Package 'AdverseEvents'

November 4, 2024

Title 'shiny' Application for Adverse Event Analysis of 'OnCore' Data
Version 0.0.3
Author Z Thompson [aut, cre], A Obermayer [aut], DT Chen [aut]
Maintainer Z Thompson < zachary.thompson@moffitt.org>
Description An application for analysis of Adverse Events, as described in Chen, et al., (2023) <doi:10.3390 cancers15092521="">. The required data for the application includes demographics, follow up, adverse event, drug administration and optional tumor measurement data. The app can produce swimmers plots of adverse events, Kaplan-Meier plots and Cox Proportional Hazards model results for the association of adverse event biomarkers and overall survival and progression free survival The adverse event biomarkers include occurrence of grade 3, low grade (1-2), and treatment related adverse events. Plots and tables of results are downloadable.</doi:10.3390>
License GPL-3
Encoding UTF-8
VignetteBuilder knitr
Suggests knitr, rmarkdown
RoxygenNote 7.3.2
Imports shiny, shinycssloaders, shinyjs, tidyverse, rio, janitor, DT, skimr, ggpubr, ggnewscale, survival, survminer, ggplot2, shinythemes, ggrepel, lubridate
NeedsCompilation no
Repository CRAN
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Description

Convert RECIST response code

Usage

Arguments

data A dataframe object that includes the column of interest to convert.

response_col A string. The column name of the column to convert.

new_col A string. The new column name of the converted response column. Defaults to

original response column name.

key A named vector key on how to convert the response IDs.

Value

Dataframe.

Examples

```
df <- data.frame(SampleID = c("A","B","C","D","E","F"),
Response = c("1","2","3","4","5","6"))
resp_code_transform(df,"Response", new_col = "Response_New")</pre>
```

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runAEapp

Run Adverse Events App

Description

Run Adverse Events App

Usage

```
runAEapp(launch.browser = TRUE)
```

Arguments

launch.browser Boolean logic to launch application in browser. Default to TRUE.

Value

An R Shiny Application.

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