

Package ‘medicalcoder’

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Title Tools for Working with ICD Codes and Comorbidity Algorithms

Version 0.6.0

Description Provides tools for working with medical coding schemas such as the International Classification of Diseases (ICD). Includes functions for comorbidity classification algorithms such as the Pediatric Complex Chronic Conditions (PCCC), Charlson, and Elixhauser indices.

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Contents

comorbidities	2
get_charlson_codes	7
get_charlson_index_scores	8
get_elixhauser_codes	9
get_elixhauser_index_scores	10

get_elixhauser_poa 10

get_icd_codes 11

get_pccc_codes 14

get_pccc_conditions 15

icd_compact_to_full 15

is_icd 16

lookup_icd_codes 19

mdcr 20

mdcr_longitudinal 21

summary.medicalcoder_comorbidities 22

summary.medicalcoder_comorbidities_with_subconditions 23

Index 25

comorbidities	<i>Comorbidities</i>
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Description

Apply established comorbidity algorithms to ICD-coded data. Supported methods include several variants of the Charlson comorbidity system, Elixhauser, and the Pediatric Complex Chronic Conditions (PCCC).

Usage

```
comorbidities(  
  data,  
  icd.codes,  
  method,  
  id.vars = NULL,  
  icdv.var = NULL,  
  icdv = NULL,  
  dx.var = NULL,  
  dx = NULL,  
  poa.var = NULL,  
  poa = NULL,  
  age.var = NULL,  
  primarydx.var = NULL,  
  primarydx = NULL,  
  flag.method = c("current", "cumulative"),  
  full.codes = TRUE,  
  compact.codes = TRUE,  
  subconditions = FALSE  
)
```

Arguments

<code>data</code>	A <code>data.frame</code> in a "long" format. The input <code>data.frame</code> is expected to have one column of ICD codes (one code per row) with additional (optional) columns for patient/encounter ids, ICD version, diagnostic/procedure status, present-on-admission flags, primary diagnostic flags, or age.
<code>icd.codes</code>	Character scalar naming the column in <code>data</code> that contains ICD codes. The codes can be full (with applicable dots, e.g., C84.2), compact (dots omitted, e.g., C842), or a mix of both full and compact codes. Use
<code>method</code>	Character string indicating the comorbidity algorithm to apply to <code>data</code> .
<code>id.vars</code>	Optional character vector of column names. When missing, the entire input <code>data</code> is treated as a single encounter from a single patient. If you want to set <code>flag.method = "current"</code> then <code>length(id.vars) >= 2</code> is expected. The last element would be the encounter order (must be sortable)."
<code>icdv.var</code>	Character scalar naming the column in <code>data</code> that indicates the ICD version (9 or 10). If present it must be integer values 9 or 10. <code>icdv.var</code> takes precedence over <code>icdv</code> if both are provided.
<code>icdv</code>	An integer value of 9L or 10L indicating that all <code>data[[icd.codes]]</code> are ICD version 9 or 10, respectively. Ignored (with a warning) if <code>icdv.var</code> is provided.
<code>dx.var</code>	Character scalar naming the column in <code>data</code> that indicates diagnostic (1) vs procedural (0) codes. If present it must be integer values 0 or 1. <code>dx.var</code> takes precedence over <code>dx</code> if both are provided.
<code>dx</code>	An integer indicating that all <code>data[[icd.codes]]</code> are diagnostic (1) or procedure (0) codes. Ignored (with a warning) if <code>dx.var</code> is provided.
<code>poa.var</code>	Character scalar naming the column with present-on-admission flags: integer 1L (present), 0L (not present), or NA. PCCC and Charlson will only flag conditions when the code is present-on-admission. Elixhauser has a mix of conditions some require present-on-admission others do not. <code>poa.var</code> takes precedence over <code>poa</code> if both are provided.
<code>poa</code>	Integer scalar 0 or 1. Use when all <code>icd.codes</code> share the same present-on-admission status. Ignored with a warning if <code>poa</code> and <code>poa.var</code> are both provided.
<code>age.var</code>	Character scalar naming the column in <code>data</code> that contains patient age in years. Only applicable to Charlson comorbidities.
<code>primarydx.var</code>	Character scalar naming the column in <code>data</code> that indicates whether <code>data[[icd.codes]]</code> are primary diagnostic codes (1L) or not (0L). Primary diagnosis is used only for Elixhauser comorbidities and is ignored when the method is PCCC or Charlson. <code>primarydx.var</code> takes precedence over <code>primarydx</code> if both are provided.
<code>primarydx</code>	An integer value of 0 or 1. If 0, treat all codes as non-primary diagnoses; if 1, treat all codes as primary diagnoses. Ignored, with a warning, if <code>primarydx.var</code> is provided.
<code>flag.method</code>	When <code>flag.method = 'current'</code> (default) only codes associated with the current <code>id.vars</code> are considered when flagging comorbidities. When <code>flag.method = 'cumulative'</code> then all prior encounters are considered when flagging comorbidities. See Details .

