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  - ❖ Today in Lab (M2D6)

# Announcements

- Final M2 quiz on Wed 11/5
- Journal club next time
  - 16-336 starting at 1:15 pm sharp
  - upload slides to Stellar in advance
  - T/R guest, then your 7 talks → 8 total
- Report due Wed 11.12.12 *main data probably 11/5*
  - best day(s) for extra OH?

R 6 + TH?  
F 7 + M 10?

other combos?  
other days?

# Troubleshooting catch-up

- Problem(s): light/dark WT backwards; magnitude lower
- Conditions for repeat: → verifying WT ✓ mutants - not yet
  - fresh antibiotics + media
  - WT/KD: fresh re streaks from freezer stocks ✓
  - M1/2: re streaks (from lig. cult.) on plain LB/antibiotic plates
- Today's samples: Today's samples re-inoculated in fresh LB/antibiotics hope for best!
- Review reasons for having signaling but no light response
  - 1) basal / background / leaky expression

$p(\text{event}) \uparrow \text{ or } \downarrow$  not binary on/off
  - 2) cross-talk when cognate partner is missing
  - 3) reminder: light turns signaling off, not on, in our system

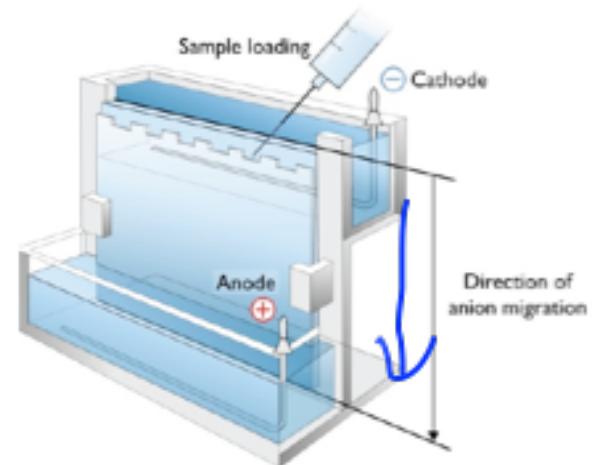
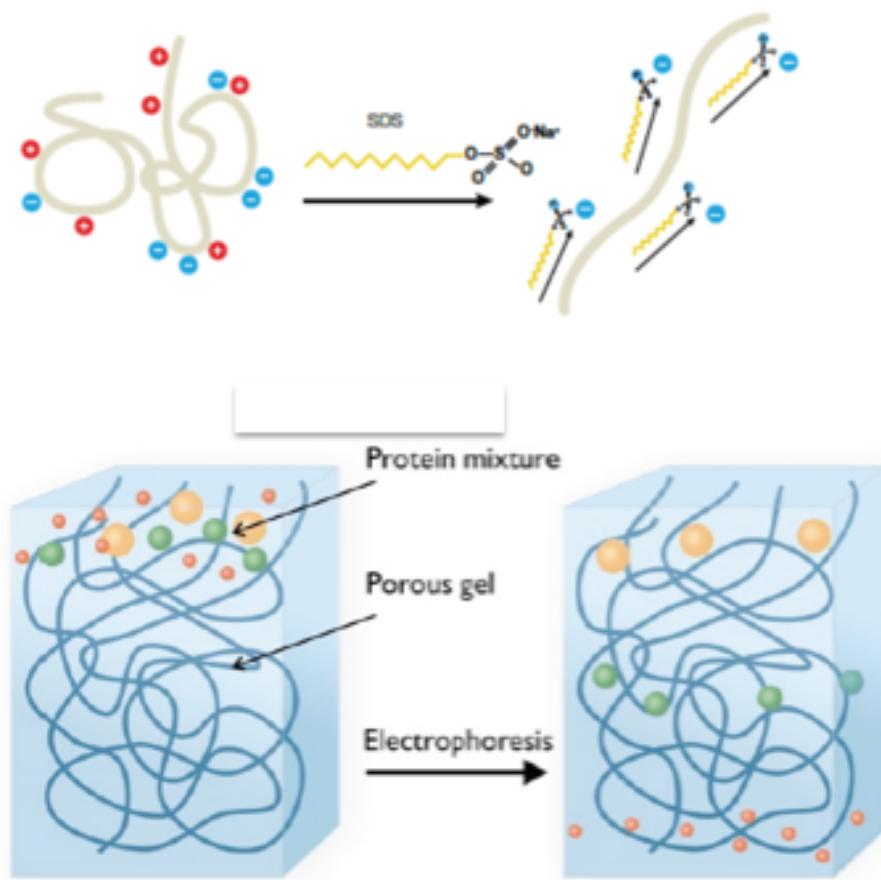


# SDS-PAGE preparation

↳ acrylamide monomer is toxic → why?

