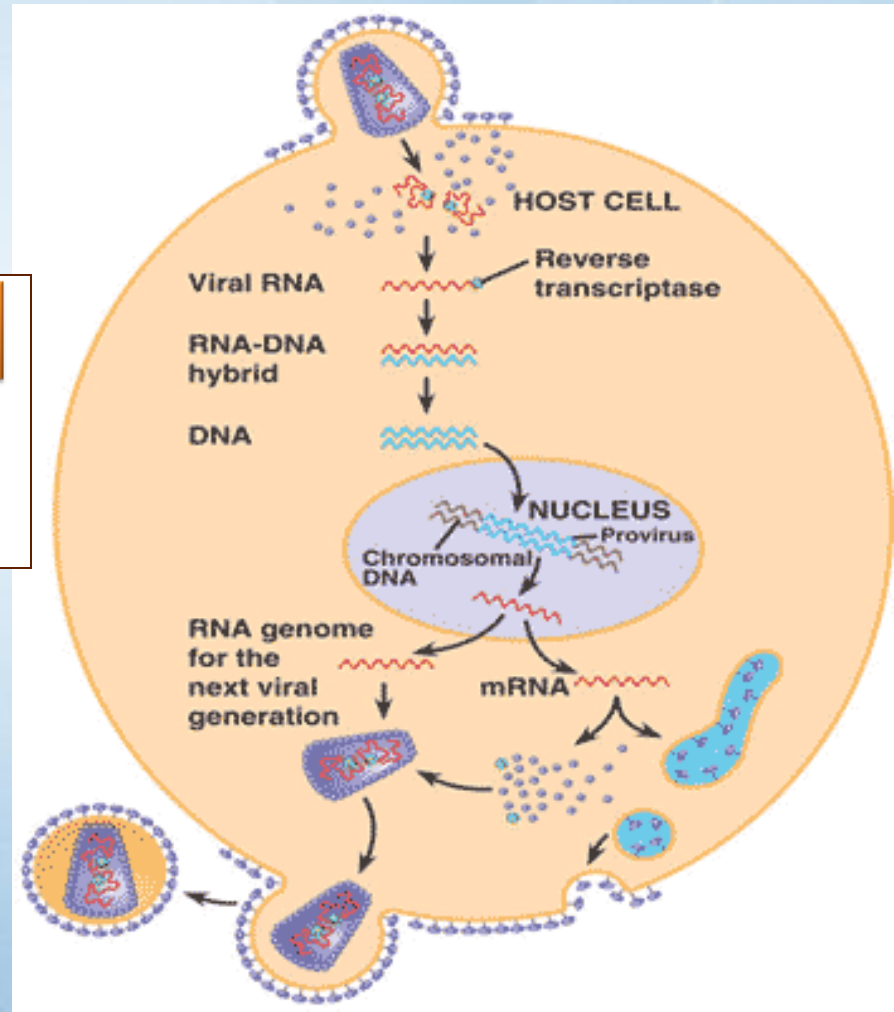
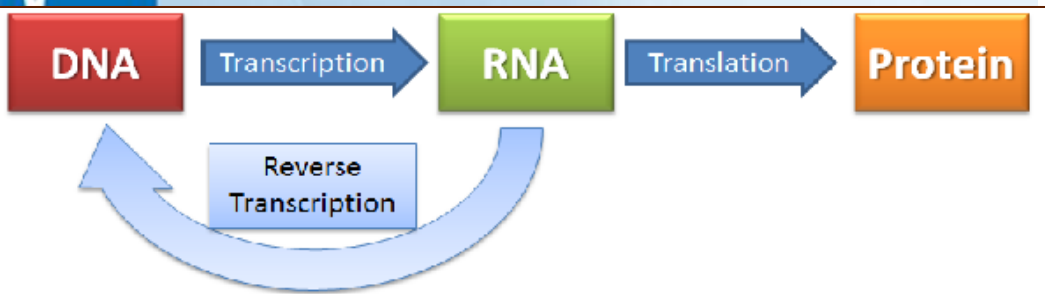
Why qRT-PCR?

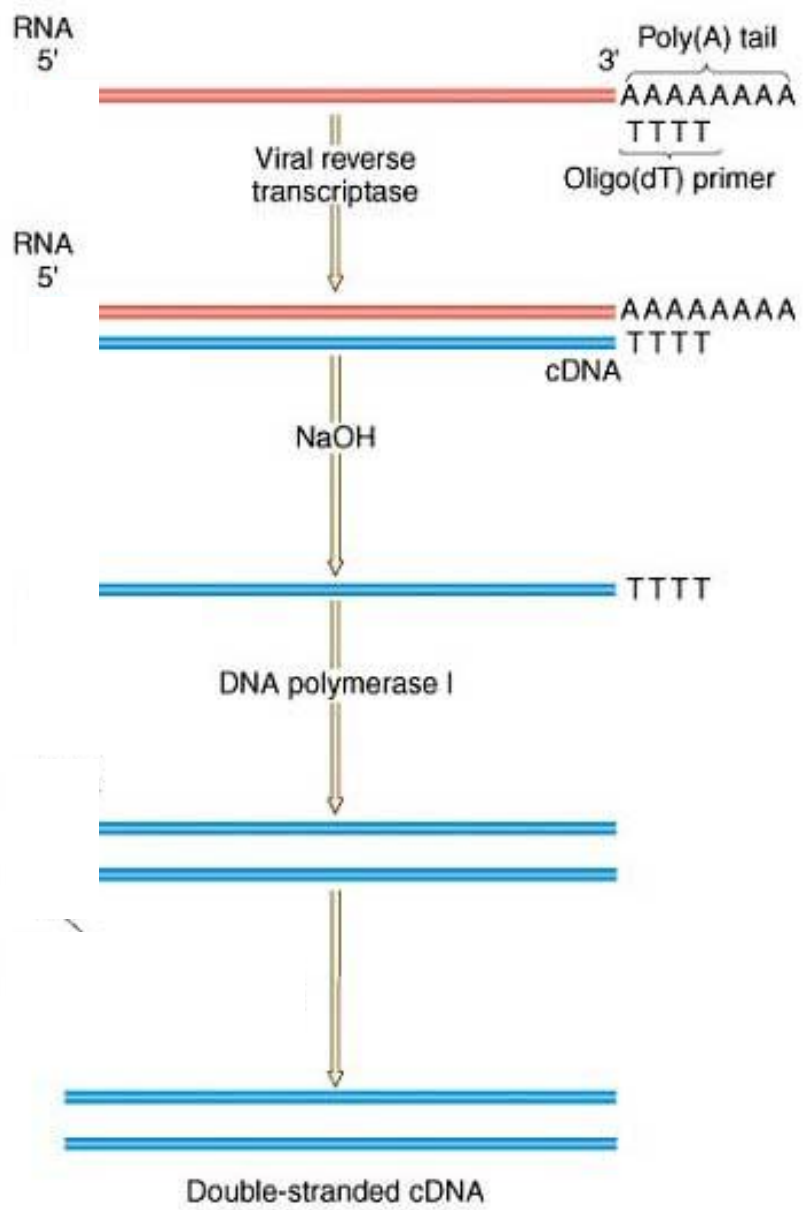


- **Gene expression analysis**
 - Cancer research
 - Drug research
- **Disease diagnosis and management**
 - Viral quantification
- **Food testing**
 - Percent *GMO* food
- **Animal and plant breeding**
 - Gene copy number

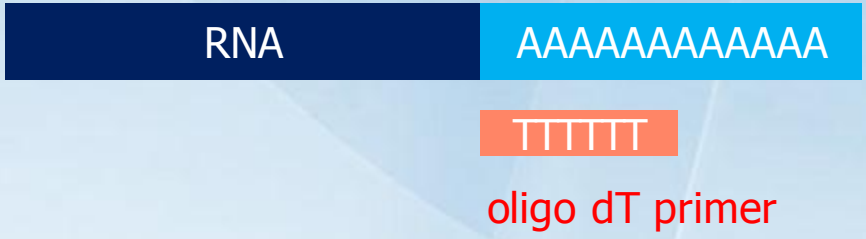
qRT-PCR is used to qualitatively detect gene expression

