

Final Project

Katrina Lee Marcy

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Project Outline

Acute myeloid leukemia (AML) is a clinically devastating cancer (<30% 5 year-survival rate) that has been extraordinarily difficult to treat due to its complex genetic subtypes. One shared hallmark of AML is the arrest of leukemic myeloblasts at an immature and self-renewing stage of development. Therapies that are able to overcome this arrest represent a powerful treatment strategy. Sykes et al developed a cellular model of HoxA9-enforced myeloid differentiation arrest to use in an unbiased phenotypic flow-cytometry based screen. In this system, 96-well and 384-well plates are treated with titrated small compounds in order to assess viability, changes in cellular phenotype (based on cell surface markers), and differentiation status (ie: are the cells able to overcome their differentiation arrest to take on a neutrophil phenotype versus a myeloblast progenitor phenotype). Compounds that allow for differentiation are considered “hits” in this system and should be further studied to identify details on mechanism of action and pharmacokinetics. Developing code to analyze the output (over 330,000 compounds will need to be tested at multiple doses) will be essential for quickly identifying “hits”.

Import File

Import Excel file for analysis. The goal will be to set up a platform to analyze flow cytometry data from 96-well and 384-well plates for cell viability (determine whether compounds are overly toxic) and cell surface markers (determine changes in differentiation status) caused by treatment with DHODH inhibitor compounds.