- You will make whole cell extracts with equal cell #s
  - Based on  $OD_{600}$  reading, normalize  
Goal:  $2^{OD}$  per sample
  - sample ① 2 mL  
② 4 mL
- Gel separates proteins based on size, shape, charge, make uniform
- Sample preparation reduces analysis complexity
  - SDS: coat proteins w/  $\rightarrow$  charge
  - $\beta$ -Me: breaks S-S bonds  $\Rightarrow$  tips in hood to air out
  - Boiling: denature higher-order structures
  - Sample Buffer has SDS,  $\beta$ -Me, plus glycerol,  $BPI\beta$  dye

# SDS-PAGE separation

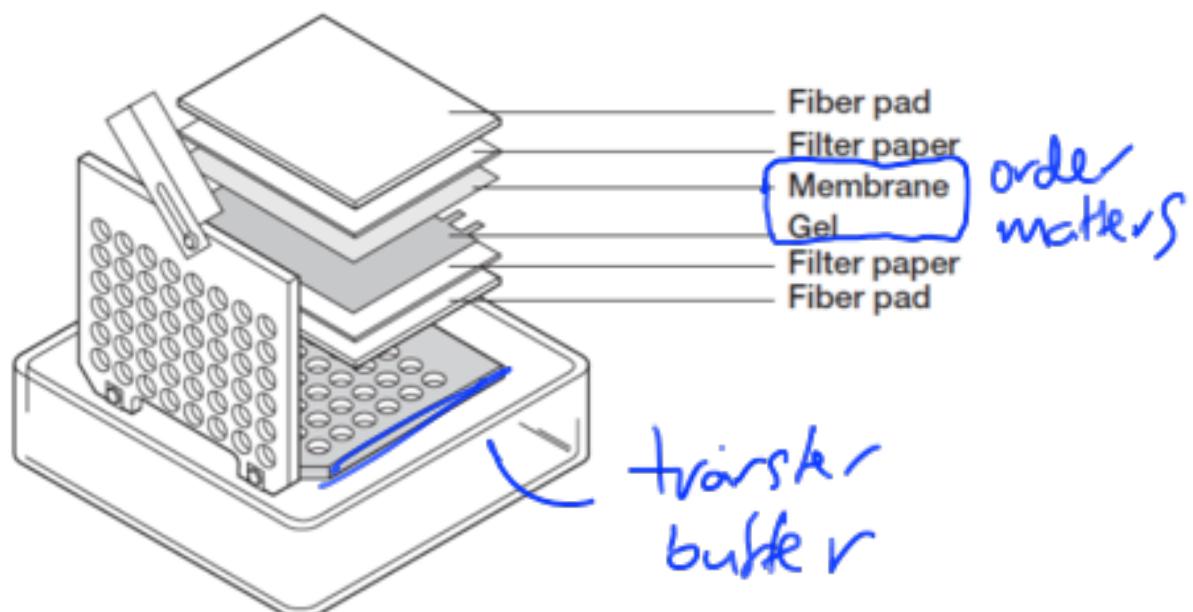


TG(S)<sup>SDS</sup>  
buffer

Slide adapted  
from Noreen L

# SDS-PAGE blotting

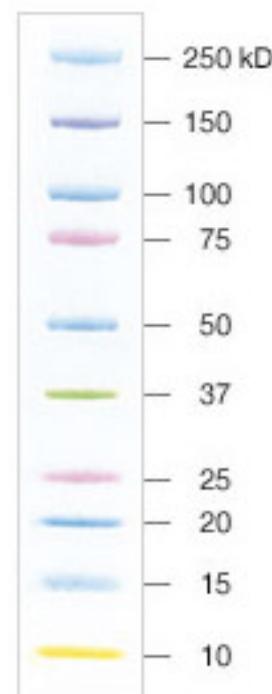
- Transfers proteins from polyacrylamide gel onto a nitrocellulose membrane
- Notes:
  - Eliminate air bubbles
  - Do not touch nitrocellulose



