FDb.UCSC.snp135common.hg19
January 17, 2024

FDb.UCSC.snp135common.hg19
UCSC common SNPs track, dbSNP build 135, genome assembly hg19

Description
FDb.UCSC.snp135common.hg19 is a FeatureDb created from the UCSC common SNPs track for dbSNP build 135 and genome assembly hg19. The track is too large to automatically construct a FeatureDb using makeFeatureDbFromUCSC, so here it has been manually retrieved (see inst/build/FDb.UCSC.snp135common.hg19.R) and saved as a FeatureDb.

Author(s)
Tim Triche, Jr.

See Also
features makeFeatureDbFromUCSC

Examples
## load the library
library(FDb.UCSC.snp135common.hg19)

## list the contents that are loaded into memory
ls('package:FDb.UCSC.snp135common.hg19')

## show the db object that is loaded by calling it's name
FDb.UCSC.snp135common.hg19

## extract features for use in annotating data
snp135common <- features(FDb.UCSC.snp135common.hg19)## need to fetch genome
met <- metadata(FDb.UCSC.snp135common.hg19)
genome(snp135common) <- met[which(met[, 'name'] == 'Genome'), 'value']
Index

* data
  FDb.UCSC.snp135common.hg19, 1
* package
  FDb.UCSC.snp135common.hg19, 1

FDb.UCSC.snp135common.hg19, 1
FDb.UCSC.snp135common.hg19-package
  (FDb.UCSC.snp135common.hg19), 1
features, /

makeFeatureDbFromUCSC, /