

# Package: ontologics (via r-universe)

October 24, 2024

**Title** Code-Logics to Handle Ontologies

**Version** 0.7.4

**Description** Provides tools to build and work with an ontology of linked (open) data in a tidy workflow. It is inspired by the Food and Agriculture Organizations (FAO) caliper platform <<https://www.fao.org/statistics/caliper/web/>> and makes use of the Simple Knowledge Organisation System (SKOS).

**URL** <https://github.com/luckinet/ontologics>

**BugReports** <https://github.com/luckinet/ontologics/issues>

**Depends** R (>= 3.5.0)

**License** GPL (>= 3)

**Encoding** UTF-8

**LazyData** true

**Roxygen** list(markdown = TRUE)

**RoxygenNote** 7.3.2

**Imports** checkmate, rlang, tibble, dplyr, tidyr, tidyselect, magrittr, purrr, stringr, readr, httr, methods, rdfliib, beeper

**Suggests** knitr, rmarkdown

**VignetteBuilder** knitr

**Repository** <https://luckinet.r-universe.dev>

**RemoteUrl** <https://github.com/luckinet/ontologics>

**RemoteRef** HEAD

**RemoteSha** a127aed8e2bac0c00984d460c7a458f2012910b8

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export_as_rdf	<i>Export an ontology as RDF</i>
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### Description

Export an ontology as RDF

### Usage

```
export_as_rdf(ontology, filename)
```

### Arguments

ontology	<a href="#">ontology(1)</a> an already loaded or created ontology object.
filename	<a href="#">character(1)</a> the filename of the exported ontology. The format of the exported ontology is guessed by the extension of the filename. The guessing is performed by the rdfliib package. Valid extensions are ".rdf" for "rdxml", ".nt" for "ntriples", ".ttl" for "turtle" or ".json" for "jsonld".

### Value

No return value, called for the side effect of exporting an ontology.

### Examples

```
ontoDir <- system.file("extdata", "crops.rds", package = "ontologics")
onto <- load_ontology(path = ontoDir)

## Not run:

  export_as_rdf(ontology = onto, filename = "onto.ttl")

## End(Not run)
```

---

get_class	<i>Get class(es) in an ontology</i>
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---

**Description**

Get class(es) in an ontology

**Usage**

```
get_class(..., regex = FALSE, external = FALSE, ontology = NULL)
```

**Arguments**

...	combination of column name and value to filter that column by. The value to filter by can be provided as regular expression, if <code>regex = TRUE</code> .
regex	<a href="#">logical(1)</a> whether or not the value in ... shall be matched in full, or whether any partial match should be returned.
external	<a href="#">logical(1)</a> whether or not the external classes (TRUE), or the harmonized classes should be returned (FALSE, default).
ontology	<a href="#">ontology(1)</a> either a path where the ontology is stored, or an already loaded ontology.

**Value**

A table of the class(es) in the ontology according to the values in ...

**Examples**

```
ontoDir <- system.file("extdata", "crops.rds", package = "ontologics")
onto <- load_ontology(path = ontoDir)

# exact classes from a loaded ontology ...
get_class(label = "class", ontology = onto)

# ... or one stored on the harddisc
get_class(id = ".xx.xx", ontology = ontoDir)

# use regular expressions ...
get_class(label = "ro", regex = TRUE, ontology = onto)

# get all sources
get_class(ontology = onto)
```

---

`get_concept`*Get a concept in an ontology*

---

**Description**

Get a concept in an ontology

**Usage**

```
get_concept(..., external = FALSE, matches = FALSE, ontology = NULL)
```

**Arguments**

<code>...</code>	combination of column name and value to filter that column by.
<code>external</code>	<code>logical(1)</code> whether or not to return merely the table of external concepts.
<code>matches</code>	<code>logical(1)</code> whether or not to include external concepts as label instead of id in the match columns of the harmonised concepts; this allows querying the external concepts in the harmonised concepts (only if <code>external = FALSE</code> ).
<code>ontology</code>	<code>ontology(1)</code> either a path where the ontology is stored, or an already loaded ontology.

