

# Package: themask (via r-universe)

October 31, 2024

**Title** Masks and houses the PEPFAR MSD-style training dataset for testing and training

**Version** 1.1.5

**Description** This package creates and hosts a masked, dummy dataset that should be used for testing, training, and demoing instead of using actual PEPFAR data.

**License** MIT + file LICENSE

**Encoding** UTF-8

**Roxygen** list(markdown = TRUE)

**RoxygenNote** 7.3.1

**Imports** base, cli, dplyr, googlesheets4, magrittr, piggyback, readr, stats, stringr, tibble, utils

**Remotes** USAID-OHA-SI/gagglr, USAID-OHA-SI/glamr, USAID-OHA-SI/gophr

**Depends** R (>= 2.10)

**LazyData** true

**Suggests** glue, googledrive, ggplot2, keyring, knitr, pak, rlang (>= 1.0.0), rmarkdown, testthat (>= 3.0.0), withr

**Config/testthat/edition** 3

**Config/testthat/parallel** true

**VignetteBuilder** knitr

**Repository** <https://usaid-oha-si.r-universe.dev>

**RemoteUrl** <https://github.com/USAID-OHA-SI/themask>

**RemoteRef** HEAD

**RemoteSha** 094cfa2e48906690c9ed49153383aba0a6a31e84

## Contents

milb . . . . .	2
minoria_geo . . . . .	3
minoria_mechs . . . . .	3

minoria_shp_ou . . . . .	4
minoria_shp_psnu . . . . .	4
minoria_shp_snu1 . . . . .	5
msk_available . . . . .	5
msk_create . . . . .	6
msk_download . . . . .	6
msk_gen_uid . . . . .	7
msk_release . . . . .	8

<b>Index</b>	<b>10</b>
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milb	<i>MiLB Information Table</i>
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### Description

This table has been extracted from Wikipedia's Minor League Baseball (MiLB) page and is used for masking geographic and partner information in PEPFAR's MER Structured Datasets (MSD).

### Usage

milb

### Format

milb:

A data frame with 120 rows and 6 columns:

**division** MLB Division (East, Central, West)

**mlb\_team** Affiliated MLB

**level** MilB team level (High A, A, AA, AAA)

**league** MilB team level (varies by level)

**city** MilB team city/location

**name** MilB team name

### Source

[https://en.wikipedia.org/wiki/Minor\\_League\\_Baseball](https://en.wikipedia.org/wiki/Minor_League_Baseball)

minoria\_geo *Minoria Geography Table*

### Description

This table mimics the structure of PEPFAR’s MER Structured Datasets (MSD). The dummy dataset for the (Kingdom of) Minoria has four region (snu1), each containing four districts (psnu). Derived from milb.

### Usage

minoria\_geo

### Format

minoria\_geo:

A data frame with 16 rows and 7 columns:

**operatingunit** Operating Unit name (Minoria)

**operatingunituid** OU unique ID (Minoria)

**country** Country (Minoria)

**snu1** Sub-National Unit 1 level below national (from MiLB league)

**snu1uid** SNU1 unique ID

**cop22\_psnu** Priority SNU (from MiLB city)

**cop22\_psnuuid** PSNU unique ID

minoria\_mechs *Minoria Mechanism Table*

### Description

This table mimics the structure of PEPFAR’s MER Structured Datasets (MSD) for mechanisms, which originally originate from FACTInfo NextGen. The dummy dataset for the (Kingdom of) Minoria has 120 (plus two dedup) mechanisms available for use. Derived from milb.

### Usage

minoria\_mechs

### Format

minoria\_mechs:

A data frame with 122 rows and 7 columns:

**mech\_code** Unique implementing mechanism code

**mech\_name** implementing mechanism code (from MiLB team name)

**prime\_partner\_name** implementing mechanism partner (from MiLB team name)

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minoria_shp_ou	<i>Minoria PSNU Shape File</i>
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### Description

This dataset is the sf file for mapping (Kingdom of) Minoria at the national level

### Usage

minoria\_shp\_ou

### Format

minoria\_shp\_ou:  
 A data frame with 16 rows and 7 columns:  
**operatingunitid** OU unique ID (Minoria)  
**operatingunit** Operating Unit name (Minoria)  
**country** Country (Minoria)  
**geometry** Country level polygon shape for mapping

### See Also

Other shp: [minoria\\_shp\\_psnu](#), [minoria\\_shp\\_snu1](#)

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minoria_shp_psnu	<i>Minoria PSNU Shape File</i>
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### Description

This dataset is the sf file for mapping (Kingdom of) Minoria by PSNU.

### Usage

minoria\_shp\_psnu

### Format

minoria\_shp\_psnu:  
 A data frame with 16 rows and 8 columns:  
**psnu** Priority SNU (from MiLB city)  
**psnuuid** PSNU unique ID  
**snu1** Sub-National Unit 1 level below national (from MiLB league)  
**snu1uid** SNU1 unique ID  
**operatingunit** Operating Unit name (Minoria)  
**operatingunitid** OU unique ID (Minoria)  
**country** Country (Minoria)  
**geometry** PSNU level polygon shape for mapping

**See Also**

Other shp: [minoria\\_shp\\_ou](#), [minoria\\_shp\\_snu1](#)

minoria\_shp\_snu1      *Minoria PSNU Shape File*

**Description**

This dataset is the sf file for mapping (Kingdom of) Minoria by SNU1.

