

GWAS Visualization in R

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We will need to get the tdt results for the data

```
library(qqman)

## 

## For example usage please run: vignette('qqman')

## 

## Citation appreciated but not required:

## Turner, (2018). qqman: an R package for visualizing GWAS results using Q-Q and manhattan plots. Journal of Statistical Software, 93(1), 1–37. doi:10.18637/jss.v093.i01

## 

library(dplyr)

## 

## Attaching package: 'dplyr'

## The following objects are masked from 'package:stats':
## 
##     filter, lag

## The following objects are masked from 'package:base':
## 
##     intersect, setdiff, setequal, union
```

```
library(ggplot2)
```

For a comparison, a sample dataset is provided in order to show what all chromosomes would look like. We will create some visualization with this as well.

```
head(gwasResults)
```

```
##   SNP CHR BP      P
## 1 rs1  1  1 0.9148060
## 2 rs2  1  2 0.9370754
## 3 rs3  1  3 0.2861395
## 4 rs4  1  4 0.8304476
## 5 rs5  1  5 0.6417455
## 6 rs6  1  6 0.5190959
```

Lets read in the TDT results - The adjusted file is a bit smaller here since the NA values cannot be adjusted and are ommitted. So lets merge and only keep values that exist in the adjusted file, but lets add the frequency or MAF

```
tdtfrq <- read.csv("data_analysis/tdtfrq.csv")
head(tdtfrq)
```

```
##   CHR     SNP A1 A2      MAF NCHROBS
## 1  21 5034209  A  G 0.003571    280
## 2  21 5034244  T  C 0.007042    284
## 3  21 5063575  T  C 0.003497    286
## 4  21 5063806  T  C 0.003448    290
## 5  21 5063842  A  G 0.010420    288
## 6  21 5089937  C  T 0.017240    290
```

```
tdtadj <- read.csv("data_analysis/tdtadj.csv", row.names=NULL)
head(tdtadj)
```

```
##   CHR     SNP    UNADJ      GC BONF HOLM SIDAK_SS SIDAK_SD FDR_BH FDR_BY
## 1  21 7946755 3.738e-05 0.005365  1    1  0.9714  0.9714 0.3734  1
## 2  21 7946763 3.738e-05 0.005365  1    1  0.9714  0.9714 0.3734  1
## 3  21 7946728 6.334e-05 0.006911  1    1  0.9976  0.9976 0.3734  1
## 4  21 7946741 6.334e-05 0.006911  1    1  0.9976  0.9976 0.3734  1
## 5  21 7946744 6.334e-05 0.006911  1    1  0.9976  0.9976 0.3734  1
## 6  21 26266437 8.770e-05 0.008081  1    1  0.9998  0.9998 0.3734  1
```

Lets merge our frequencies with the TDT results.

```
cleantdt <- merge(tdtfrq , tdtadj, by=c("CHR","SNP"))
head(cleantdt)
```

```
##   CHR     SNP A1 A2      MAF NCHROBS    UNADJ      GC BONF HOLM SIDAK_SS SIDAK_SD
## 1  21 10000132 AT  A 0.003448    290 0.3173 0.4995  1    1    1    1
## 2  21 10000176 AT  A 0.003448    290 0.3173 0.4995  1    1    1    1
## 3  21 10000297  A  G 0.006897    290 0.3173 0.4995  1    1    1    1
```

```

## 4 21 10000415 G T 0.003448    290 1.0000 1.0000    1   1   1   1
## 5 21 10000434 A G 0.003448    290 0.3173 0.4995    1   1   1   1
## 6 21 10000714 A G 0.489600    288 0.7194 0.8083    1   1   1   1
##   FDR_BH FDR_BY
## 1 0.3734      1
## 2 0.3734      1
## 3 0.3734      1
## 4 1.0000      1
## 5 0.3734      1
## 6 0.7983      1

```

Lets read in the annotation information

```

anno <- read.csv('data_analysis/query.output.genome_summary.csv')
head(anno)

```

```

##   Chr Start     End Ref Alt Func.refGene Gene.refGene
## 1 21 5034209 5034209 G A intronic ICOSLG
## 2 21 5034244 5034244 C T intronic ICOSLG
## 3 21 5063575 5063575 C T intergenic ICOSLG;LINC01678
## 4 21 5063806 5063806 C T intergenic ICOSLG;LINC01678
## 5 21 5063842 5063842 G A intergenic ICOSLG;LINC01678
## 6 21 5089937 5089937 T C intergenic ICOSLG;LINC01678
##   GeneDetail.refGene ExonicFunc.refGene AAChange.refGene X1000G_ALL
## 1
## 2
## 3 dist=22907;dist=38485
## 4 dist=23138;dist=38254
## 5 dist=23174;dist=38218
## 6 dist=49269;dist=12123
##   X1000G_AFR X1000G_AMR X1000G_EAS X1000G_EUR X1000G_SAS ExAC_Freq ExAC_AFR
## 1
## 2
## 3
## 4
## 5
## 6
##   ExAC_AMR ExAC_EAS ExAC_FIN ExAC_NFE ExAC_OTH ExAC_SAS ESP6500si_ALL
## 1
## 2
## 3
## 4
## 5
## 6
##   ESP6500si_AA ESP6500si_EA CG46 NCI60 dbSNP COSMIC_ID COSMIC_DIS ClinVar_SIG
## 1
## 2
## 3
## 4
## 5
## 6
##   ClinVar_DIS ClinVar_ID ClinVar_DB ClinVar_DBID GWAS_DIS GWAS_OR GWAS_BETA
## 1

```

```

## 2
## 3
## 4
## 5
## 6
##   GWAS_PUBMED GWAS_SNP GWAS_P SIFT_score SIFT_converted_rankscore SIFT_pred
## 1
## 2
## 3
## 4
## 5
## 6
##   Polyphen2_HDIV_score Polyphen2_HDIV_rankscore Polyphen2_HDIV_pred
## 1
## 2
## 3
## 4
## 5
## 6
##   Polyphen2_HVAR_score Polyphen2_HVAR_rankscore Polyphen2_HVAR_pred LRT_score
## 1
## 2
## 3
## 4
## 5
## 6
##   LRT_converted_rankscore LRT_pred MutationTaster_score
## 1
## 2
## 3
## 4
## 5
## 6
##   MutationTaster_converted_rankscore MutationTaster_pred MutationAssessor_score
## 1
## 2
## 3
## 4
## 5
## 6
##   MutationAssessor_score_rankscore MutationAssessor_pred FATHMM_score
## 1
## 2
## 3
## 4
## 5
## 6
##   FATHMM_converted_rankscore FATHMM_pred PROVEAN_score
## 1
## 2
## 3
## 4
## 5
## 6

```

```

## PROVEAN_converted_rankscore PROVEAN_pred VEST3_score VEST3_rankscore
## 1
## 2
## 3
## 4
## 5
## 6
## MetaSVM_score MetaSVM_rankscore MetaSVM_pred MetaLR_score MetaLR_rankscore
## 1
## 2
## 3
## 4
## 5
## 6
## MetaLR_pred M.CAP_score M.CAP_rankscore M.CAP_pred CADD_raw
## 1
## 2
## 3
## 4
## 5
## 6
## CADD_raw_rankscore CADD_phred DANN_score DANN_rankscore
## 1
## 2
## 3
## 4
## 5
## 6
## fathmm.MKL_coding_score fathmm.MKL_coding_rankscore fathmm.MKL_coding_pred
## 1
## 2
## 3
## 4
## 5
## 6
## Eigen_coding_or_noncoding Eigen.raw Eigen.PC.raw GenoCanyon_score
## 1
## 2
## 3
## 4
## 5
## 6
## GenoCanyon_score_rankscore integrated_fitCons_score
## 1
## 2
## 3
## 4
## 5
## 6
## integrated_fitCons_score_rankscore integrated_confidence_value GERP.._RS
## 1
## 2
## 3
## 4

```

```

## 5
## 6
##   GERP..._RS_rankscore phyloP100way_vertebrate phyloP100way_vertebrate_rankscore
## 1
## 2
## 3
## 4
## 5
## 6
##   phyloP20way_mammalian phyloP20way_mammalian_rankscore
## 1
## 2
## 3
## 4
## 5
## 6
##   phastCons100way_vertebrate phastCons100way_vertebrate_rankscore
## 1
## 2
## 3
## 4
## 5
## 6
##   phastCons20way_mammalian phastCons20way_mammalian_rankscore
## 1
## 2
## 3
## 4
## 5
## 6
##   SiPhy_29way_logOdds SiPhy_29way_logOdds_rankscore Interpro_domain
## 1
## 2
## 3
## 4
## 5
## 6
##   GTEx_V6_gene GTEx_V6_tissue gnomAD_exome_ALL gnomAD_exome_AFR
## 1
## 2
## 3
## 4
## 5
## 6
##   gnomAD_exome_AMR gnomAD_exome_ASJ gnomAD_exome_EAS gnomAD_exome_FIN
## 1
## 2
## 3
## 4
## 5
## 6
##   gnomAD_exome_NFE gnomAD_exome_OTH gnomAD_exome_SAS gnomAD_genome_ALL
## 1
## 2

```

```

## 3
## 4
## 5
## 6
##   gnomAD_genome_AFR gnomAD_genome_AMR gnomAD_genome_ASJ gnomAD_genome_EAS
## 1
## 2
## 3
## 4
## 5
## 6
##   gnomAD_genome_FIN gnomAD_genome_NFE gnomAD_genome_OTH Otherinfo Otherinfo.1
## 1                               hom   .
## 2                               hom   .
## 3                               hom   .
## 4                               hom   .
## 5                               hom   .
## 6                               hom   .

```

There seem to be some NA values produced by plink for TDT.. why might this be?

```

final <- merge(cleantdt, anno, by.x=c("CHR", "SNP"), by.y=c("Chr", "Start"))
head(final, 200)

```

##	CHR	SNP	A1	A2	MAF	NCHROBS	UNADJ	GC	BONF	HOLM	SIDAK_SS
## 1	21	10000132	AT	A	0.003448	290	0.317300	0.49950	1	1	1
## 2	21	10000176	AT	A	0.003448	290	0.317300	0.49950	1	1	1
## 3	21	10000297	A	G	0.006897	290	0.317300	0.49950	1	1	1
## 4	21	10000415	G	T	0.003448	290	1.000000	1.000000	1	1	1
## 5	21	10000434	A	G	0.003448	290	0.317300	0.49950	1	1	1
## 6	21	10000714	A	G	0.489600	288	0.719400	0.80830	1	1	1
## 7	21	10000838	G	A	0.003448	290	0.317300	0.49950	1	1	1
## 8	21	10001032	A	G	0.003448	290	0.317300	0.49950	1	1	1
## 9	21	10001205	T	C	0.006897	290	0.157300	0.33960	1	1	1
## 10	21	10001524	A	G	0.003448	290	0.317300	0.49950	1	1	1
## 11	21	10001527	A	T	0.003448	290	0.317300	0.49950	1	1	1
## 12	21	10001532	A	C	0.003448	290	0.317300	0.49950	1	1	1
## 13	21	10001568	G	C	0.003448	290	0.317300	0.49950	1	1	1
## 14	21	10001699	A	G	0.003472	288	0.317300	0.49950	1	1	1
## 15	21	10001937	T	C	0.006897	290	0.157300	0.33960	1	1	1
## 16	21	10002052	T	C	0.003448	290	0.317300	0.49950	1	1	1
## 17	21	10002435	AC	A	0.496600	290	0.788400	0.85620	1	1	1
## 18	21	10002450	G	A	0.496600	290	0.788400	0.85620	1	1	1
## 19	21	10002648	G	C	0.482800	290	1.000000	1.000000	1	1	1
## 20	21	10003122	C	A	0.003448	290	0.317300	0.49950	1	1	1
## 21	21	10003125	A	C	0.003448	290	0.317300	0.49950	1	1	1
## 22	21	10003217	A	G	0.003448	290	0.317300	0.49950	1	1	1
## 23	21	10003391	T	C	0.003448	290	0.317300	0.49950	1	1	1
## 24	21	10003475	A	G	0.003448	290	0.317300	0.49950	1	1	1
## 25	21	10003595	A	C	0.006897	290	1.000000	1.000000	1	1	1
## 26	21	10003938	A	G	0.003448	290	0.317300	0.49950	1	1	1
## 27	21	10004252	C	G	0.006897	290	1.000000	1.000000	1	1	1
## 28	21	10004491	T	G	0.003448	290	0.317300	0.49950	1	1	1

