

M1D1: Prepare CometChip microwell array and practice cell culture

09/11/18

1. Lab Orientation Quiz
2. Pre-lab Discussion
3. $\frac{1}{2}$ class goes to the Tissue Culture Room
4. $\frac{1}{2}$ class prepares a CometChip

Office Hours

Noreen

Monday 2pm-5pm
in 16-317

Leslie

Thursday 2-3pm
Friday 12-1pm
in 56-341c

Josephine

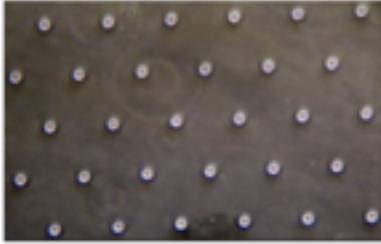
Wednesday 12-1pm
Friday 2-3pm

by appointment: nlyell@, lesliemm@, joshaw@

M1 major assignments

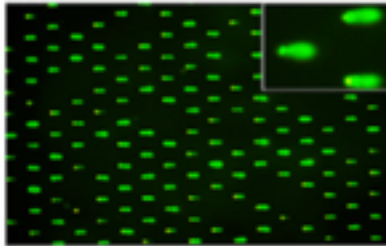
- Data summary (15%)
 - In teams, submit on Stellar
 - Draft due 10/8, final revision due 10/20
 - Bullet points, .PPTX
- Mini-presentation (5%)
 - Individual, submit video via Gmail
 - Due 10/13
- Lab quizzes –be on time!
 - M1D4 and M1D7
- Notebook (part of 10% Homework and Notebook)
 - Due 10/5 at 10pm, graded by Corban
- Blog: <https://be20109f18.blogspot.com> (part of 5% Participation)

