Haploview notes

<https://www.broadinstitute.org/haploview/>

Data: five formats, standard linkage format, completely or partially phased haplotypes, [HapMap Project](http://www.hapmap.org/" \t "_blank) data dumps, PHASE format, and PLINK outputs

Data: \*.ped

Locus: \*.map

Appears to take only binary phenotypes?

And need to check SNP formatting