## 29	21	10005091	T	C	0.003472	288	1.000000	1.000000	1	1	1
## 30	21	10005117	A	G	0.003448	290	0.317300	0.49950	1	1	1
## 31	21	10005683	G	A	0.465500	290	0.583900	0.71150	1	1	1
## 32	21	10005833	T	A	0.006897	290	0.157300	0.33960	1	1	1
## 33	21	10006639	C	T	0.493100	290	1.000000	1.000000	1	1	1
## 34	21	10006998	C	T	0.496600	290	0.928700	0.95180	1	1	1
## 35	21	10007014	C	T	0.472400	290	0.785100	0.85390	1	1	1
## 36	21	10007016	G	A	0.006897	290	1.000000	1.000000	1	1	1
## 37	21	10007253	C	T	0.006897	290	0.157300	0.33960	1	1	1
## 38	21	10007310	T	C	0.003448	290	0.317300	0.49950	1	1	1
## 39	21	10007334	A	G	0.486200	290	0.715000	0.80520	1	1	1
## 40	21	10007528	A	C	0.003448	290	0.317300	0.49950	1	1	1
## 41	21	10007647	C	T	0.003448	290	0.317300	0.49950	1	1	1
## 42	21	10007780	G	A	0.003448	290	0.317300	0.49950	1	1	1
## 43	21	10007870	G	A	0.003448	290	0.317300	0.49950	1	1	1
## 44	21	10008190	T	A	0.003448	290	0.317300	0.49950	1	1	1
## 45	21	10008297	A	G	0.006897	290	0.157300	0.33960	1	1	1
## 46	21	10008432	A	G	0.472200	288	0.649400	0.75890	1	1	1
## 47	21	10008434	GT	G	0.010420	288	0.157300	0.33960	1	1	1
## 48	21	10008438	G	T	0.003448	290	0.317300	0.49950	1	1	1
## 49	21	10008463	T	A	0.003448	290	0.317300	0.49950	1	1	1
## 50	21	10008760	G	A	0.003448	290	0.317300	0.49950	1	1	1
## 51	21	10008761	T	C	0.003448	290	0.317300	0.49950	1	1	1
## 52	21	10008985	G	A	0.006897	290	0.157300	0.33960	1	1	1
## 53	21	10009273	T	G	0.003448	290	0.317300	0.49950	1	1	1
## 54	21	10009505	C	T	0.006897	290	0.157300	0.33960	1	1	1
## 55	21	10009642	A	G	0.003448	290	0.317300	0.49950	1	1	1
## 56	21	10009665	C	T	0.003448	290	0.317300	0.49950	1	1	1
## 57	21	10009907	A	T	0.006993	286	1.000000	1.000000	1	1	1
## 58	21	10009908	A	T	0.003472	288	0.317300	0.49950	1	1	1
## 59	21	10009909	A	T	0.003472	288	0.317300	0.49950	1	1	1
## 60	21	10009914	A	T	0.003472	288	0.317300	0.49950	1	1	1
## 61	21	10010164	A	G	0.027780	288	0.004678	0.05614	1	1	1
## 62	21	10010214	A	G	0.003472	288	0.317300	0.49950	1	1	1
## 63	21	10010235	T	C	0.013790	290	0.045500	0.17680	1	1	1
## 64	21	10010539	GA	G	0.006897	290	1.000000	1.000000	1	1	1
## 65	21	10010637	C	T	0.003448	290	0.317300	0.49950	1	1	1
## 66	21	10010921	G	A	0.003448	290	0.317300	0.49950	1	1	1
## 67	21	10011051	C	G	0.003448	290	0.317300	0.49950	1	1	1
## 68	21	10011506	A	G	0.006897	290	0.157300	0.33960	1	1	1
## 69	21	10011578	T	A	0.010420	288	0.083260	0.24220	1	1	1
## 70	21	10011713	C	G	0.003448	290	0.317300	0.49950	1	1	1
## 71	21	10011772	A	G	0.003448	290	0.317300	0.49950	1	1	1
## 72	21	10011773	A	G	0.013790	290	0.045500	0.17680	1	1	1
## 73	21	10011931	C	T	0.482800	290	0.785100	0.85390	1	1	1
## 74	21	10012070	T	A	0.003448	290	0.317300	0.49950	1	1	1
## 75	21	10012101	C	T	0.003448	290	0.317300	0.49950	1	1	1
## 76	21	10012241	T	A	0.003448	290	0.317300	0.49950	1	1	1
## 77	21	10012389	A	G	0.469000	290	0.715000	0.80520	1	1	1
## 78	21	10012478	A	G	0.010340	290	1.000000	1.000000	1	1	1
## 79	21	10012532	T	C	0.003448	290	0.317300	0.49950	1	1	1
## 80	21	10012588	A	C	0.006944	288	0.317300	0.49950	1	1	1
## 81	21	10012748	T	C	0.003448	290	0.317300	0.49950	1	1	1
## 82	21	10012833	C	T	0.496600	290	1.000000	1.000000	1	1	1

## 83	21	10013140	T	C	0.006897	290	0.317300	0.49950	1	1	1
## 84	21	10013167	T	G	0.013790	290	0.045500	0.17680	1	1	1
## 85	21	10013322	G	A	0.003448	290	0.317300	0.49950	1	1	1
## 86	21	10013383	T	C	0.472400	290	0.649400	0.75890	1	1	1
## 87	21	10013904	G	C	0.003448	290	0.317300	0.49950	1	1	1
## 88	21	10014007	G	A	0.489700	290	0.857500	0.90350	1	1	1
## 89	21	10014185	C	T	0.003448	290	0.317300	0.49950	1	1	1
## 90	21	10014417	C	T	0.003448	290	0.317300	0.49950	1	1	1
## 91	21	10014535	C	T	0.003448	290	0.317300	0.49950	1	1	1
## 92	21	10014675	A	G	0.003448	290	0.317300	0.49950	1	1	1
## 93	21	10014841	T	C	0.006897	290	0.157300	0.33960	1	1	1
## 94	21	10015062	A	T	0.003448	290	0.317300	0.49950	1	1	1
## 95	21	10015095	C	T	0.003448	290	0.317300	0.49950	1	1	1
## 96	21	10015343	T	C	0.003448	290	0.317300	0.49950	1	1	1
## 97	21	10015637	C	A	0.003472	288	0.317300	0.49950	1	1	1
## 98	21	10015651	A	G	0.006944	288	1.000000	1.00000	1	1	1
## 99	21	10015659	C	A	0.003472	288	0.317300	0.49950	1	1	1
## 100	21	10015886	T	C	0.006897	290	0.157300	0.33960	1	1	1
## 101	21	10016389	T	C	0.006897	290	0.157300	0.33960	1	1	1
## 102	21	10016844	A	G	0.003448	290	0.317300	0.49950	1	1	1
## 103	21	10017177	A	G	0.020690	290	0.014310	0.09811	1	1	1
## 104	21	10017226	T	A	0.003448	290	0.317300	0.49950	1	1	1
## 105	21	10017270	A	G	0.003448	290	0.317300	0.49950	1	1	1
## 106	21	10017357	A	C	0.013790	290	0.563700	0.69660	1	1	1
## 107	21	10017391	A	T	0.003448	290	0.317300	0.49950	1	1	1
## 108	21	10017422	A	G	0.020690	290	0.014310	0.09811	1	1	1
## 109	21	10017503	T	C	0.003448	290	1.000000	1.00000	1	1	1
## 110	21	10017541	A	G	0.003448	290	0.317300	0.49950	1	1	1
## 111	21	10017965	A	G	0.003448	290	0.317300	0.49950	1	1	1
## 112	21	10018006	T	C	0.003448	290	0.317300	0.49950	1	1	1
## 113	21	10018435	G	A	0.006897	290	0.157300	0.33960	1	1	1
## 114	21	10018494	T	C	0.020690	290	0.045500	0.17680	1	1	1
## 115	21	10019021	A	T	0.003448	290	0.317300	0.49950	1	1	1
## 116	21	10019152	T	G	0.006944	288	0.157300	0.33960	1	1	1
## 117	21	10019506	A	G	0.003448	290	0.317300	0.49950	1	1	1
## 118	21	10019575	A	C	0.013790	290	0.157300	0.33960	1	1	1
## 119	21	10019675	A	G	0.003448	290	0.317300	0.49950	1	1	1
## 120	21	10019946	A	C	0.003448	290	0.317300	0.49950	1	1	1
## 121	21	10020042	A	G	0.003448	290	0.317300	0.49950	1	1	1
## 122	21	10020461	A	G	0.017240	290	0.317300	0.49950	1	1	1
## 123	21	10020539	C	T	0.003448	290	0.317300	0.49950	1	1	1
## 124	21	10020686	C	G	0.006897	290	0.157300	0.33960	1	1	1
## 125	21	10020755	T	C	0.003448	290	0.317300	0.49950	1	1	1
## 126	21	10020832	A	G	0.006897	290	0.157300	0.33960	1	1	1
## 127	21	10020891	A	C	0.003448	290	1.000000	1.00000	1	1	1
## 128	21	10021001	C	A	0.010340	290	1.000000	1.00000	1	1	1
## 129	21	10021115	C	G	0.006897	290	0.157300	0.33960	1	1	1
## 130	21	10021137	C	T	0.003448	290	0.317300	0.49950	1	1	1
## 131	21	10021258	T	A	0.006897	290	0.157300	0.33960	1	1	1
## 132	21	10021294	T	C	0.003448	290	0.317300	0.49950	1	1	1
## 133	21	10021435	C	T	0.451700	290	0.510200	0.65660	1	1	1
## 134	21	10021547	T	C	0.003448	290	0.317300	0.49950	1	1	1
## 135	21	10021582	C	T	0.003448	290	0.317300	0.49950	1	1	1
## 136	21	10021635	T	C	0.006897	290	0.157300	0.33960	1	1	1

## 137	21	10021950	T	C	0.006897	290	0.563700	0.69660	1	1	1
## 138	21	10022055	G	C	0.003448	290	0.317300	0.49950	1	1	1
## 139	21	10022419	G	A	0.003448	290	0.317300	0.49950	1	1	1
## 140	21	10022538	G	A	0.455200	290	0.574100	0.70430	1	1	1
## 141	21	10022567	G	C	0.479300	290	0.717200	0.80680	1	1	1
## 142	21	10022728	A	C	0.006897	290	0.157300	0.33960	1	1	1
## 143	21	10023143	A	G	0.006897	290	0.157300	0.33960	1	1	1
## 144	21	10023288	A	G	0.003448	290	0.317300	0.49950	1	1	1
## 145	21	10023337	T	A	0.020690	290	0.414200	0.58140	1	1	1
## 146	21	10023506	C	A	0.003448	290	0.317300	0.49950	1	1	1
## 147	21	10024005	A	G	0.003448	290	0.317300	0.49950	1	1	1
## 148	21	10024104	G	A	0.010340	290	0.563700	0.69660	1	1	1
## 149	21	10024162	C	G	0.013790	290	0.157300	0.33960	1	1	1
## 150	21	10024202	G	A	0.003448	290	0.317300	0.49950	1	1	1
## 151	21	10024370	T	C	0.003448	290	0.317300	0.49950	1	1	1
## 152	21	10024885	T	C	0.013790	290	0.563700	0.69660	1	1	1
## 153	21	10025064	A	T	0.486200	290	0.649400	0.75890	1	1	1
## 154	21	10025324	C	T	0.003448	290	0.317300	0.49950	1	1	1
## 155	21	10025596	G	A	0.003448	290	0.157300	0.33960	1	1	1
## 156	21	10025811	G	A	0.003448	290	0.317300	0.49950	1	1	1
## 157	21	10025980	T	C	0.003448	290	0.317300	0.49950	1	1	1
## 158	21	10025998	T	A	0.003448	290	0.317300	0.49950	1	1	1
## 159	21	10026037	T	C	0.006897	290	1.000000	1.00000	1	1	1
## 160	21	10026117	G	C	0.003448	290	0.317300	0.49950	1	1	1
## 161	21	10026251	A	G	0.006897	290	0.157300	0.33960	1	1	1
## 162	21	10026295	G	A	0.489700	290	0.857500	0.90350	1	1	1
## 163	21	10026366	T	C	0.003448	290	0.317300	0.49950	1	1	1
## 164	21	10026389	G	A	0.006897	290	0.157300	0.33960	1	1	1
## 165	21	10026462	C	T	0.003448	290	0.317300	0.49950	1	1	1
## 166	21	10026597	C	A	0.003448	290	0.317300	0.49950	1	1	1
## 167	21	10026608	A	T	0.006897	290	0.157300	0.33960	1	1	1
## 168	21	10026944	A	G	0.003448	290	0.317300	0.49950	1	1	1
## 169	21	10026971	G	C	0.003448	290	0.317300	0.49950	1	1	1
## 170	21	10027368	A	G	0.003448	290	0.317300	0.49950	1	1	1
## 171	21	10027586	A	G	0.003448	290	0.317300	0.49950	1	1	1
## 172	21	10027747	C	A	0.013790	290	1.000000	1.00000	1	1	1
## 173	21	10027798	T	C	0.003448	290	0.317300	0.49950	1	1	1
## 174	21	10027801	G	A	0.003448	290	0.317300	0.49950	1	1	1
## 175	21	10027872	T	C	0.479200	288	0.717200	0.80680	1	1	1
## 176	21	10027941	T	C	0.017240	290	0.179700	0.36490	1	1	1
## 177	21	10028040	G	A	0.013790	290	0.045500	0.17680	1	1	1
## 178	21	10028093	A	G	0.003448	290	0.317300	0.49950	1	1	1
## 179	21	10028262	G	A	0.003448	290	0.317300	0.49950	1	1	1
## 180	21	10028358	GA	G	0.003448	290	1.000000	1.00000	1	1	1
## 181	21	10028388	G	A	0.003448	290	0.317300	0.49950	1	1	1
## 182	21	10028489	A	G	0.003448	290	0.317300	0.49950	1	1	1
## 183	21	10028702	T	C	0.003448	290	0.317300	0.49950	1	1	1
## 184	21	10028714	G	A	0.003448	290	0.317300	0.49950	1	1	1
## 185	21	10028853	T	C	0.434500	290	0.638100	0.75080	1	1	1
## 186	21	10028988	G	A	0.003448	290	0.317300	0.49950	1	1	1
## 187	21	10029181	G	A	0.003448	290	0.317300	0.49950	1	1	1
## 188	21	10029293	C	T	0.003448	290	0.317300	0.49950	1	1	1
## 189	21	10029512	T	G	0.003448	290	0.317300	0.49950	1	1	1
## 190	21	10029516	T	C	0.020690	290	0.014310	0.09811	1	1	1

