

NET formation extended

PMA Pathway: PMA -> PKC -> NADPH oxidase -> ROS -> chromatin decondensation -> NETs release

Ionomycin Pathway: Ionomycin -> calcium level increase -> PAD4 -> histone citrullination -> NETs release

Raptinal Pathway: Raptinal -> Apoptosis induction -> Mitochondrial Outer Membrane Permeabilization (MOMP) -> caspase activation (caspase 9/3) -> gasdermin E cleavage -> GSDME-N -> membrane pores -> lytic NET release

Nigericin Pathway: Nigericin -> Potassium ions efflux with Hydrogen ions influx -> NLRP3 inflammasome -> caspase-1 directed gasdermin D cleavage -> GSDMD-N -> pore formation -> DNA release

The two pathways above are not directly associated with NETosis but do promote NET structure.

Val-boropro Pathway: pyroptosis -> DPP8/9 inhibition -> CARD8 (human) or NLRP1 (mouse) -> Caspase-1 (human), Caspase-11 (mouse) -> lytic cell death (slightly slower)

Hypothesis: Investigating different compounds with apoptosis and pyroptosis on NET formation.

Methodology:

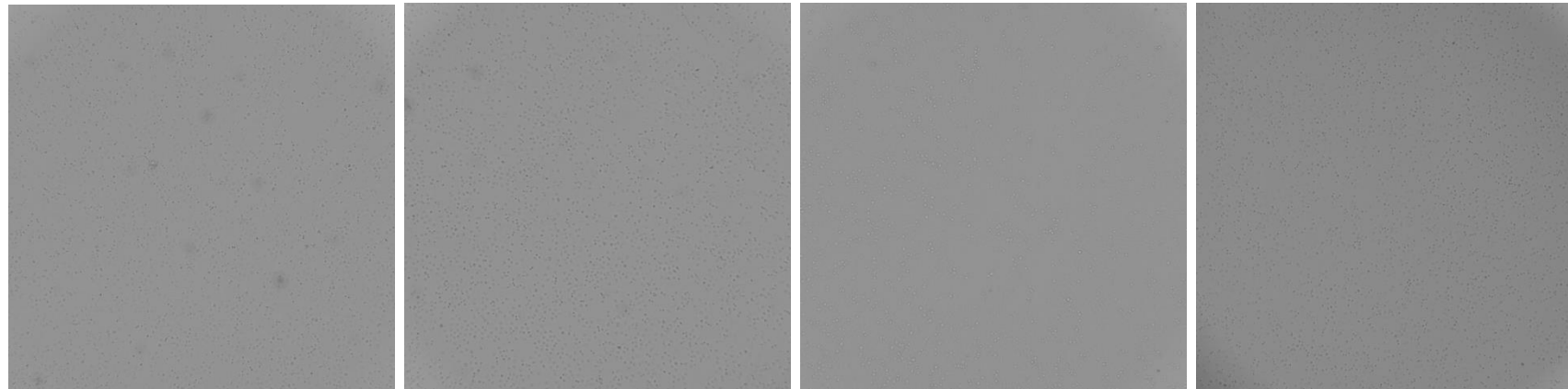
Extract the neutrophils from bone marrow

Add the compounds to the cell

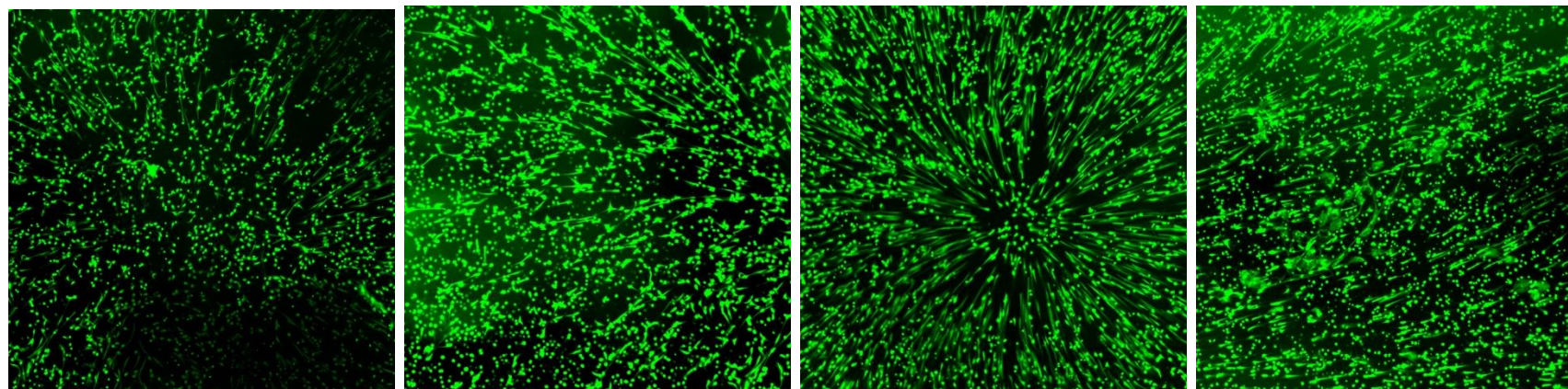
Use the device to observe the NETs

10x Microscope

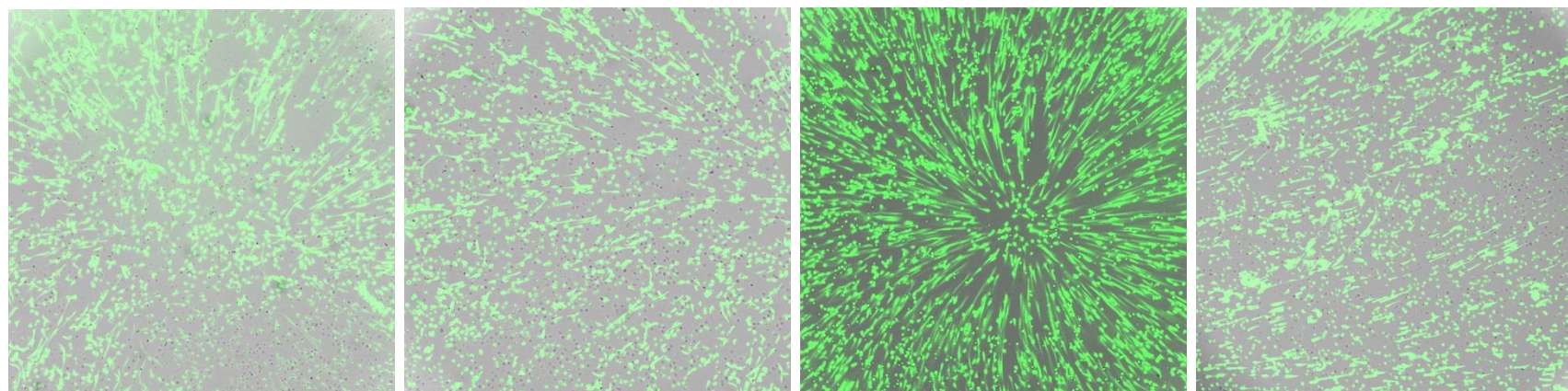
BF



GFP



Merged

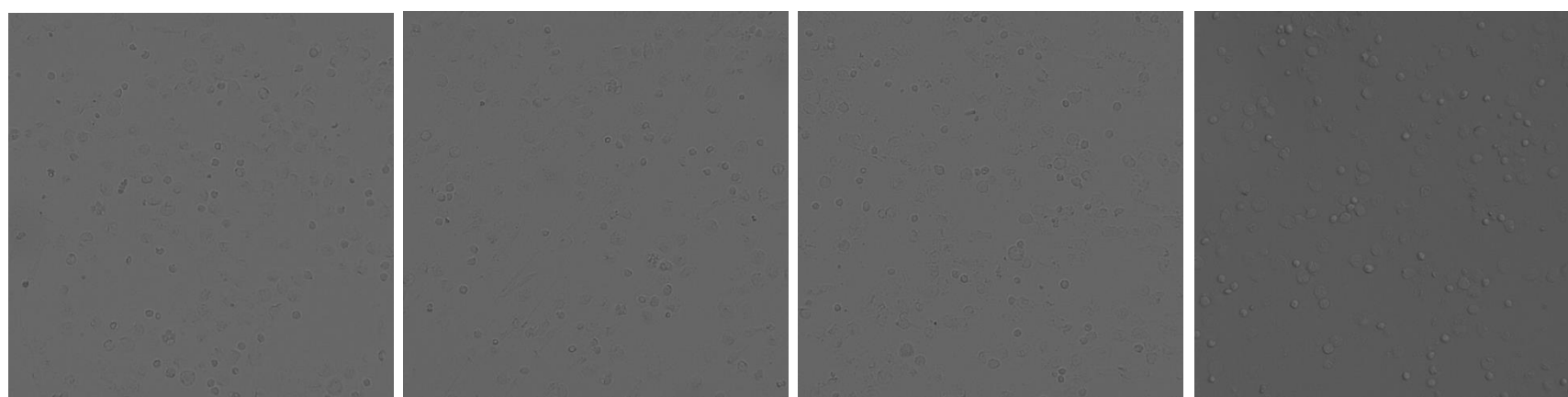


PMA

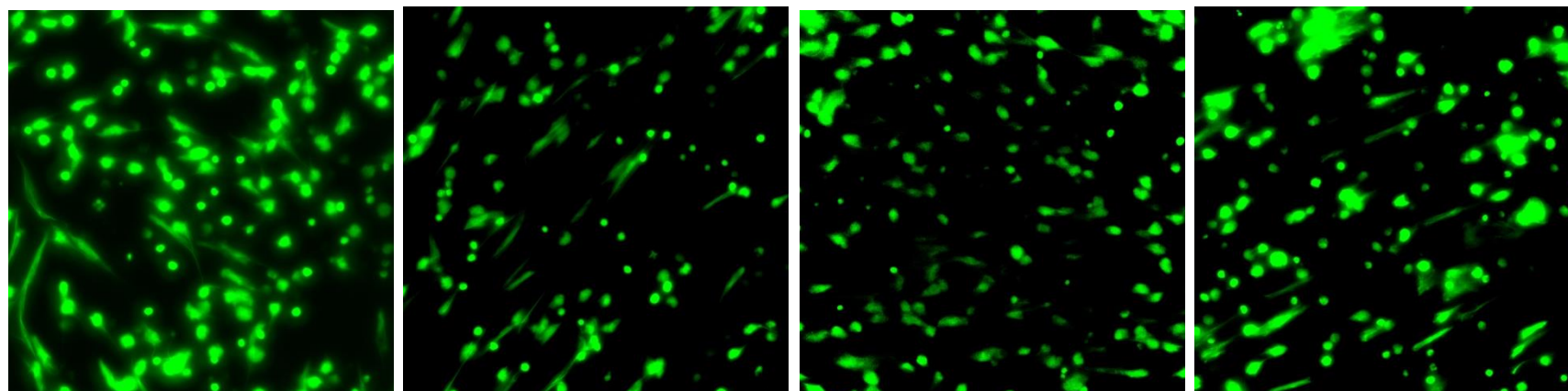
PMA

Ionomycin

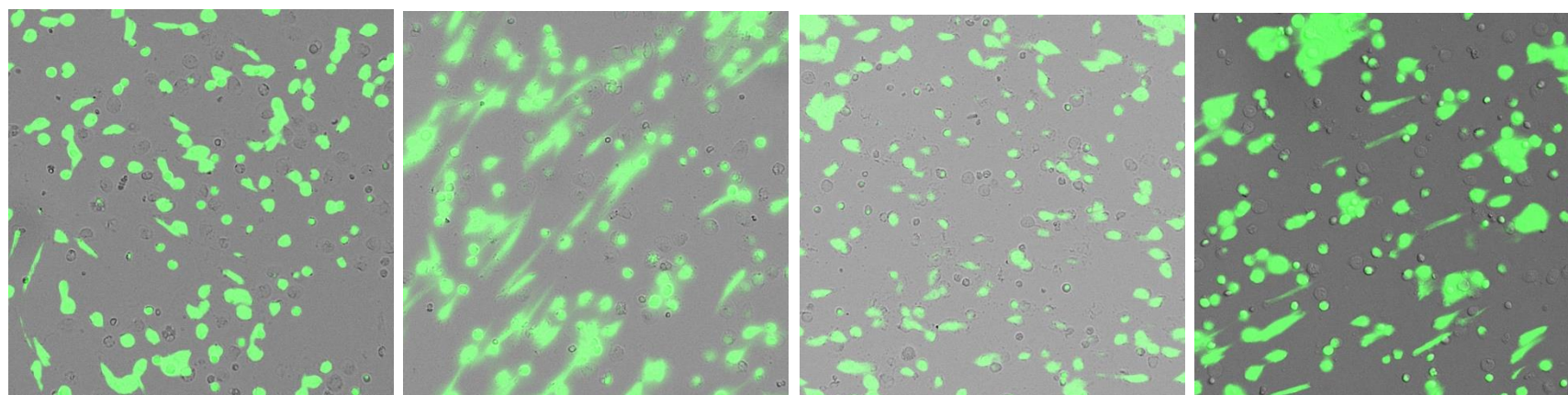
Ionomycin



BF



GFP



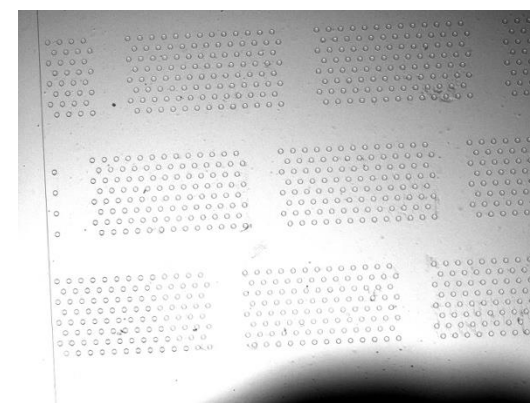
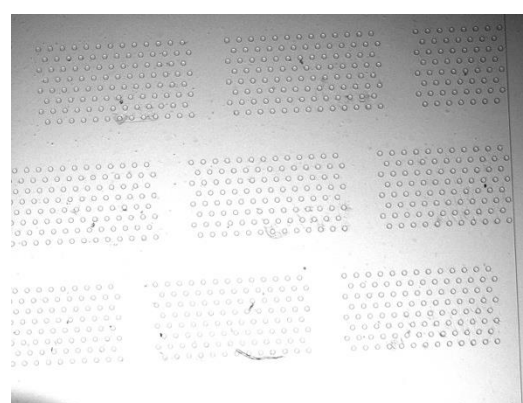
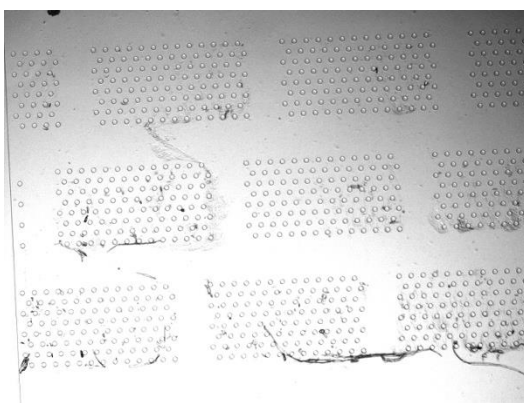
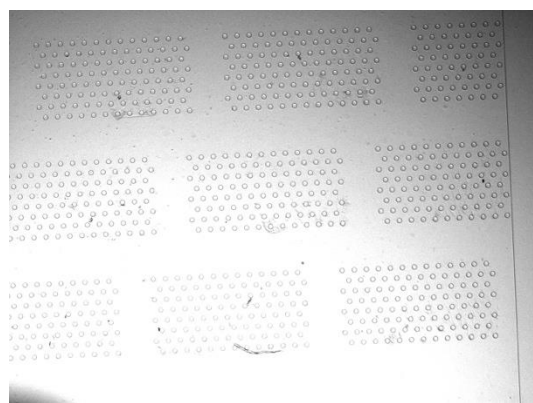
Merged

PMA

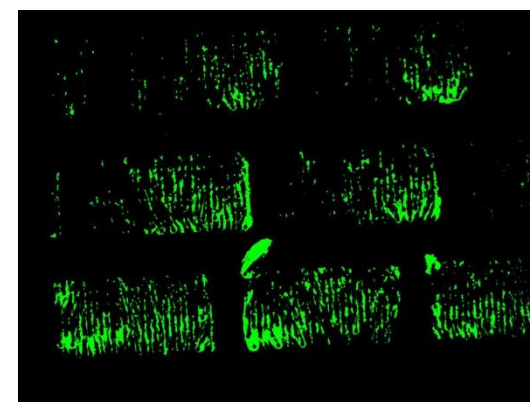
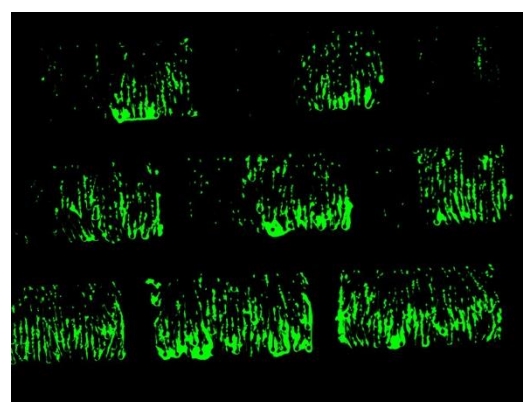
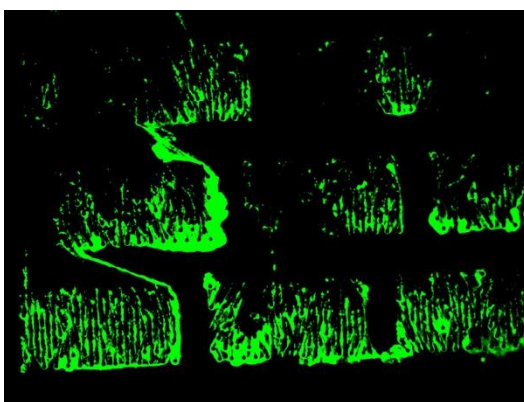
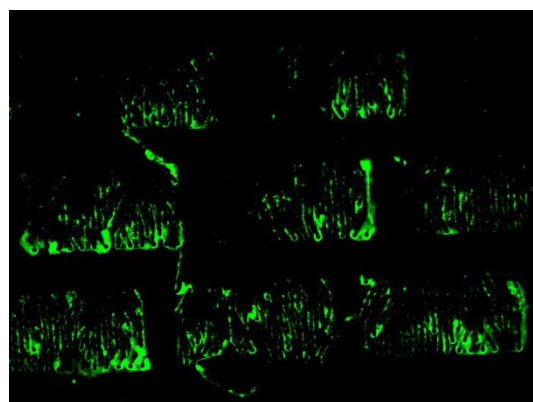
PMA

Ionomycin

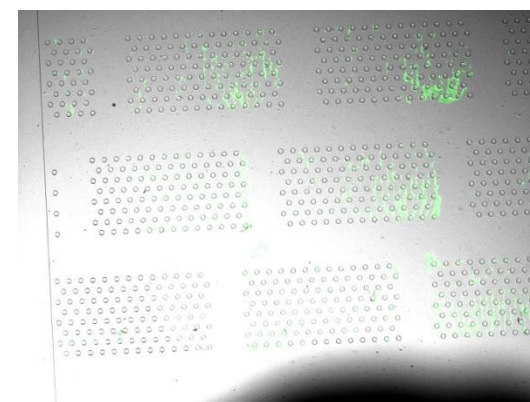
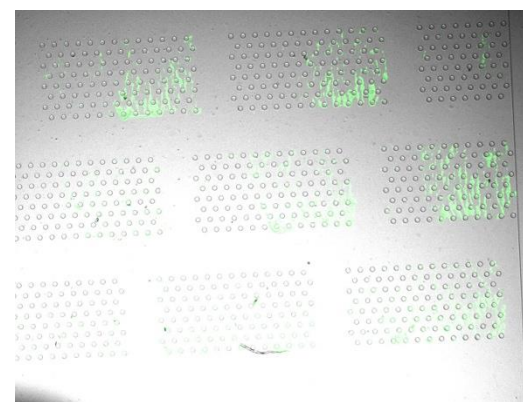
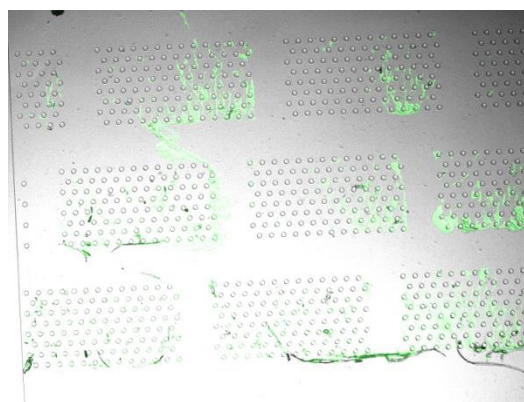
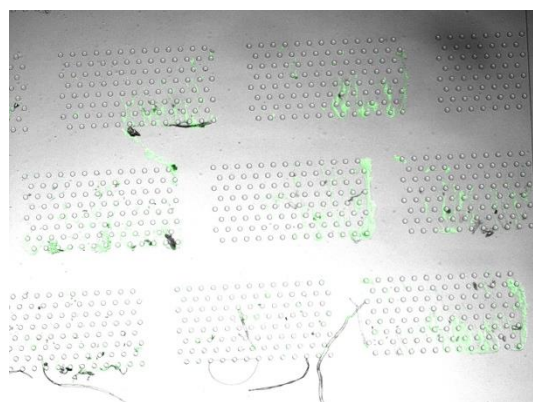
Ionomycin



10x Device
BF



GFP



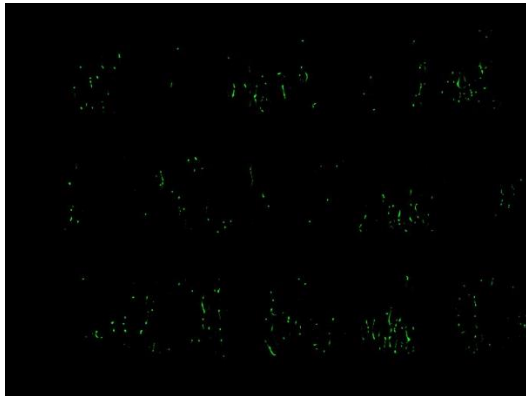
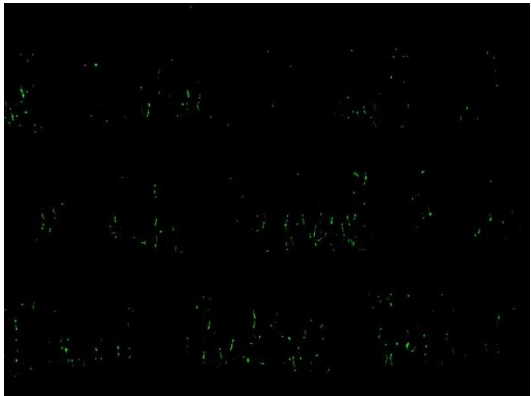
Merged

PMA

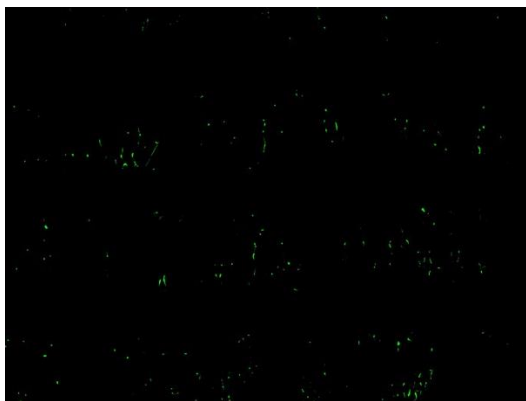
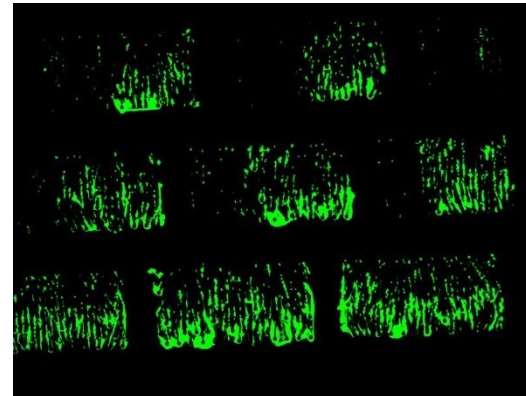
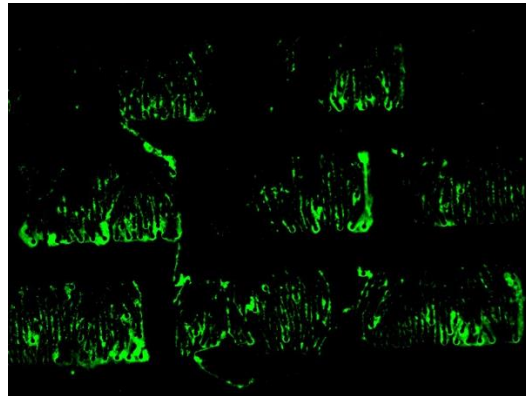
PMA

Ionomycin

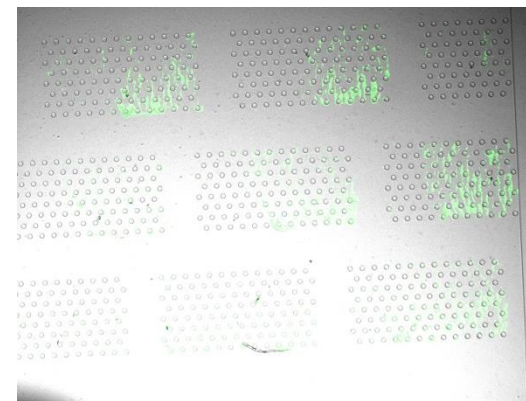
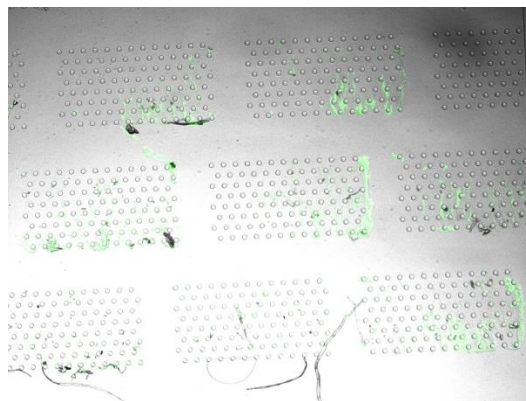
Ionomycin



GFP



GFP



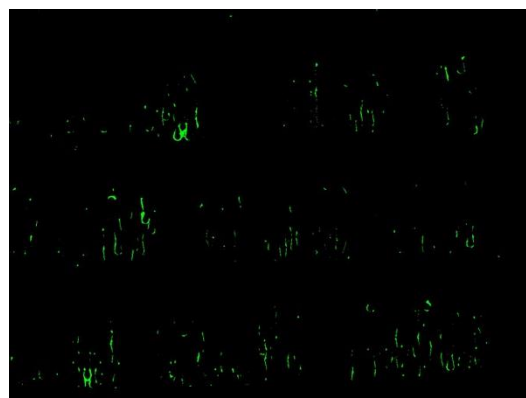
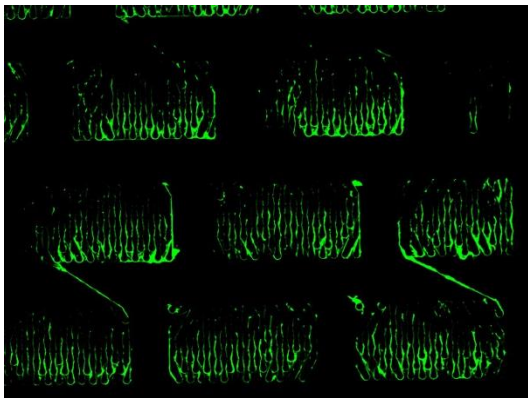
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PMA

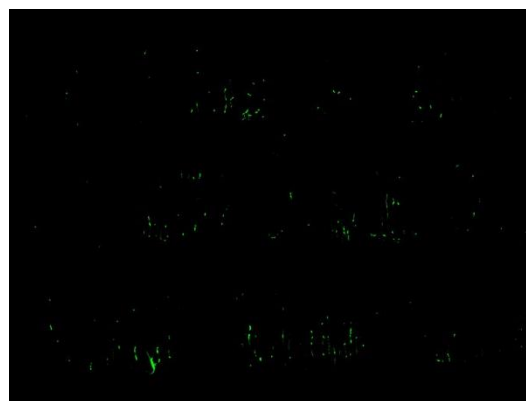
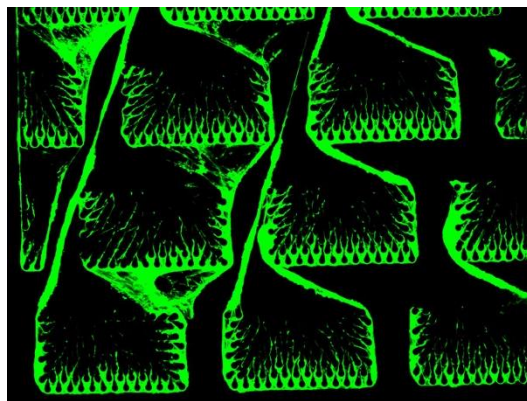
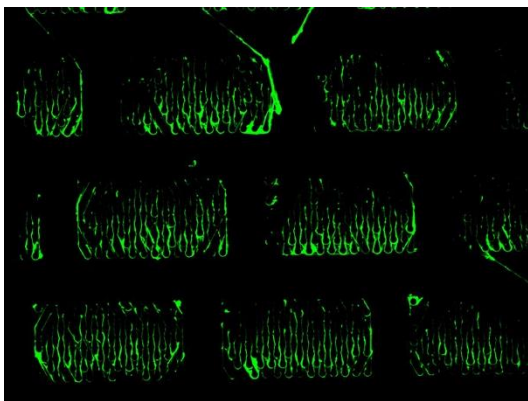
Ionomycin

10x Device



10x Device

GFP 1

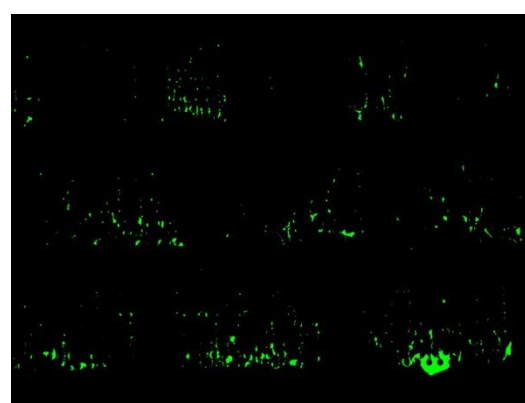
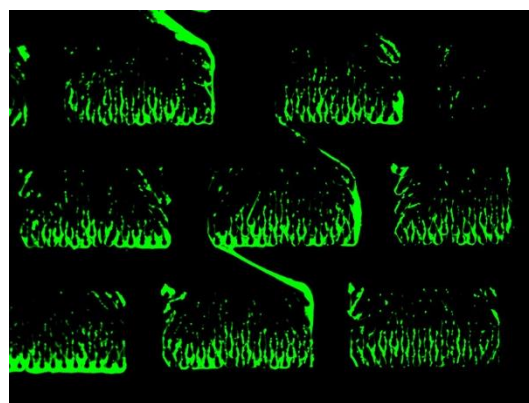
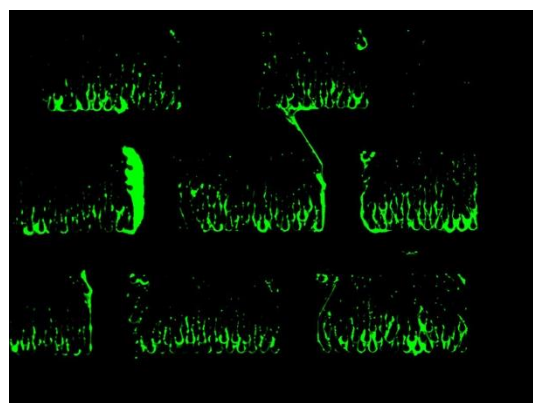


GFP 2

PMA

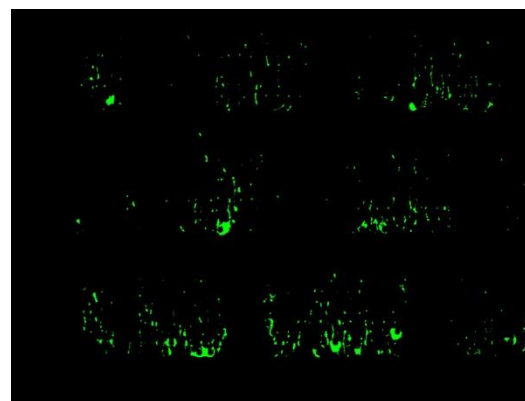
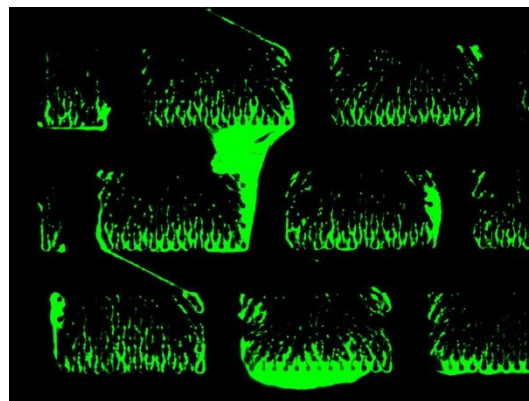
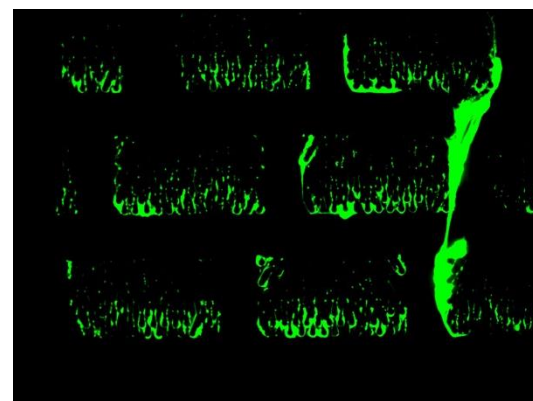
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10x Device

GFP 1

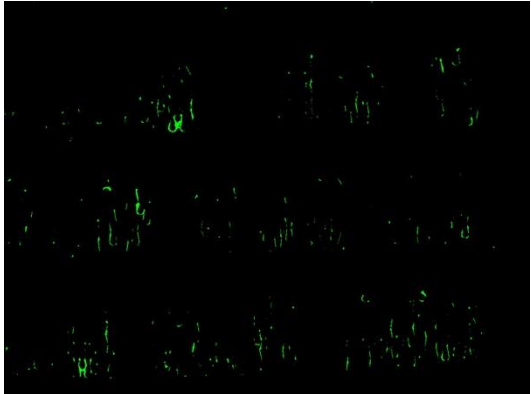


GFP 2

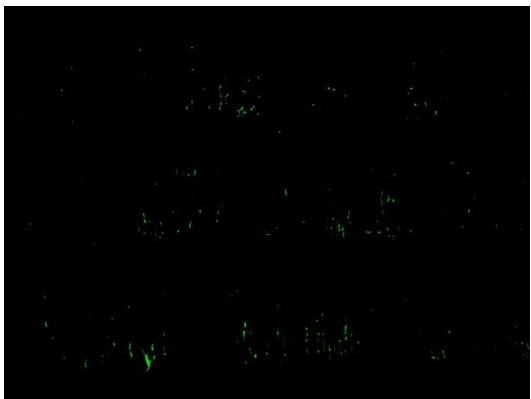
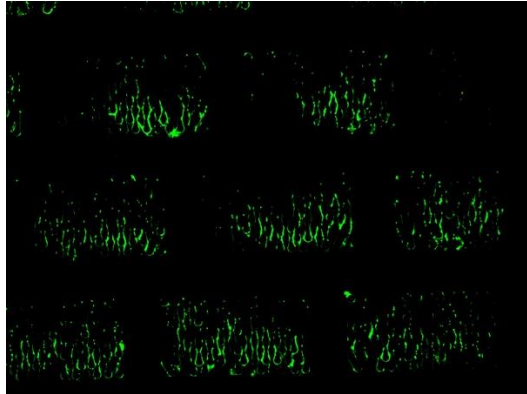
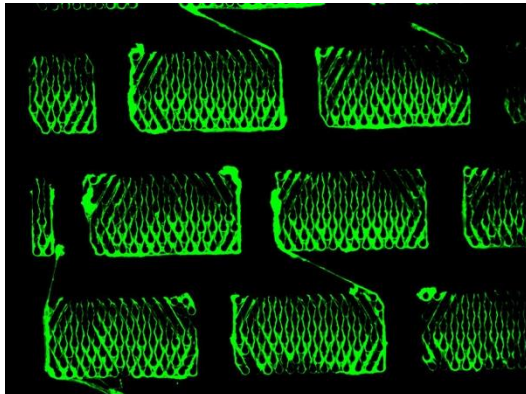
PMA

Ionomycin

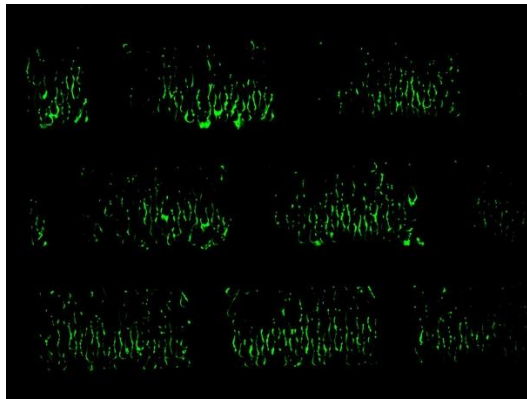
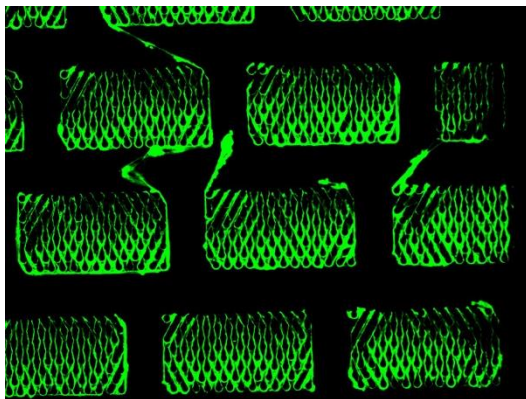
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GFP



GFP



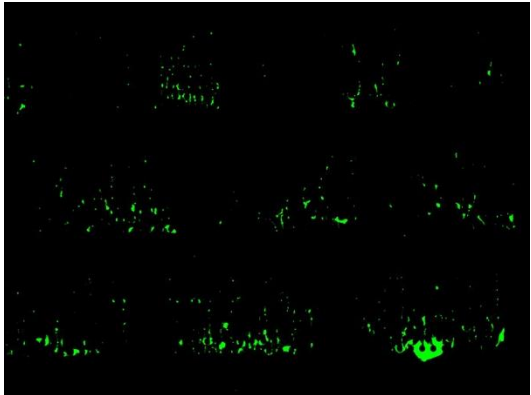
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Raptinal

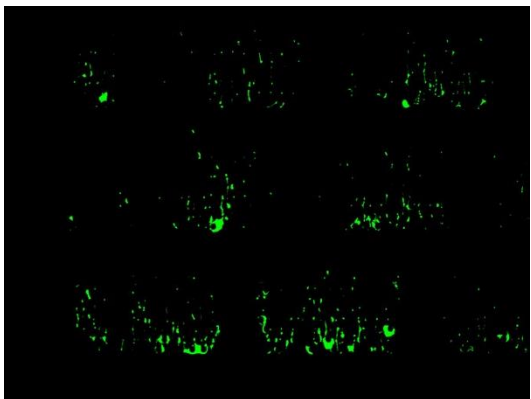
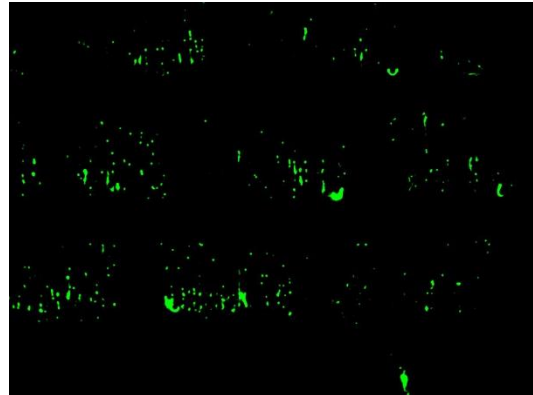
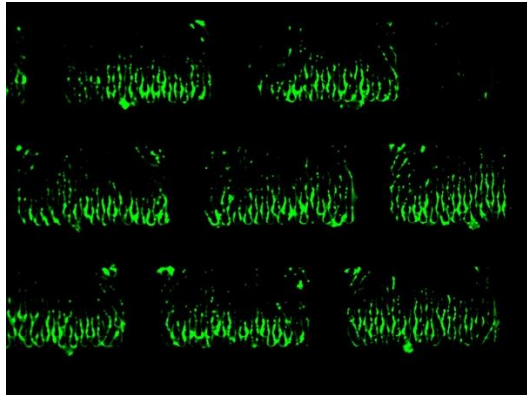
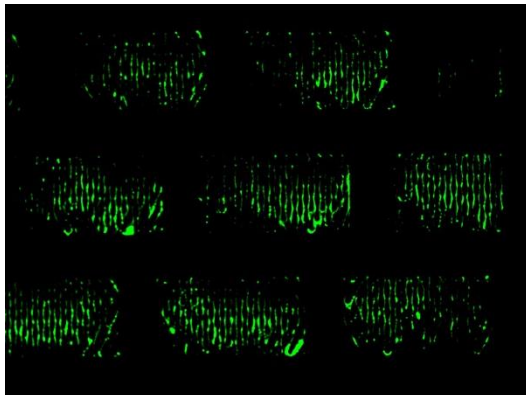
Nigericin

Val-boropro

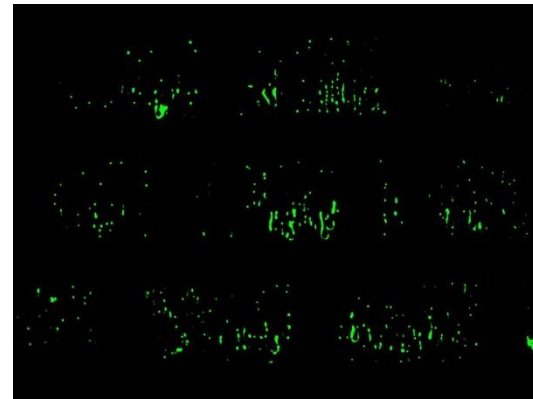
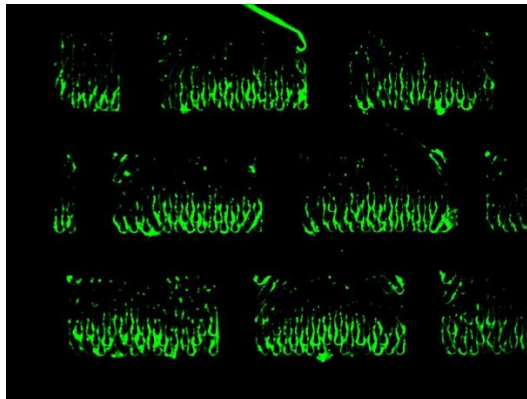
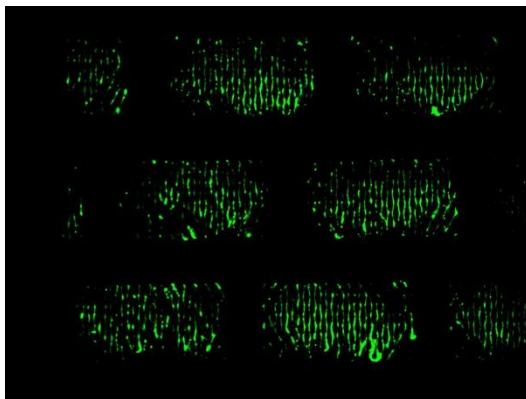
Conc.: 5 micromoles



