

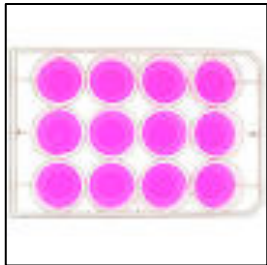
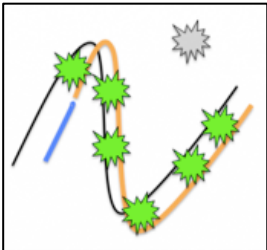
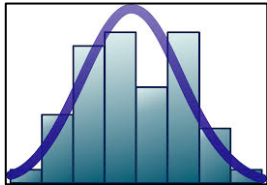
Congratulations on excellent journal club presentations!

M2D9: data analysis

04/13/2017



Today in lab



- Quiz
- Practice statistics
- Analyze qPCR data
- Analyze cell viability data

Wrapping up M2!

- Lab notebook:
 - M2D7 (RNA-Seq data analysis) graded by Rob at 10pm tonight
- Blog posts
 - on journal club due 10pm on Saturday, April 15
 - on M2 in general due 10pm on Sunday, April 23
- M2 research article 20%
 - due 10pm on Saturday, April 22
 - extra office hours: (56-302)

Monday	no office hours	Noreen
Tuesday, April 18	9:30-11:30am	Leslie
	3:00-5:00pm	Noreen
Wednesday, April 19	9:30-11:30am	Leslie
	12:00-2:00pm	Maxine
	2:00-5:00pm	Noreen
Thursday, April 20	9:30-11:00pm	Maxine

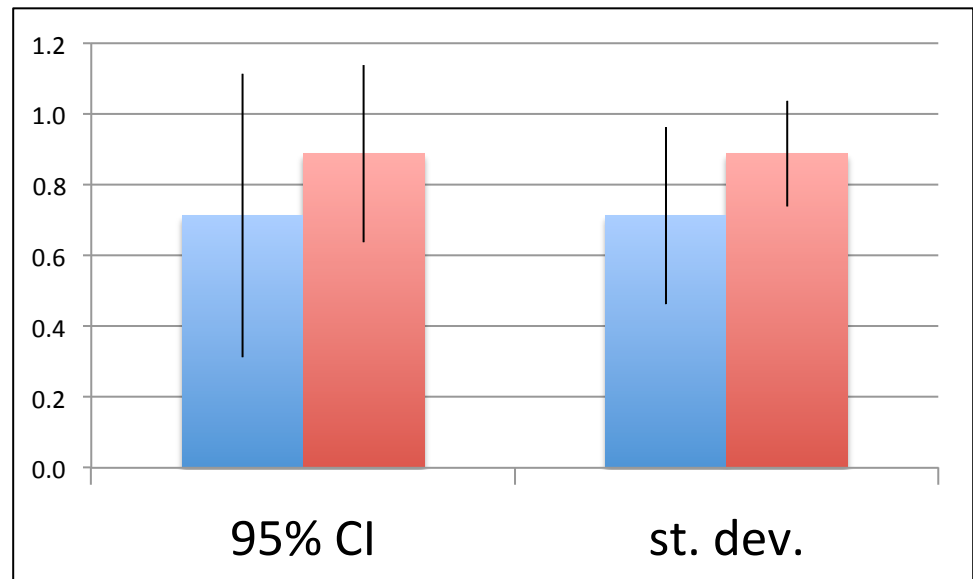
Plot error bars as 95% confidence intervals (CI)