Reverse transcription





All messenger RNAs (mRNA) have a poly A tail



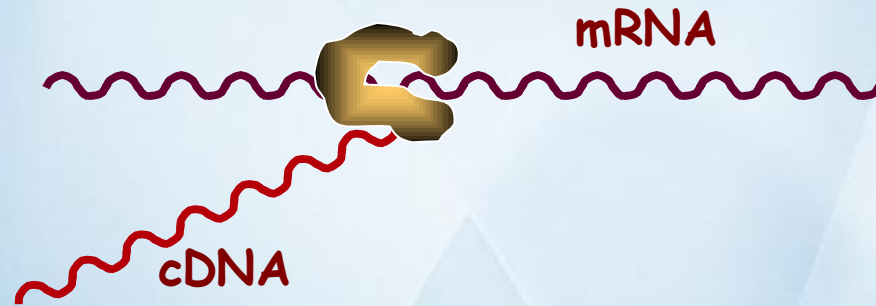
The RT reaction can be primed by a:

- ✓ target specific primer (i.e. primer targeting VHSV nucleocapsid (N) gene)
- ✓ oligo dT primer (a primer consisting of a run of T's that targets the mRNA poly A tail)
- ✓ random primers (a mix of 6 base primers consisting of random nucleotides)

RT-PCR

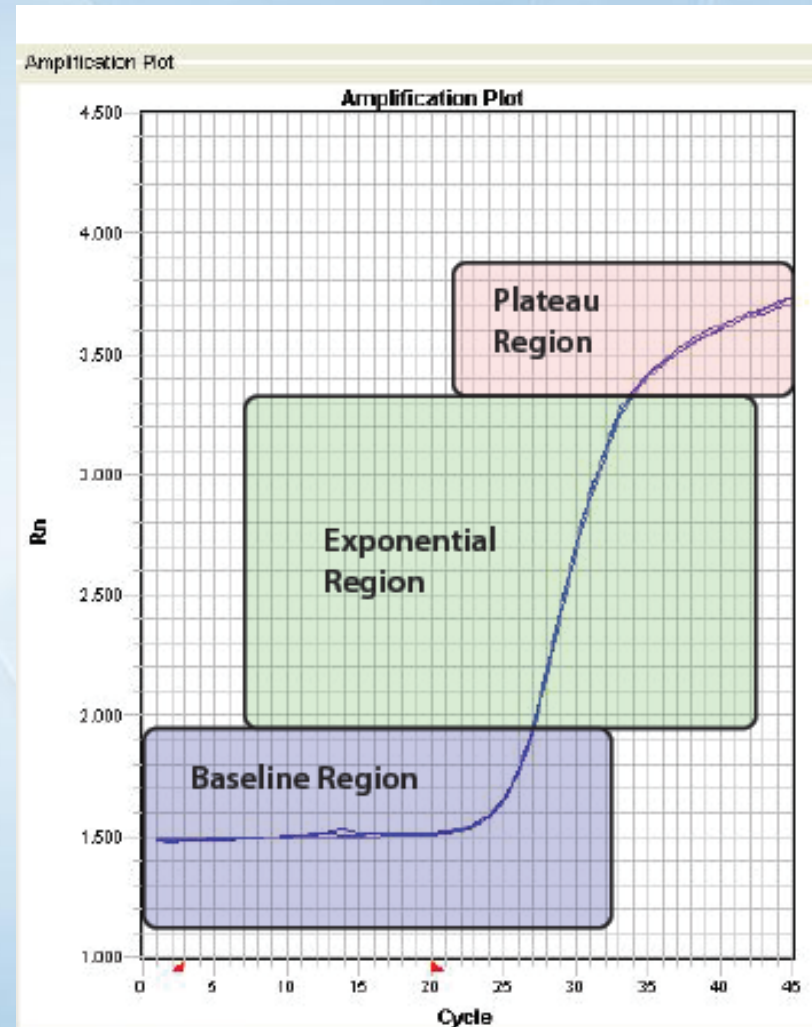
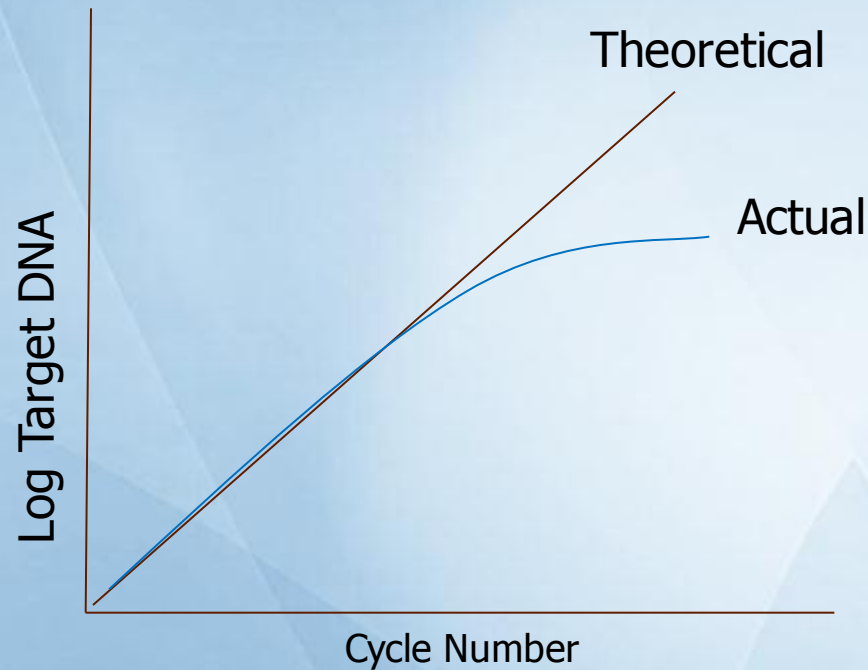


RT: Reverse transcriptase



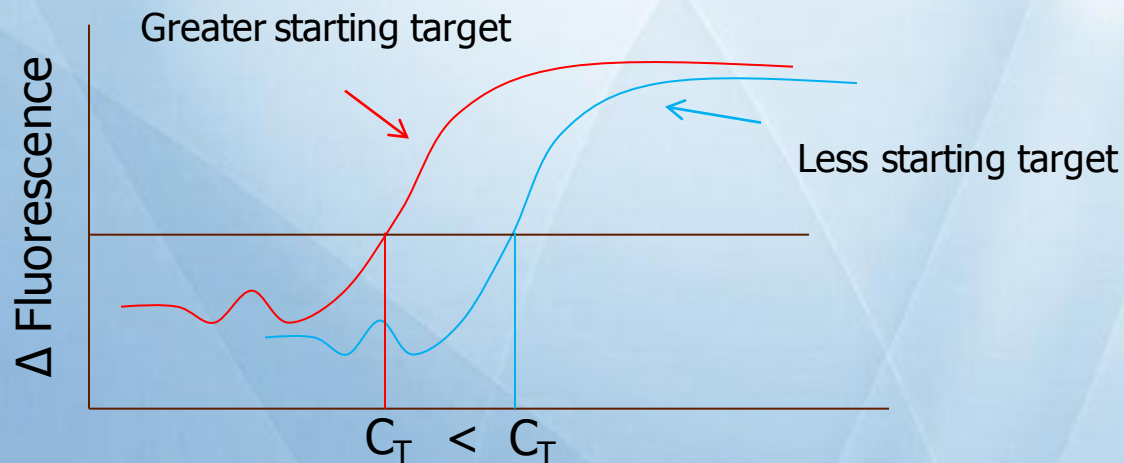
Templat for RT- PCR

How does PCR work?



Real-Time PCR

- ▶ Real-Time PCR a specialized technique that allows a PCR reaction to be visualized “in real time” as the reaction progresses.
- ▶ Quantitative PCR relies on the principal that the quantity of target at the start of the reaction is proportional to amount of product produced during the exponential phase

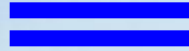


How to measure the PCR product?