Overview of Module 1: Measuring Genomic Instability



1. Optimize comet chip assay

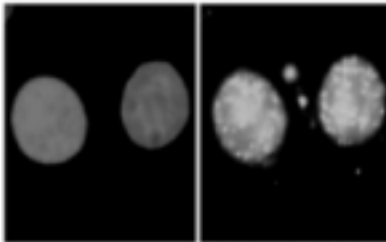
- Test loading variables



2. Use comet chip assay to measure DNA damage

- Measure effects of H_2O_2 on +/- DNA-PK cell lines

Protein involved in
DNA repair

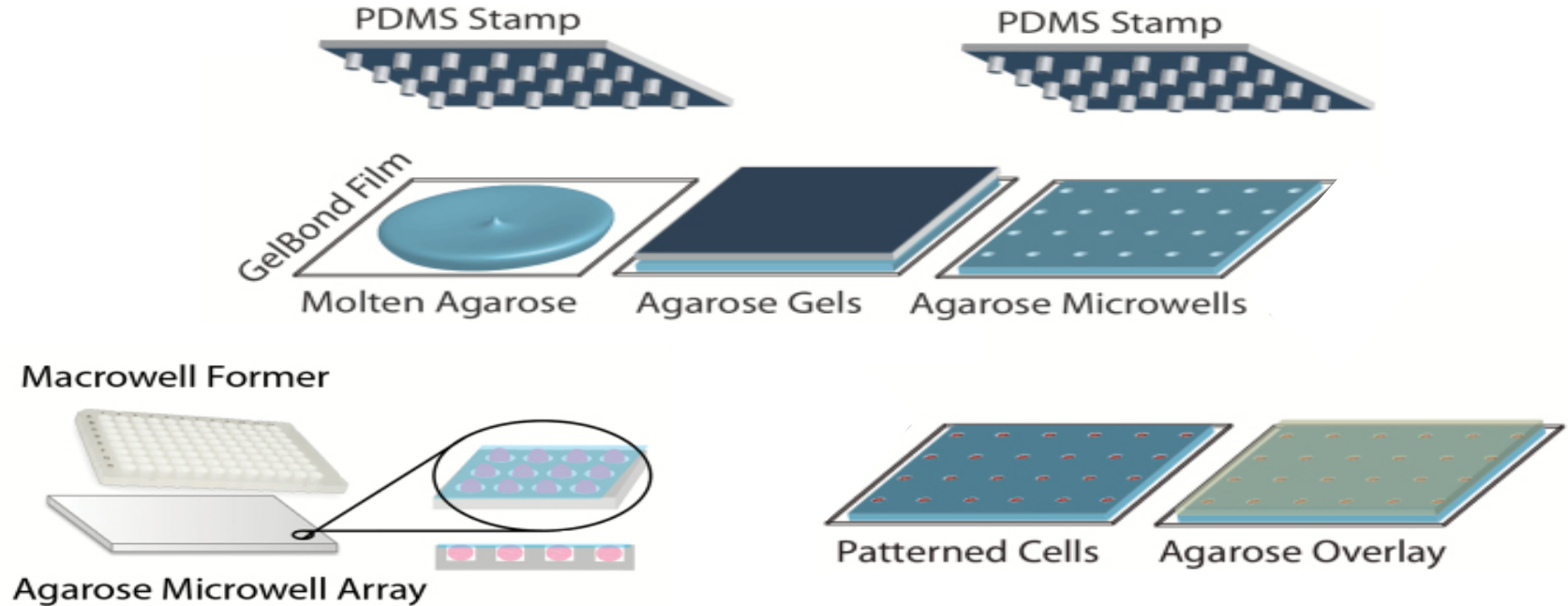


3. Use immuno-fluorescence assay to measure DNA damage

- Examine effect of H_2O_2 on γ H2AX foci formation

Overview of this week:

Create a CometChip & optimize cell loading

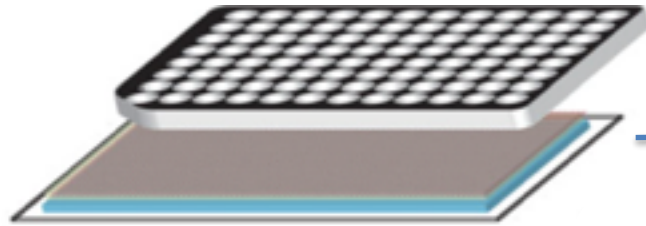


What is the minimum number of cells needed in each macrowell to obtain efficient loading?

Overview of next week:

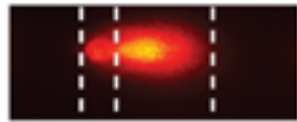
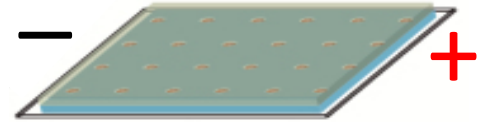
Assess DNA damage in tumor cells with & without DNAPKcs

Treat captured cells in comet chip
with H_2O_2 (oxidative damage)

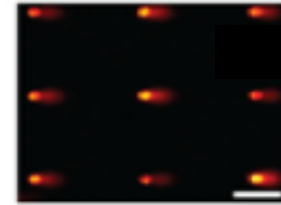
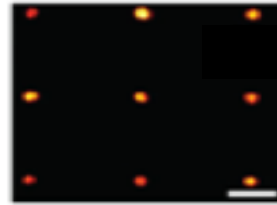


Lyse cells & unwind DNA
(DNA still captured
agarose in overlay)