- **95% CI**: the true value of the parameter will be within this interval 95% of the time if the experiment is repeated
- In Excel, find lower and upper bounds:

$$\bar{x} \pm \frac{t_{table} * stdev}{\sqrt{n}}$$

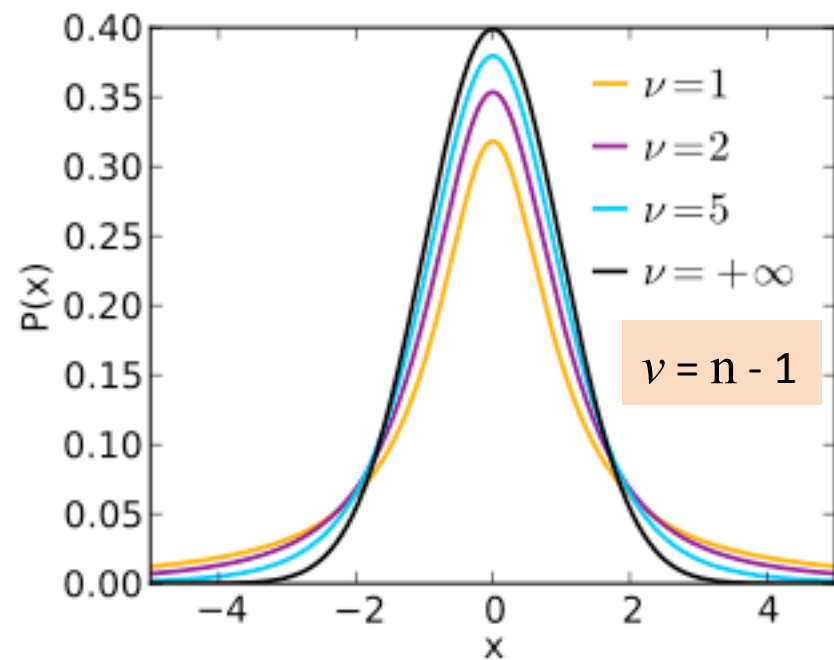
$$t_{table} = TINV(0.05, n - 1)$$

0.05 is the complement of 0.95 (of 95%)
n - 1 is the number of degrees of freedom



Student's t -distribution and t -test

- “Student”
 - pen name of William Sealy Gosset (Guinness brewery in Dublin !)
- t -distribution
 - symmetric and bell-shaped probability distribution, with heavier tails than the normal (Gaussian) distribution
 - to estimate the mean when sample size (n) is small and standard deviation unknown
- t -test
 - the test statistic follows a t -distribution under the null hypothesis
 - used to determine if two sets of data are significantly different



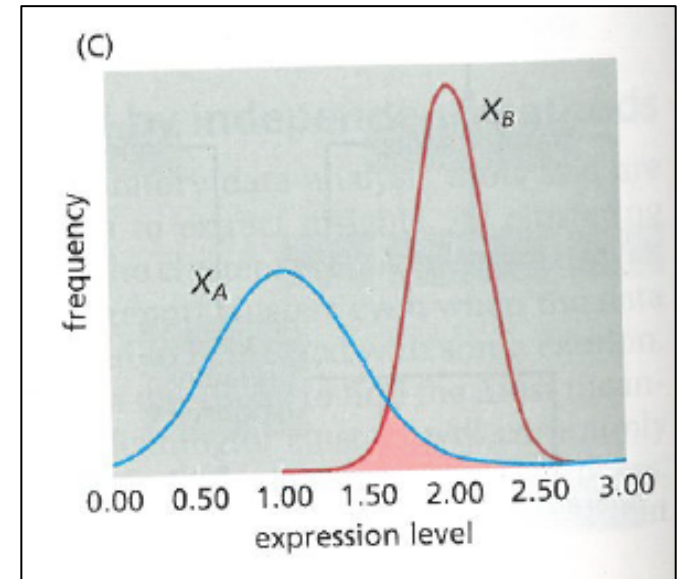
t-test in Excel

- How certain are you that two populations are different?

$$p = TTEST(array1, array2, 2, 3)$$

2-tailed

unequal variance (heteroscedastic 😊)