Initial DNA strand



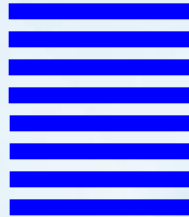
First PCR cycle



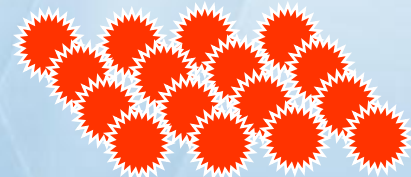
Second PCR cycle

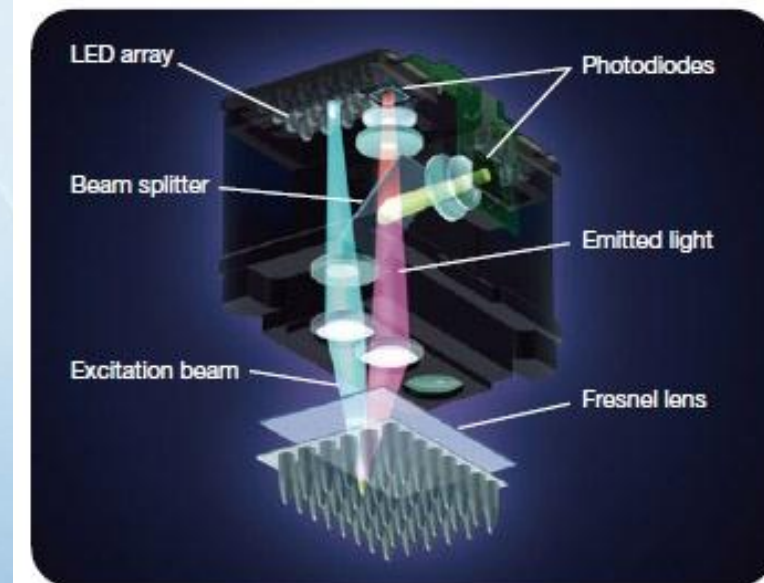
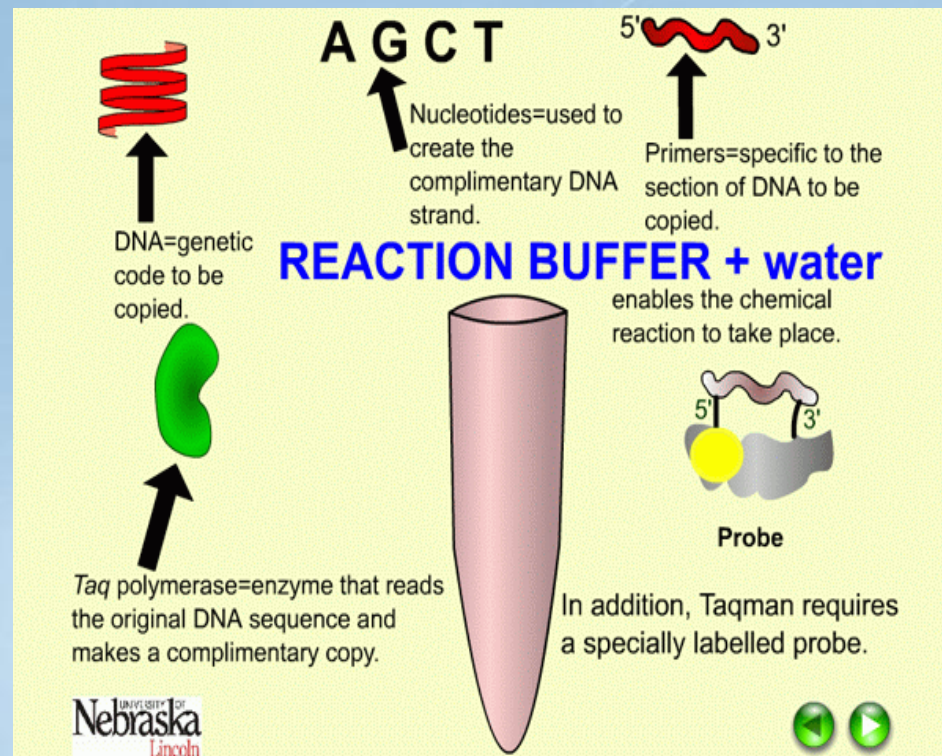