`full.codes, compact.codes`

Logical; when TRUE compare `data[[icd.codes]]` against full and/or compact ICD codes in the method's lookup tables. Full ICD codes include a decimal point (when applicable) and compact codes omit the decimal point. For example: B95.0 is the full ICD-10-CM diagnostic code for "Streptococcus, group A, as the cause of disease classified elsewhere," whereas B950 is the associated compact code.

`subconditions` Logical scalar; when TRUE, report both conditions and subconditions (PCCC only).

Details

When `flag.method = "current"`, only codes from the index encounter contribute to flags. When a longitudinal method is selected (e.g., "cumulative"), prior encounters for the same `id.vars` combination may contribute to condition flags. For the cumulative method to work the `id.vars` need to be a character vector length 2 or more. The last variable listed in the `id.vars` will be considered the encounter id and should be sortable. For example, say you have data with a hospital, patient, and encounter id. The `id.vars` could be one of two entries: `c("hospital", "patient", "encounter")` or `c("patient", "hospital", "encounter")`. In both cases the return will be the same as "encounter" within the hospital/patient id interaction is the same as "encounter" within patient/hospital interaction.

It is critically important that the `data[[tail(id.vars, 1)]]` variable can be sorted. Just because your data is sorted in temporal order does not mean that the results will be correct if the `tail(id.vars, 1)` is not in the same order as the data. For example, say you had the following:

patid	enc_id	date
P1	10823090	Aug 2023
P1	10725138	Jul 2025

`id.vars = c("patid", "enc_id")` will give the wrong result as `enc_id 10725138` would be sorted to come before `enc_id 10823090`. `id.var = c("patid", "date")` would be sufficient input, assuming that date has been correctly stored. Adding a column `enc_seq`, e.g.,

patid	enc_id	date	enc_seq
P1	10823090	Aug 2023	1
P1	10725138	Jul 2025	2

and calling `comorbidities()` with `id.vars = c("patid", "enc_seq")` will have better performance than using the date and will clear up any possible issues with non-sequential encounter ids from the source data.

Value

The return object will be slightly different depending on the value of `method` and `subconditions`.

- When `subconditions = FALSE`, a `medicalcoder_comorbidities` object (a `data.frame` with attributes) is returned. Column(s) for `id.vars`, if defined in the function call. For all method there will be the following columns:
 - `num_cmr` a count of comorbidities/conditions flagged
 - `cmrb_flag` a 0/1 integer indicator for at least one comorbidity/condition.

Additional columns:

- PCCC methods:
 - * For method = "pccc_v2.0" and method = "pccc_v2.1", there is one indicator column per condition.
 - * For method = "pccc_v3.0" and method = "pccc_v3.1", there are four columns per condition:
 - `<condition>_dxpr_or_tech`: the condition was flag due to the presence of either a diagnostic or procedure code, or was flag due to the presence of a technology dependence code along with at least one comorbidity being flagged by a diagnostic or procedure code.
 - `<condition>_dxpr_only`: the condition was flagged due to the presence of a non-technology dependent diagnostic or procedure code only.
 - `<condition>_tech_only`: the condition was flagged due to the presence of a technology dependent code only and at least one other comorbidity was flagged by a non-technology dependent code.
 - `<condition>_dxpr_and_tech`: The patient had both diagnostic or procedure codes and a technology dependence code for the condition.
- For Charlson variants, indicator columns are returned for the relevant conditions, `cci` (Charlson Comorbidity Index), and `age_score`.
- For Elixhauser variants, indicator columns are returned for all relevant comorbidities, mortality, and readmission indices.
- When `subconditions = TRUE` and the method is a PCCC variant, a list of length two is returned: the first element contains condition indicators; the second element is a named list of `data.frames` with indicators for subconditions within each condition.

References

- Pediatric Complex Chronic Conditions:
 - Feudtner, C., Feinstein, J.A., Zhong, W. et al. Pediatric complex chronic conditions classification system version 2: updated for ICD-10 and complex medical technology dependence and transplantation. *BMC Pediatr* 14, 199 (2014). <https://doi.org/10.1186/1471-2431-14-199>
 - Feinstein JA, Hall M, Davidson A, Feudtner C. Pediatric Complex Chronic Condition System Version 3. *JAMA Netw Open*. 2024;7(7):e2420579. <https://doi.org/10.1001/jamanetworkopen.2024.20579>
- Charlson Comorbidities:
 - Mary E. Charlson, Peter Pompei, Kathy L. Ales, C.Ronald MacKenzie, A new method of classifying prognostic comorbidity in longitudinal studies: Development and validation, *Journal of Chronic Diseases*, Volume 40, Issue 5, 1987, Pages 373-383, ISSN 0021-9681, [https://doi.org/10.1016/0021-9681\(87\)90171-8](https://doi.org/10.1016/0021-9681(87)90171-8).