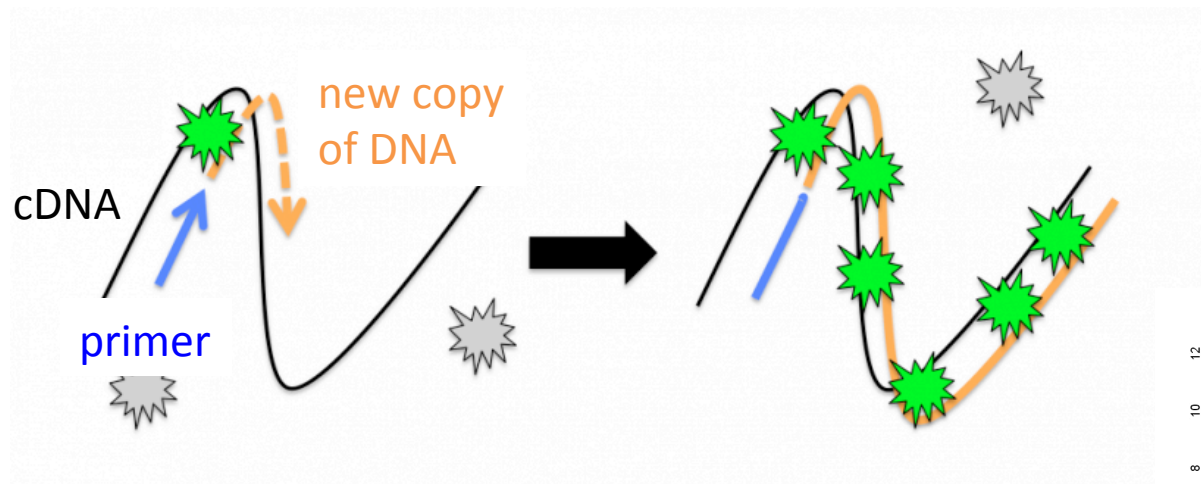
- ✧ The Student's t-test only applies to **two** data sets.
Only compare two conditions at a time.

- $p < 0.05$
 - typically considered “statistically significant”
 - the two data sets are different, have a different mean

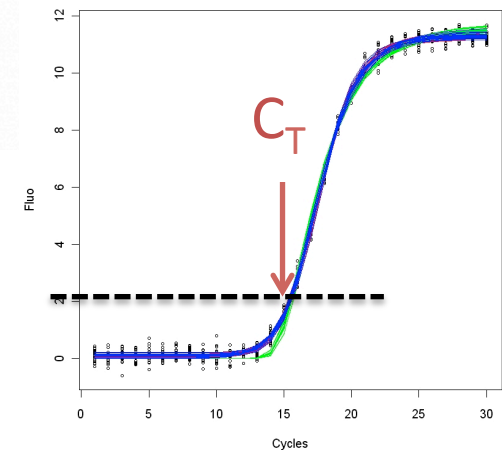
Now that you've practiced...

Analyze quantitative PCR data

- Monitor PCR as it occurs
 - using dye that is fluorescent when DNA is **double-stranded**
 - **signal** proportional to initial amount of cDNA (= original RNA)