Third PCR cycle

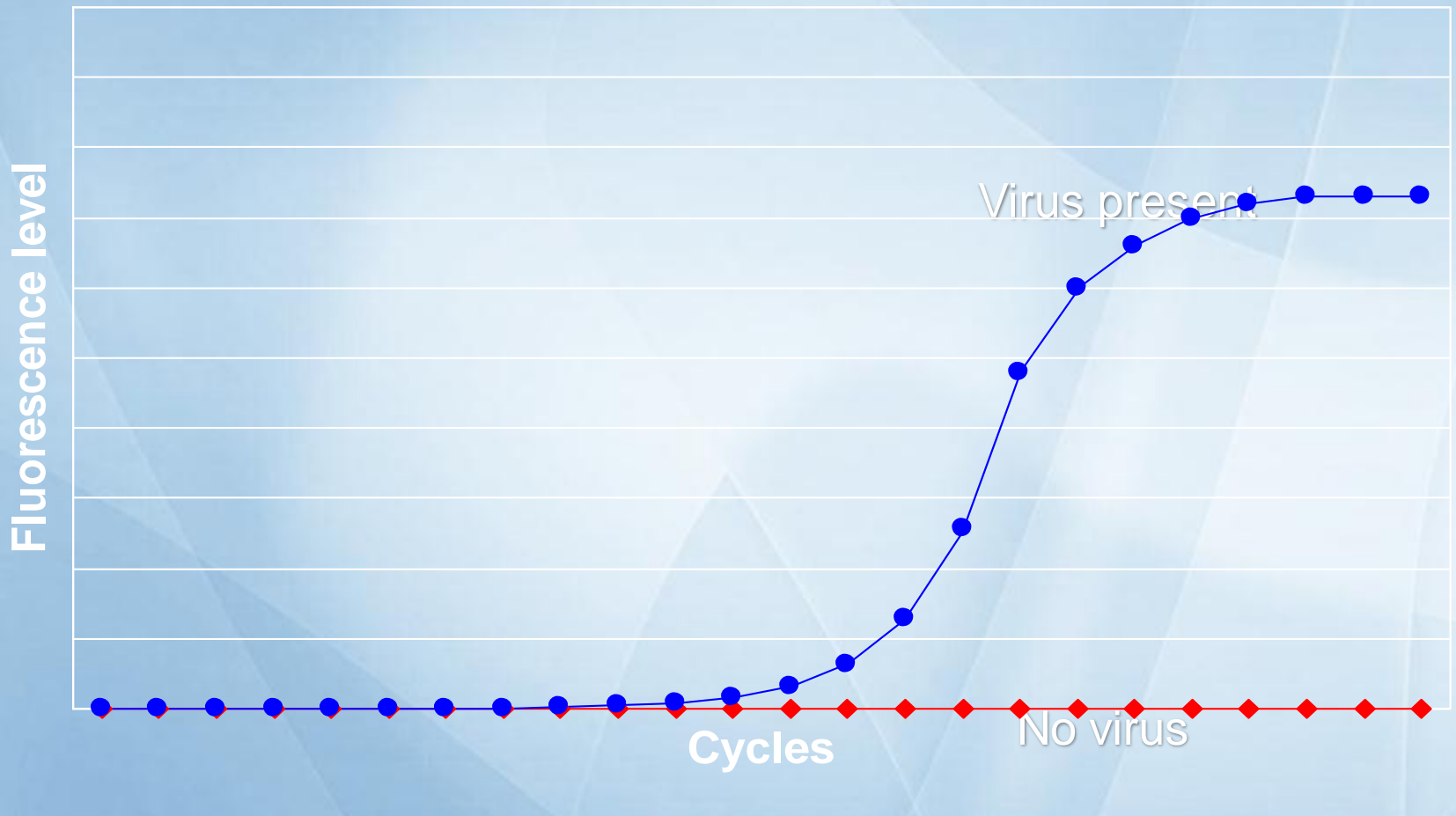


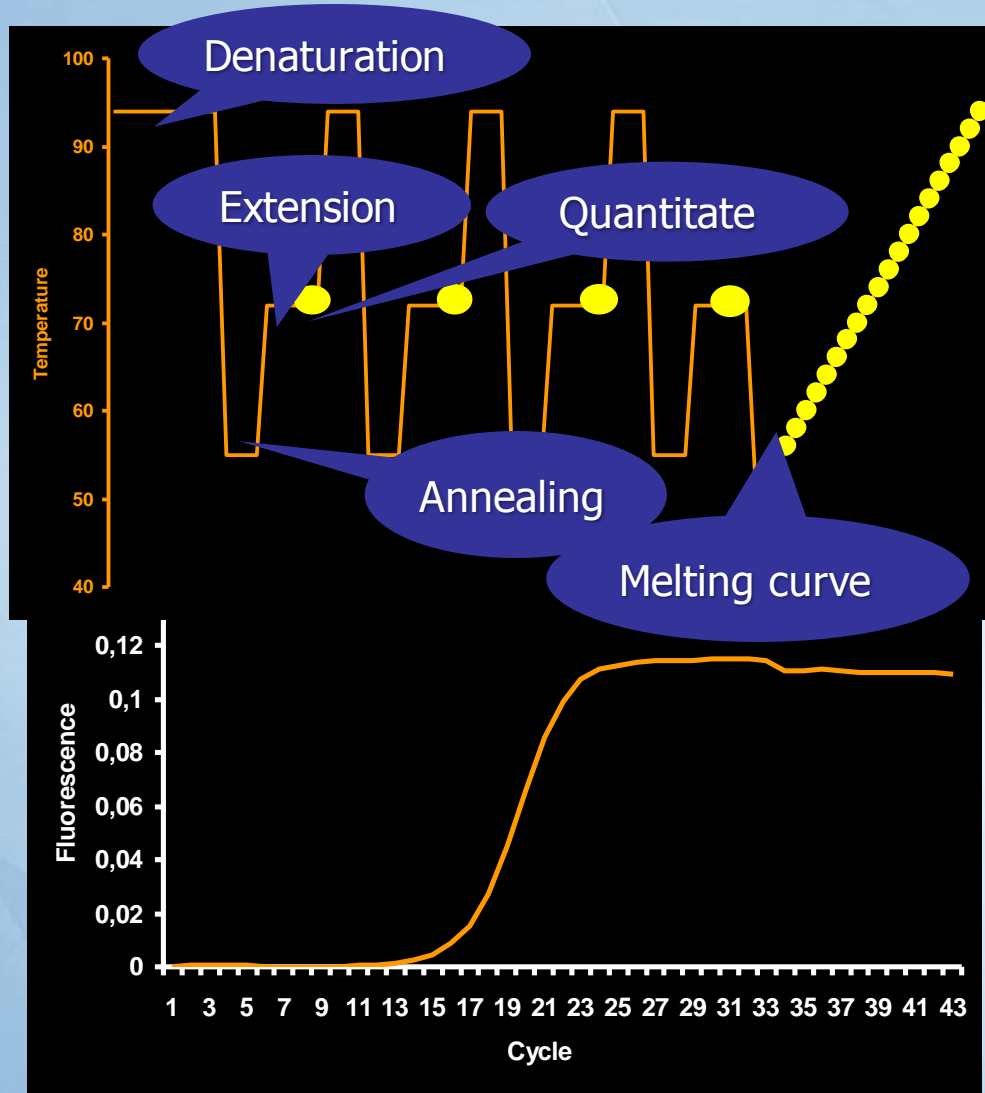
Fourth PCR cycle





Any increase in fluorescence level can be plotted onto a graph and easily interpreted

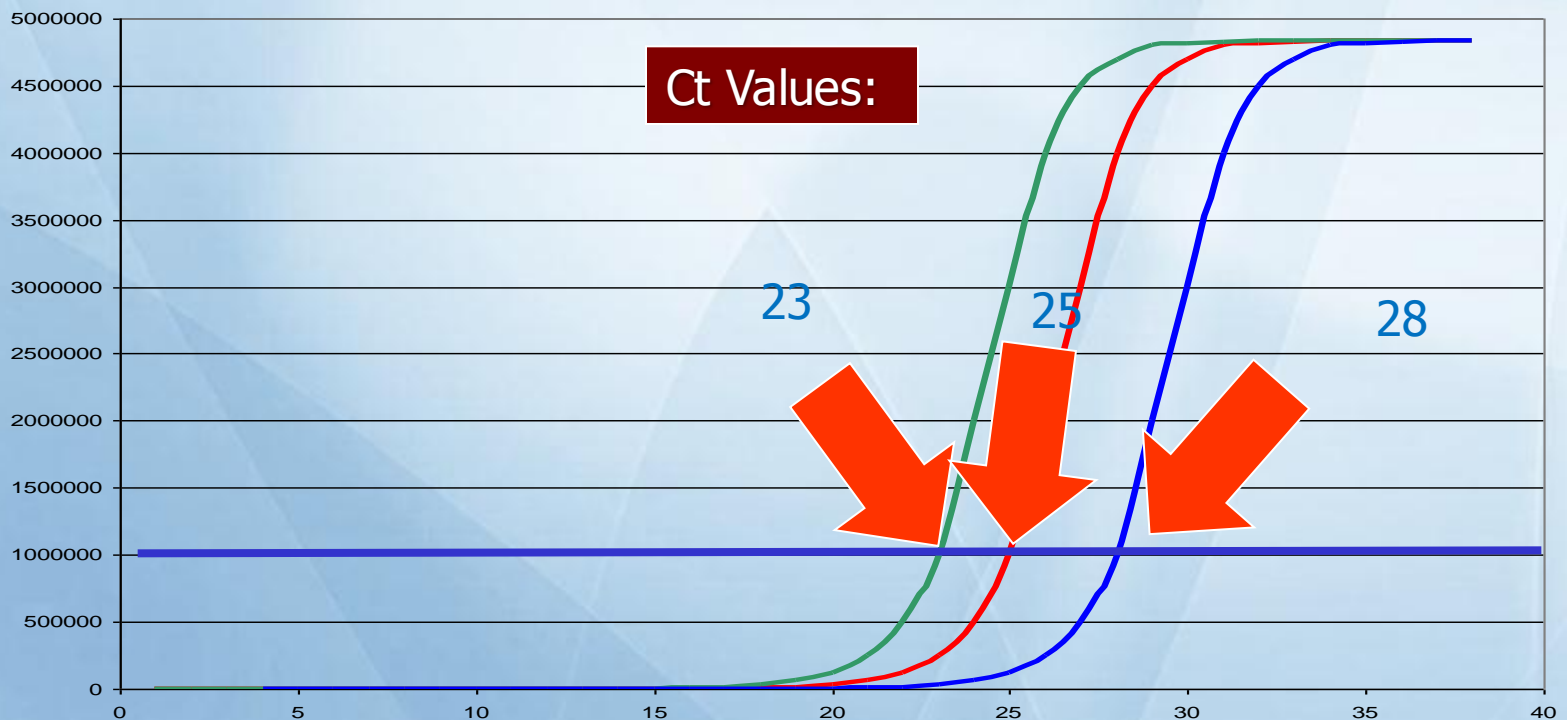




We describe the position of the lines with a value that represents the cycle number where the trace crosses a threshold.

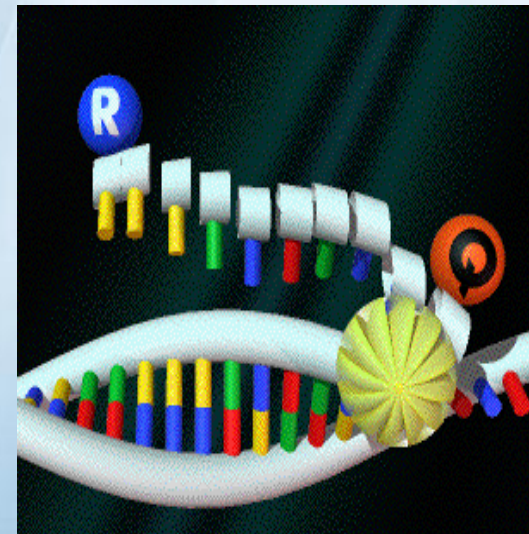
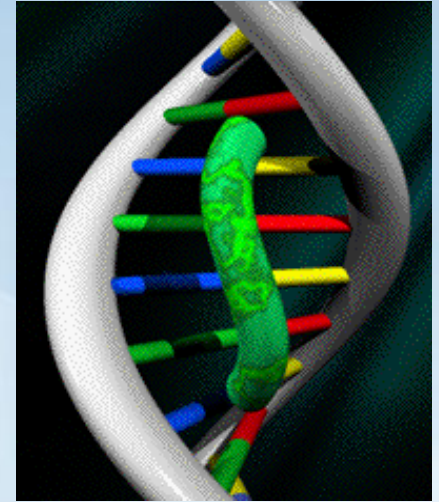
This is called the cycle threshold “Ct Value”.