Slide from Noreen L

# 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 → *Western*
- Goal: determine if mutation affected Cph8 production

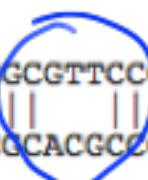


Kaleidoscope

# Sequencing food for thought

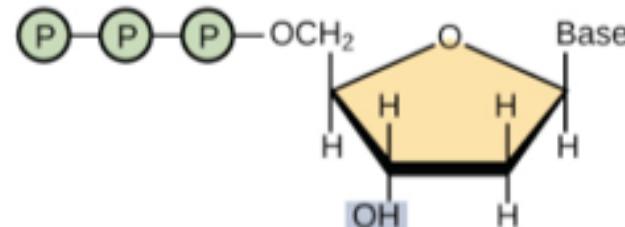
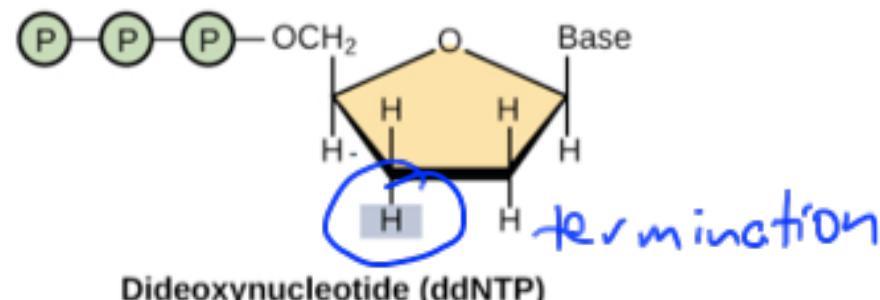
- Where in the sequence do I expect a mutation? T 541 X
- How do I interpret mutations occurring elsewhere?
- How do I interpret gaps (vs mismatches)?
- What resources may help me form a hypothesis about the behavior of a particular mutant?

Query	194	CTGGCGGANGACCGCACGCTGCTGATGCCGGGGTAAGTCACGACTTGCACGCTTCCGCTG	253
Sbjct	1570	CTGGCGGATGACCGCACGCTGCTGATGCCGGGGTAAGTCACGACTTGCACGCTTCCGCTG	1629