GFP



GFP

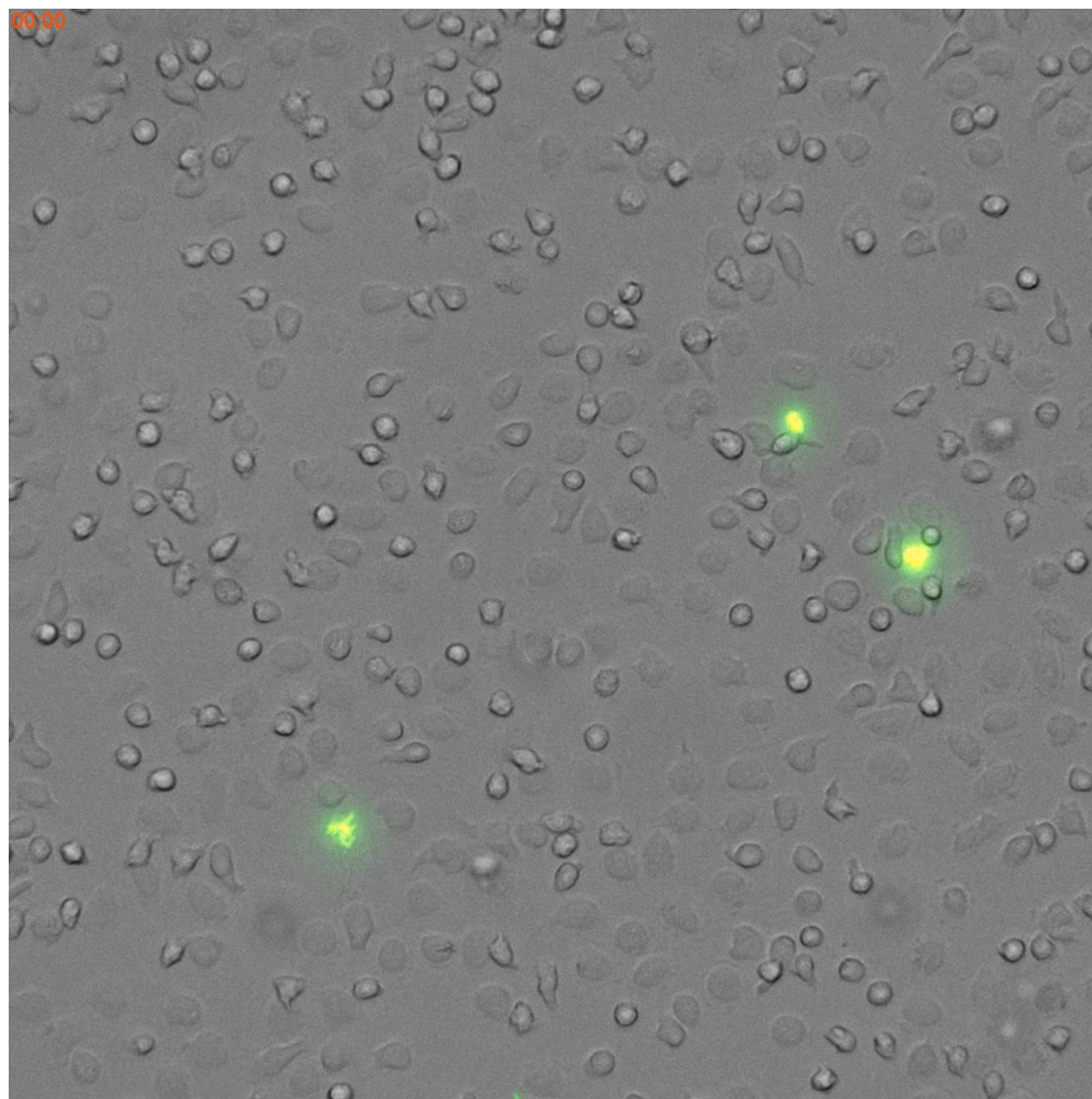


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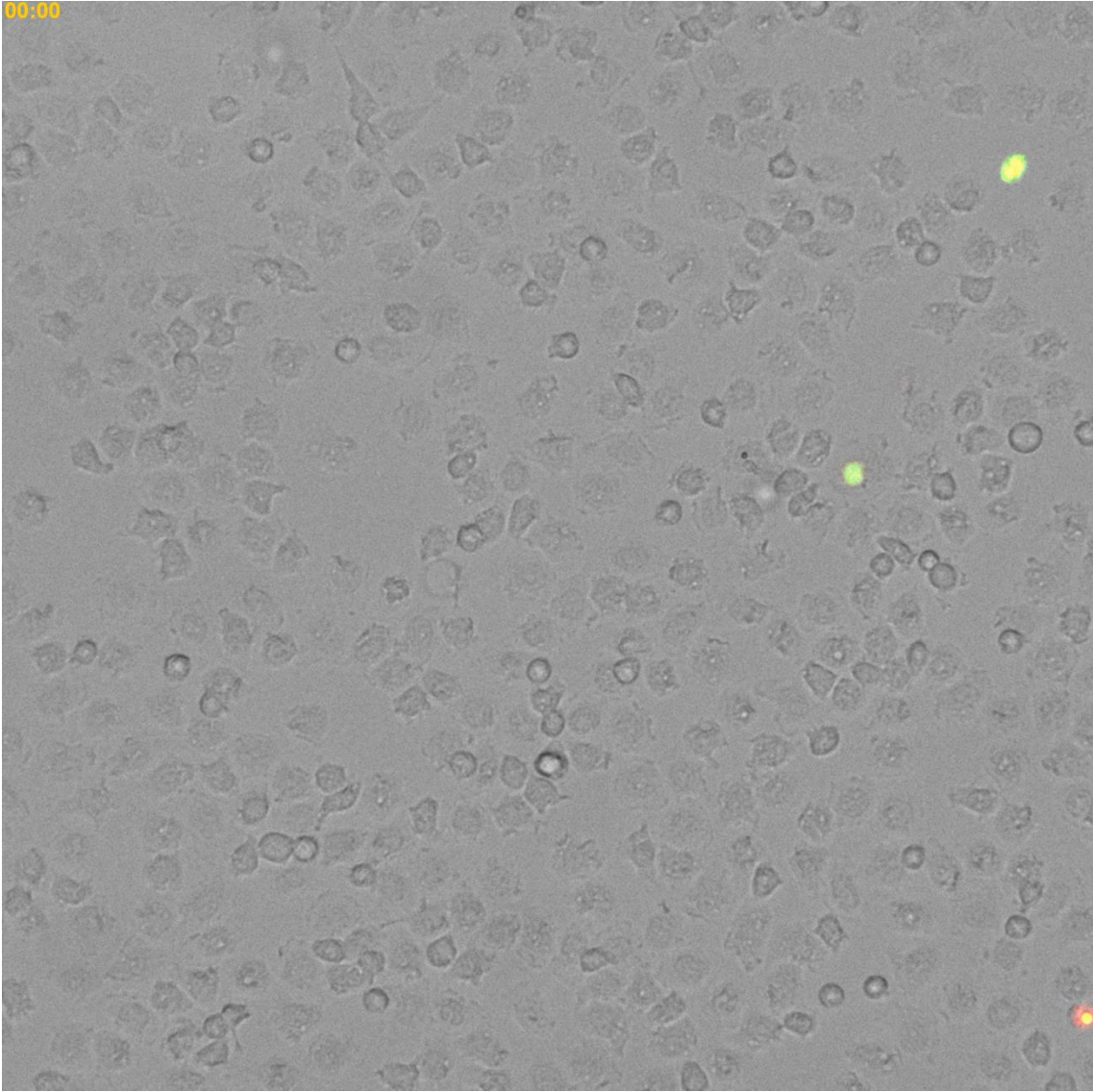
Raptinal

Nigericin 2x conc

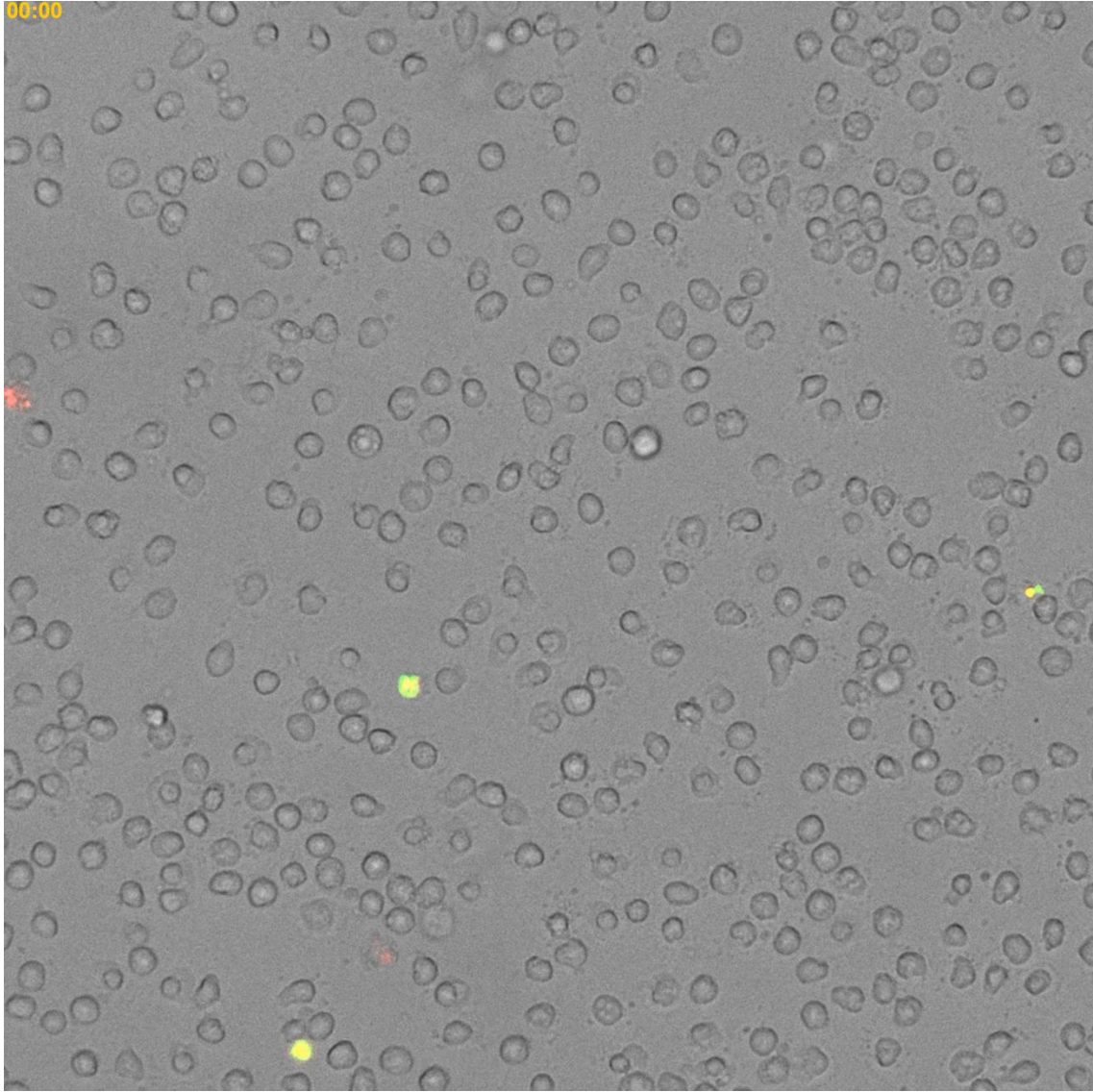
Val-boropro 10x conc



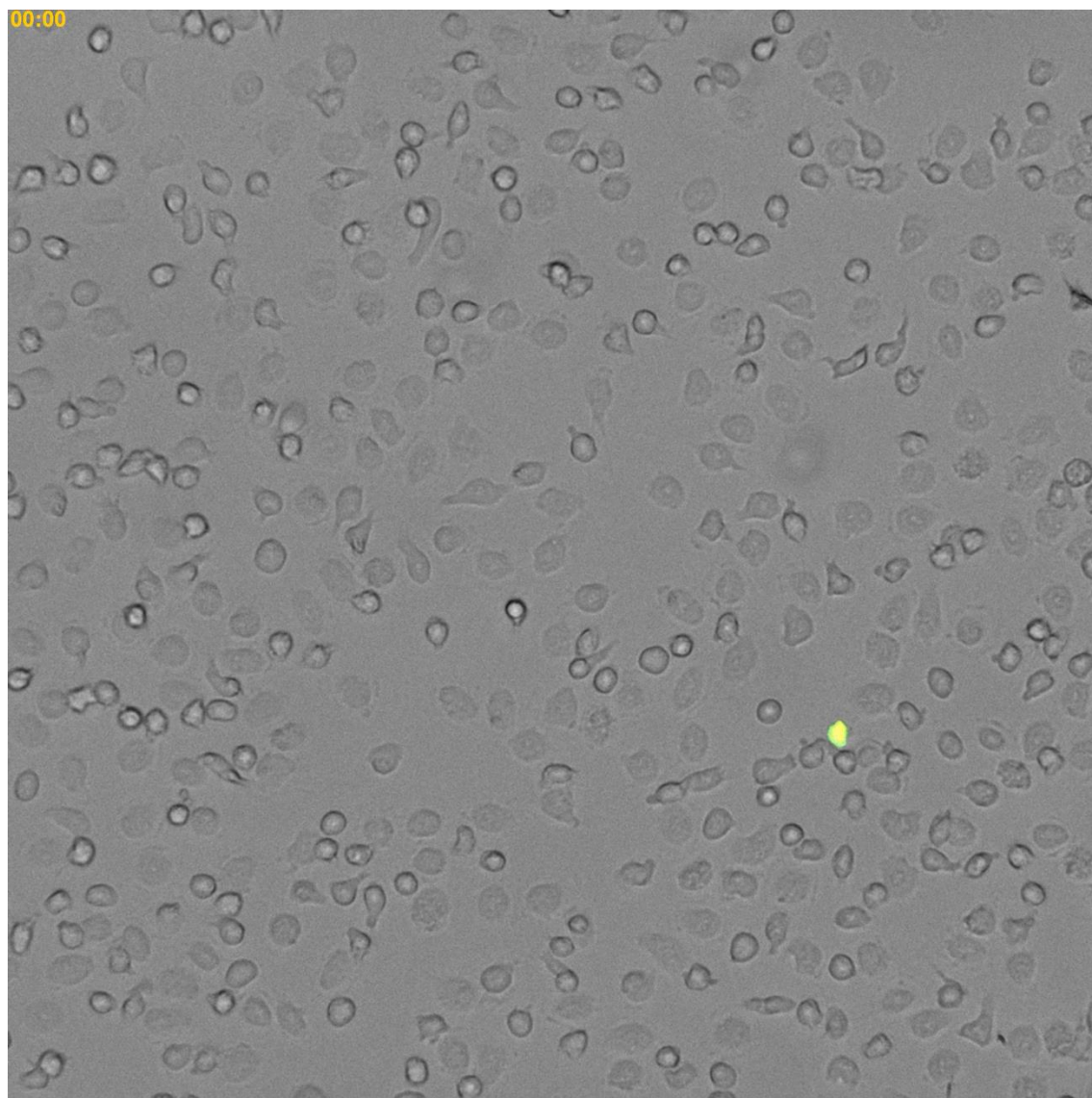
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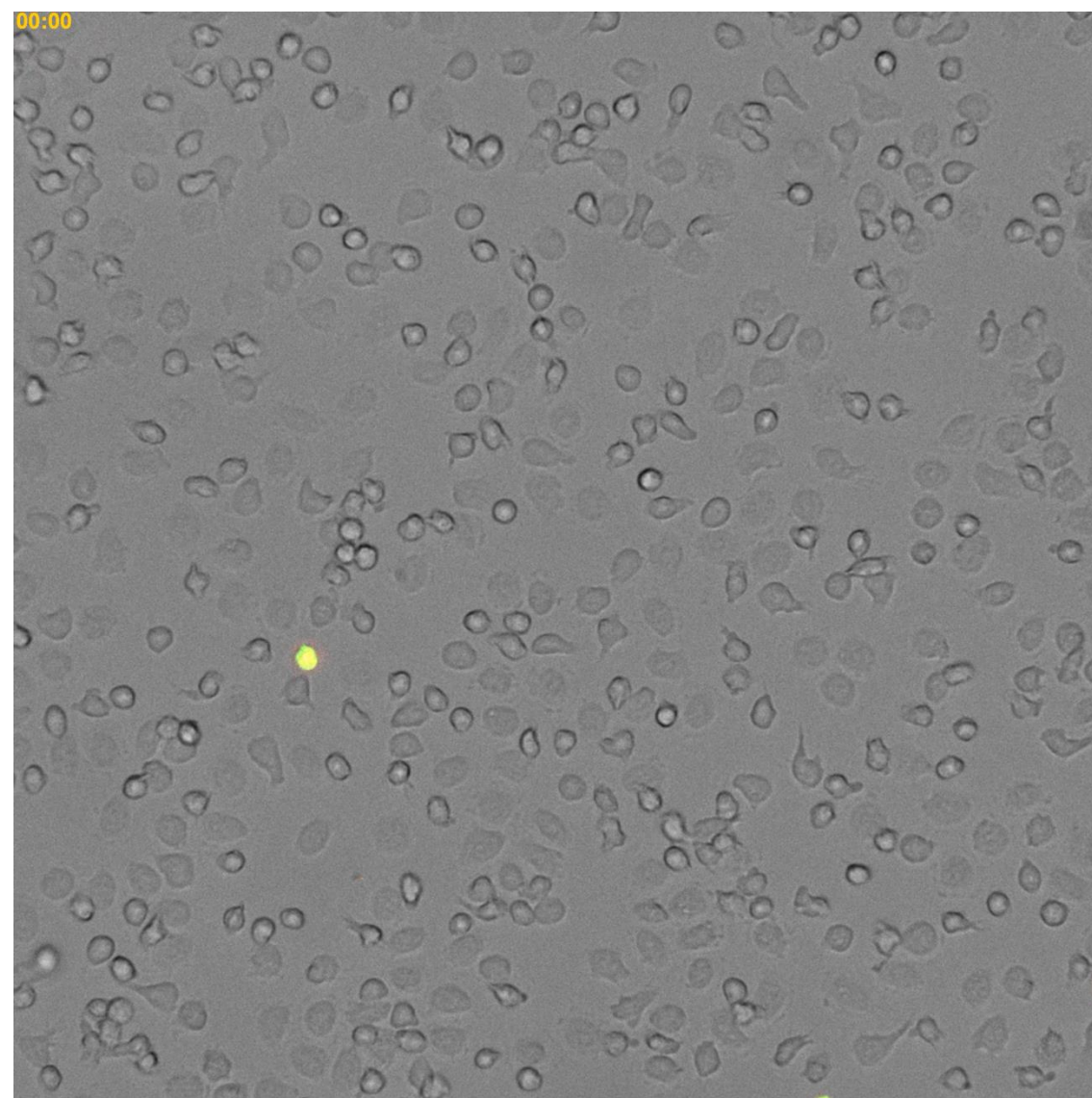
PMA



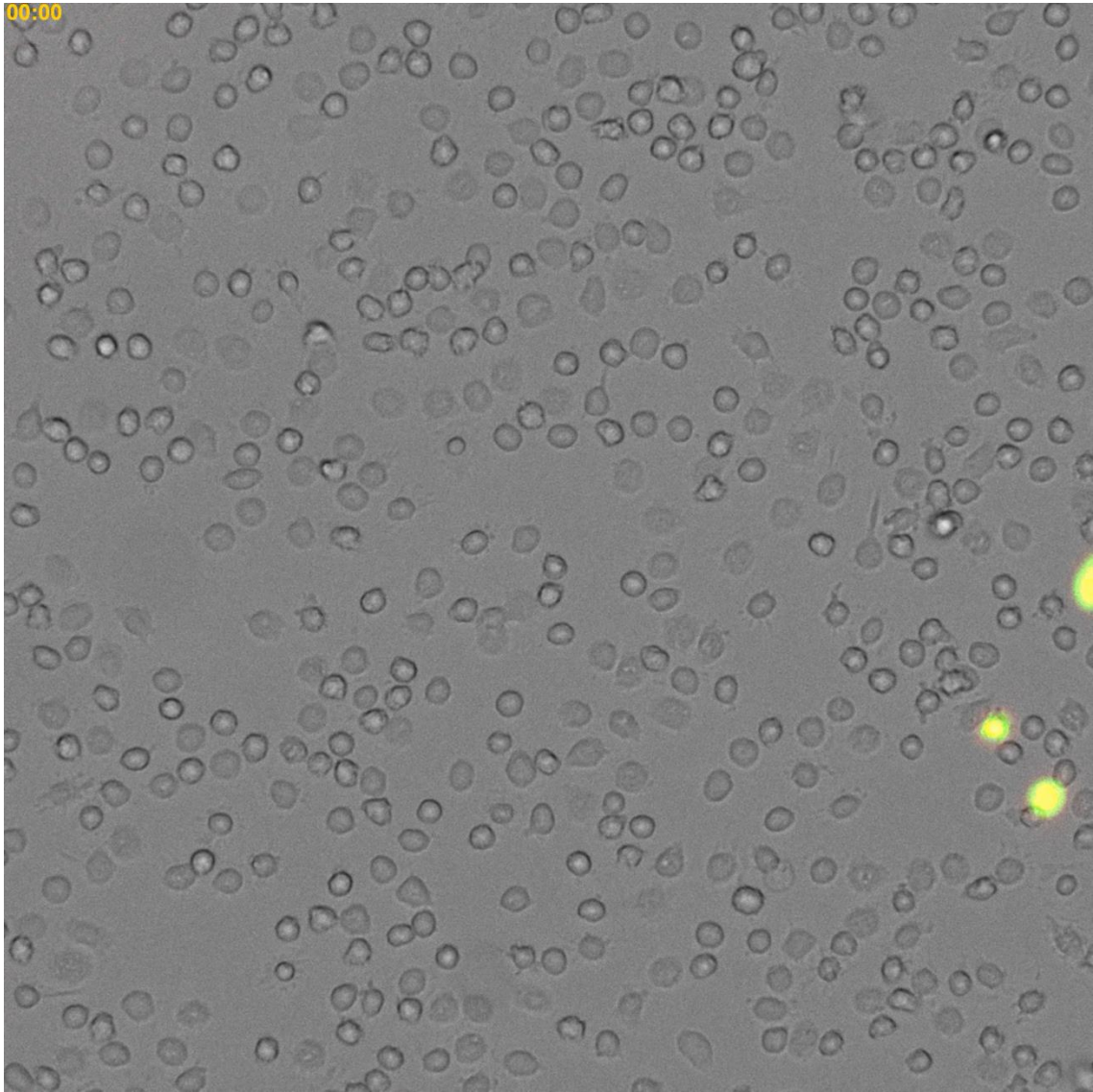
Ionomycin



Raptinal



Val-boropro



Death rate at 4 hours:

Blank: 30%

PMA: 93%

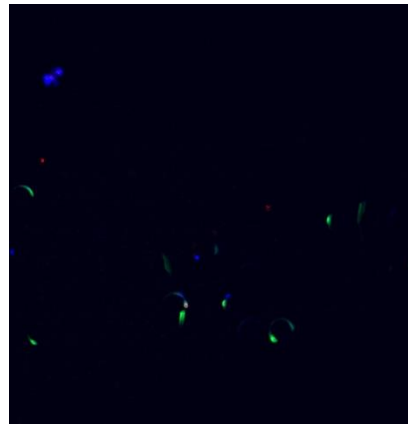
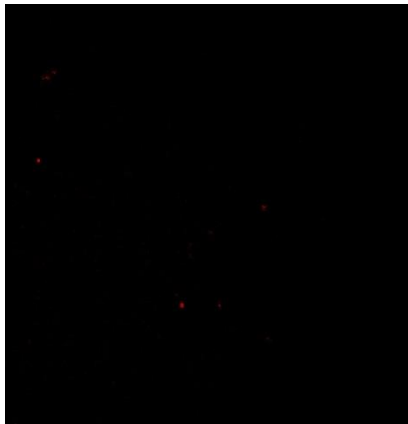
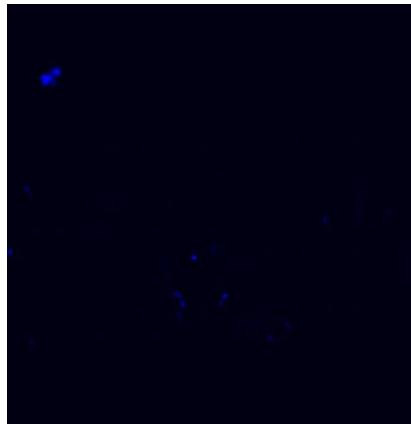
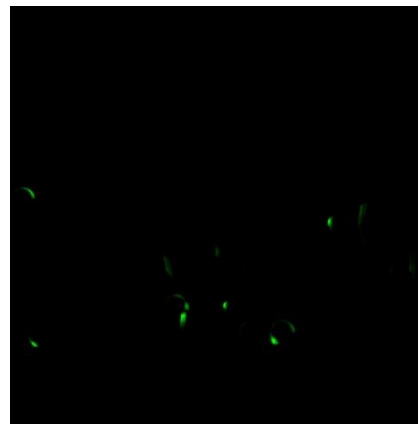
Ionomycin: 95%

Raptinal: 70%

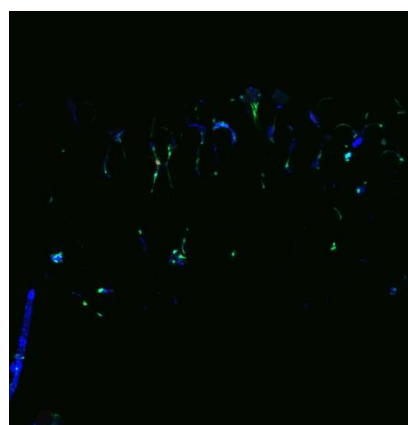
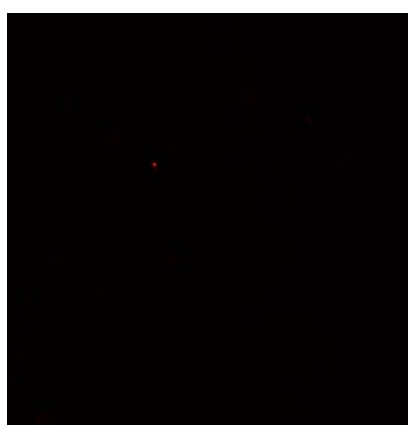
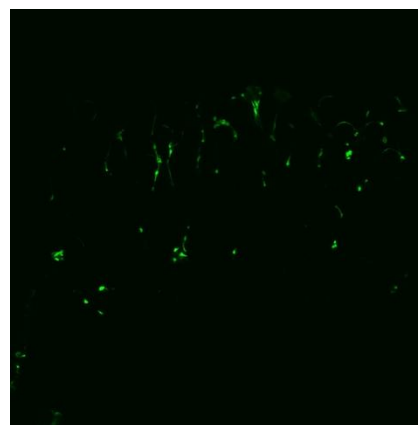
Val-boropro: 25%

Nigericin: 85%

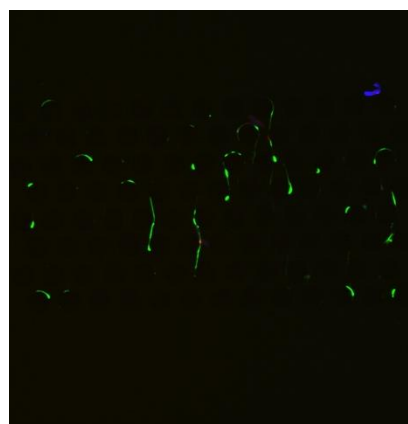
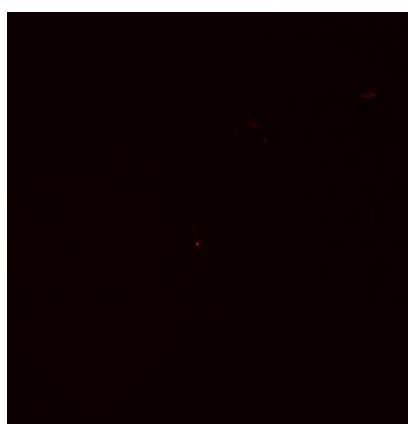
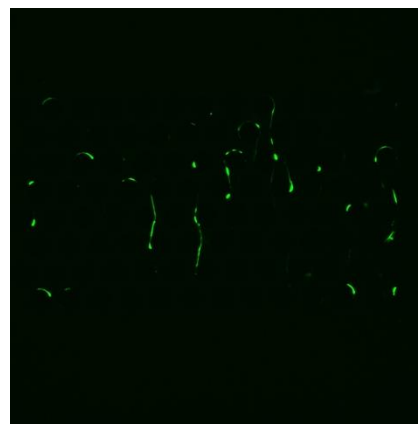
Nigericin



Blank 10x



Nigericin 10x



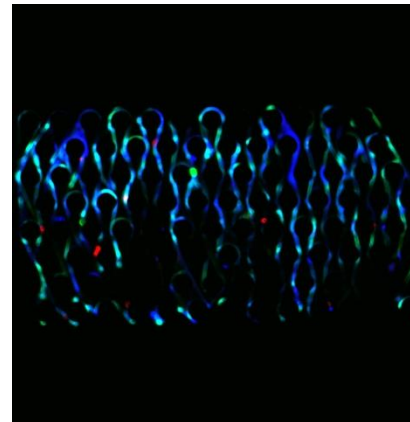
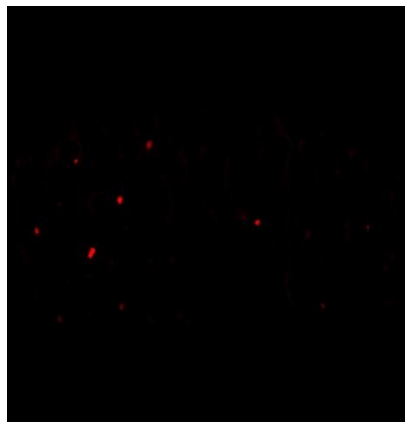
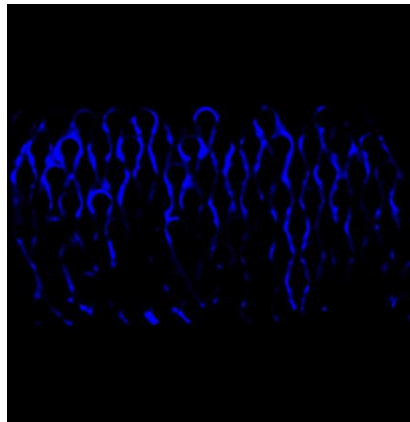
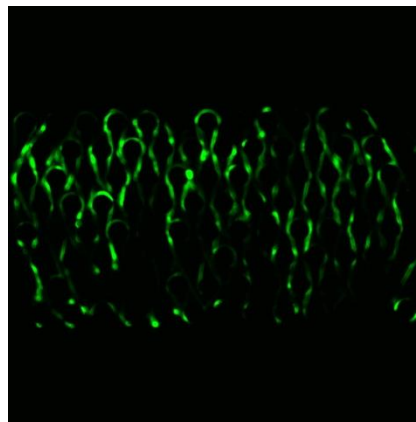
Val-boropro 10x

G

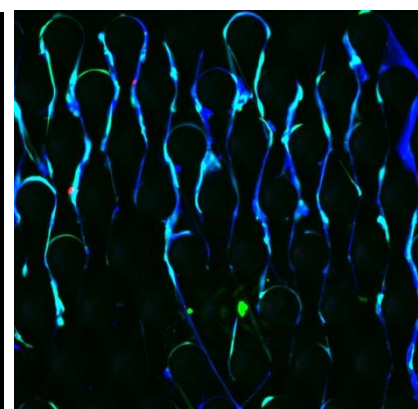
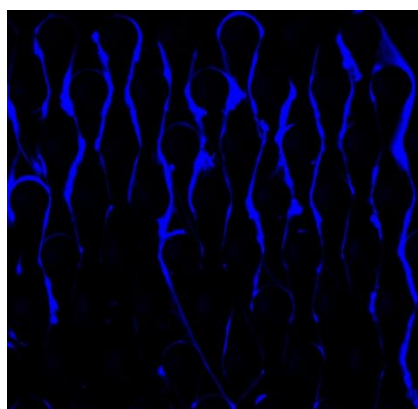
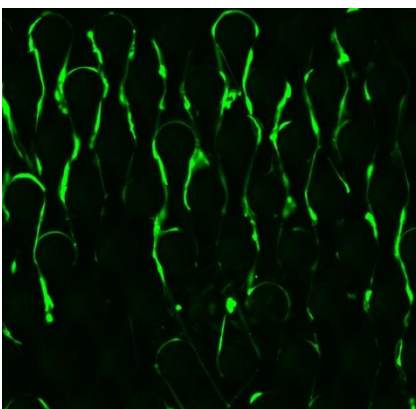
B

R

Composite



PMA 10x



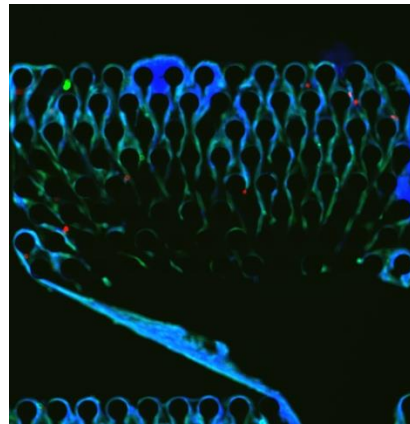
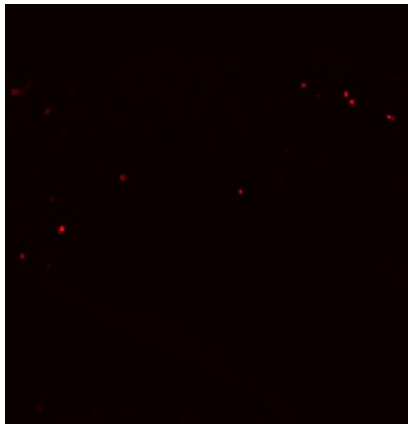
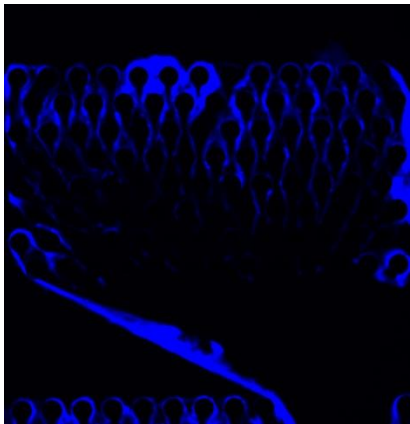
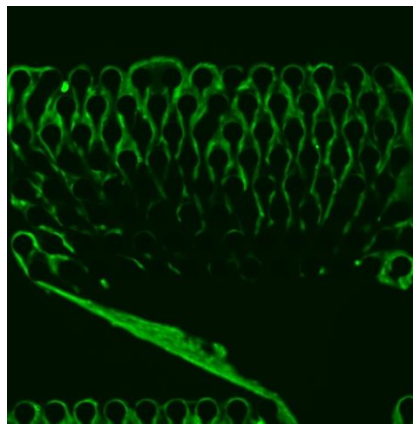
PMA 20x

G

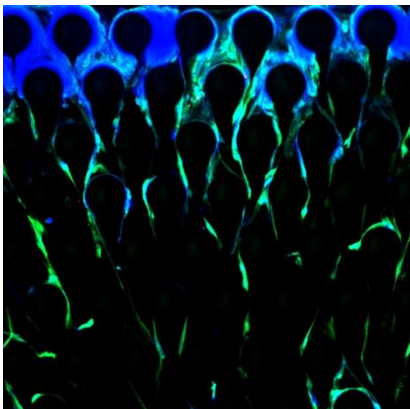
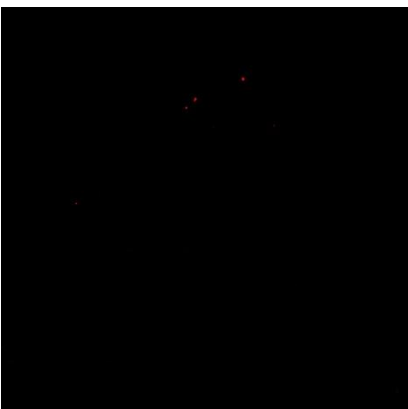
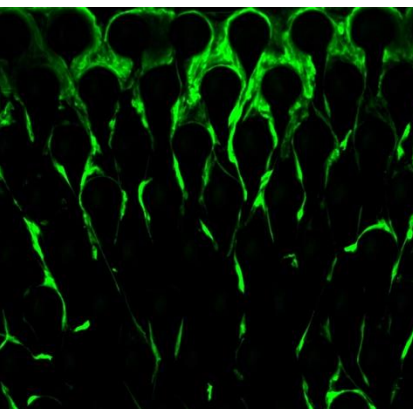
B

R

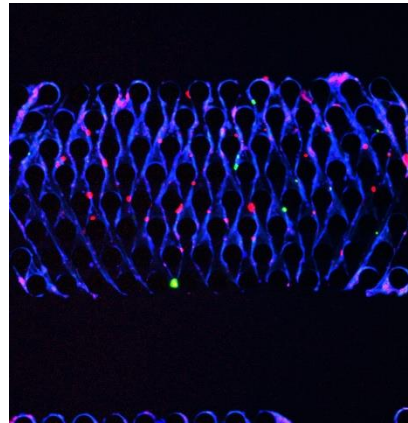
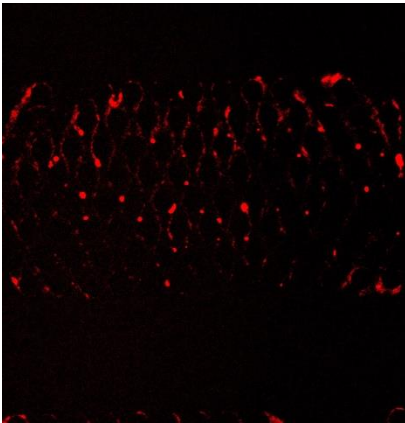
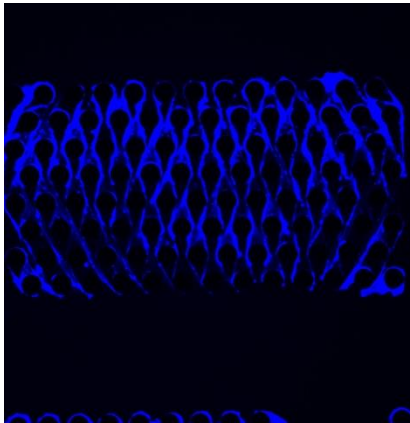
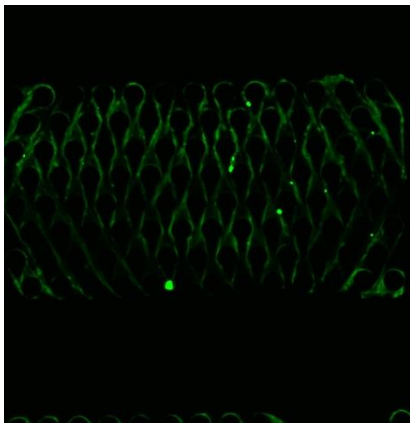
Composite



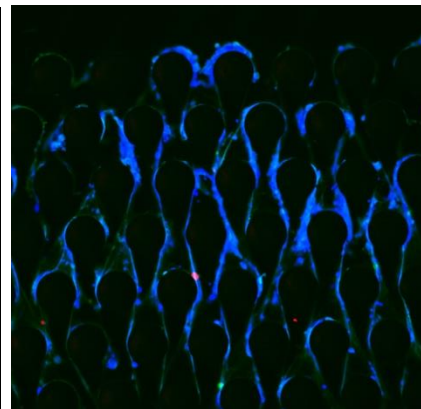
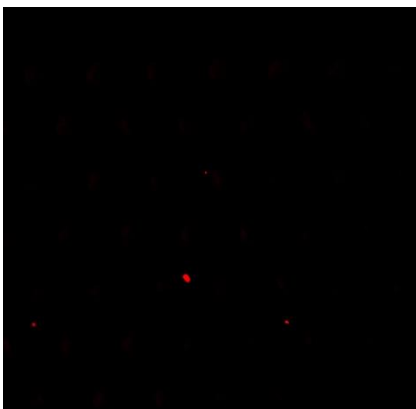
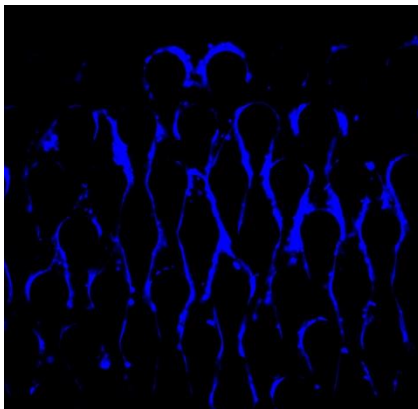
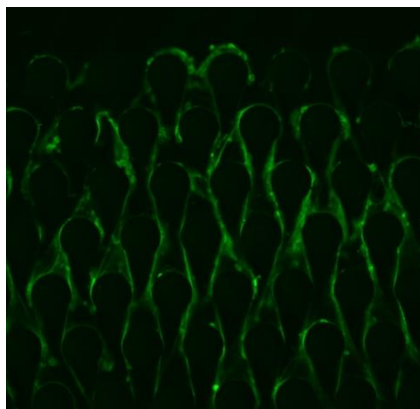
Ionomycin 10x



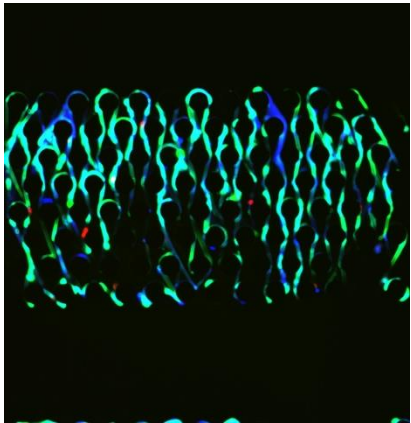
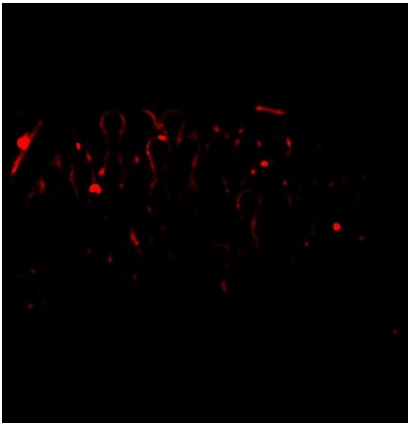
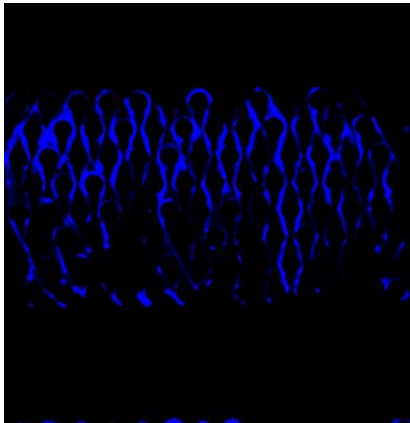
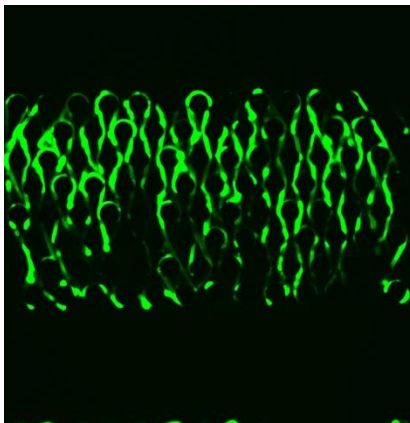
Ionomycin 20x



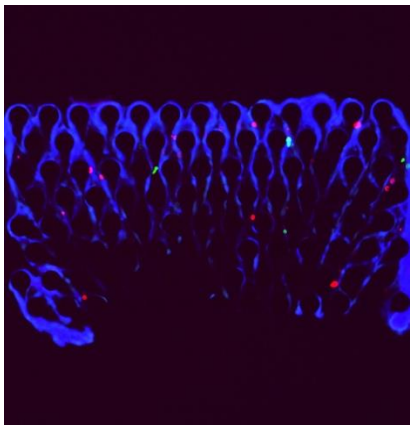
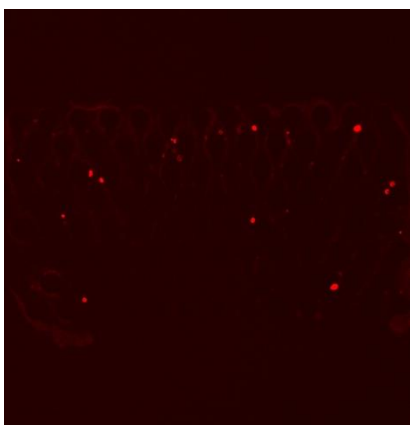
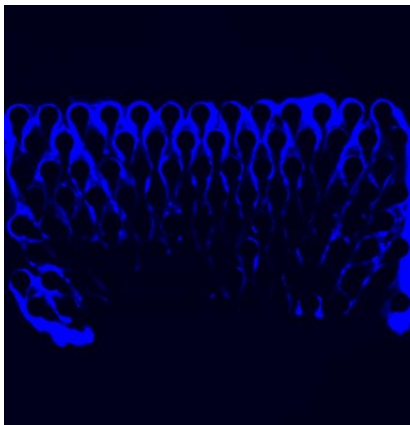
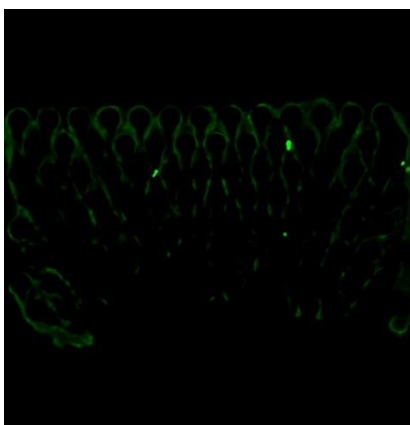
Raptinal 10x



Raptinal 20x



PMA 10x



Ionomycin 10x

NET formation pathway summarized

1. Neutrophil activation with ROS production (or PAD4 activation)
2. Histone modification and chromatin unwinding (PAD4 + MPO)
3. Nuclear rupture and NET release (cell death, not vesicle secretion)
4. NET deployment

Chromatin (in nature)

Chromatin: Complex of DNA and protein in eukaryotic cells, double helix, DNA wrapped around the histones

Chromatin decondensation: From compact chromatin fibers to a more relaxed form

Observed both in mitosis and meiosis

Occasions:

Mitosis (G1 entry, telophase)

Embryogenesis (zygote genome activation)

Cellular Reprogramming

NET formation

Driving Processes:

Histone Acetylation, Histone Phosphorylation, Histone Citrullination, Non-Histone Factors, DNA Demethylation

Chromatin decondensation in neutrophils

PAD4: nuclear enzyme, activated by calcium

Arginine residues to citrulline (histone citrullination)

Weakening histone and DNA binding, loosen up the chromatin

Neutrophil Elastase: serine protease, stored in granules

When neutrophil is activated, NE translocated to the nucleus and cleaves histones

Works complementally with PAD4

Mice deficient of NE is shown to have less chromatin decondensed (knock out potential?)

