-~~Run adjusted Manhattan plots for all cell types~~

~~-Add table of Top CpG sites for every lipid mediator~~

~~-~~Re-run Tnai and Tmem with correct CpG filter

-Re-run Bcell with batch

-Table of CpG comparison between cell types and lipid mediators

-Histograms of lipid mediators

-Add new Tmem and Tnai Manhattan plots and Q-Q plots and tables to report

-Make DMR a loop

-Finish Report assembly

-DMR results

-QC Report

Qqman package vignette:

<https://cran.r-project.org/web/packages/qqman/vignettes/qqman.html>