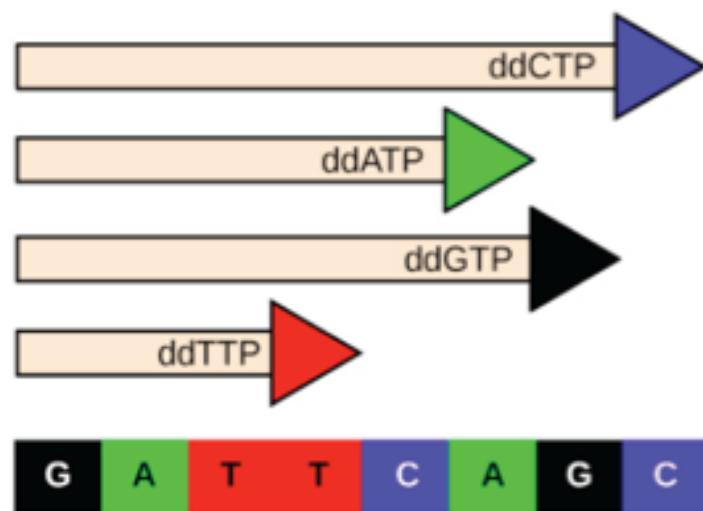
# Sanger sequencing approach

- Dye labeled dideoxynucleotides added
- Chain terminating reaction

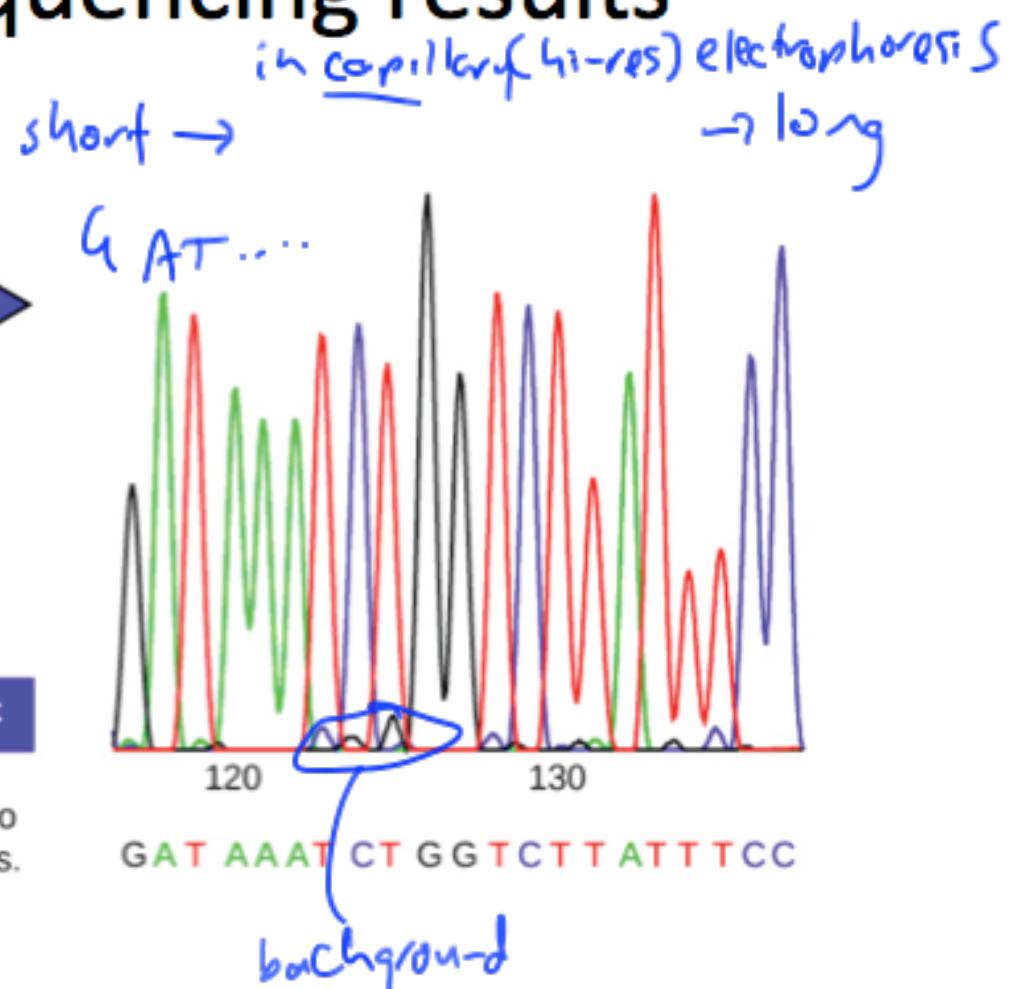


Slide from Noreen L

# Sanger sequencing results



Dye-labeled dideoxynucleotides are used to generate DNA fragments of different lengths.



Slide from Noreen L

# Sanger sequencing limitations

- Note: primer needs to anneal upstream of SOI
  - SOI = sequence of interest
- Max of ~ 1000 bp read lengths
- Sequence at beginning, end, is unreliable
  - start: need to establish baseline fluorescence
  - end: decreasing resolution in capillaries as DNA size ↑
  - both: less probability to form those products, more noise + less signal
- Note: it's okay to put "Ns" in DNA software

10 vs. 11 bp  
10% difference in length

100 vs. 101  
11  
120  
etc ...



# Today in Lab (M2D6): Workflow

Check OD of cells



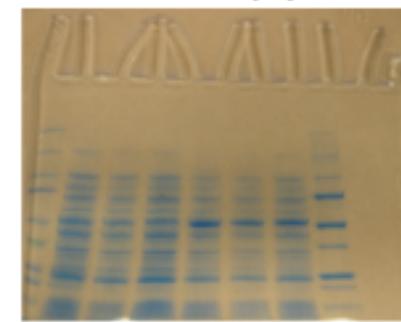
measure at a  
1: X that gives  
linear range  
on spec.  
(e.g.,  $X=10$ )  
Meanwhile



Lyse an aliquot  
and load onto  
SDS-PAGE

spin 2.0 OD, resuspend  
in 100μL → take 30μL,  
and add 30μL of 2X SB.

Also ladder & (+) control



Let run 1 hour → often more like 40'

Transfer to blot – run 1 hour



Store in milk solution  
until next time

Sequencing analysis (ApE or BLAST)

Mutant bacterial photograph – NOT YET!

Ideas/prep for Day 8 experiments... from seq/β-gal repeats to ?your idea here?