

# Package ‘pccc’

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**Title** Pediatric Complex Chronic Conditions

**Version** 1.0.6

**Description** An implementation of the pediatric complex chronic conditions (CCC) classification system using R and C++.

**Depends** R (>= 3.5.0)

**License** GPL-2

**Encoding** UTF-8

**Language** en-us

**LazyData** true

**Imports** dplyr (>= 1.0.0), Rcpp (>= 1.0.11), tibble

**Suggests** covr, knitr, rmarkdown, readr

**RoxygenNote** 7.3.2

**LinkingTo** Rcpp (>= 1.0.11)

**VignetteBuilder** knitr

**URL** <https://github.com/CUD2V/pccc>

**BugReports** <https://github.com/CUD2V/pccc/issues>

**NeedsCompilation** yes

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ccc	<i>Complex Chronic Conditions (CCC)</i>
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### Description

Generate CCC and CCC subcategory flags and the number of categories.

### Usage

```
ccc(data, id, dx_cols = NULL, pc_cols = NULL, icdv)
```

### Arguments

data	a data.frame containing a patient id and all the ICD-9-CM or ICD-10-CM codes. The data.frame passed to the function should be in wide format.
id	bare name of the column containing the patient id
dx_cols, pc_cols	column names with the diagnostic codes and procedure codes respectively. These argument are passed to <a href="#">select</a> .
icdv	ICD version 9 or 10

### Details

It is recommended that you view the codes defining the CCC via [get\\_codes](#) and make sure that the ICD codes in your data set are formatted in the same way. The ICD codes used for CCC are character strings must be formatted as follows:

- *\*Do not\** use decimal points or other separators
- ICD 9 codes: Codes less than 10 should be left padded with 2 zeros. Codes less than 100 should be left padded with 1 zero.

See ‘`vignette("pccc-overview")`’ for more details.

### Value

A data.frame with a column for the subject id and integer (0 or 1) columns for each each of the categories.

## References

See [pccc-package](#) for published paper on the topic of identifying Complex Chronic Conditions

## See Also

[get\\_codes](#) to view the ICD codes used to define the CCC. [select](#) for more examples and details on how to identify and select the diagnostic and procedure code columns.

## Examples

```
eg_data <- data.frame(id = letters,
                      dx1 = c(NA, NA, sample(get_codes(10)[["hemato_immu", "dx"]], 24)),
                      dx2 = c("A", sample(get_codes(10)[["gi", "dx"]], 25)),
                      dx3 = LETTERS,
                      pc1 = c("B", sample(get_codes(10)[["cvd", "pc"]], 25)),
                      pc2 = LETTERS,
                      other_col = LETTERS)

ccc(eg_data,
     id,
     dx_cols = dplyr::starts_with("dx"),
     pc_cols = dplyr::starts_with("pc"),
     icdv = 10)
```

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comparability

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*Multiple Cause of Death (MCOD) file extract*


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## Description

The Center for Disease Control has made publicly available death certificate data. This data set is a subset of the 2006 MCOD file for decedents aged  $\leq 21$  showing just the underlying cause of death diagnosis code in ICD-9-CM and ICD-10.

## Usage

```
comparability
```

## Format

A data frame with 65037 rows and 3 variables.

**id** Sequentially assigned patient identifier

**icd9** Underlying Cause of Death ICD 9 CM diagnosis code

**icd10** Underlying Cause of Death ICD 10 diagnosis code

## See Also

`vignette("pccc-example")` for more details about the MCOD source file.

get\_codes

*Get (view) Diagnostic and Procedure Codes***Description**

View the ICD, version 9 or 10, for the Complex Chronic Conditions (CCC) categories.

**Usage**

```
get_codes(icdv)
```

**Arguments**

icdv                      and integer value specifying ICD version. Accepted values are 9 or 10.

**Details**

The CCC categories for diagnostic and procedure codes are:

category	dx	dx_fixed	pc	pc_fixed
neuromuscul	X	X	X	
cvd	X	X	X	
respiratory	X	X	X	
renal	X		X	
gi	X		X	
hemato_immu	X		X	
metabolic	X		X	X
congeni_genetic	X			
malignancy	X		X	
neonatal	X			
tech_dep	X		X	
transplant	X		X	

The ICD codes were taken from the SAS macro provided by the reference paper.

**Value**

A matrix of character vectors. Rows are the categories and columns for diagnostic and procedure codes.