MPO: Also, a granule enzyme, associates with NE

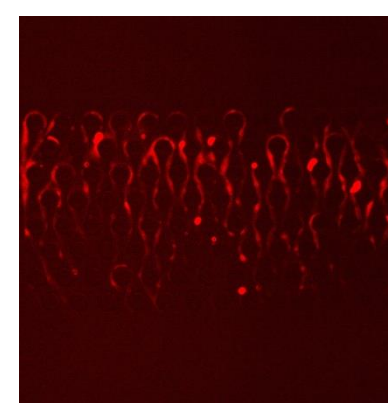
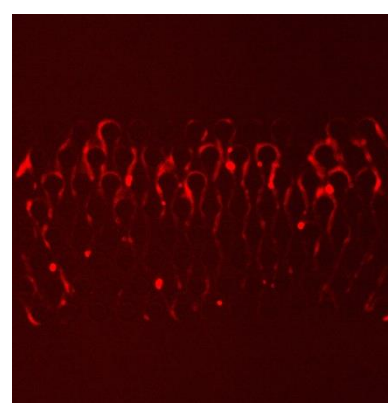
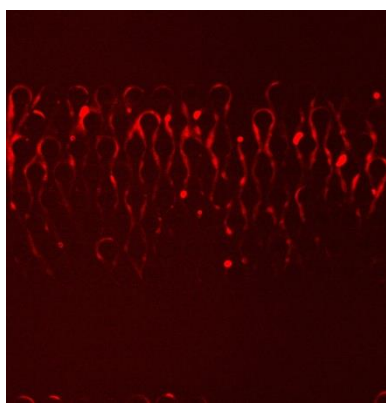
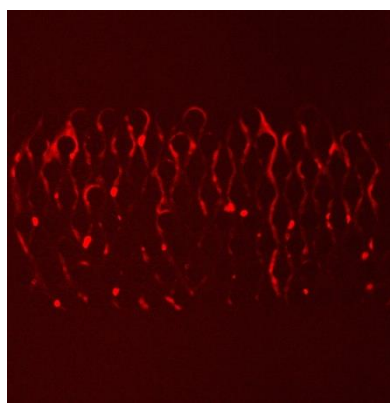
Generates oxidants from hydrogen peroxide

NETs are decorated with MPOs

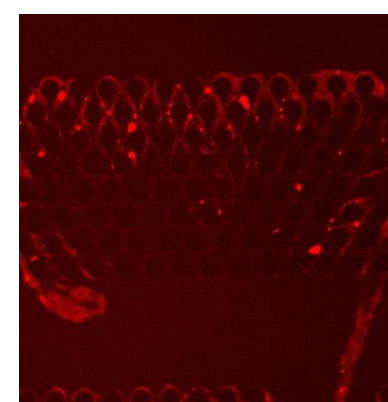
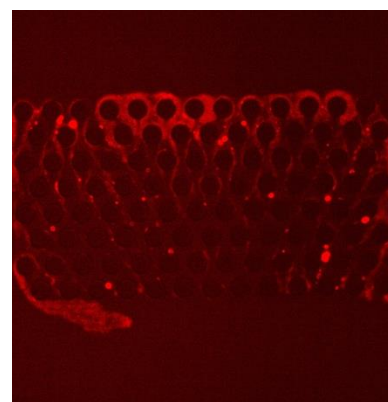
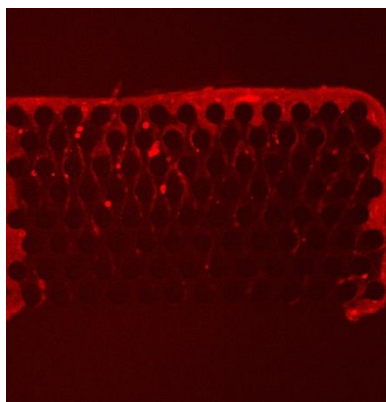
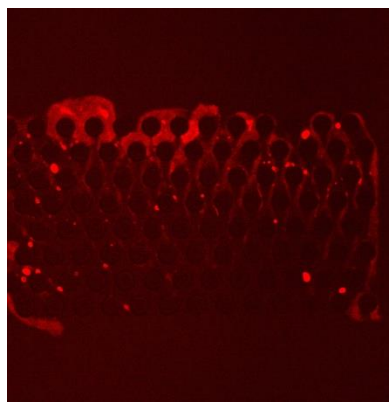
NADPH Oxidase and ROS: ROS -> PAD4

Raptinal: Lacks MPO but has PAD4, so is citrullinated (in tumor cells)

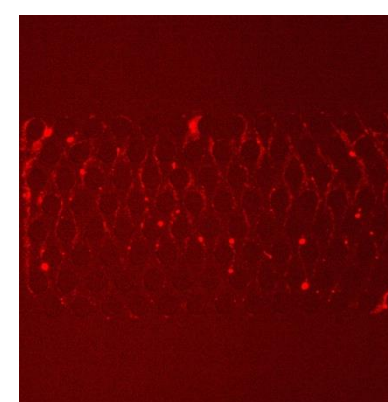
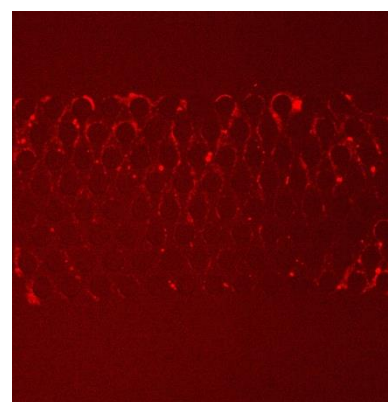
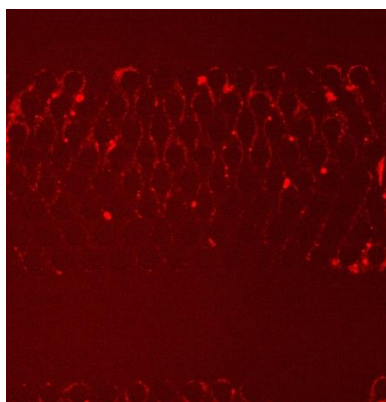
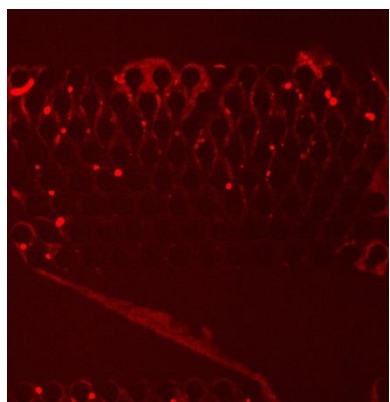
In tumor cells, NET-like structures are called NEPs



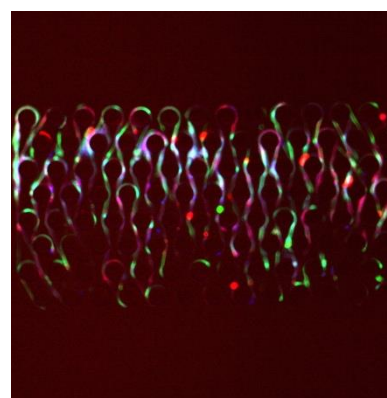
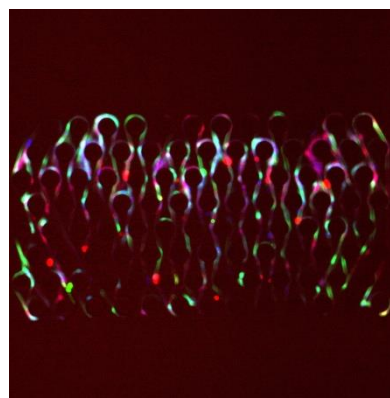
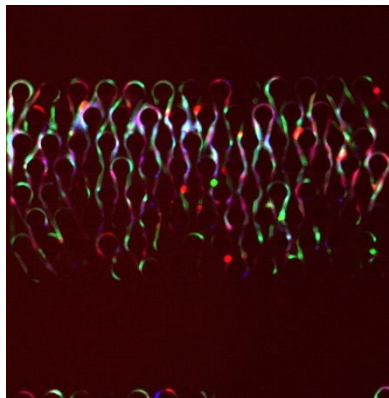
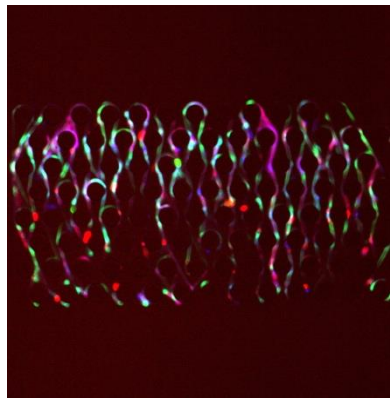
10x Device
PMA



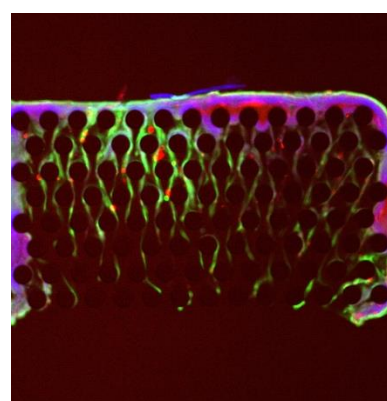
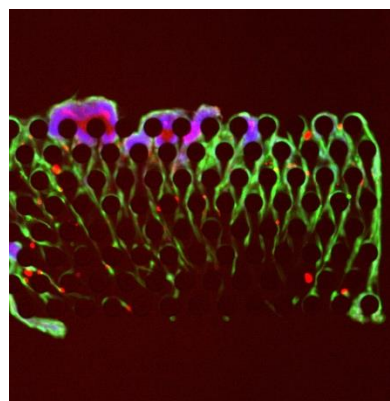
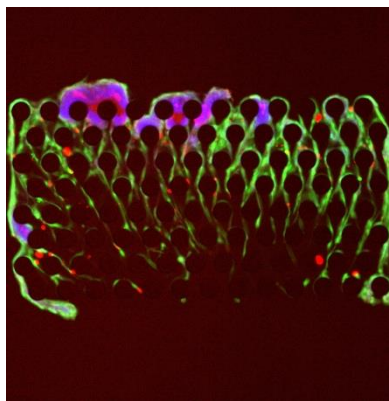
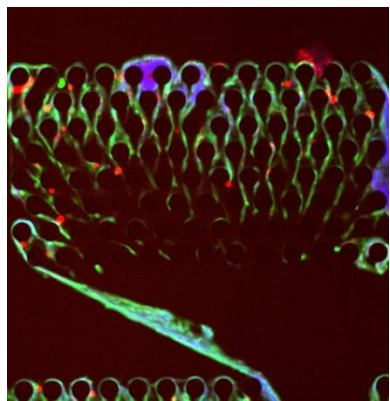
Ionomycin



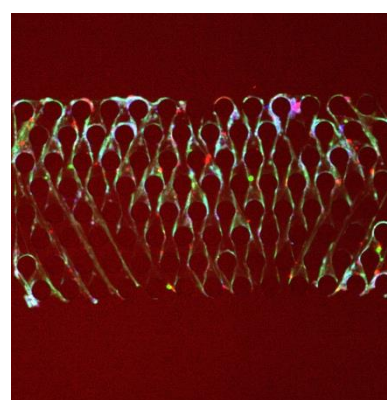
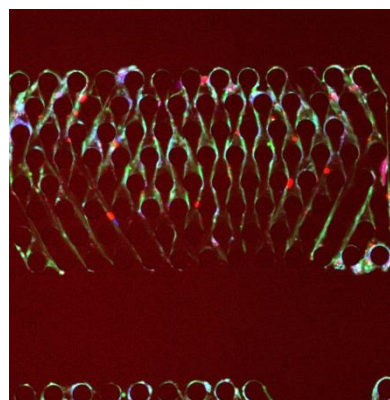
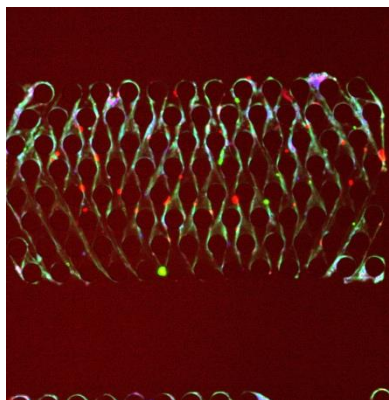
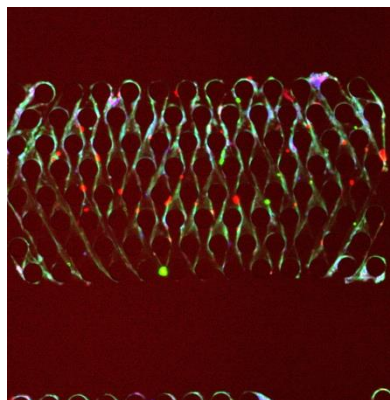
Raptinal



10x Device
PMA



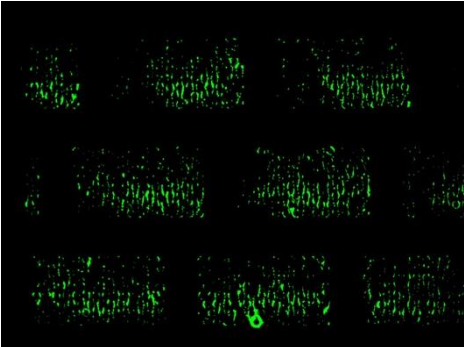
Ionomycin



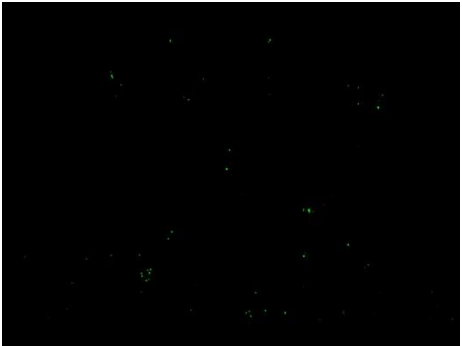
Raptinal



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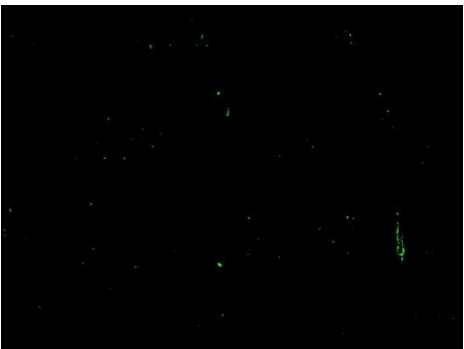
PMA 5 micromoles



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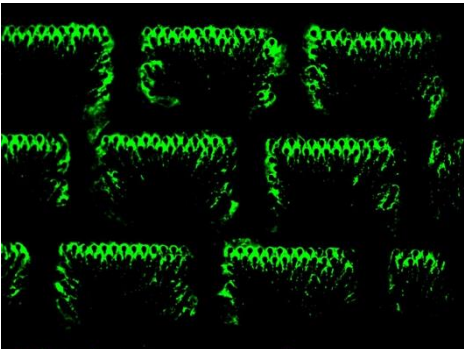
PMA 500 nanomoles



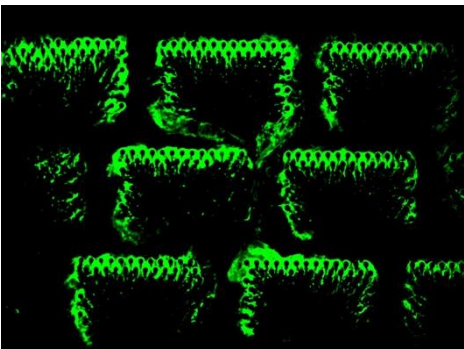
PMA 50 nanomoles



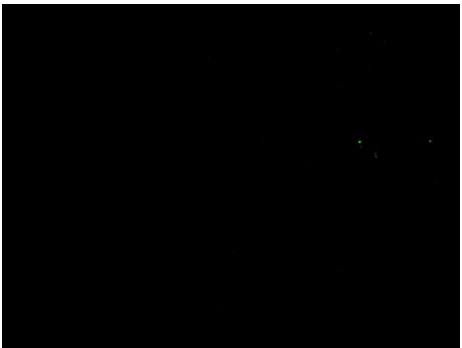
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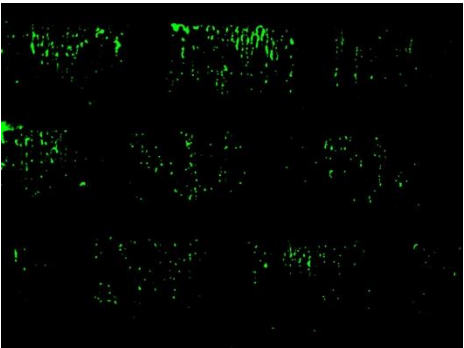
PMA 5 micromoles



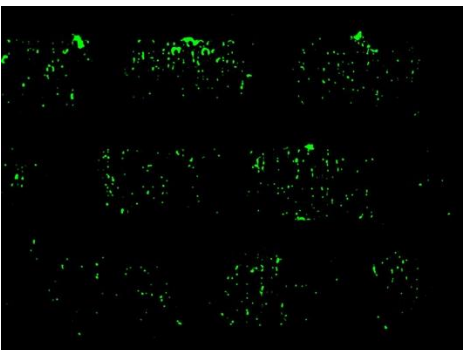
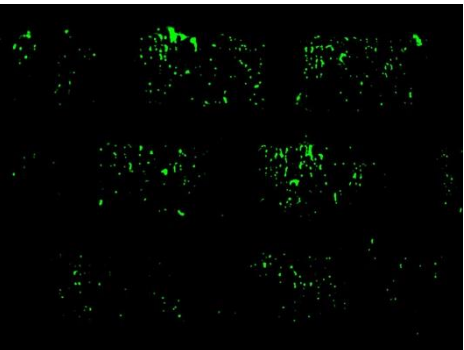
PMA 500 nanomoles



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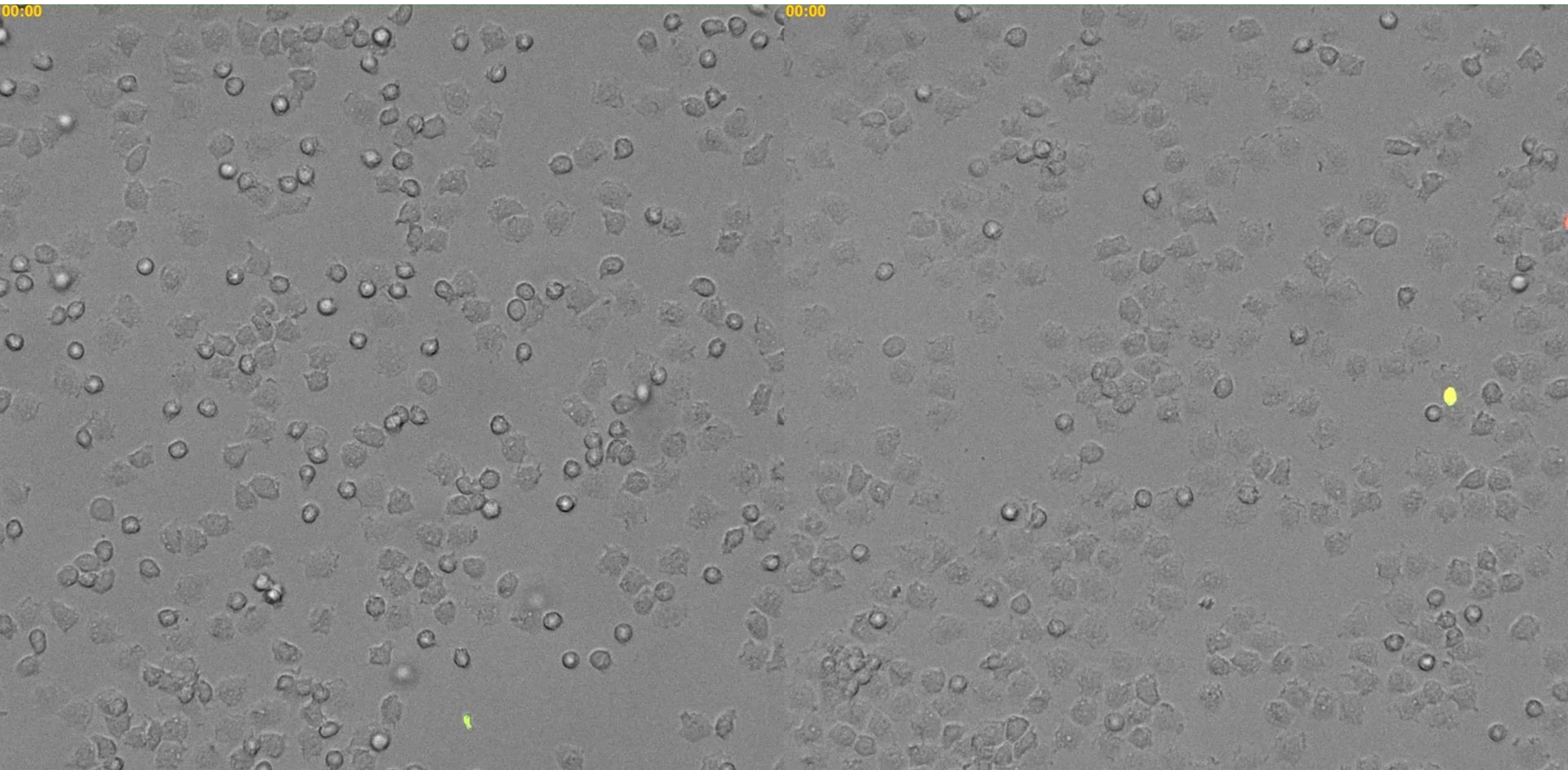


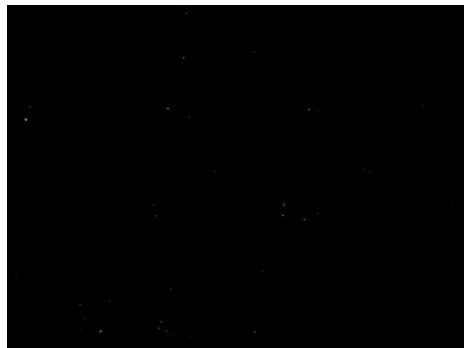
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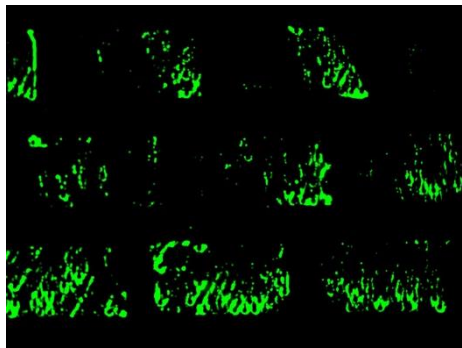
2025-06-17

PMA 500 nanomoles / 5 micromoles

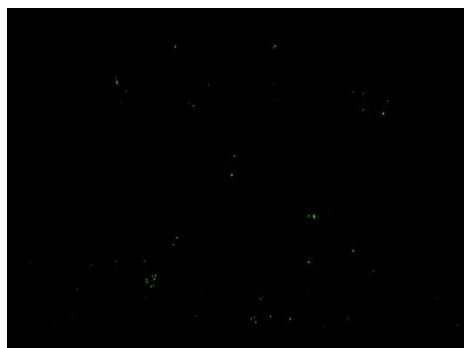
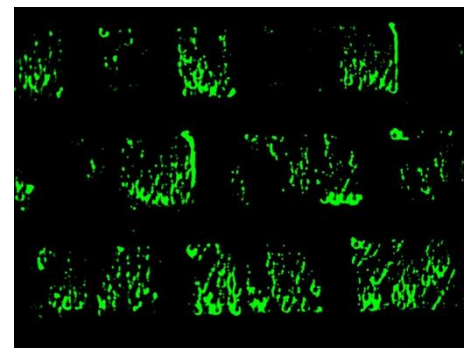




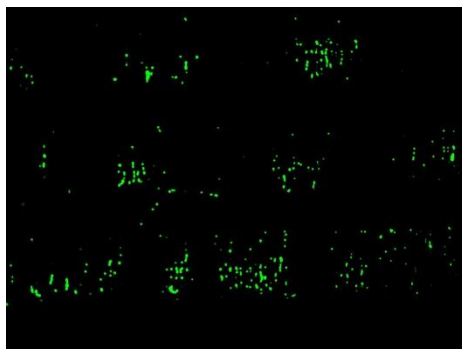
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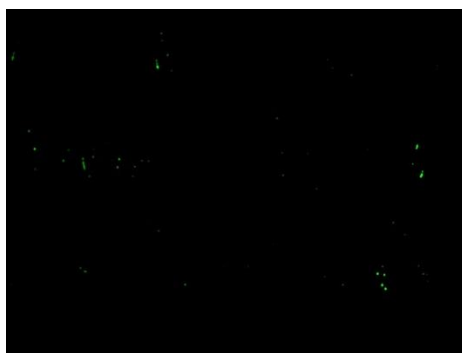
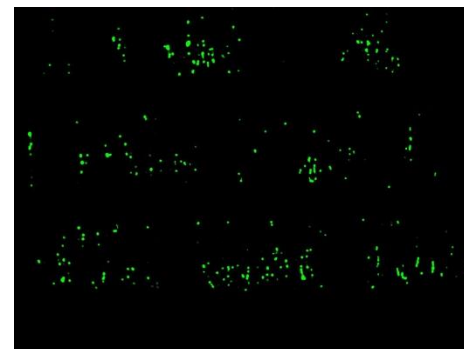
Ionomycin 5 micromoles



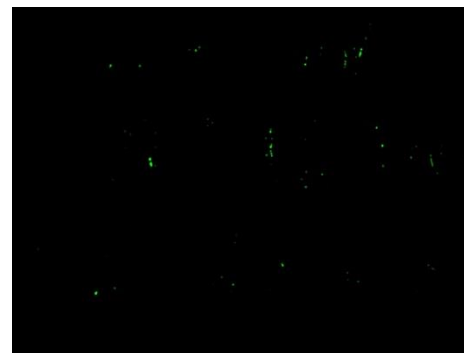
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Ionomycin 500 nanomoles

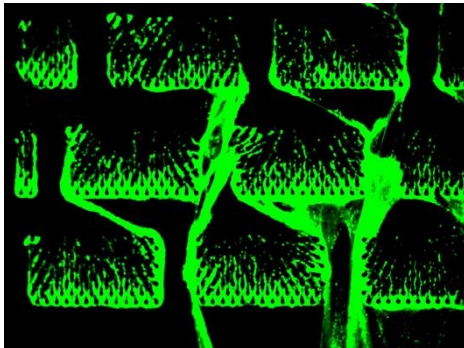
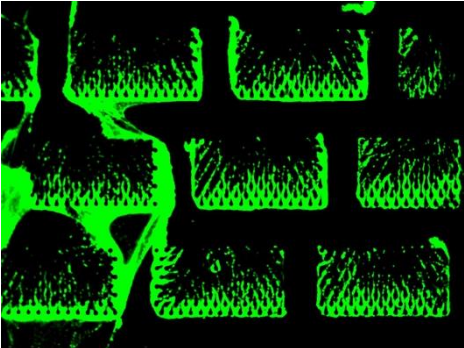


Ionomycin 50 nanomoles

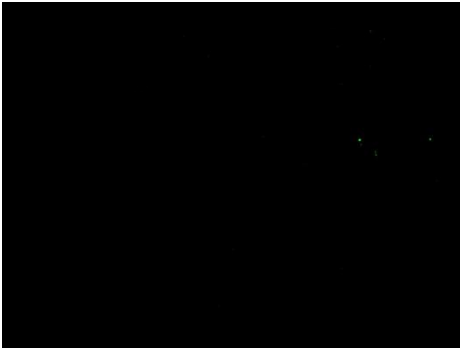




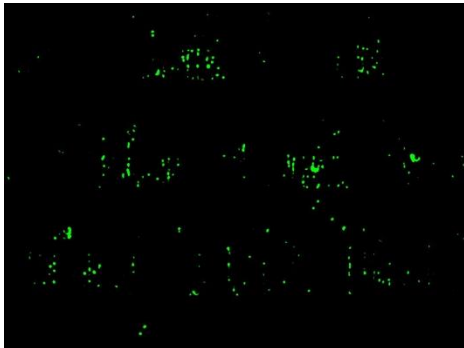
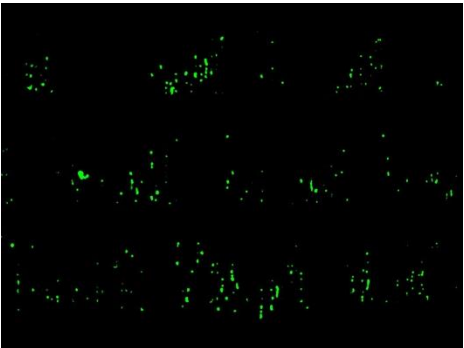
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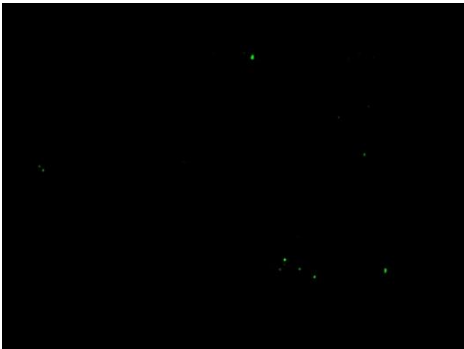
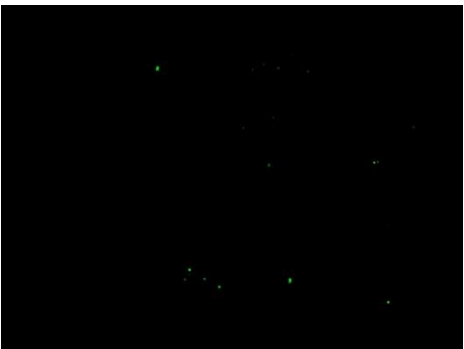
Ionomycin 5 micromoles



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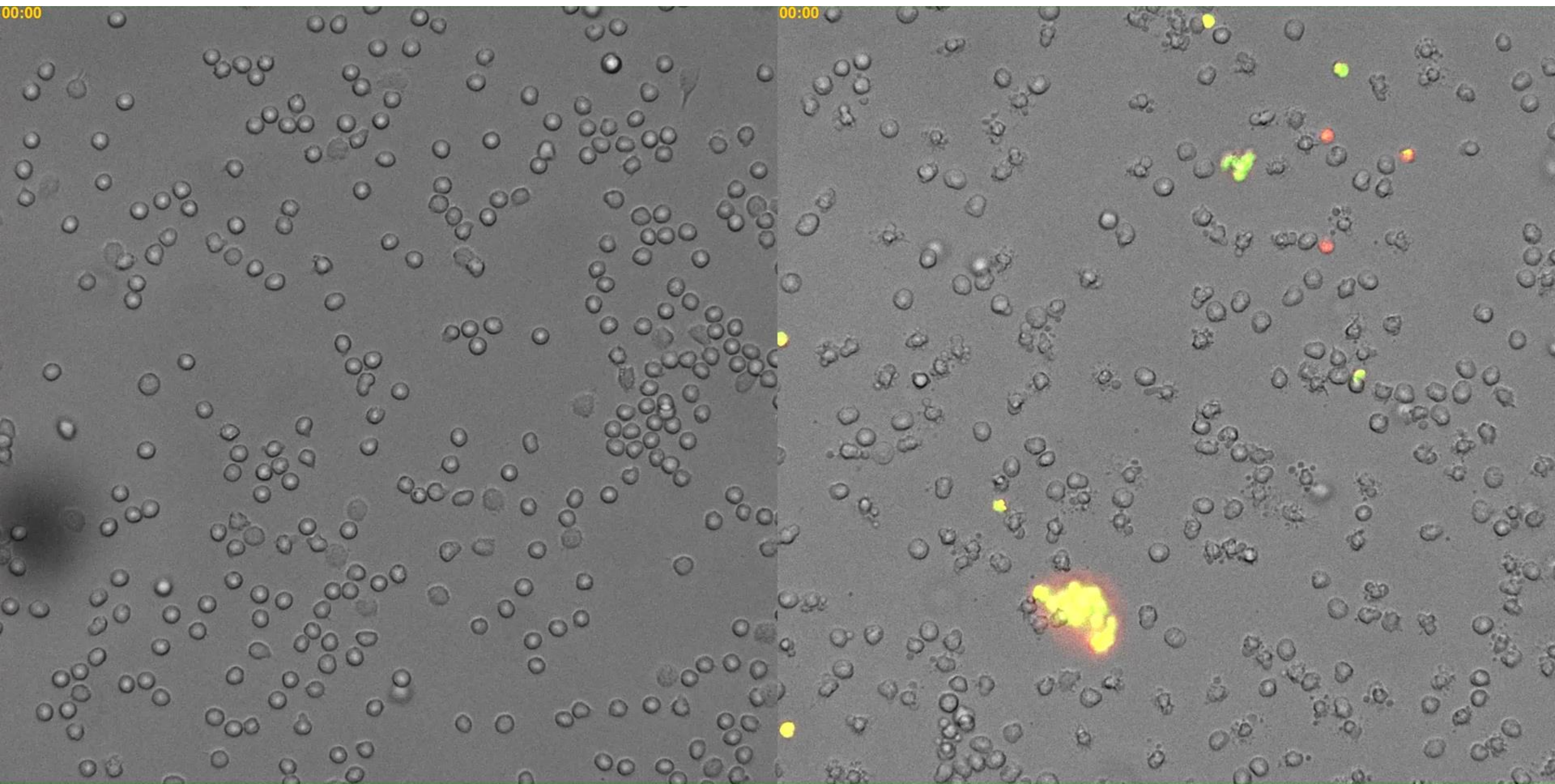
Ionomycin 500 nanomoles

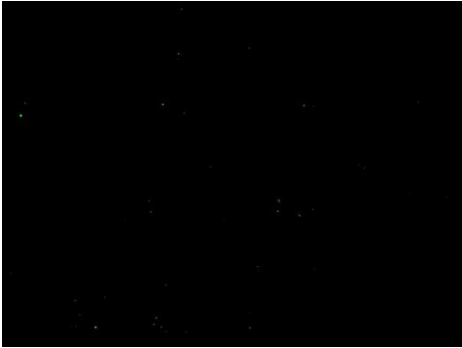


Ionomycin 50 nanomoles

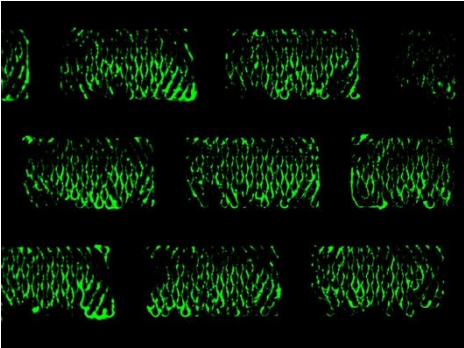
2025-06-17

Ionomycin 500 nanomoles / 5 micromoles

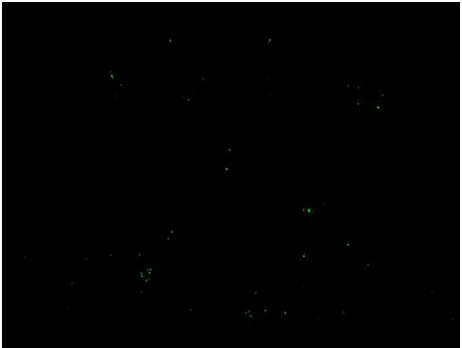




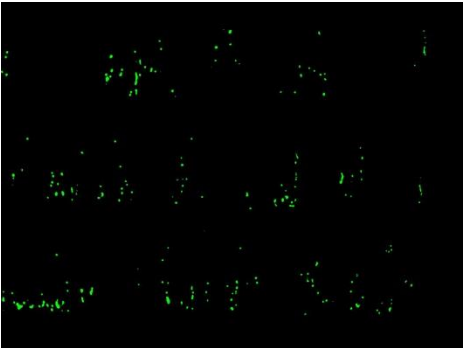
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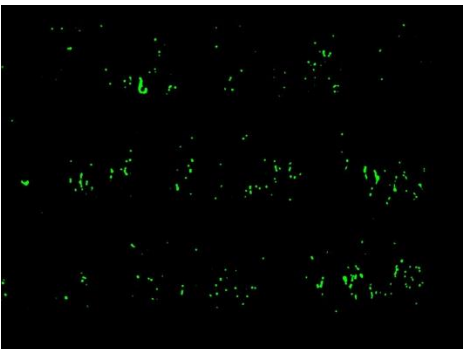
Raptinal 5 micromoles



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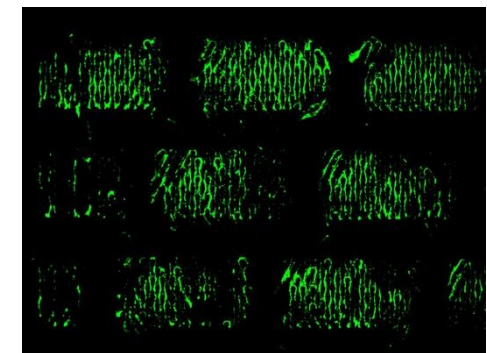
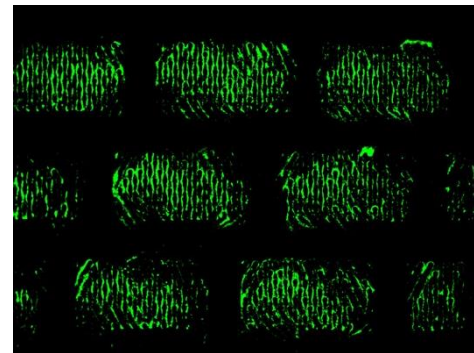
Raptinal 500 nanomoles



Raptinal 50 nanomoles



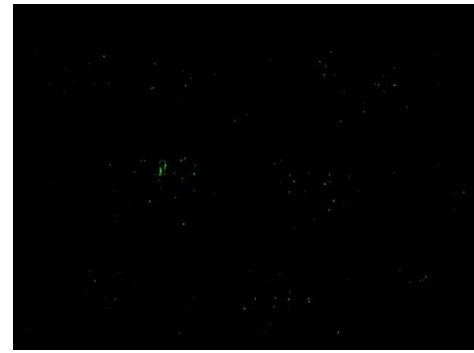
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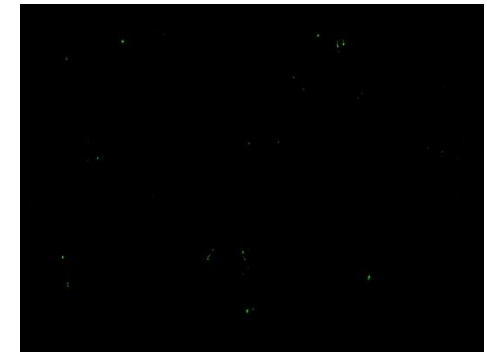
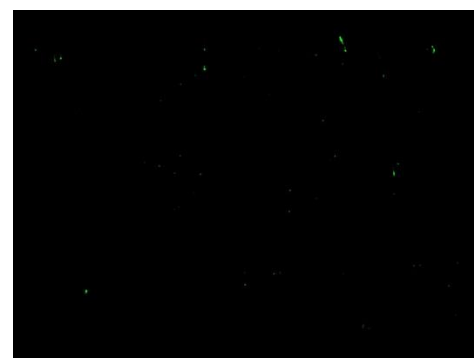
Raptinal 10 micromoles



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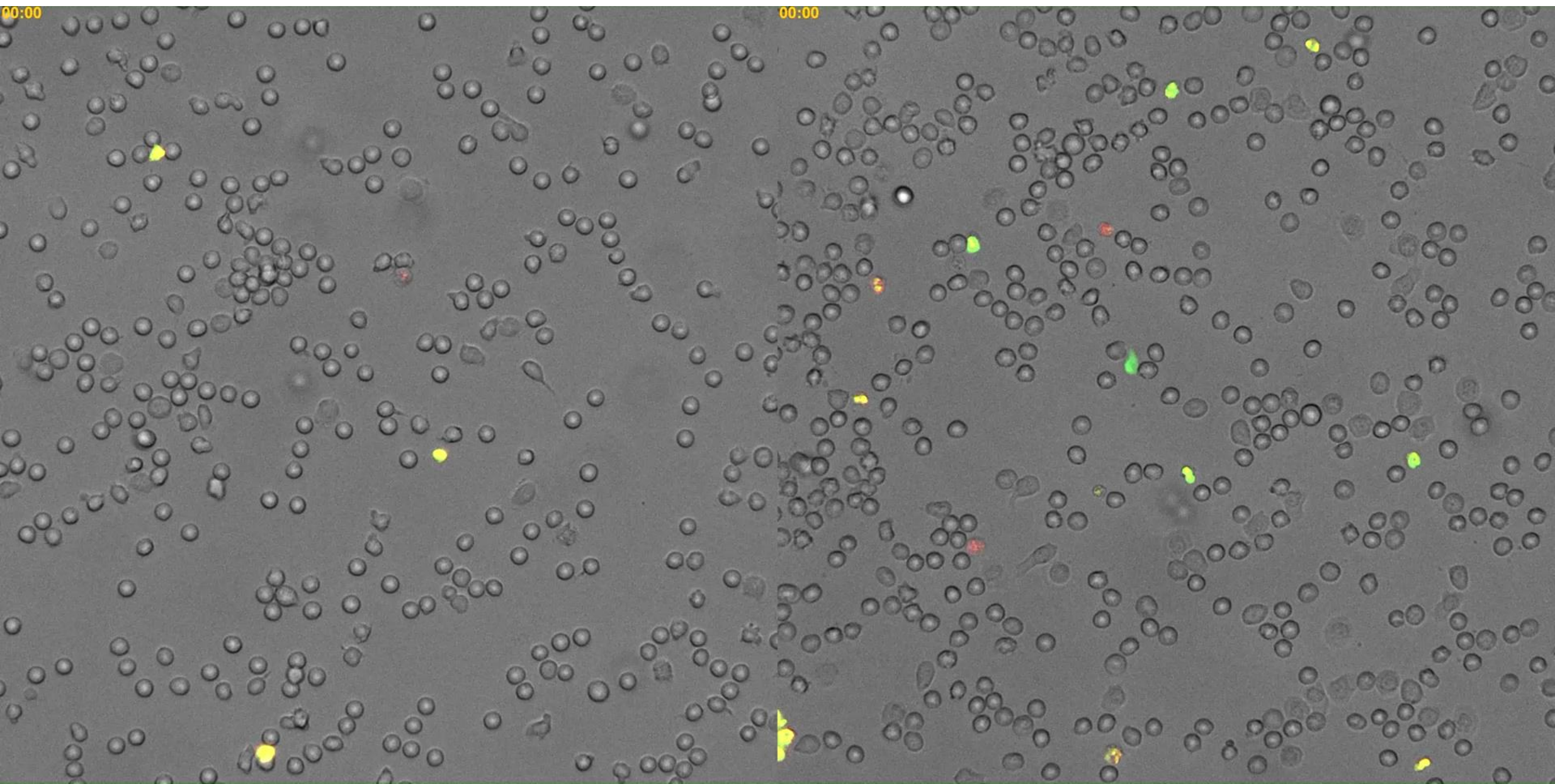
Raptinal 1 micromoles