	##	191	21	10029617	A	G	0.003448	290	0.317300	0.49950	1	1	1
	##	192	21	10029692	C	G	0.003448	290	0.317300	0.49950	1	1	1
	##	193	21	10030257	A	G	0.003448	290	0.317300	0.49950	1	1	1
	##	194	21	10030615	C	G	0.003497	286	0.317300	0.49950	1	1	1
	##	195	21	10030733	GA	G	0.003472	288	0.317300	0.49950	1	1	1
	##	196	21	10030882	A	G	0.010710	280	1.000000	1.00000	1	1	1
	##	197	21	10032811	G	A	0.003448	290	0.317300	0.49950	1	1	1
	##	198	21	10033055	T	C	0.003448	290	0.317300	0.49950	1	1	1
	##	199	21	10033264	T	A	0.006897	290	1.000000	1.00000	1	1	1
	##	200	21	10033556	A	T	0.003448	290	0.317300	0.49950	1	1	1
	##			SIDAK_SD	FDR_BH	FDR_BY		End	Ref	Alt	Func.refGene	Gene.refGene	
	##	1	1	0.3734			1	10000132	-	T	intergenic	LINC01667;BAGE	
	##	2	1	0.3734			1	10000176	-	T	intergenic	LINC01667;BAGE	
	##	3	1	0.3734			1	10000297	G	A	intergenic	LINC01667;BAGE	
	##	4	1	1.0000			1	10000415	T	G	intergenic	LINC01667;BAGE	
	##	5	1	0.3734			1	10000434	G	A	intergenic	LINC01667;BAGE	
	##	6	1	0.7983			1	10000714	G	A	intergenic	LINC01667;BAGE	
	##	7	1	0.3734			1	10000838	A	G	intergenic	LINC01667;BAGE	
	##	8	1	0.3734			1	10001032	G	A	intergenic	LINC01667;BAGE	
	##	9	1	0.3734			1	10001205	C	T	intergenic	LINC01667;BAGE	
	##	10	1	0.3734			1	10001524	G	A	intergenic	LINC01667;BAGE	
	##	11	1	0.3734			1	10001527	T	A	intergenic	LINC01667;BAGE	
	##	12	1	0.3734			1	10001532	C	A	intergenic	LINC01667;BAGE	
	##	13	1	0.3734			1	10001568	C	G	intergenic	LINC01667;BAGE	
	##	14	1	0.3734			1	10001699	G	A	intergenic	LINC01667;BAGE	
	##	15	1	0.3734			1	10001937	C	T	intergenic	LINC01667;BAGE	
	##	16	1	0.3734			1	10002052	C	T	intergenic	LINC01667;BAGE	
	##	17	1	0.8696			1	10002435	-	C	intergenic	LINC01667;BAGE	
	##	18	1	0.8696			1	10002450	A	G	intergenic	LINC01667;BAGE	
	##	19	1	1.0000			1	10002648	C	G	intergenic	LINC01667;BAGE	
	##	20	1	0.3734			1	10003122	A	C	intergenic	LINC01667;BAGE	
	##	21	1	0.3734			1	10003125	C	A	intergenic	LINC01667;BAGE	
	##	22	1	0.3734			1	10003217	G	A	intergenic	LINC01667;BAGE	
	##	23	1	0.3734			1	10003391	C	T	intergenic	LINC01667;BAGE	
	##	24	1	0.3734			1	10003475	G	A	intergenic	LINC01667;BAGE	
	##	25	1	1.0000			1	10003595	C	A	intergenic	LINC01667;BAGE	
	##	26	1	0.3734			1	10003938	G	A	intergenic	LINC01667;BAGE	
	##	27	1	1.0000			1	10004252	G	C	intergenic	LINC01667;BAGE	
	##	28	1	0.3734			1	10004491	G	T	intergenic	LINC01667;BAGE	
	##	29	1	1.0000			1	10005091	C	T	intergenic	LINC01667;BAGE	
	##	30	1	0.3734			1	10005117	G	A	intergenic	LINC01667;BAGE	
	##	31	1	0.6570			1	10005683	A	G	intergenic	LINC01667;BAGE	
	##	32	1	0.3734			1	10005833	A	T	intergenic	LINC01667;BAGE	
	##	33	1	1.0000			1	10006639	T	C	intergenic	LINC01667;BAGE	
	##	34	1	1.0000			1	10006998	T	C	intergenic	LINC01667;BAGE	
	##	35	1	0.8680			1	10007014	T	C	intergenic	LINC01667;BAGE	
	##	36	1	1.0000			1	10007016	A	G	intergenic	LINC01667;BAGE	
	##	37	1	0.3734			1	10007253	T	C	intergenic	LINC01667;BAGE	
	##	38	1	0.3734			1	10007310	C	T	intergenic	LINC01667;BAGE	
	##	39	1	0.7946			1	10007334	G	A	intergenic	LINC01667;BAGE	
	##	40	1	0.3734			1	10007528	C	A	intergenic	LINC01667;BAGE	
	##	41	1	0.3734			1	10007647	T	C	intergenic	LINC01667;BAGE	
	##	42	1	0.3734			1	10007780	A	G	intergenic	LINC01667;BAGE	
	##	43	1	0.3734			1	10007870	A	G	intergenic	LINC01667;BAGE	

## 44	1 0.3734	1 10008190	A T	intergenic LINC01667;BAGE
## 45	1 0.3734	1 10008297	G A	intergenic LINC01667;BAGE
## 46	1 0.7285	1 10008432	G A	intergenic LINC01667;BAGE
## 47	1 0.3734	1 10008434	- T	intergenic LINC01667;BAGE
## 48	1 0.3734	1 10008438	T G	intergenic LINC01667;BAGE
## 49	1 0.3734	1 10008463	A T	intergenic LINC01667;BAGE
## 50	1 0.3734	1 10008760	A G	intergenic LINC01667;BAGE
## 51	1 0.3734	1 10008761	C T	intergenic LINC01667;BAGE
## 52	1 0.3734	1 10008985	A G	intergenic LINC01667;BAGE
## 53	1 0.3734	1 10009273	G T	intergenic LINC01667;BAGE
## 54	1 0.3734	1 10009505	T C	intergenic LINC01667;BAGE
## 55	1 0.3734	1 10009642	G A	intergenic LINC01667;BAGE
## 56	1 0.3734	1 10009665	T C	intergenic LINC01667;BAGE
## 57	1 1.0000	1 10009907	T A	intergenic LINC01667;BAGE
## 58	1 0.3734	1 10009908	T A	intergenic LINC01667;BAGE
## 59	1 0.3734	1 10009909	T A	intergenic LINC01667;BAGE
## 60	1 0.3734	1 10009914	T A	intergenic LINC01667;BAGE
## 61	1 0.3734	1 10010164	G A	intergenic LINC01667;BAGE
## 62	1 0.3734	1 10010214	G A	intergenic LINC01667;BAGE
## 63	1 0.3734	1 10010235	C T	intergenic LINC01667;BAGE
## 64	1 1.0000	1 10010539	- A	intergenic LINC01667;BAGE
## 65	1 0.3734	1 10010637	T C	intergenic LINC01667;BAGE
## 66	1 0.3734	1 10010921	A G	intergenic LINC01667;BAGE
## 67	1 0.3734	1 10011051	G C	intergenic LINC01667;BAGE
## 68	1 0.3734	1 10011506	G A	intergenic LINC01667;BAGE
## 69	1 0.3734	1 10011578	A T	intergenic LINC01667;BAGE
## 70	1 0.3734	1 10011713	G C	intergenic LINC01667;BAGE
## 71	1 0.3734	1 10011772	G A	intergenic LINC01667;BAGE
## 72	1 0.3734	1 10011773	G A	intergenic LINC01667;BAGE
## 73	1 0.8680	1 10011931	T C	intergenic LINC01667;BAGE
## 74	1 0.3734	1 10012070	A T	intergenic LINC01667;BAGE
## 75	1 0.3734	1 10012101	T C	intergenic LINC01667;BAGE
## 76	1 0.3734	1 10012241	A T	intergenic LINC01667;BAGE
## 77	1 0.7946	1 10012389	G A	intergenic LINC01667;BAGE
## 78	1 1.0000	1 10012478	G A	intergenic LINC01667;BAGE
## 79	1 0.3734	1 10012532	C T	intergenic LINC01667;BAGE
## 80	1 0.3734	1 10012588	C A	intergenic LINC01667;BAGE
## 81	1 0.3734	1 10012748	C T	intergenic LINC01667;BAGE
## 82	1 1.0000	1 10012833	T C	intergenic LINC01667;BAGE
## 83	1 0.3734	1 10013140	C T	intergenic LINC01667;BAGE
## 84	1 0.3734	1 10013167	G T	intergenic LINC01667;BAGE
## 85	1 0.3734	1 10013322	A G	intergenic LINC01667;BAGE
## 86	1 0.7285	1 10013383	C T	intergenic LINC01667;BAGE
## 87	1 0.3734	1 10013904	C G	intergenic LINC01667;BAGE
## 88	1 0.9395	1 10014007	A G	intergenic LINC01667;BAGE
## 89	1 0.3734	1 10014185	T C	intergenic LINC01667;BAGE
## 90	1 0.3734	1 10014417	T C	intergenic LINC01667;BAGE
## 91	1 0.3734	1 10014535	T C	intergenic LINC01667;BAGE
## 92	1 0.3734	1 10014675	G A	intergenic LINC01667;BAGE
## 93	1 0.3734	1 10014841	C T	intergenic LINC01667;BAGE
## 94	1 0.3734	1 10015062	T A	intergenic LINC01667;BAGE
## 95	1 0.3734	1 10015095	T C	intergenic LINC01667;BAGE
## 96	1 0.3734	1 10015343	C T	intergenic LINC01667;BAGE
## 97	1 0.3734	1 10015637	A C	intergenic LINC01667;BAGE

## 98	1 1.0000	1 10015651	G A	intergenic LINC01667;BAGE
## 99	1 0.3734	1 10015659	A C	intergenic LINC01667;BAGE
## 100	1 0.3734	1 10015886	C T	intergenic LINC01667;BAGE
## 101	1 0.3734	1 10016389	C T	intergenic LINC01667;BAGE
## 102	1 0.3734	1 10016844	G A	intergenic LINC01667;BAGE
## 103	1 0.3734	1 10017177	G A	intergenic LINC01667;BAGE
## 104	1 0.3734	1 10017226	A T	intergenic LINC01667;BAGE
## 105	1 0.3734	1 10017270	G A	intergenic LINC01667;BAGE
## 106	1 0.6348	1 10017357	C A	intergenic LINC01667;BAGE
## 107	1 0.3734	1 10017391	T A	intergenic LINC01667;BAGE
## 108	1 0.3734	1 10017422	G A	intergenic LINC01667;BAGE
## 109	1 1.0000	1 10017503	C T	intergenic LINC01667;BAGE
## 110	1 0.3734	1 10017541	G A	intergenic LINC01667;BAGE
## 111	1 0.3734	1 10017965	G A	intergenic LINC01667;BAGE
## 112	1 0.3734	1 10018006	C T	intergenic LINC01667;BAGE
## 113	1 0.3734	1 10018435	A G	intergenic LINC01667;BAGE
## 114	1 0.3734	1 10018494	C T	intergenic LINC01667;BAGE
## 115	1 0.3734	1 10019021	T A	intergenic LINC01667;BAGE
## 116	1 0.3734	1 10019152	G T	intergenic LINC01667;BAGE
## 117	1 0.3734	1 10019506	G A	intergenic LINC01667;BAGE
## 118	1 0.3734	1 10019575	C A	intergenic LINC01667;BAGE
## 119	1 0.3734	1 10019675	G A	intergenic LINC01667;BAGE
## 120	1 0.3734	1 10019946	C A	intergenic LINC01667;BAGE
## 121	1 0.3734	1 10020042	G A	intergenic LINC01667;BAGE
## 122	1 0.3734	1 10020461	G A	intergenic LINC01667;BAGE
## 123	1 0.3734	1 10020539	T C	intergenic LINC01667;BAGE
## 124	1 0.3734	1 10020686	G C	intergenic LINC01667;BAGE
## 125	1 0.3734	1 10020755	C T	intergenic LINC01667;BAGE
## 126	1 0.3734	1 10020832	G A	intergenic LINC01667;BAGE
## 127	1 1.0000	1 10020891	C A	intergenic LINC01667;BAGE
## 128	1 1.0000	1 10021001	A C	intergenic LINC01667;BAGE
## 129	1 0.3734	1 10021115	G C	intergenic LINC01667;BAGE
## 130	1 0.3734	1 10021137	T C	intergenic LINC01667;BAGE
## 131	1 0.3734	1 10021258	A T	intergenic LINC01667;BAGE
## 132	1 0.3734	1 10021294	C T	intergenic LINC01667;BAGE
## 133	1 0.5954	1 10021435	T C	intergenic LINC01667;BAGE
## 134	1 0.3734	1 10021547	C T	intergenic LINC01667;BAGE
## 135	1 0.3734	1 10021582	T C	intergenic LINC01667;BAGE
## 136	1 0.3734	1 10021635	C T	intergenic LINC01667;BAGE
## 137	1 0.6348	1 10021950	C T	intergenic LINC01667;BAGE
## 138	1 0.3734	1 10022055	C G	intergenic LINC01667;BAGE
## 139	1 0.3734	1 10022419	A G	intergenic LINC01667;BAGE
## 140	1 0.6464	1 10022538	A G	intergenic LINC01667;BAGE
## 141	1 0.7966	1 10022567	C G	intergenic LINC01667;BAGE
## 142	1 0.3734	1 10022728	C A	intergenic LINC01667;BAGE
## 143	1 0.3734	1 10023143	G A	intergenic LINC01667;BAGE
## 144	1 0.3734	1 10023288	G A	intergenic LINC01667;BAGE
## 145	1 0.4848	1 10023337	A T	intergenic LINC01667;BAGE
## 146	1 0.3734	1 10023506	A C	intergenic LINC01667;BAGE
## 147	1 0.3734	1 10024005	G A	intergenic LINC01667;BAGE
## 148	1 0.6348	1 10024104	A G	intergenic LINC01667;BAGE
## 149	1 0.3734	1 10024162	G C	intergenic LINC01667;BAGE
## 150	1 0.3734	1 10024202	A G	intergenic LINC01667;BAGE
## 151	1 0.3734	1 10024370	C T	intergenic LINC01667;BAGE

