

# M1D2: Diagnostic primer design

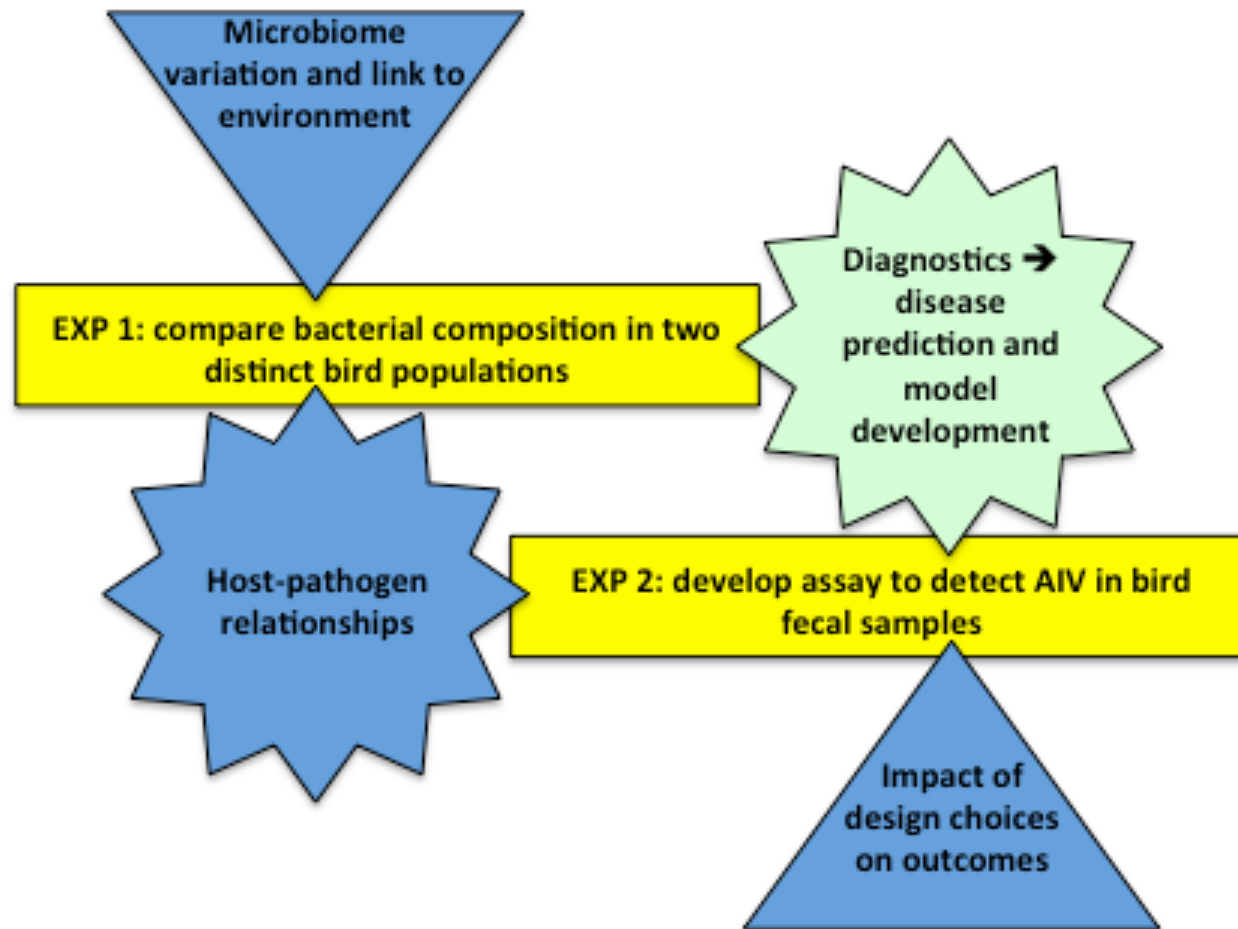
2/11/15

# Lab business

- Lab treat next time (M1D3)
  - Content will be from lecture (M1D1) and pre-labs (M1D1 and M1D2)
  - All quiz dates posted on Schedule page of wiki
- Homework due M1D2 (today)
  - Leave wiki page printout on front bench
- Office hours begin tomorrow
  - Noreen: M 2-4p, R 2-4p, and by appointment
  - Leslie: M 1-2p, R 1-2p, and by appointment

14-429b

# Module 1 conceptual overview



# AIV screening project

- Primary research question?

