cmQTL pilot, Variance Analysis

Aparna Nathan

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Data

- Two time points: 6 or 24 hours
- Four densities at each time point
 - For 6 hrs: 5000, 10000, 15000, 20000
 - For 24 hrs: 1000, 2000, 3000, 4000
- Three different aggregated measurements per well:
 - All cells
 - Isolated cells
 - Colonized cells
- Additional covariates: proportion of isolated cells/well
 - Also have well ID, but didn't use it

1. Is there a relationship between cell line and colony formation?

Mixed effects model with random effects for cell line and density

proportion.isolated ~ 1 + (1 | cell.line) + (1 | density)

```
Random effects:

Groups

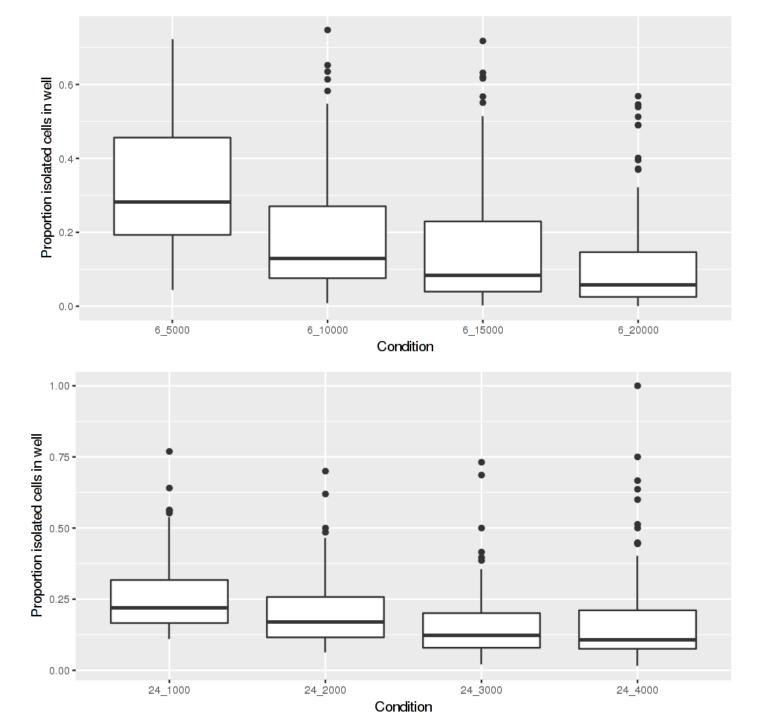
pilot.metadata.6$Metadata_line_ID

pilot.metadata.6$Metadata_plating_density (Intercept) 0.008017 0.08954

Residual

0.008556 0.09250
```

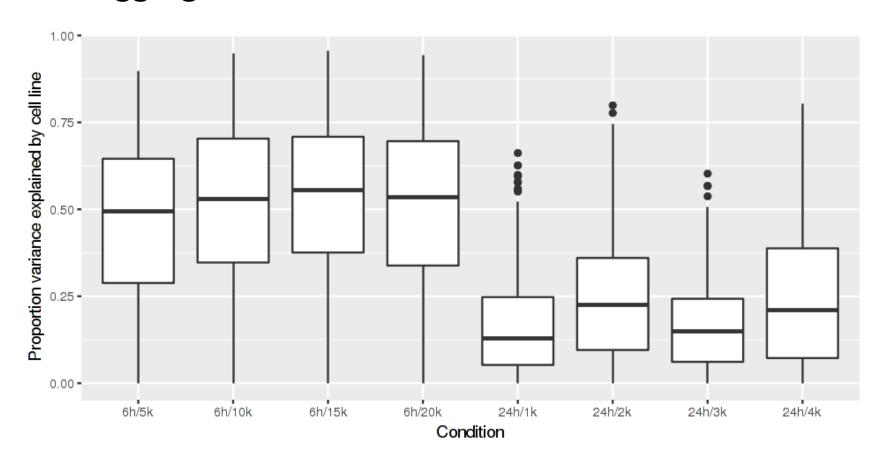
~60% of variance in proportion of isolated cells (in a well) is explained by cell line



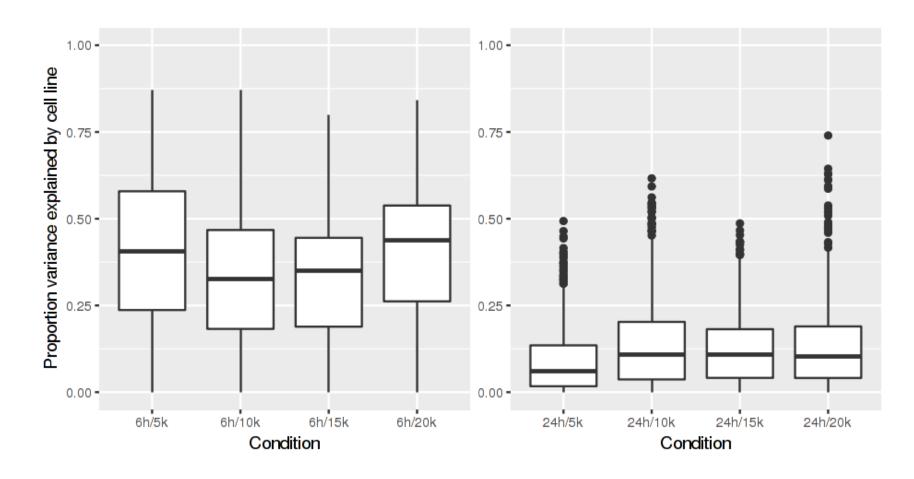
For each condition, a mixed effects model for each trait with random effect for cell line