Determine Viability Normalized to Control

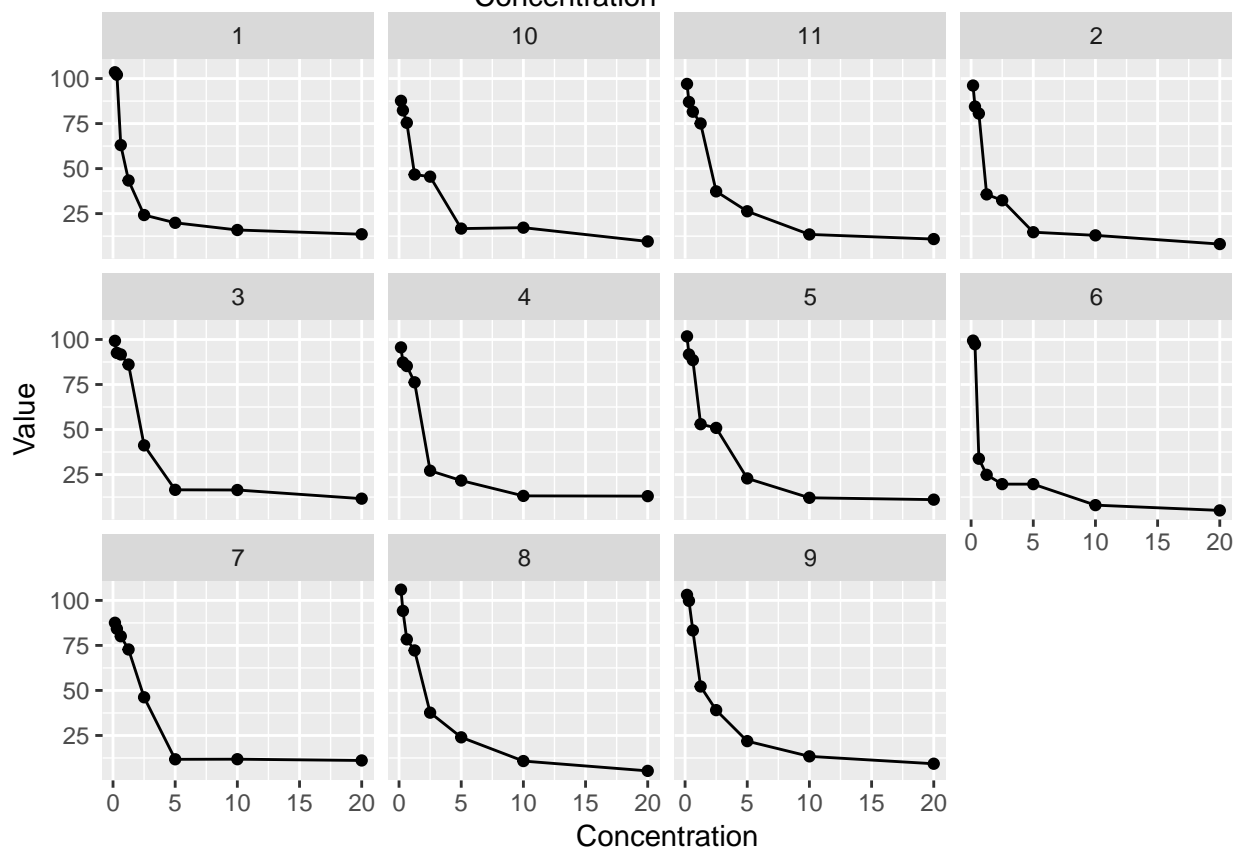
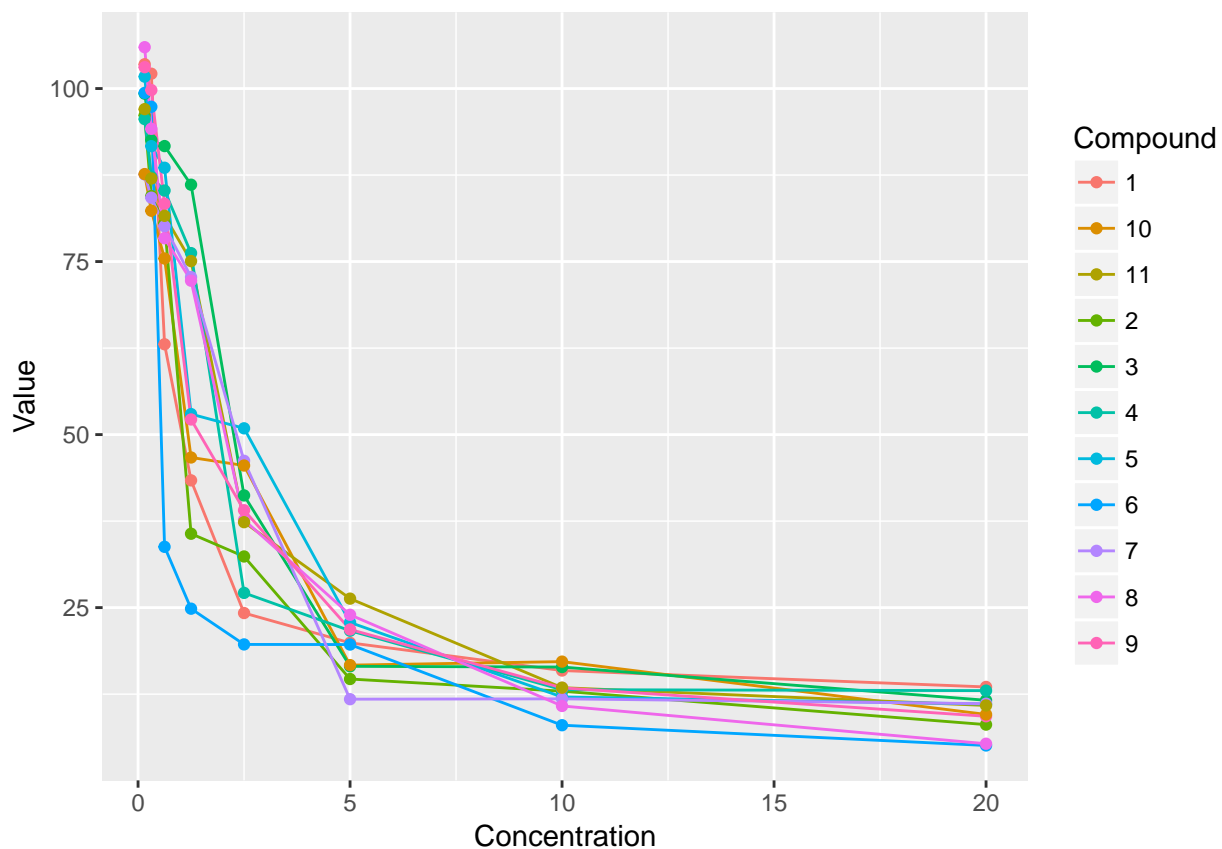
Calculate viability as a percentage normalized to an average of the DMSO wells.

| ## | Compound.1 | Compound.2 | Compound.3 | Compound.4 | Compound.5 | Compound.6 |
|------|------------|------------|------------|-------------|-------------|------------|
| ## 1 | 13.54 | 8.10 | 11.61 | 13.01 | 11.07 | 5.08 |
| ## 2 | 15.91 | 12.93 | 16.42 | 13.14 | 12.09 | 8.01 |
| ## 3 | 19.91 | 14.68 | 16.52 | 21.67 | 22.86 | 19.66 |
| ## 4 | 24.22 | 32.39 | 41.21 | 27.13 | 50.91 | 19.68 |
| ## 5 | 43.39 | 35.68 | 86.10 | 76.22 | 52.96 | 24.84 |
| ## 6 | 63.05 | 80.58 | 91.69 | 85.26 | 88.55 | 33.79 |
| ## 7 | 102.14 | 84.46 | 92.54 | 87.24 | 91.67 | 97.37 |
| ## 8 | 103.51 | 96.14 | 99.26 | 95.59 | 101.72 | 99.37 |
| ## | Compound.7 | Compound.8 | Compound.9 | Compound.10 | Compound.11 | DMSO |
| ## 1 | 11.12 | 5.34 | 9.30 | 9.57 | 10.87 | 92.00 |
| ## 2 | 11.82 | 10.79 | 13.40 | 17.20 | 13.44 | 103.55 |
| ## 3 | 11.77 | 23.97 | 21.84 | 16.69 | 26.30 | 91.14 |
| ## 4 | 46.21 | 37.67 | 39.07 | 45.53 | 37.34 | 99.37 |
| ## 5 | 72.76 | 72.23 | 52.18 | 46.69 | 75.07 | 93.57 |
| ## 6 | 80.11 | 78.37 | 83.35 | 75.44 | 81.61 | 104.88 |
| ## 7 | 84.23 | 94.18 | 99.78 | 82.35 | 87.01 | 110.41 |
| ## 8 | 87.59 | 105.97 | 103.08 | 87.64 | 97.02 | 105.09 |