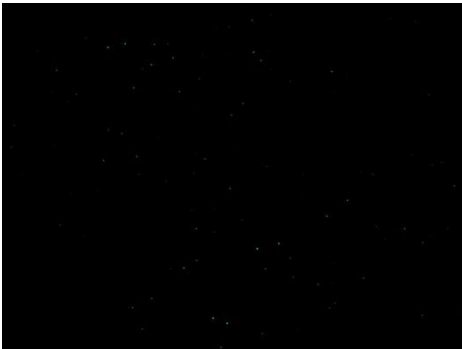


Raptinal 100 nanomoles

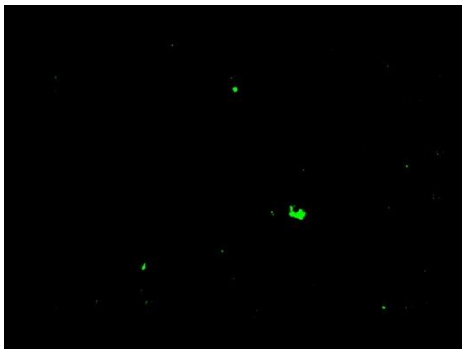
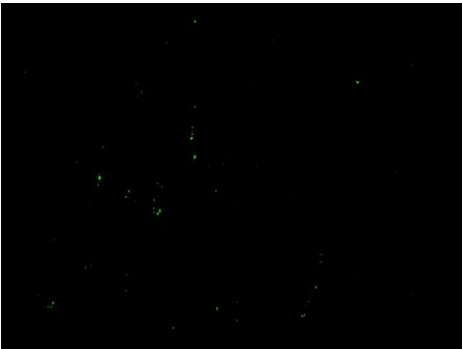
2025-06-17

Raptinal 500 nanomoles / 5 micromoles

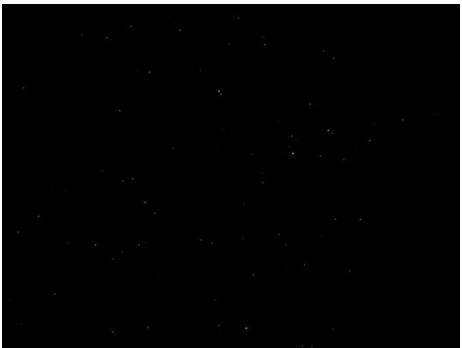




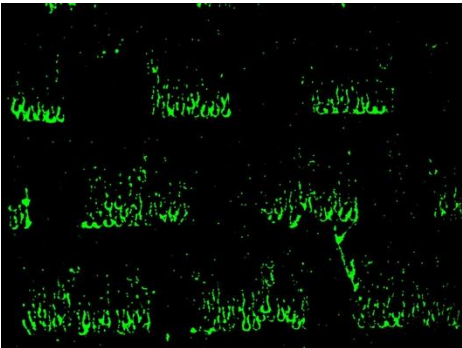
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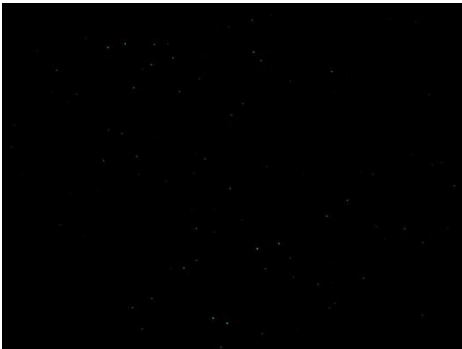
PMA PAD4-Knockout



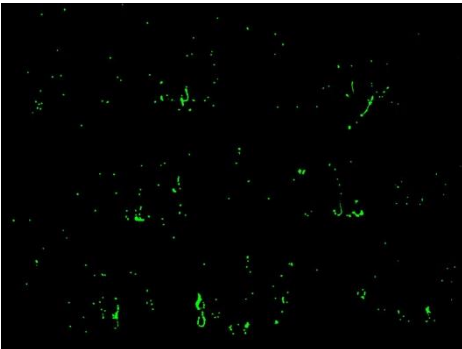
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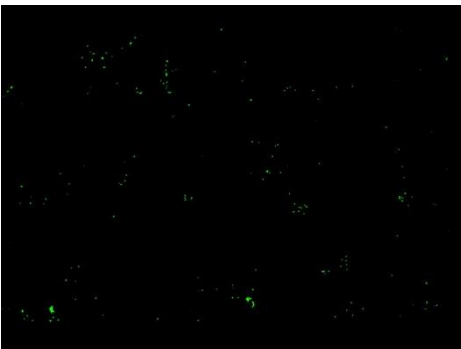
PMA Wildtype



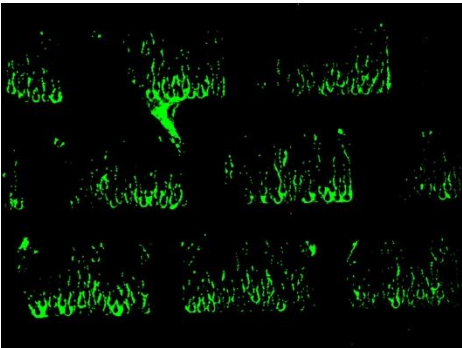
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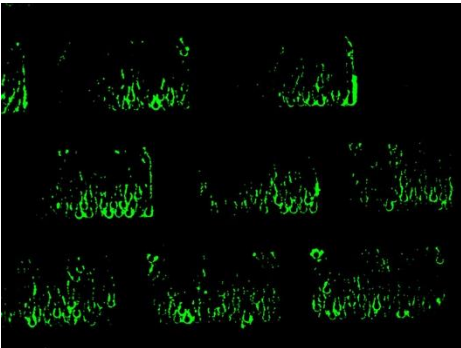
Ionomycin PAD4-Knockout



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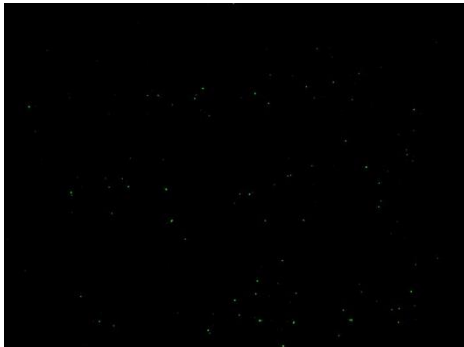


Ionomycin Wildtype





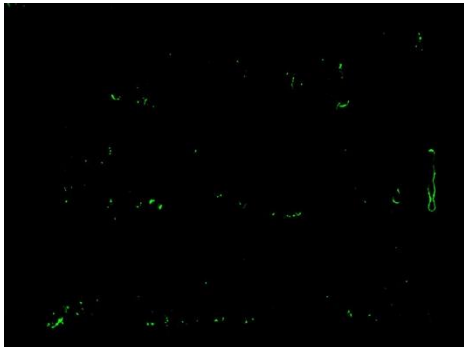
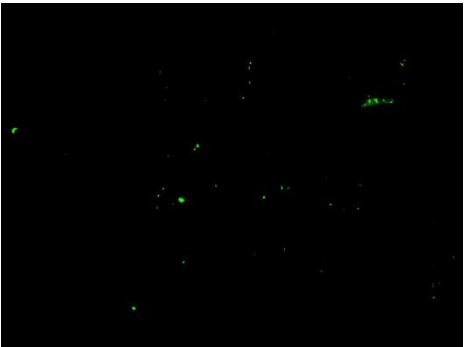
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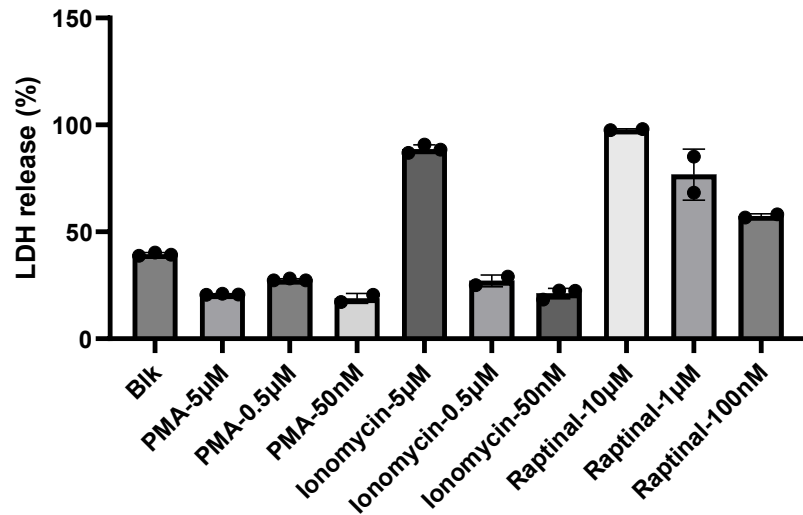
Raptinal PAD4-Knockout



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Raptinal Wildtype



ASC-critine neutrophil , 4h, LDH release (lysis death)

Normal death rates at 4h (low conc., high conc.):

PMA: 55%, 98%

Ionomycin: 100%, 100% (Both are all dead at 3:40)

Raptinal: 5%, 95%

Ionomycin: At 4h all dead, but LDH release vastly different

Raptinal: At 4h massive difference in cell death rate, but LDH release very similar (gradient among different conc.)

PMA: Needed to be repeated, but apparently there is no lysis death

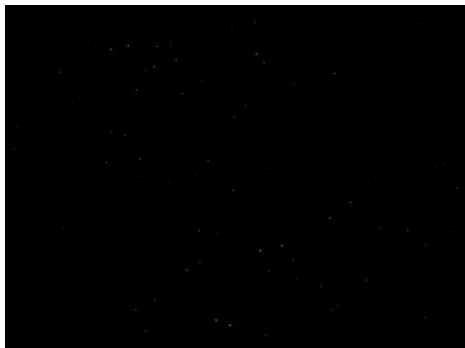
Further plans: Repeating Knockout, bacteria induction, etc.

Citations

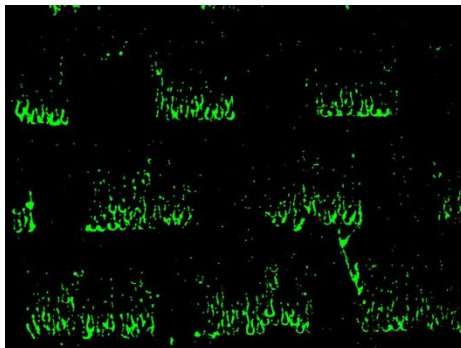
Mice NE Deficiency: (Kolaczowska, E., Jenne, C. N., Surewaard, B. G. J., Thanabalasuriar, A., Lee, W.-Y., Sanz, M.-J., Mowen, K., Opdenakker, G., & Kubes, P. (2015, March 26). Molecular mechanisms of net formation and degradation revealed by intravital imaging in the liver vasculature. *Nature News*.
<https://www.nature.com/articles/ncomms7673#:~:text=The%20enzymes%2C%20peptidyl%20arginine%20deiminase,DNase%20has%20been%20shown>)

MPO: Usher, A. K., & Stockley, R. A. (2013, November 13). *The link between chronic periodontitis and COPD: A common role for the neutrophil? - BMC medicine*. BioMed Central.
<https://bmcmmedicine.biomedcentral.com/articles/10.1186/1741-7015-11-241#:~:text=ROS%20have%20been%20implicated%20in,release%20is%20shown%20in%20Figure%C2%A04>

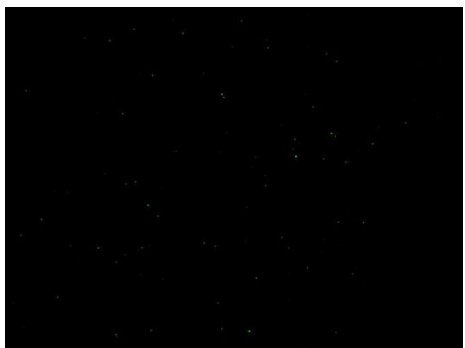
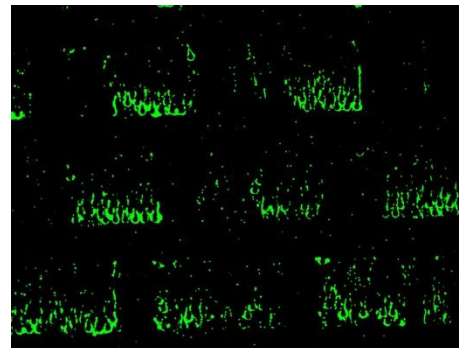
Raptinal Apoptosis: Park, W.-Y., Gray, J. M., Holewinski, R. J., Andresson, T., So, J. Y., Carmona-Rivera, C., Hollander, M. C., Yang, H. H., Lee, M., Kaplan, M. J., Cappell, S. D., & Yang, L. (2023, March 27). *Apoptosis-induced nuclear expulsion in tumor cells drives S100A4-mediated metastatic outgrowth through the rage pathway*. *Nature News*. <https://www.nature.com/articles/s43018-023-00524-z>



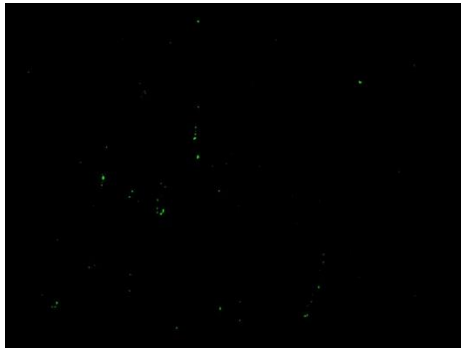
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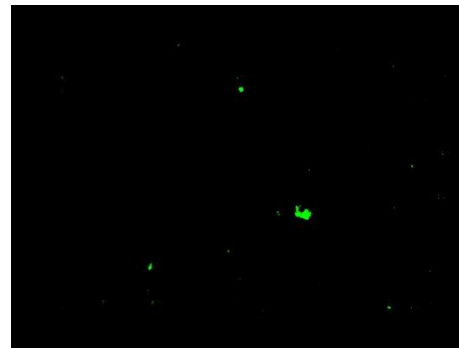
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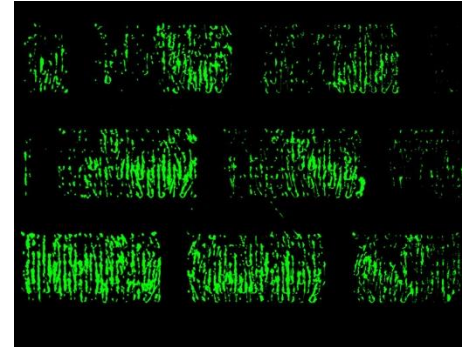
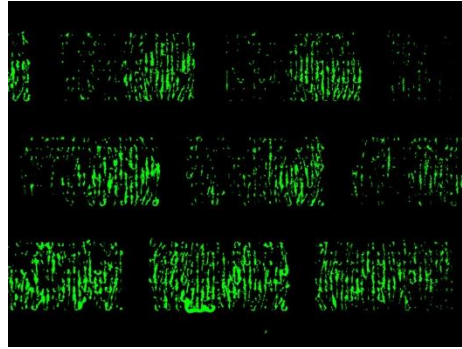


PMA PAD4-Knockout





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PMA Wildtype



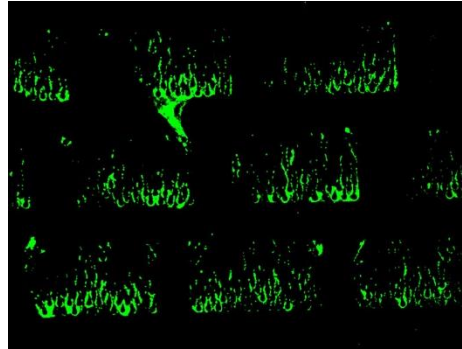
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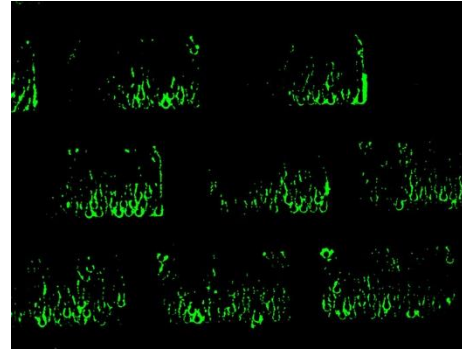
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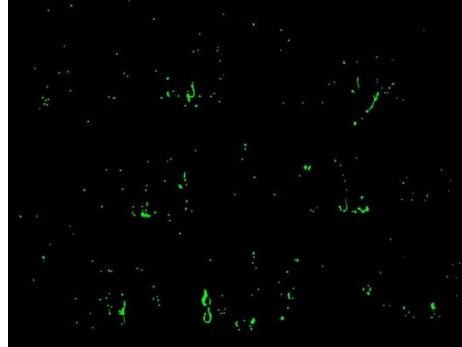
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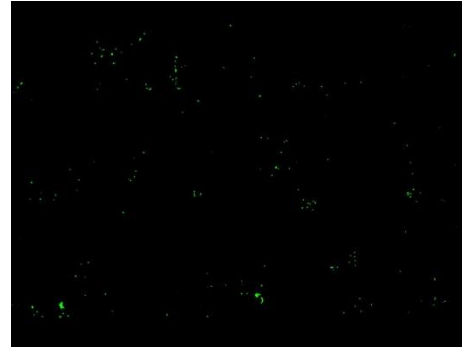
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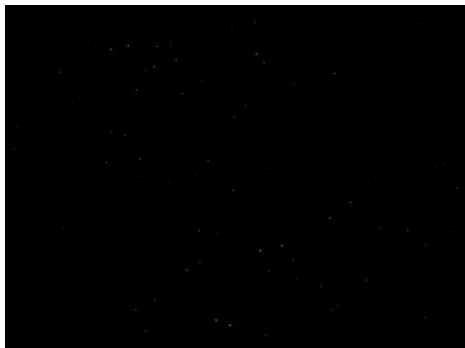


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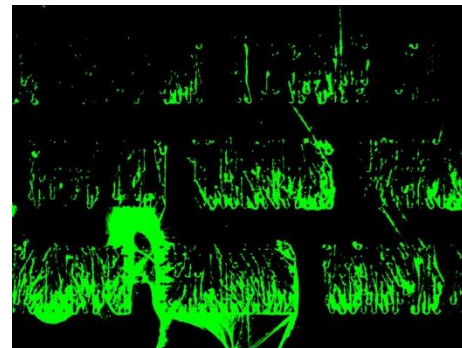
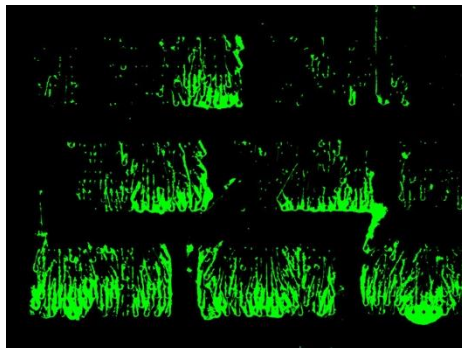


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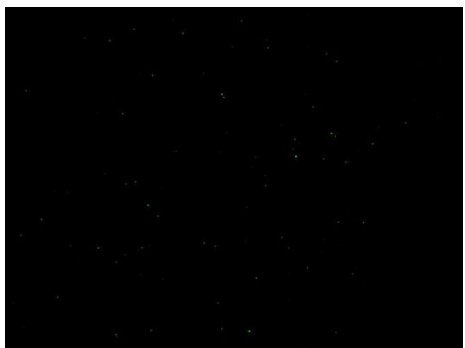




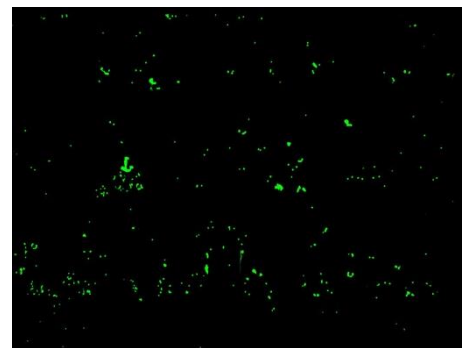
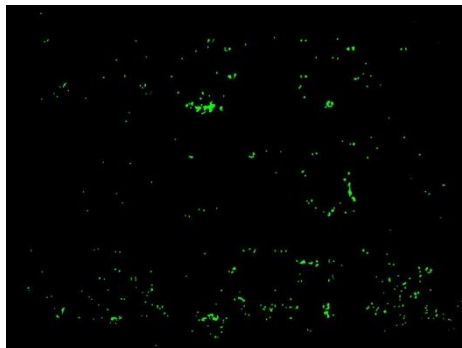
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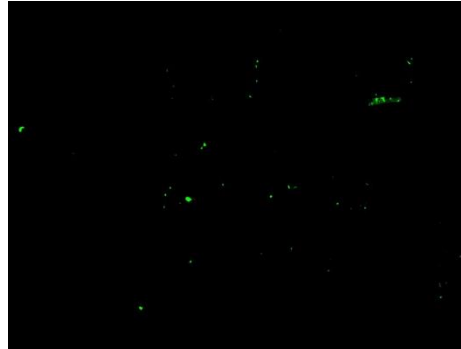
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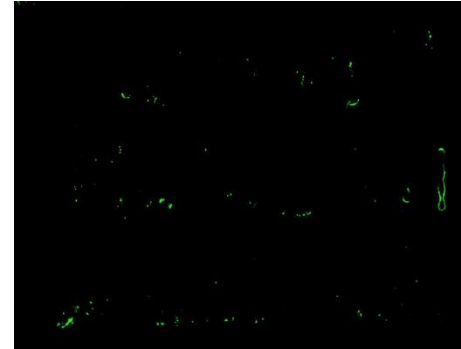
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Raptinal Wildtype



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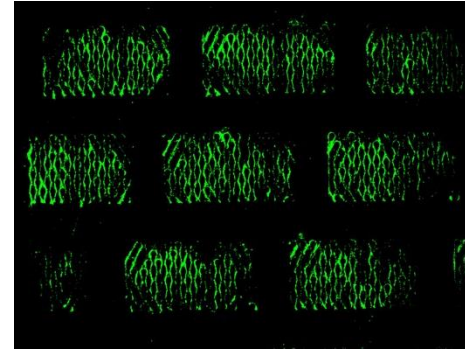
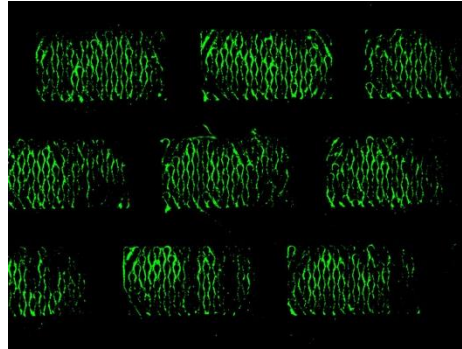


Raptinal PAD4-Knockout





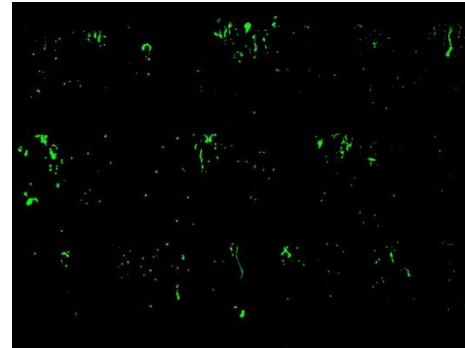
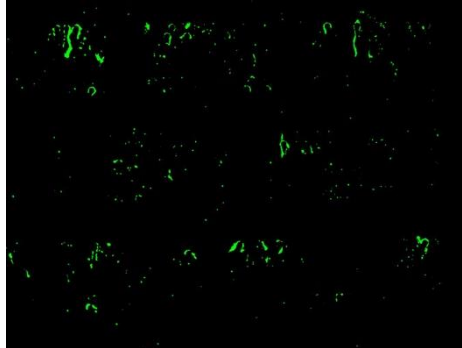
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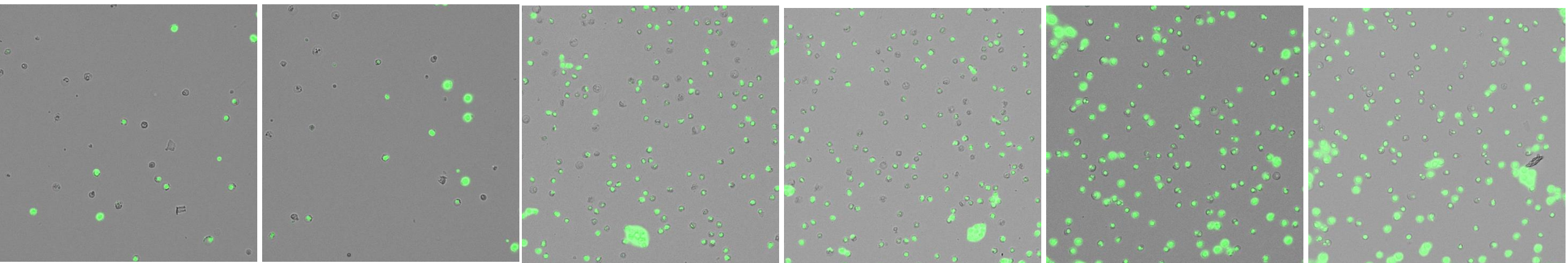
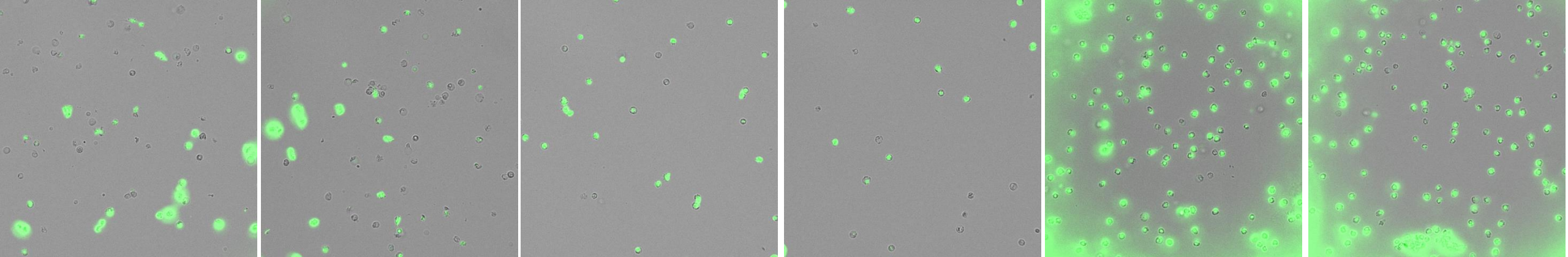
Raptinal Wildtype



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Raptinal PAD4-Knockout



PMA

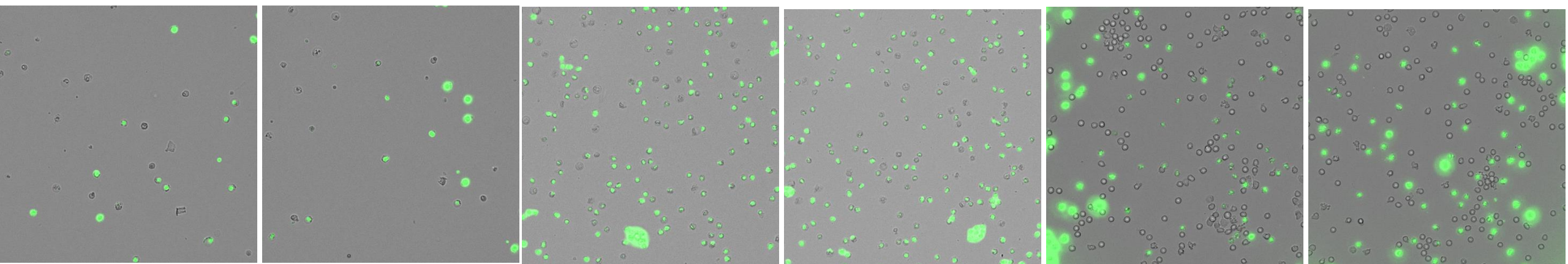
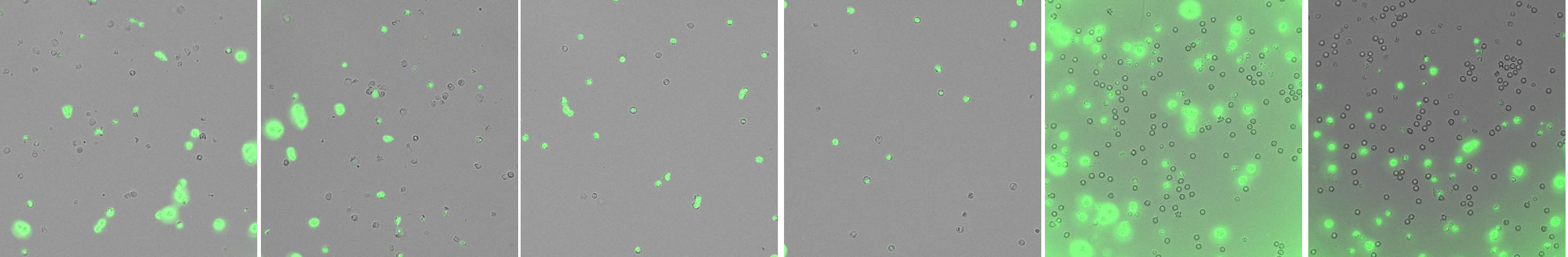
PMA

Ionomycin

Ionomycin

Raptinal

Raptinal



PMA

PMA

Ionomycin

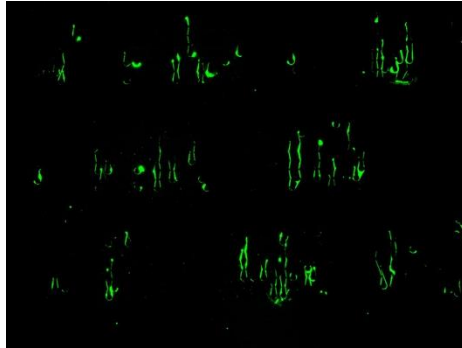
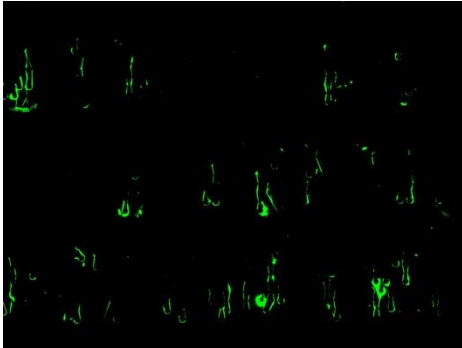
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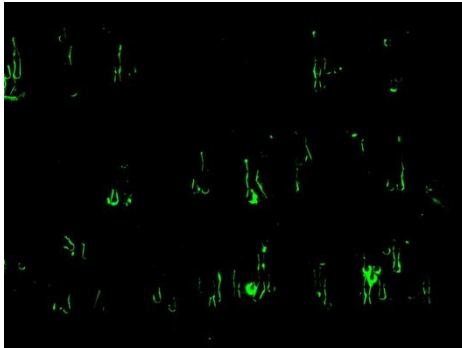
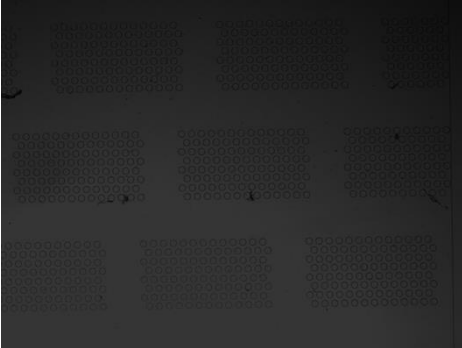
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Spontaneous Death Pico



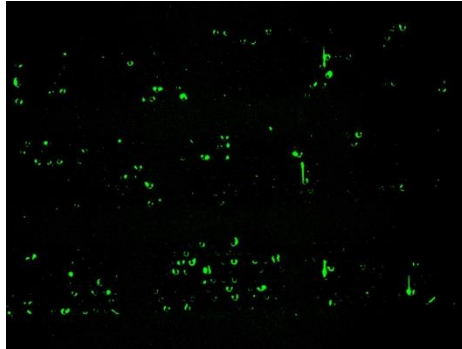
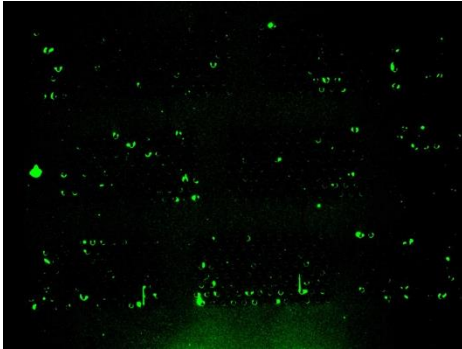
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Spontaneous Death Pico



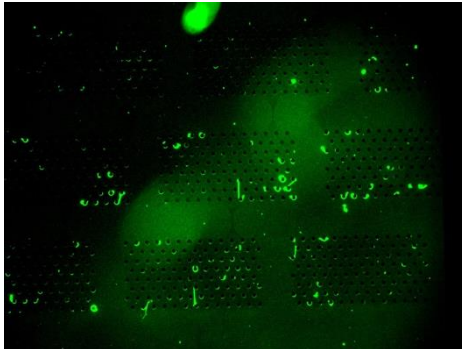
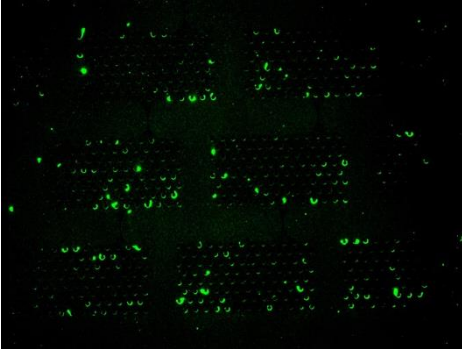
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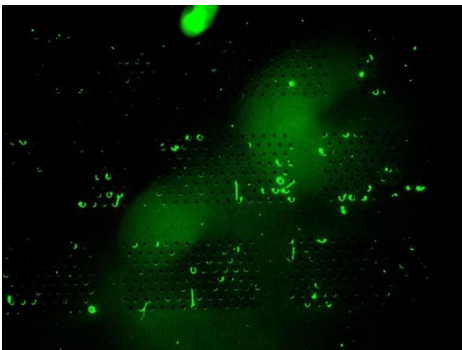
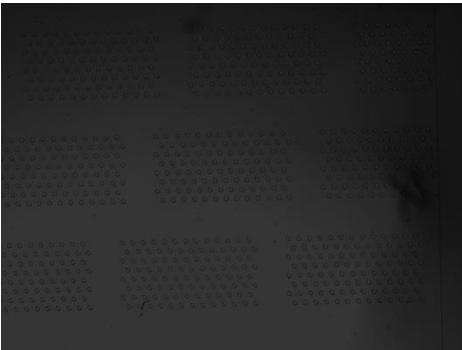
Spontaneous Death Plate
2x cells



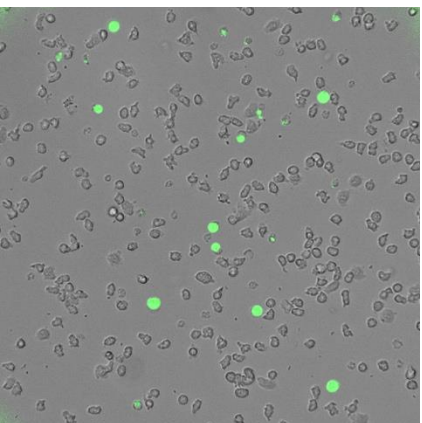
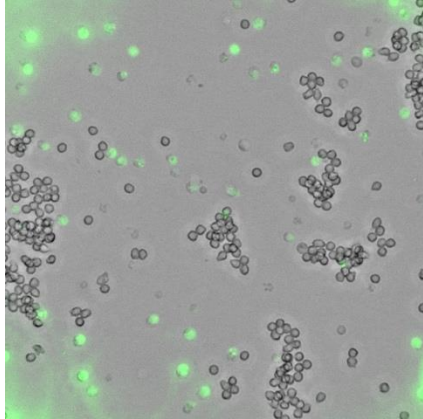
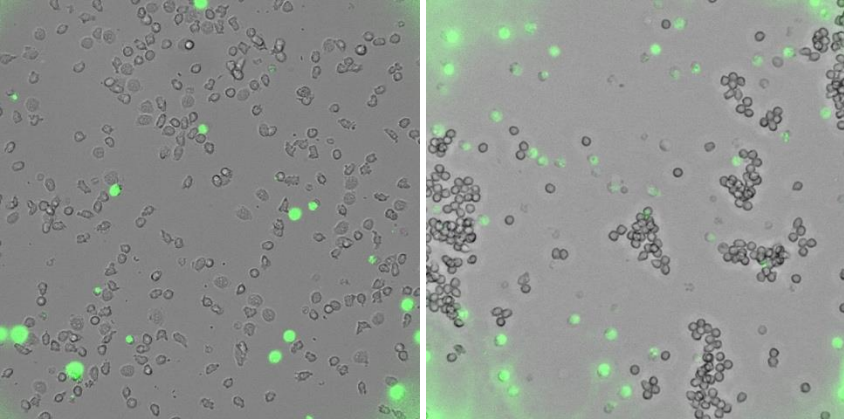
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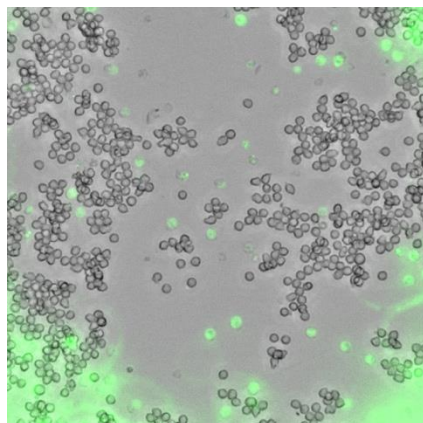
Spontaneous Death Plate
2x cells



Spontaneous Death Plate
2x cells



B2



C3

NET formation pathway summarized

- 1. Neutrophil activation with ROS production (or PAD4 activation)
- 2. Histone modification and chromatin unwinding (PAD4 + MPO)
- 3. Nuclear rupture and NET release (cell death, not vesicle secretion)
- 4. NET deployment
- **Pathway is different in mice and humans**
- **Right now, our experiments only in mice neutrophils**

Mouse neutrophil, in vitro, PMA PAD4 dependency

- Literature: “PAD4 is essential for antibacterial innate immunity mediated by neutrophil extracellular traps”
PMC , [pmc.ncbi.nlm.nih.gov/articles/PMC2931169/](https://pubmed.ncbi.nlm.nih.gov/articles/PMC2931169/).
Accessed 7 July 2025.
- Eghbalzadeh, Kaveh. “Compromised Anti-inflammatory Action of Neutrophil Extracellular Traps in PAD4-Deficient Mice Contributes to Aggravated Acute Inflammation After Myocardial Infarction”
Compromised Anti-inflammatory Action of Neutrophil,
1 Oct. 2019
www.frontiersin.org/journals/immunology/articles/10.3389/fimmu.2019.02313/full. Accessed 7 July 2025.

Neutrophils trap and kill bacteria by forming highly decondensed chromatin structures, termed neutrophil extracellular traps (NETs). We previously reported that histone hypercitullination catalyzed by peptidylarginine deiminase 4 (PAD4) correlates with chromatin decondensation during NET formation. However, the role of PAD4 in NET-mediated bacterial trapping and killing has not been tested. Here, we use PAD4 knockout mice to show that PAD4 is essential for NET-mediated antibacterial function. Unlike PAD4^{+/+} neutrophils, PAD4^{-/-} neutrophils cannot form NETs after stimulation with chemokines or incubation with bacteria, and are deficient in bacterial killing by NETs. In a mouse infectious

MI. Noteworthy, PAD4^{-/-} neutrophils were unable to release NETs upon *ex vivo* stimulation with ionomycin or PMA, but produced significantly higher amounts of reactive oxygen species (ROS). Increased levels of circulating cell-free DNA, mitochondrial DNA and cardiac troponin were found in PAD4^{-/-} mice in the acute phase of MI when compared to WT mice. Reduced cardiac expression of *IL-6*, *IL-10*, and M2 marker genes, as well as increased *TNF-α* expression, suggested a pro-inflammatory state. PAD4^{-/-} mice displayed significantly

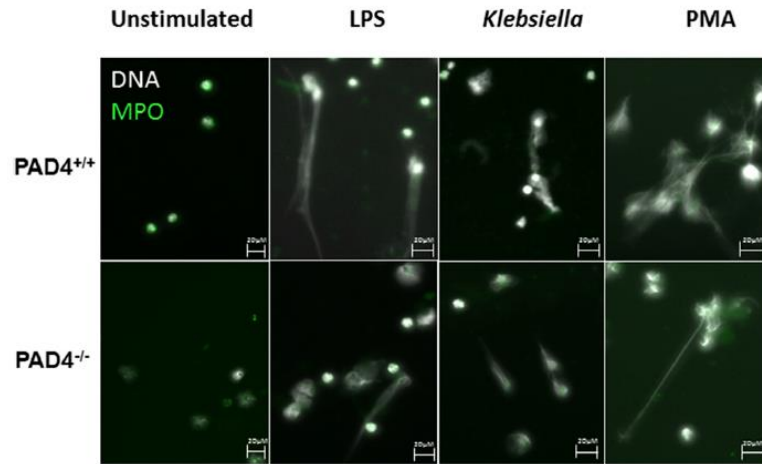


FIGURE 5. Ex vivo NET formation does not rely on PAD4. PAD4^{+/+} and PAD4^{-/-} neutrophils were stimulated with medium control, LPS, PMA, or UV-radiated *Klebsiella* for 3 h. NETs were visualized with immunofluorescence of MPO (in green) or DNA (DAPI staining in white).

During murine pneumosepsis, CitH3 levels were increased in the lungs of PAD4^{+/+} but not of PAD4^{-/-} mice. Combined light and electron microscopy showed NET-like structures surrounding *Klebsiella* in areas of CitH3 staining in the lung; however, these were also seen in PAD4^{-/-} mice with absent CitH3 lung staining. Moreover, cell-free DNA and nucleosome levels were mostly similar in both groups. Moreover, *Klebsiella* and LPS could still induce NETosis in PAD4^{-/-} neutrophils. Both

Mouse neutrophil, in
vitro, PMA PAD4
independency (?)
Pneumonia-induced

- Literature: van, "Cornelis. "Role of Peptidylarginine Deiminase 4 in Neutrophil Extracellular Trap Formation and Host Defense during *Klebsiella pneumoniae*- Induced Pneumonia-Derived Sepsis" PubMed, 9 July 2018, pubmed.ncbi.nlm.nih.gov/29987161/. Accessed 10 July 2025.
- Induced pneumonia-derived sepsis
- Not relevant

Human neutrophil, in vitro, PMA PAD4 independency

PKC is required for PMA, *C. albicans* and GBS induced NETosis. Only PMA and nigericin require calcium to make NETs

To test whether ROS were required for NETosis, we pre-incubated neutrophils with pyrocatechol before stimulation. As expected, ROS was absolutely required for PMA-induced NETosis (Figure 3D). Conversely, to induce NETOSIS.

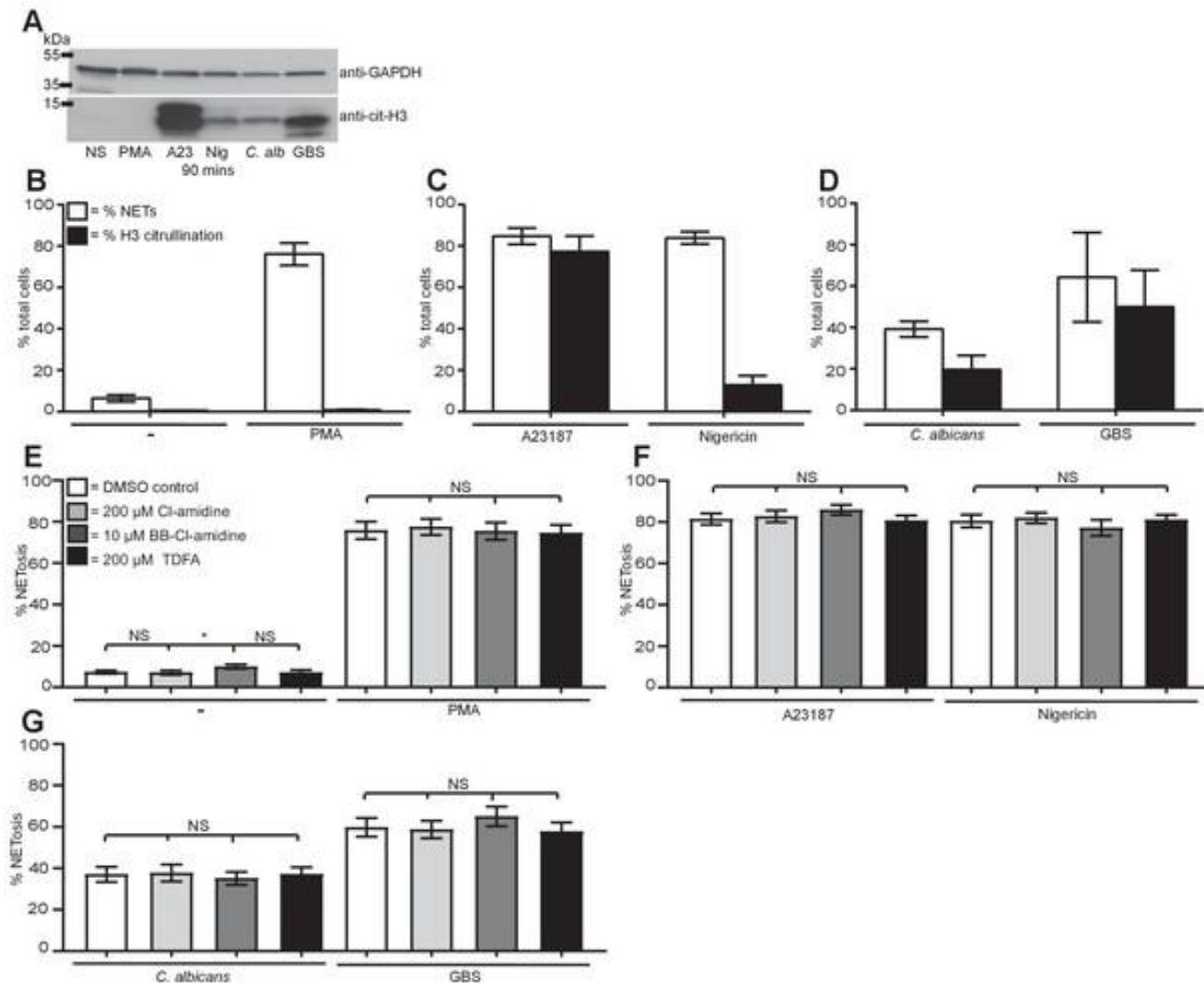
Conversely, activation with the ionophores A23187 and nigericin does not require PKC, ROS, MPO or NE or transcriptional activation and calcium only has a limited role. Histone H3 is citrullinated upon activation by these stimuli; however, pre-treatment with PAD inhibitors does not affect the ability of the neutrophils to make NETs. This suggests that citrullination (of histone H3 at least) is a consequence of NETosis, and that PAD4 is not required for NET formation. Ionophore and PMA-induced NETosis appears to be distinct, at least in the few components of the signal transduction cascade already described.

