

Turner Lab Notebook

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Tuesday 02-05-2023

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

Wednesday 03-05-2023

1. Splitting 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

Monday 19-06-2023

1. Splitting cells

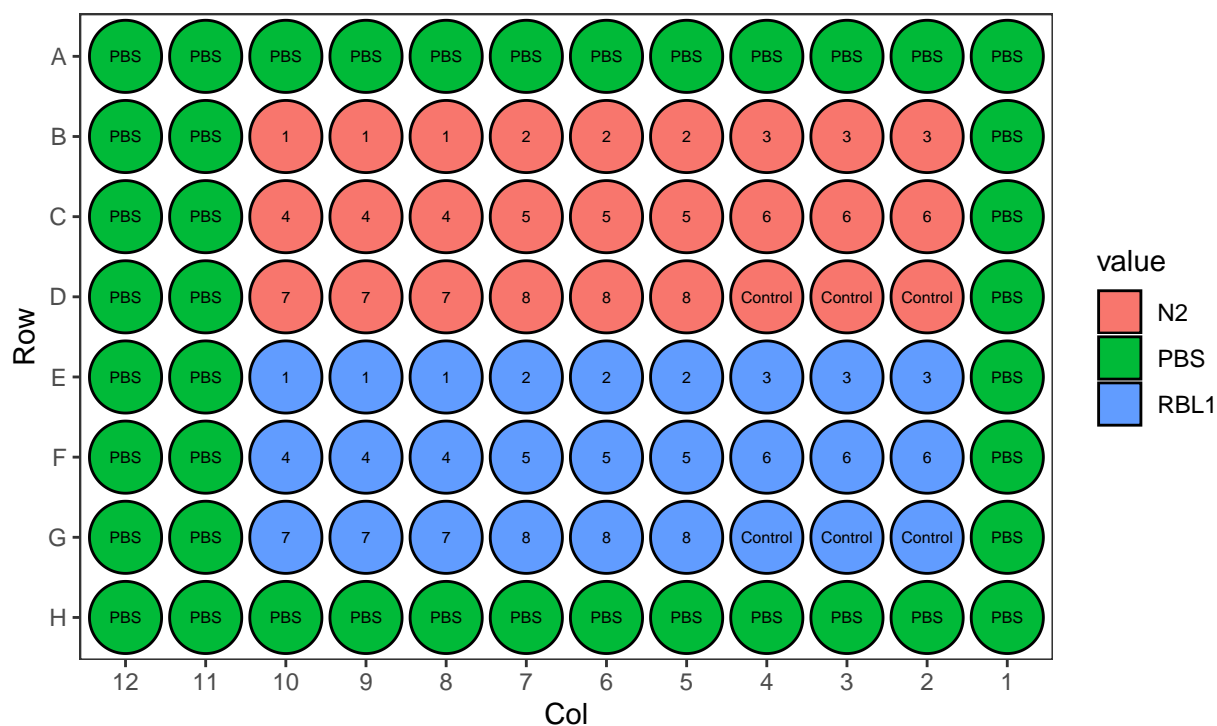
- Split N2/RBL1/JIRE
- Added 10 mL media to BLLW
 - Lots of dead cells observed

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

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 μ L to 30 μ L 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 μ M
 - 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 μ L 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 μ L	4.4 μ L
SSIV RT	1 μ L	1.1 μ L
100 mM DTT	1 μ L	1.1 μ L
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

Steps	Time
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_SPTTest3_cDNA

[High-Sensitivity DNA ScreenTape Protocol](#)

[Results](#)