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## 152      1 0.6348    1 10024885 C T intergenic LINC01667;BAGE
## 153      1 0.7285    1 10025064 T A intergenic LINC01667;BAGE
## 154      1 0.3734    1 10025324 T C intergenic LINC01667;BAGE
## 155      1 0.3734    1 10025596 A G intergenic LINC01667;BAGE
## 156      1 0.3734    1 10025811 A G intergenic LINC01667;BAGE
## 157      1 0.3734    1 10025980 C T intergenic LINC01667;BAGE
## 158      1 0.3734    1 10025998 A T intergenic LINC01667;BAGE
## 159      1 1.0000    1 10026037 C T intergenic LINC01667;BAGE
## 160      1 0.3734    1 10026117 C G intergenic LINC01667;BAGE
## 161      1 0.3734    1 10026251 G A intergenic LINC01667;BAGE
## 162      1 0.9395    1 10026295 A G intergenic LINC01667;BAGE
## 163      1 0.3734    1 10026366 C T intergenic LINC01667;BAGE
## 164      1 0.3734    1 10026389 A G intergenic LINC01667;BAGE
## 165      1 0.3734    1 10026462 T C intergenic LINC01667;BAGE
## 166      1 0.3734    1 10026597 A C intergenic LINC01667;BAGE
## 167      1 0.3734    1 10026608 T A intergenic LINC01667;BAGE
## 168      1 0.3734    1 10026944 G A intergenic LINC01667;BAGE
## 169      1 0.3734    1 10026971 C G intergenic LINC01667;BAGE
## 170      1 0.3734    1 10027368 G A intergenic LINC01667;BAGE
## 171      1 0.3734    1 10027586 G A intergenic LINC01667;BAGE
## 172      1 1.0000    1 10027747 A C intergenic LINC01667;BAGE
## 173      1 0.3734    1 10027798 C T intergenic LINC01667;BAGE
## 174      1 0.3734    1 10027801 A G intergenic LINC01667;BAGE
## 175      1 0.7966    1 10027872 C T intergenic LINC01667;BAGE
## 176      1 0.3734    1 10027941 C T intergenic LINC01667;BAGE
## 177      1 0.3734    1 10028040 A G intergenic LINC01667;BAGE
## 178      1 0.3734    1 10028093 G A intergenic LINC01667;BAGE
## 179      1 0.3734    1 10028262 A G intergenic LINC01667;BAGE
## 180      1 1.0000    1 10028358 - A intergenic LINC01667;BAGE
## 181      1 0.3734    1 10028388 A G intergenic LINC01667;BAGE
## 182      1 0.3734    1 10028489 G A intergenic LINC01667;BAGE
## 183      1 0.3734    1 10028702 C T intergenic LINC01667;BAGE
## 184      1 0.3734    1 10028714 A G intergenic LINC01667;BAGE
## 185      1 0.7168    1 10028853 C T intergenic LINC01667;BAGE
## 186      1 0.3734    1 10028988 A G intergenic LINC01667;BAGE
## 187      1 0.3734    1 10029181 A G intergenic LINC01667;BAGE
## 188      1 0.3734    1 10029293 T C intergenic LINC01667;BAGE
## 189      1 0.3734    1 10029512 G T intergenic LINC01667;BAGE
## 190      1 0.3734    1 10029516 C T intergenic LINC01667;BAGE
## 191      1 0.3734    1 10029617 G A intergenic LINC01667;BAGE
## 192      1 0.3734    1 10029692 G C intergenic LINC01667;BAGE
## 193      1 0.3734    1 10030257 G A intergenic LINC01667;BAGE
## 194      1 0.3734    1 10030615 G C intergenic LINC01667;BAGE
## 195      1 0.3734    1 10030733 - A intergenic LINC01667;BAGE
## 196      1 1.0000    1 10030882 G A intergenic LINC01667;BAGE
## 197      1 0.3734    1 10032811 A G intergenic LINC01667;BAGE
## 198      1 0.3734    1 10033055 C T intergenic LINC01667;BAGE
## 199      1 1.0000    1 10033264 A T intergenic LINC01667;BAGE
## 200      1 0.3734    1 10033556 T A intergenic LINC01667;BAGE
##                               GeneDetail.refGene ExonicFunc.refGene AAChange.refGene X1000G_ALL
## 1      dist=179071;dist=413388 . .
## 2      dist=179115;dist=413344 . .
## 3      dist=179236;dist=413223 0.0018
## 4      dist=179354;dist=413105

```

## 5	dist=179373;dist=413086	.
## 6	dist=179653;dist=412806	.
## 7	dist=179777;dist=412682	.
## 8	dist=179971;dist=412488	0.001
## 9	dist=180144;dist=412315	.
## 10	dist=180463;dist=411996	.
## 11	dist=180466;dist=411993	.
## 12	dist=180471;dist=411988	.
## 13	dist=180507;dist=411952	0.0022
## 14	dist=180638;dist=411821	.
## 15	dist=180876;dist=411583	.
## 16	dist=180991;dist=411468	.
## 17	dist=181374;dist=411085	0.52
## 18	dist=181389;dist=411070	.
## 19	dist=181587;dist=410872	0.47
## 20	dist=182061;dist=410398	.
## 21	dist=182064;dist=410395	.
## 22	dist=182156;dist=410303	.
## 23	dist=182330;dist=410129	.
## 24	dist=182414;dist=410045	.
## 25	dist=182534;dist=409925	0.001
## 26	dist=182877;dist=409582	.
## 27	dist=183191;dist=409268	.
## 28	dist=183430;dist=409029	.
## 29	dist=184030;dist=408429	.
## 30	dist=184056;dist=408403	.
## 31	dist=184622;dist=407837	.
## 32	dist=184772;dist=407687	.
## 33	dist=185578;dist=406881	.
## 34	dist=185937;dist=406522	.
## 35	dist=185953;dist=406506	.
## 36	dist=185955;dist=406504	.
## 37	dist=186192;dist=406267	.
## 38	dist=186249;dist=406210	.
## 39	dist=186273;dist=406186	.
## 40	dist=186467;dist=405992	.
## 41	dist=186586;dist=405873	.
## 42	dist=186719;dist=405740	.
## 43	dist=186809;dist=405650	.
## 44	dist=187129;dist=405330	.
## 45	dist=187236;dist=405223	.
## 46	dist=187371;dist=405088	.
## 47	dist=187373;dist=405086	.
## 48	dist=187377;dist=405082	.
## 49	dist=187402;dist=405057	.
## 50	dist=187699;dist=404760	0.0004
## 51	dist=187700;dist=404759	0.0002
## 52	dist=187924;dist=404535	0.0008
## 53	dist=188212;dist=404247	.
## 54	dist=188444;dist=404015	.
## 55	dist=188581;dist=403878	.
## 56	dist=188604;dist=403855	.
## 57	dist=188846;dist=403613	.
## 58	dist=188847;dist=403612	.