**Usage**

minoria\_shp\_snu1

**Format**

minoria\_shp\_snu1:

A data frame with 4 rows and 6 columns:

**snu1uid** SNU1 unique ID

**snu1** Sub-National Unit 1 level below national (from MiLB league)

**operatingunit** Operating Unit name (Minoria)

**operatingunituid** OU unique ID (Minoria)

**country** Country (Minoria)

**geometry** SNU1 level polygon shape for mapping

**See Also**

Other shp: [minoria\\_shp\\_ou](#), [minoria\\_shp\\_psnu](#)

msk\_available      *Check the latest version available*

**Description**

This function is used to check what masked version is currently available and will flag if there it is up to date or you should run `msk_create` yourself. It will also list all available historic releases that can be downloaded in `msk_download` by specifying the version in the tag param.

**Usage**

`msk_available()`

**See Also**

Other download: [msk\\_download\(\)](#)

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`msk_create`*Create the Masked Dataset*

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**Description**

This function is used to create a masked dataset for use in testing and training. The data are subset to 16 PSNUs and masked across geographic and mechanism variables. Either the PSNUxIM or NAT\_SUBNAT Structured Datasets can be masked.

**Usage**

```
msk_create(filepath, output_folder)
```

**Arguments**

`filepath` path to the PSD file (PSNUxIM or NAT\_SUBNAT)  
`output_folder` location where you want to store the new file (default = does not export the data)

**Value**

dataframe with converted geography + mech info

**Examples**

```
## Not run:  
#create a masked dataset  
library(glamr)  
library(themask)  
  
#store path to latest MSD  
path <- si_path() %>% return_latest("PSNU_IM")  
  
#create a masked dataset from the PSNUxIM MSD  
msk_create(path, "project1/data")  
  
## End(Not run)
```

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`msk_download`*Download Masked Dataset*

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**Description**

This function download a masked dataset from GitHub ([USAID-OHA-SI/themask](https://github.com/USAID-OHA-SI/themask)) for use in training or testing. Recommend running `msk_available` to see what version is available to download.

**Usage**

```
msk_download(folderpath, tag = "latest", launch = FALSE)
```

**Arguments**

folderpath	where should the file be downloaded to?
tag	version tag, default = "latest"
launch	whether to launch Windows Explorer to the location after the download completes (default = FALSE)

**References**

msk\_available

**See Also**

Other download: [msk\\_available\(\)](#)

**Examples**

```
## Not run:  
#check available version  
msk_available()  
  
#download to your downloads folder  
msk_download("~/Downloads")  
  
## End(Not run)  
## Not run:  
#download an older version  
msk_download("~/Downloads", tag = "2023.06.27c")  
  
## End(Not run)
```

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msk\_gen\_uid

*Generate a Unique ID*

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**Description**

This function is used to create a unique ID (UID) to mimic the UIDs created and used by DATIM for the MER and other PEPFAR Structured Datasets.

**Usage**

```
msk_gen_uid(codeSize = 11)
```

**Arguments**

codeSize	character length for UID output (default = 11)
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**Value**

random alphanumeric string

**Examples**

```
msk_gen_uid()
```

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msk\_release

*Upload a New Masked Dataset to GitHub*

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**Description**

This function is used to upload the new masked dataset to GitHub (USAID-OHA-SI/themask) so that others can use it. This function is for package developers use only. The data from `msk_create()` are subset to 16 PSNUs and masked across geographic and mechanism variables. Either the PSNUxIM or NAT\_SUBNAT Structured Datasets can be masked.

**Usage**

```
msk_release(filepath, output_folder)
```

**Arguments**

filepath            path to the PSD file (PSNUxIM or NAT\_SUBNAT) or masked file  
output\_folder      location where you want to store the new file (default does not save the file)

**Value**

dataframe with converted geography + mech info

**References**

msk\_create

**Examples**

```
## Not run:  
#create and upload a new release  
library(glamr)  
library(themask)  
  
#store path to latest MSD  
path <- si_path() %>% return_latest("PSNU_IM")  
  
#create a masked dataset from the PSNUxIM MSD  
msk_release(path, "project1/data")  
  
## End(Not run)
```



```
## Not run:
#upload release from an existing masked dataset
library(glamr)
library(themask)

#store path to masked dataset
path_msk <- return_latest("project1/data","TRAINING")

#create a masked dataset from the PSNUxIM MSD
msk_release(path_msk)

## End(Not run)
```

# Index

## \* datasets

- milb, 2
- minoria\_geo, 3
- minoria\_mechs, 3
- minoria\_shp\_ou, 4
- minoria\_shp\_psnu, 4
- minoria\_shp\_snu1, 5

## \* download

- msk\_available, 5
- msk\_download, 6

## \* shp

- minoria\_shp\_ou, 4
- minoria\_shp\_psnu, 4
- minoria\_shp\_snu1, 5

- milb, 2
- minoria\_geo, 3
- minoria\_mechs, 3
- minoria\_shp\_ou, 4, 5
- minoria\_shp\_psnu, 4, 4, 5
- minoria\_shp\_snu1, 4, 5, 5
- msk\_available, 5, 7
- msk\_create, 6
- msk\_create(), 8
- msk\_download, 5, 6
- msk\_gen\_uid, 7
- msk\_release, 8