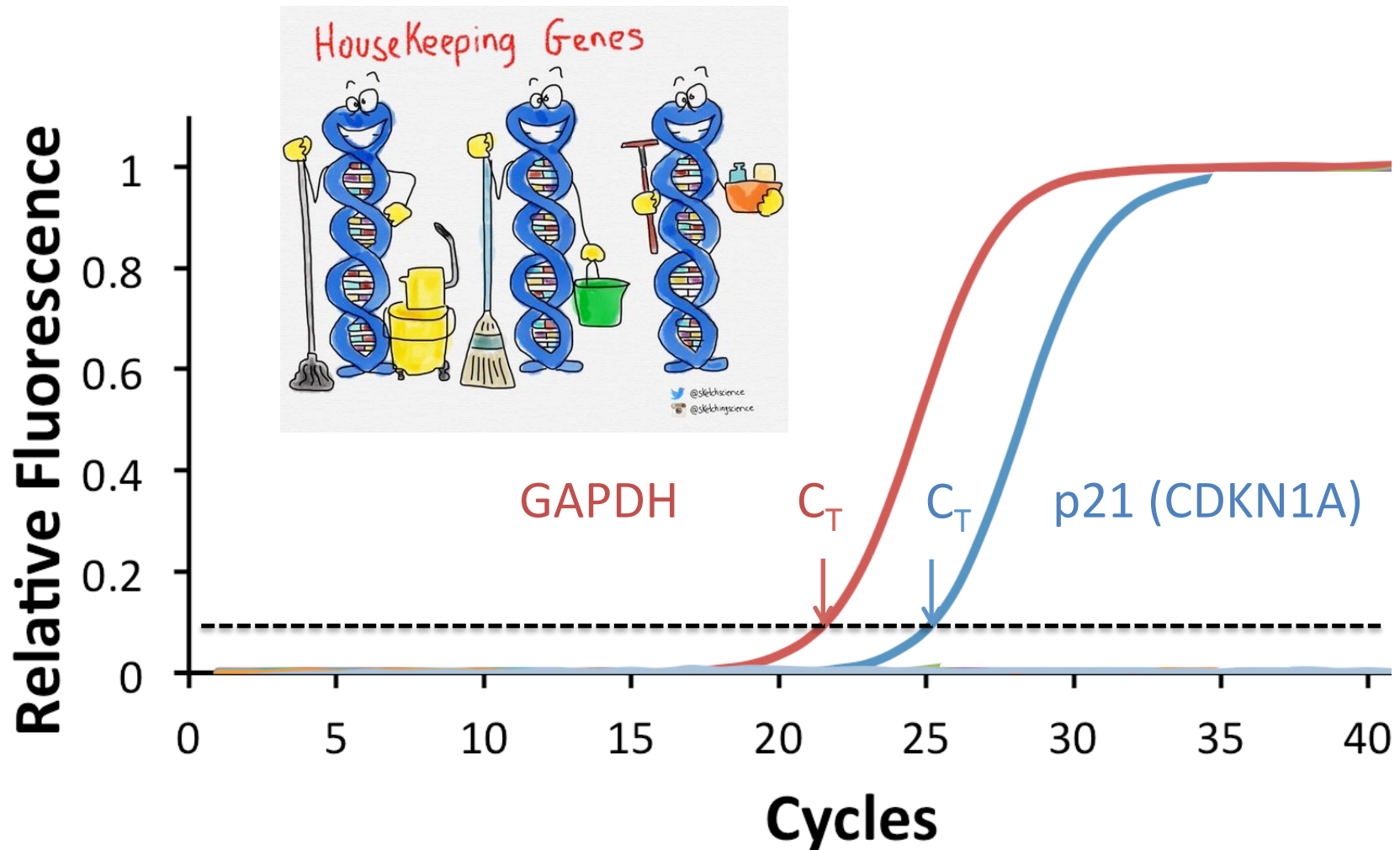
- Plot fluorescence vs. cycle number
- Extract C_T threshold cycle



0

40 cycles

C_T value \sim amount of cDNA template present at the start of the amplification reaction



The students' Ct values were ~ 40 (end of qPCR!)
Most likely the RNA had been degraded (by RNases),
the cDNA was of poor quality.

What do we calculate from these C_T values?

$$\Delta C_T = C_T (\text{p21}) - C_T (\text{GAPDH})$$

	DLD-1	DLD-1 + etoposide	BRCA2-/-	BRCA2-/- + etoposide
1	$2^{-\Delta C_T}$	$2^{-\Delta C_T}$	$2^{-\Delta C_T}$	$2^{-\Delta C_T}$
2	$2^{-\Delta C_T}$	$2^{-\Delta C_T}$	$2^{-\Delta C_T}$	$2^{-\Delta C_T}$
3	$2^{-\Delta C_T}$	$2^{-\Delta C_T}$	$2^{-\Delta C_T}$	$2^{-\Delta C_T}$
average				