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- Quan H, Li B, Couris CM, Fushimi K, Graham P, Hider P, Januel JM, Sundararajan V. Updating and validating the Charlson comorbidity index and score for risk adjustment in hospital discharge abstracts using data from 6 countries. *Am J Epidemiol*. 2011 Mar 15;173(6):676-82. <https://doi.org/10.1093/aje/kwq433>. Epub 2011 Feb 17. PMID: 21330339.
- Glasheen WP, Cordier T, Gumpina R, Haugh G, Davis J, Renda A. Charlson Comorbidity Index: ICD-9 Update and ICD-10 Translation. *Am Health Drug Benefits*. 2019 Jun-Jul;12(4):188-197. PMID: 31428236; PMCID: PMC6684052.
- Elixhauser Comorbidities:
 - Agency for Healthcare Research and Quality (AHRQ). Elixhauser Comorbidity Software Refined for ICD-10-CM Diagnoses, v2025.1 [Internet]. 2025. Available from: https://www.hcup-us.ahrq.gov/toolssoftware/comorbidityicd10/comorbidity_icd10.jsp

See Also

- `vignettes(topic = "comorbidities", package = "medicalcoder")`
- `vignettes(topic = "pccc", package = "medicalcoder")`
- `vignettes(topic = "charlson", package = "medicalcoder")`
- `vignettes(topic = "elixhauser", package = "medicalcoder")`

Examples

```
pccc_v3.1_results <-
  comorbidities(data = mdcr,
    icd.codes = "code",
    id.vars = "patid",
    dx.var = "dx",
    method = "pccc_v3.1",
    flag.method = 'current',
    poa = 1)
summary(pccc_v3.1_results)

pccc_v3.1_subcondition_results <-
  comorbidities(data = mdcr,
    icd.codes = "code",
    id.vars = "patid",
    dx.var = "dx",
    method = "pccc_v3.1",
    flag.method = 'current',
    poa = 1,
    subconditions = TRUE)
summary(pccc_v3.1_subcondition_results)
```

```

charlson_results <-
  comorbidities(data = mdcr,
                icd.codes = "code",
                id.vars = "patid",
                dx.var = "dx",
                method = "charlson_quan2011",
                flag.method = 'current',
                poa = 1)
summary(charlson_results)

elixhauser_results <-
  comorbidities(data = mdcr,
                icd.codes = "code",
                id.vars = "patid",
                dx.var = "dx",
                method = "elixhauser_ahrq2025",
                primarydx = 1,
                flag.method = 'current',
                poa = 1)
summary(elixhauser_results)

```

get_charlson_codes	<i>Get Charlson Codes</i>
--------------------	---------------------------

Description

Retrieve a copy of internal lookup tables for the ICD codes used in assessing Charlson comorbidities.

Usage

```
get_charlson_codes()
```

Value

A data.frame with the following columns:

- `icdv`: Integer vector indicating if the code is from ICD-9 or ICD-10
- `dx`: Integer vector. 1 if the code is a diagnostic, (ICD-9-CM, ICD-10-CM, WHO, CDC Mortality), or 0 if the code is procedural (ICD-9-PCS, ICD-10-PCS)
- `full_code`: Character vector with the ICD code and any relevant decimal point
- `code`: Character vector with the compact ICD code
- `condition`: Character vector of the conditions
- `charlson_<variant>`: Integer vector indicating if the code is part of the <variant> of the Charlson comorbidities.

See Also

- [get_charlson_index_scores\(\)](#) for a lookup table of the by comorbidity index scores.
- [get_icd_codes\(\)](#) for the lookup table of all ICD codes.
- [get_pccc_codes\(\)](#) for the lookup table of ICD codes used for the PCCC.
- [get_elixhauser_codes\(\)](#) for the lookup table of ICD codes used for the Elixhauser comorbidities.
- [comorbidities\(\)](#) for applying comorbidity algorithms to a data set.

Examples

```
head(get_charlson_codes())  
str(get_charlson_codes())
```

```
get_charlson_index_scores  
  Get Charlson Index Scores
```

Description

Retrieve a copy of internal lookup tables of index scores used in assessing Charlson comorbidities.

Usage

```
get_charlson_index_scores()
```

Value

A data.frame with the following columns:

- condition: Character vector of the conditions
- index: Character vector indicating if the score is for the mortality or the readmission index score
- charlson_<variant>: the index scores for the variant

See Also

- [get_charlson_codes\(\)](#) for a lookup table of the ICD codes mapping to the Charlson comorbidities.
- [comorbidities\(\)](#) for applying comorbidity algorithms to a data set.

Examples

```
head(get_charlson_index_scores())  
str(get_charlson_index_scores())
```

`get_elixhauser_codes` *Get Elixhauser Codes*

Description

Retrieve copy of internal lookup tables for the ICD codes used in assessing Elixhauser comorbidities.

Usage

```
get_elixhauser_codes()
```

Value

A data.frame with the following columns:

- `icdv`: Integer vector indicating if the code is from ICD-9 or ICD-10
- `dx`: Integer vector. 1 if the code is a diagnostic, (ICD-9-CM, ICD-10-CM, WHO, CDC Mortality), or 0 if the code is procedural (ICD-9-PCS, ICD-10-PCS)
- `full_code`: Character vector with the ICD code and any relevant decimal point
- `code`: Character vector with the compact ICD code omitting any relevant decimal point
- `condition`: Character vector of the conditions
- `elixhauser_<variant>`: Integer vector indicating if the code is part of the <variant> of the Elixhauser comorbidities.