Can we improve the detection of AIV in gulls?

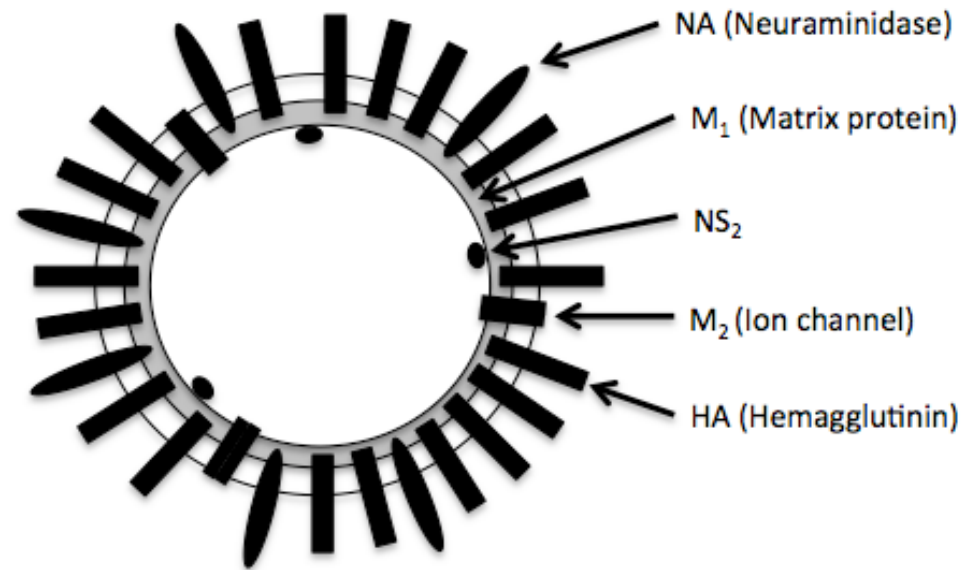
- Broader impact?

Improving the screen better predict disease model development



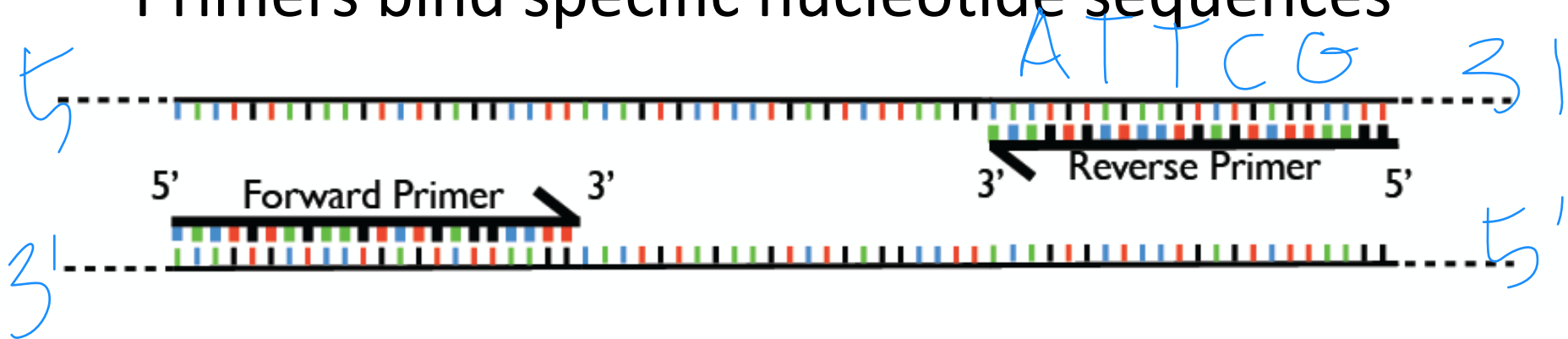
# How will you develop your AIV screen?