**Value**

A table of a subset of the ontology according to the values in ...

**Examples**

```
ontoDir <- system.file("extdata", "crops.rds", package = "ontologics")
onto <- load_ontology(path = ontoDir)

# exact matches from a loaded ontology ...
get_concept(label = "FODDER CROPS", ontology = onto)

# ... or a path
get_concept(label = c("FODDER CROPS", "CEREALS"), ontology = ontoDir)

# ignore queries that would not be valid in filter()
get_concept(label != 'Bioenergy woody' & has_broader == '.01', ontology = onto)

# extract concepts based on regular expressions
library(stringr)
get_concept(str_detect(label, "crop") & str_detect(id, ".03$"), ontology = ontoDir)
```

---

get_source	<i>Get source(e) in an ontology</i>
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---

## Description

Get source(e) in an ontology

## Usage

```
get_source(..., regex = FALSE, ontology = NULL)
```

## Arguments

...	combination of column name and value to filter that column by. The value to filter by can be provided as regular expression, if <code>regex = TRUE</code> .
regex	<code>logical(1)</code> whether or not the value in ... shall be matched in full, or whether any partial match should be returned.
ontology	<code>ontology(1)</code> either a path where the ontology is stored, or an already loaded ontology.

## Value

A table of the source(s) in the ontology according to the values in ...

## Examples

```
ontoDir <- system.file("extdata", "crops.rds", package = "ontologics")
onto <- load_ontology(path = ontoDir)

# exact sources from a loaded ontology ...
get_source(label = "harmonised", ontology = onto)

# ... or one stored on the harddisc
get_source(version = "0.0.1", ontology = ontoDir)

# get all sources
get_source(ontology = onto)
```

---

load_ontology	<i>Load an ontology</i>
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---

**Description**

Load an ontology

**Usage**

```
load_ontology(path = NULL)
```

**Arguments**

path	<code>character(1)</code> the path where the ontology to load is stored.
------	---

**Value**

A table of the full ontology (i.e., where attribute and mapping tables are joined).

**Examples**

```
# load an already existing ontology
load_ontology(path = system.file("extdata", "crops.rds", package = "ontologies"))
```

---

make_tree	<i>Make a tree of an ontology</i>
-----------	-----------------------------------

---

**Description**

Make a tree of an ontology

**Usage**

```
make_tree(..., reverse = FALSE, ontology = NULL)
```

**Arguments**

...	<code>character(1)</code> the concepts that shall be the target, combination of ' <i>column name = value</i> '.
reverse	<code>logical(1)</code> whether or not to make a tree that gives the parents, instead of the children, of target concepts.
ontology	<code>ontology(1)</code> either a path where the ontology is stored, or an already loaded ontology.

---

new_class	<i>Add a new valid class to an ontology</i>
-----------	---

---

**Description**

Add a new valid class to an ontology

**Usage**

```
new_class(new, target, description = NULL, ontology = NULL)
```

**Arguments**

new	<a href="#">character(1)</a> the new class label.
target	<a href="#">character(1)</a> the class into which the new class shall be nested.
description	<a href="#">character(1)</a> a verbatim description of the new class.
ontology	<a href="#">ontology(1)</a> either a path where the ontology is stored, or an already loaded ontology.

**Value**

the updated ontology that contains the new class(es) defined here.

**Examples**

```
ontoDir <- system.file("extdata", "crops.rds", package = "ontologics")
onto <- load_ontology(path = ontoDir)

onto <- new_class(new = "use type", target = "class", description = "something",
  ontology = onto)
```

---

new_concept	<i>Add a new concept to an ontology</i>
-------------	---

---

**Description**

This adds a new concept to an existing ontology to semantically integrate and thus harmonise it with the already existing ontology.

