Description

Package provides clinical, expression, cnv and mutation data from Genome Cancer Browser for 12 cohorts types. Converting those datasets from data.frames to Bioconductor classes is explained here convertTCGA.

Usage

- `clinical.cb`
- `cnv.cb`
- `mutation.cb`
- `expression.cb1`
- `expression.cb2`

Value

Data frames PANCAN12 data.

Source

[https://genome-cancer.ucsc.edu/proj/site/hgHeatmap/](https://genome-cancer.ucsc.edu/proj/site/hgHeatmap/)
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