Agarose Electrophoresis

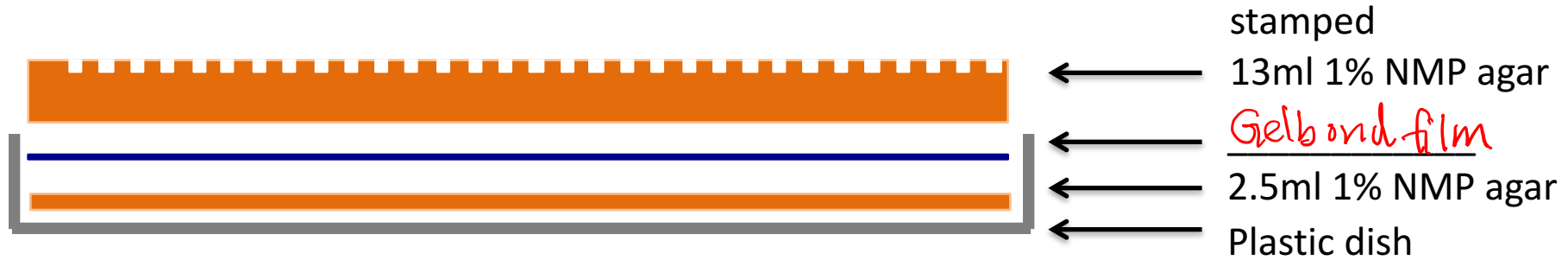


Analysis
via
Matlab

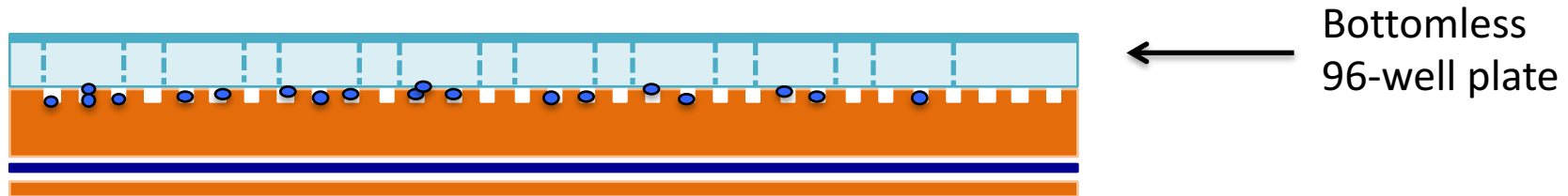


Stain DNA and image via
fluorescence microscopy

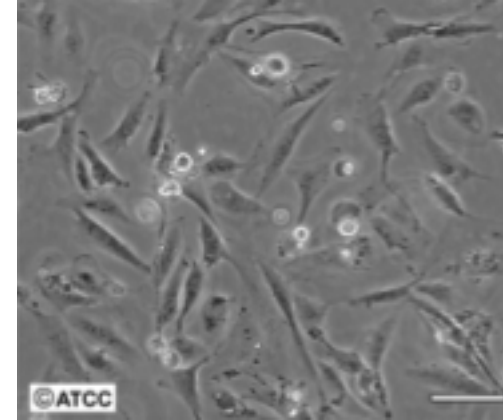
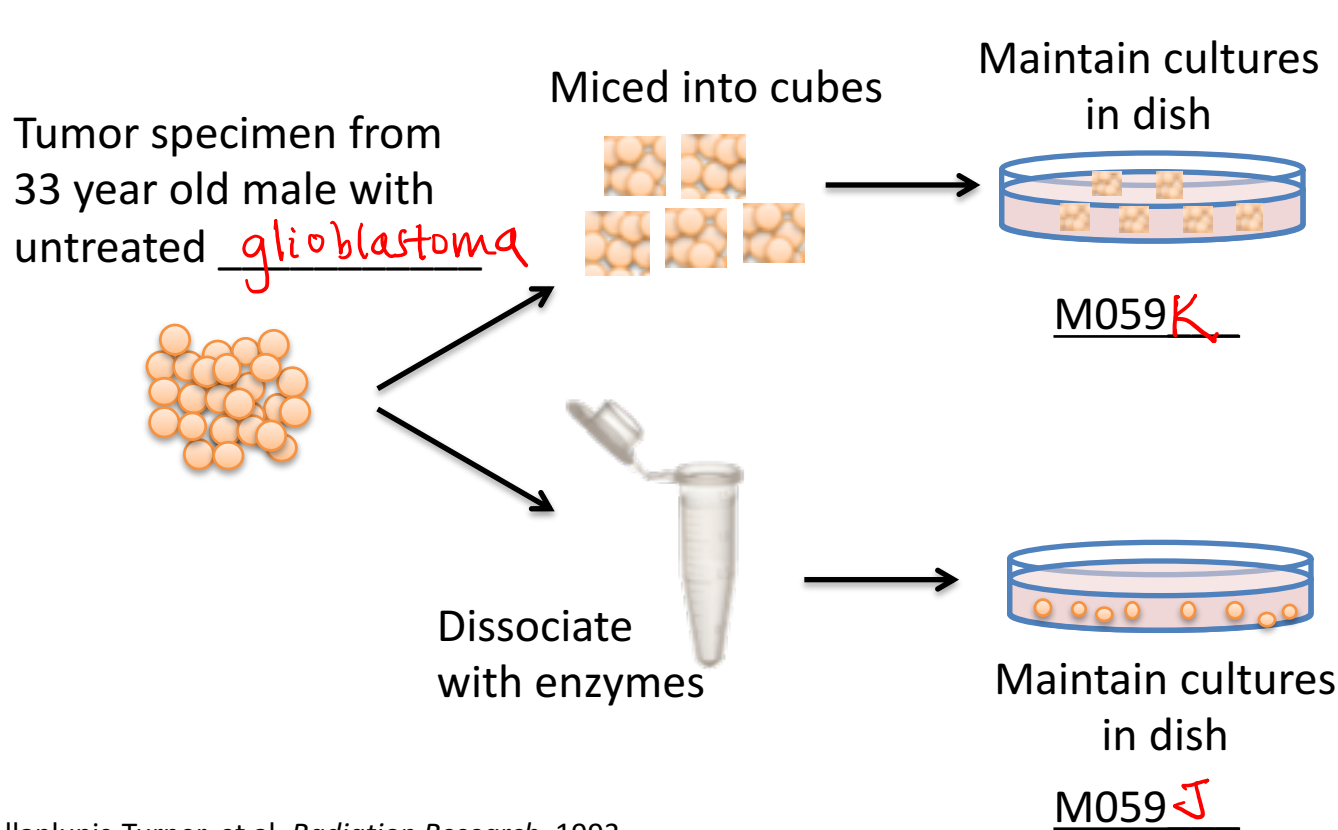
Today, make a CometChip