See Also

- [get_elixhauser_index_scores\(\)](#) for the lookup table of the condition by condition scores for mortality and readmission indices.
- [get_elixhauser_poa\(\)](#) for the lookup table of the conditions which do not require associated ICD codes to be present-on-admission to flag the comorbidity.
- [get_icd_codes\(\)](#) for the lookup table of all ICD codes.
- [get_pccc_codes\(\)](#) for the lookup table of ICD codes used for the PCCC.
- [get_charlson_codes\(\)](#) for the lookup table of ICD codes used for the Charlson comorbidities.
- [comorbidities\(\)](#) for applying comorbidity algorithms to a data set.

Examples

```
head(get_elixhauser_codes())  
str(get_elixhauser_codes())
```

get_elixhauser_index_scores

Get Elixhauser Index Scores

Description

Functions to get a copy of internal lookup tables for the ICD codes and index scores used in assessing Elixhauser comorbidities.

Usage

```
get_elixhauser_index_scores()
```

Value

A data.frame with the following columns:

- condition: Character vector of the conditions
- index: Character vector indicating if the score is for the mortality or the readmission index score
- elixhauser_<variant>: integer vector of the scores

See Also

- [get_elixhauser_codes\(\)](#) for the lookup table of ICD codes mapping to the Elixhauser comorbidities.
- [get_elixhauser_poa\(\)](#) for the lookup table of the conditions which do and do not require associated ICD codes to be present-on-admission to flag the comorbidity.
- [comorbidities\(\)](#) for applying comorbidity algorithms to a data set.

Examples

```
head(get_elixhauser_index_scores())
str(get_elixhauser_index_scores())
```

get_elixhauser_poa

Get Elixhauser Present-on-Admission Requirements

Description

Retrieve a copy of internal lookup table with details on which Elixhauser comorbidities do and do not require the associated ICD codes to be present-on-admission to be flagged.

Usage

```
get_elixhauser_poa()
```

Value

A data.frame with the following columns:

- condition: Character vector of the conditions
- desc: Character vector with a verbose description of the condition
- poa_required: Integer indicators if the code needs to present on admission to be considered a comorbidity
- elixhauser_<variant>: indicators for the Elixhauser <variant>

See Also

- [get_elixhauser_index_scores\(\)](#) for the lookup table of the condition by condition scores for mortality and readmission indices.
- [get_elixhauser_codes\(\)](#) for the lookup table of ICD codes mapping to the Elixhauser comorbidities.
- [comorbidities\(\)](#) for applying comorbidity algorithms to a data set.

Examples

```
head(get_elixhauser_poa())
str(get_elixhauser_poa())
```

get_icd_codes

Get ICD Codes

Description

Retrieve a copy of the internal look up table for all known ICD codes.

Usage

```
get_icd_codes(with.descriptions = FALSE, with.hierarchy = FALSE)
```

Arguments

with.descriptions

Logical scalar, if TRUE include the description of the codes.

with.hierarchy Logical scalar, if TRUE include the ICD hierarchy.

Details

Sources:

There are three sources of ICD codes.

- cms: Codes from the ICD-9-CM, ICD-9-PCS, ICD-10-CM, and ICD-10-PCS standards.
- who: Codes from World Health Organization.
- cdc: Codes from CDC Mortality coding standard.

Fiscal and Calendar Years:

When reporting years there is a mix of fiscal and calendar years.

Fiscal years are the United States Federal Government fiscal years, running from October 1 to September 30. For example, fiscal year 2013 started October 1 2012 and ended on September 30 2013.

Calendar years run January 1 to December 31.

Within the ICD data there are columns `known_start`, `known_end`, `assignable_start`, `assignable_end`, `desc_start` and `desc_end`. For ICD codes with `src == "cms"`, these are fiscal years. For codes with `src == "cdc"` or `src == "who"` these are calendar years.

`known_start` is the first fiscal or calendar year (depending on source) that the `medicalcoder` package as definitive source data for. ICD-9-CM started in the United States in fiscal year 1980. Source information that could be downloaded from the CDC and CMS and added to the source code for the `medicalcoder` package goes back to 1997. As such 1997 is the "known start"

`known_end` is the last fiscal or calendar year (depending on source) for which we have definitive source data for. For ICD-9-CM and ICD-9-PCS that is 2015. For ICD-10-CM and ICD-10-PCS, which are active, it is just the last year of known data. ICD-10 from the WHO ends in 2019.

Header and Assignable Codes:

"Assignable" indicates that the code is the most granular for the source. Ideally codes are reported with the greatest level of detail but that is not always the case. Also, the greatest level of detail can differ between sources. Example: C86 is a header code for cms and who because codes C86.0, C86.1, C86.2, C86.3, C86.4, C86.5, and C86.6 all exist in both standards. No code with a fifth digit exists in the who so all these four digit codes are 'assignable.' In the cms standard, C86.0 was assignable through fiscal year 2024. In fiscal year 2025 codes C86.00 and C86.01 were added making C86.0 a header code and C86.00 and C86.01 assignable codes.