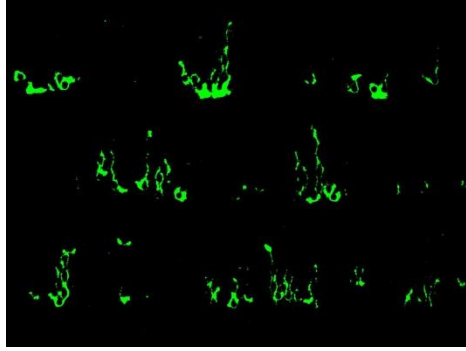
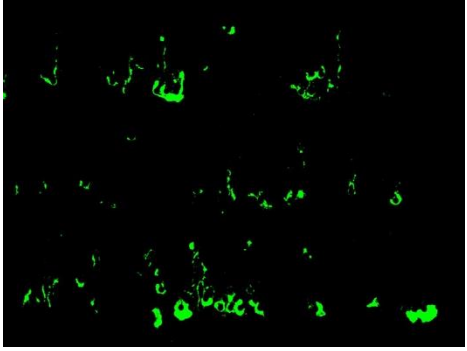
We next explored if citrullination was required for NET formation. Neutrophils were pre-treated with three inhibitors of PAD proteins: Cl-amidine and BB-Cl-amidine, both of which inhibit PAD2 and PAD4, and Thr-Asp-F-amidine (TDFA), a potent specific PAD4 inhibitor. Treating neutrophils with Cl-amidine, BB-Cl-amidine or TDFA did not induce NETosis spontaneously (Figure 6E). PMA induced NETosis was not affected by these inhibitors, consistent with the data obtained with GSK199, another PAD4-selective inhibitor (Figure 6E) (Lewis et al., 2015). Importantly, in response to A23187 or nigericin stimulation, NETosis remained intact after incubation with the three inhibitors (Figure 6F).

• Literature: Kenny, Elaine F. “Diverse stimuli engage different neutrophil extracellular trap pathways” ELife, 2 June 2017, elifesciences.org/articles/24437. Accessed 7 July 2025.

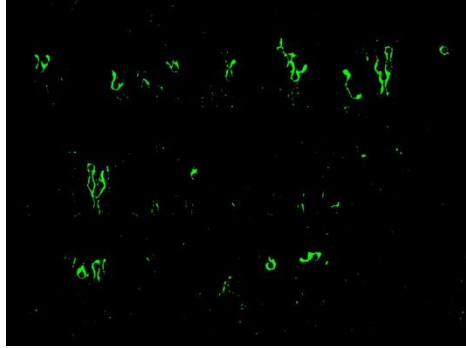
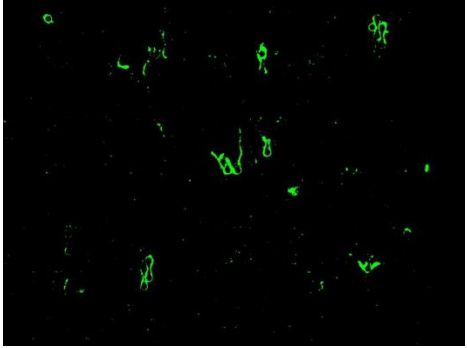
Human neutrophil, in vitro, PMA PAD4 independency

- Literature: Kenny, Elaine F. “Diverse stimuli engage different neutrophil extracellular trap pathways” ELife, 2 June 2017, [elifesciences.org/articles/24437](https://doi.org/10.7554/elifesciences.org/articles/24437). Accessed 7 July 2025.
- (Figure 6E is PAD4 knockout)

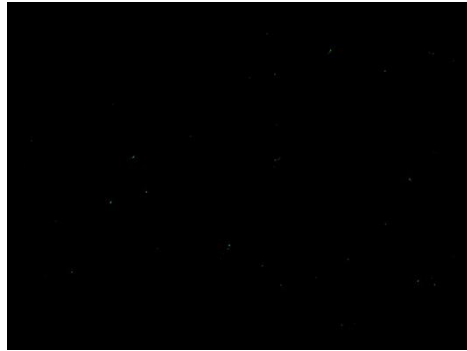
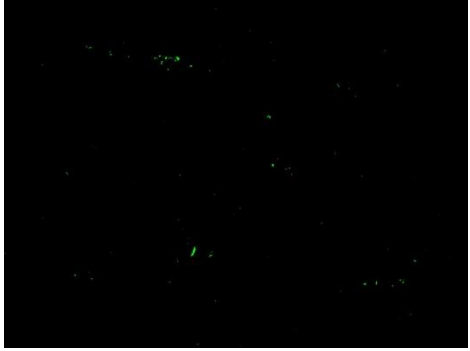




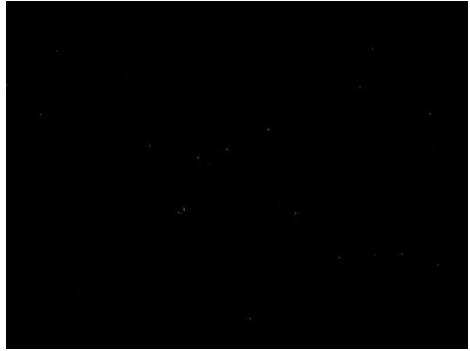
Spontaneous Death
- Percoll, 24h



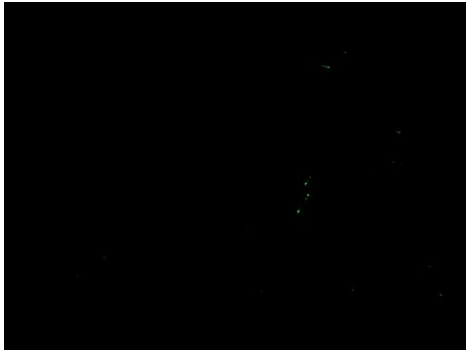
Spontaneous Death
- Percoll, 48h



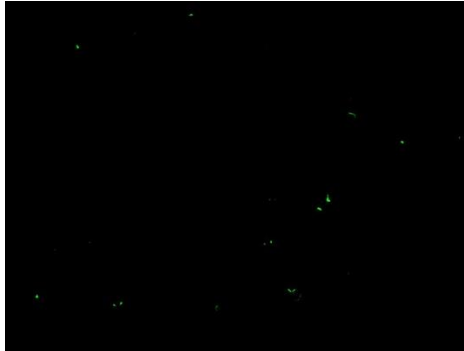
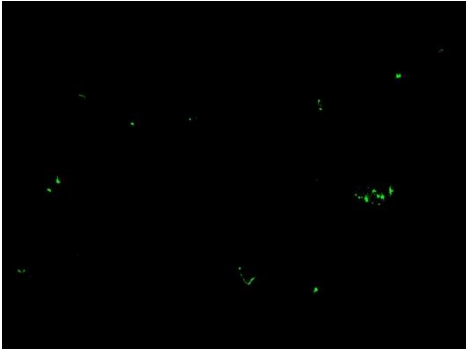
Spontaneous Death
- Percoll – Histopaque
- 2x cells, 24h



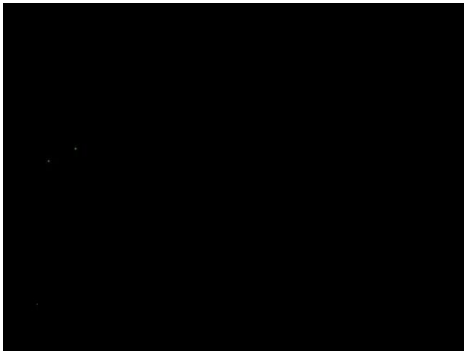
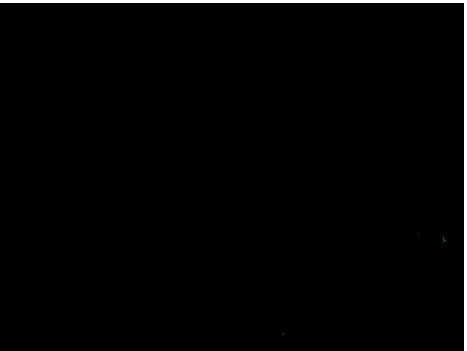
Spontaneous Death
- Percoll – Histopaque, 48h



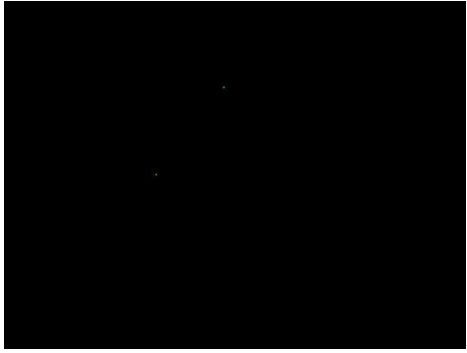
Spontaneous Death
- Percoll – Negative Selection, 24h



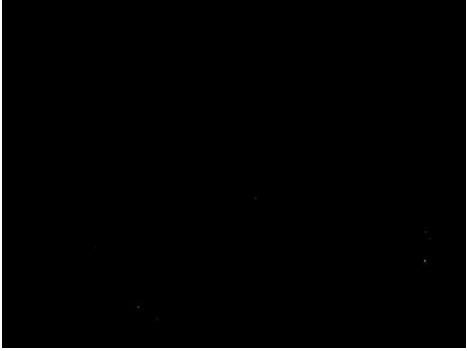
Spontaneous Death
- Percoll – Negative Selection
- 2x cells, 24h



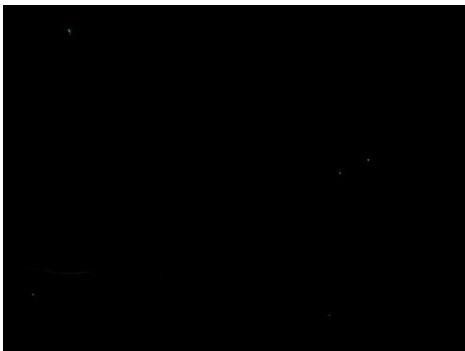
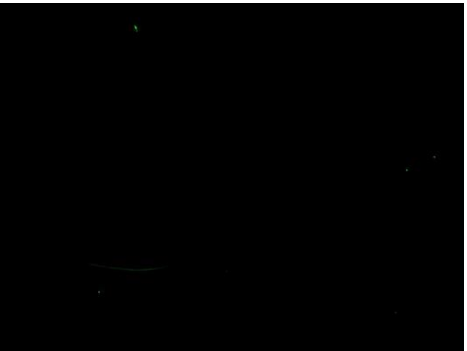
Spontaneous Death
- Percoll – Negative Selection
- 48h



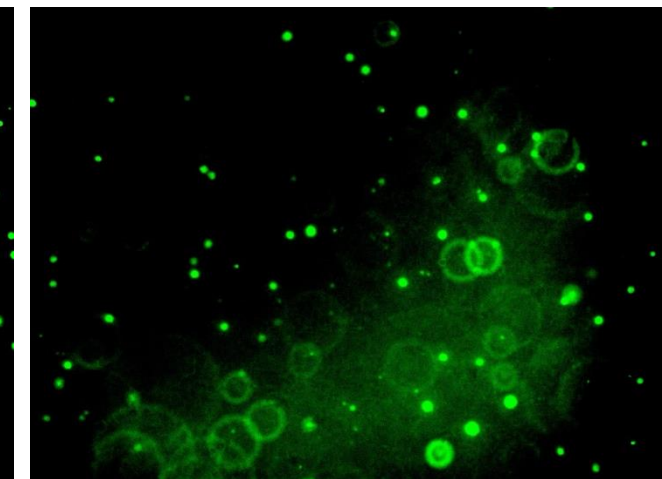
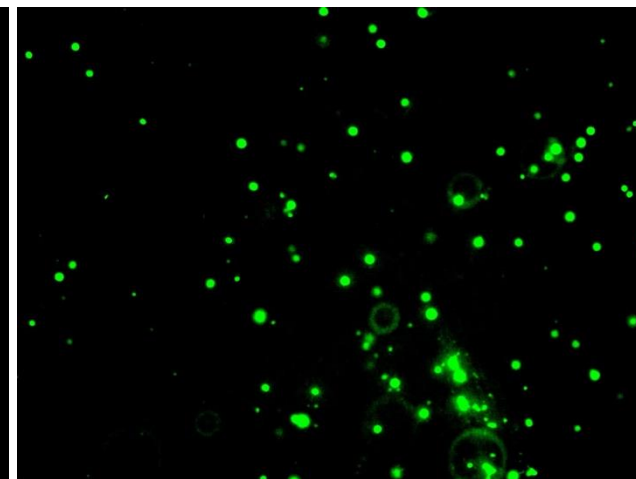
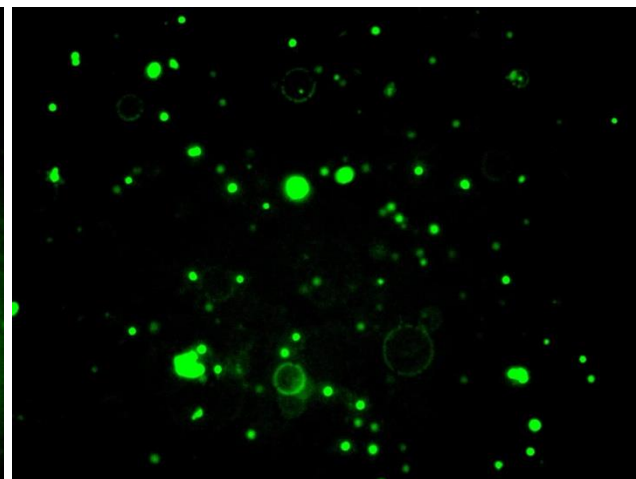
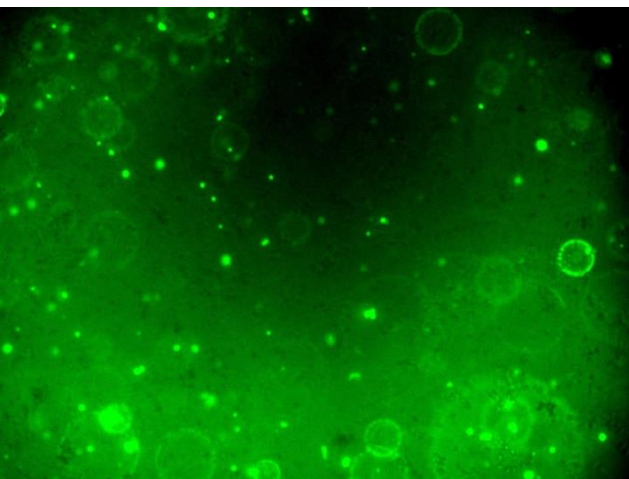
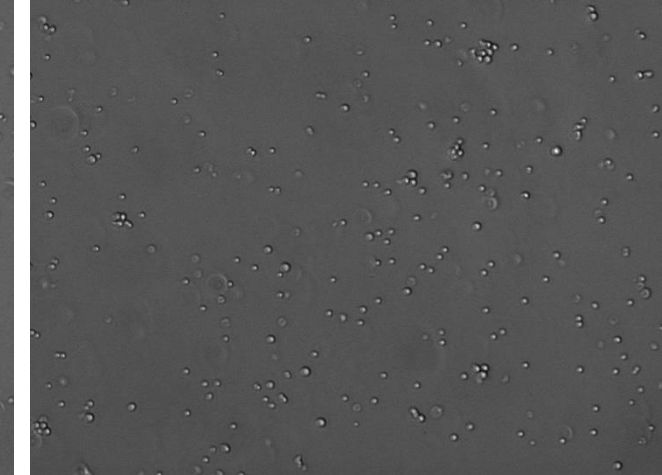
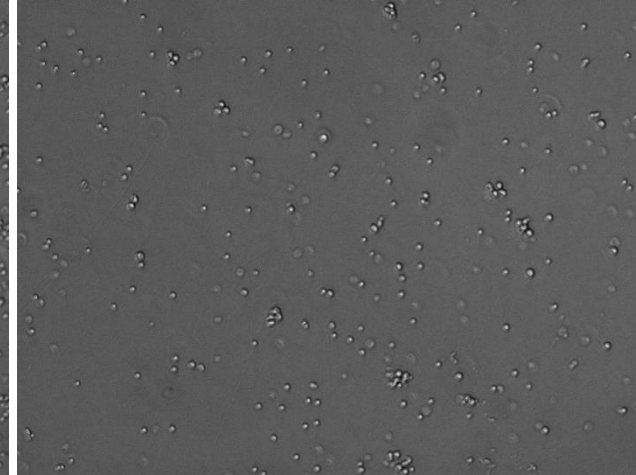
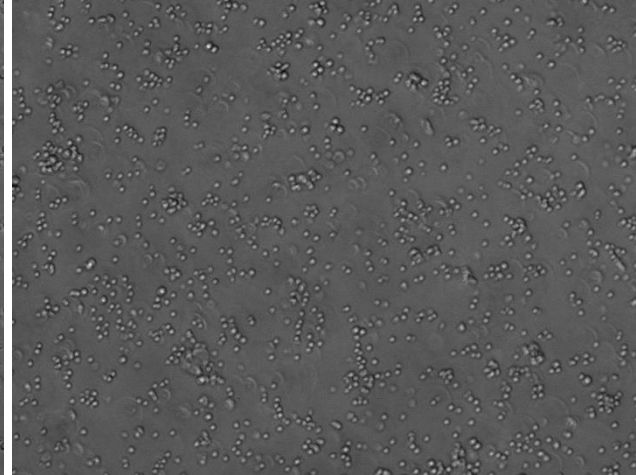
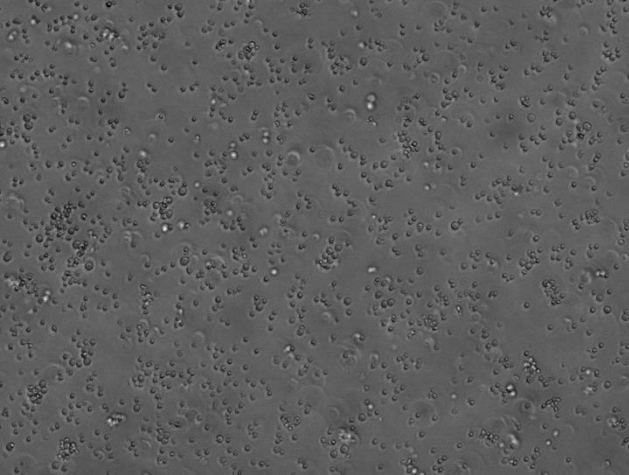
Spontaneous Death
- Percoll – Histopaque
- Negative Selection, 24h



Spontaneous Death
- Percoll – Histopaque
- Negative Selection
- 2x cells, 24h



Spontaneous Death
- Percoll – Histopaque
- Negative Selection
- 2x cells, 48h

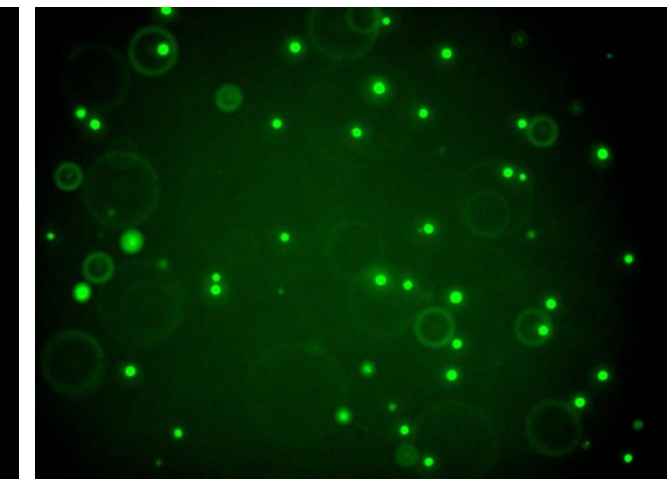
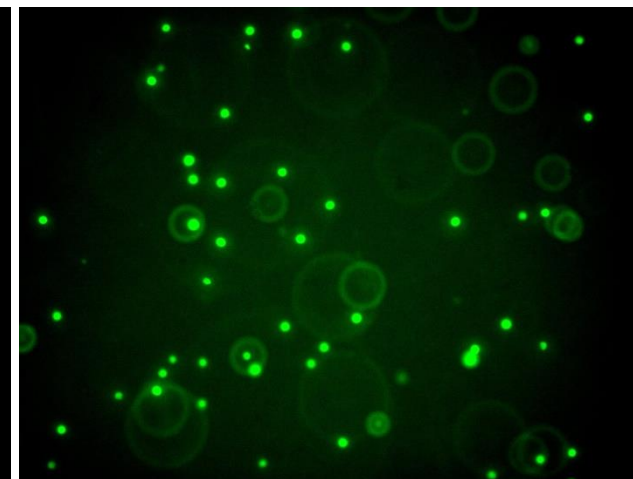
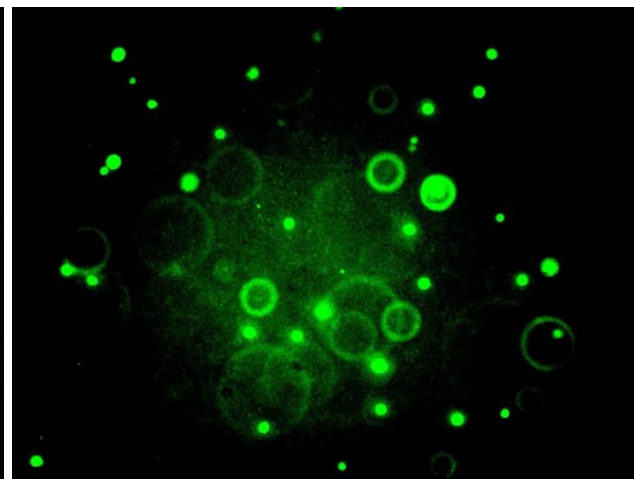
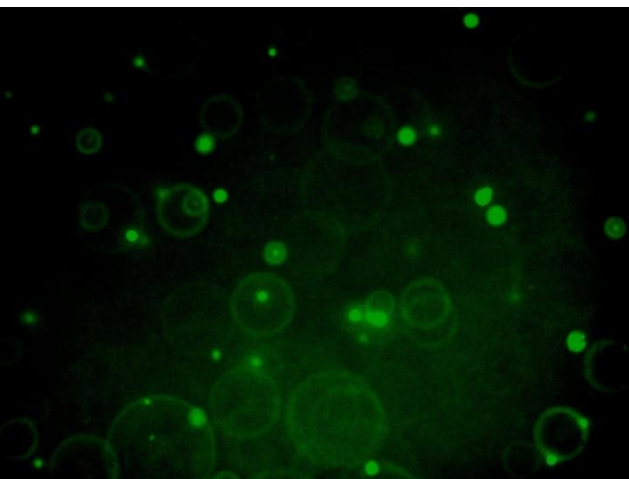
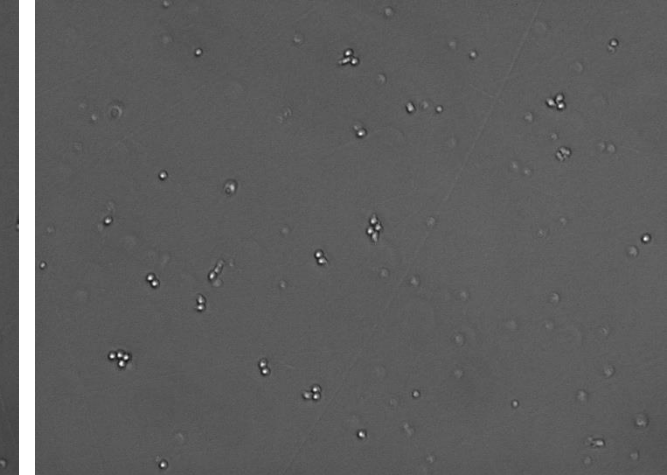
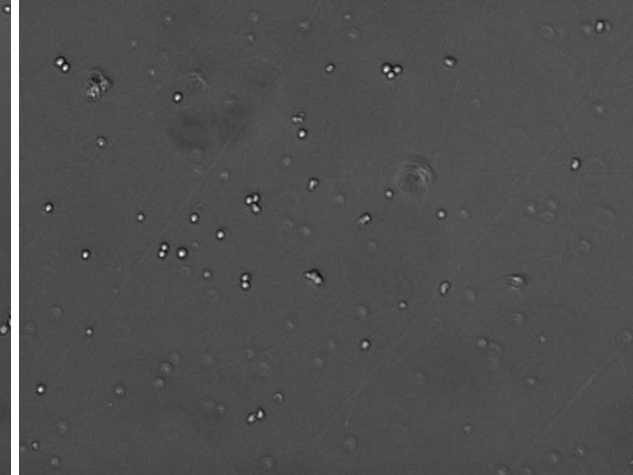
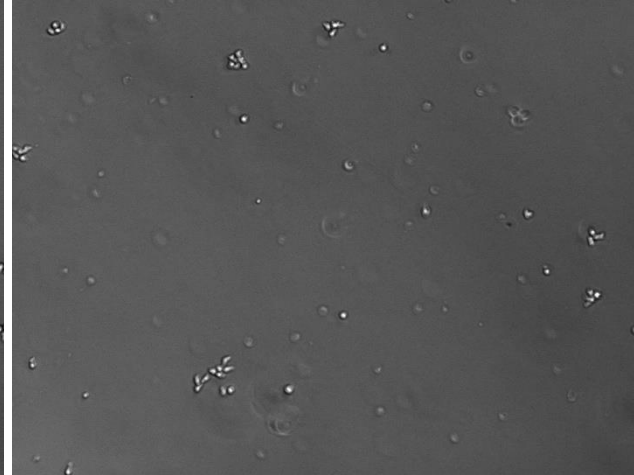
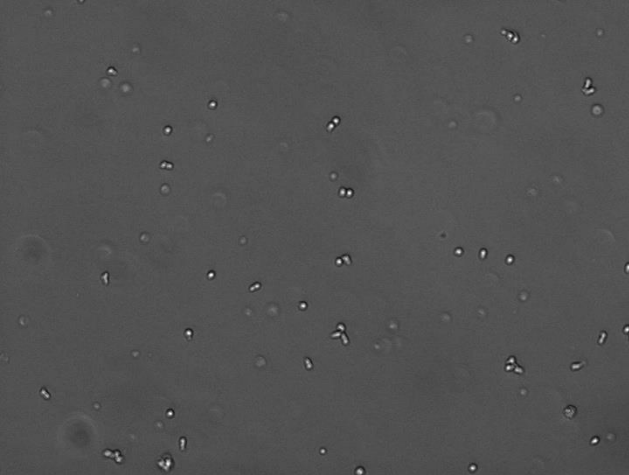


Percoll

Percoll

Percoll - Histopaque

Percoll - Histopaque

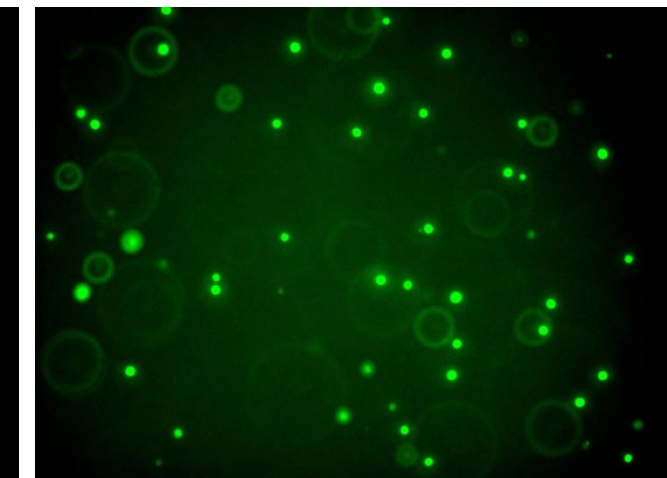
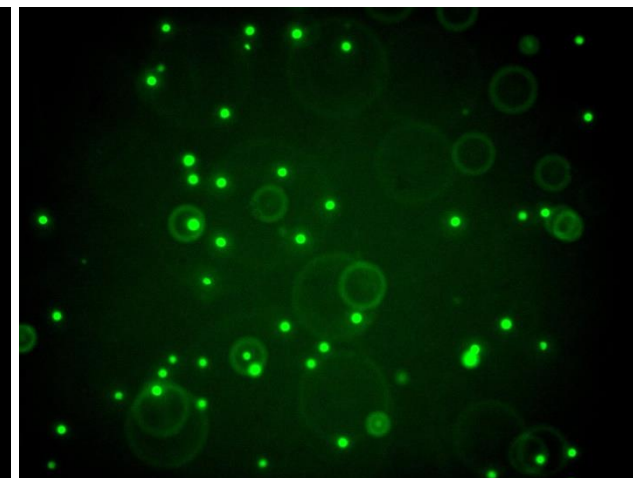
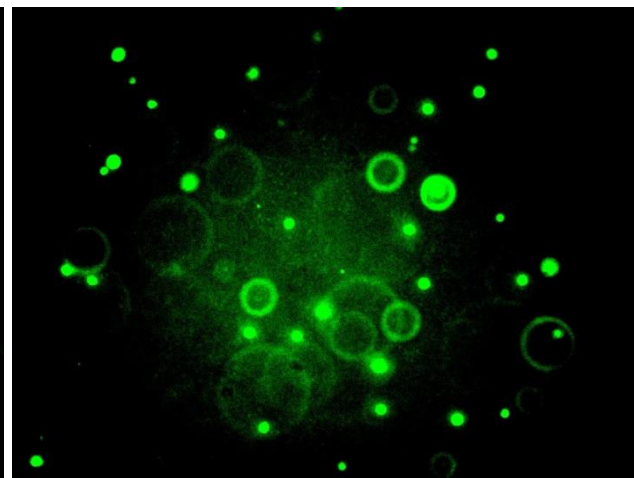
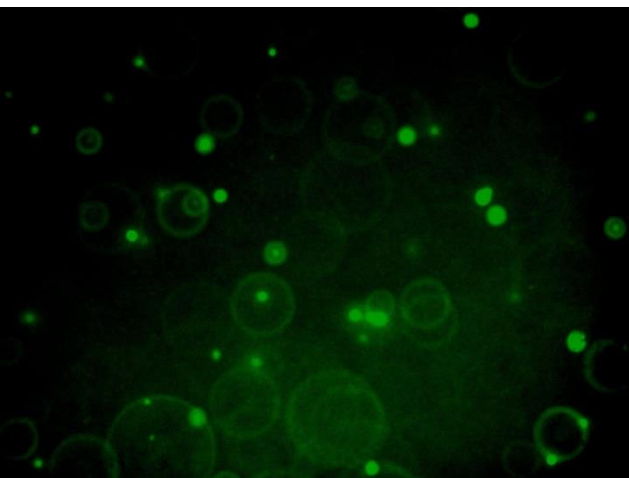
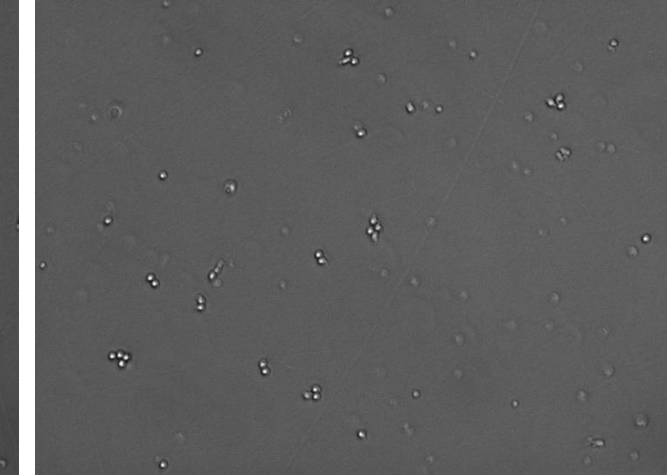
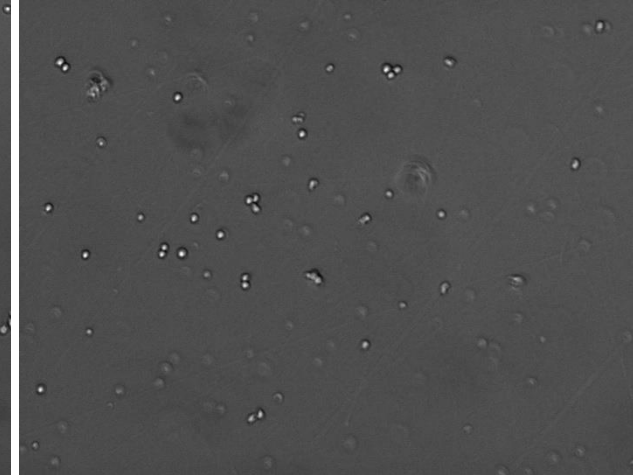
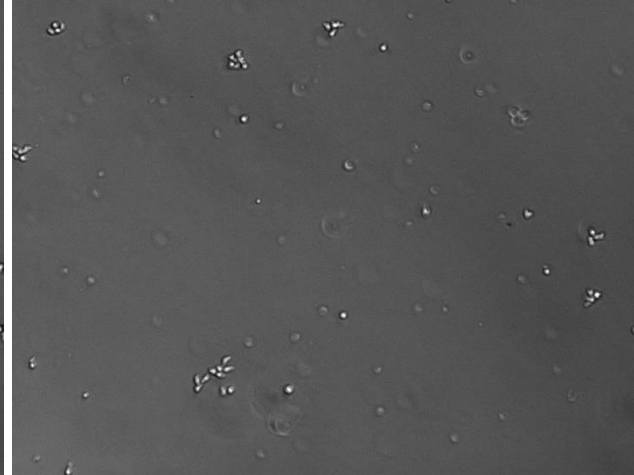
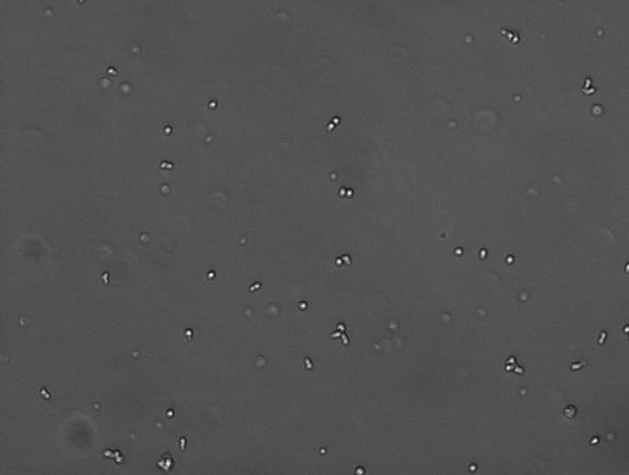


Percoll – Negative Selection

Percoll – Negative Selection

Percoll – Histopaque
- Negative Selection

Percoll – Histopaque
- Negative Selection

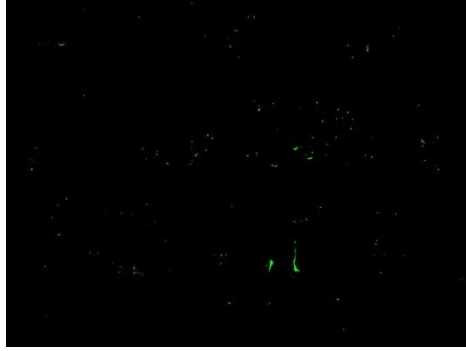


10

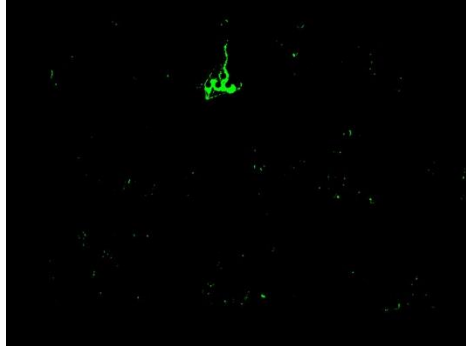
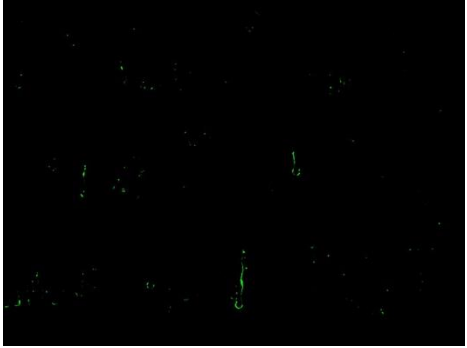
Percoll – Negative Selection

Percoll – Histopaque
- Negative Selection

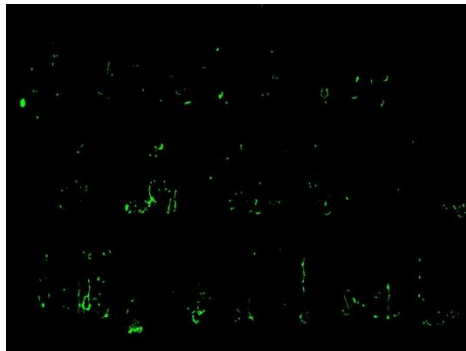
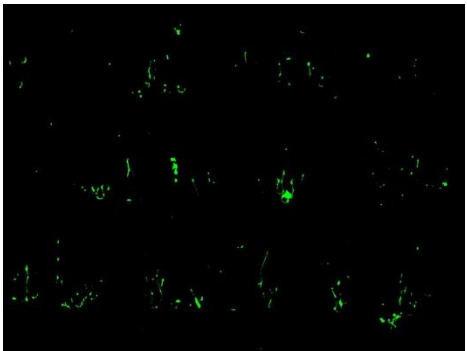
Percoll – Histopaque
- Negative Selection



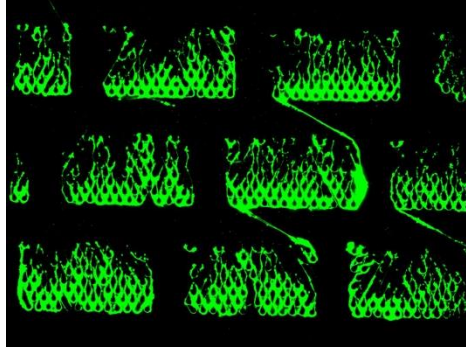
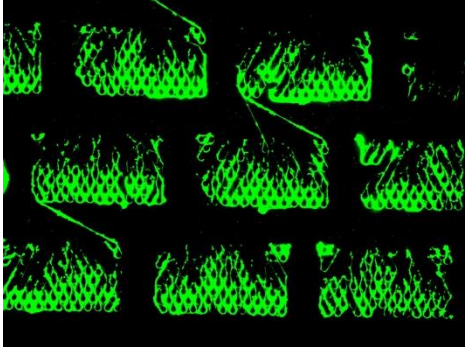
S. Aureus
ATCC 10390



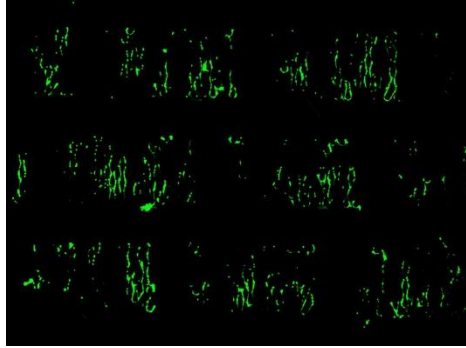
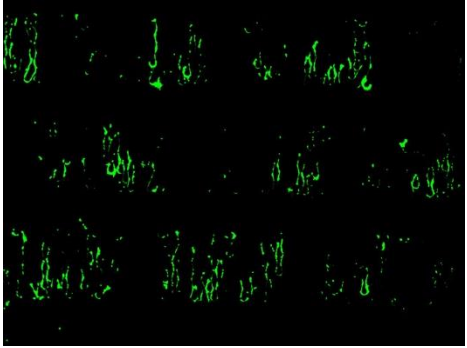
S. Aureus
ATTC BAA-1717



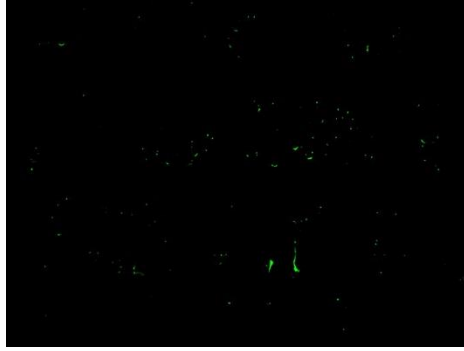
K12
ATCC MG1655 700926



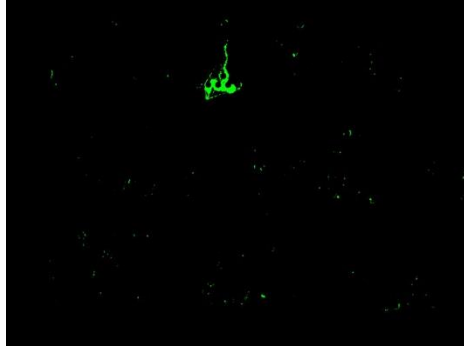
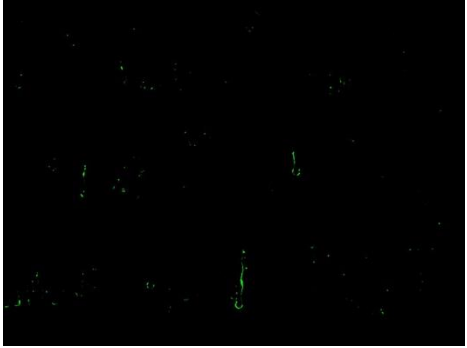
Bacillus cereus
ATCC 14579



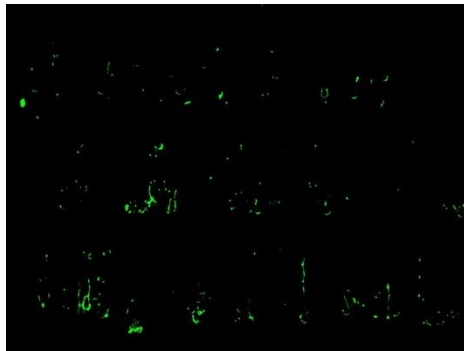
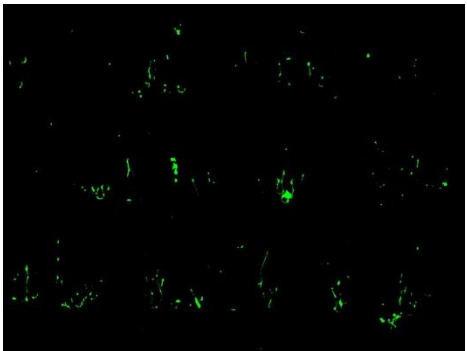
A7
ATCC 700928



S. Aureus
ATCC 10390



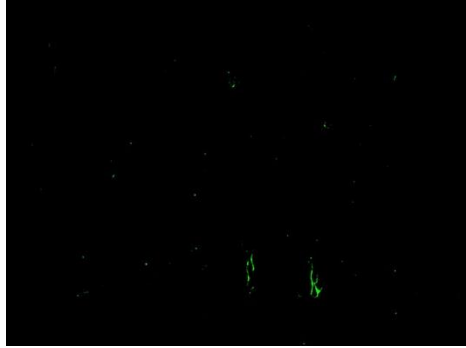
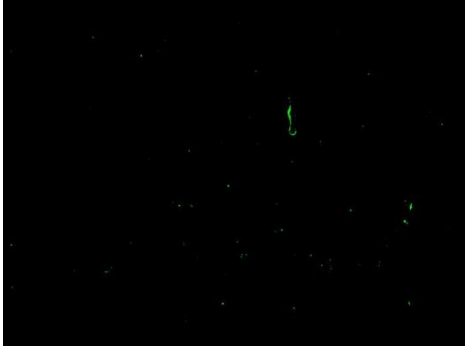
S. Aureus
ATTC BAA-1717



