Download all Data and accessory files---done

(Write script for accessory file download in future)---done

CHIP-seg project

Read Vervoort paper focusing on RNA-pol 2 analysis--in progress

Try to recreate that analysis until receive further instruction---in progress

Restricted Analysis to ALL cell lines w distinct genotypes?

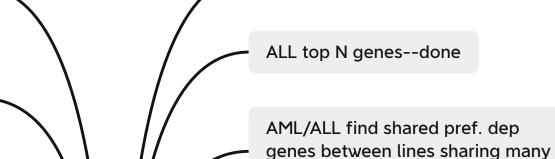
5. what tissues contain more selective expression of genes? What gene types are these?

4. - subset on whether these are ubiquitously expressed or also selectively expressed.

3. - How often do transcription factors and/or chromatin proteins display selectivity in dependency.

2. What functional classes do all selective dependencies belong to

Make graph visualization---in progress



DepMap

Restricted Analysis to---Molm14 Molm13 Mv411 Thp1 Nomol

AML top N genes--done

genes---done

Monomac1----done (are these the "distinct genotype cell lines?)

1. Expand to all other tissue types---in progress

Upset Plots---done