Ct values are directly related to the starting quantity of DNA, by way of the formula:



Quantitative PCR – in depth

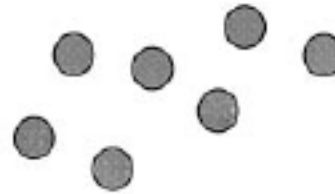
- ▶ Major assay types
 - ▶ **Hydrolysis probes**
 - ▶ Basis of TaqMan® chemistry
 - ▶ Uses two primers and an internal hydrolysis probe
 - ▶ Most commonly used for fish health diagnostics
 - ▶ **SYBR® green dye**
 - ▶ Increased fluorescence when bound to dsDNA
 - ▶ Slightly lower specificity
 - ▶ Costs less
 - ▶ May not be as sensitive as the 5' nuclease assays



SYBRGreen

A

SYBR
Green I



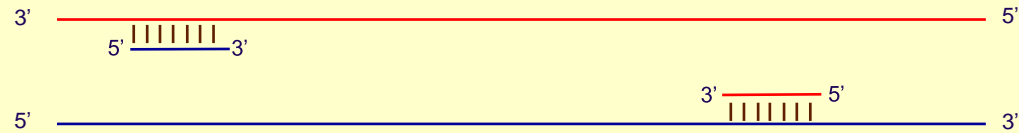
Target

B

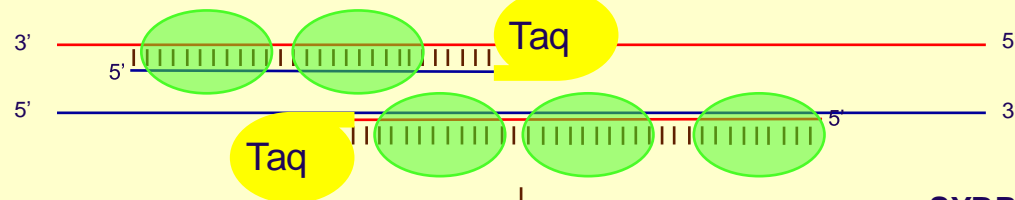


Fluorescent dye

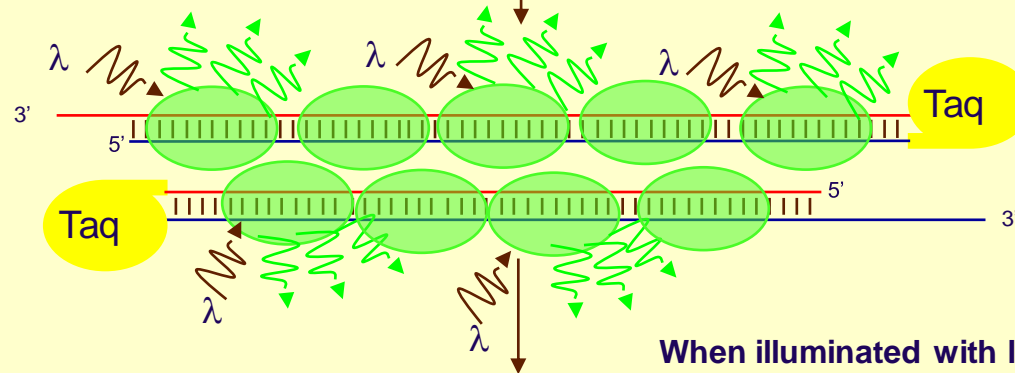




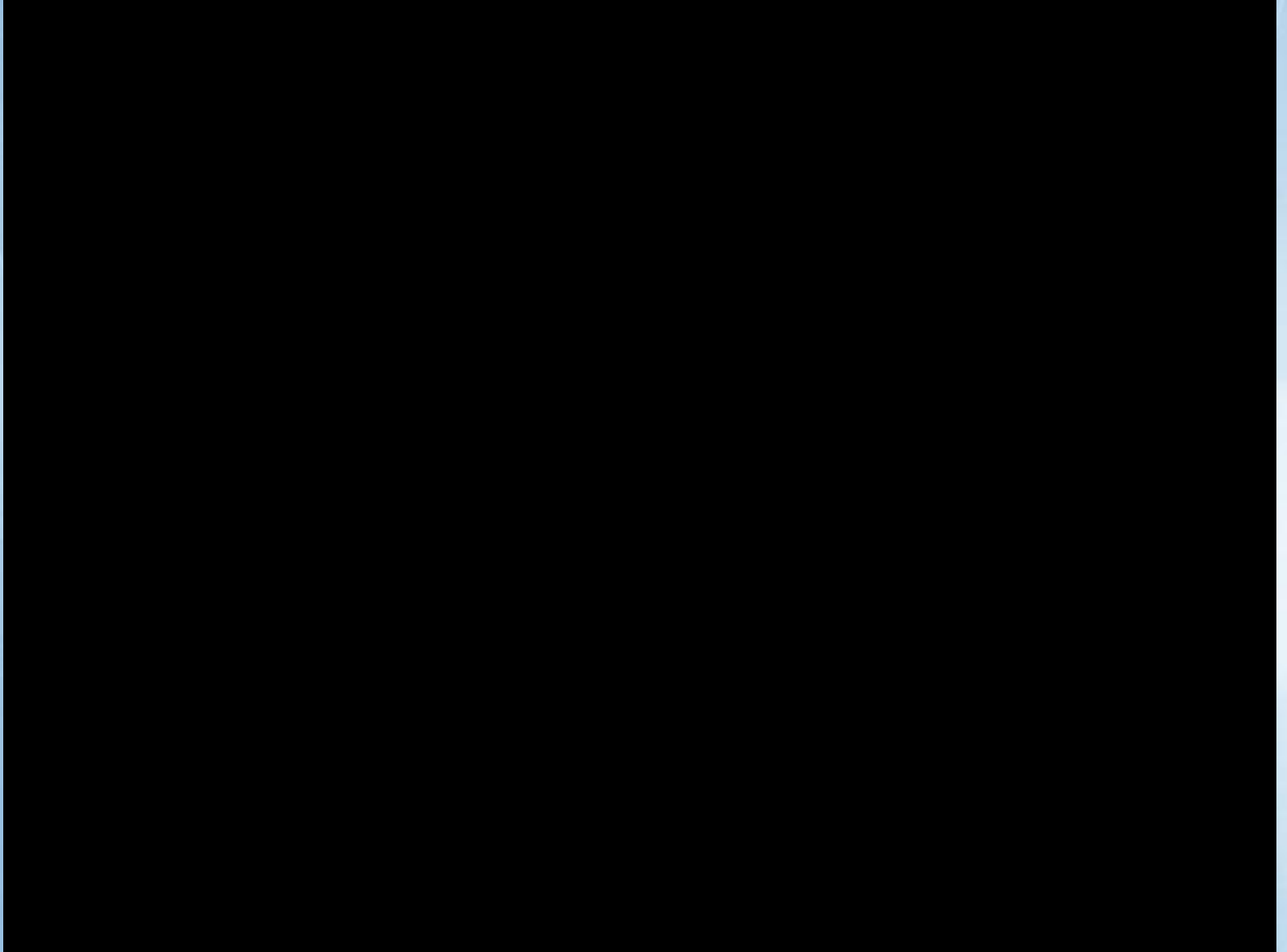
PCR makes more double-stranded DNA



SYBR Green dye binds to dsDNA



When illuminated with light at 490nm, the SYBR+DNA complex fluoresces at 520nm.