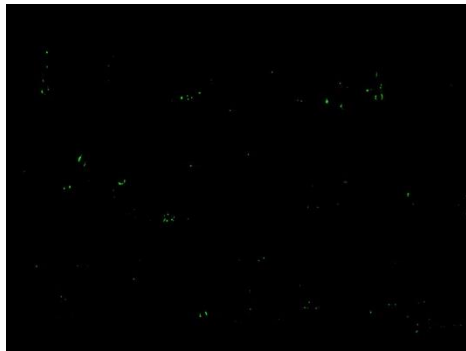
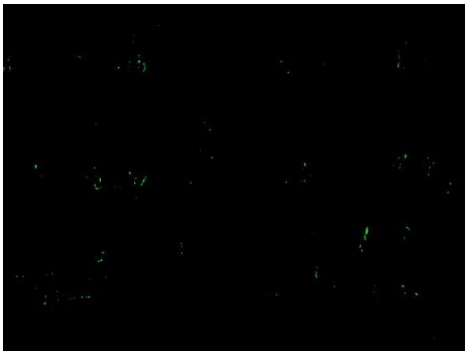
K12
ATCC MG1655 700926



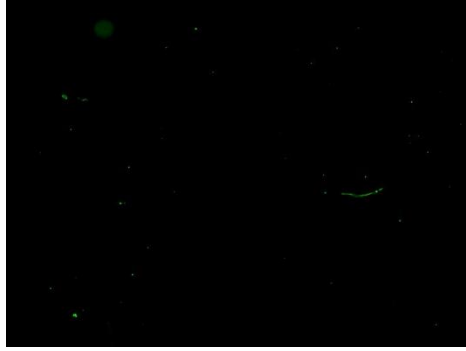
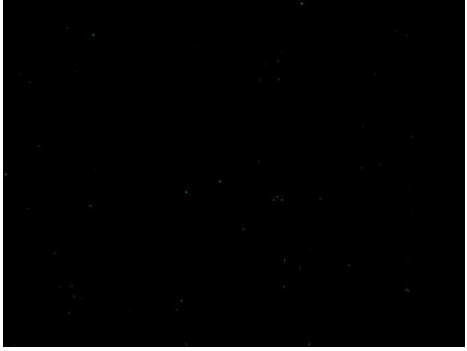
S. Aureus
ATCC 10390



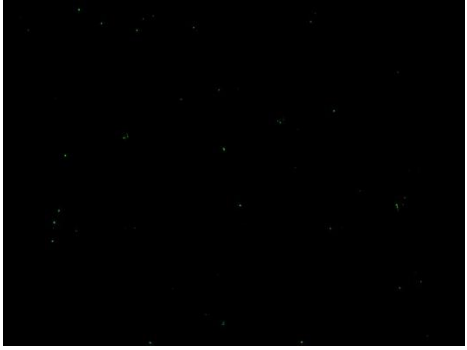
S. Aureus
ATTC BAA-1717



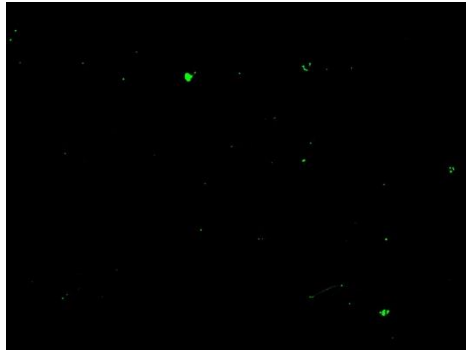
K12
ATCC MG1655 700926



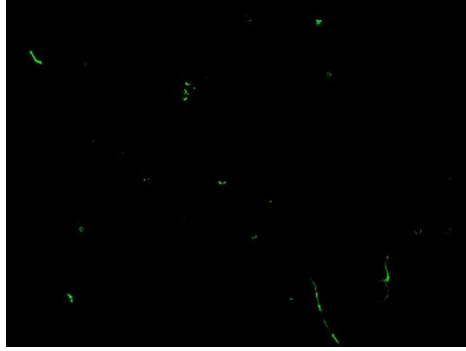
S. Aureus
ATCC 10390



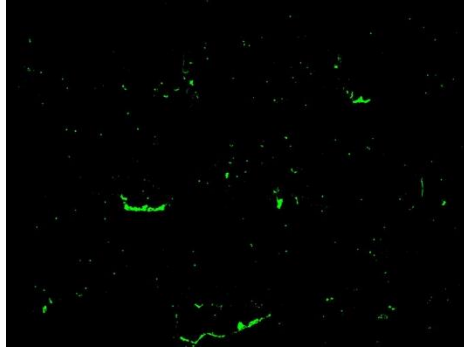
S. Aureus
ATTC BAA-1717



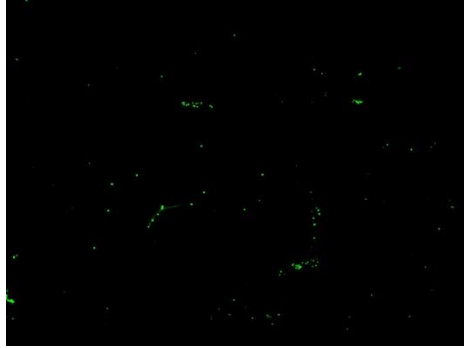
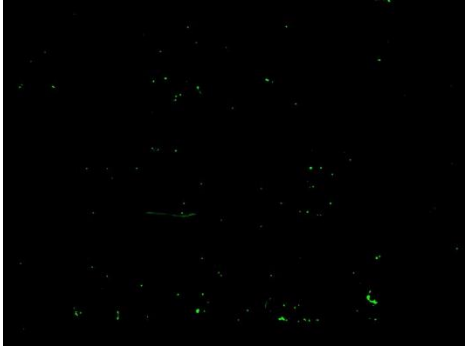
K12
ATCC MG1655 700926



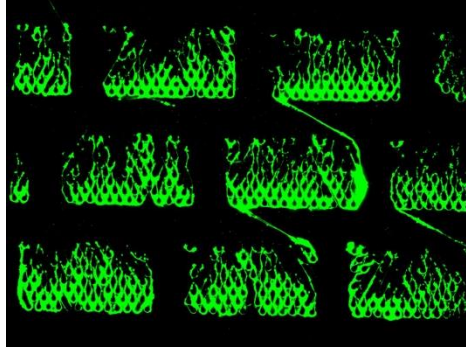
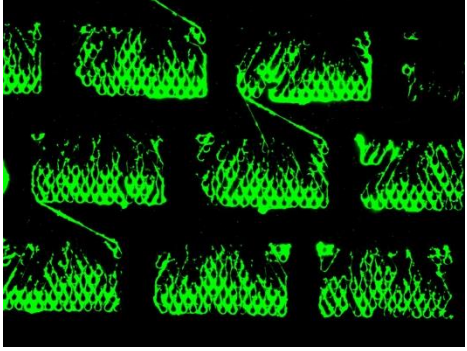
S. Aureus
ATTC BAA-1717



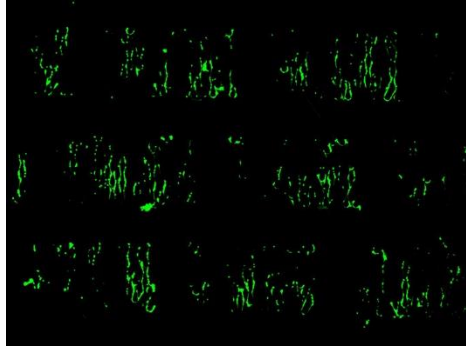
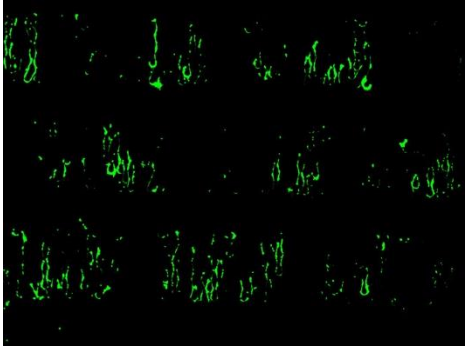
S. Aureus
ATCC 10390



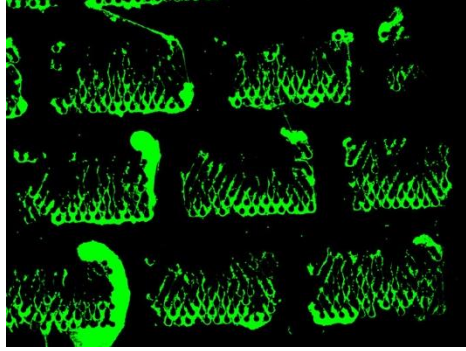
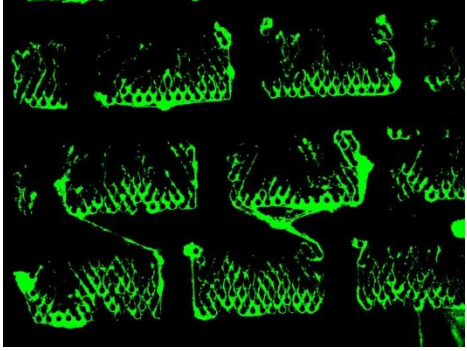
S. Aureus
ATTC BAA-1717



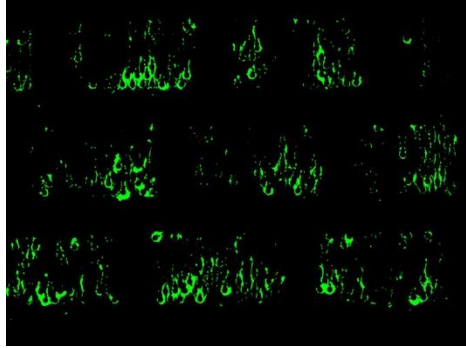
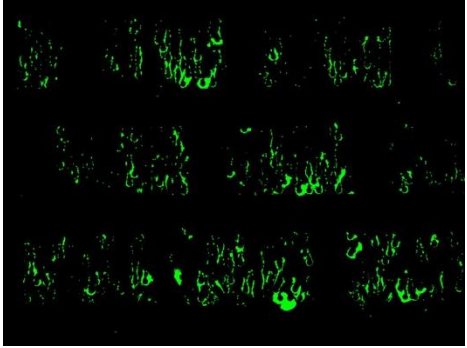
Bacillus cereus
ATCC 14579



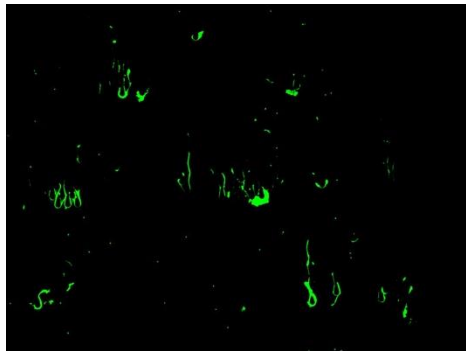
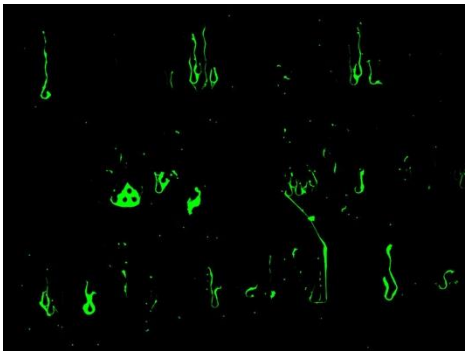
A7
ATCC 700928



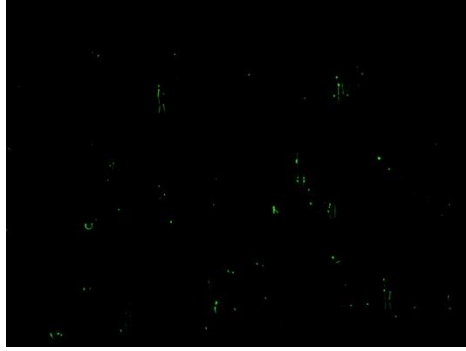
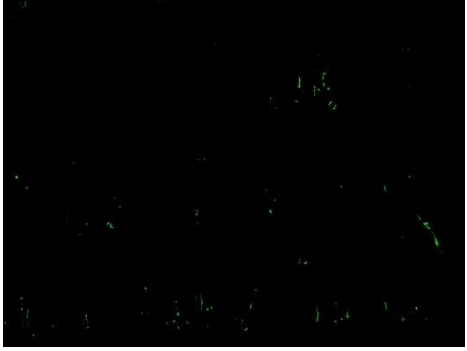
Bacillus cereus
ATCC 14579



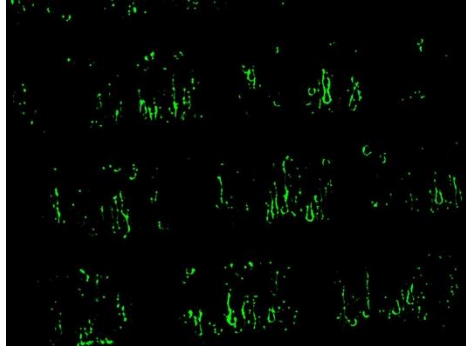
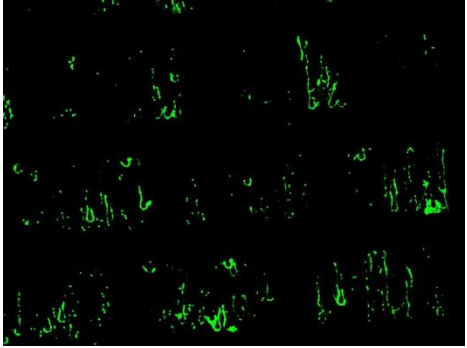
A7
ATCC 700928



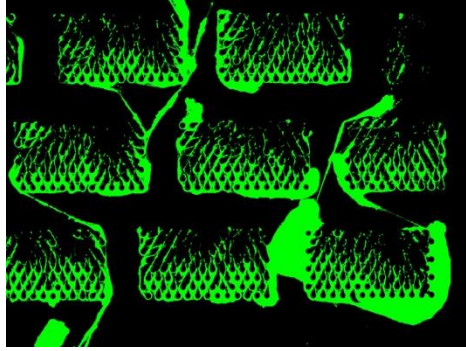
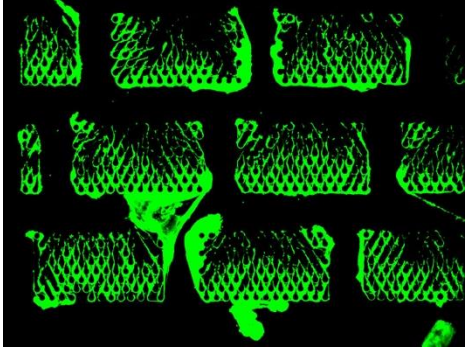
Triton



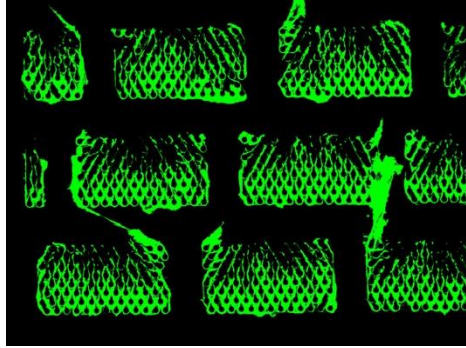
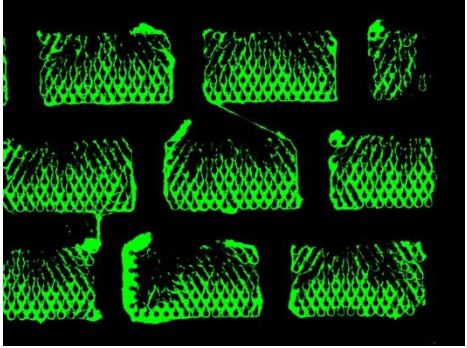
Blank



A7
ATCC 700928



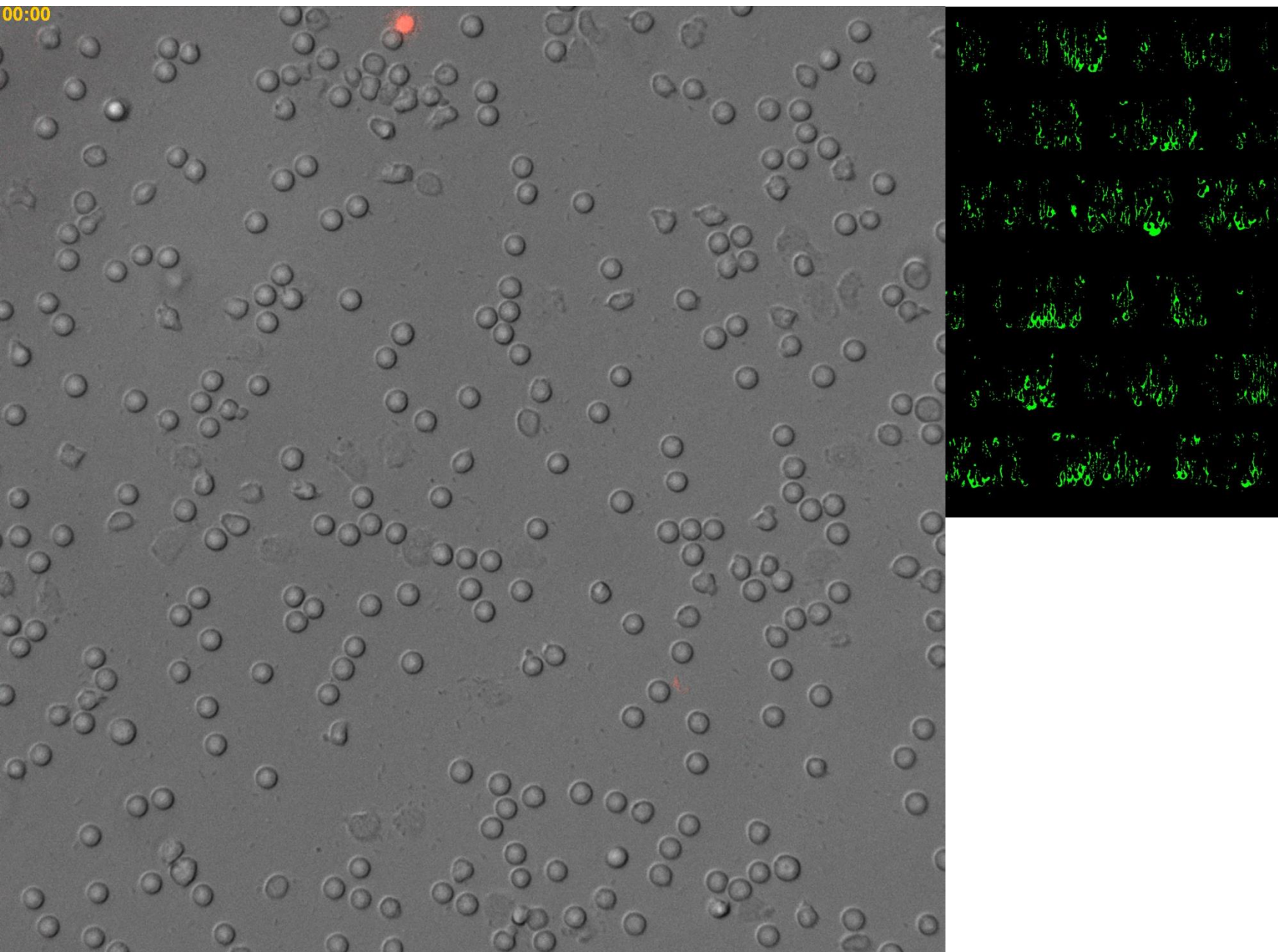
Bacillus cereus
ATCC 14579



Ionomycin

2025-07-15

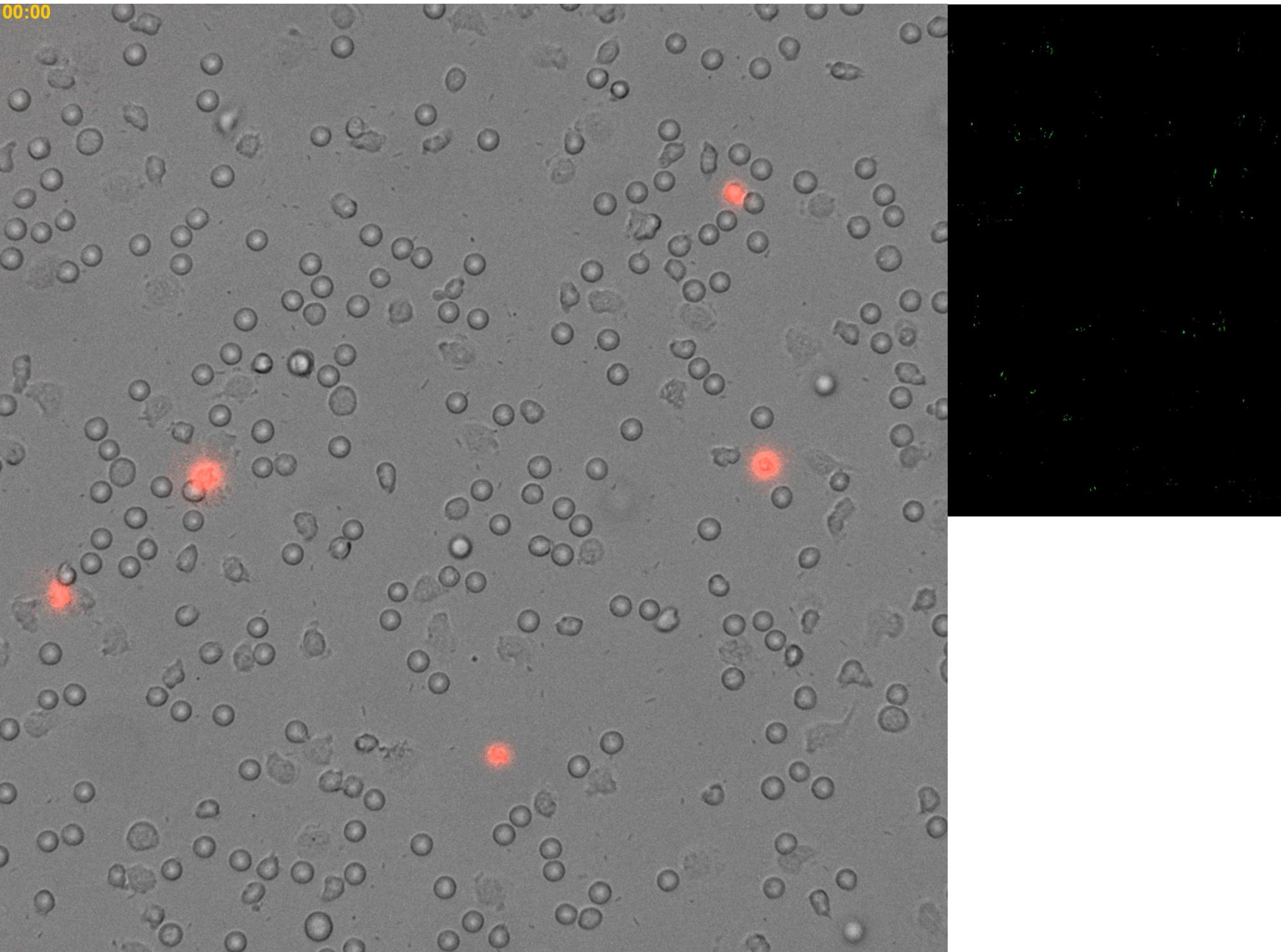
A7



2025-07-15

K12

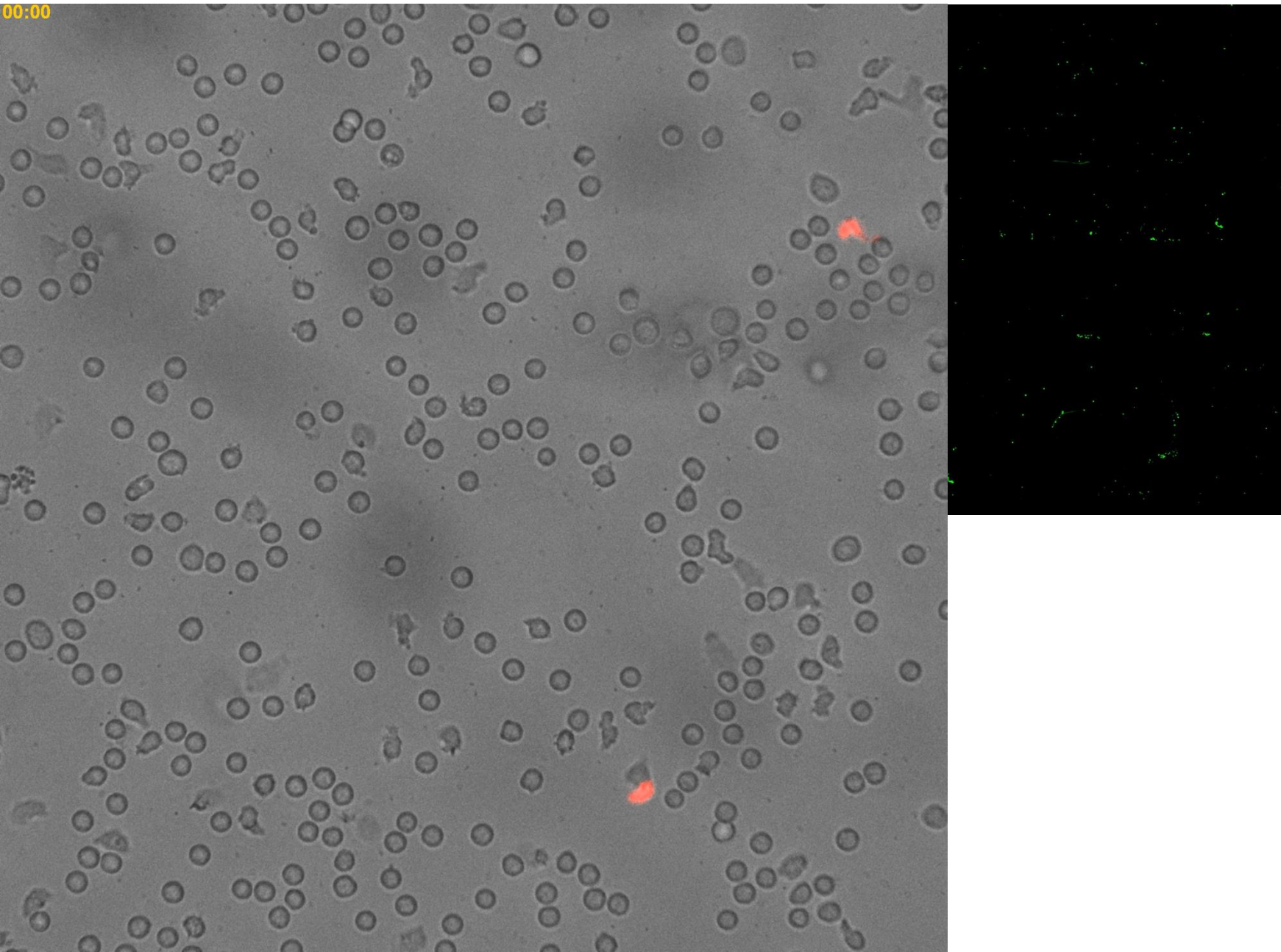
00:00



2025-07-15

Sa

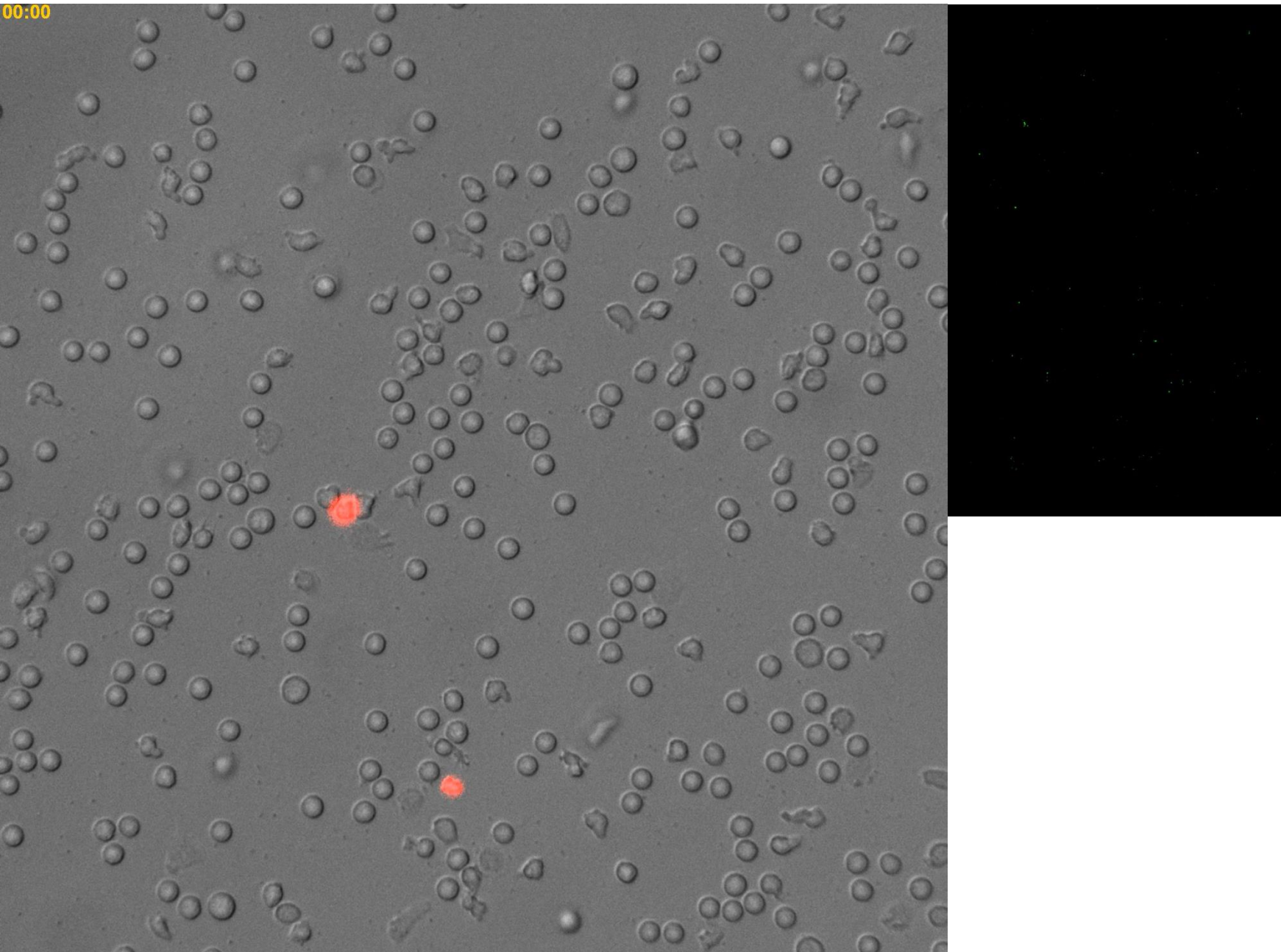
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2025-07-15

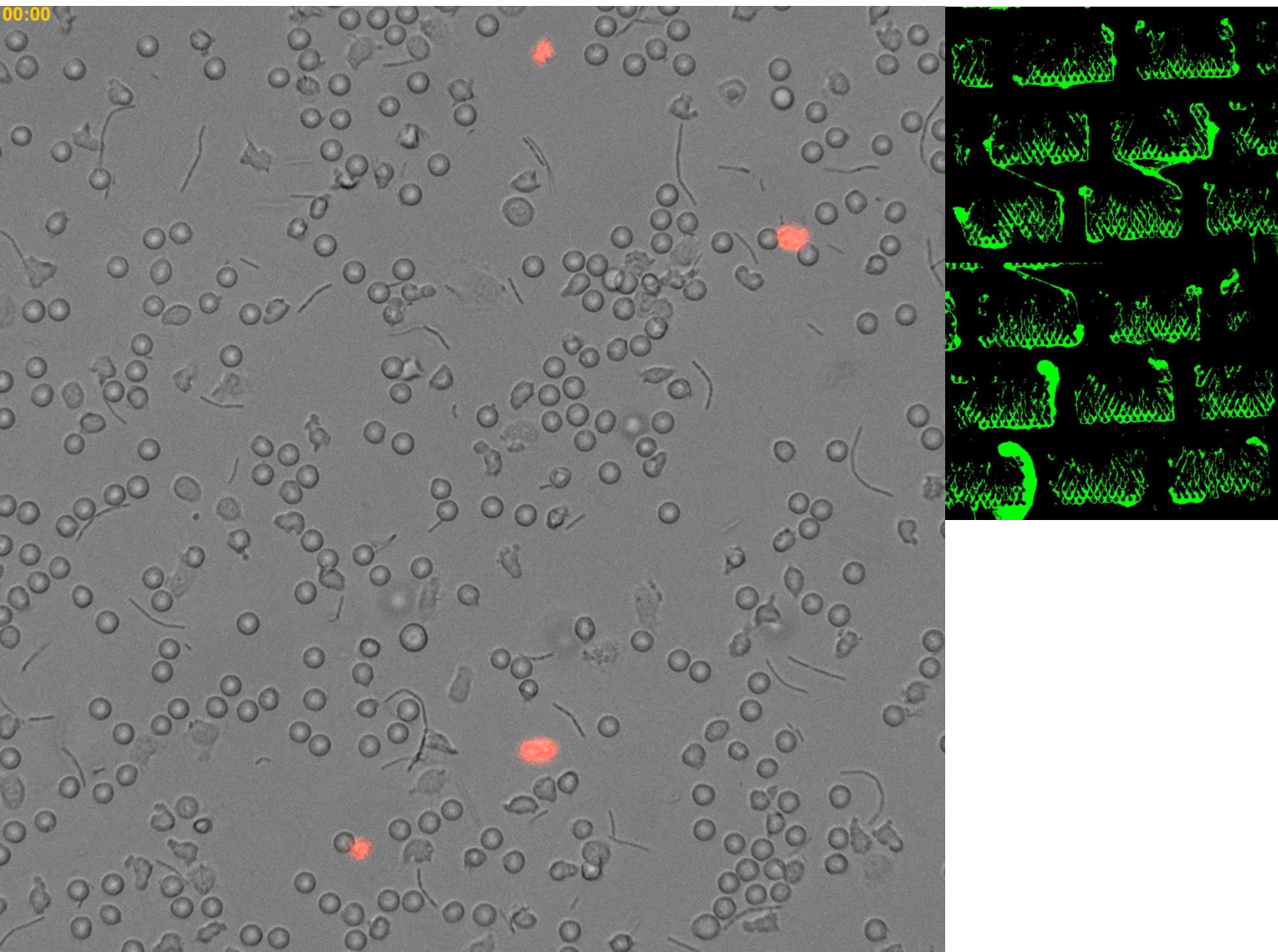
10390

00:00



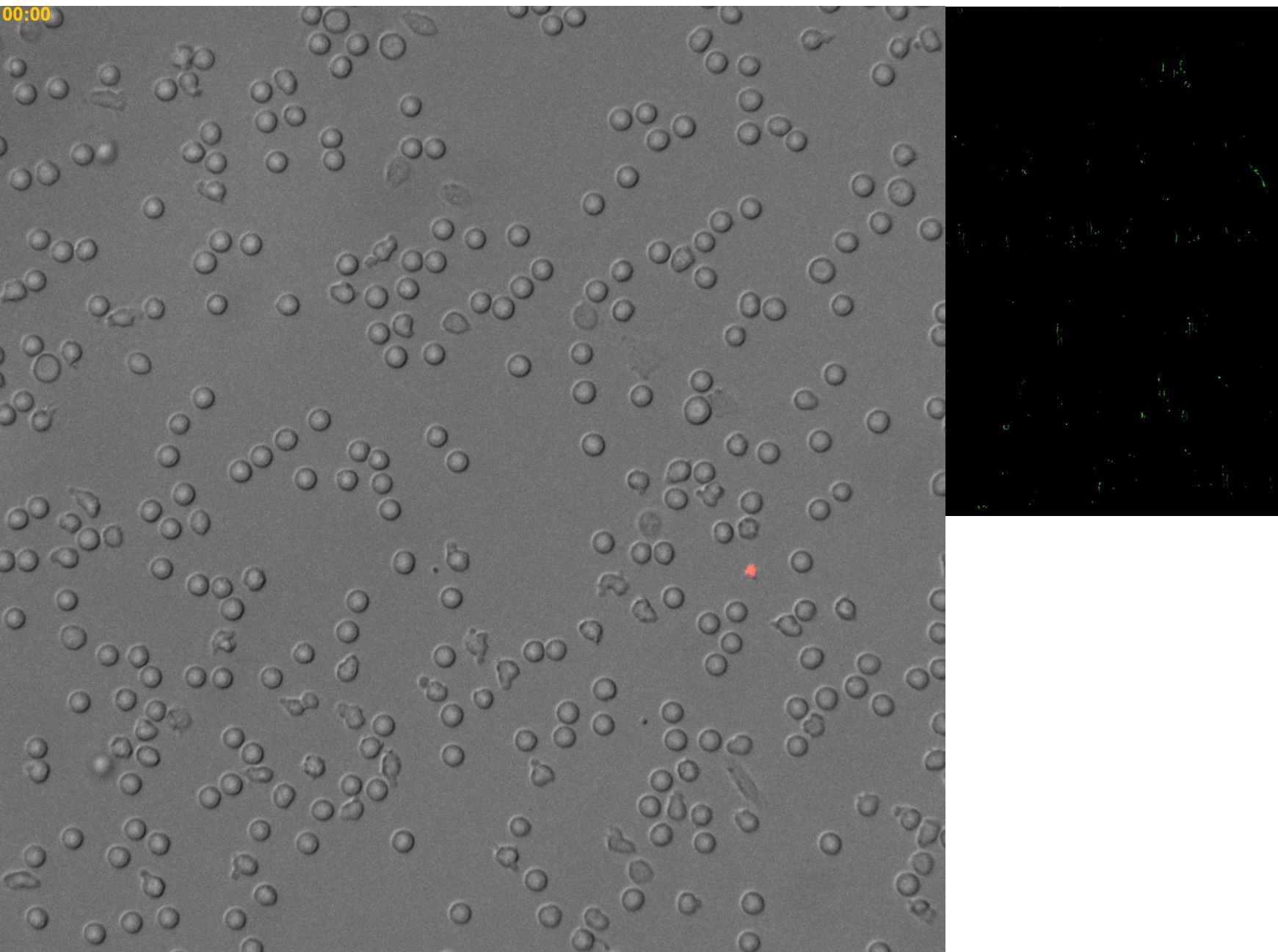
2025-07-15

Bacillus



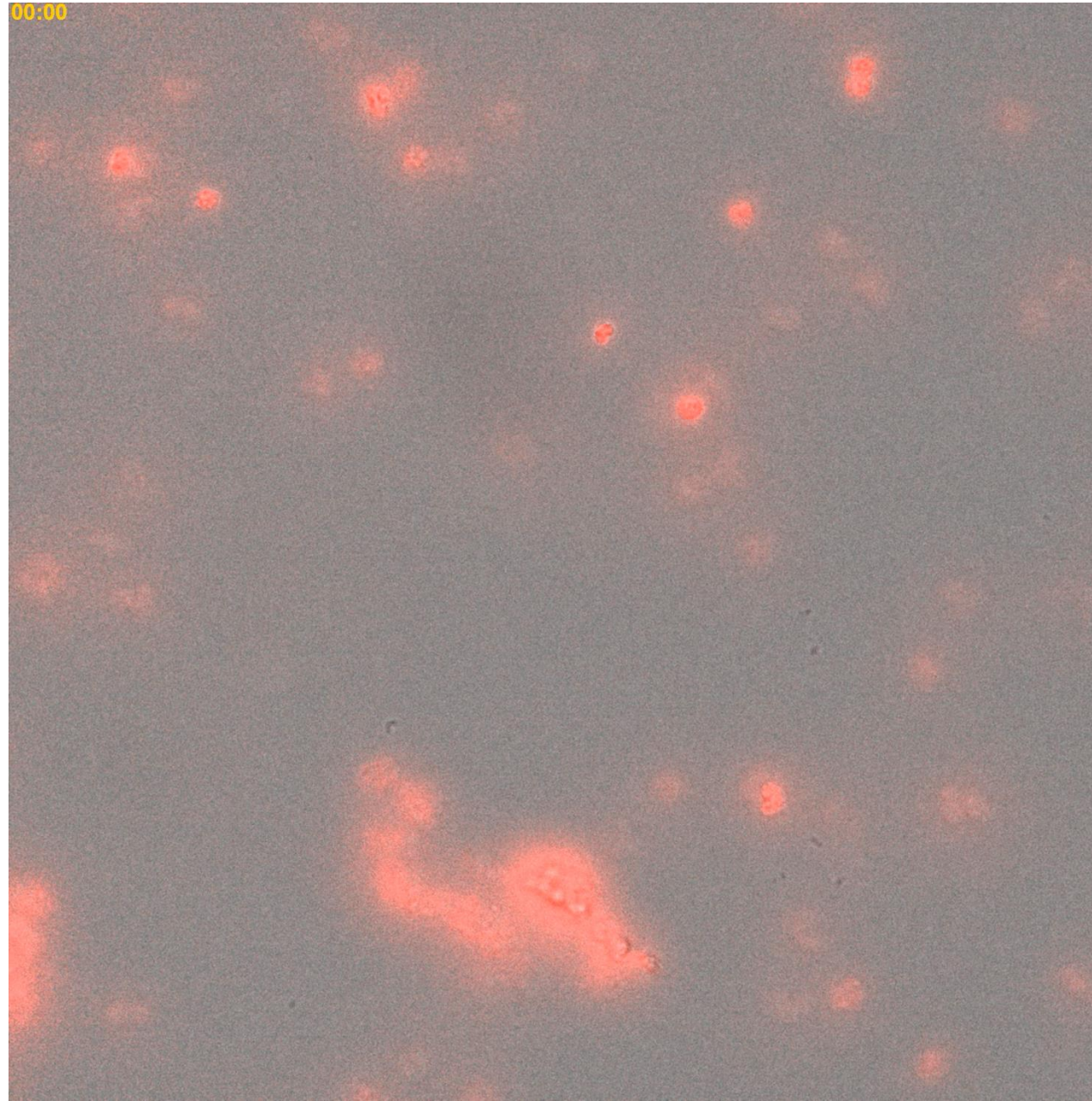
2025-07-15

Blank (8h NET)