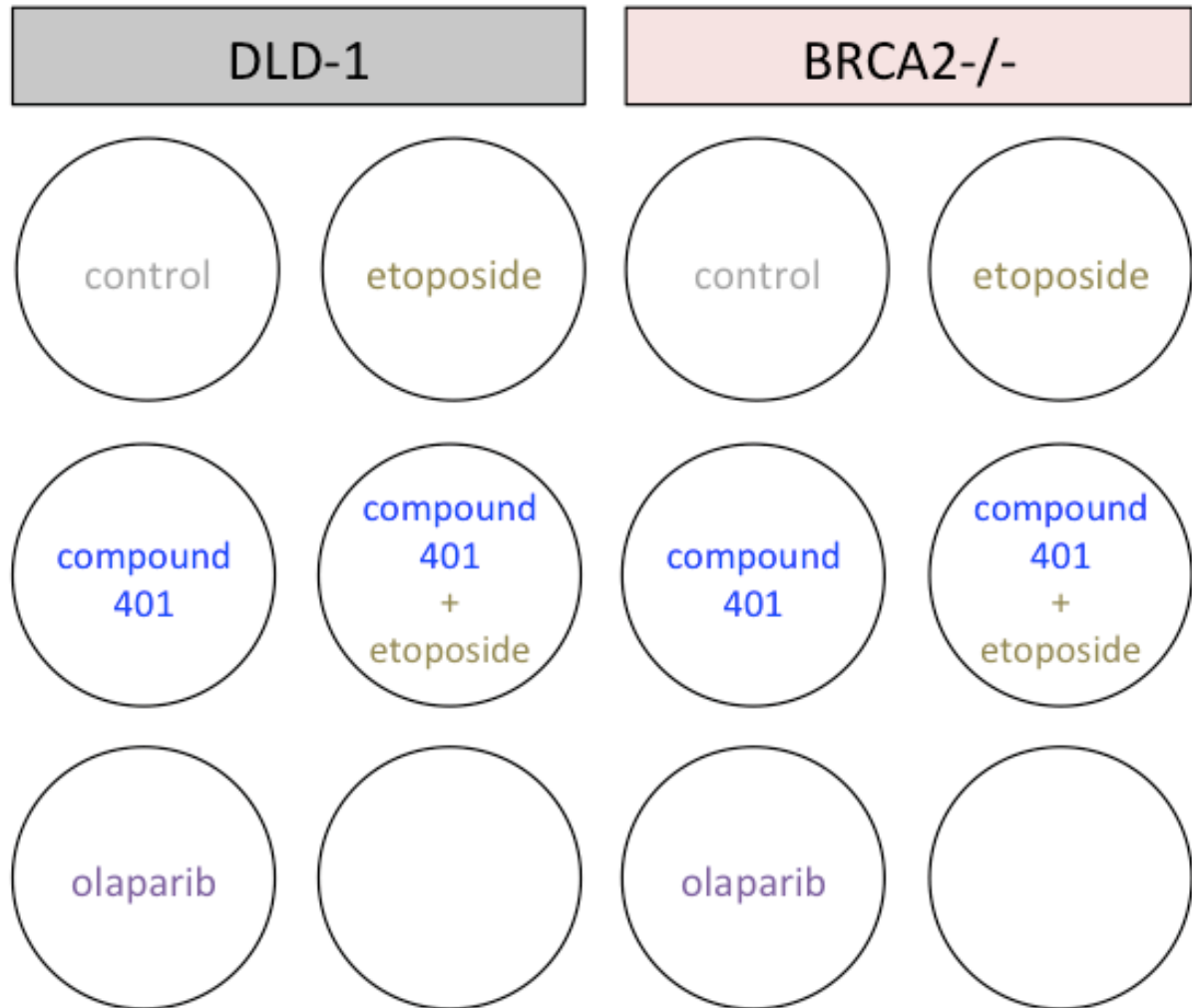
- Compare p21 mRNA levels in 4 conditions