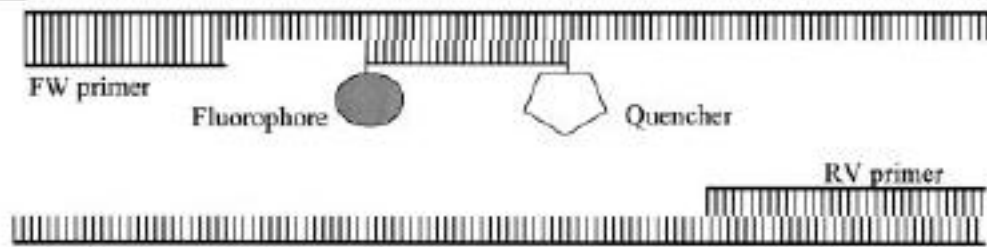


BackMan vs TaqMan

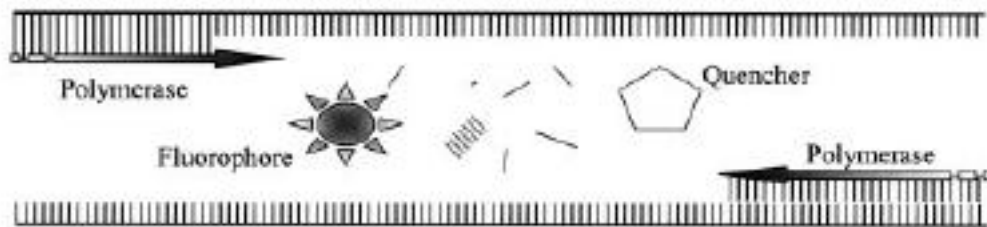


TaqMan

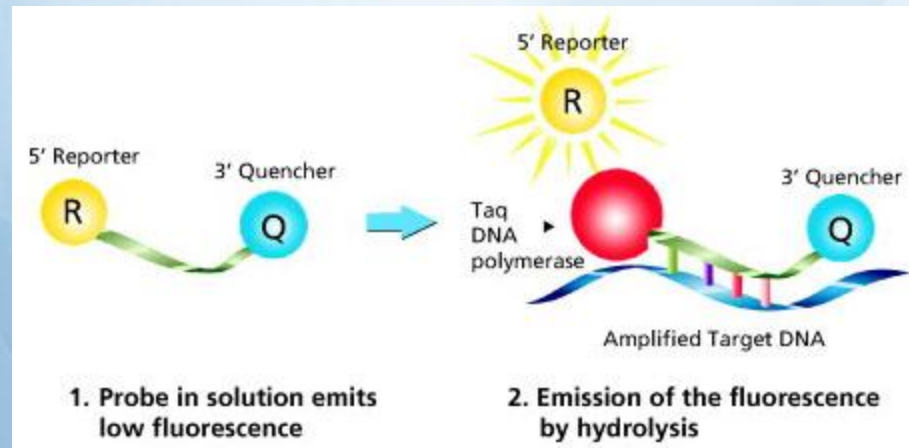
A

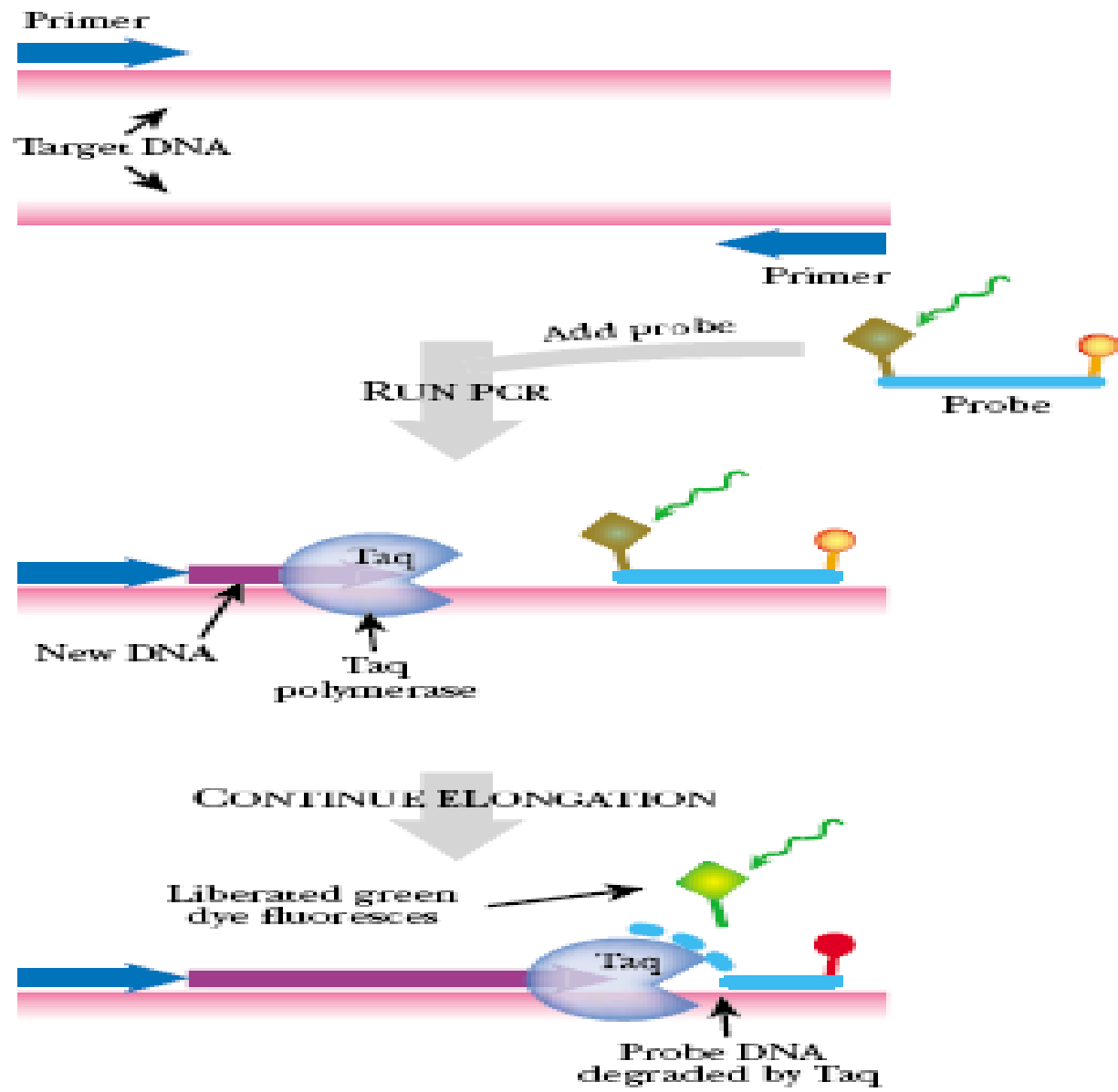


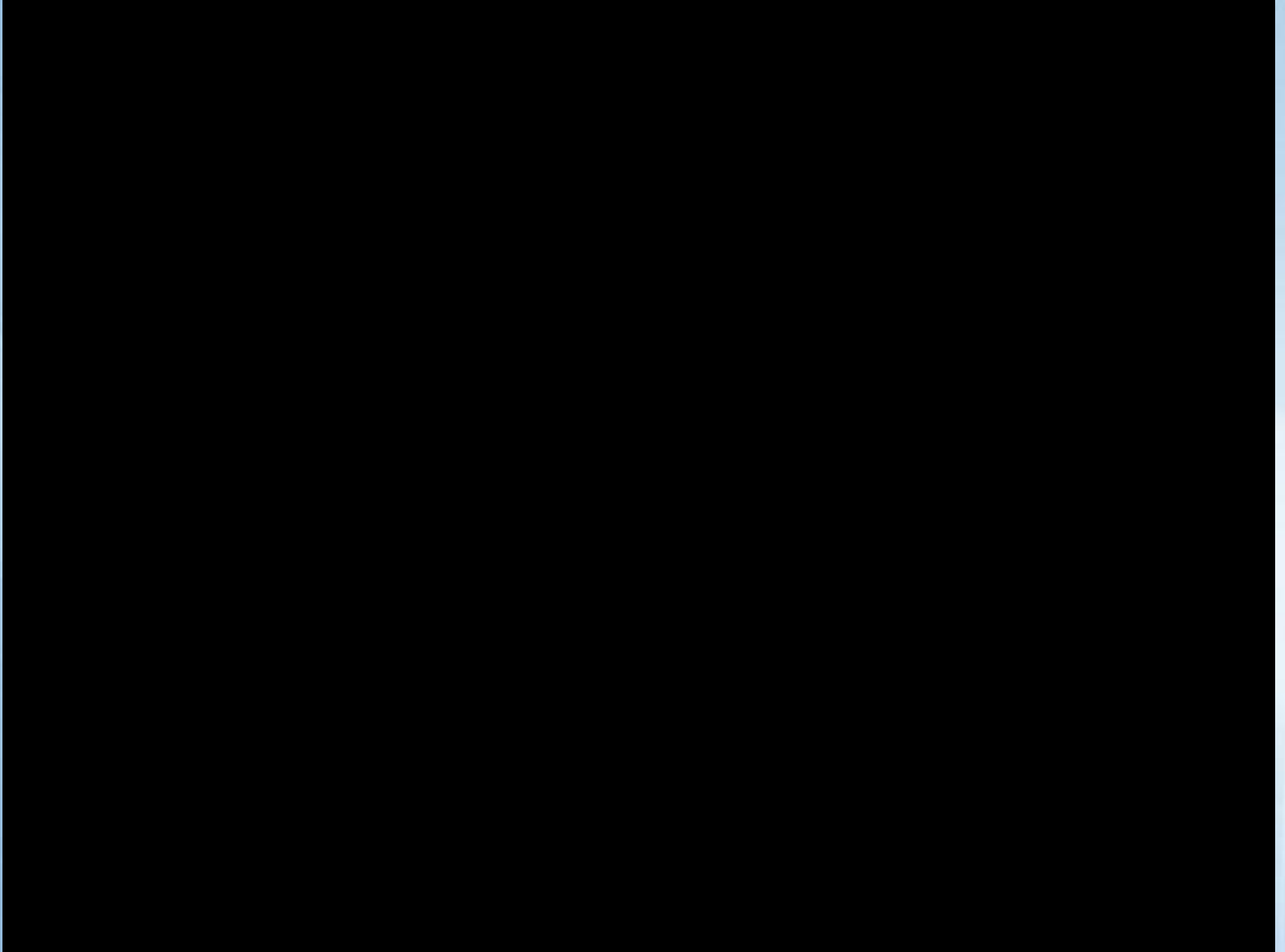
B



Probe hydrolysis







Taqman vs. SYBR Green

TaqMan Probe

Advantages:

- Increased specificity
- Use when the most accurate quantitation of PCR product accumulation is desired.
- Option of detecting multiple genes in the same well (multiplexing).

Disadvantages:

- Relative high cost of labeled probe.

● SYBR Green

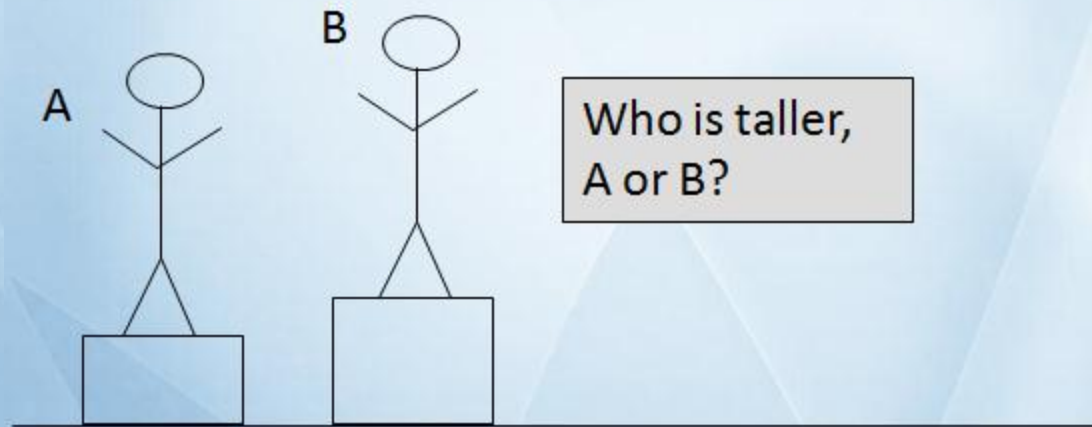
● Advantages:

- Relative low cost of primers.
- No fluorescent-labeled probes required.

● Disadvantages:

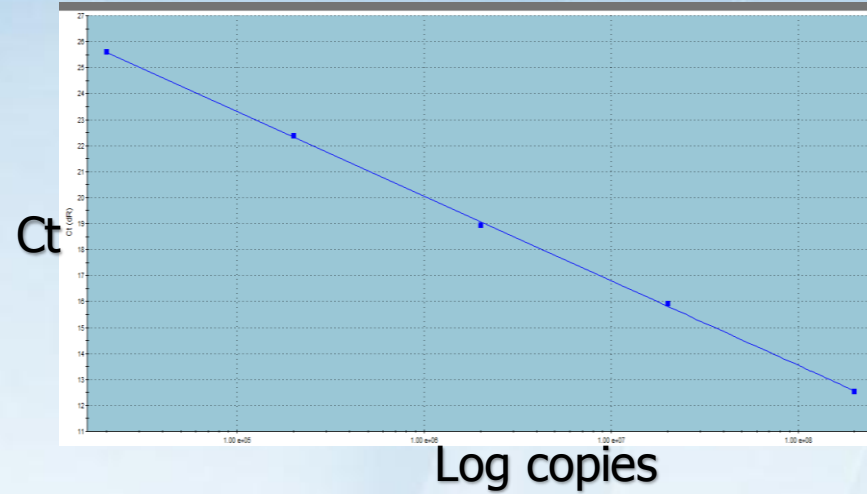
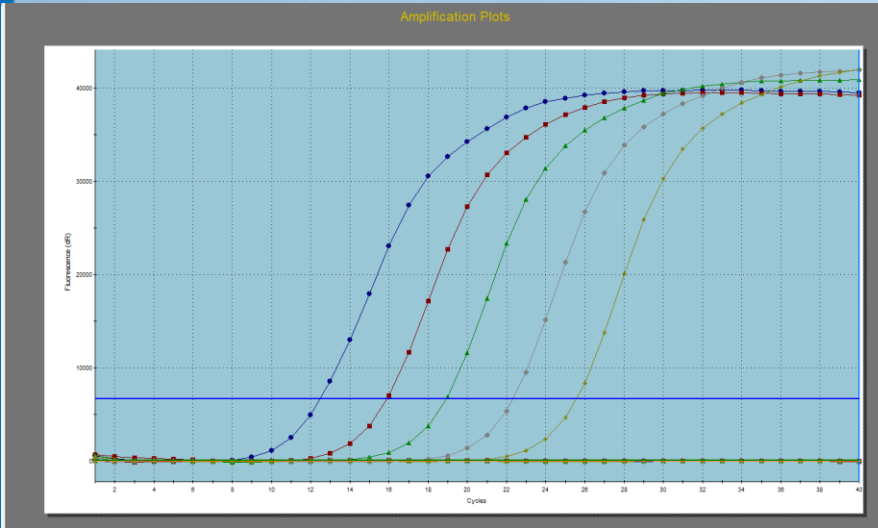
- Less specific - only primers determine specificity.
- Specific and non-specific double-stranded PCR products generate the same fluorescence signal upon binding SYBR Green I dye.
- Not possible to multiplex multiple gene targets.

Normalization



Normalization: corrects for variation in template quantity and/or template quality

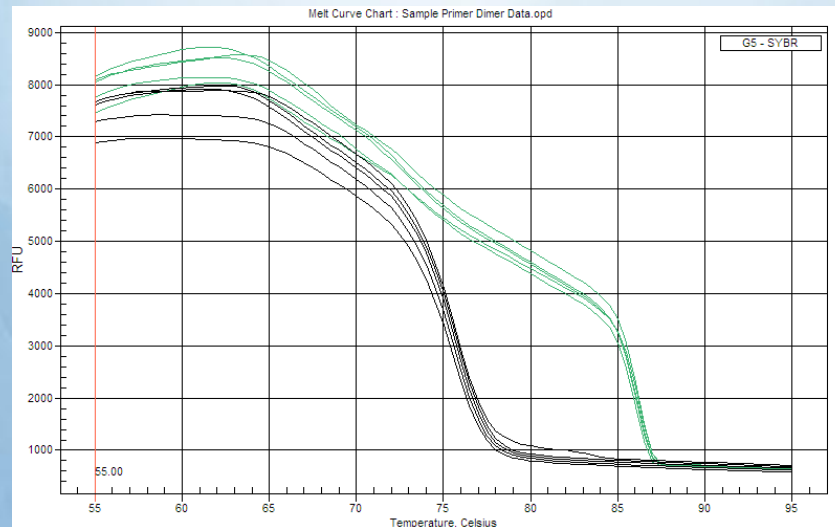
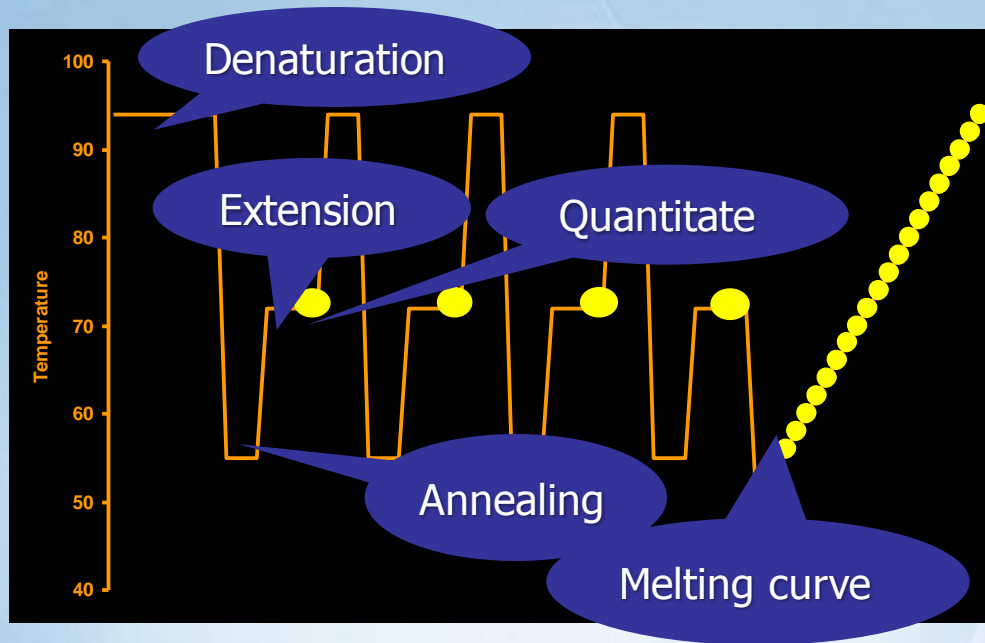
Standard Curves