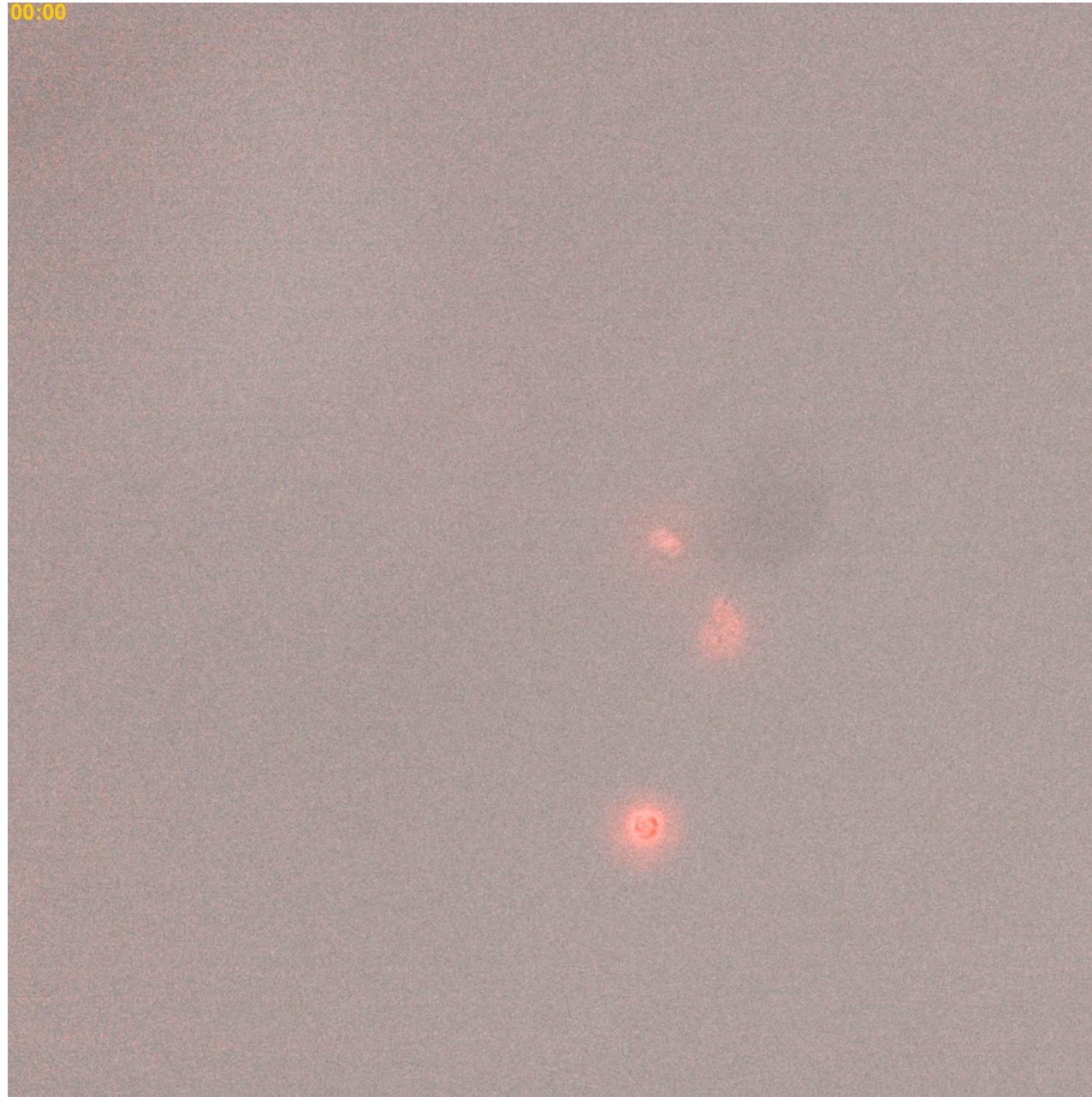
2025-07-15

Triton



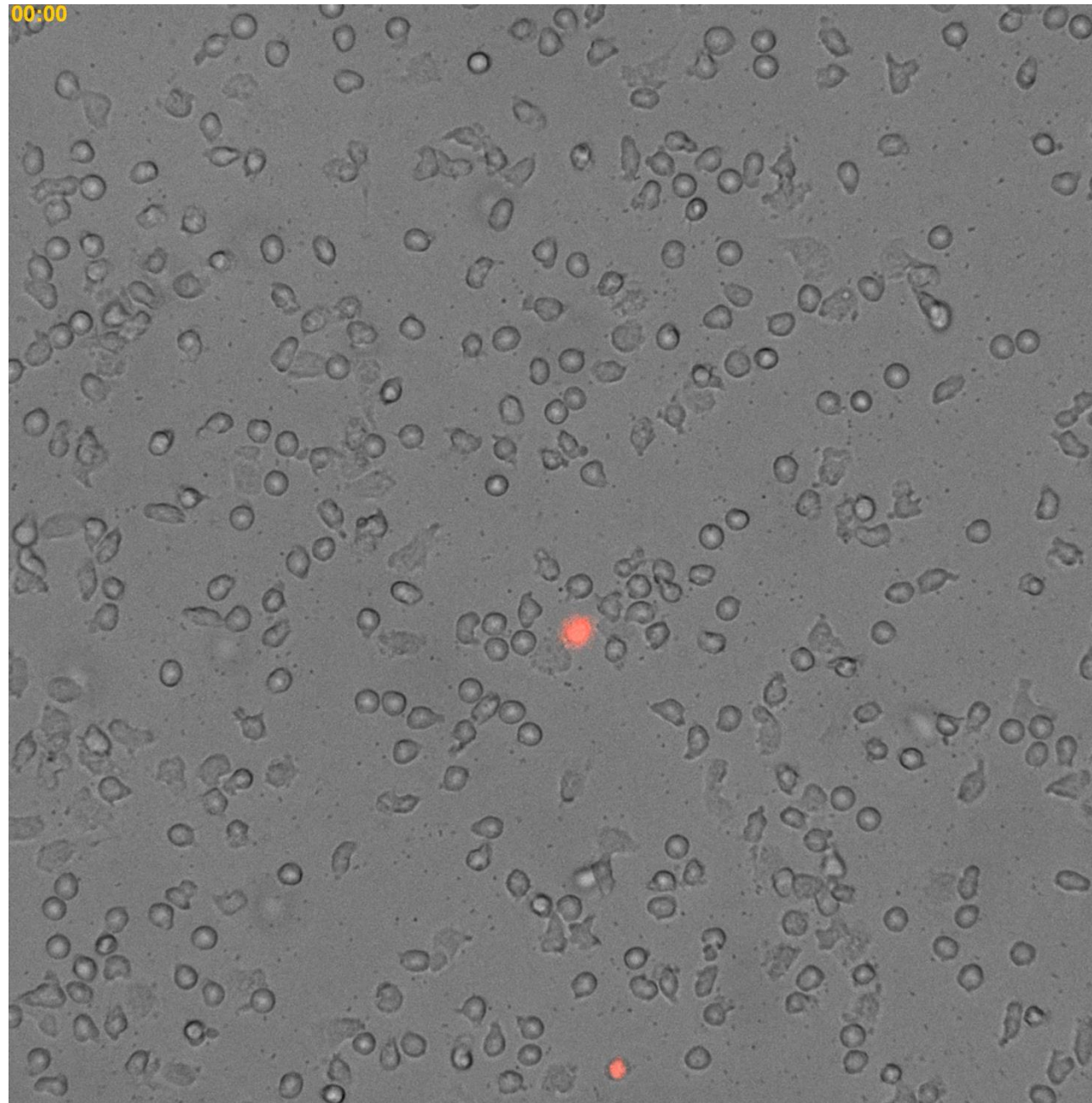
2025-07-15

Lysis Buffer



2025-07-17

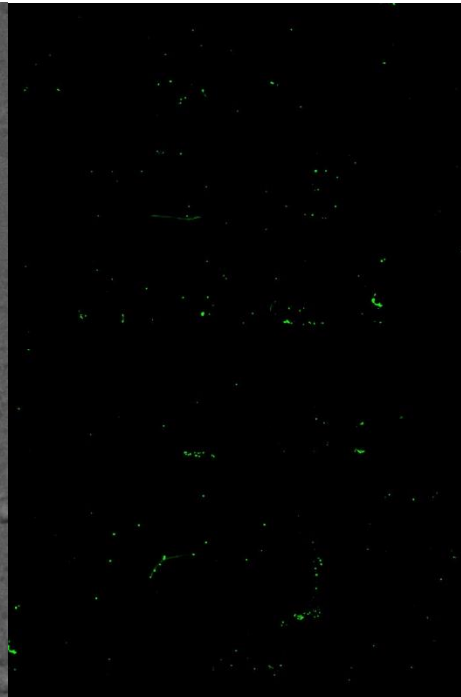
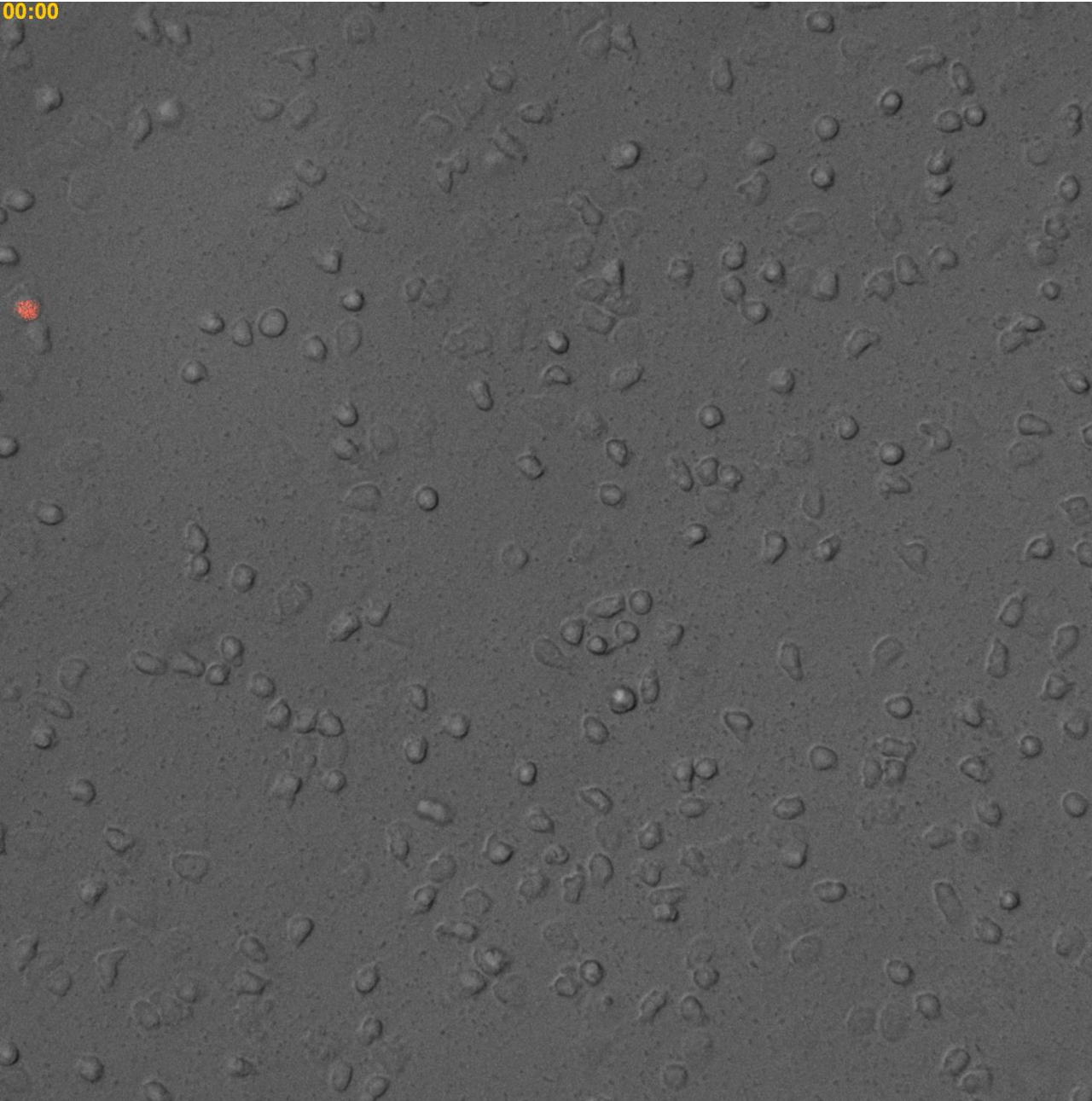
Sa MOI=50



2025-07-17

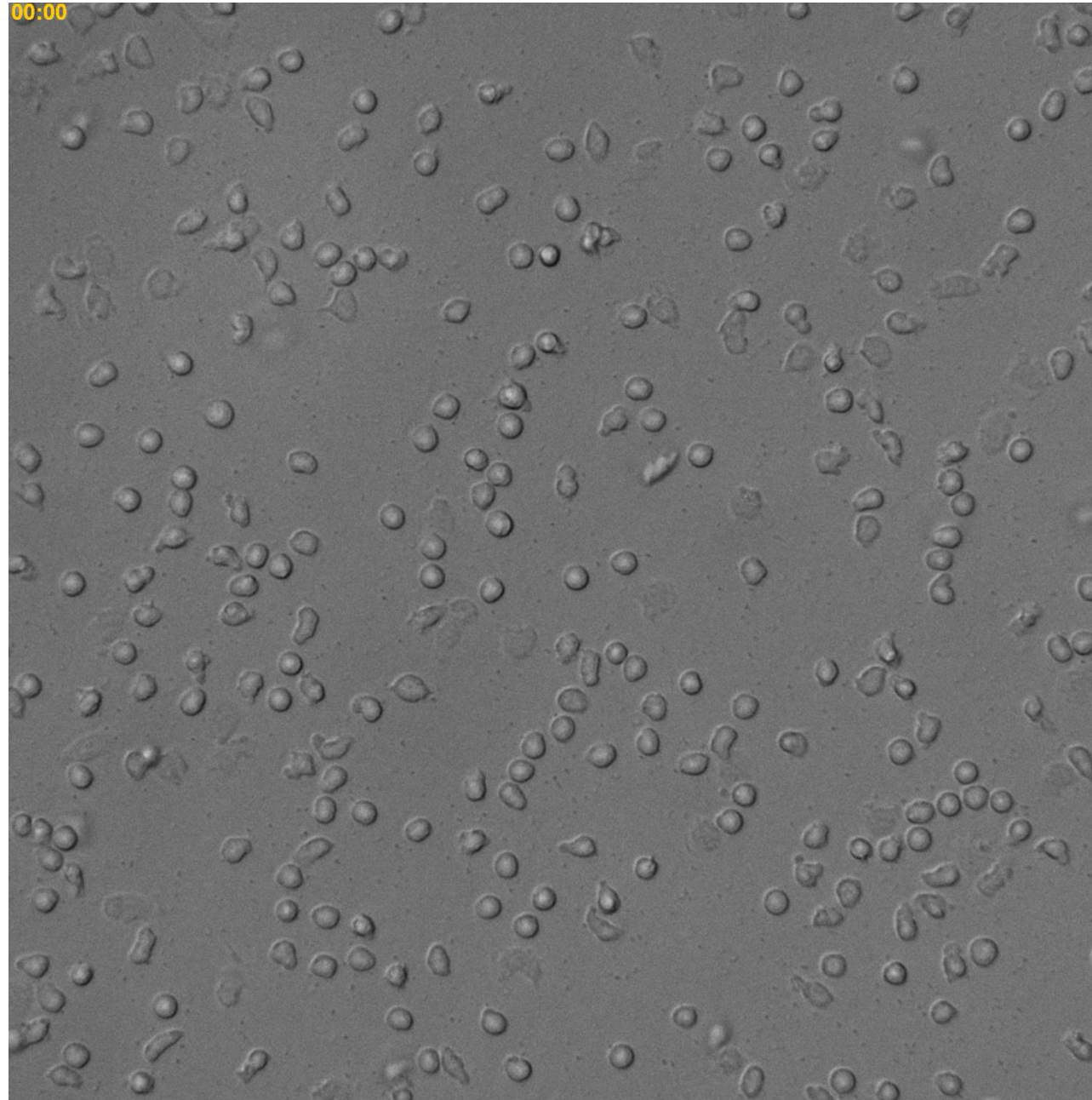
Sa MOI=200

00:00



2025-07-17

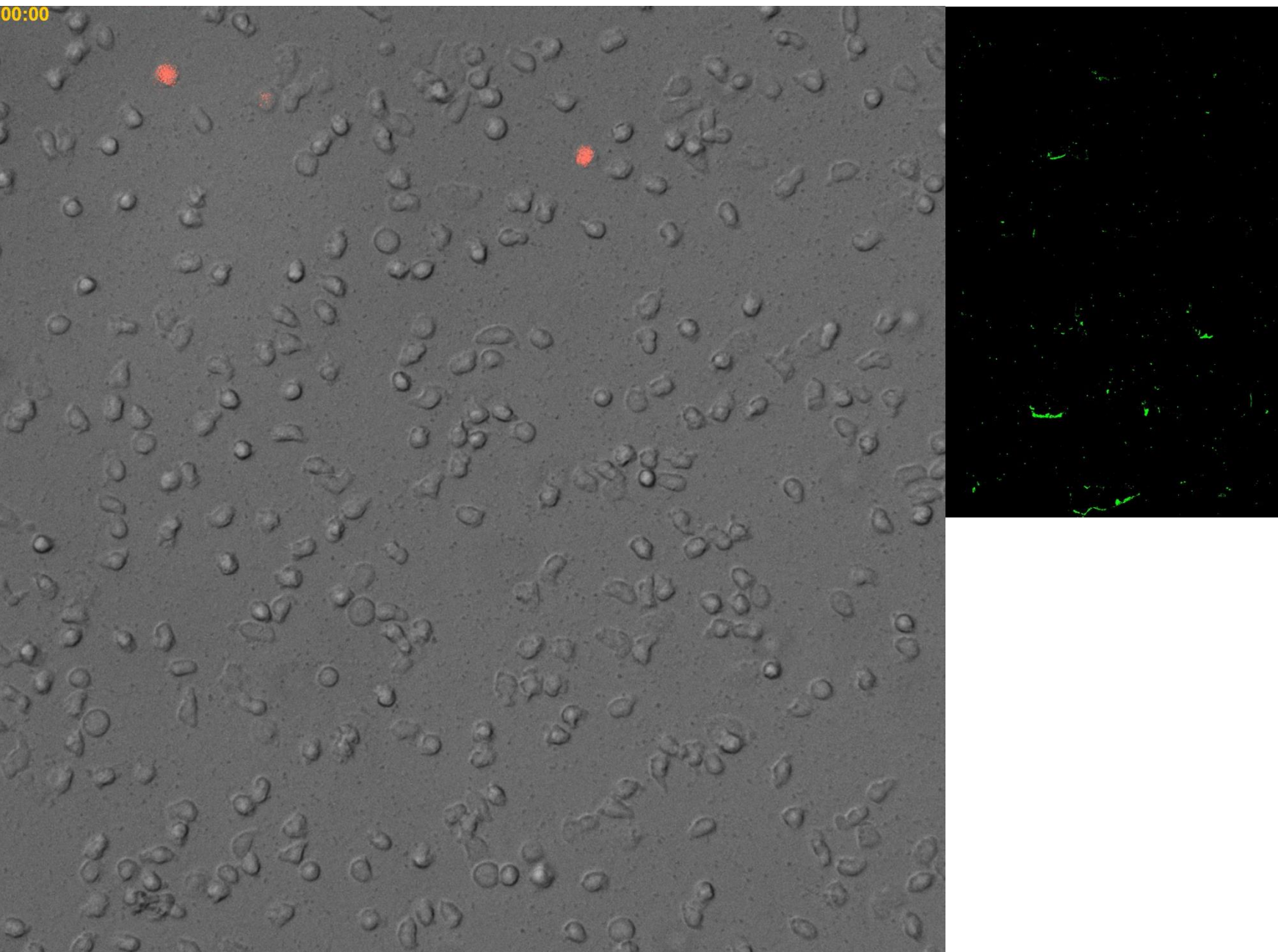
10390 MOI=35



2025-07-17

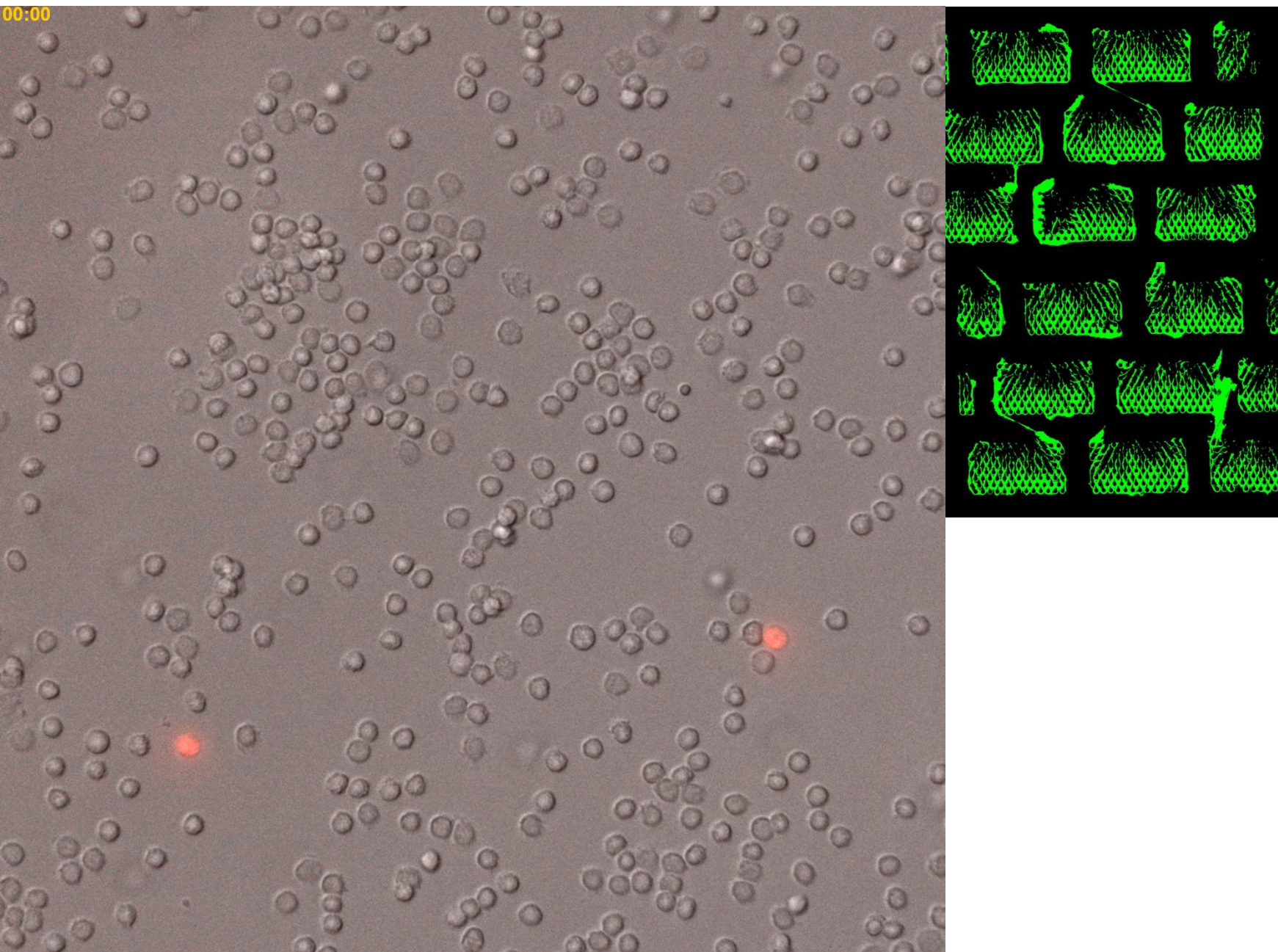
10390 MOI=140

00:00



2025-07-17

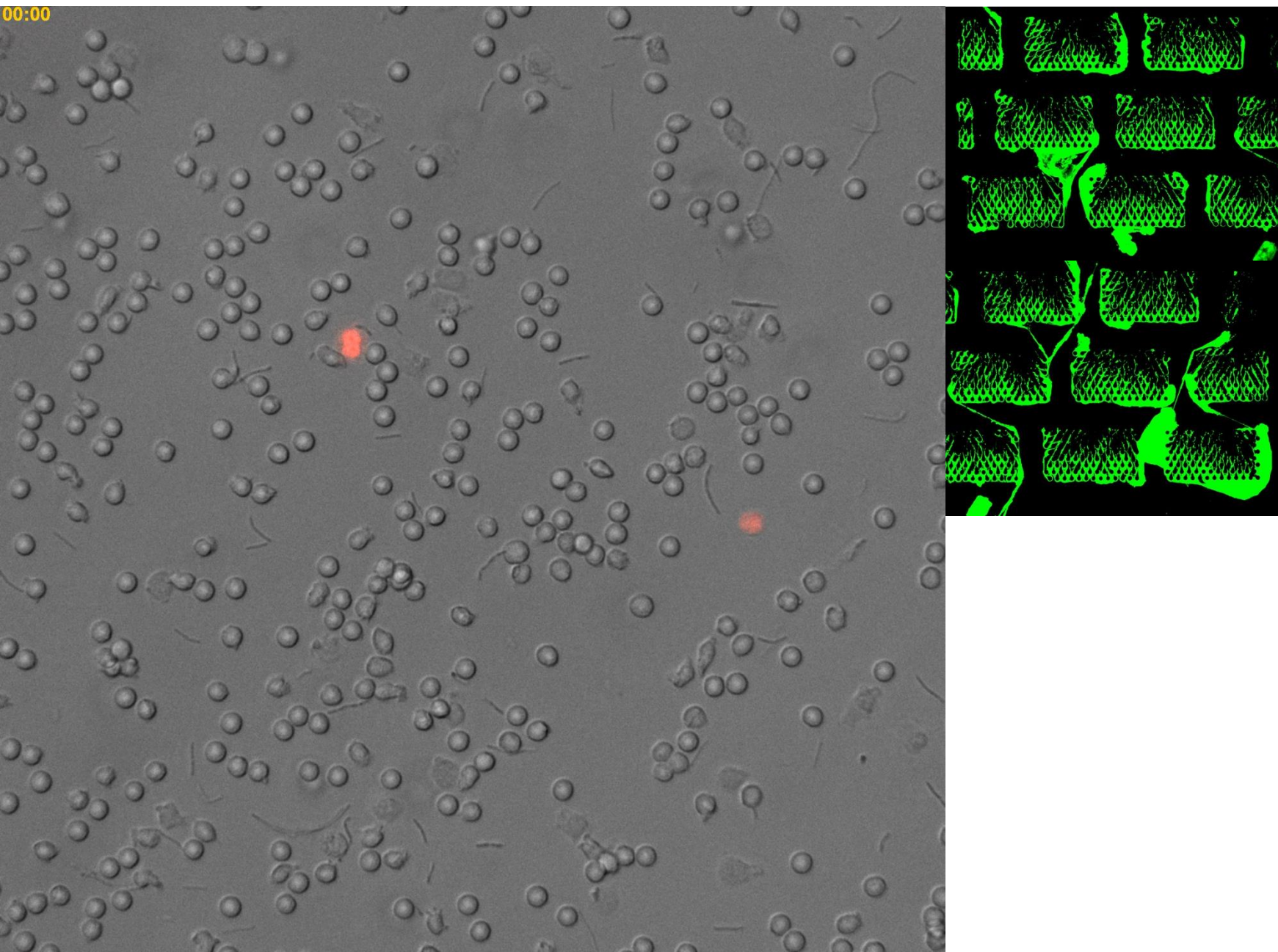
Ionomycin



2025-07-17

Bacillus MOI=5

00:00



Protocol Change

After we get our NETs from ionomycin:

We add 400 μL of bacteria at a concentration of $5 \times 10^8/\text{mL}$ through the device.

Then we wash it with 200 μL of PBS, repeated 8 times.

Add 50 μL of DNase through the device.

Incubate the device at 37°C for 30 minutes.

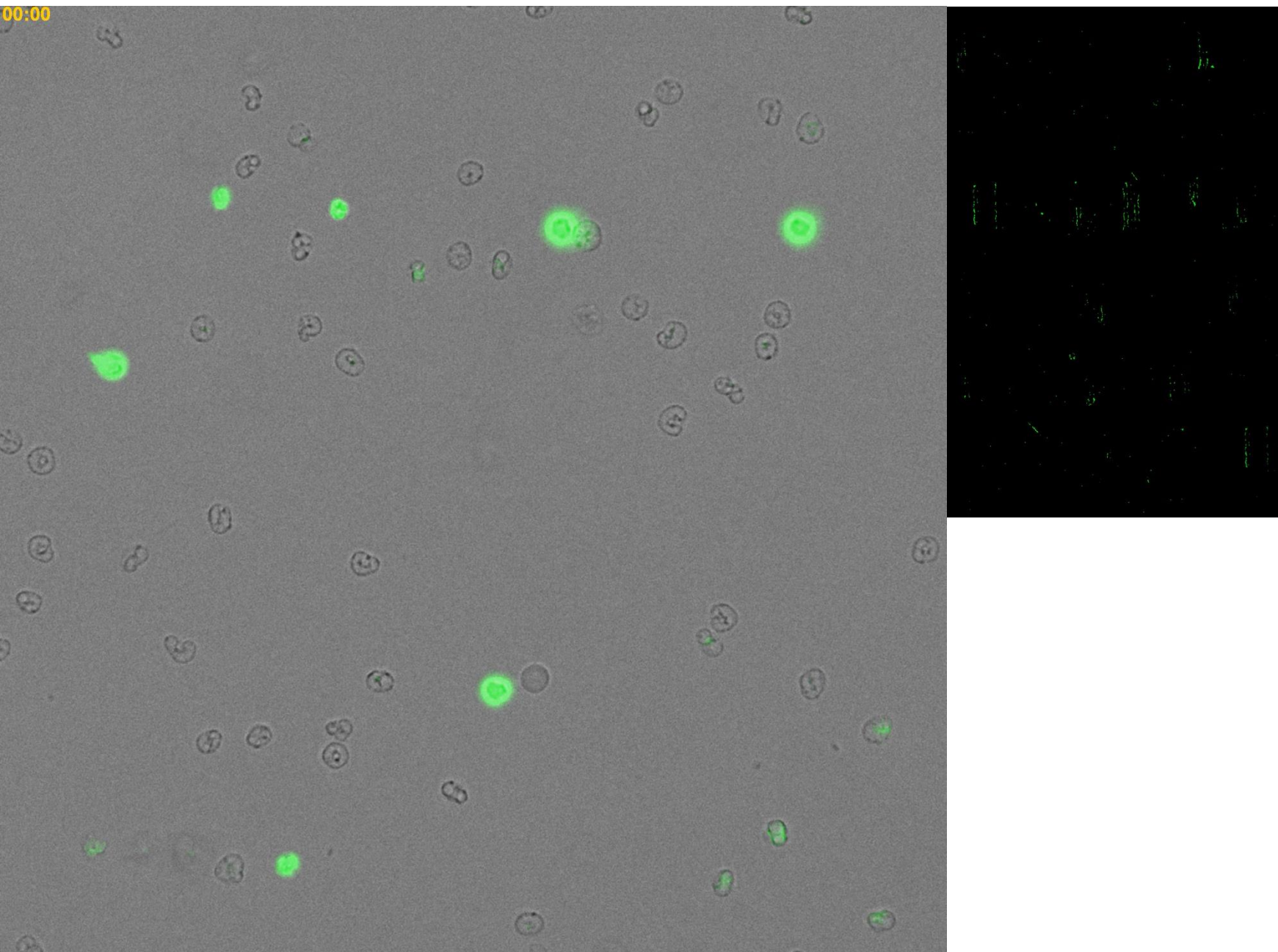
Place the device under the microscope.

Use 1 mL of PBS to collect the bacteria from the device.

Dilute the collected bacteria and spread on an agar plate for counting.

2025-07-22

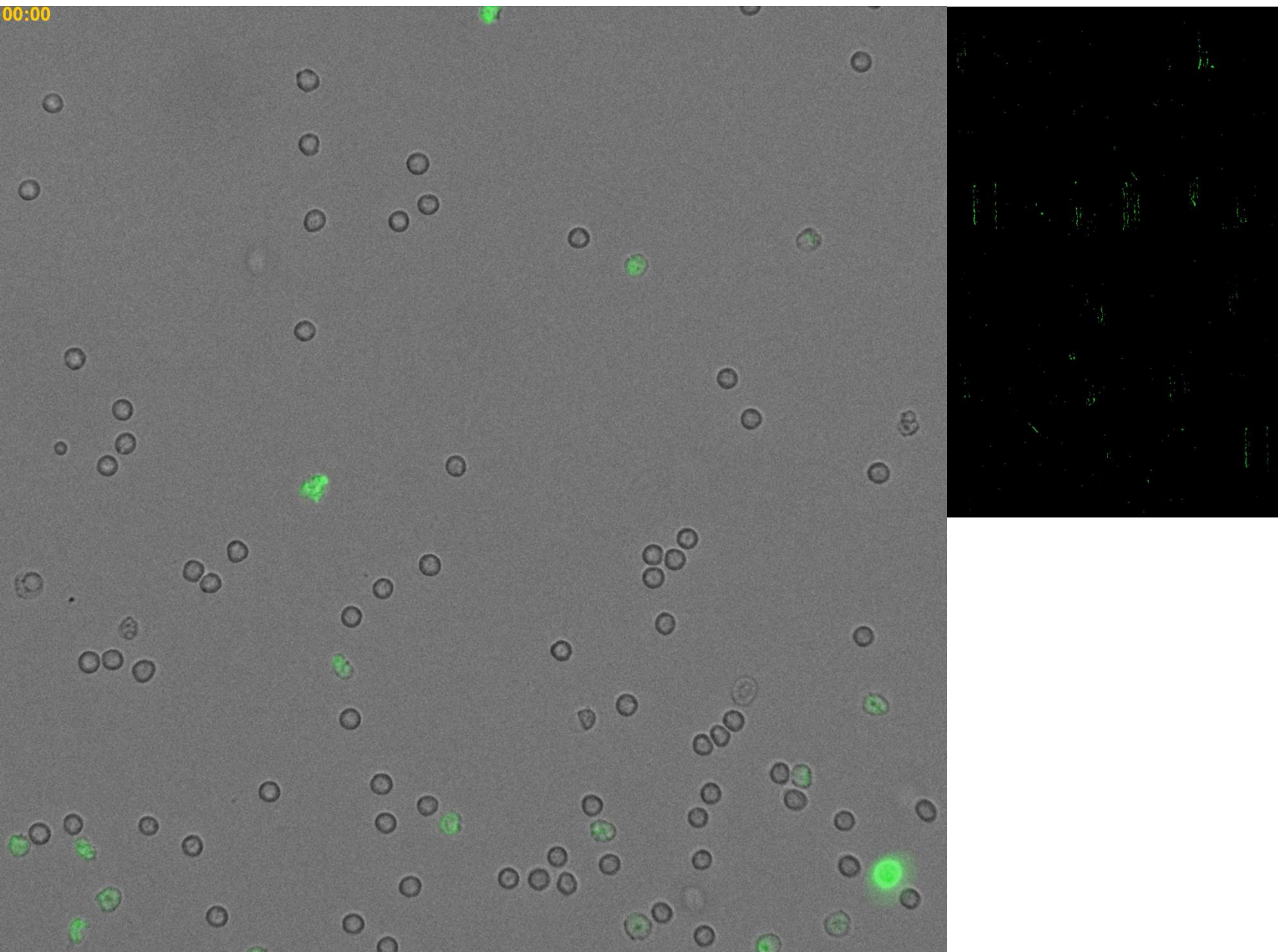
Triton

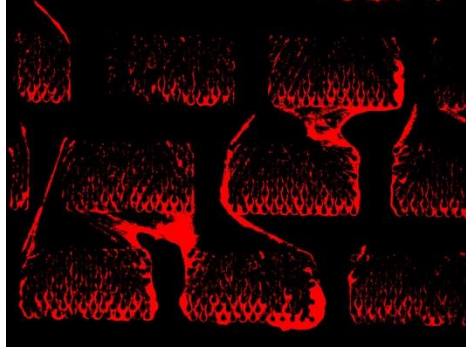


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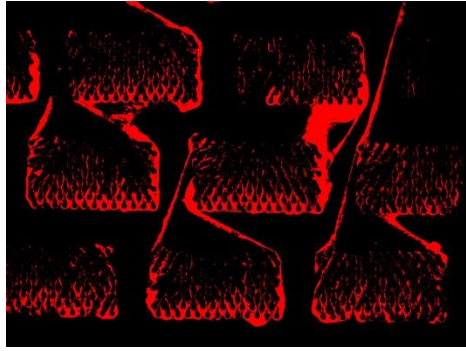
Triton

00:00





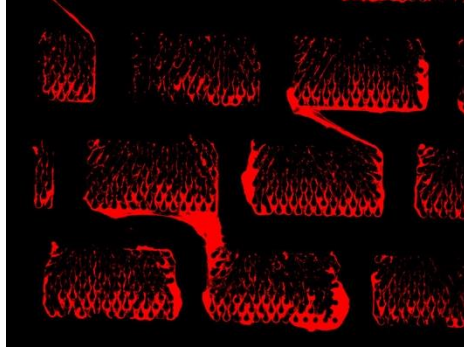
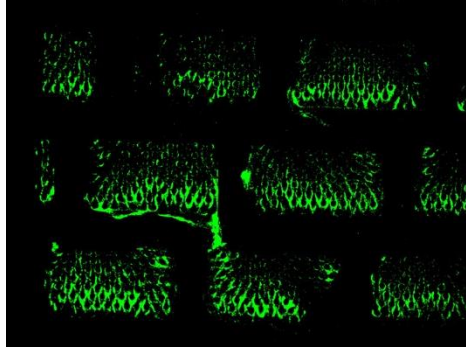
Ionomycin View 1



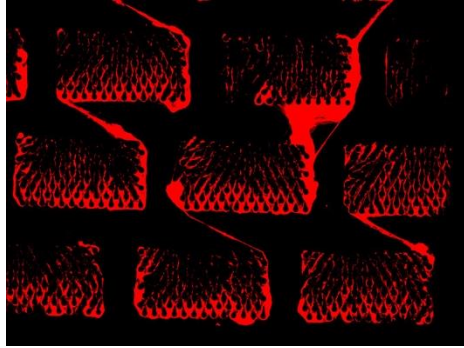
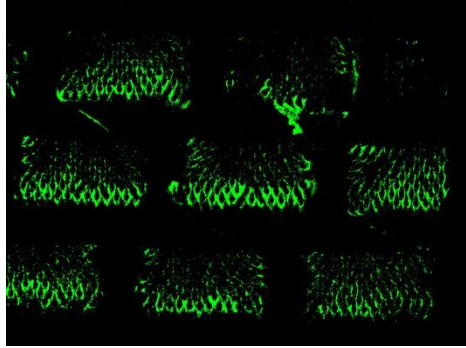
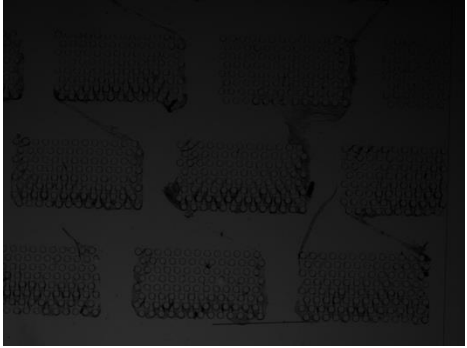
Ionomycin View 2

BF

Cy5



Ionomycin View 1
After adding bacteria
and washing



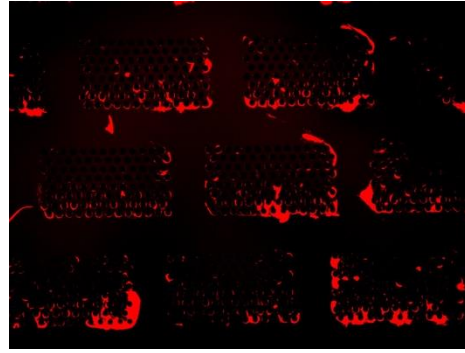
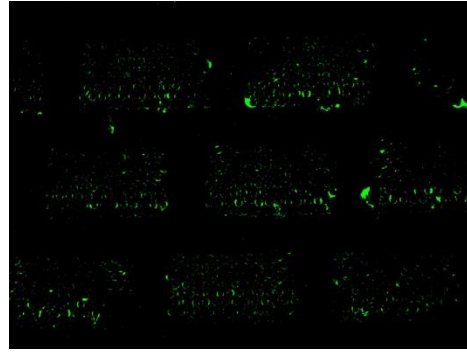
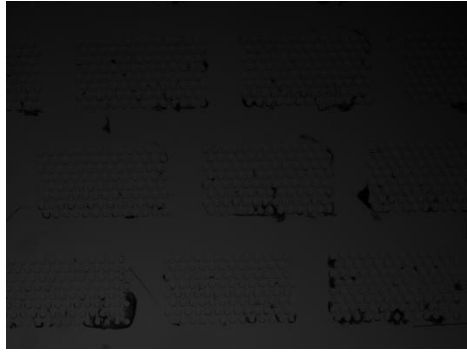
Ionomycin View 2
After adding bacteria
And washing

BF

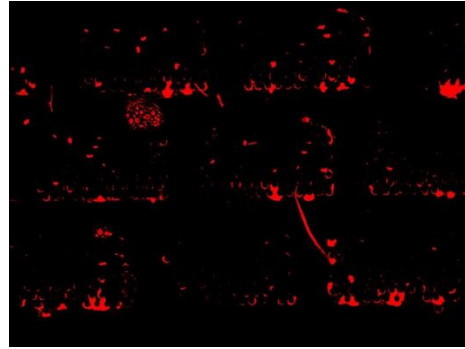
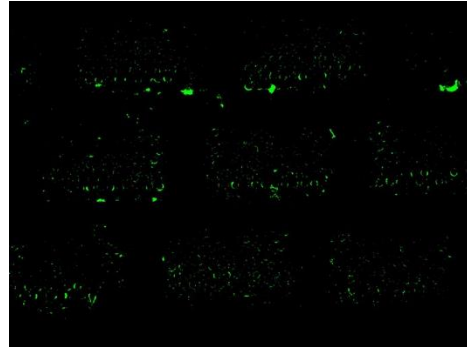
GFP

Cy5

GFP is bacteria (they are green in nature), and Cy5 is the NET.
The red staining is weaker than the green staining,
but same luminance settings have been used (from before)



Ionomycin
Before flushing

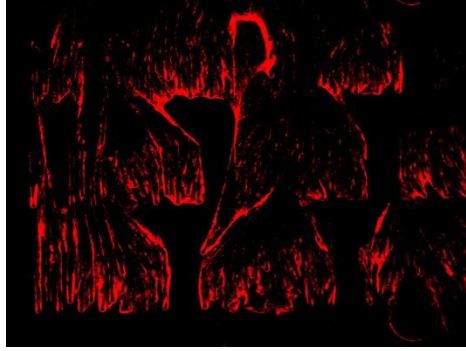
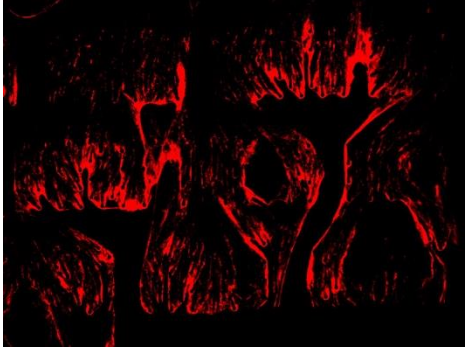


Ionomycin View 2
After flushing

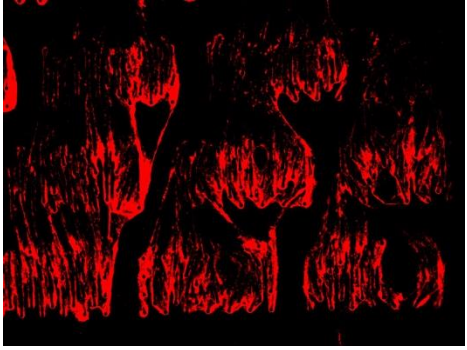
BF

GFP

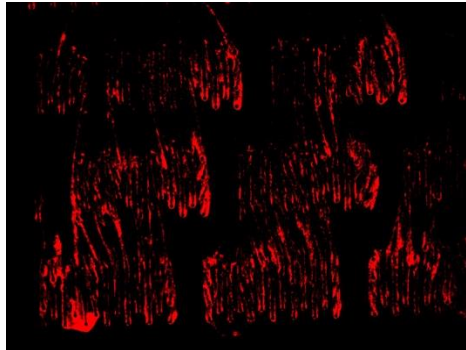
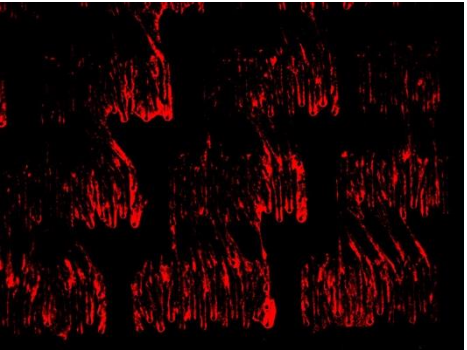
Cy5



Ionomycin



Ionomycin

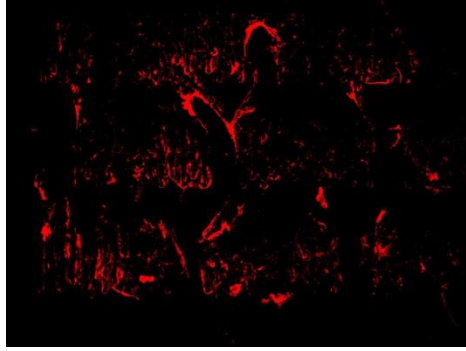
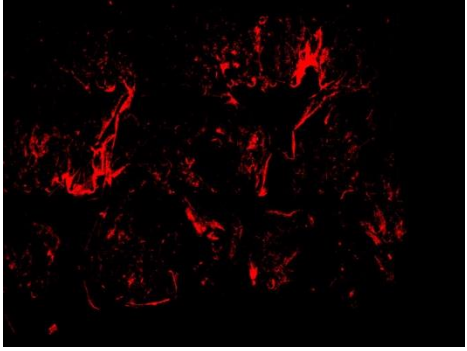


Ionomycin

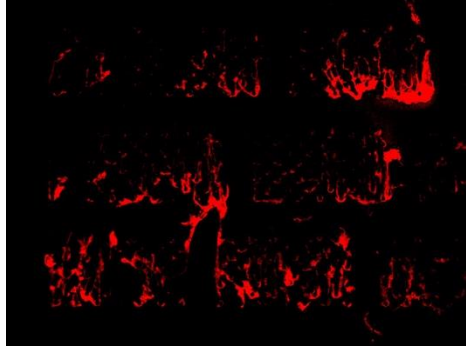
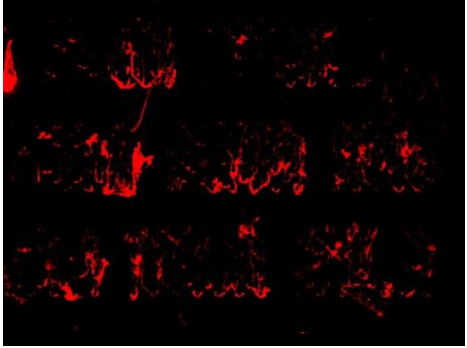
Cy5

