

- Announcements
- Quiz
- Pre-lab Lecture
  - ❖ Today in Lab: M2D3
  - ❖ Shape effects in DNA gels
  - ❖ Diagnostic digest preview
  - ❖ Introductory statistics *why? class-wide  
WT + ref. mutants*

# Announcements

- Quiz, then load gel, and then pre-lab lecture
  - don't add loading dye to entire reaction!
- Purpose of shifting  $K_D$  for a sensor
  - what would happen if you used fluo-3 w/dendrites?

*everything bright / always on*

- what about purpose of shifting cooperativity?

*steep vs gradual change useful  
in different biological contexts*

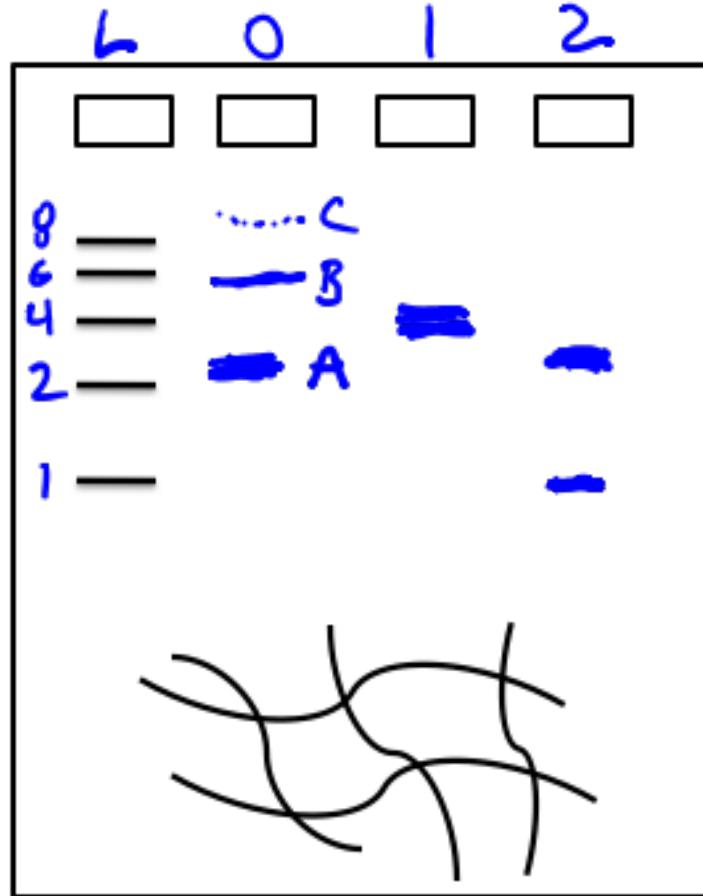
# Today in Lab: M2D3

- Set up gel of digested SDM product
  - ~~mark your area with colored tape~~
  - runs 45 min, we will photograph and post it
- Meanwhile...
  - pre-lab lecture
  - label tubes two ways (sticker top, sticker/marker side)
  - start FNT! *M2D4 tends to run long - prepare*
  - practice analysis *optional for now*
- Briefly analyze gel for DNA size and amount
- Finally, bacterial transformation – be gentle!
  - includes 30 min incubation step

# Polymerase error rates

- *Taq* polymerase ~ 1 in  $10^5$  # bp
  - Standard version has no proofreading capability  
↳ exonucleases
- *Pfu* polymerase ~ 1 in  $10^6$ 
  - Standard version requires longer extension times

