

Package: socialrisk (via r-universe)

September 13, 2024

Type Package

Title Identifying Patient Social Risk from Administrative Health Care Data

Version 0.5.1

Description Social risks are increasingly becoming a critical component of health care research. One of the most common ways to identify social needs is by using ICD-10-CM ``Z-codes." This package identifies social risks using varying taxonomies of ICD-10-CM Z-codes from administrative health care data. The conceptual taxonomies come from: Centers for Medicare and Medicaid Services (2021) [<https://www.cms.gov/files/document/zcodes-infographic.pdf>](https://www.cms.gov/files/document/zcodes-infographic.pdf), Reidhead (2018) [<https://web.mhanet.com/>](https://web.mhanet.com/), A Arons, S DeSilvey, C Fichtenberg, L Gottlieb (2018) [<https://sirennetwork.ucsf.edu/tools-resources/resources/compendium-medical-terminology-codes-social-risk-factors>](https://sirennetwork.ucsf.edu/tools-resources/resources/compendium-medical-terminology-codes-social-risk-factors).

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Encoding UTF-8

LazyData true

RoxygenNote 7.1.2

Imports dplyr, magrittr, stringr, rlang, tidyselect, tidyr,

URL <https://github.com/WYATTBENSKEN/multimorbidity>

BugReports <https://github.com/WYATTBENSKEN/multimorbidity/issues>

Suggests rmarkdown, knitr

VignetteBuilder knitr

Depends R (>= 3.5.0)

NeedsCompilation no

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Contents

clean_data	2
i10_wide	3
socialrisk	3

Index	5
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clean_data	<i>Prepare our administrative data for analysis</i>
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Description

clean_data returns a dataset which has been transformed and cleaned for subsequent functions in this package.

Usage

```
clean_data(dat = NULL, style = "long", id = NULL, prefix_dx = "dx")
```

Arguments

dat	dataset
style	long, the default, is one diagnosis column per row whereas wide is multiple diagnosis columns
id	unique patient identifier variable name
prefix_dx	the variable prefix for the diagnosis columns (defaults to "dx"), in quotes

Details

This function takes our raw administrative data, in a number of different forms, and prepares it in a way which allows the other functions in this package to easily work with it. It is recommended to run this package on all data regardless of setup.

Value

dataframe with multiple rows per patient, which has re-structured their administrative data

Examples

```
clean_data(dat = i10_wide, id = patient_id, style = "wide", prefix_dx = "dx")
```

`i10_wide`*Example administrative data.*

Description

A dataset with fake patient data for 5 patients with ICD-10 diagnosis codes.

Usage

```
data(i10_wide)
```

Format

A data frame with 29 rows and 11 variables:

patient_id patient_id

sex patient's sex (male or female)

date_of_serv the date of service for the fake claim

dx1 first diagnosis

dx2 second diagnosis

dx3 third diagnosis

dx4 fourth diagnosis

dx5 fifth diagnosis

visit_type inpatient (ip) or outpatient(ot)

hcpcs HCPCS code

icd_version Which version of ICD the row is. 9 = ICD-9, 0 = ICD-10

Source

This was created by the package author.

`socialrisk`*Social Risk*

Description

`socialrisk` returns a summary dataset containing indicators of social risk, which vary based on the taxonomy command, for each patient.

Usage

```
socialrisk(dat = NULL, id = NULL, dx = "dx", taxonomy = "cms")
```

Arguments

<code>dat</code>	dataset which has been properly prepared in long format
<code>id</code>	variable of the unique patient identifier
<code>dx</code>	the column with the diagnoses (defaults to 'dx')
<code>taxonomy</code>	the taxonomy one wishes to use for social risk, with options of "cms" (default), "mha", and "siren"

Details

This function uses data which has been properly prepared to identify and flag social risks.

Value

dataframe with one row per patient, a column for their patient id, a column with whether they have any social risk, a column with the number of social risk domains, and columns with indicator variables for each social risk

Examples

```
data <- clean_data(dat = i10_wide, id = patient_id, style = "wide", prefix_dx = "dx")
socialrisk(dat = data, id = patient_id, dx = dx, taxonomy = "cms")
```

Index

* **datasets**
 i10_wide, 3

clean_data, 2

i10_wide, 3

socialrisk, 3