

# Digression

costume day



just pretty much  
always awesome day



- Announcements
- Lab Quiz
- Pre-lab Lecture
  - ❖ Recap SDS-PAGE
  - ❖ Sequence analysis preview
  - ❖ Today in Lab (M2D6)

# Announcements

- Next time I'll need to leave early and Shannon (T/R section leader for Mods 2 and 3) will come help out
- Please email us your experimental plans as you develop them
- Next journal club in 1 week
- Report due M 11.12.12 → fill out survey for extended OH

# Mutant Miller assay data

Mutant 1 B-Gal Average (Dark)	Mutant 1 B-Gal Average (Light)	Mutant 1 T541X; X =	Mutant 2 B-Gal Average (Dark)	Mutant 2 B-Gal Average (Light)	Mutant 2 T541X; X =
<del>1.2-fold</del>					
2328.25	1928.66	1.2	--	--	
1417.88	1409.76	1.0	1450.07	454.78	3.2

Wild Type System B-Gal (Dark)	Wild Type System B-Gal (Light)	H243A Mutant B-Gal (Dark)	H243A Mutant B-Gal (Light)
<del>2.6-fold</del>		<del>1.2-fold</del>	
1099.13	414.85	54.54	45.27
1099.13	414.85	54.54	45.27

Comparisons to make?  
Representing data?

Go, Team Pinkle!

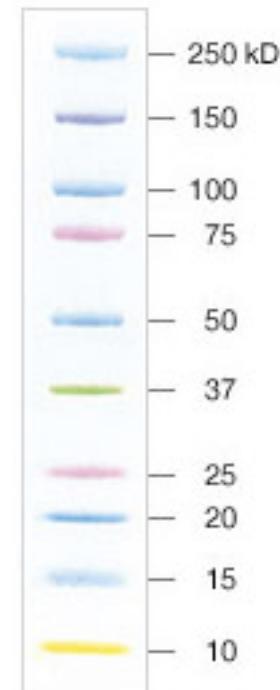
# SDS-PAGE preparation

acrylamide monomer neurotoxic

- You will make whole cell extracts with equal cell #s
  - Based on OD<sub>600</sub> reading, normalize  $\textcircled{1} \text{ OD}=1.0$   $\textcircled{2} \text{ OD}=2.5$   
Goal: 2 OD  
per sample  $\textcircled{1} \text{ 2 mL}$   
 $\textcircled{2} \text{ 4 mL}$
- Gel separates proteins based on size, shape, charge
- Sample preparation reduces analysis complexity
  - SDS: coat proteins w/(-) charge
  - $\beta$ -Me: breaks SS bonds  $\rightarrow$  tips in hood
  - Boiling: denature higher order
  - Sample Buffer has SDS,  $\beta$ -Me, plus glycerol,  $\beta$ PB dye

# SDS-PAGE visualization, analysis

- Determine size in comparison to ladder (pre-stained)
- Visualize specific protein amount with antibody and staining
  - antibody conjugated to enzyme
  - enzyme catalyzes colorimetric rxn
  - more detail next time
- Goal: determine if mutation affected Cph8 production



Kaleidoscope

# Sequencing food for thought

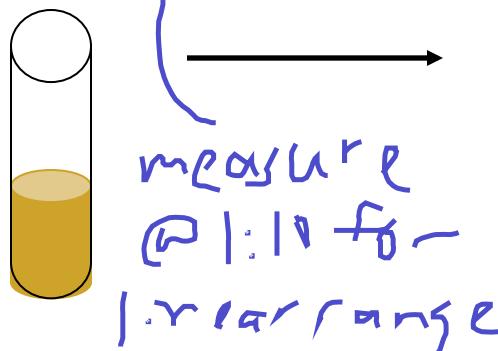
- Where in the sequence do I expect a mutation?
- How do I interpret mutations occurring elsewhere?
- How do I interpret gaps (vs mismatches)?
- What resources may help me form a hypothesis about the behavior of a particular mutant?

Query	194	CTGGCGGANGACCGCACGCTGCTGATGGCGGGGTAAGTCACGACTTGC	CGTTCCGCTG	253
Sbjct	1570	CTGGCGGATGACCGCACGCTGCTGATGGCGGGGTAAGTCACGACTTGC	CACGCCCGCTG	1629



# Today in Lab (M2D6): Workflow

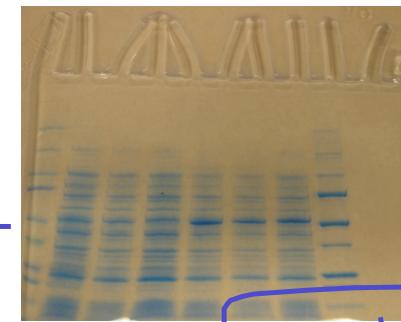
Check OD of cells



Lyse an aliquot  
and load onto  
SDS-PAGE

Spin 20D, resus. 100 μL  
30 μL + 30 μL SBS

Also ladder & (+) control



Let run 1 hour

Transfer to blot – run 1 hour

Meanwhile...

Sequencing analysis (ApE or BLAST)

Mutant bacterial photograph *maybe*

Store in milk solution  
until next time

Ideas/prep for experiments next time... from seq/β-gal repeats to ?your idea here?