- NA and HA are variable and used to define AIV subtypes
- M1 (MA) is highly conserved
- Which is a better target for screening bird samples for AIV?



# How will we target M1?

- Primers bind specific nucleotide sequences



- Forward primer binds the  $\alpha$ -sense strand
- Reverse primer binds sense strand

make sure to use  
reverse complement

# Primer sensitivity

- In diagnostic assays, sensitivity is important for determining presence/absence
  - Some false positives okay, but want as few as possible
- Specificity is important in identifying strain or subtype information
- Is your goal to increase sensitivity or specificity?

# Primer design considerations

## 1. Length

>17 bp

10 bp  $4^{10} = 10^4$   $3 \cdot 10^9$  bp

## 2. Melting temperature

55-60°C  $58 \text{ C} \pm 3$

$T_m = 50\%$  bound/unbound

## 3. GC content

$\uparrow$  GC  $\uparrow$   $T_m$

40-50%

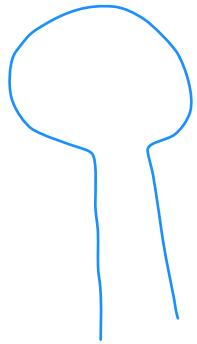
GC clamp 3'

# Primer design considerations

## 5. Long repeats

non specific binding  
T stretches → poly A tails

## 6. Secondary structure



energetically  
favorable

# Primer design considerations

## 6. Primer dimers

homodimers  
heterodimers

**OligoAnalyzer 3.1**

Instructions | Definitions | Feedback

Sequence: 5'-CCTCTCCGGAAACCAACCCCTG # Bases: 21

Target Type: DNE

Oligo Conc: 0.25  $\mu$ M

Na<sup>+</sup> Conc: 50 mM

Mg<sup>++</sup> Conc: 0 mM

Analyze

Hairpin

Self-Dimer

Hetero-Dimer

NCBI Blast

TM Mismatch

Clear Sequence

Results | 5' mods | Internal Mods | 3'

**HOMO-DIMER ANALYSIS ?**

Dimer Sequence

5'- CCTCTCCGGAAACCAACCCCTG -3'

Maximum Delta G -43.97 kcal/mole

Delta G -12.9 kcal/mole

Base Pairs 6

IDT Website: "You can also compare the value of the maximum delta g (the delta g for a perfect duplex) to that of each individual self dimer. If the values are within 10% of each other, you should redesign. Heterodimer analysis works the same way."

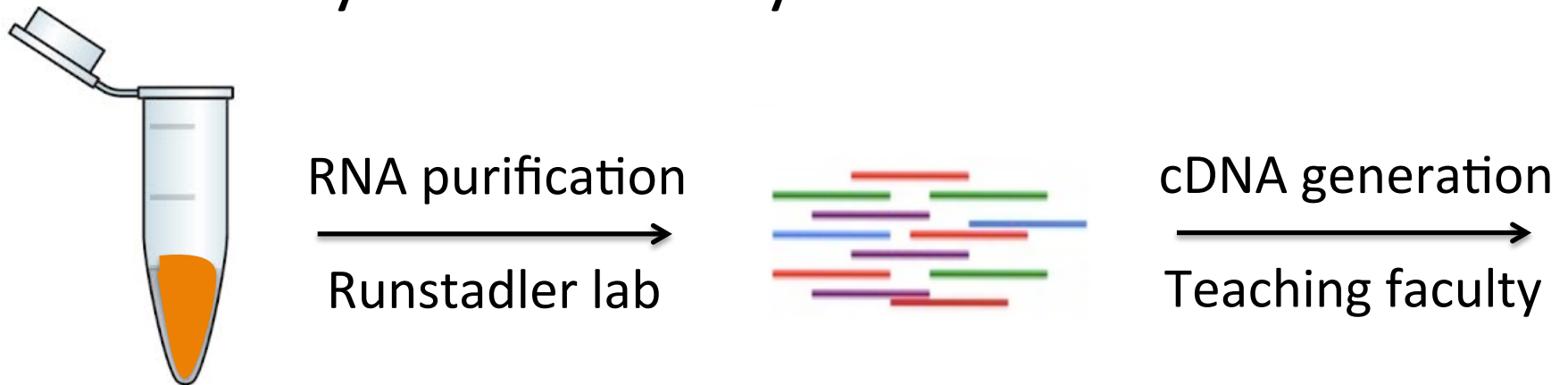
# You have your primers, now what?