Value

a `data.frame`

The default return has the following columns:

- `icdv`: Integer vector indicating if the code is from ICD-9 or ICD-10
- `dx`: Integer vector. 1 if the code is a diagnostic, (ICD-9-CM, ICD-10-CM, WHO, CDC Mortality), or 0 if the code is procedural (ICD-9-PCS, ICD-10-PCS)
- `full_code`: Character vector with the ICD code and any relevant decimal point
- `code`: Character vector with the compact ICD code omitting any relevant decimal point
- `src`: Character vector reporting the source of the information. See Details.
- `known_start`: Integer vector reporting the first known year of use. See Details.

- `known_end`: Integer vector reporting the last known year of use. See Details.
- `assignable_start`: Integer vector reporting the first known year the code was assignable. See Details.
- `assignable_end`: Integer vector reporting the last known year the code was assignable. See Details.

When `with.descriptions = TRUE` there are the following additional columns:

- `desc`: Character vector of descriptions. For `cms` codes descriptions from CMS are used preferentially over CDC.
- `desc_start`: Integer vector of the first year the description was used.
- `desc_end`: Integer vector of the last year the description was used.

When `with.hierarchy = TRUE` there are the following additional columns:

- `chapter`
- `subchapter`
- `category`
- `subcategory`
- `subclassification`
- `subsubclassification`
- `extension`

See Also

[is_icd\(\)](#), [lookup_icd_codes\(\)](#), [vignette\(topic = "icd", package = "medicalcoder"\)](#)

Examples

```
icd_codes <- get_icd_codes()
str(icd_codes)

# Explore the change in the assignable year for C86 code between CMS and
# WHO
subset(get_icd_codes(), grepl("^C86$", full_code))
subset(get_icd_codes(), grepl("^C86\\.\\.\\d$", full_code))
subset(get_icd_codes(), grepl("^C86\\.\\.0(\\d|$)", full_code))

is_icd("C86", headerok = FALSE) # FALSE
is_icd("C86", headerok = TRUE)  # TRUE
is_icd("C86", headerok = TRUE, src = "cdc") # Not a CDC mortality code

lookup_icd_codes("^C86\\.\\.0\\d*", regex = TRUE)
```

get_pccc_codes

*Pediatric Complex Chronic Conditions ICD Codes***Description**

Retrieve a copy of internal lookup tables for the ICD codes mapping to the Pediatric Complex Chronic Conditions (PCCC) conditions and subconditions by variant.

Usage

```
get_pccc_codes()
```

Value

a `data.frame` with the following columns

- `icdv`: Integer vector indicating if the code is from ICD-9 or ICD-10.
- `dx`: Integer vector. 1 if the code is a diagnostic, (ICD-9-CM, ICD-10-CM, WHO, CDC Mortality), or 0 if the code is procedural (ICD-9-PCS, ICD-10-PCS).
- `full_code`: Character vector with the ICD code and any relevant decimal point.
- `code`: Character vector with the compact ICD code omitting any relevant decimal point.
- `condition`: Character vector of the conditions.
- `subcondition`: Character vector of the subconditions.
- `transplant_flag`: Integer vector indicating if the code is associated with a transplant.
- `tech_dep_flag`: Integer vector indicating if the code is associated with technology dependence.
- `pccc_<variant>`: Integer vector indicating if the code is part of the v2.0, v2.1, v3.0, or v3.1 variant.

See Also

- [get_pccc_conditions\(\)](#) for a reference of the PCCC conditions and subconditions.
- [get_icd_codes\(\)](#) for the lookup table of all ICD codes.
- [comorbidities\(\)](#) for applying comorbidity algorithms to a data set.

Examples

```
head(get_pccc_codes())
str(get_pccc_codes())
```

get_pccc_conditions	<i>Pediatric Complex Chronic Condition and Subconditions</i>
---------------------	--

Description

Retrieve a copy of internal lookup tables for the syntax valid and human readable labels of the Pediatric Complex Chronic Conditions (PCCC) conditions and subconditions.

Usage

```
get_pccc_conditions()
```

Value

a `data.frame` with the following columns

- `condition`: (character) syntax valid name for the condition
- `subconditions`: (character) syntax valid name for the subcondition
- `conditions_label`: (character) human readable label for the condition
- `subconditions_label`: (character) human readable label for the subcondition

See Also

- [get_pccc_codes\(\)](#) for the lookup table of ICD codes used for the PCCC.
- [comorbidities\(\)](#) for applying comorbidity algorithms to a data set.

Examples

```
get_pccc_conditions()
```

icd_compact_to_full	<i>Convert ICD Compact Codes to Full Codes</i>
---------------------	--

Description

Take an assumed ICD compact code string and convert to a full code based on the ICD version (9 or 10) and type (diagnostic or procedure). This method only formats strings and does not validate the code(s).