```

## 59 dist=188848;dist=403611 .  

## 60 dist=188853;dist=403606 .  

## 61 dist=189103;dist=403356 .  

## 62 dist=189153;dist=403306 0.0004  

## 63 dist=189174;dist=403285 .  

## 64 dist=189478;dist=402981 .  

## 65 dist=189576;dist=402883 .  

## 66 dist=189860;dist=402599 .  

## 67 dist=189990;dist=402469 .  

## 68 dist=190445;dist=402014 .  

## 69 dist=190517;dist=401942 .  

## 70 dist=190652;dist=401807 .  

## 71 dist=190711;dist=401748 .  

## 72 dist=190712;dist=401747 .  

## 73 dist=190870;dist=401589 .  

## 74 dist=191009;dist=401450 .  

## 75 dist=191040;dist=401419 .  

## 76 dist=191180;dist=401279 .  

## 77 dist=191328;dist=401131 .  

## 78 dist=191417;dist=401042 .  

## 79 dist=191471;dist=400988 .  

## 80 dist=191527;dist=400932 .  

## 81 dist=191687;dist=400772 0.0006  

## 82 dist=191772;dist=400687 .  

## 83 dist=192079;dist=400380 .  

## 84 dist=192106;dist=400353 .  

## 85 dist=192261;dist=400198 .  

## 86 dist=192322;dist=400137 .  

## 87 dist=192843;dist=399616 0.0002  

## 88 dist=192946;dist=399513 .  

## 89 dist=193124;dist=399335 .  

## 90 dist=193356;dist=399103 0.0018  

## 91 dist=193474;dist=398985 .  

## 92 dist=193614;dist=398845 .  

## 93 dist=193780;dist=398679 0.0008  

## 94 dist=194001;dist=398458 .  

## 95 dist=194034;dist=398425 .  

## 96 dist=194282;dist=398177 .  

## 97 dist=194576;dist=397883 .  

## 98 dist=194590;dist=397869 0.0006  

## 99 dist=194598;dist=397861 .  

## 100 dist=194825;dist=397634 0.0014  

## 101 dist=195328;dist=397131 .  

## 102 dist=195783;dist=396676 .  

## 103 dist=196116;dist=396343 .  

## 104 dist=196165;dist=396294 .  

## 105 dist=196209;dist=396250 .  

## 106 dist=196296;dist=396163 .  

## 107 dist=196330;dist=396129 .  

## 108 dist=196361;dist=396098 .  

## 109 dist=196442;dist=396017 0.0006  

## 110 dist=196480;dist=395979 .  

## 111 dist=196904;dist=395555 .  

## 112 dist=196945;dist=395514 .

```

## 113 dist=197374;dist=395085	0.0006
## 114 dist=197433;dist=395026	.
## 115 dist=197960;dist=394499	.
## 116 dist=198091;dist=394368	.
## 117 dist=198445;dist=394014	.
## 118 dist=198514;dist=393945	.
## 119 dist=198614;dist=393845	.
## 120 dist=198885;dist=393574	.
## 121 dist=198981;dist=393478	.
## 122 dist=199400;dist=393059	.
## 123 dist=199478;dist=392981	0.0002
## 124 dist=199625;dist=392834	.
## 125 dist=199694;dist=392765	0.0002
## 126 dist=199771;dist=392688	.
## 127 dist=199830;dist=392629	.
## 128 dist=199940;dist=392519	0.0008
## 129 dist=200054;dist=392405	.
## 130 dist=200076;dist=392383	.
## 131 dist=200197;dist=392262	.
## 132 dist=200233;dist=392226	.
## 133 dist=200374;dist=392085	.
## 134 dist=200486;dist=391973	0.013
## 135 dist=200521;dist=391938	.
## 136 dist=200574;dist=391885	0.0012
## 137 dist=200889;dist=391570	.
## 138 dist=200994;dist=391465	.
## 139 dist=201358;dist=391101	.
## 140 dist=201477;dist=390982	.
## 141 dist=201506;dist=390953	.
## 142 dist=201667;dist=390792	.
## 143 dist=202082;dist=390377	.
## 144 dist=202227;dist=390232	.
## 145 dist=202276;dist=390183	.
## 146 dist=202445;dist=390014	.
## 147 dist=202944;dist=389515	.
## 148 dist=203043;dist=389416	.
## 149 dist=203101;dist=389358	0.0004
## 150 dist=203141;dist=389318	0.0004
## 151 dist=203309;dist=389150	.
## 152 dist=203824;dist=388635	.
## 153 dist=204003;dist=388456	.
## 154 dist=204263;dist=388196	.
## 155 dist=204535;dist=387924	.
## 156 dist=204750;dist=387709	.
## 157 dist=204919;dist=387540	.
## 158 dist=204937;dist=387522	.
## 159 dist=204976;dist=387483	.
## 160 dist=205056;dist=387403	.
## 161 dist=205190;dist=387269	.
## 162 dist=205234;dist=387225	.
## 163 dist=205305;dist=387154	0.0006
## 164 dist=205328;dist=387131	0.0006
## 165 dist=205401;dist=387058	.
## 166 dist=205536;dist=386923	.

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## 167 dist=205547;dist=386912          .
## 168 dist=205883;dist=386576          0.0006
## 169 dist=205910;dist=386549          .
## 170 dist=206307;dist=386152          .
## 171 dist=206525;dist=385934          .
## 172 dist=206686;dist=385773          .
## 173 dist=206737;dist=385722          .
## 174 dist=206740;dist=385719          .
## 175 dist=206811;dist=385648          .
## 176 dist=206880;dist=385579          .
## 177 dist=206979;dist=385480          0.0002
## 178 dist=207032;dist=385427          .
## 179 dist=207201;dist=385258          .
## 180 dist=207297;dist=385162          .
## 181 dist=207327;dist=385132          .
## 182 dist=207428;dist=385031          .
## 183 dist=207641;dist=384818          .
## 184 dist=207653;dist=384806          .
## 185 dist=207792;dist=384667          .
## 186 dist=207927;dist=384532          .
## 187 dist=208120;dist=384339          .
## 188 dist=208232;dist=384227          .
## 189 dist=208451;dist=384008          .
## 190 dist=208455;dist=384004          .
## 191 dist=208556;dist=383903          .
## 192 dist=208631;dist=383828          .
## 193 dist=209196;dist=383263          .
## 194 dist=209554;dist=382905          0.0012
## 195 dist=209672;dist=382787          .
## 196 dist=209821;dist=382638          .
## 197 dist=211750;dist=380709          .
## 198 dist=211994;dist=380465          .
## 199 dist=212203;dist=380256          0.0008
## 200 dist=212495;dist=379964          .

##      X1000G_AFR X1000G_AMR X1000G_EAS X1000G_EUR X1000G_SAS ExAC_Freq ExAC_AFR
## 1          .          .          .          .          .          .
## 2          .          .          .          .          .          .
## 3          0.0008     0.0014     .          .          0.003     0.0041   .
## 4          .
## 5          .
## 6          .
## 7          .
## 8          .          0.0029     0.001     0.002     .
## 9          .
## 10         .
## 11         .
## 12         .
## 13         .          0.0029     .          .          0.002     0.0072   .
## 14         .
## 15         .
## 16         .
## 17         0.61       0.5        0.49       0.48       0.51      .
## 18         .
## 19         0.53       0.45       0.46       0.44       0.46      .

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## 25	.	.	.	0.002	0.0031	.	.	.
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## 50	.	0.0014	.	0.001
## 51	.	.	.	0.001
## 52	.	0.0014	.	0.003
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## 62	.	.	0.002
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## 81	.	.	0.003	.	.	.
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86
## 87	0.001	.
88
89
## 90	.	0.0014	.	0.007	0.001	.
91
92
## 93	.	0.0014	.	0.003	.	.
94
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## 98	.	0.0014	.	0.002	.	.
99
## 100	0.0072	.
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## 109	0.0008	0.0029
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## 113	.	0.0014	.	0.002	.	.
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## 123	0.0008
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## 125	.	.	.	0.001	.	.
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## 128	.	.	.	0.003	0.001	.	.
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## 134	0.045	0.0029	.	.	0.001	.	.
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## 136	.	0.0043	.	0.003	.	.	.
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## 149	.	.	0.002
## 150	.	.	0.002
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## 163	0.0008	.	.	0.002	.	.	.
## 164	.	.	.	0.003	.	.	.
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## 168	.	.	.	0.003	.	.	.
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## 177	.	.	.	0.001	.	.	.
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## 193 . . . . . .
## 194 . 0.0014 . 0.005 . .
## 195 . . . . . .
## 196 . . . . . .
## 197 . . . . . .
## 198 . . . . . .
## 199 . 0.0014 . 0.003 . .
## 200 . . . . . .

## ExAC_AMR ExAC_EAS ExAC_FIN ExAC_NFE ExAC_OTH ExAC_SAS ESP6500si_ALL
## 1 . . . . . .
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```

	ESP6500si_AA	ESP6500si_EA	CG46	NCI60	dbSNP	COSMIC_ID
197
198
199
200
1
2
## 3	rs531431278	.
4
5
## 6	rs4109702	.
7
## 8	rs567189015	.
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## 13	rs574696181	.
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## 17	rs71253558	.
## 18	rs2319203	.
## 19	rs2945210	.
20
21
## 22	COSN8003327
23
24
## 25	rs555964136	.
26
## 27	rs768174516	.
28
## 29	rs369796600	.
30
31
32
## 33	rs2677954	.
## 34	.	.	0.043	.	rs2689070	COSN17946623
35
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## 39	.	.	0.033	.	rs376209411	COSN17949250
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## 50 . . . rs577472844
## 51 . . . rs546435512
## 52 . . . rs563259499
## 53
## 54 . . . .
## 55 . . . .
## 56 . . . .
## 57 . . . .
## 58 . . . .
## 59 . . . .
## 60 . . . .
## 61 . 0.011 . rs542179028
## 62 . . . rs187461713
## 63 . . . rs71233904
## 64
## 65 . . . .
## 66 . . . .
## 67 . . . .
## 68 . . . .
## 69 . . . .
## 70 . . . rs368119387
## 71 . . . .
## 72 . . . rs375766055
## 73 . . . rs4058854
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## 81 . . . rs573665656
## 82
## 83 . . . .
## 84 . . . .
## 85
## 86
## 87 . . . rs531345805
## 88 . 0.022 . rs2260269
## 89
## 90 . . . rs538642460
## 91
## 92
## 93 . . . rs527289068
## 94 . . . .
## 95 . . . .
## 96 . . . .
## 97
## 98 . . . rs529977713
## 99
## 100 . . . rs548638402
## 101
## 102 . . . rs866421787
## 103 . 0.011 . rs375557516

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## 104	.	0.011	. rs868400103	.
105
## 106	.	.	. rs866552418	.
107
## 108	.	.	. rs371004994	.
## 109	.	.	. rs553312128	.
## 110	.	.	. rs866237684	.
111
## 112	.	.	. rs372462526	COSN24335658
## 113	.	.	. rs535794846	.
## 114	.	.	. rs866797096	.
## 115	.	.	. rs867002641	.
116
117
## 118	.	.	. rs770910751	COSN5135258;COSN21887995
## 119	.	0.011	. rs2945203	.
## 120	.	0.011	. .	.
121
## 122	COSN10020725
## 123	.	.	. rs576659356	.
124
## 125	.	.	. rs180725506	.
## 126	.	.	. rs866126735	.
127
## 128	.	.	. rs530259540	.
129
## 130	.	.	. rs866884174	.
131
132
## 133	.	0.043	. rs4506328	.
## 134	.	.	. rs570081054	.
## 135	.	.	. rs866530850	.
## 136	.	.	. rs532521475	.
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## 140	.	.	. rs4098754	.
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## 145	.	.	. rs4593657	.
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## 149	.	.	. rs561884554	.
## 150	.	.	. rs544543474	.
## 151	.	.	. rs868233999	.
## 152	.	.	. rs868652834	.
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## 154	.	.	. rs373314016	.
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## 157	.	.	. rs768835634	.

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## 158 . . . .
## 159 . . . .
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## 162 . . . .
## 163 . . . rs563054122 .
## 164 . . . rs544292479 .
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## 168 . . . rs571402066 .
## 169 .
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## 171 . . . .
## 172 . . . .
## 173 . . . rs767315388 .
## 174 . . . rs373810525 COSN10020730
## 175 .
## 176 . . . rs751502631 .
## 177 . . . rs571890012 .
## 178 .
## 179 . . . rs374014051 .
## 180 . . . .
## 181 . . . .
## 182 . . . rs796318422 .
## 183 . . . .
## 184 . . . rs796930527 COSN15749470
## 185 . . . rs1832925 .
## 186 .
## 187 . . . rs371904116 .
## 188 . . . rs369316415 .
## 189 .
## 190 . . . rs796968718 COSN1881641
## 191 . . . rs375196402 .
## 192 .
## 193 .
## 194 . . . rs551833403 .
## 195 . . . rs797009701 .
## 196 . . . rs2988252 .
## 197 . . . .
## 198 . . . rs372580324 .
## 199 . . . rs569049492 .
## 200 .

##      COSMIC_DIS ClinVar_SIG ClinVar_DIS ClinVar_ID ClinVar_DB ClinVar_DBID
## 1 . . . .
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## 22  1(pancreas)
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## 34  1(skin)
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## 39  1(skin)
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## 118 1(breast) . . . . . .
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## 122  1(kidney)
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## 190 1(liver)
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##      GWAS_DIS GWAS_OR GWAS_BETA GWAS_PUBMED GWAS_SNP GWAS_P SIFT_score
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##      SIFTConverted_rankscore SIFT_pred Polyphen2_HDIV_score
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##      Polyphen2_HDIV_rankscore Polyphen2_HDIV_pred Polyphen2_HVAR_score
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##      Polyphen2_HVAR_rankscore Polyphen2_HVAR_pred LRT_score
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##      LRTConvertedRankScore LRT_Pred MutationTaster_Score
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##      MutationTaster_converted_rankscore MutationTaster_pred
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##      MutationAssessor_score MutationAssessor_score_rankscore
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##      MutationAssessor_pred FATHMM_score FATHMM_converted_rankscore FATHMM_pred
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##      PROVEAN_score PROVEAN_converted_rankscore PROVEAN_pred VEST3_score
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##      VEST3_rankscore MetaSVM_score MetaSVM_rankscore MetaSVM_pred MetaLR_score
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##      MetaLR_rankscore MetaLR_pred M.CAP_score M.CAP_rankscore M.CAP_pred
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##      fathmm.MKL_coding_score fathmm.MKL_coding_rankscore fathmm.MKL_coding_pred
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##      GenoCanyon_score_rankscore integrated_fitCons_score
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##      phyloP100way_vertebrate_rankscore phyloP20way_mammalian
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##      phyloP20way_mammalian_rankscore phastCons100way_vertebrate

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##      SiPhy_29way_logOdds_rankscore Interpro_domain GTEx_V6_gene GTEx_V6_tissue
## 1          . . . .
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## 200
##      gnomAD_exome_ALL gnomAD_exome_AFR gnomAD_exome_AMR gnomAD_exome_ASJ
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##      gnomAD_exome_EAS gnomAD_exome_FIN gnomAD_exome_NFE gnomAD_exome_OTH
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## 200
##      gnomAD_exome_SAS gnomAD_genome_ALL gnomAD_genome_AFR gnomAD_genome_AMR
## 1          .           0.0020        0.0074          0
## 2          .           3.229e-05       0            0
## 3          .           0.0014        0.0003        0.0024
## 4          .
## 5          .           0.0013        0.0002          0
## 6          .           0.5038        0.5532        0.4973
## 7          .           0.0002          0            0.0012

```

## 8	.	0.0022	0.0006	0.0024
## 9	.	0.0024	0.0008	0.0024
## 10	.			
## 11	.			
## 12	.	0.0003	0.0002	0
## 13	.	0.0013	0.0008	0.0012
## 14	.	0.0012	0.0007	0
## 15	.	0.0012	0.0031	0.0024
## 16	.	0.0007	0	0
## 17	.	0.5106	0.5523	0.5025
## 18	.	0.5113	0.5537	0.5025
## 19	.	0.5110	0.5547	0.5013
## 20	.	0	0	0
## 21	.	0	0	0
## 22	.	0.0031	0.0099	0.0012
## 23	.	3.228e-05	0.0001	0
## 24	.	0.0001	0	0
## 25	.	0.0010	0.0001	0
## 26	.	6.456e-05	0.0001	0
## 27	.	0.0072	0.0008	0.0061
## 28	.			
## 29	.	0.0020	0.0013	0
## 30	.	0.0007	0.0001	0
## 31	.			
## 32	.	0.0018	0.0005	0.0012
## 33	.	0.5105	0.5525	0.5038
## 34	.	0.5107	0.5530	0.5012
## 35	.			
## 36	.	0.0022	0.0011	0.0012
## 37	.	3.227e-05	0	0
## 38	.	0.0003	0.0001	0.0012
## 39	.	0.5120	0.5557	0.5037
## 40	.	3.228e-05	0	0
## 41	.			
## 42	.	0.0005	0.0001	0
## 43	.	0.0011	0.0001	0.0024
## 44	.			
## 45	.	0.0051	0.0185	0.0012
## 46	.			
## 47	.	0.0003	0	0
## 48	.			
## 49	.			
## 50	.	0.0003	0	0.0012
## 51	.	0.0018	0.0003	0
## 52	.	0.0016	0.0003	0
## 53	.			
## 54	.	0.0057	0.0209	0
## 55	.	0.0002	0	0
## 56	.			
## 57	.	0.0017	0.0060	0
## 58	.	0.0008	0.0031	0
## 59	.	0.0002	0	0
## 60	.	0.0002	0.0005	0
## 61	.	0.0101	0.0027	0.0027

