**Description**

This annotation package stores phastCons conservation scores from UCSC for the human genome (hg38), calculated from genome-wide multiple alignments with other 99 vertebrate species. The data are stored in the form of Rle objects and are loaded automatically as an object of class GScores. 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 GScores objects are found in the GenomicScores software package.

**Format**

**phastCons100way.UCSC.hg38**  
GScores object containing phastCons conservation scores from UCSC for the human genome.

**Author(s)**

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**Source**

UCSC Genome Browser (URL: http://genome.ucsc.edu) [March, 2018, accessed]
See Also

GScores GenomicScores

Examples

library(GenomicRanges)
library(phastCons100way.UCSC.hg38)

ls("package:phastCons100way.UCSC.hg38")

phast <- phastCons100way.UCSC.hg38
phast
citation(phast)
gscores(phast, GRanges("chr7:117592326-117592330"))
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