Usage

```
icd_compact_to_full(x, icdv, dx)
```

Arguments

x	Character vector
icdv	Integer vector of allowed ICD versions. Use 9L and/or 10L. Defaults to both.
dx	Integer vector indicating allowed code type(s): 1L for diagnostic (ICD-9-CM, ICD-10-CM, CDC mortality, WHO), 0L for procedural (ICD-9-PCS, ICD-10-PCS). Defaults to both.

Value

A character vector the same length as x.

See Also

- [get_icd_codes\(\)](#) to retrieve the internal lookup table of ICD codes.
- [lookup_icd_codes\(\)](#) for retrieving details on a specific set of ICD codes.
- [is_icd\(\)](#) to test if a string is a known ICD code.

Other ICD tools: [is_icd\(\)](#), [lookup_icd_codes\(\)](#)

is_icd	<i>Is ICD</i>
--------	---------------

Description

Answer the question "is the character string x a valid ICD code?" ICD codes should be character vectors. `is_icd` will assess for both "full codes" (decimal point present when appropriate) and "compact codes" (decimal point omitted).

ICD-10 code "C00" is a header code because the four-character codes C00.0, C00.1, C00.2, C00.3, C00.4, C00.5, C00.6, C00.7, C00.8, and C00.9 exist. Those four-character codes are assignable (as of 2025) because no five-character descendants (e.g., C00.40) exist.

When the source is the World Health Organization (WHO) or CDC Mortality, years refer to calendar years. CDC/CMS sources use the U.S. federal fiscal year, which starts on October 1 (e.g., fiscal year 2024 runs 2023-10-01 to 2024-09-30).

Usage

```
is_icd(  
  x,  
  icdv = c(9L, 10L),  
  dx = c(1L, 0L),  
  src = c("cms", "who", "cdc"),  
  year,  
  headerok = FALSE,  
  ever.assignable = missing(year),  
  warn.ambiguous = TRUE,  
  full.codes = TRUE,  
  compact.codes = TRUE  
)
```


Arguments

<code>x</code>	Character vector of ICD codes (full or compact form).
<code>icdv</code>	Integer vector of allowed ICD versions. Use 9L and/or 10L. Defaults to both.
<code>dx</code>	Integer vector indicating allowed code type(s): 1L for diagnostic (ICD-9-CM, ICD-10-CM, CDC mortality, WHO), 0L for procedural (ICD-9-PCS, ICD-10-PCS). Defaults to both.
<code>src</code>	Character vector of code sources. One or more of "cms", "who", "cdc". Defaults to all.
<code>year</code>	Numeric scalar. Calendar or fiscal year to reference. Default is the most current year available per source. For ICD-9, the latest year is 2015; ICD-10 source are updated annually. Calendar year for WHO and CDC mortality. Fiscal year for CMS.
<code>headerok</code>	Logical scalar. If FALSE (default), only assignable codes are considered valid; if TRUE, header codes are also accepted.
<code>ever.assignable</code>	Logical scalar. If TRUE then ignore year and return TRUE if the x was ever an assignable code.
<code>warn.ambiguous</code>	Logical scalar. If TRUE (default), warn when a code matches more than one ICD version and/or type (e.g., both CM and PCS).
<code>full.codes</code>	Logical scalar. If TRUE (default), match codes that include the decimal point where applicable.
<code>compact.codes</code>	Logical scalar. If TRUE (default), match codes without the decimal point.

Details

Similarly for ICD-9-CM: "055" is a header for measles; 055.0, 055.1, 055.2, 055.8, and 055.9 are assignable. Codes 055.3–055.6 do not exist. Code 055.7 is a header because 055.71 and 055.72 exist.

Some codes change status across years. For example, ICD-9-CM 516.3 was assignable in fiscal years 2006–2011, then became a header in 2012–2015.

Value

A logical vector the same length as `x`.

See Also

- [get_icd_codes\(\)](#) to retrieve the internal lookup table of ICD codes.
- [lookup_icd_codes\(\)](#) for retrieving details on a specific set of ICD codes.
- [icd_compact_to_full\(\)](#) converts a string from a compact format to the full format based on ICD version and type (diagnostic or procedure).

Other ICD tools: [icd_compact_to_full\(\)](#), [lookup_icd_codes\(\)](#)