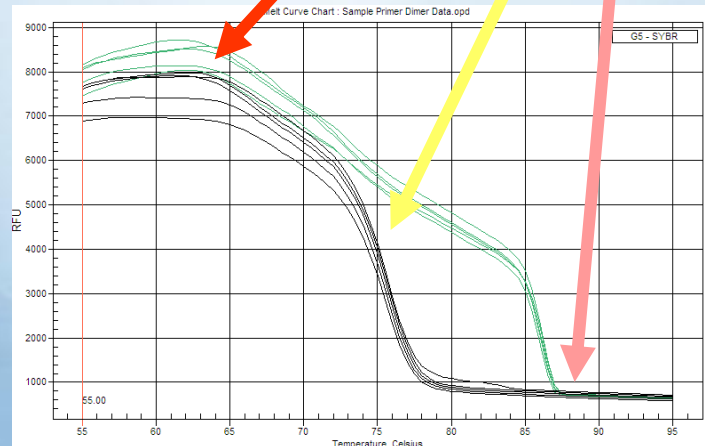
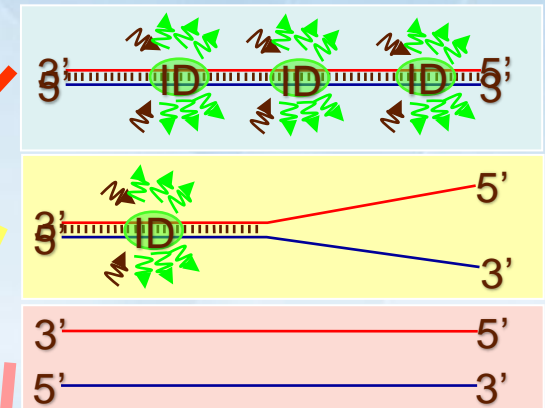
- Use at least 5 concentrations for a standard curve
- Use serial dilutions that are one order of magnitude apart
_ 1:10, 1:100, 1:1000,...

Housekeeping Genes

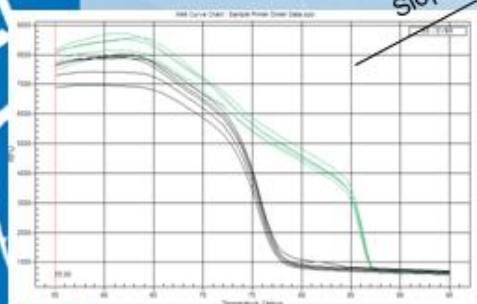
Gene	Genomic structure / pseudogenes	Regulation e.g.
β -actin	multigene family; > 20 genes; 1 active locus 20 pseudogenes	\uparrow : hormones of thyroid gland \uparrow : stomach tumor
γ -actin	multigene family; pseudogenes	
GAPDH	multigene family; 10-30 genes; > 200 in mouse mostly pseudogenes	\uparrow : lung, pancreatic, colon cancer \uparrow : insulin, EGF
5.8S, 18S, 28S RNA	pseudogenes	
β 2-microglobulin	no pseudogenes	\uparrow : Non-Hodgkin lymphoma abnormal expression in tumors
G6PDH	no pseudogenes	\uparrow : kidney, stomach tumor \uparrow : hormones, oxidant stress, growth factors
PBGD	no pseudogenes	
aldolase	pseudogenes	
HPRT	pseudogenes	
U3, U8, ...	Pseudogenes	
ornithin decarboxylase		\uparrow : tumors
...		



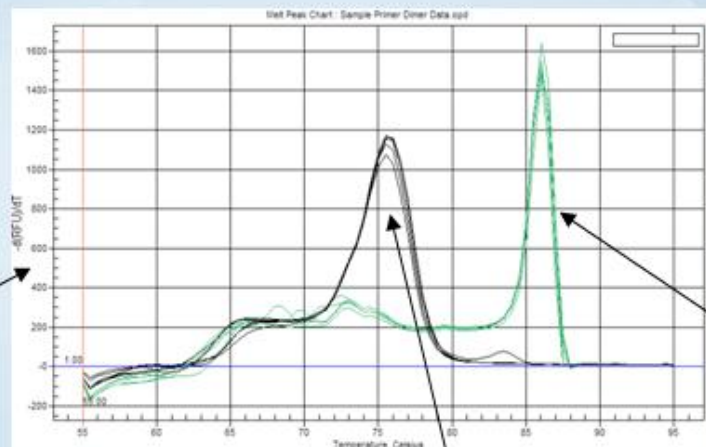
- **Melt curves can tell us what products are in a reaction.**
- PCR products that are shorter will melt at lower temperatures.
- Different PCR products will therefore have different shaped curves.



- For convenience, we typically view the derivative (*slope*) of the actual melt curve data.
- The resulting graph looks like a chromatogram, with peaks that represent different PCR products.



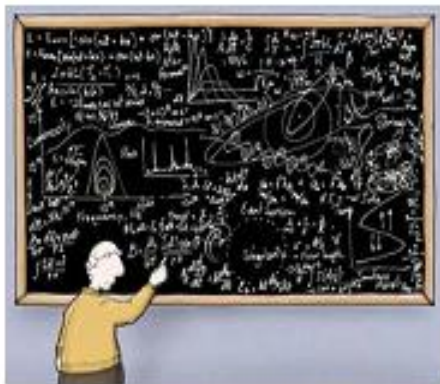
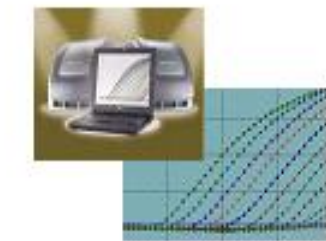
Slope vs. Temp



These PCR products melt at a relatively low temperature, are likely primer-dimers.

These PCR products melt at a higher temperature, and are likely to be the main PCR product.

Steps towards successful Real-time QPCR experiments



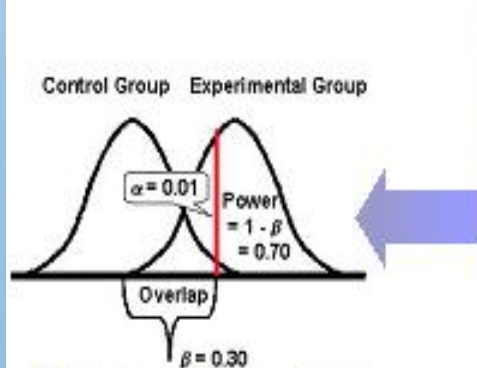
1. Experimental Design



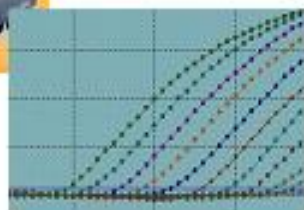
2. Sample preparation and purification



3. RNA/DNA quantification and quality



6. Post-run Analysis



5. Real time QPCR



cDNA



Total RNA

4. Reverse Transcription

