# M1D4: Treat cells and perform high-throughput genome damage assay

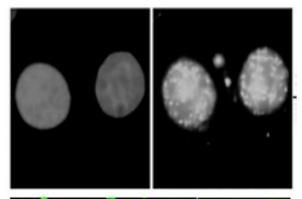
1. Quiz

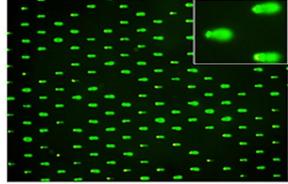
- 2. Prelab
  - 1. Review H2AX analysis

3. Perform CometChip experiment



### Mod 1 Overview





#### 1. Use repair foci experiment to measure DNA breaks

• Examine effect of  $H_2O_2$  +/- As on double strand DNA breaks by measuring  $\gamma$ H2AX foci formation

# 2. Use high-throughput genome damage assay to measure DNA damage

• Measure effects of  $H_2O_2$  +/- As on DNA damage by measuring DNA migration in agarose matrix

# Notes on fluorescence imaging and analysis

#### • Imaging set up:

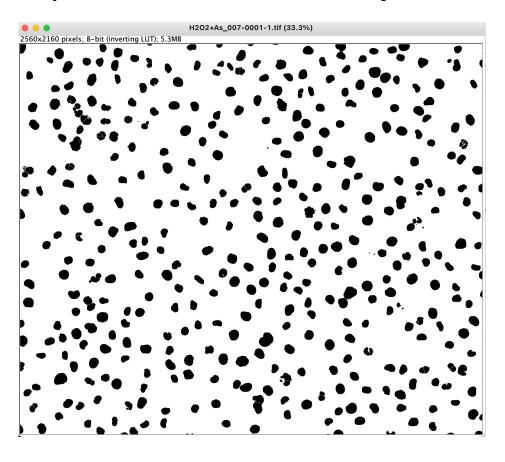
- Experimental condition (presumably the most damage/H2AX foci)
- Set exposure time for each channel with this condition (we did 50ms)
  - Prevents saturation in the image (i.e. "signal blow out") and allows for cleaner analysis
- Images from all 4 conditions are collected under these parameters to ensure comparability in analysis

#### • Image Presentation:

- Images kept well below saturation threshold can be difficult to visualize by eye
- The signal intensity can be adjusted manually to provide more contrast
  - Be sure to keep adjustment parameters as the same range so that your images can still be compared

### Fun with foci maxima...

My nuclei were masked just fine...

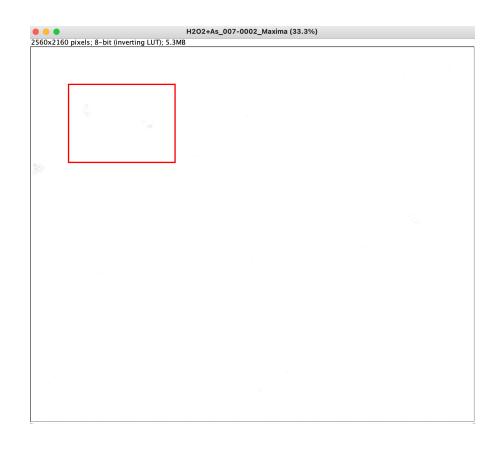


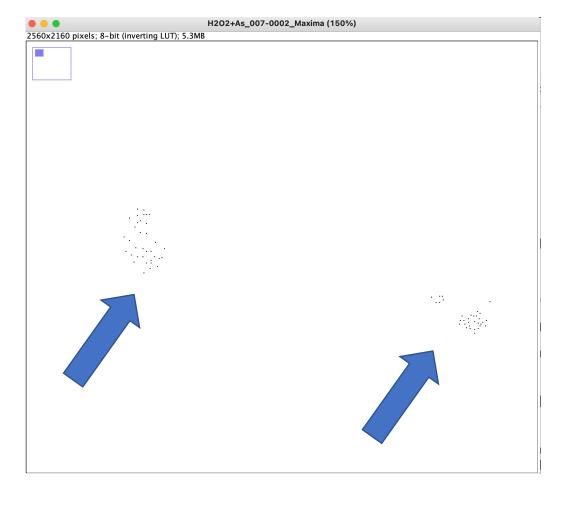
• The nuclei outlines seem accurate...



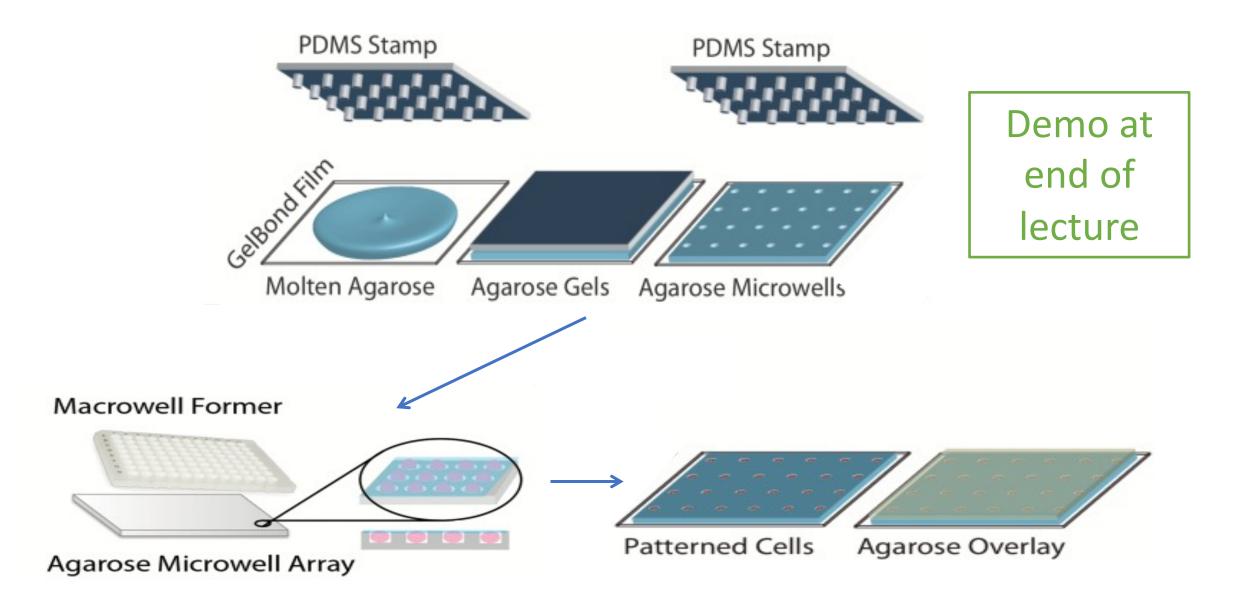
### Fun with foci maxima...

• But the foci analysis is giving me a white screen.

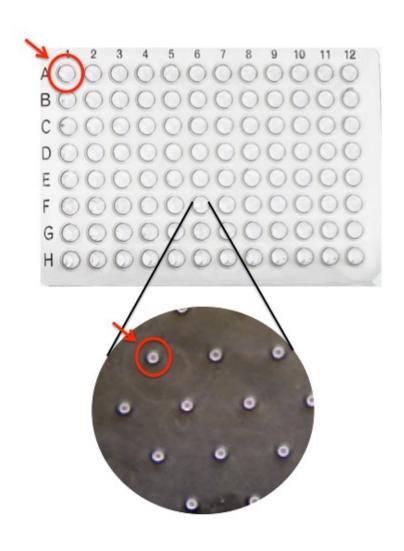




### Overview of the CometChip assay: pouring and loading cells

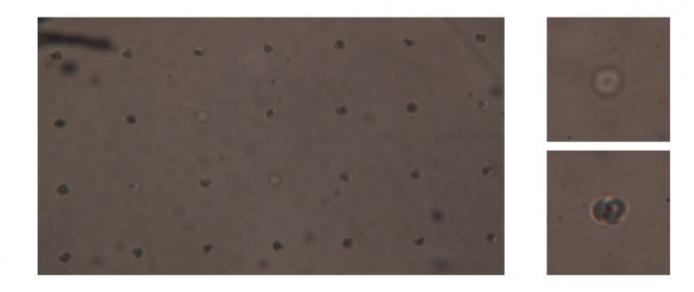


# Loading cells into CometChip wells



 How many cells are in a microwell?

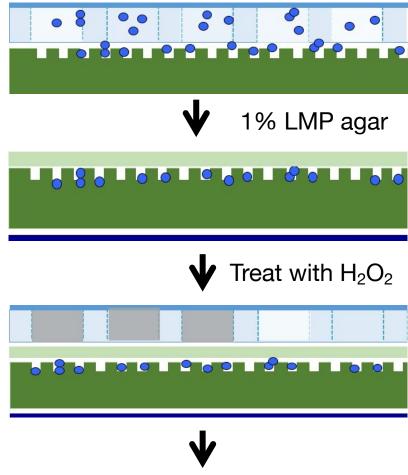
 How many cells are in a macrowell?



## Overview of the CometChip assay: treating cells

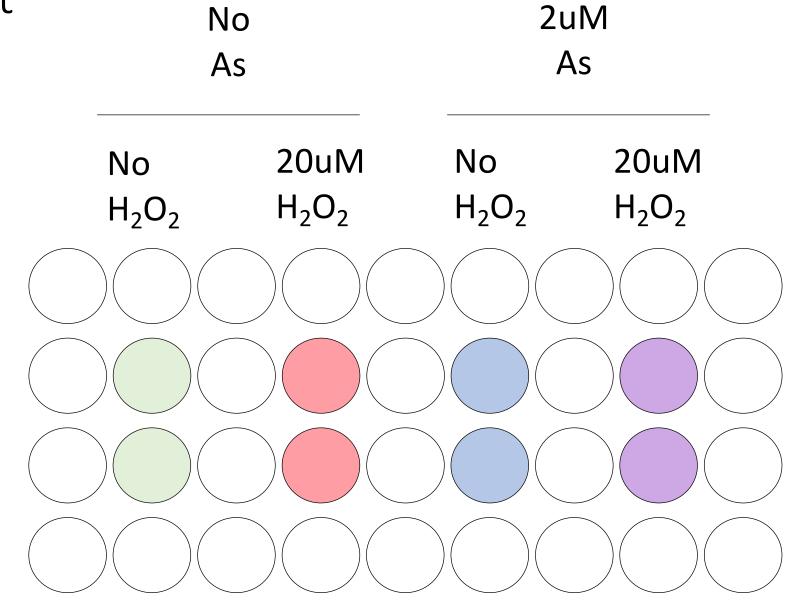


Treat with As for 24hrs

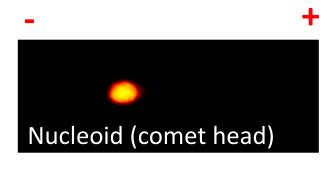


Omin recovery
Place directly in lysis
buffer

# Macrowell layout

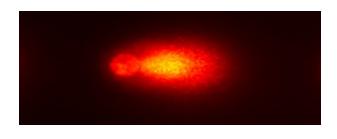


# Output of the alkaline CometChip assay



### **No Damage**

- Supercoiled nucleoid
- Little or no migration



### **High Damage**

- SSBs, abasic sites, alkali labile sites, sites of incomplete excision repair
- forms a "comet tail"

- \* Nuclear DNA normally supercoiled
  - \* DNA breaks and fragmentation releases tension
    - \* Unwound DNA will migrate in response to electrical current to create comet

# For Today

- Perform CometChip experiment
- With any extra time, continue H2AX analysis
- At 4:30pm, Demo of CometChip Electrophoresis

### For M1D5

#### Group

Revise methods and add in M1D3 (I'm uploading Noreen's comments to Stellar)

#### **Individual**

- Read paper linked on M1D5 and prepare for group discussion
- Write summary for BE Comm Lab visit