

Turner Lab Notebook

Callum Malcolm

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Wednesday 15-03-2023

Cell Culture

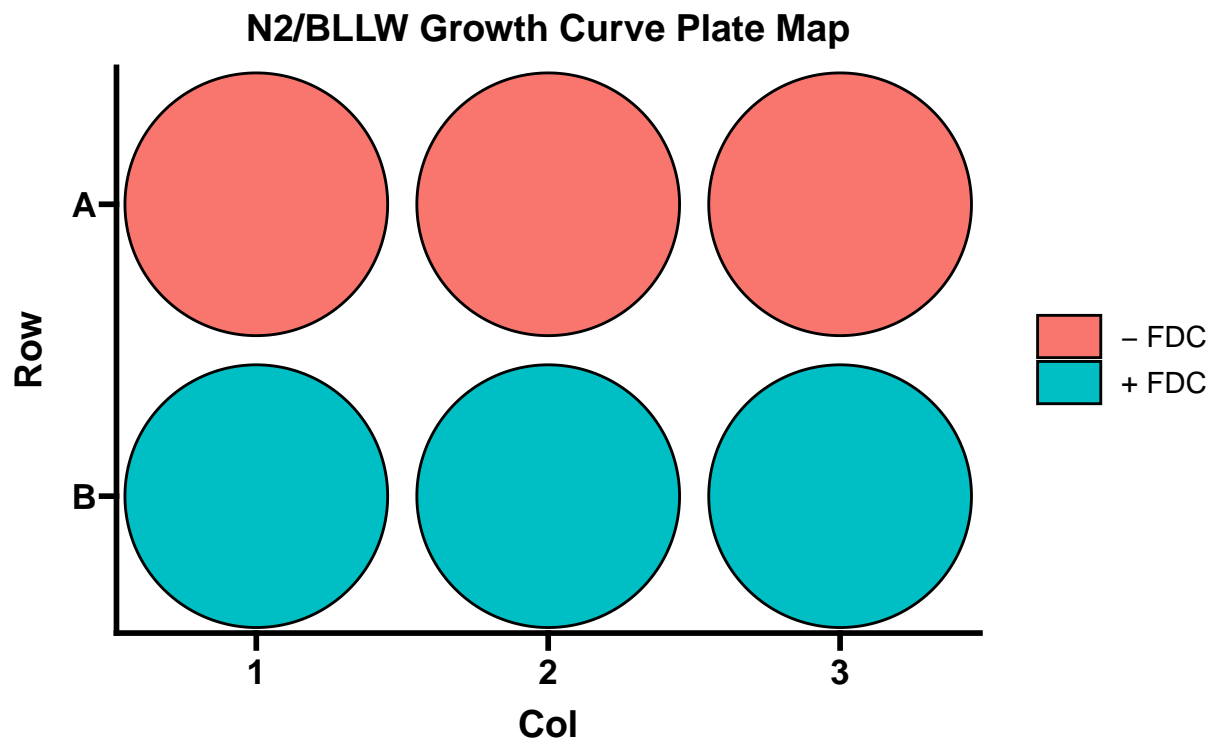
- Given N2/RBL1/BLLW PDX line from Chris Steele

N2 Growth Curve

Looking to compare N2 growth on and off feeders

- Seeded 2mL of N2/BLLW cell suspension at 0.25×10^6 cells/mL

96 Well Plate Map



Thursday 16-03-2023

N2/BLLW Growth Curve Day 1

- Cells gently mixed
- 30 μ L cell suspension taken and counted
- Only counted 1 square and multiplied by 4

| Cell Line | FDC Status | Count | Cell Number |
|-----------|------------|-------|-------------|
| N2 | + FDC | 25 | 2000000 |
| N2 | + FDC | 26 | 2080000 |
| N2 | + FDC | 12 | 960000 |
| N2 | - FDC | 15 | 1200000 |
| N2 | - FDC | 14 | 1120000 |
| N2 | - FDC | 6 | 480000 |
| BLLW | + FDC | 6 | 480000 |
| BLLW | + FDC | 6 | 480000 |
| BLLW | + FDC | 5 | 400000 |
| BLLW | - FDC | 3 | 240000 |
| BLLW | - FDC | 3 | 240000 |
| BLLW | - FDC | 6 | 480000 |

Friday 17-03-2023

N2/BLLW Growth Curve Day 2

- Cells gently mixed
- 30 μ L cell suspension taken and counted
- Only counted 1 square and multiplied by 4

| Cell Line | FDC Status | Count | Cell Number |
|-----------|------------|-------|-------------|
| N2 | + FDC | 36 | 2880000 |
| N2 | + FDC | 28 | 2240000 |
| N2 | + FDC | 38 | 3040000 |
| N2 | - FDC | 61 | 4880000 |
| N2 | - FDC | 38 | 3040000 |
| N2 | - FDC | 16 | 1280000 |
| BLLW | + FDC | 3 | 240000 |
| BLLW | + FDC | 5 | 400000 |
| BLLW | + FDC | 5 | 400000 |
| BLLW | - FDC | 1 | 80000 |
| BLLW | - FDC | 0 | 0 |
| BLLW | - FDC | 1 | 80000 |

Split Cells

| Cell Line | Cells/mL | Cells/mL Seeded | CS volume | Final Volume | Seeded Cells/mL |
|-----------|-------------------|-----------------|-----------|--------------|-----------------|
| N2 | 8.8×10^5 | 3×10^5 | 6.8 mL | 12 mL | 3×10^5 |
| RBL1 | 8.6×10^5 | 3×10^5 | 11.6 mL | 12 mL | 5×10^5 |

Saturday 18-03-2023

N2/BLLW Growth Curve Day 3

- Counted by Chris

Sunday 18-03-2023

N2/BLLW Growth Curve Day 4

- Counted by Chris

Monday 20-03-2023

N2 Growth Curve Day 5

- BLLW were all dead, experiment stopped
- Cells gently mixed
- 30 μ L cell suspension taken and counted
- Only counted 1 square and multiplied by 4

| Cell Line | FDC Status | Count | Cell Number |
|-----------|------------|-------|-------------|
| N2 | + FDC | 146 | 11680000 |
| N2 | + FDC | 112 | 8960000 |
| N2 | + FDC | 132 | 10560000 |
| N2 | - FDC | 112 | 8960000 |
| N2 | - FDC | 155 | 12400000 |
| N2 | - FDC | 141 | 11280000 |

Split Cells

- Split cell lines N2/RBL1/JIRE
- Dirty split 1:2

Seed BLLW

- Given BLLW aliquot from Chris Steel
- Thawed and resuspended in 12 mL media
- Added to T75 pre-seeded with FDC cells (from Chris Steel)

Tuesday 21-03-2023

Large Infection in TC

- N2 Growth Curve stopped

Friday 24-03-2023

Seed Cells - N2/RBL1/JIRE

- Vials given by Chris Steel
- Thawed and resuspended in 12 mL media
- Added to T75 pre-seeded with FDC cells (from Chris Steel)

Monday 27-03-2023

Split Cells

- Split cell lines N2/RBL1/JIRE
- Dirty split 1:3

Wednesday 29-03-2023

Split Cells

- Split by Chris Steel.

Friday 31-03-2023

Split Cells

- Split cell lines N2/RBL1/JIRE
- Dirty split 1:4

Monday 03-04-2023

Large Infection in TC

Seed Cells - N2/RBL1/JIRE

- Vials given by Chris Steel
- Thawed and resuspended in 12 mL media
- Added to T75 pre-seeded with FDC cells (from Chris Steel)

Seed EC50

DNW - N2/JIRE/RBL1

- Vin/Dactolisib

Wednesday 05-04-2023

Seed Cells - N2/RBL1/JIRE

- Vials given by Chris Steel
- Thawed and resuspended in 12 mL media
- Added to T75 pre-seeded with FDC cells (from Chris Steel)

Thursday 06-04-2023

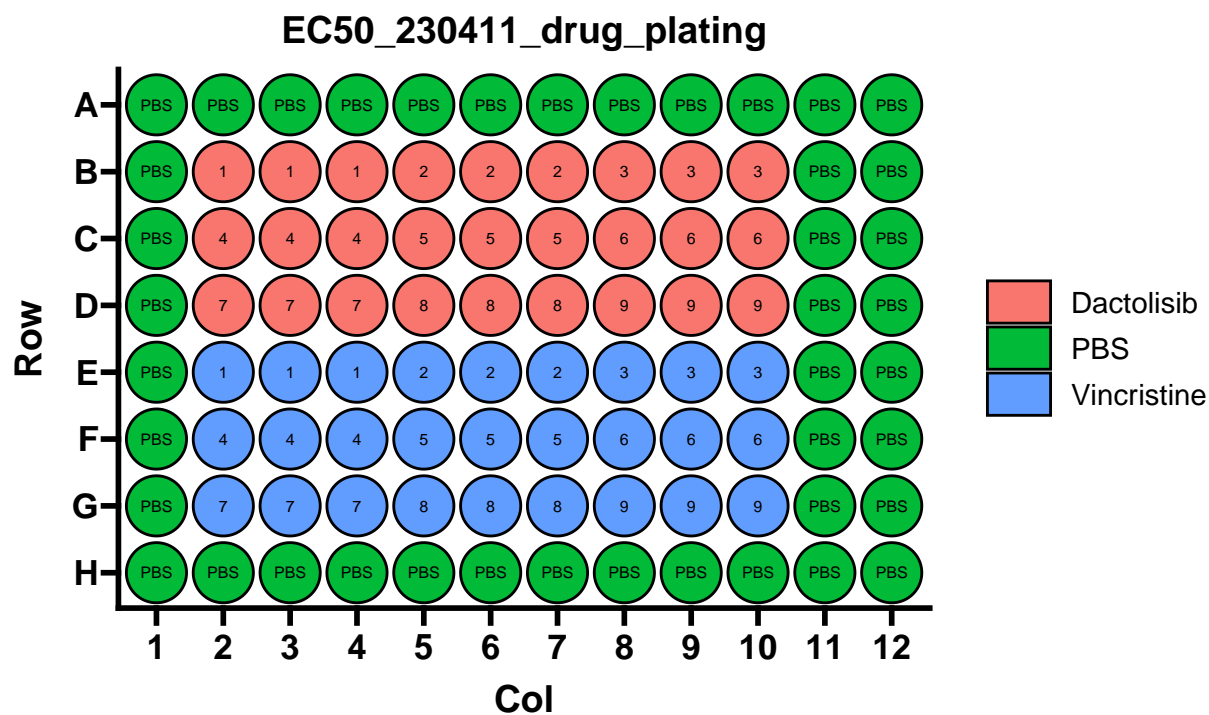
Collect EC50 Plate

- Plate ID: EC50_230619_Plate1
- DNW

Tuesday 11-04-2023

Seed EC50_230411

96 Well Plate Map



Drug Dilutions

| Drug | Dilution | Concentration | Units | Starting_Concentration |
|------------|----------|---------------|-------|------------------------|
| Dactolisib | 1 | 20 | uM | 20000 |
| Dactolisib | 2 | 2 | uM | 2000 |
| Dactolisib | 3 | 0.2 | uM | 200 |
| Dactolisib | 4 | 0.02 | uM | 20 |
| Dactolisib | 5 | 0.002 | uM | 2 |
| Dactolisib | 6 | 0.0002 | uM | 0.2 |
| Dactolisib | 7 | 2e-05 | uM | 0.02 |
| Dactolisib | 8 | 2e-06 | uM | 0.002 |
| Dactolisib | 9 | 2e-07 | uM | 0.0002 |

