

- Announcements
- Lab Quiz
- Pre-lab Lecture
 - ❖ Where we are/going
 - ❖ Mutant phenotypes
 - ❖ Today in Lab (M2D5)

Announcements

- Methods sections
 - see general comments, apply to many of you
 - review writing guidelines
 - breakdown a published journal article as a model
 - meet with me or WAC for help revising
- Plus one more iPad experiment

• Plus music: oops

Where we are/going

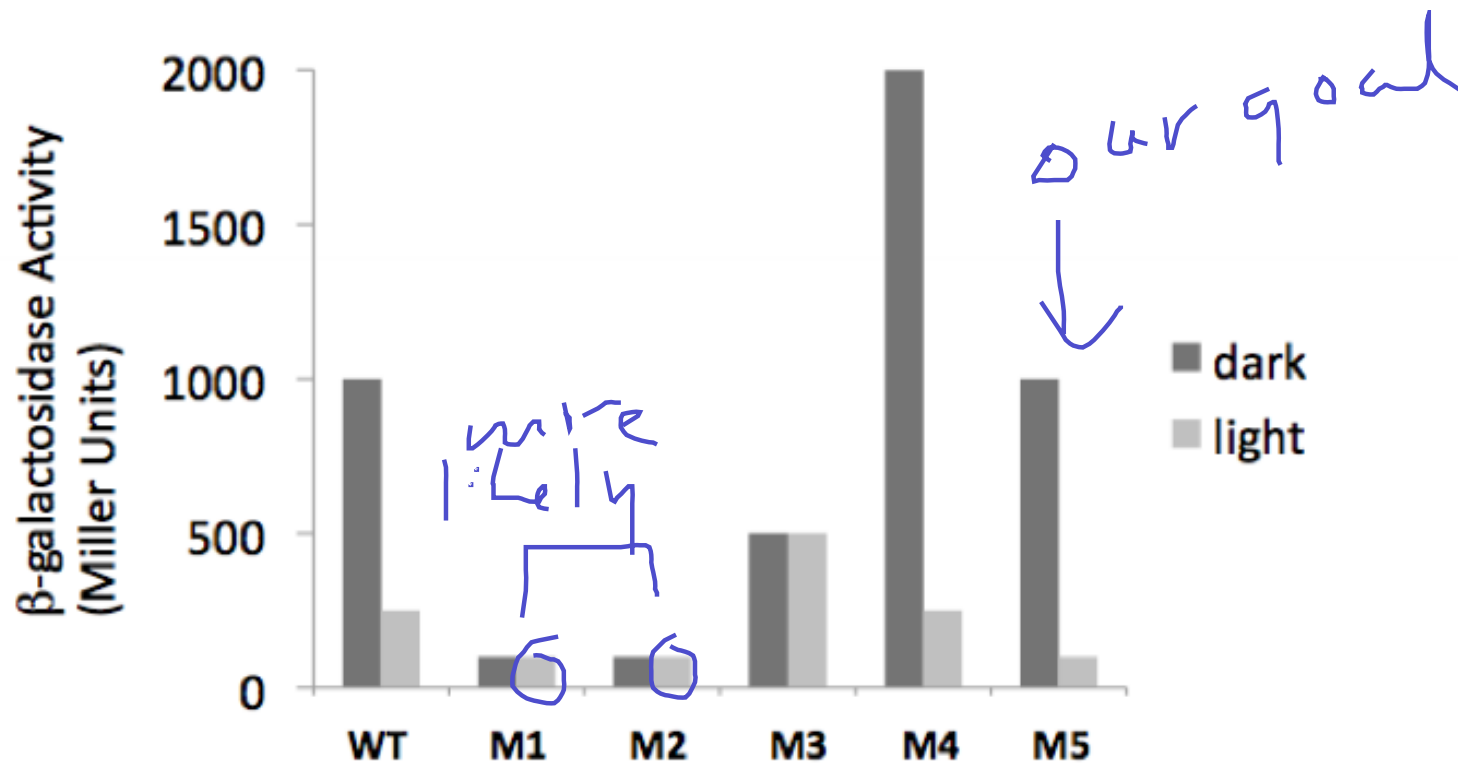
- Each colony you chose was re-streaked, then grown O/N in light and dark. Now what?

- 1 β -gal assay — to compare mutant to original
* perform as similarly as possible (OD)
 - 2 Sequence DNA — what is each mutant?
* are there other mutations?
 - 3 is [Cph8] protein itself changed?
+ * potential for artifacts
 - 4 new photo — compare to original
+ ideally, would run same day, with same light/media conditions, cell & D
- LATER

Chance for design/investigation

- Day 7 = Western = lots of long incubations
- Chance to design and carry out an additional experiment that could shed light on your results
- Start thinking about it now (or rather as soon as you have your first batch of mutant data)
- Also a day to share, collaborate with other groups

β -gal assay of K-P+ candidates



Cph1/EnvZ	A553	G554	V555	S556	H557
EnvZ	A239T	G240E	V241G	S242D	H243A

Slide from N. Kuldell

Today in Lab:M2D5

- Miniprep each mutant candidate (light *or* dark)
- Resuspend in sterile water, mix an aliquot with sequencing primer → ideally done by 3 pm!
- Linda will discuss paper structure @ 4 pm