# DNA EP: shape-dependence



e.g. 4Kbp

Plasmid versus linear samples

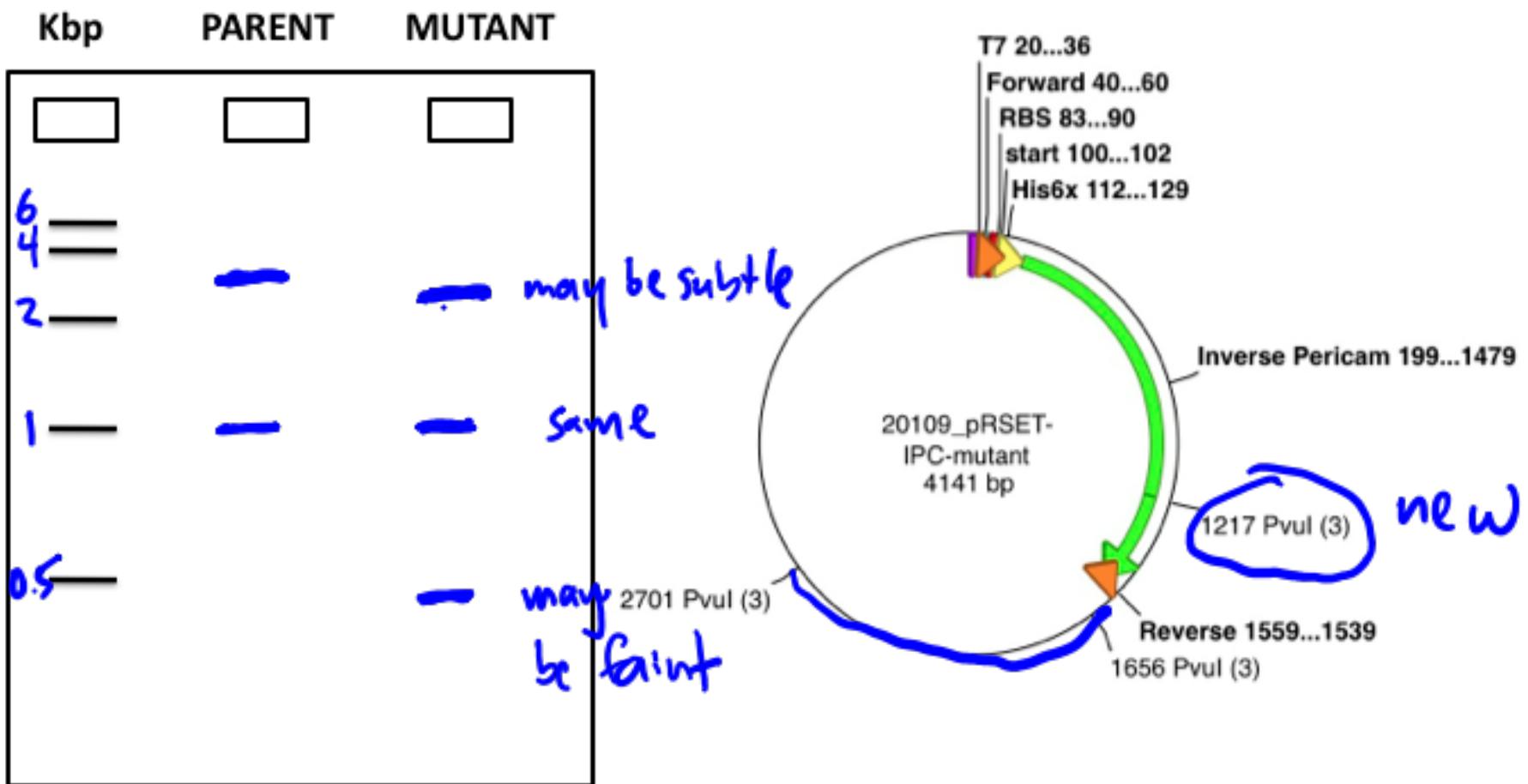
1-cut: linear DNA runs w/ ladder

2-cut: DNA must sum to whole  
e.g., 3+1 Kbp

- uncut:
  - A) supercoiled - faster
  - B) relaxed or nicked circular - slower
  - C) may be high MW concatemers

Remember to wear **nitrile** gloves.