- You will screen fecal samples collected from 'scallop pile' in Nantucket



# Experimental overview

- While you were away...

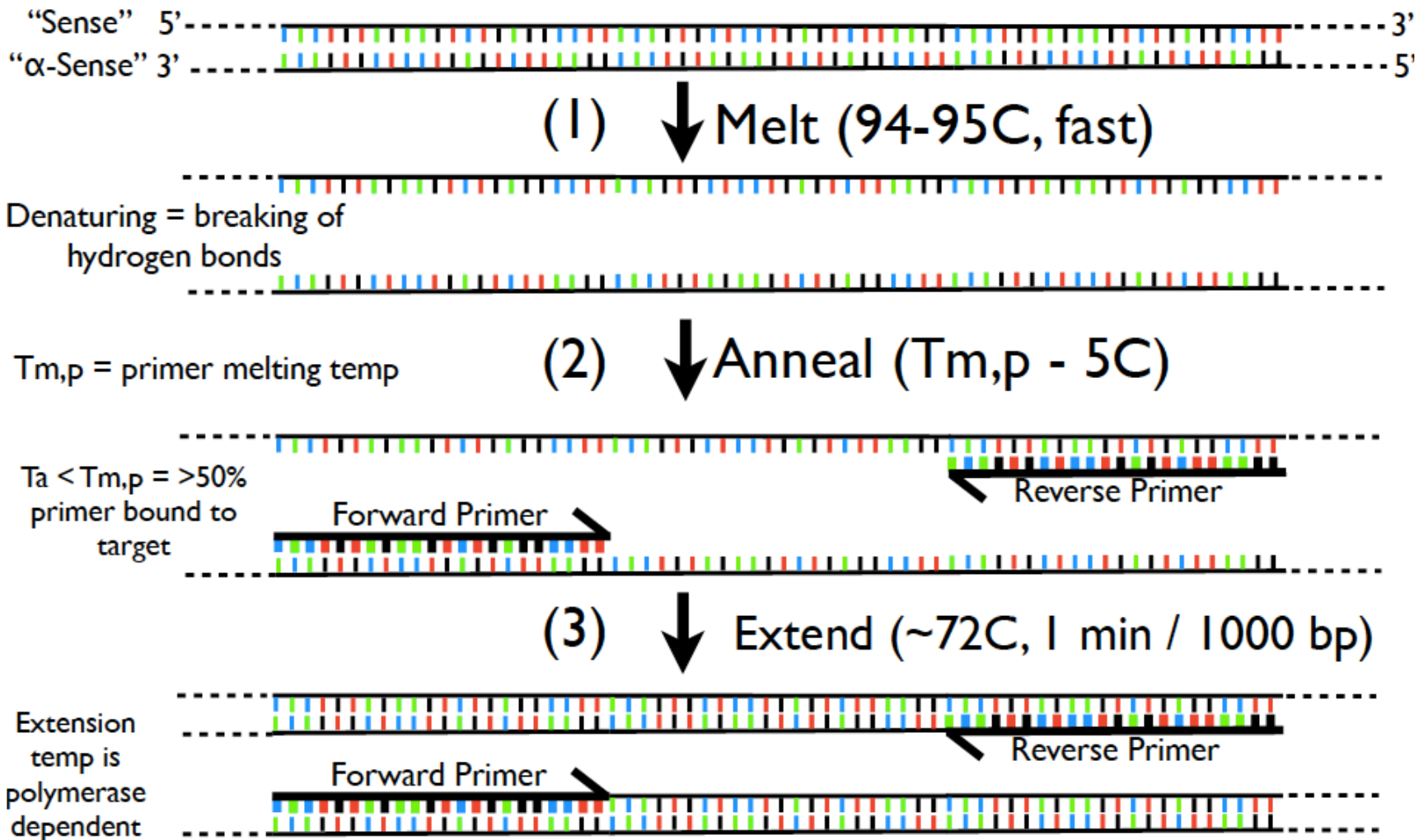


- Your part...

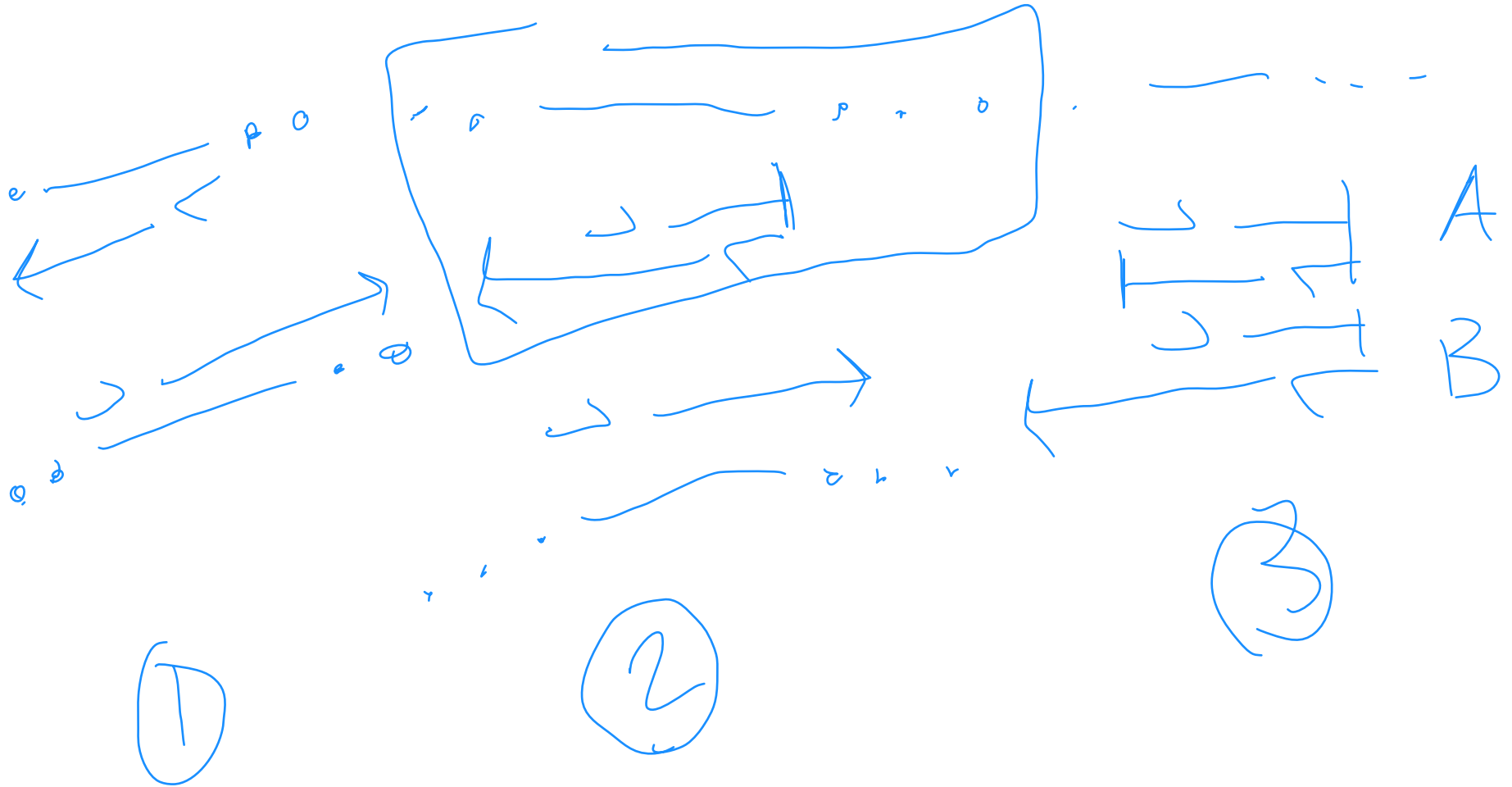
Screen cDNA pool using  
your primers.



