Package 'helpeRs'

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Title Helper functions for high-quality presentation of quantitative research
Version 0.0
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Description Helper functions for high-quality presentation of quantitative research.
Depends R (>= $3.3.3$)
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Usage
<pre>getTableEntry()</pre>

2 fixZeroEndings

Arguments

x Description

Details

getTableEntry implements...

Value

z Description

Examples

```
# Perform analysis
TableEntry <- getTableEntry()
print( TableEntry )</pre>
```

fixZeroEndings

Implements...

Description

Implements...

Usage

```
fixZeroEndings(...)
```

Arguments

Х

Description

Details

fixZeroEndings implements...

Value

z Description

Examples

```
# Perform analysis
fixZeroEndings <- getTableEntry()
print( fixZeroEndings )</pre>
```

FullTransformer 3

FullTransformer

Implements...

Description

Implements...

Usage

```
FullTransformer(...)
```

Arguments

Х

Description

Details

FullTransformer implements...

Value

z Description

Examples

```
# Perform analysis
TableEntry <- FullTransformer()
print( TableEntry )</pre>
```

 ${\tt GetTableEntry}$

Implements...

Description

Implements...

Usage

```
\texttt{getTableEntry}(\dots)
```

Arguments

Χ

Description

Details

getTableEntry implements...

4 Stargazer2FullTable

Value

z Description

Examples

```
# Perform analysis
TableEntry <- getTableEntry()
print( TableEntry )</pre>
```

Stargazer2FullTable

Implements...

Description

Implements...

Usage

```
Stargazer2FullTable(...)
```

Arguments

Χ

Description

Details

Tables2Tex implements...

Value

z Description

Examples

```
# Perform analysis
Stargazer2FullTable()
print( TableEntries2Tex )
```

Tables2Tex 5

Tables2Tex

Implements...

Description

Implements...

Usage

```
Tables2Tex(...)
```

Arguments

Χ

Description

Details

Tables2Tex implements...

Value

z Description

Examples

```
# Perform analysis
Tables2Tex()
print( TableEntries2Tex )
```

vcovCluster

Implements...

Description

```
Implements...
```

Implements...

Usage

```
vcovCluster(...)
vcovCluster(...)
```

Arguments

Х

Description

WidenMargins

Details

```
vcovCluster implements... vcovCluster implements...
```

Value

- z Description
- z Description

Examples

```
# Perform analysis
TableEntry <- vcovCluster()
print( TableEntry )

# Perform analysis
TableEntry <- vcovCluster()
print( TableEntry )</pre>
```

WidenMargins

Implements...

Description

Implements...

Usage

```
getTableEntry(...)
```

Arguments

x

Description

Details

getTableEntry implements...

Value

z Description

Examples

```
# Perform analysis
TableEntry <- getTableEntry()
print( TableEntry )</pre>
```

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