# Diagnostic DNA gels



# Statistics review: basics

- Essential concepts: standard deviation ( $s$ ), mean ( $\bar{x}$ ), sample size  $n$ , degrees of freedom  $DOF$
- Normal (Gaussian) distribution



1  $s$  includes  
68 %  
of the data

x-axis: measured value (e.g., intensity)

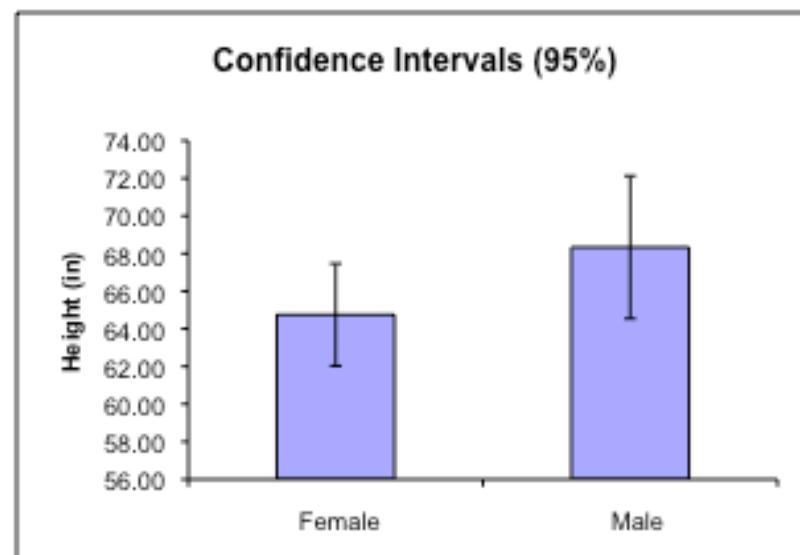
y-axis: # or fraction of samples w/ that value

## Confidence intervals (CI): principle

- Example:  $\bar{x} = 60$  (sample/measured mean) and 95% CI calculated to be  $\pm 3$  from real data
- Meaning: 95% of the time our population/true  $\propto n$  mean  $\mu$  lies in the range, here  $60 \pm 3$ 
  - subtly different from 95% likely that the range  $60 \pm 3$  contains the population (true) mean  $\mu$ , which we can't say
- 90% CI:  $\mu = \bar{x} \pm a$  where  $a < 3$ ,  $a > 3$ , or  $a = 3$ ?  
*trade-off between precision and confidence*
- Consider betting example
- What about  $n$ ? *as n increases for given C.I., more precise*

# Calculating confidence intervals (CI) *per a population*

$$\mu = \bar{x} \pm \frac{t s}{\sqrt{n}}$$



- $t$  is tabulated by DOF vs CI%
    - DOF =  $n - 1$  why?  $\sum \text{errors}(x_n - \bar{x}) = 0 \rightarrow \text{constraint}$
  - In Excel, us  $TINV$  function
    - input  $p$ -value =  $(100-\text{CI})/100$
- ; if C.I. = 95%  
 $p = 0.05$

# Introduction to t-test

- Every statistical test
  - Has *assumptions*
  - Asks *a specific question*
  - Requires *human interpretation (algo)*
- Some t-test assumptions
  - normal distribution (cf. Mann-Whitney test)
  - equal variances (type 2 in Excel; type 3 unequal)
- Posing a question *are mean male and female heights different at a confidence level of 95%?*

# Calculating t-test significance

$$t_{\text{calc}} = \frac{\bar{x}_1 - \bar{x}_2}{\text{pooled } s} \sqrt{\frac{n_1 n_2}{n_1 + n_2}}$$

$$\text{DOF} = n_1 + n_2 - 2$$

$t_{\text{table}}$  listed by DOF  
vs CL (confidence level)

- If  $t_{\text{calc}} > t_{\text{table}}$  difference is significant at that CL
- In Excel, use *TTEST* function
- Excel returns *p*-value  $\rightarrow$  confidence level (CL)  
 $\text{e.g., } p=0.01 \rightarrow CL=99\%$
- 1-tailed vs. 2-tailed test
  - 1- one-sided; hypothesis in advance
  - 2- full distribution; no a priori hypothesis

## Practice assignment for today

- Female heights: 61, 65, 61, 68, 65, 63, 61, 67, 60, 63, 64
- Male heights: 72, 72, 70, 65, 72, 69
- Calculate 95% CI for each mean
- Plot means on bar graph with CI error bars
- Apply t-test to the means
  - for multiple comparisons, ANOVA is better
  - comparing many means requires correction
  - remember,  $p = 0.05$  means 1 in 20 false positives!