Next time, load cells

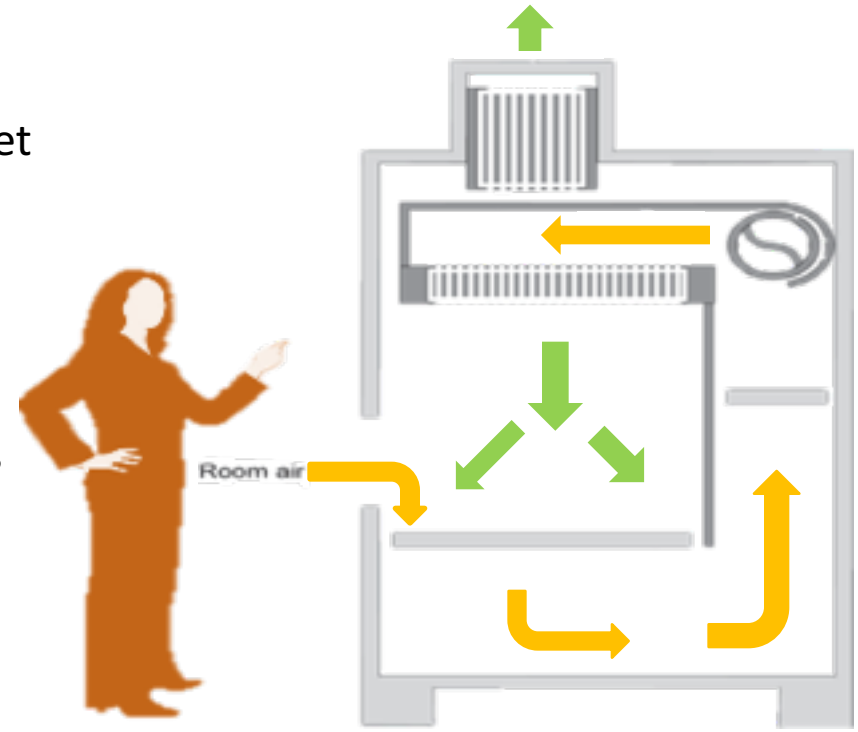


Background: Two glial cell lines— M059J (-DNAPKcs) and M059K (+DNAPKcs)



Tissue culture sterile technique

- **70% ethanol** everything:
 - Wipe cabinet before and after use
 - Wipe everything that enters the cabinet
 - Do not spray cells with EtOH
- **Do not disturb air flow:**
 - Do not block grille or slots
 - Minimize side-to-side arm movements
 - Work > 6" away from sash
 - Leave blower *on always*
- Do not talk into incubator!
- Only open sterile media in hood



Mammalian cell culture medium

What do cells need to survive?