**Usage**

```
new_concept(
  new,
  broader = NULL,
  description = NULL,
  class = NULL,
  ontology = NULL
)
```

**Arguments**

new	<a href="#">character(.)</a> the english label(s) of new concepts that shall be included in the ontology.
broader	<a href="#">data.frame(.)</a> the english label(s) of already harmonised concepts to which the new concept shall be semantically linked via a <b>skos:broader</b> relation, see Details.
description	<a href="#">character(.)</a> a verbatim description of the new concept(s).
class	<a href="#">character(.)</a> the class(es) of the new labels.
ontology	<a href="#">ontology(1)</a> either a path where the ontology is stored, or an already loaded ontology.

**Value**

returns invisibly a table of the new harmonised concepts that were added to the ontology, or a message that nothing new was added.

**Examples**

```
ontoDir <- system.file("extdata", "crops.rds", package = "ontologies")
onto <- load_ontology(path = ontoDir)

# add fully known concepts
concepts <- data.frame(
  old = c("Bioenergy woody", "Bioenergy herbaceous"),
  new = c("acacia", "miscanthus")
)

onto <- new_source(
  version = "0.0.1",
  name = "externalDataset",
  description = "a vocabulary",
  homepage = "https://www.something.net",
  license = "CC-BY-0",
  ontology = onto
)

onto <- new_concept(
```



```
    new = concepts$new,  
    broader = get_concept(label = concepts$old, ontology = onto),  
    class = "crop",  
    ontology = onto  
  )  
  
  # add concepts where the nesting is clear, but not the new class  
  concepts <- data.frame(  
    old = c("Barley", "Barley"),  
    new = c("food", "bio-energy")  
  )  
  
  onto <- new_concept(  
    new = concepts$new,  
    broader = get_concept(label = concepts$old, ontology = onto),  
    ontology = onto  
  )  
  
  # define that class ...  
  onto <- new_class(  
    new = "use type", target = "class",  
    description = "the way a crop is used", ontology = onto  
  )  
  
  # ... and set the concepts again  
  onto <- new_concept(  
    new = concepts$new,  
    broader = get_concept(label = concepts$old, ontology = onto),  
    class = "use type",  
    ontology = onto  
  )  
)
```

---

new\_mapping

*Add a new mapping to an ontology*

---

## Description

Extend an ontology by creating mappings between classes and concepts of external vocabularies and the harmonised classes and concepts.

## Usage

```
new_mapping(  
  new = NULL,  
  target,  
  source = NULL,  
  lut = NULL,  
  match = NULL,  
  certainty = NULL,  
)
```

```

    type = "concept",
    ontology = NULL,
    verbose = FALSE,
    beep = NULL
  )

```

### Arguments

new	<a href="#">character(.)</a> the english external label(s) that shall be mapped to labels that do already exist in the ontology.
target	<a href="#">data.frame(.)</a> the already harmonised English label(s) to which the external labels shall be mapped; derive with <code>get_concept()</code> .
source	<a href="#">character(1)</a> any character uniquely identifying the source dataset of the new label.
lut	<a href="#">character(.)</a> in case the terms used for mapping are from a look up table (i.e. a standardised set of terms with a description), provide this table with column names 'label' and 'description' here.
match	<a href="#">character(1)</a> the <b>skos mapping property</b> used to describe the link, possible values are "close", "exact", "broad" and "narrow".
certainty	<a href="#">integerish(1)</a> the certainty of the match. Possible values are between 1 and 4, with meaning <ul style="list-style-type: none"> <li>• 1 = probably unreliable</li> <li>• 2 = unclear, assigned according to a given definition</li> <li>• 3 = clear, assigned according to a given definition</li> <li>• 4 = original, harmonised term (can't be assigned by a user)</li> </ul>
type	<a href="#">character(1)</a> whether the new labels are mapped to a "concept", or to a "class".
ontology	<a href="#">ontology(1)</a> either a path where the ontology is stored, or an already loaded ontology.
verbose	<a href="#">logical(1)</a> whether or not to give detailed information on the process of this function.
beep	<a href="#">integerish(1)</a> Number specifying what sound to be played to signal the user that a point of interaction is reached by the program, see <a href="#">beep</a> .