| Drug | Dilution | Concentration | Units | Starting_Concentration |
|-------------|----------|---------------|-------|------------------------|
| Vincristine | 1 | 30 | nM | 30000 |
| Vincristine | 2 | 3 | nM | 3000 |
| Vincristine | 3 | 0.3 | nM | 300 |
| Vincristine | 4 | 0.03 | nM | 30 |
| Vincristine | 5 | 0.003 | nM | 3 |
| Vincristine | 6 | 0.0003 | nM | 0.3 |
| Vincristine | 7 | 3e-05 | nM | 0.03 |
| Vincristine | 8 | 3e-06 | nM | 0.003 |
| Vincristine | 9 | 3e-07 | nM | 0.0003 |

- 3 plates seeded
 - Plate 1: N2

- Plate 2: JIRE
- Plate 3: RBL1

Wednesday 12-04-2023

Cell Culture

- Thaw RBL1

Side Population - N2_230412

- Hoescht
 - 2.5 / 2.75 / 3
- DNW

Tuesday 02-05-2023

Seeding Cells - N2/RBL1/RAMOS

1. Thawed N2, RBL1, and RAMOS into T75 flasks
Note: N2 and RBL1 flasks seeded with feeders

Wednesday 03-05-2023

Split Cells

| Cell Line | Cells/mL | Cells/mL Seeded | CS volume | Final Volume | Last day split | P number |
|-----------|----------------------|---------------------|-----------|--------------|----------------|----------|
| N2 | 8.93x10 ⁵ | 3x10 ⁵ . | 6.71 mL | 12 mL | Tues 02-05-23 | 2 |
| RBL1 | 7.73x10 ⁵ | 3x10 ⁵ . | 7.76 mL | 12 mL | Tues 02-05-23 | 2 |

2. Expanded N2-Barcodes to T175
3. Seeded feeders into 96-well
 - a. Diluted feeders to 3.2x10⁴ cells/mL
 - b. Add 100μL cell suspension to each well

Thursday 04-05-2023

Monday 19-06-2023

Split Cells

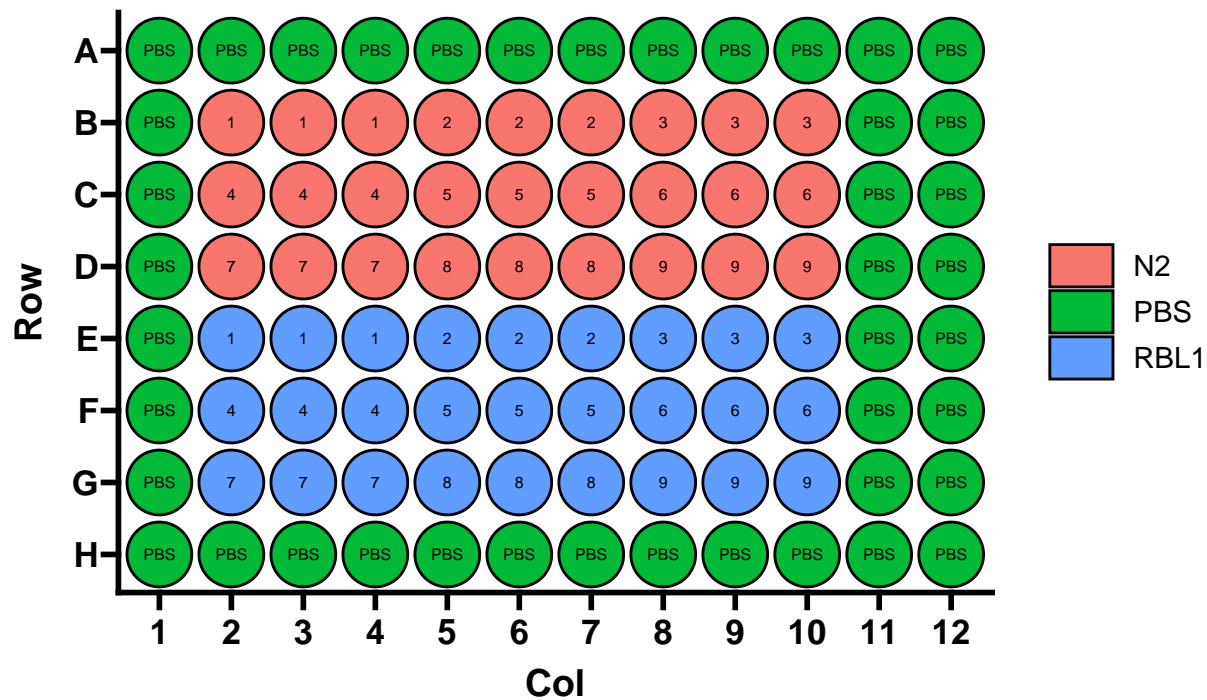
- Split N2/RBL1/JIRE
- Added 10 mL media to BLLW
 - Lots of dead cells observed

Seed EC50

- Plate ID: EC50_230619_Plate1
- Cyclophosphamide (Stock: 20mM) EC50 for N2 and RBL1

96 Well Plate Map

EC50_230619_plate1



Drug Dilutions

| Drug | Dilution | Concentration | Units | Starting_Concentration |
|------------------|----------|---------------|-------|------------------------|
| Cyclophosphamide | 1 | 20 | uM | 20000 |
| Cyclophosphamide | 2 | 2 | uM | 2000 |
| Cyclophosphamide | 3 | 0.2 | uM | 200 |
| Cyclophosphamide | 4 | 0.02 | uM | 20 |
| Cyclophosphamide | 5 | 0.002 | uM | 2 |
| Cyclophosphamide | 6 | 0.0002 | uM | 0.2 |
| Cyclophosphamide | 7 | 2e-05 | uM | 0.02 |
| Cyclophosphamide | 8 | 2e-06 | uM | 0.002 |
| Cyclophosphamide | 9 | 2e-07 | uM | 0.0002 |

Saturday 20-05-2023

Tuesday 20-06-2023

RNA Extraction for SP Test 3 Sample

- Sample collected:
- Low number of cells so used Pico-Pure Kit
- 1 sample (n = 1)

Note before starting:

- Pre-heat block to 42C
- Pre-make DNase treatment master mix (1.1n)

| Component | Volume | MM volume |
|-----------------------|------------|--------------|
| DNase I Stock | 5 μ L | 5.5 μ L |
| DNA Disgestion Buffer | 35 μ L | 38.5 μ L |

RNA Extraction Pico-Pure Protocol

1. Extract cells with 100 μ L of Extraction Buffer (XB)
 - Resuspend the cell pellet gently by pipetting
 - DO NOT VORTEX
2. Incubate at 42C for 30min
3. Centrifuge sample at 3000xg for two minutes
4. Collect supernatant
 - Can stop here and freeze RNA at 70C
5. Pre-condition the RNA Purification Column
 - Add 250 μ L Conditioning Buffer onto the purification column filter membrane
 - Incubate RNA Purification Column with CB for 5min at RT
 - Centrifuge purification column at 16000xg for 1 min
6. Pipette 100 μ L of 70% Ethanol (EtOH) into cell extract
 - Mix well by pipetting
 - DO NOT VORTEX
7. Add cell extract to column
8. Centrifuge column for 2 minutes at 100 x g and then 16000 x g for 30s
9. Pipette 100 μ L Wash Buffer 1 (W1) into the purification column and centrifuge for 1min at 8000 x g
10. Add 40 μ L of DNase treatment master mix to purification tube
 - Add directly onto membrane
11. Incubate at RT for 15 min
12. Pipette 40 μ L Wash Buffer 1 (W1) onto purification column membrane and centrifuge at 8000 x g for 15 seconds
13. Pipette 100 μ L Wash Buffer 1 (W2) into purification column membrane and centrifuge for one minute at 8000 x g
14. Pipette 100 μ L Wash Buffer 1 (W2) into purification column membrane and centrifuge for two minutes at 16000 x g
 - Check if any wash buffer remains in column and centrifuge again at 16000 x g for one minute to clear all the liquid

15. Transfer purification column to new 0.5mL tube
 - Insert purification column/0.5mL assembly into 1.5mL tube for centrifugation
16. Pipette EB directly onto membrane of the purification column
 - Gently touch tip to surface of membrane to ensure maximum absorption of EB into membrane
 - Use 11 μL to 30 μL EB
17. Incubate purification column for one minute at room temperature
18. Centrifuge column for one minute at 1000 x g to distribute EB in column
19. Centrifuge column for one minute at 16000 x g to elute RNA
20. Started PCR and completed Tapestation

Tapestation__230620

High-Sensitivity RNA ScreenTape Protocol

Results

- Inconclusive, proceeded with PCR

Reverse Transcription

Notes Before Starting:

- Label PCR strip tubes
- Thaw RT reagents
- Uses WILDseq specific RT primer
- Dilute RT primer to 2 μM
 - Do this in a PCR hood
 - **IMPORTANT** Do not mix primers and carefully note which RT primer is added to which sample
- Make RT and ExoI master mixes (MM) before starting
 - MM amounts are 1.1n

Protocol:

Note: Used “rt” protocol in callum folder on thermocycler for heating steps

1. In PCR strip tubes diluted 1-5ug RNA in a total volume of 10 μL of RNase/DNase-free water
2. Added 1 ul of RT primer
3. Added 1 ul of dNTPs
4. Denatured at 65 C for 5 mins in PCR machine
5. Added straight to ice for 2 mins
6. Spun to get liquid to bottom of tube
7. Prepared master mix of RT enzyme and buffers (Made prior to starting)

| Component | Volume | MM volume |
|----------------|-----------------|-------------------|
| 5x SSIV Buffer | 4 μL | 4.4 μL |
| SSIV RT | 1 μL | 1.1 μL |
| 100 mM DTT | 1 μL | 1.1 μL |

| Component | Volume | MM volume |
|-----------|-----------|-------------|
| RNase Out | 1 μ L | 1.1 μ L |

8. Added 7 μ L of RT MM prepared above to each sample and mix
 - Spun briefly to get liquid to bottom of the tube
9. In PCR Machine: incubated at 53C for 10 mins followed by 80C for 10 mins
10. Added 3 μ L of ExoI MM to each sample

| Component | Volume | MM volume |
|----------------------------|-----------|-------------|
| Thermolabile Exonuclease I | 1 μ L | 1.1 μ L |
| NEBuffer r3.1* | 2 μ L | 2.2 μ L |

- Most PCR buffers are compatible
11. In PCR Machine: heat at 37C for 4 mins followed by 80C for 1 min
 - Spin briefly to get liquid to bottom of the tube
 12. Add 1 μ L of RNase H and incubate at 37C for 20 mins
 13. Dilute final cDNA 1 in 2 with DNase/RNase-free water

PCR Amplification of cDNA

1. Prepare mastermix of below reagents at 1.1x the number of samples

| Component | Volume | MM volume |
|--------------------------------|-----------------------------|-----------|
| Q5 Master Mix | 25 μ L | |
| P5_R1_fwd (100 μ M) | 2.5 μ L | |
| P7_rev | 2.5 μ L | |
| DNase/RNase H20 | 13.5 μ L | |
| Q5 polymerase | 0.5 μ L | |
| Total Master Mix volume | 40 μL | |
| cDNA from above | 10 μ L | |

2. Perform PCR using the following parameters:

| Steps | Time |
|--------------------|------------|
| Step 1: 98C | 30s |
| Step 2: 98C | 10s |
| Step 3: 61C | 30s |
| Step 4: 72C | 30s |
| Step 5: 72C | 2 min |
| Step 6: 12C | Hold |

20-25 cycles of steps 2-4

Ran overnight and left at 12C

Wednesday 21-06-2023

Tapestation_230619_SPTest3_cDNA

High-Sensitivity DNA ScreenTape Protocol

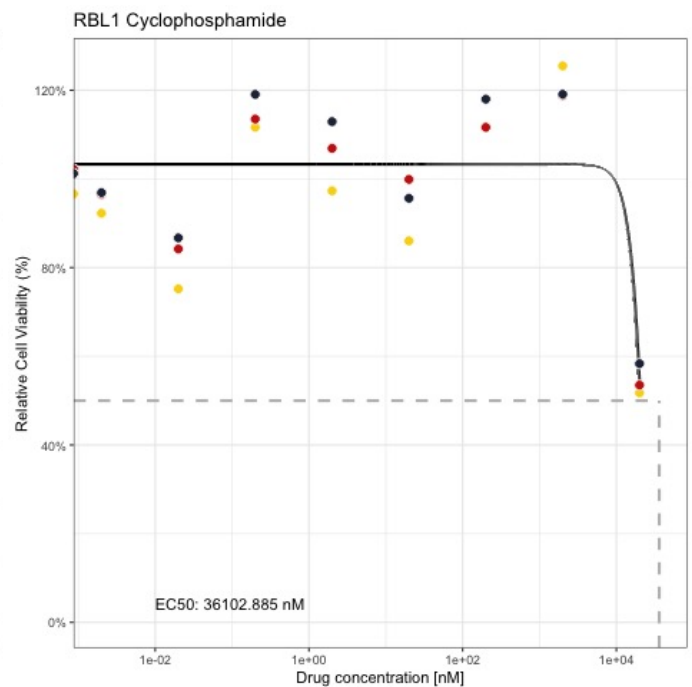
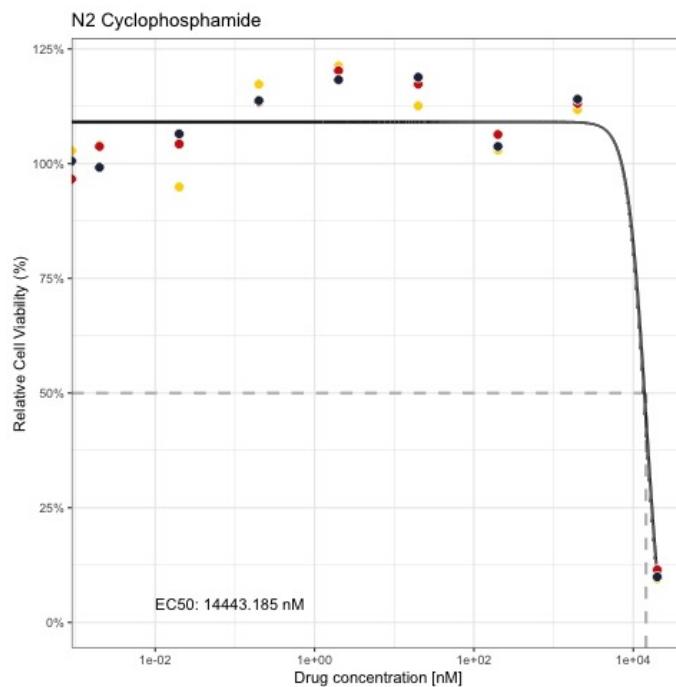
Results

- * Did not work, bands present but faint and not at 200bp
- * Potential solutions:
 - Repeat PCR with increased cycles
 - Repeat RNA extraction with trizol protocol (Anna)

Thursday 22-06-2023

Collect EC50 Plate

- Plate ID: EC50_230619_Plate1



Friday 23-06-23

Split Cells

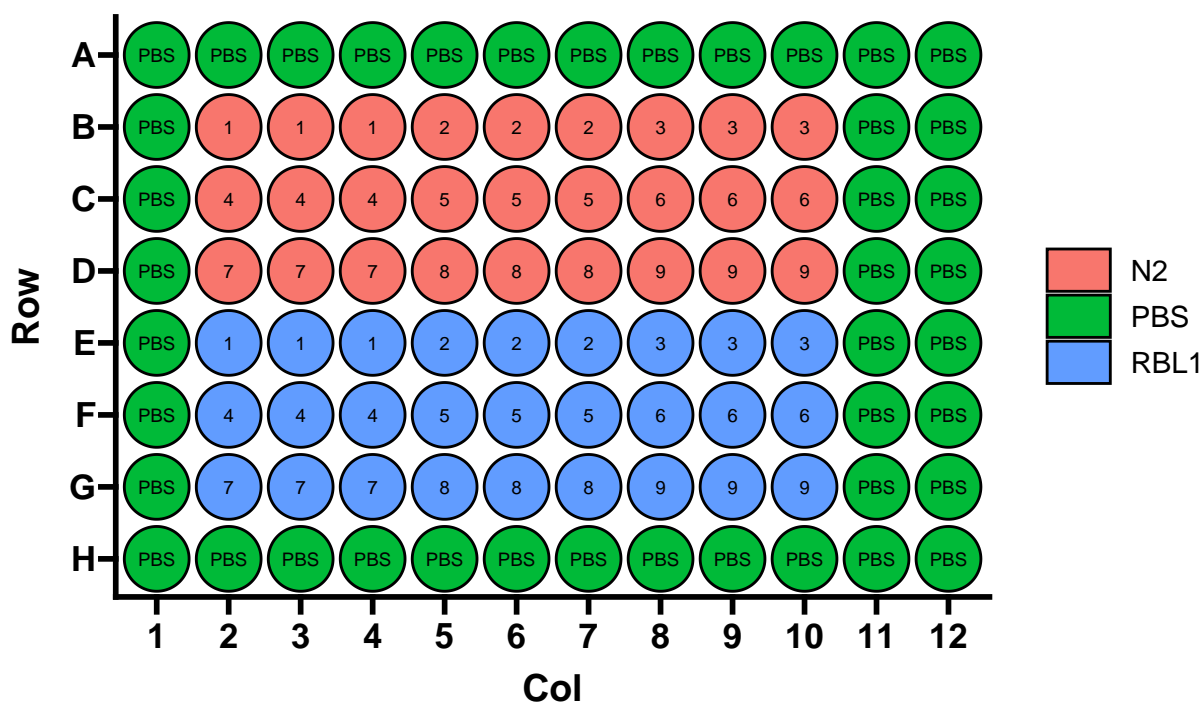
1. Threw out BLLW (all dead)
2. Split JIRE 1/2
3. Split N2/RBL1 1/4