Examples

```
#####
# Some ICD-9 diagnostic codes
x <- c("136.2", "718.60", "642.02")

is_icd(x, icdv = 9, dx = 1)
is_icd(x, icdv = 9, dx = 0)
is_icd(x, icdv = 10, dx = 1)
is_icd(x, icdv = 10, dx = 0)

is_icd(x, icdv = 9, dx = 1, headerok = TRUE)
is_icd(x, icdv = 9, dx = 1, year = 2006)

#####
# ICD code with, or without a dot. The ICD-9 diagnostic code 799.3 and ICD-9
# procedure code 79.93 both become 7993 when assessed against the ICD code look
# up tables. As such "7993" is a valid ICD-9 diagnostic and procedure code,
# whereas 799.3 is only a valid dx code, and 79.93 is only a valid pr code.
# Further, codes such as ".7993", "7.993", "7993.", are all non-valid codes.

x <- c("7993", ".7993", "7.993", "79.93", "799.3", "7993.")
data.frame(
  x,
  dx = is_icd(x, icdv = 9, dx = 1),
  pr = is_icd(x, icdv = 9, dx = 0)
)

#####
# example of a ICD-9 code that was assignable, but became a header when
# more descriptive codes were introduced: ICD-9 diagnostic code 516.3
lookup_icd_codes(paste0("516.3", c("", as.character(0:9))))

# ICD-9 code 516.3 was an assignable code through fiscal year 2011.
is_icd("516.3")

# If `year` is omitted, and `ever.assignable = FALSE` then the `year` is
# implied to be the max `known_end` year for ICD codes matched by `icdv`,
# `dx`, and `src`.
is_icd("516.3", ever.assignable = FALSE)

# when `year` is provided then `ever.assignable` is `FALSE` by default and
# the return is TRUE when 516.3 was assignable and FALSE otherwise.
is_icd("516.3", year = 2015)
is_icd("516.3", year = 2011)

# when year is a non-assignable year, but `ever.assignable = TRUE` the return
# will be TRUE. Useful if you know the data is retrospective and collected
# through fiscal year 2015.
is_icd("516.3", year = 2015, ever.assignable = TRUE)

#####
# Consider the string E010
```

```
# - This could be a ICD-9-CM full code
# - Could be a ICD-10-CM compact code
lookup_icd_codes("E010")
subset(get_icd_codes(with.descriptions = TRUE), grepl("^E010$", code))

is_icd("E010")
is_icd("E010", icdv = 9) # FALSE because it is a header code and was never assignable
is_icd("E010", icdv = 9, ever.assignable = TRUE) # FALSE
is_icd("E010", icdv = 9, headerok = TRUE) # TRUE
```

lookup_icd_codes	<i>Lookup ICD Codes</i>
------------------	-------------------------

Description

Functions for working with ICD codes.

ICD-10 code "C00" is a header code because the four-character codes C00.0, C00.1, C00.2, C00.3, C00.4, C00.5, C00.6, C00.7, C00.8, and C00.9 exist. Those four-character codes are assignable (as of 2025) because no five-character descendants (e.g., C00.40) exist.

When the source is the World Health Organization (WHO) or CDC Mortality, years refer to calendar years. CDC/CMS sources use the U.S. federal fiscal year, which starts on October 1 (e.g., fiscal year 2024 runs 2023-10-01 to 2024-09-30).

Usage

```
lookup_icd_codes(
  x,
  regex = FALSE,
  full.codes = TRUE,
  compact.codes = TRUE,
  ...
)
```

Arguments

x	Character vector of ICD codes (full or compact form).
regex	Logical scalar. If TRUE, treat x as regular expressions; if FALSE, use exact matching.
full.codes	Logical scalar. If TRUE (default), match codes that include the decimal point where applicable.
compact.codes	Logical scalar. If TRUE (default), match codes without the decimal point.
...	Passed to <code>base::grep()</code> when <code>regex = TRUE</code>

Details

ICD codes should be character vectors. These tools work with either "full codes" (decimal point present when appropriate) or "compact codes" (decimal point omitted).

Similarly for ICD-9-CM: "055" is a header for measles; 055.0, 055.1, 055.2, 055.8, and 055.9 are assignable. Codes 055.3–055.6 do not exist. Code 055.7 is a header because 055.71 and 055.72 exist.

Some codes change status across years. For example, ICD-9-CM 516.3 was assignable in fiscal years 2006–2011, then became a header in 2012–2015.

Value

A data frame with one or more rows per input, including columns

- `match_type`: did the input match a full or compact code
- `icdv`: icd version (9 or 10)
- `dx`: diagnostic code (1) or procedure code (0)
- `full_code`: the full code string
- `code`: the compact codes string
- `src`: the source - CMS, CDC, or WHO.
- year ranges (`known_*`, `assignable_*`).

See Also

- [get_icd_codes\(\)](#) to retrieve the internal lookup table of ICD codes.
- [is_icd\(\)](#) to test if a string is a known ICD code.
- [icd_compact_to_full\(\)](#) converts a string from a compact format to the full format based on ICD version and type (diagnostic or procedure).

Other ICD tools: [icd_compact_to_full\(\)](#), [is_icd\(\)](#)

mdcr

Synthetic Data

Description

Synthetic Data

Usage

mdcr

Format

mdcr is a `data.frame` with 4 columns, one for a patient id and 41 for diagnostic codes and 41 possible procedure codes. Each row is for one patient id.

- patid: patient identifier, integer values
- icdv: ICD version; integer values, 9 or 10
- dx: indicator column for ICD diagnostic (1) or procedure (0) codes
- code: ICD code; character values

See Also

Other datasets: [mdcr_longitudinal](#)

mdcr_longitudinal	<i>Synthetic Longitudinal Data</i>
-------------------	------------------------------------

Description

Synthetic Longitudinal Data

Usage

```
mdcr_longitudinal
```

Format

mdcr_longitudinal is a `data.frame` with four columns. The codes are expected to be treated as diagnostic codes but there are a few ICD-9 codes which could match to procedure codes as well.