## 62	.	0.0026	0.0006	0
## 63	.	0.0033	0.0022	0.0171
## 64	.	0.0004	0.0001	0
## 65	.	3.228e-05	0	0.0012
## 66	.	0.0002	0.0002	0
## 67	.	0.0005	0	0
## 68	.	0.0054	0.0173	0.0024
## 69	.	0.0021	0.0007	0.0012
## 70	.	0.0021	0.0010	0
## 71	.	0.0009	0.0003	0
## 72	.	0.0170	0.0491	0.0121
## 73	.	0.5090	0.5516	0.5013
## 74	.	0.0007	0.0003	0
## 75	.			
## 76	.	9.684e-05	0	0
## 77	.			
## 78	.	0.0006	0.0003	0
## 79	.			
## 80	.	0.0002	0	0
## 81	.	0.0015	0.0011	0.0012
## 82	.			
## 83	.	0.0015	0.0009	0.0036
## 84	.	0.0009	0.0006	0
## 85	.			
## 86	.			
## 87	.			
## 88	.	0.5077	0.5615	0.4925
## 89	.			
## 90	.	0.0017	0.0003	0
## 91	.			
## 92	.	0	0	0
## 93	.	0.0002	0.0002	0.0012
## 94	.	6.455e-05	0	0
## 95	.	0.0003	0.0001	0
## 96	.	0.0015	0.0053	0.0012
## 97	.			
## 98	.	0.0013	0.0006	0
## 99	.			
## 100	.			
## 101	.			
## 102	.	0.0180	0.0688	0.0085
## 103	.	0.0336	0.1096	0.0099
## 104	.	0.0224	0.0877	0.0049
## 105	.	0	0	0
## 106	.	0.0024	0.0007	0.0036
## 107	.	0.0016	0.0001	0
## 108	.	0.0315	0.1098	0.0123
## 109	.	0.0003	0.0003	0
## 110	.	0.0049	0.0173	0.0024
## 111	.	0	0	0
## 112	.	0.0079	0.0293	0.0024
## 113	.	0.0010	0.0003	0.0012
## 114	.	0.0044	0.0010	0.0048
## 115	.	0.0006	0.0006	0.0012

## 116				
## 117	.	3.227e-05	0	0
## 118	.	0.0010	0.0007	0
## 119	.	0.0221	0.0875	0.0049
## 120	.	0.0033	0.0003	0
## 121	.	9.682e-05	0	0
## 122	.	0.0030	0.0011	0.0036
## 123	.	9.688e-05	0	0
## 124	.	6.456e-05	0	0
## 125	.	0.0009	0.0009	0.0012
## 126	.	0.0030	0.0105	0
## 127	.	0.0002	0	0
## 128	.	0.0013	0.0002	0
## 129	.	0.0001	0	0
## 130	.	0.0046	0.0163	0.0036
## 131	.	0.0005	0.0005	0
## 132	.	3.227e-05	0	0
## 133	.	0.5089	0.5921	0.4211
## 134	.	0.0066	0.0243	0.0036
## 135	.	0.0019	0.0063	0
## 136	.	0.0007	0.0001	0
## 137	.	0.0011	0.0003	0
## 138				
## 139				
## 140	.	0.5095	0.5923	0.4256
## 141				
## 142	.	3.227e-05	0	0
## 143				
## 144	.	9.682e-05	0	0
## 145	.	0.0141	0.0199	0.0101
## 146	.	0.0005	0.0001	0
## 147	.	0.0003	0.0002	0
## 148	.	0.0035	0.0009	0
## 149	.	0.0010	0.0003	0.0012
## 150	.	0.0003	0	0
## 151	.	0.0007	0	0.0012
## 152	.	0.0018	0.0002	0.0012
## 153				
## 154	.	0.0287	0.0851	0.0024
## 155				
## 156	.	0.0004	0	0
## 157	.	0.0051	0.0181	0.0048
## 158	.	0.0010	0	0
## 159	.	0.0007	0	0
## 160	.	3.227e-05	0.0001	0
## 161	.	0.0065	0.0237	0.0012
## 162				
## 163	.	0.0003	0.0001	0.0012
## 164	.	0.0012	0.0002	0
## 165	.	0.0011	0.0002	0
## 166	.	6.455e-05	0	0
## 167	.	0.0014	0.0007	0.0012
## 168	.	0.0011	0	0
## 169				

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## 170 . 0.0003 0.0003 0
## 171 . 0.0023 0.0074 0.0012
## 172 . 0.0030 0.0015 0
## 173 . 0.0006 0.0021 0
## 174 . 0.0085 0.0091 0
## 175
## 176 . 0.0032 0.0013 0.0073
## 177 . 6.462e-05 0.0001 0
## 178
## 179 . 0.0083 0.0104 0
## 180 . 0 0 0
## 181 . 0.0047 0.0002 0
## 182 . 0.0096 0.0187 0
## 183 . 0.0004 0.0002 0
## 184 . 0.0098 0.0123 0
## 185 . 0.4904 0.5508 0.4809
## 186
## 187 . 0.0073 0.0006 0.0012
## 188 . 0.0162 0.0219 0
## 189
## 190 . 0.0332 0.1149 0.0111
## 191 . 0.0106 0.0136 0
## 192 . 3.229e-05 0 0
## 193
## 194 . 0.0032 0.0010 0.0024
## 195 . 0.0074 0.0006 0
## 196 . 0.5607 0.6752 0.5102
## 197 . 3.227e-05 0 0
## 198 . 3.229e-05 0 0
## 199 . 0.0034 0.0007 0
## 200
##      gnomAD_genome_ASJ gnomAD_genome_EAS gnomAD_genome_FIN gnomAD_genome_NFE
## 1          0             0                 0             0
## 2          0             0.0006            0             0
## 3          0             0                 0.0029        0.0017
## 4
## 5          0             0                 0.0023        0.0019
## 6          0.4661        0.4938            0.4870        0.4786
## 7          0             0                 0             0.0003
## 8          0             0.0050            0.0029        0.0026
## 9          0.0068        0.0006            0.0032        0.0031
## 10
## 11
## 12          0             0                 0             0.0003
## 13          0.0243        0                 0.0003        0.0013
## 14          0             0                 0.0006        0.0019
## 15          0             0                 0             0.0003
## 16          0             0                 0.0003        0.0014
## 17          0.4892        0.5019            0.4936        0.4921
## 18          0.4893        0.5019            0.4949        0.4922
## 19          0.4892        0.5013            0.4934        0.4909
## 20          0             0                 0             0
## 21          0             0                 0             0
## 22          0.0067        0                 0             0.0005

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## 23	0	0	0	0
## 24	0	0	0	0.0003
## 25	0	0	0.0003	0.0017
## 26	0	0	0	6.661e-05
## 27	0.0139	0	0.0137	0.0101
## 28				
## 29	0	0.0314	0	0
## 30	0	0	0.0037	0.0006
## 31				
## 32	0.0069	0	0.0003	0.0030
## 33	0.4895	0.5026	0.4919	0.4916
## 34	0.4894	0.5019	0.4921	0.4922
## 35				
## 36	0	0	0.0032	0.0029
## 37	0	0	0	6.659e-05
## 38	0.0033	0	0	0.0003
## 39	0.4895	0.5026	0.4951	0.4924
## 40	0	0	0	6.66e-05
## 41				
## 42	0.0033	0	0	0.0008
## 43	0	0	0.0006	0.0018
## 44				
## 45	0	0	0	0
## 46				
## 47	0	0	0	0.0006
## 48				
## 49				
## 50	0	0	0	0.0004
## 51	0	0	0.0064	0.0020
## 52	0	0	0.0064	0.0015
## 53				
## 54	0	0	0	0
## 55	0	0	0.0003	0.0003
## 56				
## 57	0	0.0019	0	0.0006
## 58	0	0	0	0.0002
## 59	0	0.0008	0.0027	0.0002
## 60	0	0	0	8.691e-05
## 61	0	0	0.0114	0.0151
## 62	0.0103	0	0.0090	0.0021
## 63	0.0035	0.0026	0.0029	0.0030
## 64	0	0	0	0.0007
## 65	0	0	0	0
## 66	0	0	0	0.0002
## 67	0	0.0025	0	0.0007
## 68	0	0	0.0011	0.0007
## 69	0	0	0.0026	0.0030
## 70	0.0033	0.0063	0.0038	0.0019
## 71	0	0.0012	0.0003	0.0015
## 72	0.0138	0.0257	0.0017	0.0036
## 73	0.4852	0.5027	0.4916	0.4887
## 74	0	0	0.0006	0.0011
## 75				
## 76	0	0	0	0.0002

## 77				
## 78	0	0	0.0006	0.0010
## 79				
## 80	0	0	0.0006	0.0001
## 81	0	0	0.0006	0.0021
## 82				
## 83	0.0066	0	0.0006	0.0021
## 84	0	0	0.0011	0.0013
## 85				
## 86				
87
## 88	0.4698	0.5020	0.4912	0.4809
## 89				
## 90	0	0	0	0.0032
## 91				
## 92	0	0	0	0
## 93	0	0	0	0.0001
## 94	0	0	0	0.0001
## 95	0	0	0	0.0006
## 96	0	0	0	0
## 97				
## 98	0	0	0.0017	0.0019
## 99				
100
## 101				
## 102	0	0	0	0.0001
## 103	0.0207	0.0334	0.0047	0.0059
## 104	0	0	0	0.0003
## 105	0	0	0	0
## 106	0	0	0.0052	0.0029
## 107	0	0	0.0046	0.0021
## 108	0.0139	0.0296	0.0009	0.0039
## 109	0	0.0006	0.0003	0.0002
## 110	0.0033	0	0	0.0002
## 111	0	0	0	0
## 112	0.0033	0	0	6.661e-05
## 113	0	0	0.0006	0.0016
## 114	0.0067	0	0.0017	0.0075
## 115	0	0	0	0.0007
## 116				
## 117	0	0	0	6.661e-05
## 118	0	0	0.0003	0.0017
## 119	0	0	0	0.0001
## 120	0	0	0.0106	0.0038
## 121	0	0.0012	0	6.66e-05
## 122	0	0	0.0050	0.0038
## 123	0	0	0	0.0001
## 124	0	0	0	0.0001
## 125	0	0	0.0009	0.0009
## 126	0	0.0013	0	6.664e-05
## 127	0	0.0043	0	0
## 128	0	0	0.0009	0.0024
## 129	0	0	0	0.0003
## 130	0.0033	0	0	0

## 131	0	0	0	0.0008
## 132	0.0033	0	0	0
## 133	0.5106	0.4676	0.4716	0.4763
## 134	0	0	0	6.684e-05
## 135	0.0034	0	0	0.0001
## 136	0	0	0.0011	0.0011
## 137	0	0	0.0006	0.0019
## 138				
## 139				
## 140	0.5110	0.4695	0.4778	0.4759
## 141				
## 142	0	0	0	6.66e-05
## 143				
## 144	0	0	0	0.0002
## 145	0.0077	0.0209	0.0167	0.0098
## 146	0	0	0.0003	0.0009
## 147	0	0	0	0.0005
## 148	0	0	0.0091	0.0044
## 149	0	0	0.0041	0.0009
## 150	0	0	0	0.0007
## 151	0	0	0	0.0013
## 152	0.01	0	0.0009	0.0031
## 153				
## 154	0.01	0.0854	0.0049	0.0027
## 155				
## 156	0	0	0.0009	0.0006
## 157	0	0	0	0
## 158	0	0	0.0046	0.0008
## 159	0.0034	0	0.0023	0.0009
## 160	0	0	0	0
## 161	0	0	0	6.663e-05
## 162				
## 163	0	0	0.0009	0.0003
## 164	0	0	0.0020	0.0018
## 165	0	0	0.0011	0.0017
## 166	0	0	0	0.0001
## 167	0	0	0.0006	0.0021
## 168	0	0	0.0061	0.0007
## 169				
## 170	0	0	0	0.0005
## 171	0.0068	0	0	0.0002
## 172	0	0	0.0023	0.0047
## 173	0	0	0	0.0001
## 174	0.0033	0.0866	0.0044	0.0025
## 175				
## 176	0	0	0.0009	0.0051
## 177	0	0	0	6.672e-05
## 178				
## 179	0.0033	0.0774	0.0049	0.0019
## 180	0	0	0	0
## 181	0.0066	0.0703	0.0030	0.0016
## 182	0.0066	0.0674	0.0035	0.0015
## 183	0	0.0019	0	0.0004
## 184	0.0033	0.0892	0.0041	0.0029

## 185	0.3934	0.4972	0.4846	0.4551
## 186				
## 187	0.0066	0.1017	0.0044	0.0027
## 188	0.0067	0.1205	0.0098	0.0050
## 189				
## 190	0.0105	0.0339	0.0009	0.0042
## 191	0.0066	0.0910	0.0064	0.0029
## 192	0	0	0	6.668e-05
## 193				
## 194	0	0	0.0032	0.0049
## 195	0.0067	0.0955	0.0064	0.0033
## 196	0.5252	0.5490	0.5172	0.5079
## 197	0	0	0	6.66e-05
## 198	0	0	0	6.66e-05
## 199	0.0066	0	0.0011	0.0060
## 200				
##	gnomAD_genome_OTH	Otherinfo	Otherinfo.1	
## 1	0	unknown	.	
## 2	0	unknown	.	
## 3	0.0031	hom	.	
## 4		hom	.	
## 5	0.0020	hom	.	
## 6	0.4857	hom	.	
## 7	0	hom	.	
## 8	0.0041	hom	.	
## 9	0.0041	hom	.	
## 10		hom	.	
## 11		hom	.	
## 12	0.0010	hom	.	
## 13	0.0041	hom	.	
## 14	0.0010	hom	.	
## 15	0.0021	hom	.	
## 16	0	hom	.	
## 17	0.4968	het	.	
## 18	0.4978	het	.	
## 19	0.4967	hom	.	
## 20	0	hom	.	
## 21	0	hom	.	
## 22	0	hom	.	
## 23	0	hom	.	
## 24	0	hom	.	
## 25	0.0031	hom	.	
## 26	0	hom	.	
## 27	0.0094	hom	.	
## 28		hom	.	
## 29	0.0010	hom	.	
## 30	0	hom	.	
## 31		het	.	
## 32	0.0020	hom	.	
## 33	0.4957	hom	.	
## 34	0.4968	hom	.	
## 35		het	.	
## 36	0.0021	hom	.	
## 37	0	hom	.	