Seed EC50

- EC50 Plate ID: EC50_230623_Plate1

96 Well Plate Map

EC50_230623_plate1



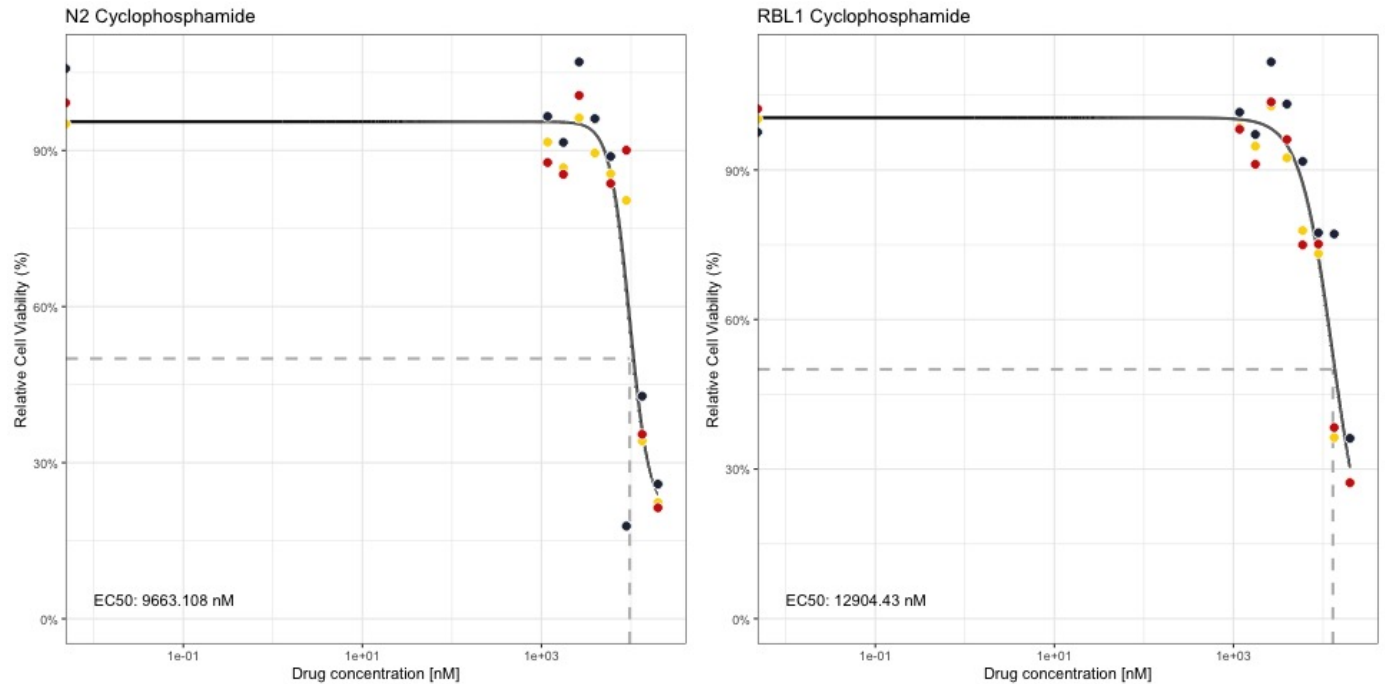
Drug Dilutions

| Drug | Dilution | Concentration | Units | Starting_Concentration |
|------------------|----------|---------------|-------|------------------------|
| Cyclophosphamide | 1 | 20 | uM | 20000 |
| Cyclophosphamide | 2 | 13.3 | uM | 13300 |
| Cyclophosphamide | 3 | 8.89 | uM | 8890 |
| Cyclophosphamide | 4 | 5.93 | uM | 5930 |
| Cyclophosphamide | 5 | 3.95 | uM | 3950 |
| Cyclophosphamide | 6 | 2.63 | uM | 2630 |
| Cyclophosphamide | 7 | 1.76 | uM | 1760 |
| Cyclophosphamide | 8 | 1.17 | uM | 1170 |
| Cyclophosphamide | 9 | 0.78 | uM | 780 |

Monday 26-06-23

Collect EC50 Plate

- Plate ID: EC50_230623_Plate1



Break for writing MRes Report: 27/06/2023 - 24/07/2023

Monday 24-07-2023

Seeding Cells - RAMOS/FDC

Made New Media: AR-1

1. Made new media: AR-1

| Solution | ID code | Volume | % Total volume |
|---------------|---------|--------|----------------|
| Advanced RMPI | — | 500 mL | 78% |
| Glutamax | — | 6.5 mL | 20% |
| Pen-Strep | — | 6.5 mL | 1% |
| FBS | — | 128 mL | 1% |

2. Seeded Feeders

- 2 vial of FDC cells thawed
- Resuspended in 12 mL of media
- 6 mL of FDC cell suspension were added to 2 separate T75 flasks (Seeding RBL1/BLW on 25-07-2023)
- Remaining 12 mL of Media were added to T175 (Seeding N2 on 25-07-2023)

3. Seeded Ramos WILDseq Barcode Pool 2

- Thawed a vial of RAMOS pool 2 (previously barcoded DATE)
- Resuspended in 5 mL
- Spun down 90* for 5min
- Resuspended in 12 mL of AR-1 media
- Seeded into T75

Experimental Plans

1. Barcode BLLW/RBL1
2. Set up Drug pressure experiment for RAMOS barcoded cells
 - EC50's for cyclophosphamide and methotrexate need to be completed
 - Drugs to test (EC50):

| Drug | Concentration | Replicates |
|--------------|---------------|------------|
| DMSO | 1% | 3 |
| Doxorubicin | 200nM | 3 |
| Methotrexate | 100nM | 3 |
| Vincristine | 1nM | 3 |

3. Do Side Population Experiment for N2 Barcoded Cells
 - Grow up sorted side pop to avoid issues with low cell numbers

Tuesday 25-07-2023

Seed Cells - N2-BC4/RBL1/BLLW/RAMOS-BC1

- Got vials from Chris
- Thawed, resuspended in 6 mL AR-1
- Spun down at 90* for 5 min
- Discarded supernatant and resuspended pellet in 12 mL of AR-1
- Added RBL1/BLLW to T75's pre-seeded with FDC (24-07-2023)
- Added RAMOS-BC1 to new T75
 - RAMOS-BC2 from 24-07-2023 was dead
 - Was taken from -80, New RAMOS vial from LN
- Added N2 to T125 seeded with FDC (24-07-2023)

Wednesday 26-07-2023

Cell Culture

- Checked cells, all flasks looked healthy
- Added media
 - N2 - 20ml of AR-1
 - RAMOS - 10 mL of AR-1
 - RBL1 - 10ml of AR-1
 - BLLW - 10ml of AR-1

Thursday 27-07-2023

Split Cells - RAMOS

- Split Ramos 1:2
- Added 10ml of AR-1 to N2
- Plan to split RBL1/BLLW tomorrow

Lucy Drug Randomizer

- Randomized drugs for Lucy Hare mouse experiment

| Drug | Identification |
|------------------|----------------|
| Alectinib | A |
| Erda | B |
| Vehicle | C |
| Alectinib + Erda | D |

Friday 28-07-2023

| Drug | Dilution | Concentration | Units | Starting Concentration |
|------------------|----------|---------------|-------|------------------------|
| Cyclophosphamide | 1 | 20 | uM | 20000 |
| Cyclophosphamide | 2 | 13.3 | uM | 13300 |
| Cyclophosphamide | 3 | 8.89 | uM | 8890 |
| Cyclophosphamide | 4 | 5.93 | uM | 5930 |
| Cyclophosphamide | 5 | 3.95 | uM | 3950 |
| Cyclophosphamide | 6 | 2.63 | uM | 2630 |
| Cyclophosphamide | 7 | 1.76 | uM | 1760 |
| Cyclophosphamide | 8 | 1.17 | uM | 1170 |
| Cyclophosphamide | 9 | 0.78 | uM | 780 |

Friday 06-10-2023

Cell Culture

Made New Media: AR-2

1. Made new media: AR-2

| Solution | ID code | Volume | % Total volume |
|---------------|---------|--------|----------------|
| Advanced RMPI | — | 500 mL | 78% |
| Glutamax | — | 6.5 mL | 20% |
| Pen-Strep | — | 6.5 mL | 1% |
| FBS | — | 128 mL | 1% |

Monday 09-10-2023

Cell Culture

Seeded A20

- Got A20 from Swetha box: 2nd rack in shared -80
- AR-2

Seeded FDC

- Seeded feeders, taken from feeder box, irradiated by Chris
- Will seed N4 PDX tomorrow

Tuesday 10-10-2023

Cell Culture

Seeded N4

- Given by Chris -AR-2
- Seeded onto FDC cells from 09-10
- Future experiments:
 - Barcode PDX
 - EC50: Vincristine/Doxorubicin/Methotrexate/Cyclophosphamide (Active)

Seeded RAMOS-BC

- Seeded barcoded Ramos cells - Pool #4
 - From Chris
 - LN tank 1 - Rack 5 - box 2nd from Bottom
- AR-2
- Future Experiments:
 - Combination Drug pressure: Vin/Dox/Meth

Checked A20

- Look healthy
- ~40% confluency

Wednesday 11-10-2023

Cell Culture

A20 Culture

- ~70% confluent
- Split A20 in half
 - Half in 1xT75 and other half in 1xT175
 - T175 to be frozen down

RAMOS-BC4 Culture

- ~60% confluent
- Spun down at 400xg for 5 min and resuspended in 20ml AR-2

N4 Culture

- Look healthy, attached to feeders with some floaters
- Will add media tomorrow
- Plans:
 - Freeze down aliquots ($2-3 \times 10^6$ per vial needed, ideally freeze down 3x vials)
 - Plan barcoding experiment

Thursday 12-10-2023

Cell Culture

Ramos-BC4

- Infected, will need to grab another vial

- Is the Stock infected? Will thaw 2 different pools this time

A20

- 1 x T75 + 1 x T175 both about ~60% confluent
- Media is slightly turbid, Jamie did a visual check and cells seem clean (A20 have a tendency towards slight turbidity - Jamie)
- Will add 10mL of AR-2 and observe till tomorrow
- If clean:
 - Freeze down 3 vials from T175
 - Set up EC50s with left over cells (Vin/Dox/Meth)
 - Split

N4

- Look healthy, media clean
- About ~40-50% confluent
- Will add 10mL of AR-2 and observe on Friday

Friday 13-10-2023

Cell Culture

A20

- Freeze down (10% DMSO in FBS)
- A20 count: 2.91×10^6 /mL
 - 2mL + 3mL Media
- 3 vials frozen and put in LN Tank 6, Rack 5, Box 6