- patid: patient identifier, integer values
- date: date the diagnostic code was recorded
- icdv: ICD version 9 or 10, integer valued
- code: ICD codes; character values

See Also

Other datasets: [mdcr](#)

summary.medicalcoder_comorbidities
Summaries of Comorbidities

Description

Build summaries (counts and percentages) for each comorbidity and other summary statistics by method.

Usage

```
## S3 method for class 'medicalcoder_comorbidities'  
summary(object, ...)
```

Arguments

object	a medicalcoder_comorbidities object generated by calling comorbidities()
...	additional parameters, not currently used

Value

either a list or a data data.frame

Examples

```
pccc_v3.1_results <-  
  comorbidities(data = mdcr,  
                icd.codes = "code",  
                id.vars = "patid",  
                dx.var = "dx",  
                method = "pccc_v3.1",  
                flag.method = 'current',  
                poa = 1)  
summary(pccc_v3.1_results)  
  
charlson_results <-  
  comorbidities(data = mdcr,  
                icd.codes = "code",  
                id.vars = "patid",  
                dx.var = "dx",  
                method = "charlson_quan2011",  
                flag.method = 'current',  
                poa = 1)  
summary(charlson_results)  
  
elixhauser_results <-  
  comorbidities(data = mdcr,  
                icd.codes = "code",  
                id.vars = "patid",
```

```

        dx.var = "dx",
        method = "elixhauser_ahrq2025",
        primarydx = 1,
        flag.method = 'current',
        poa = 1)
summary(elixhauser_results)

```

```
summary.medicalcoder_comorbidities_with_subconditions
```

Summaries of Comorbidities with Subconditions

Description

Build summaries (counts and percentages) for each Pediatric Complex Chronic Condition (PCCC) condition and subcondition.

Usage

```
## S3 method for class 'medicalcoder_comorbidities_with_subconditions'
summary(object, ...)
```

Arguments

object	a medicalcoder_comorbidities_with_subcondtions object generated by calling <code>comorbidities()</code> with <code>subconditions = TRUE</code> . This is currently only applicable to PCCC.
...	additional parameters, not currently used

Value

a data.frame with five columns.

1. condition the primary condition
2. subcondition the subcondition(s) within the condition. There will be a row where subcondition is NA which is used to report the count and percent_of_cohort for the condition overall.
3. count the number of rows in object with the applicable condition and subcondition.
4. percent_of_cohort: a numeric value within [0, 100] for the percent of rows in object with the flagged condition and subcondition.
5. percent_of_those_with_condition: a numeric value within [0, 100] for the subset of rows in object with the primary condition and the flagged subcondition. Will be NA for the primary condition.

See Also

`comorbidities()`, `vignette(topic = "pccc", package = "medicalcoder")`

Examples

```
pccc_v3.1_subcondition_results <-  
  comorbidities(data = mdcr,  
    icd.codes = "code",  
    id.vars = "patid",  
    dx.var = "dx",  
    method = "pccc_v3.1",  
    flag.method = 'current',  
    poa = 1,  
    subconditions = TRUE)  
summary(pccc_v3.1_subcondition_results)
```


Index

* ICD tools

icd_compact_to_full, [15](#)

is_icd, [16](#)

lookup_icd_codes, [19](#)

* datasets

mdcr, [20](#)

mdcr_longitudinal, [21](#)

* icd tools

get_icd_codes, [11](#)

base::grep(), [19](#)

comorbidities, [2](#)

comorbidities(), [8–11](#), [14](#), [15](#), [22](#), [23](#)

get_charlson_codes, [7](#)

get_charlson_codes(), [8](#), [9](#)

get_charlson_index_scores, [8](#)

get_charlson_index_scores(), [8](#)

get_elixhauser_codes, [9](#)

get_elixhauser_codes(), [8](#), [10](#), [11](#)

get_elixhauser_index_scores, [10](#)

get_elixhauser_index_scores(), [9](#), [11](#)

get_elixhauser_poa, [10](#)

get_elixhauser_poa(), [9](#), [10](#)

get_icd_codes, [11](#)

get_icd_codes(), [8](#), [9](#), [14](#), [16](#), [17](#), [20](#)

get_pccc_codes, [14](#)

get_pccc_codes(), [8](#), [9](#), [15](#)

get_pccc_conditions, [15](#)

get_pccc_conditions(), [14](#)

icd_compact_to_full, [15](#), [17](#), [20](#)

icd_compact_to_full(), [17](#), [20](#)

is_icd, [16](#), [16](#), [20](#)

is_icd(), [13](#), [16](#), [20](#)

lookup_icd_codes, [16](#), [17](#), [19](#)

lookup_icd_codes(), [13](#), [16](#), [17](#)

mdcr, [20](#), [21](#)

mdcr_longitudinal, [21](#), [21](#)

summary.medicalcoder_comorbidities, [22](#)

summary.medicalcoder_comorbidities_with_subconditions,
[23](#)