trait ~ 1 + (1 | cell.line)

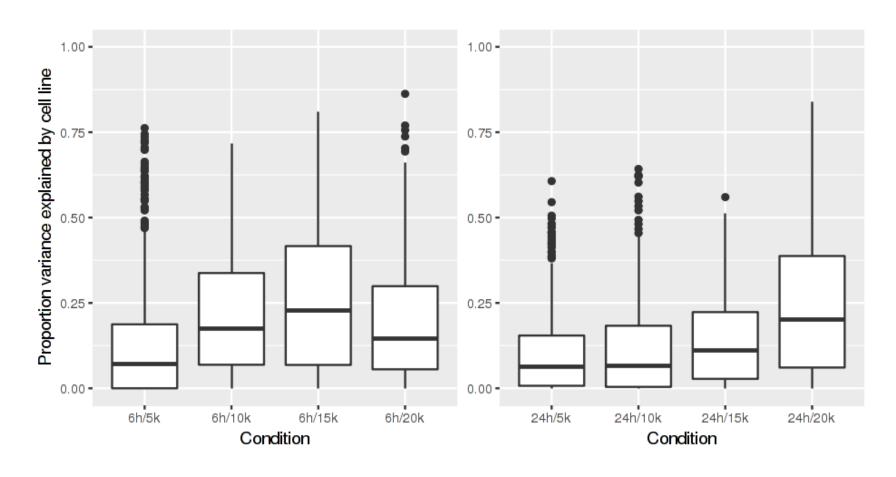
First, in data aggregated from all cells in each well:



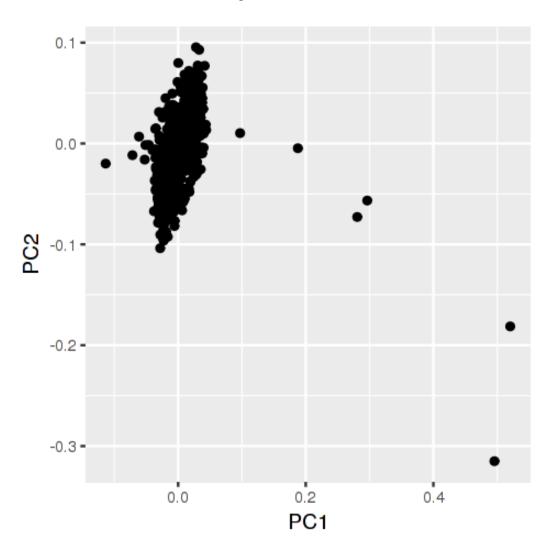
Only aggregated over isolated cells in each well:

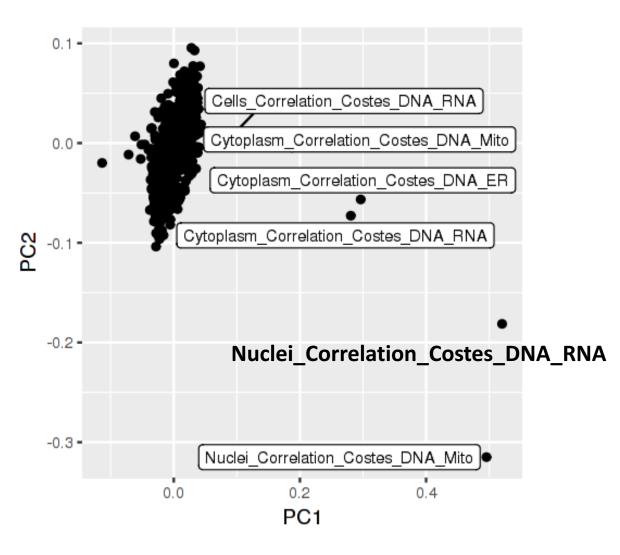


Only aggregated over colonized cells in each well:



3. Are any traits outliers?





Conclusions

- Cell line does influence proportion of isolated vs. colonized cells
- More variance is explained by cell line at 6 hours than at 24 hours
- More variance is explained by cell line in isolated cells than in colonized cells
- Aggregating all cells/well, all densities have similar variance explained by cell line
- Chosen condition (6h/10k) is comparable between isolated and colonized cells
- Outlier traits are measurements of correlated signal between different channels