EC50_231013

EC50 Plate ID: EC50_231013_Plate1

- Doxorubicin top/Vincristine bottom

Doxorubicin Stock: 10mM DMSO

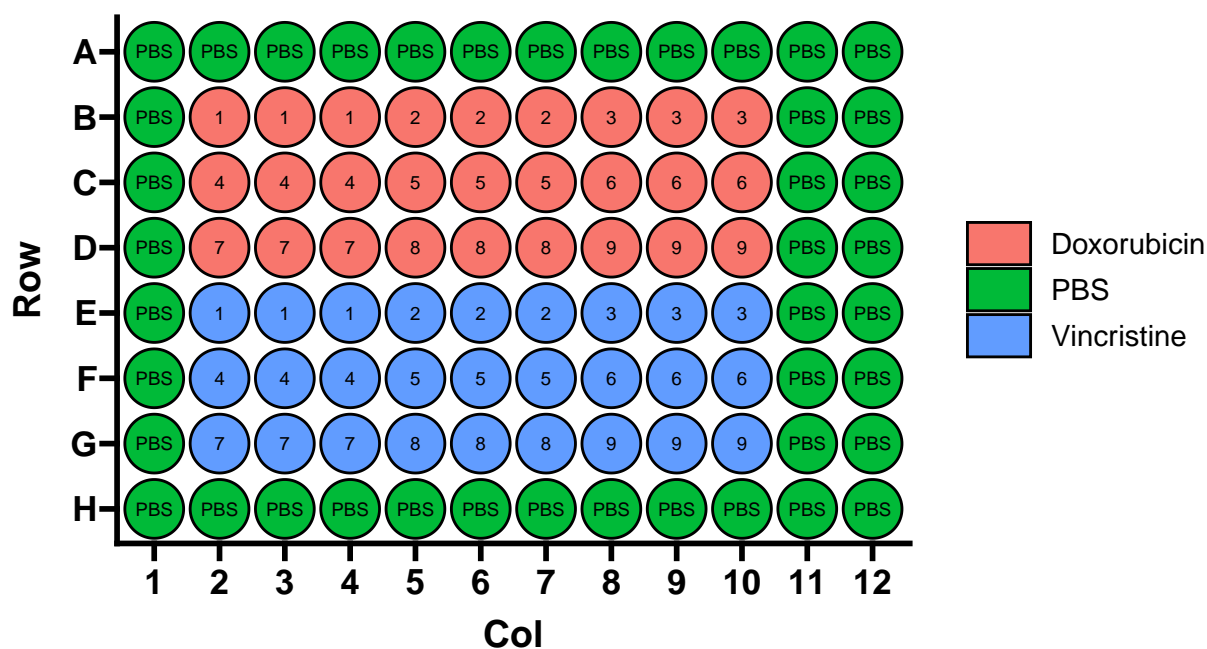
| Drug | Dilution | Concentration | Units | Starting_Concentration |
|-------------|----------|---------------|-------|------------------------|
| Doxorubicin | 1 | 300 | nM | 300000 |
| Doxorubicin | 2 | 200 | nM | 200000 |
| Doxorubicin | 3 | 133 | nM | 133000 |
| Doxorubicin | 4 | 88.9 | nM | 88900 |
| Doxorubicin | 5 | 59.3 | nM | 59300 |
| Doxorubicin | 6 | 39.5 | nM | 39500 |
| Doxorubicin | 7 | 26.3 | nM | 26300 |
| Doxorubicin | 8 | 17.6 | nM | 17600 |
| Doxorubicin | 9 | 11.7 | nM | 11700 |

Vincristine stock: 30 μ M

| Drug | Dilution | Concentration | Units | Starting_Concentration |
|-------------|----------|---------------|-------|------------------------|
| Vincristine | 1 | 10 | nM | 10000 |
| Vincristine | 2 | 7.14 | nM | 7140 |
| Vincristine | 3 | 5.1 | nM | 5100 |
| Vincristine | 4 | 3.64 | nM | 3640 |
| Vincristine | 5 | 2.6 | nM | 2600 |
| Vincristine | 6 | 1.86 | nM | 1860 |
| Vincristine | 7 | 1.33 | nM | 1330 |
| Vincristine | 8 | 0.949 | nM | 949 |
| Vincristine | 9 | 0.678 | nM | 678 |

96 Well Plate Map

EC50_231013_plate1



EC50 Plate ID: EC50_231013_Plate2

- Methotrexate top/Cyclophosphamide bottom

Methotrexate Stock: 10mM DMSO

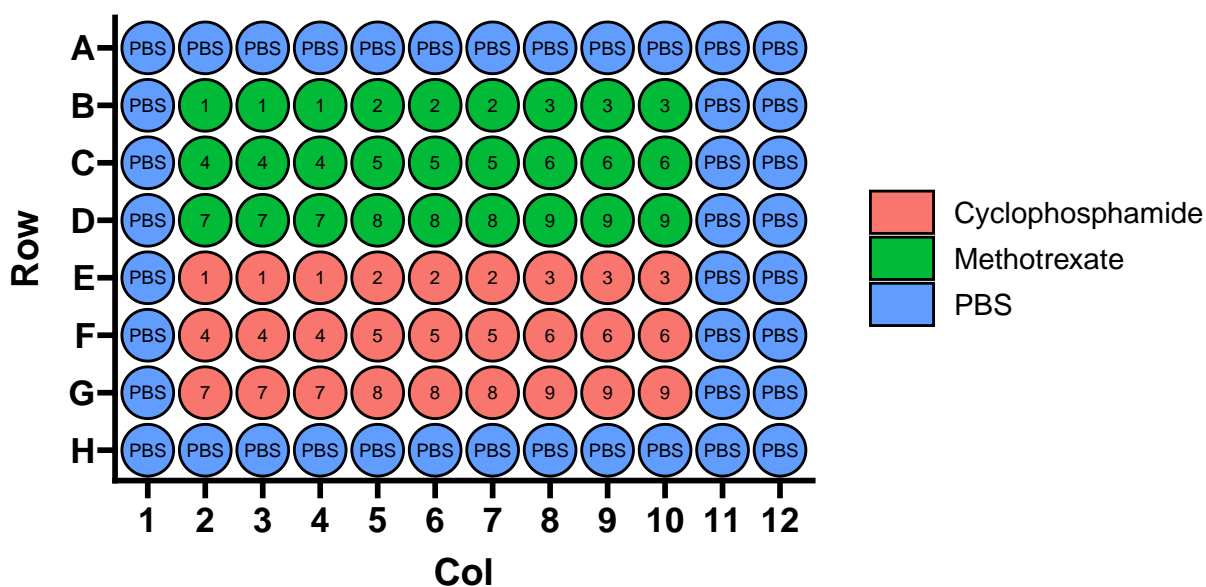
| Drug | Dilution | Concentration | Units | Starting_Concentration |
|--------------|----------|---------------|-------|------------------------|
| Methotrexate | 1 | 500 | nM | 500000 |
| Methotrexate | 2 | 250 | nM | 250000 |
| Methotrexate | 3 | 125 | nM | 125000 |
| Methotrexate | 4 | 62.5 | nM | 62500 |
| Methotrexate | 5 | 31.2 | nM | 31200 |
| Methotrexate | 6 | 15.6 | nM | 15600 |
| Methotrexate | 7 | 7.81 | nM | 7810 |
| Methotrexate | 8 | 3.91 | nM | 3910 |
| Methotrexate | 9 | 1.95 | nM | 1950 |

Cyclophosphamide stock: 20 mM DMSO

| Drug | Dilution | Concentration | Units | Starting_Concentration |
|------------------|----------|---------------|-------|------------------------|
| Cyclophosphamide | 1 | 20 | uM | 20000 |
| Cyclophosphamide | 2 | 14.3 | uM | 14300 |
| Cyclophosphamide | 3 | 10.2 | uM | 10200 |
| Cyclophosphamide | 4 | 7.29 | uM | 7290 |
| Cyclophosphamide | 5 | 5.21 | uM | 5210 |
| Cyclophosphamide | 6 | 3.72 | uM | 3720 |
| Cyclophosphamide | 7 | 2.66 | uM | 2660 |
| Cyclophosphamide | 8 | 1.9 | uM | 1900 |
| Cyclophosphamide | 9 | 1.36 | uM | 1360 |