## 38	0.0010	hom	.
## 39	0.4967	hom	.
## 40	0	hom	.
## 41		hom	.
## 42	0	hom	.
## 43	0.0010	hom	.
## 44		hom	.
## 45	0	hom	.
## 46		het	.
## 47	0	unknown	.
## 48		hom	.
## 49		hom	.
## 50	0.0010	hom	.
## 51	0.0021	hom	.
## 52	0.0010	hom	.
## 53		hom	.
## 54	0.0010	hom	.
## 55	0	hom	.
## 56		hom	.
## 57	0	hom	.
## 58	0	hom	.
## 59	0	hom	.
## 60	0	hom	.
## 61	0.0100	hom	.
## 62	0.0045	hom	.
## 63	0.0090	hom	.
## 64	0	unknown	.
## 65	0	hom	.
## 66	0	hom	.
## 67	0	hom	.
## 68	0.0041	hom	.
## 69	0.0031	hom	.
## 70	0.0020	hom	.
## 71	0	hom	.
## 72	0.0073	hom	.
## 73	0.4955	hom	.
## 74	0.0010	hom	.
## 75		hom	.
## 76	0	hom	.
## 77		het	.
## 78	0	hom	.
## 79		hom	.
## 80	0.0010	hom	.
## 81	0.0031	hom	.
## 82		hom	.
## 83	0.0010	hom	.
## 84	0	hom	.
## 85		hom	.
## 86		het	.
## 87	.	hom	.
## 88	0.4876	hom	.
## 89		hom	.
## 90	0.0020	hom	.
## 91		hom	.

## 92	0	hom	.
## 93	0	hom	.
## 94	0	hom	.
## 95	0	hom	.
## 96	0.0010	hom	.
## 97		hom	.
## 98	0.0020	hom	.
## 99		hom	.
## 100	.	hom	.
## 101		hom	.
## 102	0.0021	hom	.
## 103	0.0137	hom	.
## 104	0.0062	hom	.
## 105	0	hom	.
## 106	0.0031	hom	.
## 107	0	hom	.
## 108	0.0156	hom	.
## 109	0.0010	hom	.
## 110	0.0020	hom	.
## 111	0	hom	.
## 112	0.0010	hom	.
## 113	0.0010	hom	.
## 114	0.0041	hom	.
## 115	0.0010	hom	.
## 116		hom	.
## 117	0	hom	.
## 118	0	hom	.
## 119	0.0010	hom	.
## 120	0.0062	hom	.
## 121	0	hom	.
## 122	0.0062	hom	.
## 123	0.0010	hom	.
## 124	0	hom	.
## 125	0.0010	hom	.
## 126	0.0020	hom	.
## 127	0	hom	.
## 128	0	hom	.
## 129	0	hom	.
## 130	0.0010	hom	.
## 131	0	hom	.
## 132	0	hom	.
## 133	0.4718	het	.
## 134	0	hom	.
## 135	0	hom	.
## 136	0	hom	.
## 137	0	hom	.
## 138		hom	.
## 139		hom	.
## 140	0.4731	het	.
## 141		het	.
## 142	0	hom	.
## 143		hom	.
## 144	0	hom	.
## 145	0.0150	hom	.

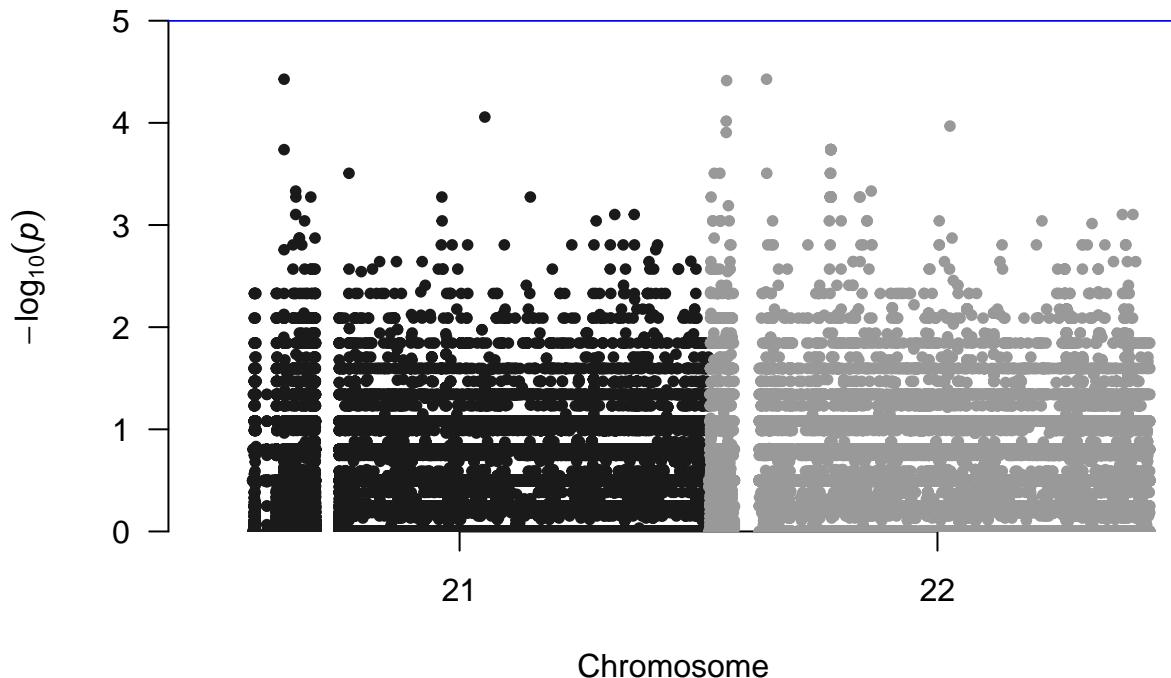
## 146	0.0010	hom	.
## 147	0	hom	.
## 148	0.0031	hom	.
## 149	0	hom	.
## 150	0	hom	.
## 151	0.0031	hom	.
## 152	0	hom	.
## 153		het	.
## 154	0.0146	hom	.
## 155		hom	.
## 156	0.0010	hom	.
## 157	0.0010	hom	.
## 158	0.0020	hom	.
## 159	0.0010	hom	.
## 160	0	hom	.
## 161	0	hom	.
## 162		het	.
## 163	0.0010	hom	.
## 164	0	hom	.
## 165	0.0020	hom	.
## 166	0	hom	.
## 167	0.0031	hom	.
## 168	0.0010	hom	.
## 169		hom	.
## 170	0	hom	.
## 171	0	hom	.
## 172	0.0021	hom	.
## 173	0	hom	.
## 174	0.0042	hom	.
## 175		het	.
## 176	0.0031	hom	.
## 177	0	hom	.
## 178		hom	.
## 179	0.0073	hom	.
## 180	0	unknown	.
## 181	0.0053	hom	.
## 182	0.0052	hom	.
## 183	0	hom	.
## 184	0.0052	hom	.
## 185	0.4585	hom	.
## 186		hom	.
## 187	0.0083	hom	.
## 188	0.0103	hom	.
## 189		hom	.
## 190	0.0106	hom	.
## 191	0.0083	hom	.
## 192	0	hom	.
## 193		hom	.
## 194	0.0031	hom	.
## 195	0.0083	unknown	.
## 196	0.5188	hom	.
## 197	0	hom	.
## 198	0	hom	.
## 199	0.0051	hom	.

```
## 200
```

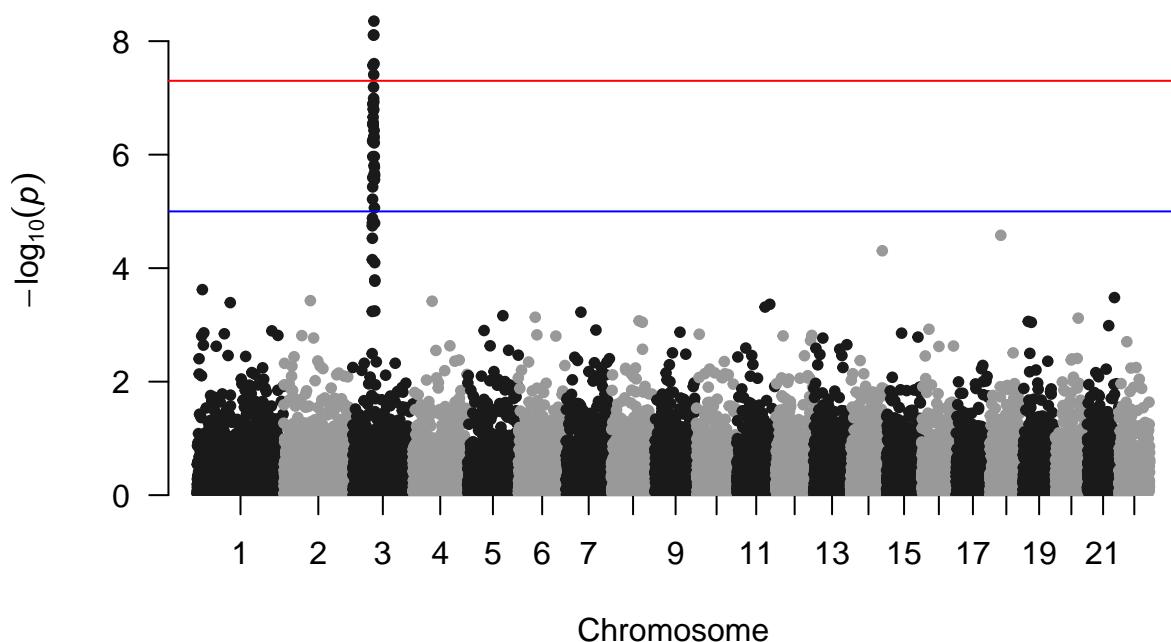
```
hom
```

Make the Manhattan plot on the final dataset Make the Manhattan plot on the gwasResults dataset

```
manhattan(final, chr="CHR", bp="SNP", snp="dbSNP", p="UNADJ" )
```