Food:

- DMEM (Dulbecco's Modified Eagle's medium) : F12 (defined)
 - amino acids
 - sugars
 - salts
 - phenol red – pH indicator
 - pH buffers
 - vitamins
- FBS (fetal bovine serum) (undefined)
 - growth factors
 - lipids
 - cholesterol
 - cytokines



Non-food:

- antibiotics:
 - penicillin
 - streptomycin
- prevent bacterial growth



Mammalian cell culture terminology

- Confluence

density

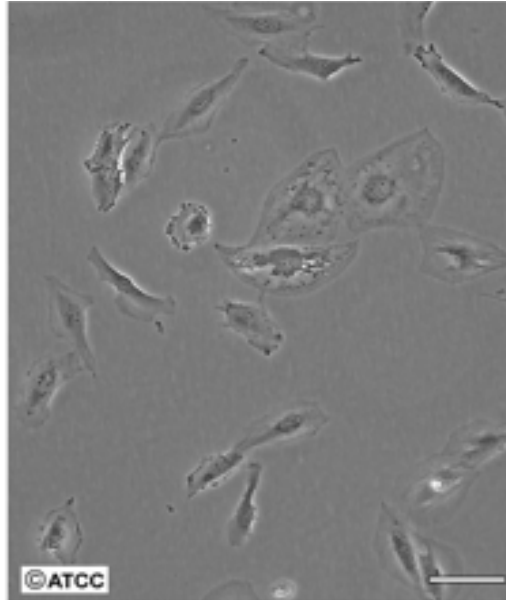
- Splitting (subculturing)

*~80% confluent,
put cells on new dish*

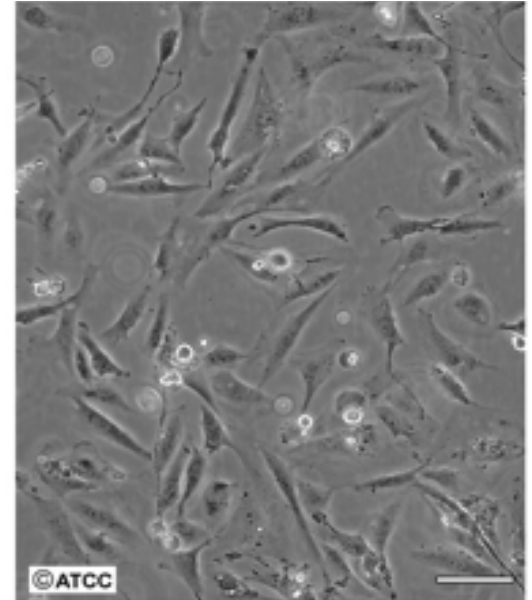
- Seeding

*~20-40% of confluent
culture on to new dish*

Low Density



High Density



General steps for splitting cells +WHY?

1. Look at cells, estimate confluence

Be sure
not to mix
the two
cell lines!

T25

2. Rinse with PBS

Wash away dead cells, debris, anti-trypsin agents

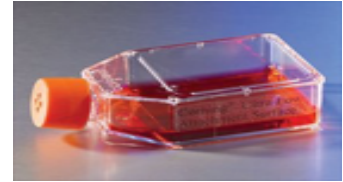
3. Detach cells with trypsin (enzyme)