96 Well Plate Map

EC50_231013_plate2



Monday 16-10-2023

Cell Culture

A20

- Split 1/5
- AR-2

N4

- Split 1/5
- AR-2

Collect EC50_231013

Plate ID: EC50_231013_Plate1

- Vincristine DNW

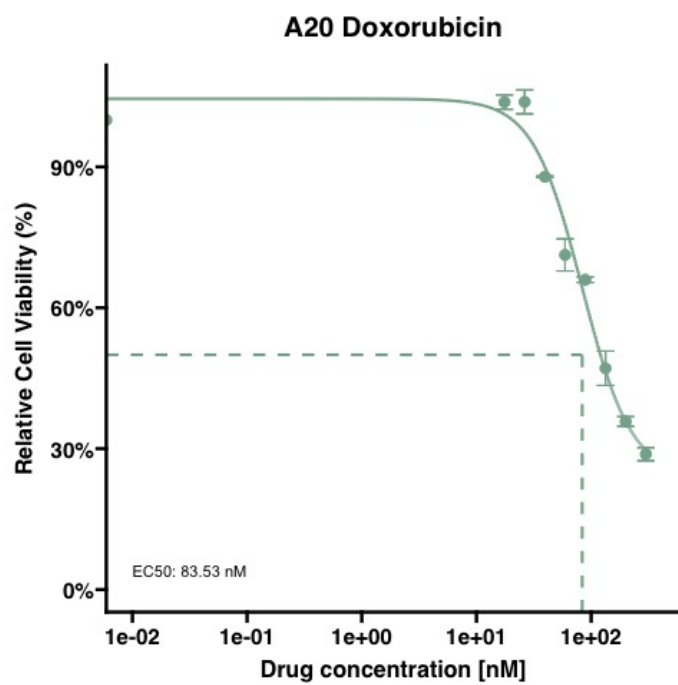


Figure 1: A20 Doxorubicin Results

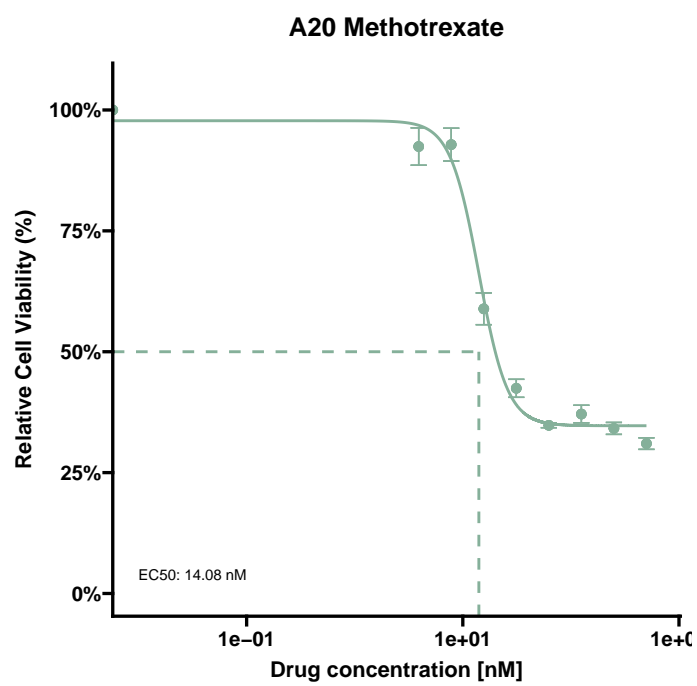


Figure 2: A20 Methotrexate Results

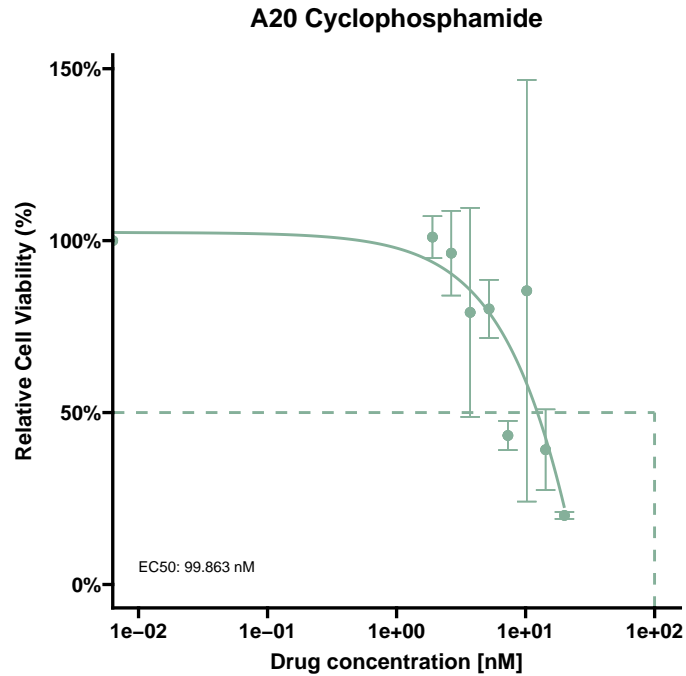


Figure 3: A20 Cyclophosphamide Results

Plate ID: EC50_231013_Plate2

Thursday 19-10-2023

Cell Culture

A20

- Split 1/5
- AR-2

N4

- Split 1/5
- AR-2

Friday 20-10-2023

Cell Culture

A20

- Split 1/5
- AR-2

N4

- Split 1/5
- AR-2

Sunday 22-10-2023

Cell Culture

RAMOS Barcode

- LN Tank 6, Rack 5, Box 6
- Thawed Ramos-BC 1, 5, 6

Monday 23-10-2023

Cell Culture

N4

- ~40-50% confluent
- Added 10mL AR-2

A20

- ~60-70% confluent
- Added 10mL AR-2

Ramos-BC 1

- ~50% confluent
- Added 10mL AR-2

Ramos-BC 5

- ~50% confluent
- Added 10mL AR-2

Ramos-BC 6

- ~50% confluent
- Added 10mL AR-2

Seed Feeders (FDC)

- Count: 4.19×10^5
- Dilute to 4.8×10^4
- Resuspended in 5mL AR-2, added 3.73 mL
- Seeded 1x 6-well plate
 - 2 mL/well
- Seeded 1x 96-well plate (15 wells)
 - 90 μ L/well

Tuesday 23-10-2023

Cell Culture

N4

- No split, maintained

A20

- No split, maintained

Ramos-BC 1

- ~60% confluent
- Split 1/3
- AR-2

Ramos-BC 5

- ~60% confluent
- Split 1/3
- AR-2

Ramos-BC 6

- ~60% confluent
- Split 1/3
- AR-2

FDC

- Looked attached, healthy
- Media fine

Transduce A20/N4 with WILDseq Library

1. Made 2mL cell suspension of 4×10^6 cells/mL
 - A20: 1.91×10^6 count
 - N4: 1.3×10^5 count (took 2 mL of CS from flask)
2. Add 0.5mL of cell suspension to 4 wells of 12 well plate
3. Add virus mixture (listed below) to respective wells:

| Well ID | Virus | Media | HEPES |
|---------|------------|-------------|------------|
| A1 | 25 μ L | 900 μ L | 50 μ L |
| A2 | 10 μ L | 930 μ L | 50 μ L |
| A3 | 1 μ L | 948 μ L | 50 μ L |
| A4 | 0 μ L | 950 μ L | 50 μ L |

4. Centrifuged cells at 600 x g for 1.5 hours at 32C
5. Resuspended cells and transferred to 6 well plate
 - N4 were put on feeder plate which was seeded 23-10-2023

Wednesday 25-10-23

Attempt 1: A20/N4 Barcoding - Media change

- Changed media
- AR-2

Made New Media: AR-3

1. Made new media: AR-3

| Solution | ID code | Volume | % Total volume |
|---------------|---------|--------|----------------|
| Advanced RMPI | — | 500 mL | 78% |
| Glutamax | — | 6.5 mL | 20% |
| Pen-Strep | — | 6.5 mL | 1% |
| FBS | — | 128 mL | 1% |

Thursday 26-10-23

Attempt 1: A20/N4 Barcoding - Sort

- DNW
- No cells were positive
- Possible causes:
 - Virus volume too low
 - Spin speed too low
 - packaging envelope incorrect for mouse cells (GALV)

Friday 27-10-23

Cell Culture

A20

- Maintained
- No Split

N4

- Split 1/4
- AR-3

Ramos-BC 1

- ~60% confluent
- Split 1/4
- AR-3

Ramos-BC 5

- ~60% confluent
- Split 1/4
- AR-3

Ramos-BC 6

- ~60% confluent
- Split 1/4
- AR-3

EC50_231027

Overview

- Treated A20 with Doxorubicin, Vincristine, Cyclophosphamide, Methotrexate
- Concentrations were the same as EC50_231013
- Repeat for biological replicates
- Made an error for plate 1

EC50 Protocol:

1. Create 12 ml cell suspension of 2.5×10^6 cells
 - A20 Cell count: 2.5×10^6
 - 8.33mL AR-3 added to 1.66mL of flask CS
2. Add 90 μ L of CS to respective wells of a 96-well plate
3. Dilute drug to starting concentration indicated in DMSO
4. Serial dilute into DMSO 8 times
5. Add 10uL of diluted drug to respective eppendorf containing 990uL media
6. Add 10 μ L of drugged media to respective well

EC50 Plate ID: EC50_231027_Plate1

- Doxorubicin top/Vincristine bottom

Doxorubicin Stock: 10mM DMSO

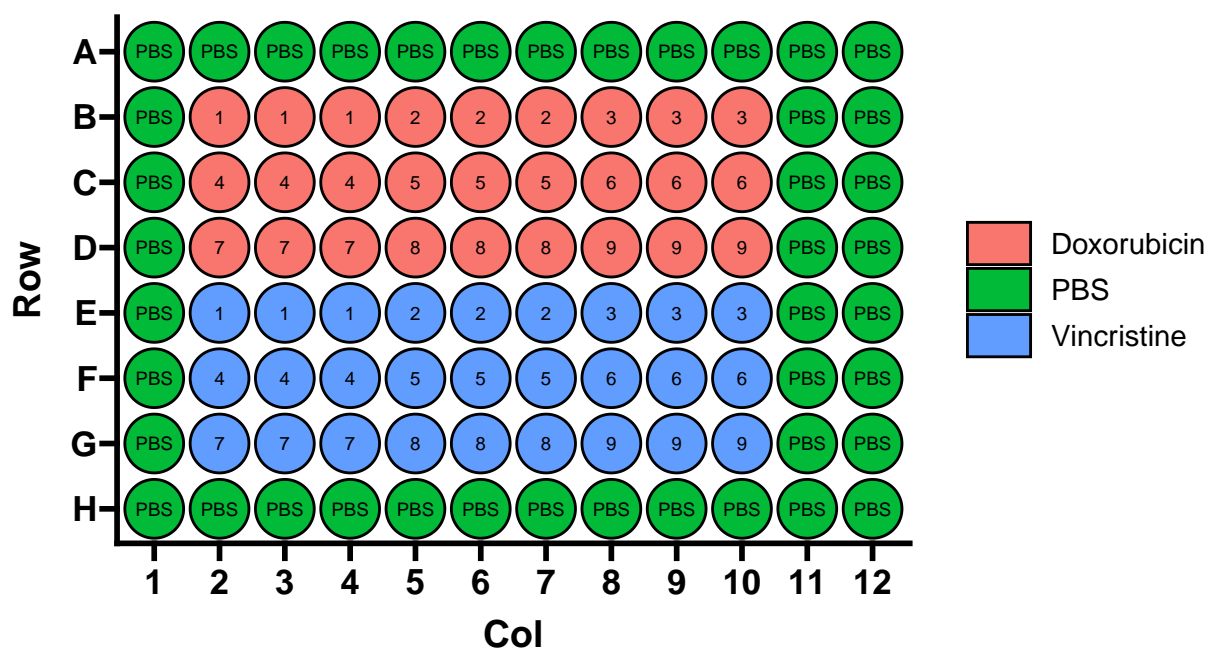
| Drug | Dilution | Concentration | Units | Starting_Concentration |
|-------------|----------|---------------|-------|------------------------|
| Doxorubicin | 1 | 300 | nM | 300000 |
| Doxorubicin | 2 | 200 | nM | 200000 |
| Doxorubicin | 3 | 133 | nM | 133000 |
| Doxorubicin | 4 | 88.9 | nM | 88900 |
| Doxorubicin | 5 | 59.3 | nM | 59300 |
| Doxorubicin | 6 | 39.5 | nM | 39500 |
| Doxorubicin | 7 | 26.3 | nM | 26300 |
| Doxorubicin | 8 | 17.6 | nM | 17600 |
| Doxorubicin | 9 | 11.7 | nM | 11700 |

Vincristine stock: 30 μ M DMSO

| Drug | Dilution | Concentration | Units | Starting_Concentration |
|-------------|----------|---------------|-------|------------------------|
| Vincristine | 1 | 10 | nM | 10000 |
| Vincristine | 2 | 7.14 | nM | 7140 |
| Vincristine | 3 | 5.1 | nM | 5100 |
| Vincristine | 4 | 3.64 | nM | 3640 |
| Vincristine | 5 | 2.6 | nM | 2600 |
| Vincristine | 6 | 1.86 | nM | 1860 |
| Vincristine | 7 | 1.33 | nM | 1330 |
| Vincristine | 8 | 0.949 | nM | 949 |
| Vincristine | 9 | 0.678 | nM | 678 |

96 Well Plate Map

EC50_231013_plate1



NOTE: Made an error by adding Methotrexate and Cyclophosphamide treatments to PLate 1 (as well as listed treatments) - Disregard results

EC50 Plate ID: EC50_231027_Plate2

- Methotrexate top/Cyclophosphamide bottom

Methotrexate Stock: 10mM DMSO

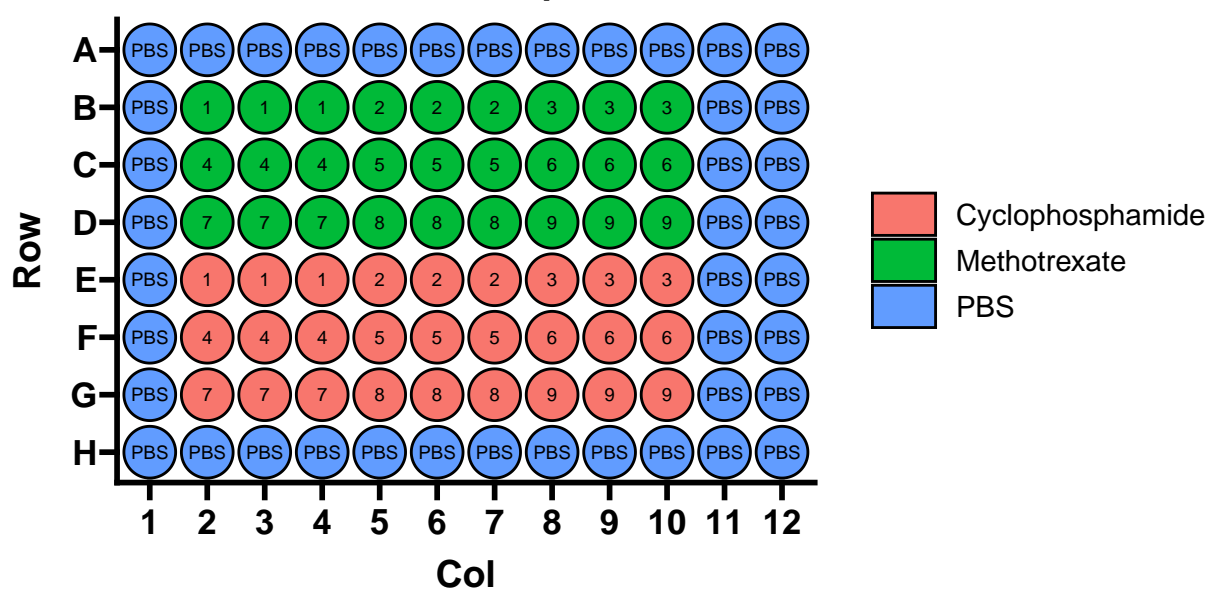
| Drug | Dilution | Concentration | Units | Starting_Concentration |
|--------------|----------|---------------|-------|------------------------|
| Methotrexate | 1 | 500 | nM | 500000 |
| Methotrexate | 2 | 250 | nM | 250000 |
| Methotrexate | 3 | 125 | nM | 125000 |
| Methotrexate | 4 | 62.5 | nM | 62500 |
| Methotrexate | 5 | 31.2 | nM | 31200 |
| Methotrexate | 6 | 15.6 | nM | 15600 |
| Methotrexate | 7 | 7.81 | nM | 7810 |
| Methotrexate | 8 | 3.91 | nM | 3910 |
| Methotrexate | 9 | 1.95 | nM | 1950 |

Cyclophosphamide stock: 20 mM DMSO

| Drug | Dilution | Concentration | Units | Starting_Concentration |
|------------------|----------|---------------|-------|------------------------|
| Cyclophosphamide | 1 | 20 | uM | 20000 |
| Cyclophosphamide | 2 | 14.3 | uM | 14300 |
| Cyclophosphamide | 3 | 10.2 | uM | 10200 |
| Cyclophosphamide | 4 | 7.29 | uM | 7290 |
| Cyclophosphamide | 5 | 5.21 | uM | 5210 |
| Cyclophosphamide | 6 | 3.72 | uM | 3720 |
| Cyclophosphamide | 7 | 2.66 | uM | 2660 |
| Cyclophosphamide | 8 | 1.9 | uM | 1900 |
| Cyclophosphamide | 9 | 1.36 | uM | 1360 |

96 Well Plate Map

EC50_231013_plate2



Monday 27-10-23

Cell Culture

A20

- Split 1/2
- AR-3

N4

- Split 1/2
- AR-3

Ramos-BC 1

- Split 1/2
- AR-3

Ramos-BC 5

- Split 1/2
- AR-3

Ramos-BC 6

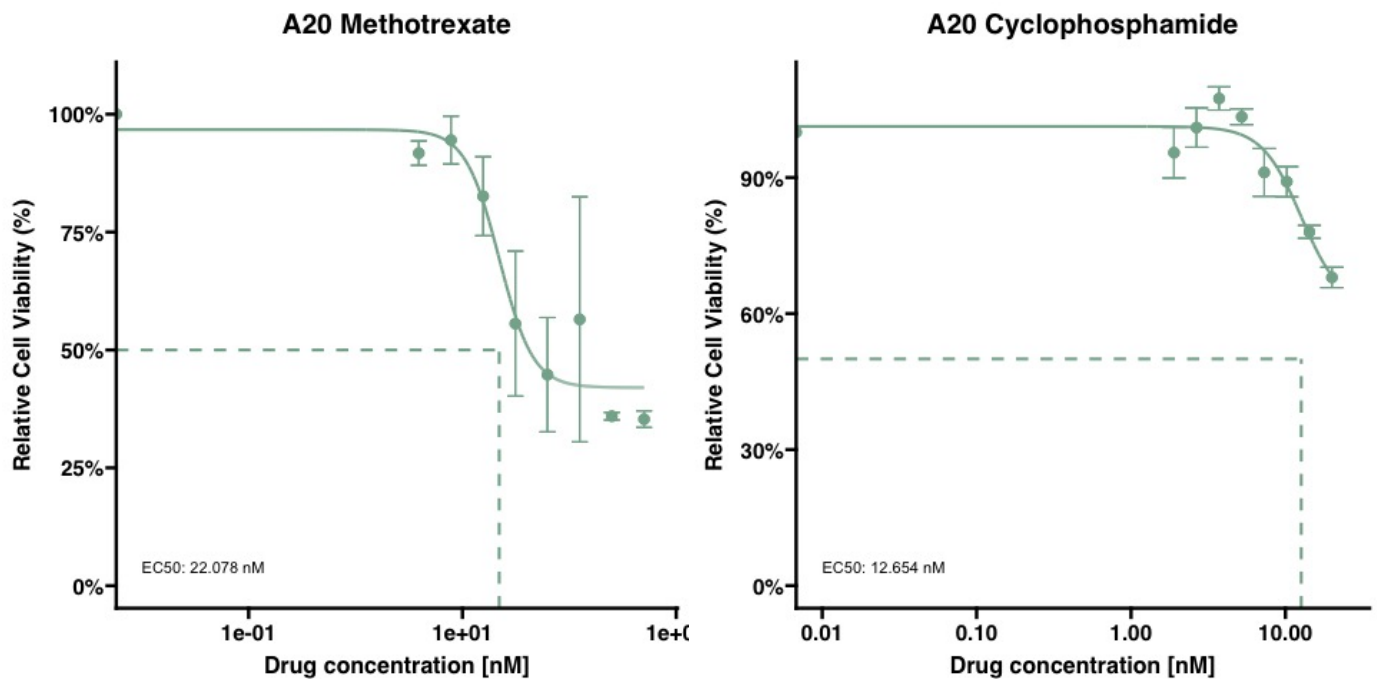
- Split 1/2
- AR-3

Seed Feeders

- Repeating barcoding of A20/N4
- Count: 3.64×10^5
- Dilute to 4.8×10^4
- Resuspended in 5mL AR-2, added 2.58 mL
- Seeded 1x 6-well plate
 - 2 mL/well
- Seeded 1x 96-well plate (15 wells)
 - 90 μ L/well

Collect EC50_231027

Plate ID: EC50_231027_Plate



Tuesday 31-10-2023

Attempt 2: Transduce A20/N4 with WILDseq Library

1. Made 2mL cell suspension of 4×10^6 cells/mL
 - A20: 2.18×10^6 count
 - N4: 3.06×10^5 count (took 2 mL of CS from flask)

2. Add 0.5mL of cell suspension to 4 wells of 12 well plate
3. Add virus mixture (listed below) to respective wells:

| Well ID | Virus | Media | HEPES |
|---------|-------------|-------------|------------|
| A1 | 250 μ L | 225 μ L | 25 μ L |
| A2 | 150 μ L | 150 μ L | 25 μ L |
| A3 | 50 μ L | 50 μ L | 25 μ L |
| A4 | 0 μ L | 0 μ L | 25 μ L |

4. Centrifuged cells at 1000 x g for 1.5 hours at 0C

- Mistakenly set centrifuge to 0C

5. Resuspended cells and transferred to 6 well plate

- N4 were put on feeder plate which was seeded 23-10-2023

Drug Pressure Experiment 1: RAMOS Day 1

Overview

- Designing combination treatment to reflect common BL treatments in clinic
- Plan to analyse differences in barcode representation between baseline, DMSO, and treatment-resistant
- Drug concentrations:

| Drug | Treatment Day | Concentration | Stock |
|------------------|---------------|---------------|---------------------------|
| Doxorubicin | Day 2 | 80nM | 100 μ M working stock |
| Vincristine | Day 1 | 10nM | 30 μ M |
| Methotrexate | Day 1 | 20nM | 100 μ M working stock |
| Cyclophosphamide | Day 2/3/4 | 20 μ M | 20 μ M working stock |

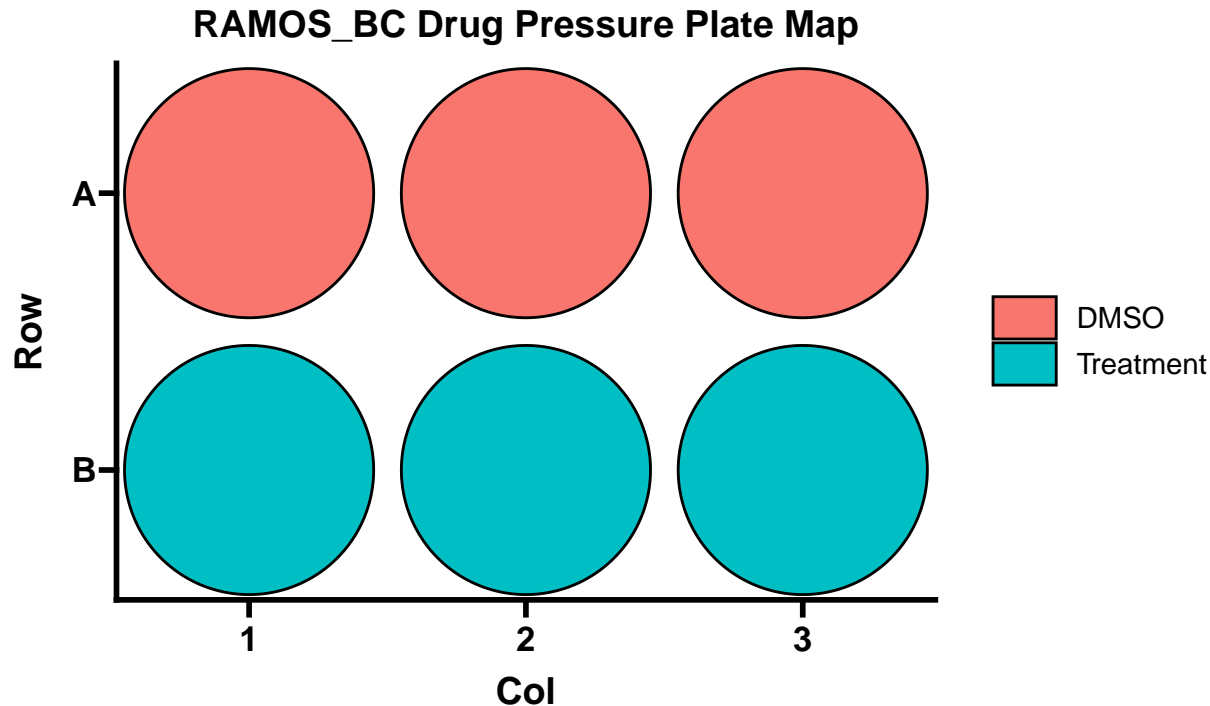
- Plan:
 1. Collect sample at baseline
 2. Complete 4 day treatment course of RAMOS-BC cells (with corresponding DMSO controls)
 3. Allow treatment(T) cells to recover and collect sample
 4. Complete 4 day treatment course again and collect sample
- Possible future experiments:
 - EC50's of T vs WT cells (are T RAMOS cells drug resistant?)

Day 1

Seeded RAMOS BC

- Seeded 3x 6 well plates with RAMOS-BC 1, 5, 6 respectively

96 Well Plate Map



- Seeded 0.25×10^6 cells/well in 3mL AR-3
- Treatment wells

| Drug | Treatment Day | Concentration | Stock | Stock Volume | Media Volume |
|--------------|---------------|---------------|---------------------------|-------------------|--------------|
| DMSO | Day 1 | - | - | 14.4 μ L | 27 mL AR-3 |
| Vincristine | Day 1 | 10nM | 30 μ M | 9 μ L | 27 mL AR-3 |
| Methotrexate | Day 1 | 20nM | 100 μ M working stock | 5.4 μ L of WS | 27 mL AR-3 |

Collected RAMOS Baseline

- Took 1mL CS from respective flask and transferred to eppendorf
- Spun down 2500 x g at 4C
- Removed Media
- Snap froze in LN
- Put in -80 CM Box 1

Wednesday 01-11-2023

Drug Pressure Experiment 1: RAMOS Day 2

- Changed media
- Spun down at 600 x g for 5 min
- Resuspended in Day 2 Treatment:

| Drug | Treatment Day | Concentration | Stock | Stock Volume | Media Volume |
|------------------|------------------|---------------|------------------------------|-----------------------------------|--------------|
| DMSO | Day 1 | - | - | 48.60 μ L | 27 mL AR-3 |
| Doxorubicin | Day 1 | 80nM | 100 μ M working stock | 21.60 μ L of working stock | 27 mL AR-3 |
| Cyclophosphamide | Day 1 | 20 μ M | 20mM | 27 μ L | 27 mL AR-3 |

Attempt 2: Barcoding A20/N4

- Sanity check
- Ran samples on E6 flow
- A20 negative
- Suggests envelope protein is not compatible with mouse cells
- N4 dead
- Re-make WILDseq virus with VSVG packaging envelope

Thursday 02-11-2023

Cell Culture

A20

- Split 1/2
- AR-3

N4

- Split 1/2
- AR-3

Ramos-BC 1

- Split 1/4
- AR-3

Ramos-BC 5

- Split 1/4
- AR-3

Ramos-BC 6

- Split 1/4
- AR-3

Seeded Feeders

- 1 vial
- Split into 2 T75 flasks

Drug Pressure Experiment 1: RAMOS Day 3

- Changed media
- Spun down at 600 x g for 5 min
- Resuspended in Day 3 Treatment:

| Drug | Treatment Day | Concentration | Stock | Stock Volume | Media Volume |
|------------------|------------------|---------------|-------|--------------|--------------|
| DMSO | Day 1 | - | - | 27 μ L | 27 mL AR-3 |
| Cyclophosphamide | Day 1 | 20 μ M | 20mM | 27 μ L | 27 mL AR-3 |

Friday 03-11-2023

Cell Culture

A20

- Split 1/2
- AR-3

N4

- Split into two new feeder flasks
 - Seeded 02-11-2023
 - N4.1 / N4.2
 - 1 flask to transduce, 1 flask to freeze down
- AR-3

Ramos-BC 1

- Add 10mL AR-3

Ramos-BC 5

- Add 10mL AR-3

Ramos-BC 6

- Add 10mL AR-3

Drug Pressure Experiment 1: RAMOS Day 4

- Treatment cells too sick to continue
- Changed media
- Spun down at 600 x g for 5 min
- Resuspended both DMSO and Treatment cells in fresh AR-3
- Allow to grow

Monday 06-11-2023

Cell Culture

A20

- Split 1/2
- AR-3

N4.1

- Healthy/growing
- Leave till tomorrow

N4.2

- Healthy/growing
- Leave till tomorrow

Ramos-BC 1

- Froze down
- 3x vials
- FBS +10% DMSO

Ramos-BC 5

- Froze down
- 3x vials
- FBS +10% DMSO

Ramos-BC 6

- Froze down
- 3x vials
- FBS +10% DMSO

Seed Feeders

- For WILDseq transduction of N4 (tuesday?)
- 1x 4 wells of 6-well plate, 15wells of 96-well plate
- Dilute in 8ml
- 2mL/well of 6well
- 90 μ L/well of 96well plate

Drug Pressure Experiment 1: RAMOS Day 7

- Split DMSO cells 1/6
- Add 1mL AR-3 to treatment cells
 - treatment cells still sick, allow to grow for a few more days
 - Perhaps transfer to smaller well?

Tuesday 07/11/2023

Made New Media: AR-4

1. Made new media: AR-4

| Solution | ID code | Volume | % Total volume |
|---------------|---------|--------|----------------|
| Advanced RMPI | — | 500 mL | 78% |
| Glutamax | — | 6.5 mL | 20% |
| Pen-Strep | — | 6.5 mL | 1% |
| FBS | — | 128 mL | 1% |

Cell Culture

A20

- Split 1/6
- AR-4

N4.1

- Healthy/growing
- Leave till tomorrow

N4.2

- Froze down
- 2x vials
- FBS +10% DMSO

Drug Pressure Experiment 1: RAMOS Day 5

- Split D 1/6

Attempt 3: Transduce N4 with WILDseq Library

1. Made 2mL cell suspension of 4×10^6 cells/mL
 - N4: took 10ml of 20ml CS, spun down and resuspended in 2mL AR-4
2. Add 0.5mL of cell suspension to 4 wells of 12 well plate
3. Add virus mixture (listed below) to respective wells:

| Well ID | Virus | Media | HEPES |
|---------|-------------|-------------|------------|
| A1 | 250 μ L | 225 μ L | 25 μ L |
| A2 | 150 μ L | 150 μ L | 25 μ L |
| A3 | 50 μ L | 50 μ L | 25 μ L |
| A4 | 0 μ L | 0 μ L | 25 μ L |

4. Centrifuged cells at 1000 x g for 1.5 hours at 32C
5. Resuspended cells and transferred to 6 well plate
 - N4 were put on feeder plate which was seeded 06-11-2023

Wednesday 07/11/2023

Attempt 3: Barcoding N4

- Changed Media
- Cells look healthy ## Drug Pressure Experiment 1: RAMOS Day 6
- Checked treatment cells with trypan blue
- Everything was dead
- Ended experiment: will reconsider approach
 - Treat Ramos with individual agents as well as combination
 - Treat for 3 days and recover or 12 days and recover(?)

Thursday 07/11/2023

Attempt 3: Barcoding N4

- Booked sort for tomorrow
- Changed media
- Sanity check on E6 Fortessa ## Cell Culture

A20
N4