Analyze cell viability data

- etoposide:
creates double-stranded
breaks

- compound 401:
inhibitor of DNA-PK
NHEJ

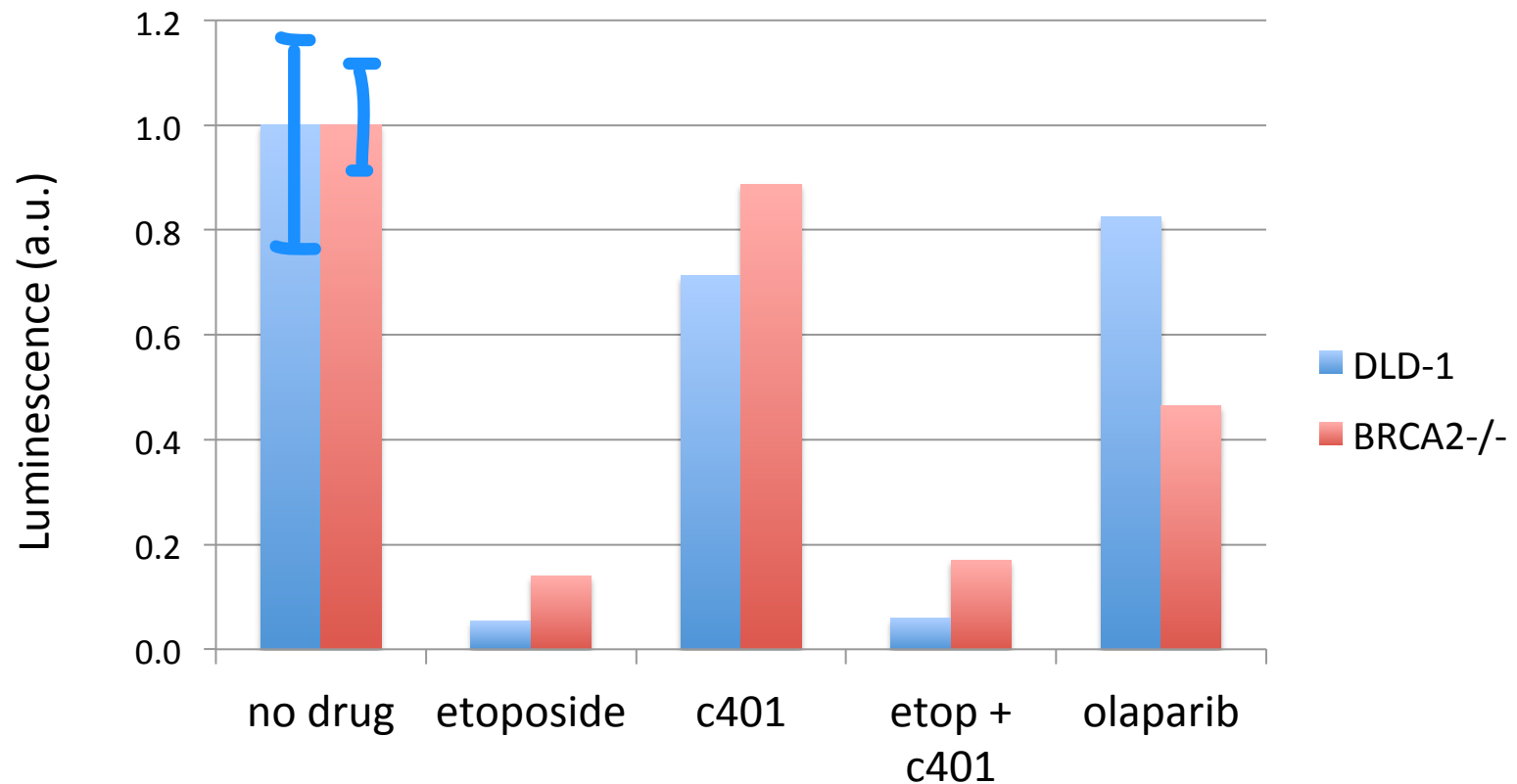
- olaparib:
inhibitor of PARP
BER



Analyze cell viability data, **with error bars**

using the class's 13 replicates

[http://engineerbiology.org/wiki/Talk:20.109\(S17\):Module_2](http://engineerbiology.org/wiki/Talk:20.109(S17):Module_2)



Today in lab

- Practice statistics
- Analyze qPCR data
- Analyze CellTiter Glo cell viability data
- ❖ Review Prof. Samson's M2L8 lecture for ideas
 - In Discussion / Future Directions, propose 2 years of research
 - to further elucidate DLD-1 / and BRCA2-/- genotype
 - and/or to continue answer "How does DNA repair affect the ability of chemotherapeutic drugs to kill cancer cells?" and "How does chemotherapy affect gene expression?"

M2: Experimental overview