break substrate adhesions

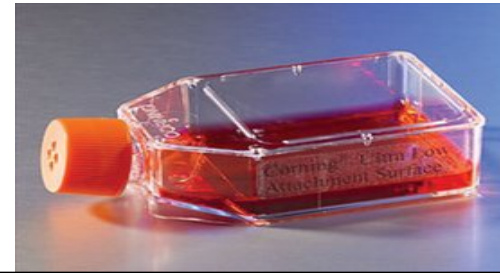
4. Count cells

5. "Seed" new culture vessel

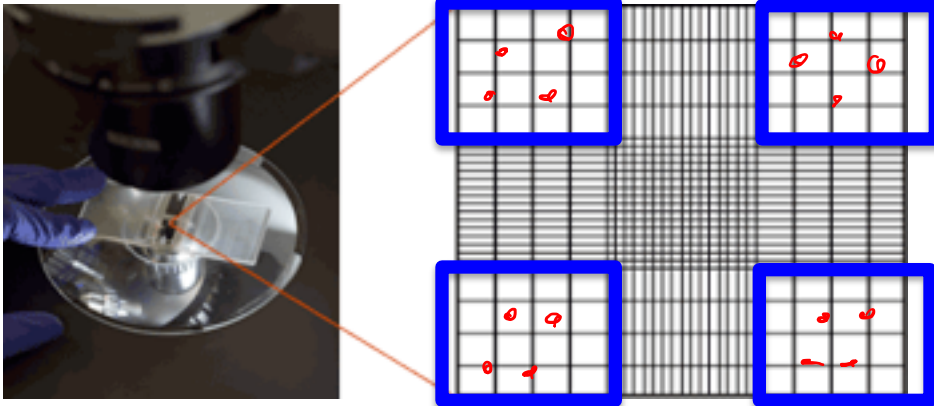
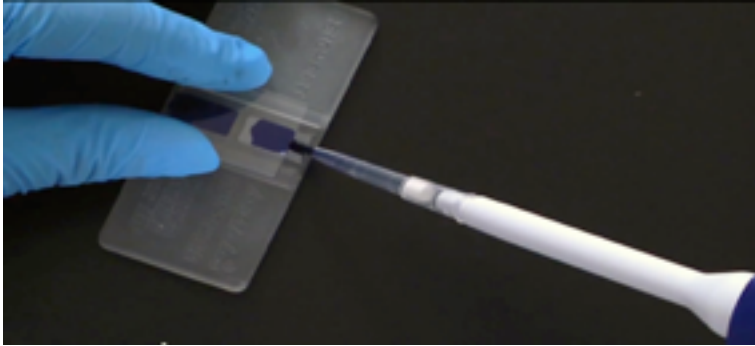
give room to grow



T75



Counting Cells



- Hemocytometer:
- Trypan blue:
stains dead cells
- # cells / mL = 10,000 x
average of 4 corners

$$16 / 4 = 4$$

$$4 \times 10,000 = 40 \text{ k cells/mL}$$

What should go in your notebook?

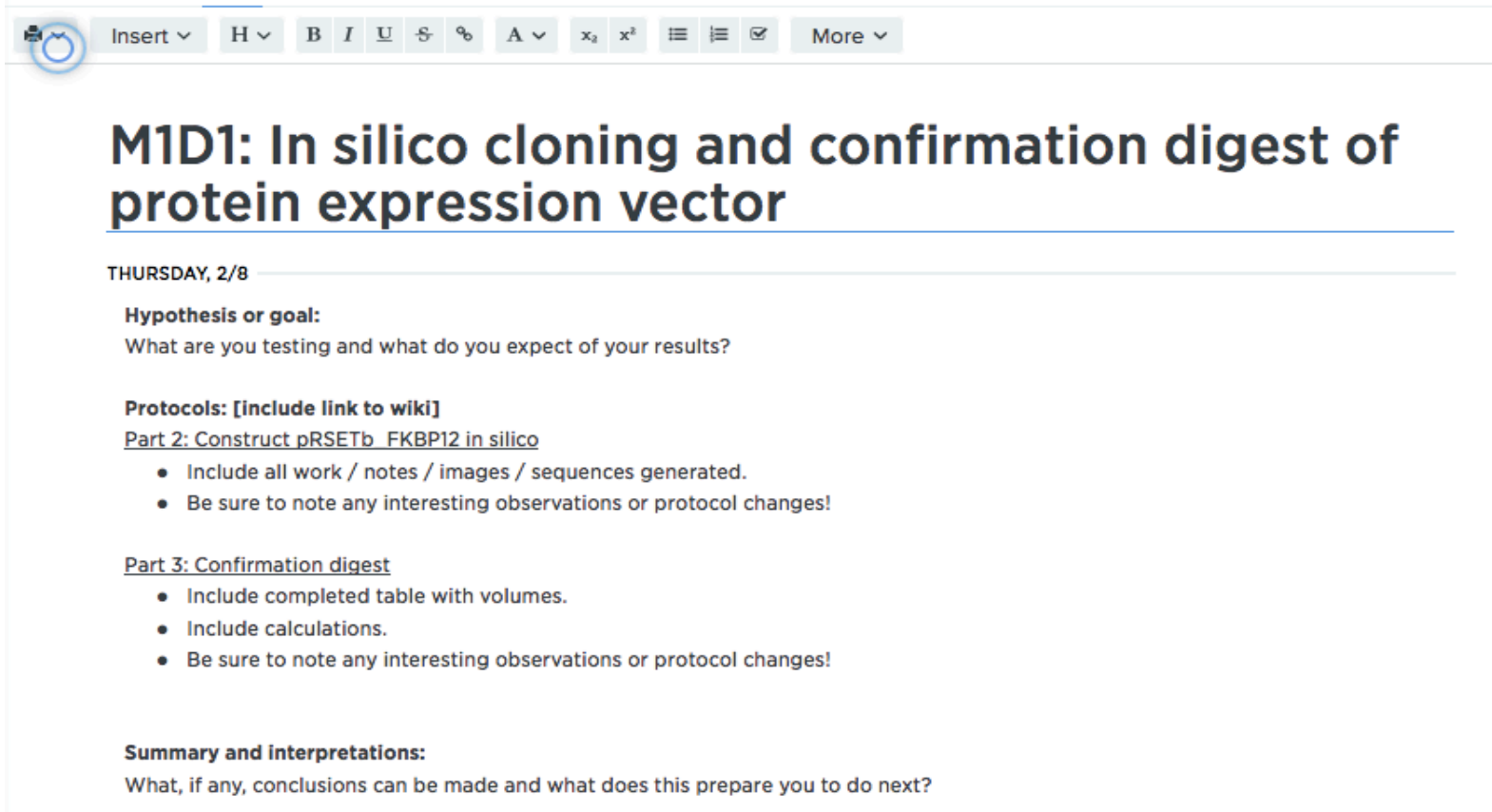
Laboratory notebook entry component:

	Points:		
	Complete	Partial	Incomplete
Date of experiment (include Module#/Day#) and Title for experiment	1	0.5	0
Hypothesis or goal / purpose	1	0.5	0
Protocols (link to appropriate wiki sections)	1	0.5	0
Notes on protocol changes / clarifications	1	0.5	0
Observations	2	1	0
*Visual details			
*Qualitative information			
*Raw data			
Data analysis	3	1.5	0
*Calculations			
*Graphs and Tables			
Summary and interpretation of data	3	1.5	0
*What did you learn?			
*How does this information fit into the larger scope of the project?			
Information is clear	2	1	0
All days represented	1	0.5	0
OVERALL /15			

Due 10pm after each module, as posted on wiki

[http://engineerbiology.org/wiki/20.109\(F18\):_Assignments](http://engineerbiology.org/wiki/20.109(F18):_Assignments)

How should you format your notebook?



The screenshot shows a digital notebook interface with a toolbar at the top containing icons for insert, heading, bold, italic, underline, strikethrough, link, text color, background color, list, indent, and a 'More' dropdown. The main content area displays a notebook page with a title, a date, and several sections of text and lists.

M1D1: In silico cloning and confirmation digest of protein expression vector

THURSDAY, 2/8

Hypothesis or goal:
What are you testing and what do you expect of your results?

Protocols: [include link to wiki]
Part 2: Construct pRSETb_FKBP12 in silico

- Include all work / notes / images / sequences generated.
- Be sure to note any interesting observations or protocol changes!

Part 3: Confirmation digest

- Include completed table with volumes.
- Include calculations.
- Be sure to note any interesting observations or protocol changes!

Summary and interpretations:
What, if any, conclusions can be made and what does this prepare you to do next?

How should you organize your notebook?

- Entitle your project “20.109(F18)_YourName”
 - Make each module a new folder
 - Make each day a new entry within module folder
- Share the project with Josephine and Corban
 - Right-click and choose ‘settings’
 - Add collaborators by email address

Today in lab:

1. 3 teams into tissue culture room to split cells (Red, Orange, Green)
 2. 2 teams start preparing CometChip (Yellow and Blue)
 3. Make sure to keep notes in Benchling!
- Watch Engleward lab JOVE video during downtime
(<https://www.jove.com/video/50607/cometchip-high-throughput-96-well-platform-for-measuring-dna-damage>)
 - M1D2HW: Create a template for your benchling notebook and make a M1D2 entry from it.