* Did not work, bands present but faint and not at 200bp

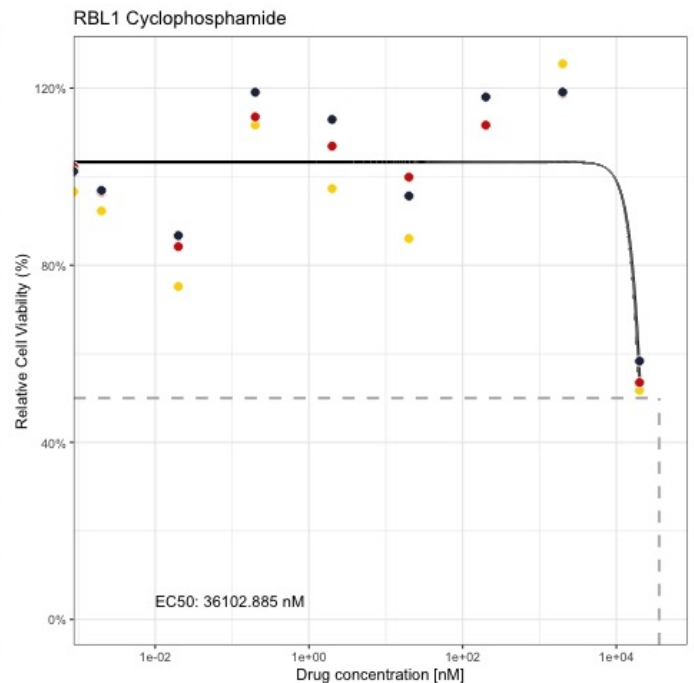
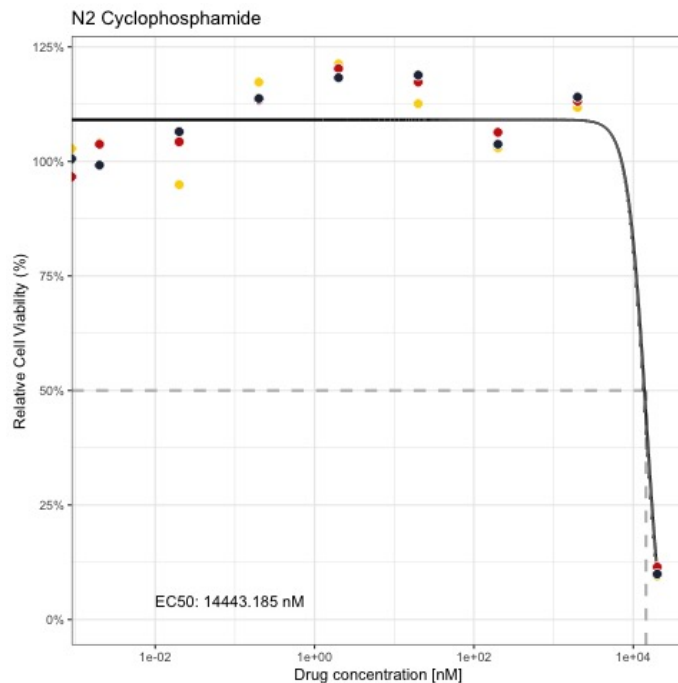
* Potential solutions:

- Repeat PCR with increased cycles
- Repeat RNA extration 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)

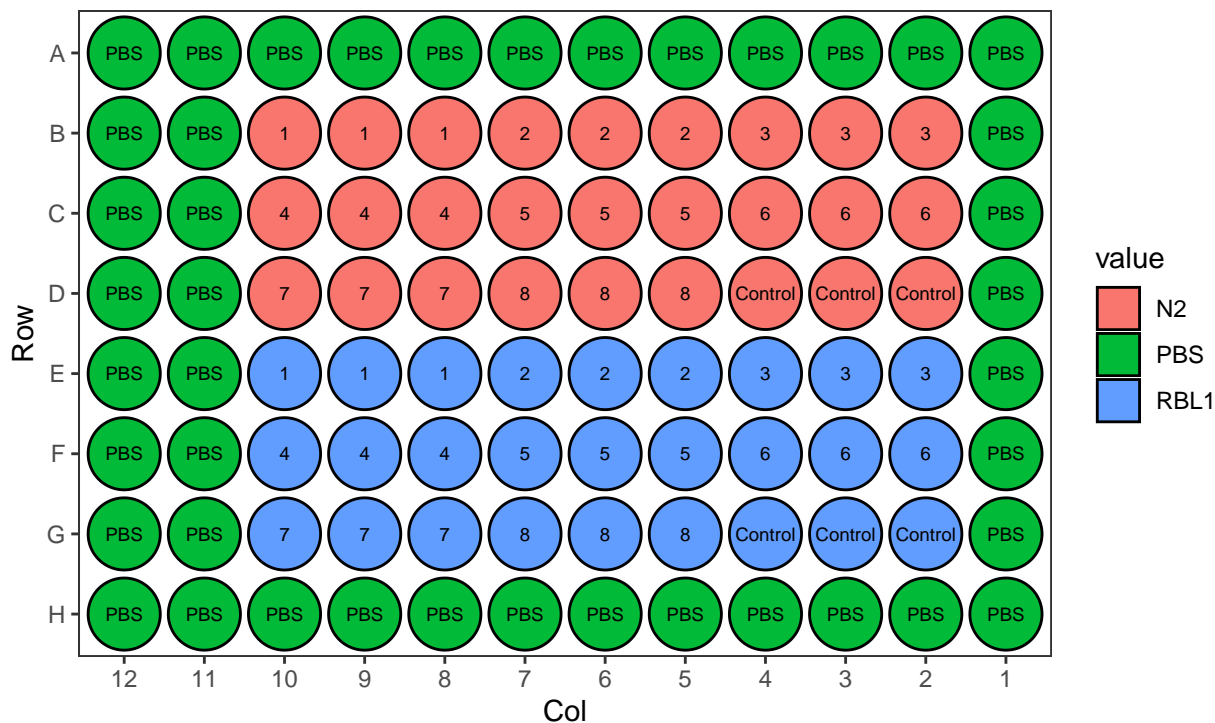
2. Split JIRE 1/2
3. Split N2/RBL1 1/4

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