# Polymerase chain reaction (PCR)



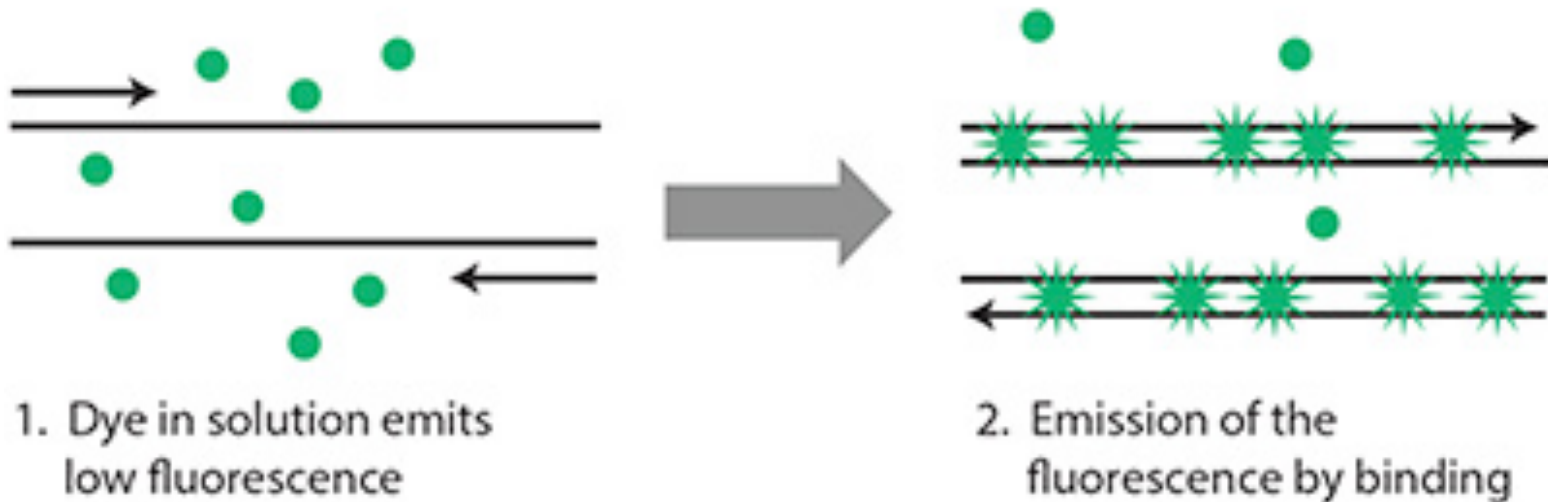
# Draw the first three cycles



In which cycle do you first see your amplicon?

# Quantitative PCR

- SYBR green dye binds double stranded DNA
- Amount of fluorescence is proportional to the amount of PCR product



# PCR versus qPCR

- Utility of PCR?

cloning

- Utility of qPCR?

# Today

- Explore current AIV primers
- Develop your AIV screening assay
  - Specifically, you will design your primers
  - Enter sequences on M1D2 Talk page
- Homework
  - Prepare for paper discussion
  - Discuss your primer design choices
  - Post final primer design (by M1D4)
  - Choose journal article presentation day

(by M1D4)