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Text

Query results may be downloaded in formatted Excel and tab-delimited text formats

Query Messages

Occasionally you will find messages with information about the query.

NOTE: currently, for 1000 Genomes data, genes entered in the *Enter Genomic Features* section will only retrieve variants that are SNPs in dbSNP.

The exact SQL query passed to MySQL.

Query Used

```
1 | SELECT QTBL.* FROM bioq_1000genomes_ip1r.vcf_sites AS QTBL
2 |   INNER JOIN my_vars MV ON (QTBL.pos_global = MV.pos_global)
3 |   LIMIT 1000
```

Copy to advanced query section

The query can be pasted into the advanced queries section where it can be revised using powerful SQL to filter rows or add and remove columns, possibly from other databases.

Additional information about the query and genomic features entered.

Additional Information

Summary

SNP/Gene/Region Queries

SNPs

Genes

Program Parameters

This table shows the values of various parameters selected for this query, as well as some parameters that are for internal use only.

name	value	description
add-ld-proxies	0	Add LD proxies to list of SNPs
bioq-db	run_info_bioq	Name of dbSNP Web Tool MySQL database - reference tables
bioq-ld-db	bioq_hapmap_pr28_dbsnp132	Database for LD estimates
bioq-ld-table	ceu_plink_ld	Table for LD estimates with columns source,proxy,rsq