**References**

Feudtner C, et al. Pediatric complex chronic conditions classification system version 2: updated for ICD-10 and complex medical technology dependence and transplantation, BMC Pediatrics, 2014, 14:199, DOI: 10.1186/1471-2431-14-199

**Examples**

```
# All ICD 9 codes for CCC
get_codes(9)

# All ICD 10 codes for CCC
get_codes(10)

# Get all the codes for ICD 9 related to malignancy
get_codes(9)["malignancy", ]
```

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pccc\_icd10\_dataset      *Randomly Generated ICD 10 Sample Data Set*

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**Description**

This dataset was produced from a tool available at [https://github.com/magic-lantern/icd\\_file\\_generator](https://github.com/magic-lantern/icd_file_generator). ICD codes were taken from CMS. The code source, for both the diagnosis and produced codes can be found at <https://www.cms.gov/medicare/coding-billing/icd-10-codes/icd-10-cm-icd-10-pcs-gem-archive> with a copy of the downloaded data on the package github page, <https://github.com/CUD2V/pccc>

**Usage**

```
pccc_icd10_dataset
```

**Format**

A data frame with 1000 rows and 31 variables. There is a patient identifier, ten diagnosis codes, ten procedure codes, and ten "other data" values, specifically:

**id** Sequentially assigned patient identifier

**dx1** a ICD 10 diagnosis code

**dx2** a ICD 10 diagnosis code

**dx3** a ICD 10 diagnosis code

**dx4** a ICD 10 diagnosis code

**dx5** a ICD 10 diagnosis code

**dx6** a ICD 10 diagnosis code

**dx7** a ICD 10 diagnosis code

**dx8** a ICD 10 diagnosis code

**dx9** a ICD 10 diagnosis code

**dx10** a ICD 10 diagnosis code

**pc1** a ICD 10 procedure codes

**pc2** a ICD 10 procedure codes

**pc3** a ICD 10 procedure codes

- pc4** a ICD 10 procedure codes
- pc5** a ICD 10 procedure codes
- pc6** a ICD 10 procedure codes
- pc7** a ICD 10 procedure codes
- pc8** a ICD 10 procedure codes
- pc9** a ICD 10 procedure codes
- pc10** a ICD 10 procedure codes
- g1** Random data to simulate other data often present in export of patient data with 20% of values missing.
- g2** Random data to simulate other data often present in export of patient data with 20% of values missing.
- g3** Random data to simulate other data often present in export of patient data with 20% of values missing.
- g4** Random data to simulate other data often present in export of patient data with 20% of values missing.
- g5** Random data to simulate other data often present in export of patient data with 20% of values missing.
- g6** Random data to simulate other data often present in export of patient data with 20% of values missing.
- g7** Random data to simulate other data often present in export of patient data with 20% of values missing.
- g8** Random data to simulate other data often present in export of patient data with 20% of values missing.
- g9** Random data to simulate other data often present in export of patient data with 20% of values missing.
- g10** Random data to simulate other data often present in export of patient data with 20% of values missing.

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pccc\_icd9\_dataset

*Randomly Generated ICD 9 Sample Data Set*


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## Description

This dataset was produced from a tool available at [https://github.com/magic-lantern/icd\\_file\\_generator](https://github.com/magic-lantern/icd_file_generator). ICD codes were taken from CMS. The ICD 9 diagnosis and procedure codes were generated with 20 missing values. Code source: <https://www.cms.gov/medicare/coding-billing/icd-10-codes/icd-9-cm-diagnosis-procedure-codes-abbreviated-and-full-code-titles>

## Usage

pccc\_icd9\_dataset

**Format**

A data frame with 1000 rows and 31 variables. There is a patient identifier, ten diagnosis codes, ten procedure codes, and ten "other data" values, specifically:

**id** Sequentially assigned patient identifier

**dx1** a ICD 9 diagnosis code

**dx2** a ICD 9 diagnosis code

**dx3** a ICD 9 diagnosis code

**dx4** a ICD 9 diagnosis code

**dx5** a ICD 9 diagnosis code

**dx6** a ICD 9 diagnosis code

**dx7** a ICD 9 diagnosis code

**dx8** a ICD 9 diagnosis code

**dx9** a ICD 9 diagnosis code

**dx10** a ICD 9 diagnosis code

**pc1** a ICD 9 procedure codes

**pc2** a ICD 9 procedure codes

**pc3** a ICD 9 procedure codes

**pc4** a ICD 9 procedure codes

**pc5** a ICD 9 procedure codes

**pc6** a ICD 9 procedure codes

**pc7** a ICD 9 procedure codes

**pc8** a ICD 9 procedure codes

**pc9** a ICD 9 procedure codes

**pc10** a ICD 9 procedure codes

**g1** Random data to simulate other data often present in export of patient data with 20% of values missing.

**g2** Random data to simulate other data often present in export of patient data with 20% of values missing.

**g3** Random data to simulate other data often present in export of patient data with 20% of values missing.

**g4** Random data to simulate other data often present in export of patient data with 20% of values missing.

**g5** Random data to simulate other data often present in export of patient data with 20% of values missing.

**g6** Random data to simulate other data often present in export of patient data with 20% of values missing.

**g7** Random data to simulate other data often present in export of patient data with 20% of values missing.

- g8** Random data to simulate other data often present in export of patient data with 20% of values missing.
- g9** Random data to simulate other data often present in export of patient data with 20% of values missing.
- g10** Random data to simulate other data often present in export of patient data with 20% of values missing.

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test\_helper

*Test Helper*

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### Description

Tool to help access internal variables to use in testthat scripts, or anytime access via `:::` would be needed.

### Usage

```
test_helper(var)
```

### Arguments

var                      bare name of the internal variable to be accessed.

### Value

Object from internal PCCC name space (if it exists)



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