

Generated on July 31, 2015

Genome Browser

Browse related tracks on the genome browser by clicking on the link(s) below:

View tracks for this study in the region around the INS gene

H3K4me1, H3K4me2, H3K4me3, and

H3K27me3 Peak Calls

Lists of Locations

Use the following form(s) to refine the parameters and add the list of genomic sequences corresponding to peak calls to a strategy. Depending on your choices, these searches may be slow.

H3K4me2 Histone Modification in Normal Human Islets (GLITR Pipeline:Pooled Unique-Best-Match Reads)

Retrieve:

Whole Genome

Peaks in a Region of Interest (specify below):

chr5

Enter a region (e.g., chr:start-stop) or enter just the chromosome (e.g., chr12 or chrX) to search for peaks on a single chromosome. Select the "Whole Genome" option or leave the text box blank to return all results from this analysis.

Find Locations

H3K4me3 Histone Modification in Normal Human Islets (GLITR Pipeline:Pooled Unique-Best-Match Reads)

H3K27me3 Histone Modification in Normal Human Islets (GLITR Pipeline:Pooled Unique-Best-Match Reads)

H3K4me1 Histone Modification in Normal Human Islets (GLITR Pipeline:Pooled Unique-Best-Match Reads)

Repositories

Kaestner Lab

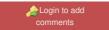


Stock #: Not provided

Availability Notes: Not provided

Comments

There are no comments for this entry.



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