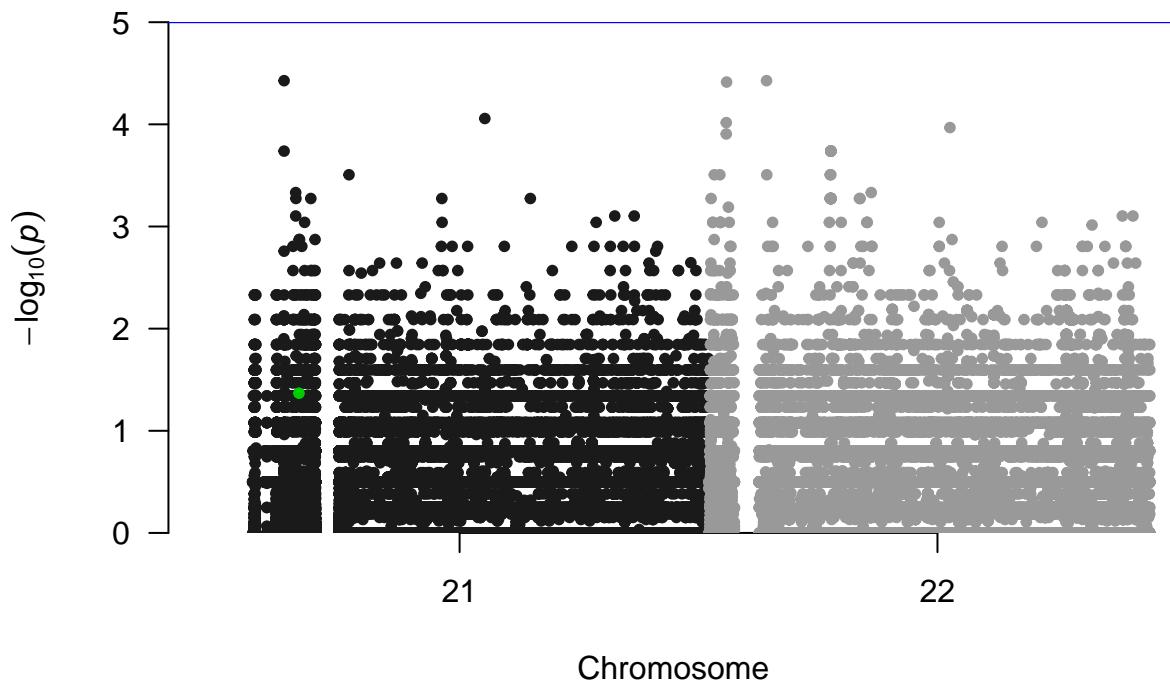


```
manhattan(gwasResults)
```

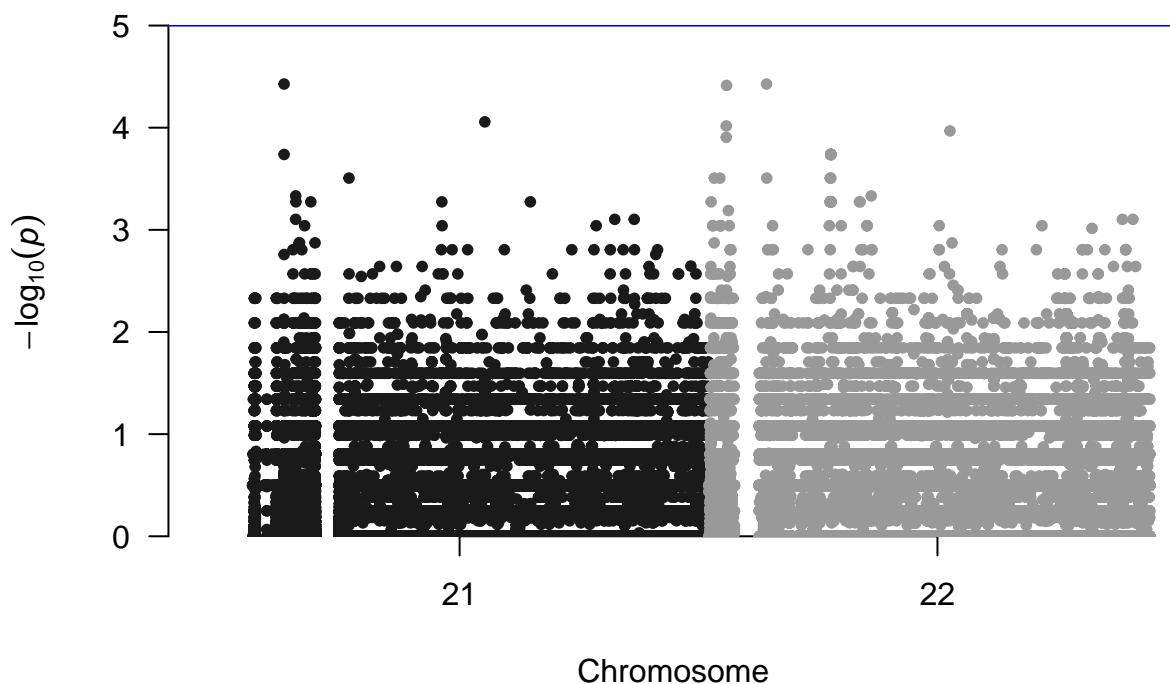


A list of SNP of interest is provided with the library: Let's highlight them, with a bit of customization on the plot

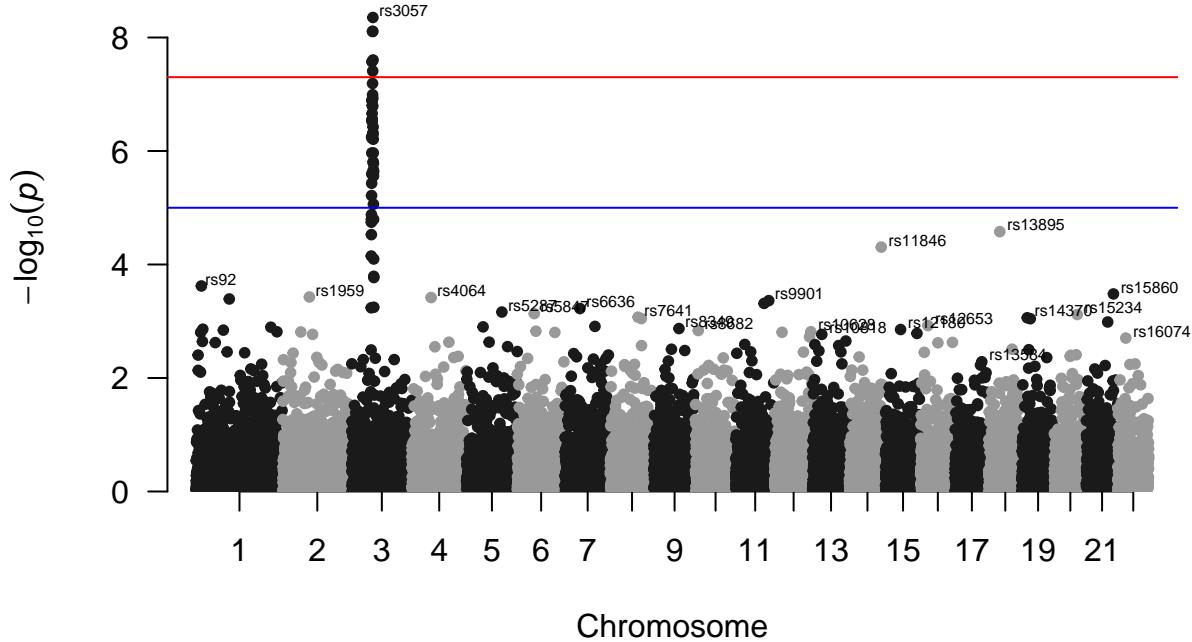
```
manhattan(final, chr="CHR", bp="SNP", snp="dbSNP", p="UNADJ", highlight = c("rs370508396"))
```



```
manhattan(final, chr="CHR", bp="SNP", snp="dbSNP", p="UNADJ", annotatePval = 0.5)
```



```
manhattan(gwasResults, annotatePval = 0.05)
```



We have tons of other metadata, what else would be interesting to explore?

circular version with CMplot

The CMplot library by Lilin Yin is a good choice if you want to make a circular version of your manhattanplot. I believe than doing a circular version makes sense: it gives less space to all the non significant SNPs that do not interest us, and gives more space for the significant association. Moreover, the CMplot makes their realization straightforward.

```
library("CMplot")

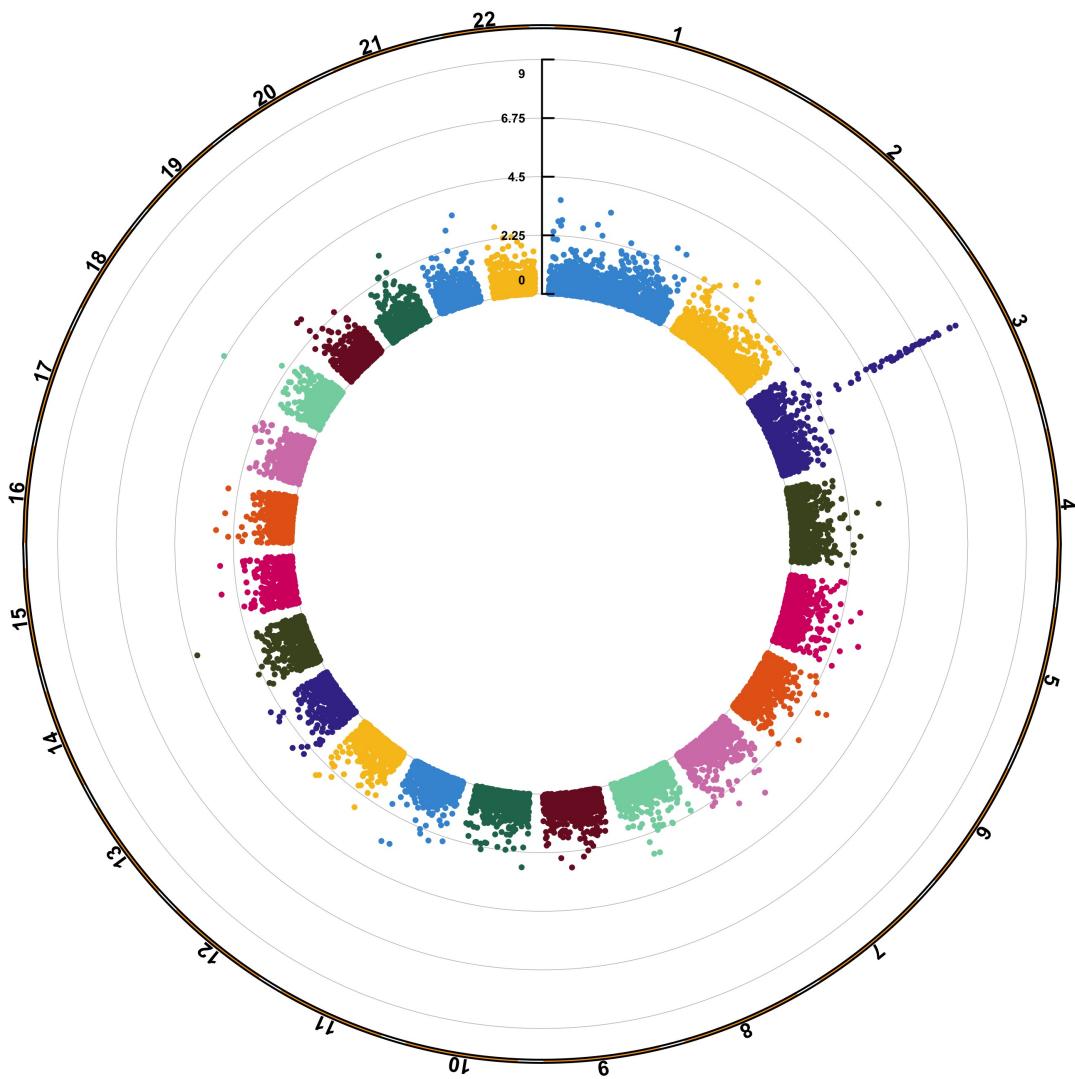
## Much appreciate for using CMplot.

## Full description, Bug report, Suggestion and the latest codes:

## https://github.com/YinLiLin/CMplot

CMplot(gwasResults, plot.type="c", r=1.6, cir.legend=TRUE,
       outward=TRUE, cir.legend.col="black", cir.chr.h=.1 ,chr.den.col="orange", file="jpg",
       memo="", dpi=300, chr.labels=seq(1,22))

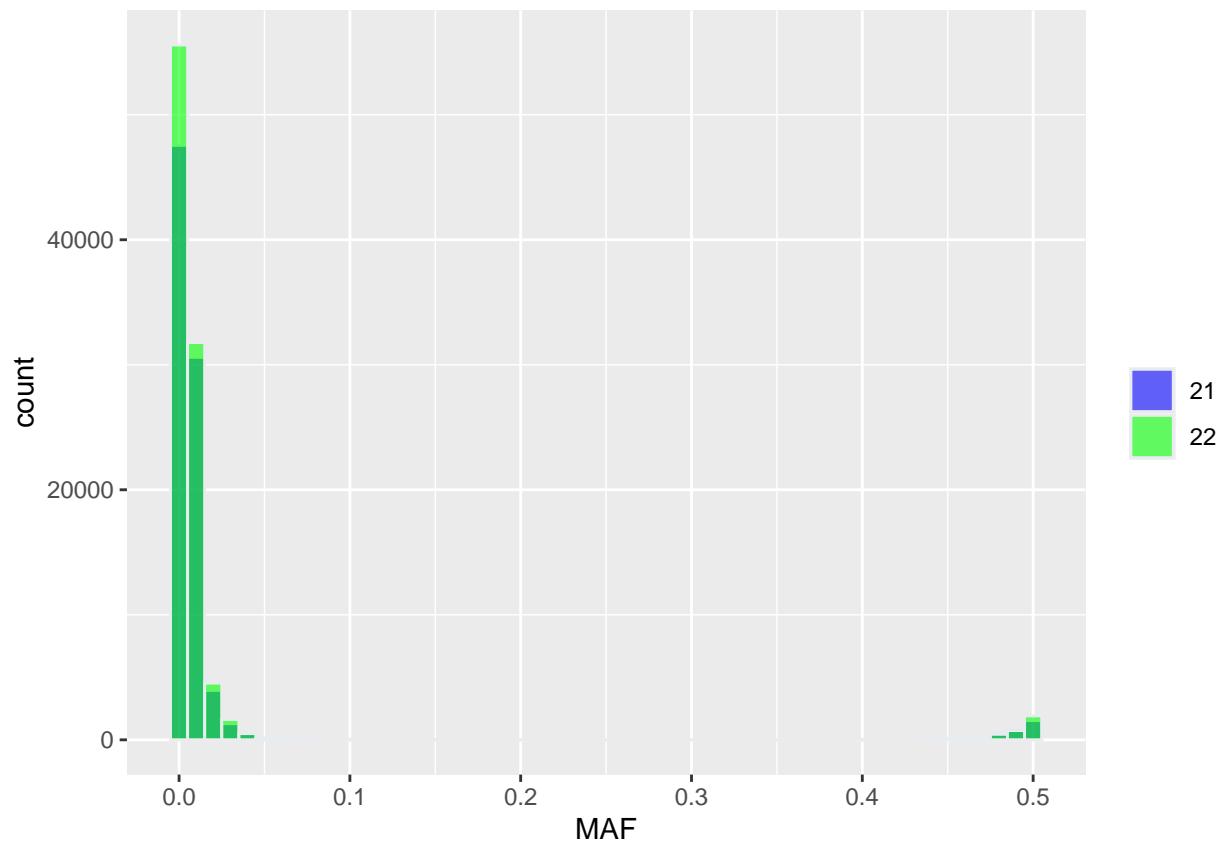
## Circular-Manhattan Plotting P.
## Plots are stored in: /Users/keithmitchell/Desktop/Repositories/2021-July-Genome-Wide-Association-St
```



Explore the metadata:

- MAF

```
p <- ggplot(final, aes(x=MAF, fill=as.factor(CHR))) + geom_histogram( color="#e9ecf", binwidth = 0.01,
  scale_fill_manual(values=c("blue", "green")) +
  labs(fill=""))
p
```



Isolating a gene of interest: