## MafDb.ESP6500SI.V2.SSA137

April 10, 2016

MafDb.ESP6500SI.V2.SSA137-package

Annotation package for minor allele frequency data from the NHLBI ESP project

#### **Description**

This annotation package stores minor allele frequency (MAF) data values frozen from the NHLBI Exome Sequencing (ESP) project. The data is stored in the form of a SQLite database and is loaded automatically in the form of a MafDb object. The name of the exposed object matches the name of the package and part of the filename that contained the data imported into the package. The class definition and methods to access MafDb objects are found in the VariantFiltering software package.

#### **Format**

MafDb . ESP6500SI . V2 . SSA137 MafDb object containing MAF values from 6500 ESP exomes downloaded in October 2015

## Author(s)

R. Castelo

## Source

Exome Variant Server, NHLBI GO Exome Sequencing Project (ESP), Seattle, WA (URL: http://evs.gs.washington.edu/EVS) [October, 2015, accessed]

#### See Also

makeMafDbPackageESP MafDb.ALL.wgs.phase1.release.v3.20101123 MafDb.ALL.wgs.phase3.release.v5a.20130502 MafDb-class snpid2maf VariantFiltering

#### **Examples**

```
library(MafDb.ESP6500SI.V2.SSA137)

ls("package:MafDb.ESP6500SI.V2.SSA137")

MafDb.ESP6500SI.V2.SSA137

knownVariantsMAFcols(MafDb.ESP6500SI.V2.SSA137)

## lookup allele frequencies for rs1129038, an SNP associated to blue and brown eye colors
## as reported in Eiberg et al. Blue eye color in humans may be caused by a perfectly associated
## founder mutation in a regulatory element located within the HERC2 gene inhibiting OCA2 expression.
## Human Genetics, 123(2):177-87, 2008 [http://www.ncbi.nlm.nih.gov/pubmed/18172690]
snpid2maf(MafDb.ESP6500SI.V2.SSA137, "rs1129038")
```

makeMafDbPackageESP

Make a MafDb annotation data package from the NHLBI GO ESP Project

#### Description

This function creates an annotation data package for a MafDb object. Its primary purpose is to ease the task of fetching and packaging newer minor allele frequency (MAF) data. Note that MAF values are processed and stored in a way to reduce their space in disk. Please consult the manual page for the MafDb-class to know the details of these processing steps.

### Usage

makeMafDbPackageESP(destDir=path.expand("~"), MafDbURL=MafDbESPdefaultURL, MafDbPkgName=MafDbESPdefagenome="hg19", version=NULL, author=NULL, maintainer=NULL, license=NULL, yieldSize=1

#### **Arguments**

destDir Destination directory for the newly created package. MafDbURL URL to the source VCF file(s). By default, it points to the URL holding the data currently stored in the package that defines this function. Name of the newly created package. This will define as well the name of the MafDbPkgName exported MafDb object. Version of the human genome, following UCSC nomenclature. Necessary for genome the internal call to the readVcf() function from the VariantAnnotation package. version Version to put on the new package. By default, the version corresponds to the version of the package that defines this function, bumping the second version number. author Author to put on the new package. By default, the author corresponds to the author of the package that defines this function.

maintainer	Maintainer to put on	the new package.	By default,	the maintainer corresponds
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to the maintainer of the package that defines this function.

license License to put on the new package. By default, the license corresponds to the

license of the package that defines this function.

yieldSize In the case source tabix VCF files, they are not read at once, but scanned in

batches whose size is determined by this argument. By default is set to one

million variants but it may be reduced to lower main memory requirements.

#### Value

Path to the folder containing the created data package.

### Author(s)

R. Castelo

#### See Also

```
MafDb-class snpid2maf MafDb.ESP6500SI.V2.SSA137
```

## **Examples**

```
MafDbESPdefaultURL ## default URL from where makeMafDbPackageESP() fetches the MAF data

MafDbESPdefaultPkgName ## default name for the package that makeMafDbPackageESP() creates

## Not run:

## the previous two default values can be overridden when calling makeMafDbPackageESP()

makeMafDbPackageESP()

## End(Not run)
```

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