Viability Curves

```
## -----  
## data.table + dplyr code now lives in dtplyr.  
## Please library(dtplyr)!  
## -----  
  
##  
## Attaching package: 'data.table'  
  
## The following objects are masked from 'package:dplyr':  
##  
##   between, first, last  
  
## The following object is masked from 'package:purrr':  
##  
##   transpose  
  
##      Concentration      DMSO    variable    value Compound  
## 1:      20.00000    91.99632 Compound.1    13.540708        1  
## 2:      10.00000   103.54761 Compound.1    15.909614        1  
## 3:       5.00000    91.13960 Compound.1    19.911454        1  
## 4:       2.50000    99.37328 Compound.1    24.218031        1  
## 5:       1.25000    93.56601 Compound.1    43.387764        1  
## 6:       0.62500   104.87580 Compound.1    63.051978        1  
## 7:       0.31250   110.41283 Compound.1   102.138914        1  
## 8:       0.15625   105.08855 Compound.1   103.507360        1  
## 9:      20.00000    91.99632 Compound.2     8.095676        2  
## 10:     10.00000   103.54761 Compound.2    12.925483        2  
## 11:      5.00000    91.13960 Compound.2    14.684913        2  
## 12:      2.50000    99.37328 Compound.2    32.388454        2  
## 13:      1.25000    93.56601 Compound.2    35.677323        2  
## 14:      0.62500   104.87580 Compound.2    80.577277        2  
## 15:      0.31250   110.41283 Compound.2    84.458372        2  
## 16:      0.15625   105.08855 Compound.2    96.136155        2  
## 17:     20.00000    91.99632 Compound.3    11.614535        3  
## 18:     10.00000   103.54761 Compound.3    16.415593        3  
## 19:      5.00000    91.13960 Compound.3    16.519089        3  
## 20:      2.50000    99.37328 Compound.3    41.208602        3  
## 21:      1.25000    93.56601 Compound.3    86.102806        3  
## 22:      0.62500   104.87580 Compound.3    91.685833        3  
## 23:      0.31250   110.41283 Compound.3    92.542548        3  
## 24:      0.15625   105.08855 Compound.3    99.258280        3  
## 25:     20.00000    91.99632 Compound.4    13.005980        4  
## 26:     10.00000   103.54761 Compound.4    13.143974        4  
## 27:      5.00000    91.13960 Compound.4    21.665133        4  
## 28:      2.50000    99.37328 Compound.4    27.127415        4  
## 29:      1.25000    93.56601 Compound.4    76.218951        4  
## 30:      0.62500   104.87580 Compound.4    85.263339        4  
## 31:      0.31250   110.41283 Compound.4    87.235511        4  
## 32:      0.15625   105.08855 Compound.4    95.589926        4  
## 33:     20.00000    91.99632 Compound.5    11.068307        5  
## 34:     10.00000   103.54761 Compound.5    12.091766        5  
## 35:      5.00000    91.13960 Compound.5    22.855336        5  
## 36:      2.50000    99.37328 Compound.5    50.908464        5
```