Cy5

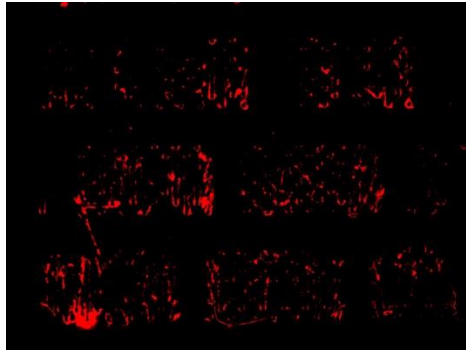
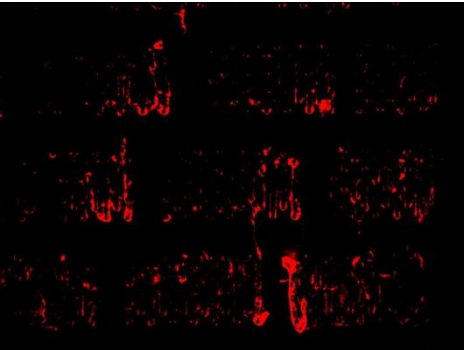
2025-07-25, Bacteria induced NETs, 3h, 1% FBS, 400 microliters of cells



Ionomycin
After adding enzyme 2



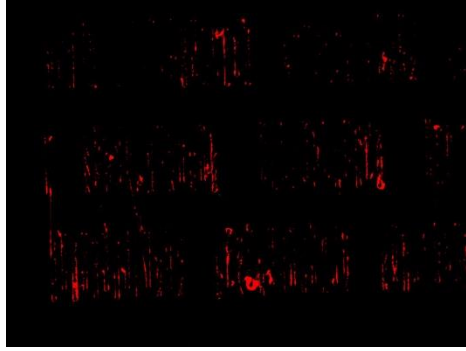
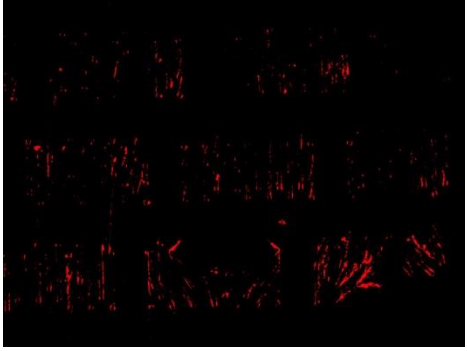
Ionomycin
After adding enzyme 3



Ionomycin
After adding enzyme 4

Cy5

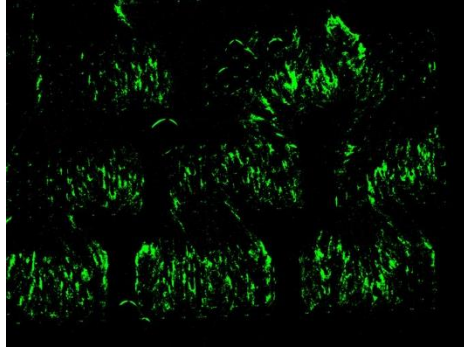
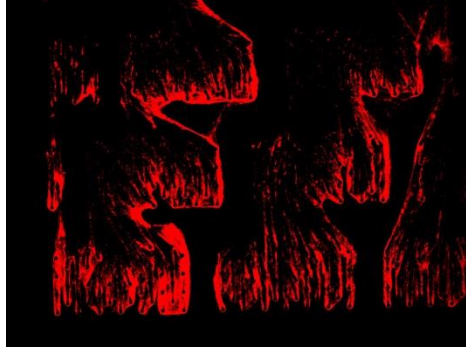
Cy5



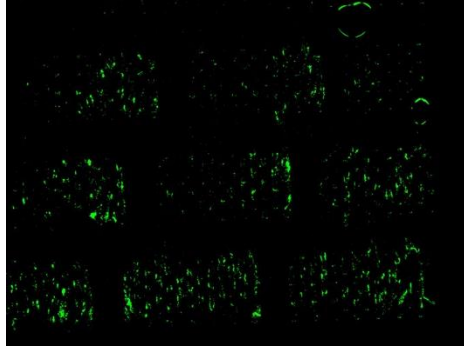
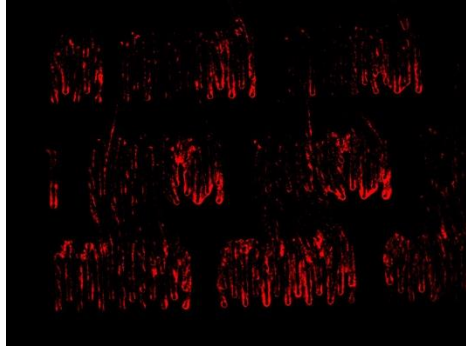
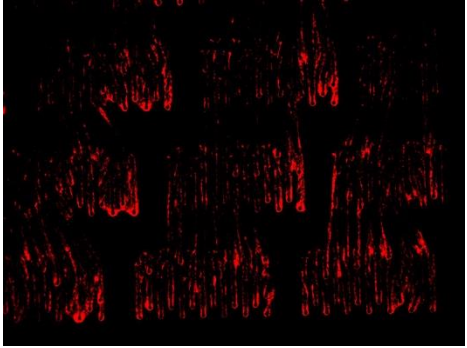
Ionomycin
Before adding DNase



Ionomycin
Before adding DNase



Ionomycin View 1

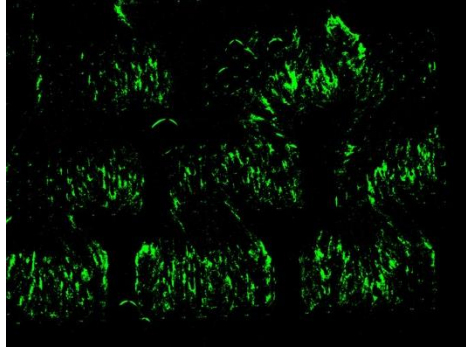
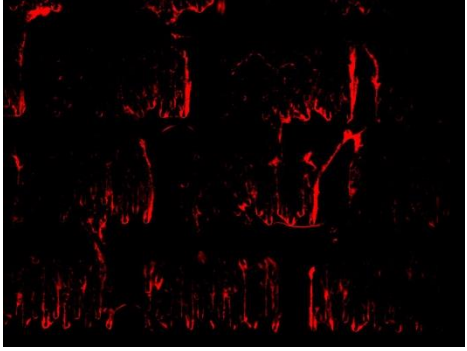


Ionomycin View 2

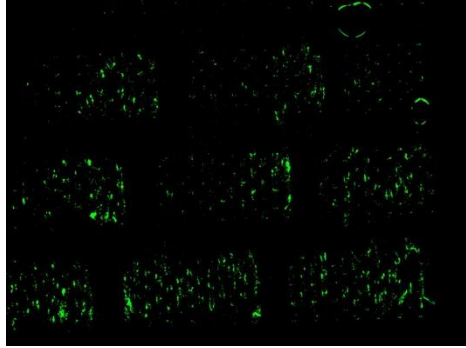
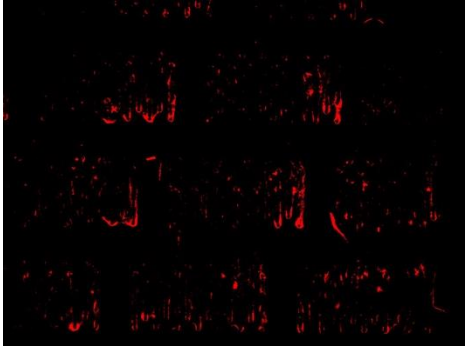
Cy5

Cy5

Bacteria



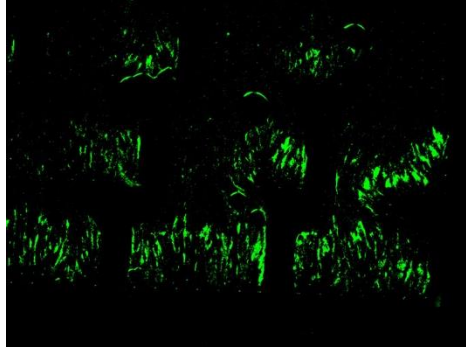
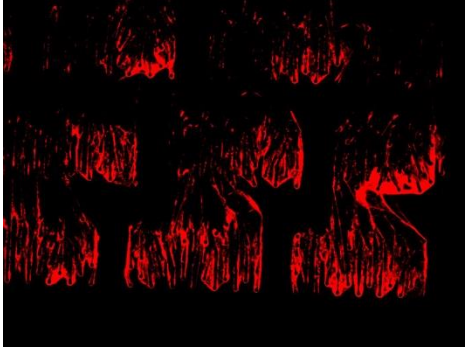
Ionomycin View 1



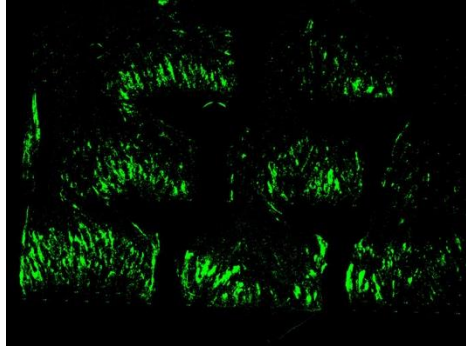
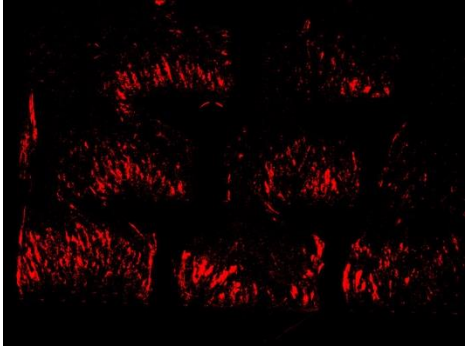
Ionomycin View 2

Cy5

Bacteria



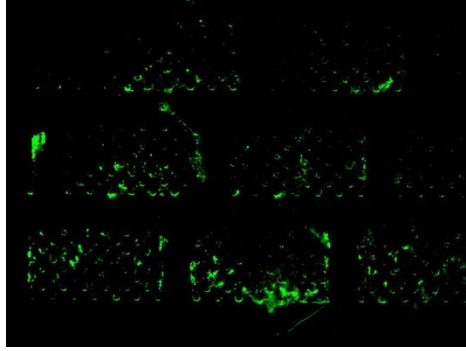
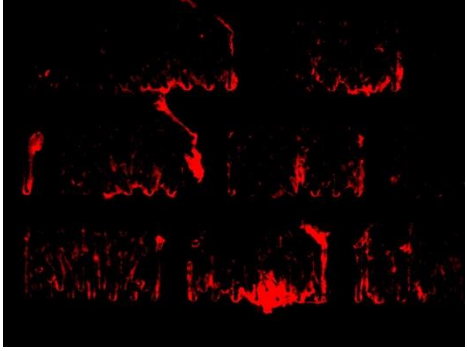
Ionomycin View 1



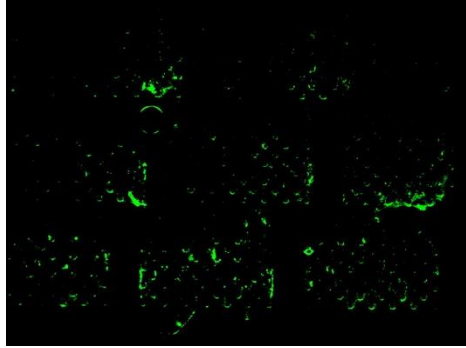
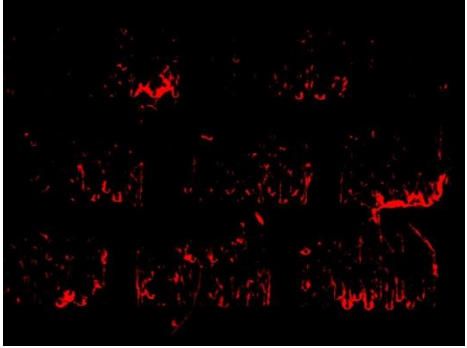
Ionomycin View 2

Cy5

Bacteria



Ionomycin View 1

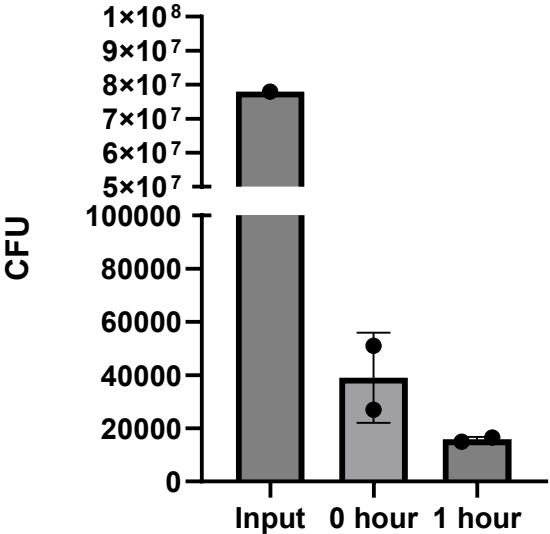


Ionomycin View 2

Cy5

Bacteria

20250727 Bacteria trapping and killing



Project Summary

2025

Timeline

All systems in vitro mouse neutrophils:

Pre-06-03: Training

06-03 NETs in plate wells (PMA, Iono)

06-05 NETs in devices (PMA, Iono)

Conclusion: Both produce NETs in wells and devices, Ionomycin more

06-10 NETs in devices repeat (PMA, Iono, Raptinal, Nigericin, Val-boropro)

06-12 NETs in devices repeat (PMA, Iono, Raptinal, Nigericin, Val-boropro)

Cell death videos (PMA, Iono, Raptinal, Nigericin, Val-boropro)

Conclusion:

PMA, Ionomycin and Raptinal all consistently produces NETs

Nigericin has some NETs. Val-boropro has too low of cell death rate.

Timeline

06-13 Cell death videos (PMA, Iono, Raptinal, Nigericin, Val-boropro)
Images of different staining (PMA, Iono, Raptinal, Nigericin, Val-boropro)

06-17 NETs of concentration gradient (PMA, Iono, Raptinal)
Cell death videos of concentration gradient (PMA, Iono, Raptinal)
Conclusion: NETs decrease with the concentration decrease

06-24 LDH release (PMA, Iono, Raptinal)

06-27 NETs of PAD4 Knockout and WT (PMA, Iono, Raptinal)

07-01 NETs of PAD4 Knockout and WT (PMA, Iono, Raptinal)

Cell death video (PMA, Iono, Raptinal)

Conclusion: In WT NETs, in PAD4 KO no NETs (or less NETs)

Ionomycin kills all in both KO and WT, but only in WT NETs are observed

Raptinal has a big difference in death rate of KO and WT, and a huge difference in NET formation, but LDH remains the same

PMA has a big difference in death rate of KO and WT, but no lysis death at all

Timeline

07-03 NETs of Spontaneous Death

07-09 NETs of Spontaneous Death Repeat

NETs of different protocols - Percoll, Histopaque, Negative Selection

Conclusion: Only Percoll has a little NETs (possible due to RBCs)

Percoll + Histopaque, + Negative Selection, + Histopaque + Negative Selection, all no NETs

07-10 NETs of bacteria induction (S. Aureus 10390, S. Aureus BAA, K12, Bacillus Cereus, A7)

07-15 NETs of bacteria induction Repeat (S. Aureus 10390, S. Aureus BAA, K12, Bacillus Cereus, A7, Triton)

Cell death videos (S. Aureus 10390, S. Aureus BAA, K12, Bacillus Cereus, A7, Triton, Lysis buffer)

07-17 NETs of bacteria induction Repeat (S. Aureus 10390, S. Aureus BAA, K12, Bacillus Cereus, A7)

Cell death videos (S. Aureus 10390 (two different MOI), S. Aureus BAA (two different MOI), Bacillus Cereus, Ionomycin)

Conclusion: Bacillus Cereus can induce lot of NETs (close to Ionomycin)

A7 has NETs (bit less than Raptinal)

S. Aureus 10390, S. Aureus BAA, K12 have no NETs in our system

NETs of 1% FBS and 10% FBS (Bacillus Cereus, Ionomycin)

Conclusion: FBS does not affect NET formation

Timeline

07-22 NETs of Triton

Conclusion: Despite total death in 5 minutes, no NETs

07-22 NETs for bacteria trapping and killing (Ionomycin, 3h, 4h, 5h)

07-29 NETs for bacteria trapping and killing repeat (Ionomycin, 3h, 4h, 5h)

07-31 NETs for bacteria trapping and killing repeat (Ionomycin, 4h, 5h)

ImageJ Macros

20250620_organize_images.bat:

Takes all current directory tif images

Automatically categorizes them into the subfolders of cell video

/* A folder named Template must be provided in the same directory with the correct structure

This can be accomplished by either generating a Template folder by using create_template_folders, or by copying the Template folder from the disk. */

20250620_composite_macro_workflow.jim:

Takes three subfolders, "BF", "PI", "Speck", *number of* images

Processes them into a video automatically

20250620_convert_to_gfp.jim:

Takes all current directory tif images

Converts them to GFP and store them in the subdirectory GFP under the current directory

Same exposure time / gain (no automatic adjustment for individual images, quantification purposes)

(This can be freely edited in the raw script)

20250703_convert_gfp_bf_manual.ijm:

Takes a folder, where there are subfolders containing a bright field image and a GFP image

Loads the GFP image and allows you to set the brightness manually

Outputs the composite image

Protocol Change

After we get our NETs from ionomycin:

Add **300 µL of bacteria** at a concentration of **$5 \times 10^8/\text{mL}$** through the device.

Wash it with **200 µL of PBS, repeated 8 times.**

Add **50 µL of DNase** through the device. **Don't collect the liquid sucked out (at this moment nothing is digested yet)**

Incubate the device at 37°C for 10 minutes.

Add **50 µL of DNase** through the device. **Collect the liquid sucked out.**

Incubate the device at 37°C for 10 minutes.

Place the device under the microscope.

Use 1 mL of PBS to collect the bacteria from the device.

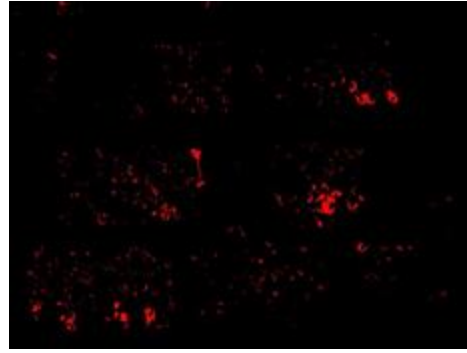
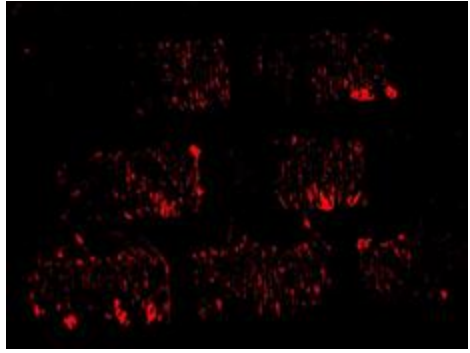
Dilute the collected bacteria and spread on an agar plate for counting.

Protocol Change

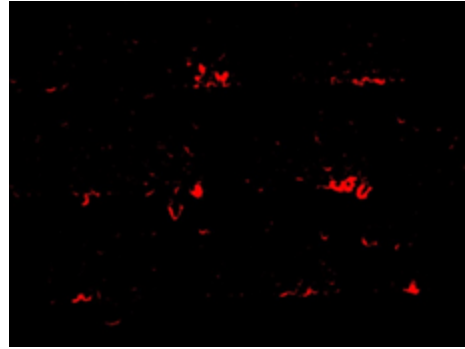
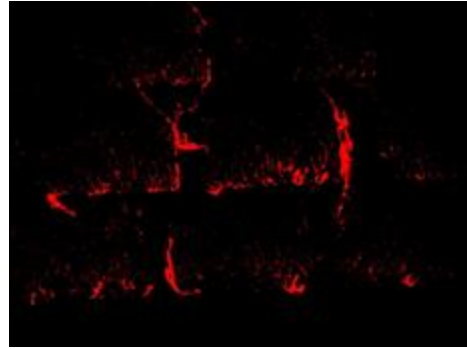
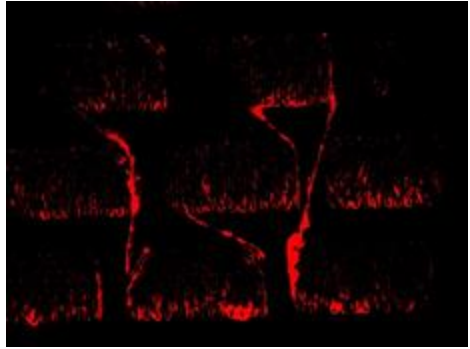
We include **two sets of devices** in our batch:

We culture the cells for 4 hours after adding Ionomycin, then:

1. We **directly add** the cells, bacteria and DNase (and culture) two times and see it under the microscope.
1. We add the cells and bacteria, then we **incubate the device for an hour**, take it out and add the DNase.



Ionomycin
High conc.
DNase

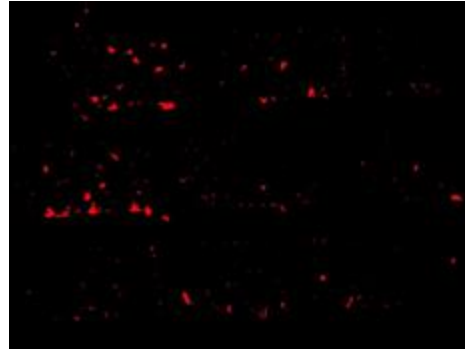
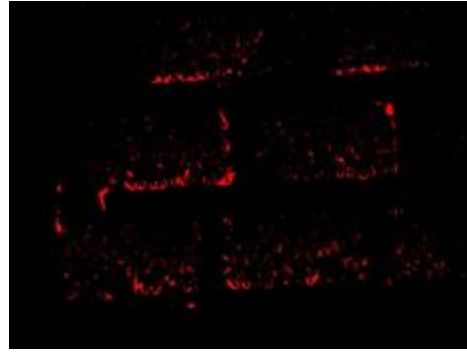
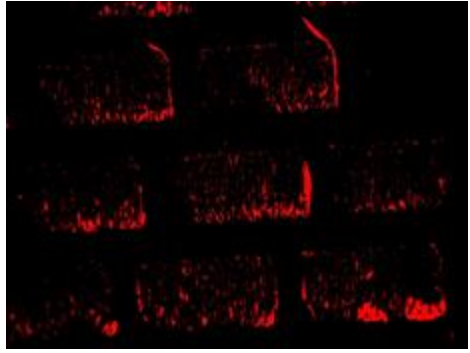


Ionomycin
Low conc.
DNase

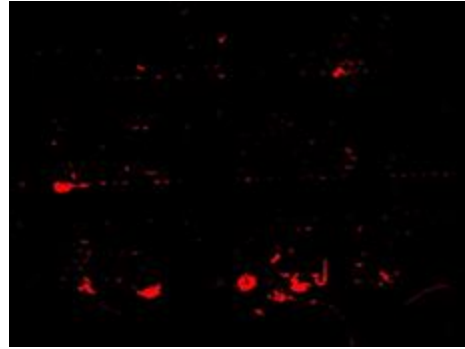
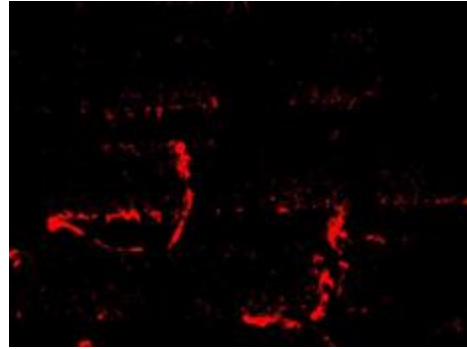
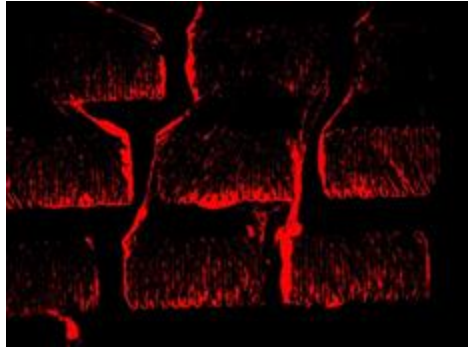
0
min

10 min
1*DNase

20 min
2*DNase



Ionomycin
High conc.
DNase

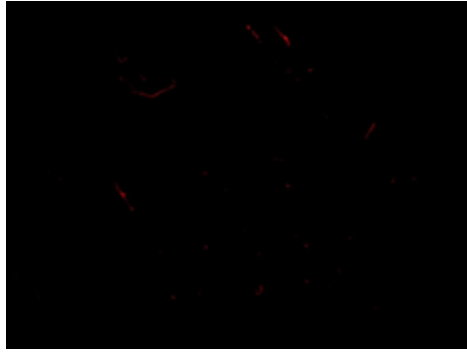
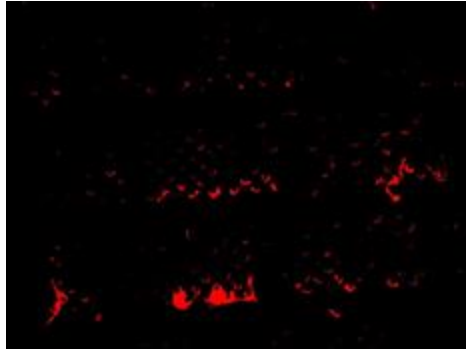


Ionomycin
Low conc.
DNase

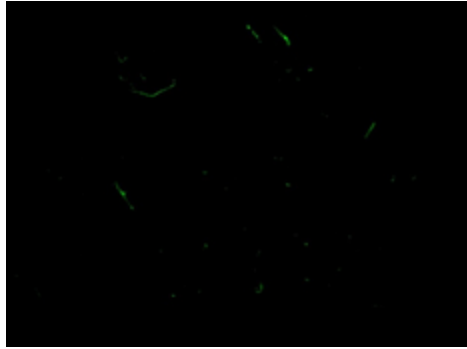
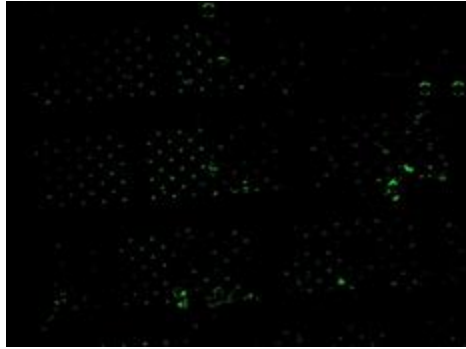
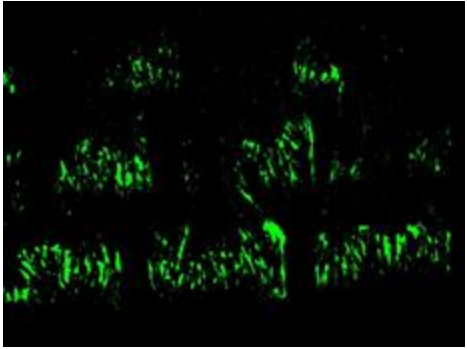
0
min

10 min
1*DNase

20 min
2*DNase



NETs

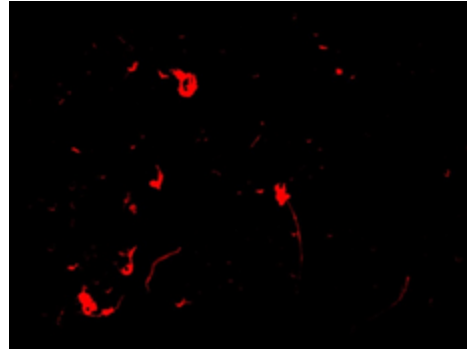
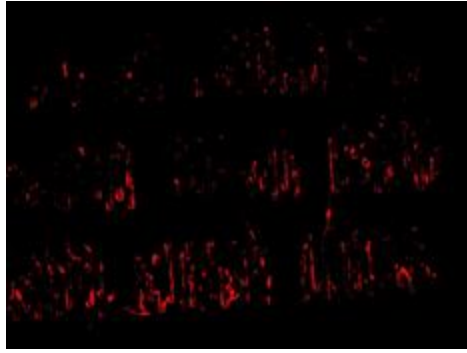


Bacteria

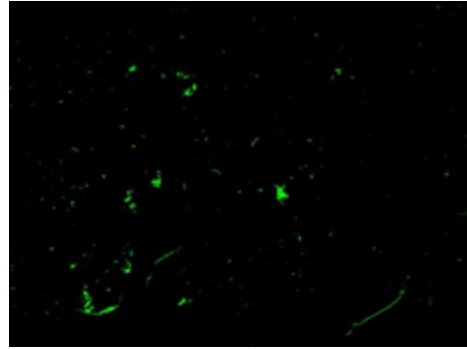
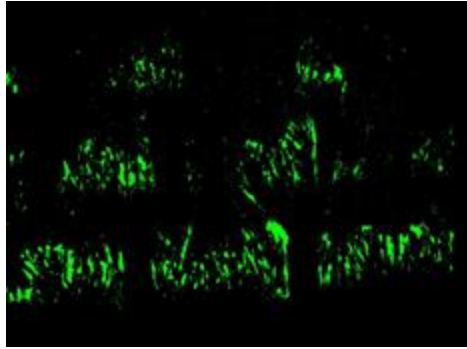
0
min

Group 1
Add enzymes
immediately

Group 2
Incubate 1 hour
more



NETs

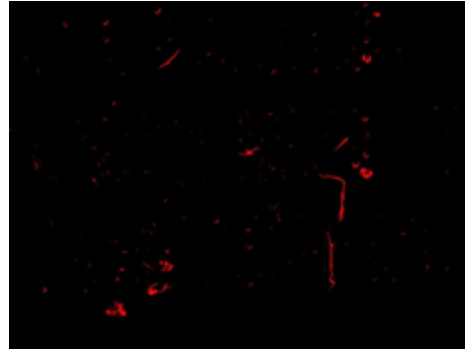
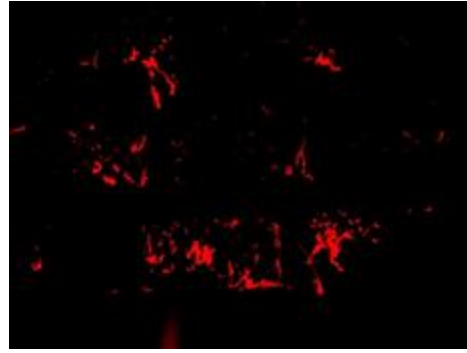
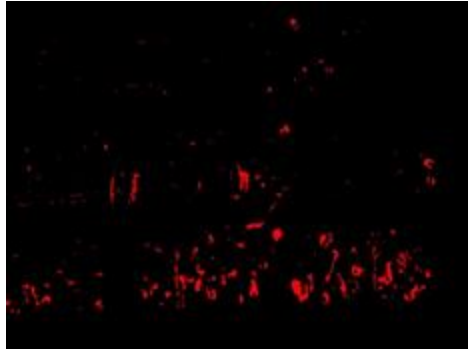


Bacteria

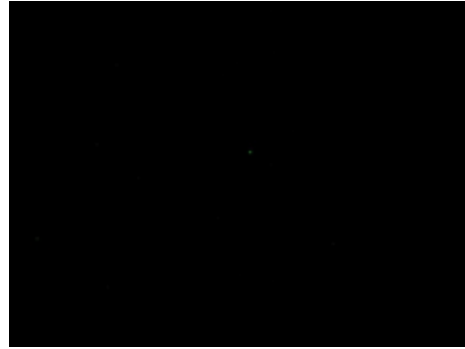
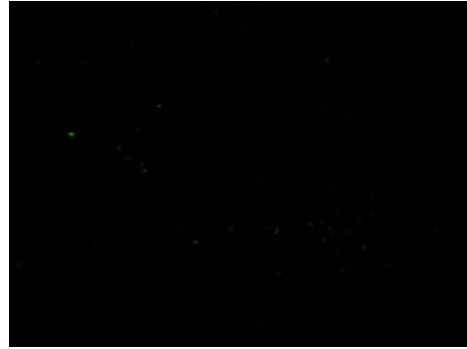
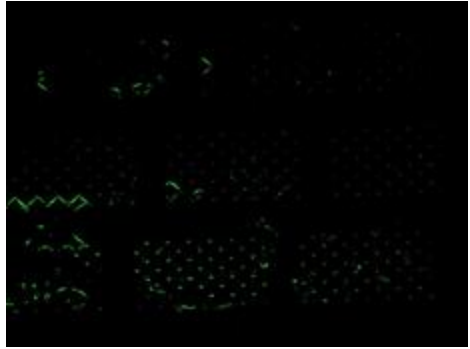
0
min

Group 1
Add enzymes
immediately

Group 2
Incubate 1 hour
more



NETs

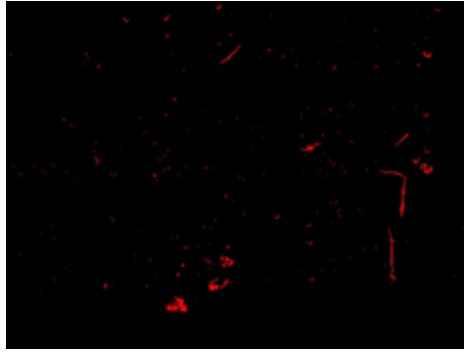
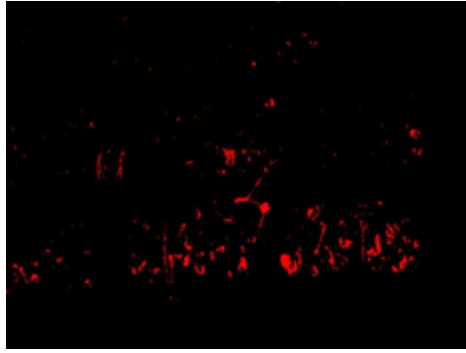
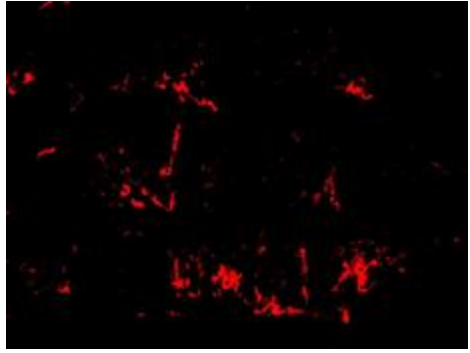


Bacteria

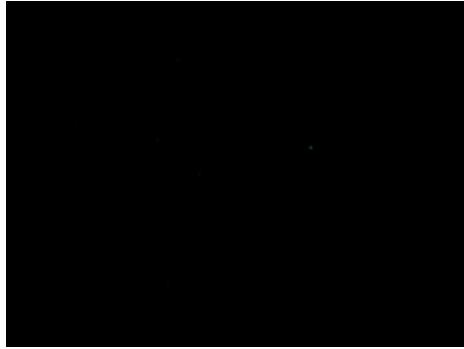
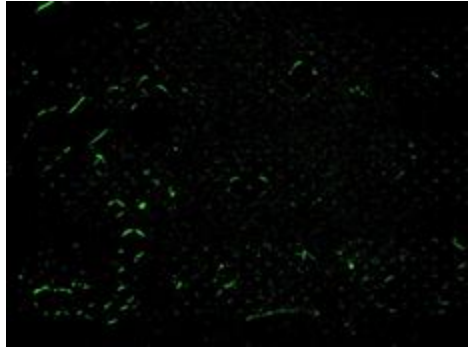
0
min

10 min
1*DNase

20 min
2*DNase



NETs

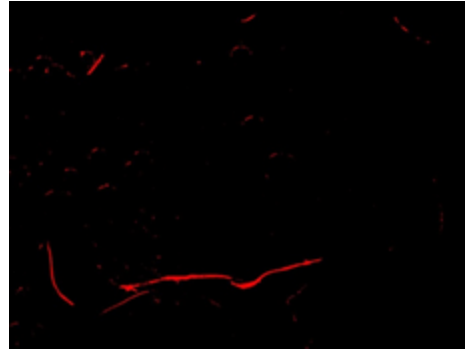
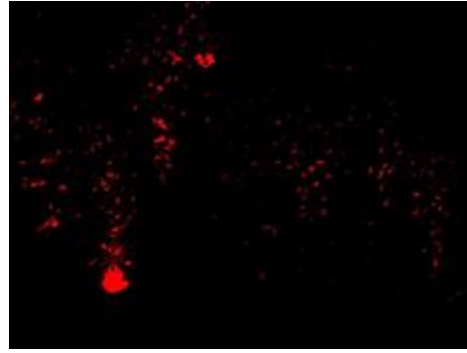
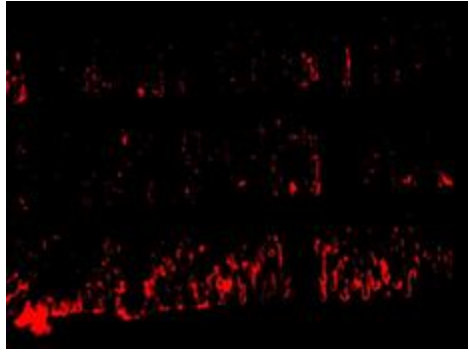


Bacteria

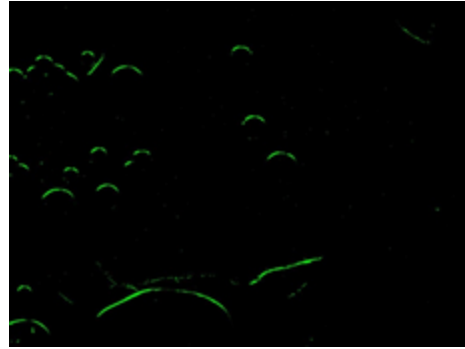
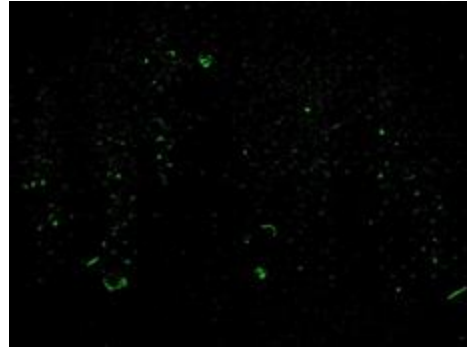
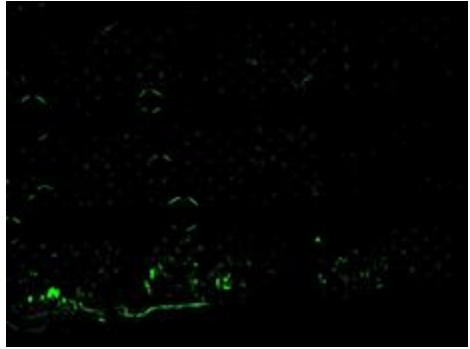
0
min

10 min
1*DNase

20 min
2*DNase



NETs

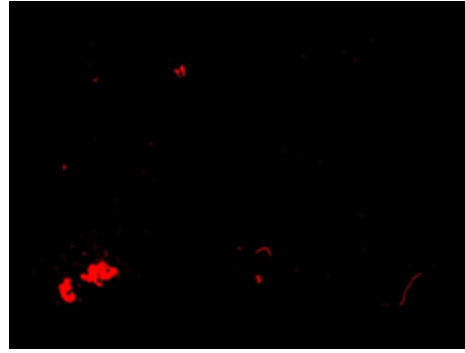
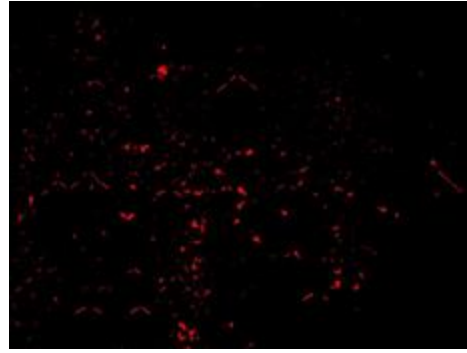
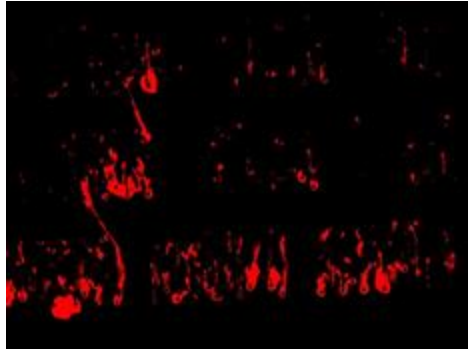


Bacteria

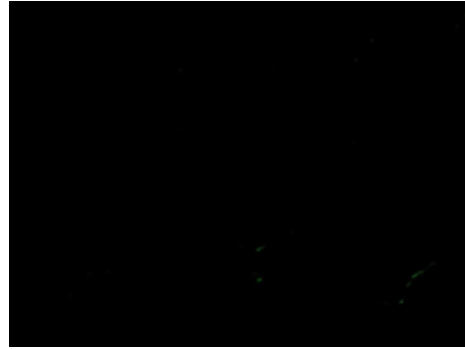
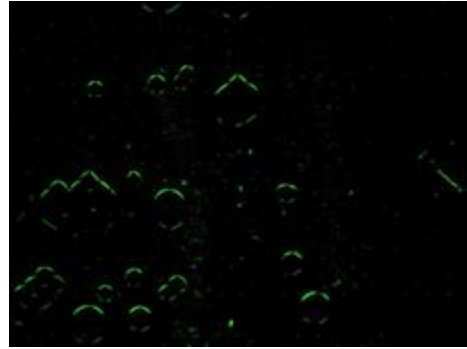
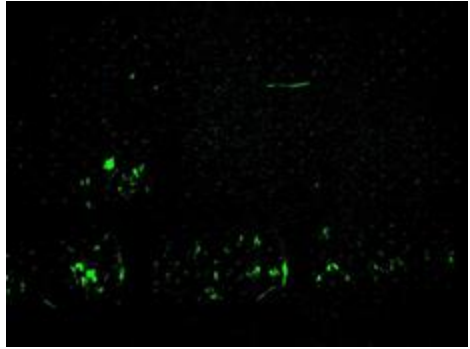
0
min

10 min
1*DNase

20 min
2*DNase



NETs



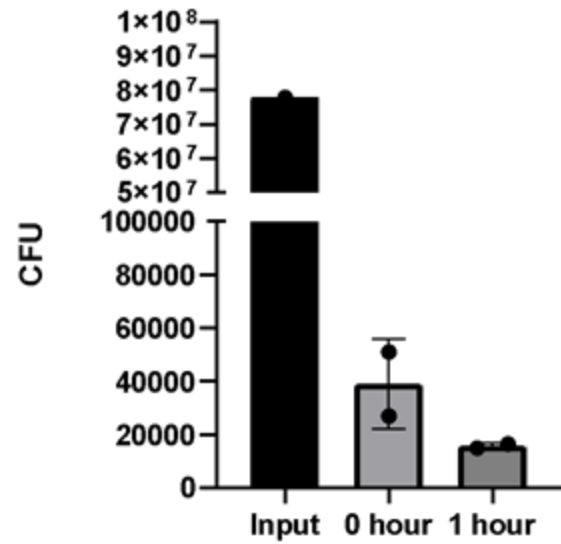
Bacteria

0
min

10 min
1*DNase

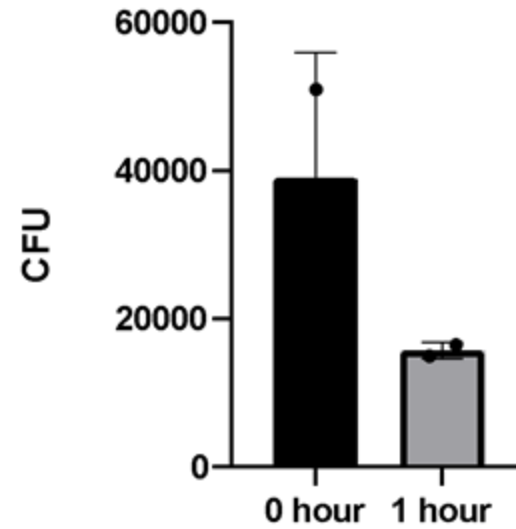
20 min
2*DNase

20250727 Bacteria trapping and killing



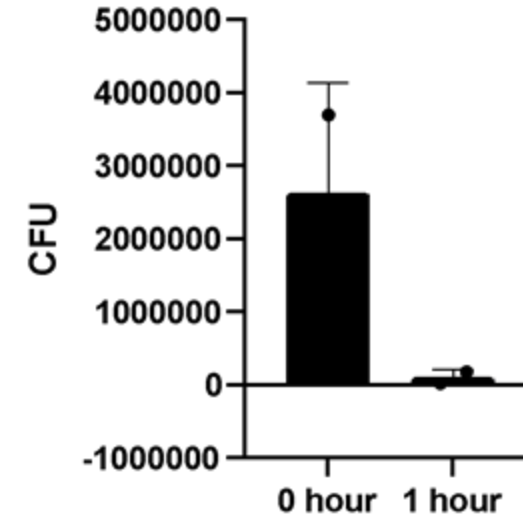
Once digestion
(100ul/1.5ml)

20250729



Twice digestion
(100ul+100ul/1.5ml)

20250801 CFU



Twice digestion
(100ul+100ul/1.5ml)