| | | | | | |
|--------|---------------|-----------|-------------|------------|----------|
| ## 37: | 1.25000 | 93.56601 | Compound.5 | 52.961132 | 5 |
| ## 38: | 0.62500 | 104.87580 | Compound.5 | 88.552208 | 5 |
| ## 39: | 0.31250 | 110.41283 | Compound.5 | 91.674333 | 5 |
| ## 40: | 0.15625 | 105.08855 | Compound.5 | 101.724931 | 5 |
| ## 41: | 20.00000 | 91.99632 | Compound.6 | 5.082797 | 6 |
| ## 42: | 10.00000 | 103.54761 | Compound.6 | 8.009430 | 6 |
| ## 43: | 5.00000 | 91.13960 | Compound.6 | 19.664213 | 6 |
| ## 44: | 2.50000 | 99.37328 | Compound.6 | 19.681463 | 6 |
| ## 45: | 1.25000 | 93.56601 | Compound.6 | 24.844756 | 6 |
| ## 46: | 0.62500 | 104.87580 | Compound.6 | 33.785649 | 6 |
| ## 47: | 0.31250 | 110.41283 | Compound.6 | 97.366605 | 6 |
| ## 48: | 0.15625 | 105.08855 | Compound.6 | 99.373275 | 6 |
| ## 49: | 20.00000 | 91.99632 | Compound.7 | 11.120055 | 7 |
| ## 50: | 10.00000 | 103.54761 | Compound.7 | 11.821527 | 7 |
| ## 51: | 5.00000 | 91.13960 | Compound.7 | 11.769779 | 7 |
| ## 52: | 2.50000 | 99.37328 | Compound.7 | 46.210902 | 7 |
| ## 53: | 1.25000 | 93.56601 | Compound.7 | 72.757590 | 7 |
| ## 54: | 0.62500 | 104.87580 | Compound.7 | 80.105796 | 7 |
| ## 55: | 0.31250 | 110.41283 | Compound.7 | 84.228381 | 7 |
| ## 56: | 0.15625 | 105.08855 | Compound.7 | 87.591996 | 7 |
| ## 57: | 20.00000 | 91.99632 | Compound.8 | 5.335787 | 8 |
| ## 58: | 10.00000 | 103.54761 | Compound.8 | 10.792318 | 8 |
| ## 59: | 5.00000 | 91.13960 | Compound.8 | 23.965041 | 8 |
| ## 60: | 2.50000 | 99.37328 | Compound.8 | 37.666743 | 8 |
| ## 61: | 1.25000 | 93.56601 | Compound.8 | 72.234361 | 8 |
| ## 62: | 0.62500 | 104.87580 | Compound.8 | 78.369365 | 8 |
| ## 63: | 0.31250 | 110.41283 | Compound.8 | 94.181233 | 8 |
| ## 64: | 0.15625 | 105.08855 | Compound.8 | 105.974011 | 8 |
| ## 65: | 20.00000 | 91.99632 | Compound.9 | 9.303128 | 9 |
| ## 66: | 10.00000 | 103.54761 | Compound.9 | 13.402714 | 9 |
| ## 67: | 5.00000 | 91.13960 | Compound.9 | 21.843376 | 9 |
| ## 68: | 2.50000 | 99.37328 | Compound.9 | 39.069687 | 9 |
| ## 69: | 1.25000 | 93.56601 | Compound.9 | 52.184913 | 9 |
| ## 70: | 0.62500 | 104.87580 | Compound.9 | 83.348666 | 9 |
| ## 71: | 0.31250 | 110.41283 | Compound.9 | 99.775759 | 9 |
| ## 72: | 0.15625 | 105.08855 | Compound.9 | 103.081877 | 9 |
| ## 73: | 20.00000 | 91.99632 | Compound.10 | 9.567617 | 10 |
| ## 74: | 10.00000 | 103.54761 | Compound.10 | 17.197562 | 10 |
| ## 75: | 5.00000 | 91.13960 | Compound.10 | 16.691582 | 10 |
| ## 76: | 2.50000 | 99.37328 | Compound.10 | 45.532429 | 10 |
| ## 77: | 1.25000 | 93.56601 | Compound.10 | 46.693882 | 10 |
| ## 78: | 0.62500 | 104.87580 | Compound.10 | 75.442732 | 10 |
| ## 79: | 0.31250 | 110.41283 | Compound.10 | 82.348206 | 10 |
| ## 80: | 0.15625 | 105.08855 | Compound.10 | 87.637994 | 10 |
| ## 81: | 20.00000 | 91.99632 | Compound.11 | 10.867065 | 11 |
| ## 82: | 10.00000 | 103.54761 | Compound.11 | 13.437213 | 11 |
| ## 83: | 5.00000 | 91.13960 | Compound.11 | 26.299448 | 11 |
| ## 84: | 2.50000 | 99.37328 | Compound.11 | 37.344756 | 11 |
| ## 85: | 1.25000 | 93.56601 | Compound.11 | 75.068997 | 11 |
| ## 86: | 0.62500 | 104.87580 | Compound.11 | 81.606486 | 11 |
| ## 87: | 0.31250 | 110.41283 | Compound.11 | 87.005520 | 11 |
| ## 88: | 0.15625 | 105.08855 | Compound.11 | 97.021619 | 11 |
| ## | Concentration | DMSO | variable | value | Compound |



Calculate IC50

Calculate the concentration at which 50% of the cells are dead. This value can be used in further validation studies.

1. Take concentration value from each graph corresponding to 50% on y-axis
2. Convert Concentration in μM to $\log\text{M}$

Calculate Differentiation

Determine which compounds cause differentiation of myeloblasts by calculating the percent of cells double-positive for Cd11b and GR-1, two cell surface markers of mature myeloid cells.