### Value

No return value, called for the side effect of adding new mappings to an ontology.

## Examples

```
ontoDir <- system.file("extdata", "crops.rds", package = "ontologics")
onto <- load_ontology(path = ontoDir)

mapping <- data.frame(old = c("BIOENERGY CROPS", "Bioenergy woody",
                             "Other bioenergy crops"),
                     new = c("bioenergy plants", "Wood plantation for fuel",
                             "Algae for bioenergy"),
                     type = c("close", "broader", "broader"))

onto <- new_source(name = "externalDataset",
                  version = "0.0.1",
                  description = "a vocabulary",
                  homepage = "https://www.something.net",
                  license = "CC-BY-0",
                  ontology = onto)

onto <- get_concept(label = mapping$old, ontology = onto) %>%
  new_mapping(new = mapping$new,
             target = .,
             match = mapping$type,
             source = "externalDataset",
             certainty = 3,
             ontology = onto)
```

---

new\_source

*Add a new valid source to an ontology*

---

## Description

Add a new valid source to an ontology

## Usage

```
new_source(
  ontology = NULL,
  name = NULL,
  version = NULL,
  date = NULL,
  description = NULL,
  homepage = NULL,
  uri_prefix = NULL,
  license = NULL,
  notes = NULL
)
```

## Arguments

ontology	<a href="#">ontology(1)</a> either a path where the ontology is stored, or an already loaded ontology into which the new source should be included.
name	<a href="#">character(1)</a> the name of the new source (must not contain empty spaces).
version	<a href="#">character(1)</a> an optional version of the new source (any value is allowed, but should be a value that follows <a href="#">semantic versioning</a> ). Either version or date need to be given.
date	<a href="#">character(1)</a> an optional date at which that version of an external vocabulary has been created. Should be a value of the form YYYY-MM-DD. Either version or date need to be given.
description	<a href="#">character(1)</a> a verbatim description of the new source.
homepage	<a href="#">character(1)</a> the homepage of the new source, typically the place where additional information or meta-data could be retrieved in a non-formalised way.
uri_prefix	<a href="#">character(1)</a> the basic uniform resource locator (URL) all concepts of a new source have in common and which is thus the basis to construct the concept specific URI.
license	<a href="#">character(1)</a> the licenses under which the new source is published.
notes	<a href="#">character(1)</a> any notes on the new source that don't fit into any of the other meta-data fields here.

## Details

Fundamentally, there are two types of sources that can be defined with this function.

- *attribute collections*: where a collection of terms or concepts are associated as a descriptive attribute to the harmonised concepts, and
- *linked open data*: where the concepts that occur in another vocabulary or ontology and which are themselves part of linked datasets (and hence have a valid URI) are associated as related concepts to the harmonised concepts.

In the latter case, each mapped concept should be provided by its ID and the source needs to have a URL that allows in combination with the concept IDs to construct the URI under which the mapped concepts are stored in the semantic web.

## Value

the updated ontology that contains the new source defined here.

**Examples**

```
ontoDir <- system.file("extdata", "crops.rds", package = "ontologics")
onto <- load_ontology(path = ontoDir)

onto <- new_source(name = "externalDataset",
                  version = "0.0.1",
                  description = "a vocabulary",
                  homepage = "https://www.something.net",
                  license = "CC-BY-0",
                  ontology = onto)
```

---

onto-class	<i>Ontology class (S4) and methods</i>
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---

**Description**

Ontology class (S4) and methods

**Slots**

sources [data.frame\(.\)](#)

classes [data.frame\(.\)](#)

concepts [data.frame\(.\)](#)

---

show, onto-method	<i>Print onto in the console</i>
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---

**Description**

Print onto in the console

**Usage**

```
## S4 method for signature 'onto'
show(object)
```

**Arguments**

object            object to show.

---

start_ontology	<i>Start an ontology</i>
----------------	--------------------------

---

**Description**

Start an ontology

**Usage**

```
start_ontology(  
    name = NULL,  
    version = NULL,  
    path = NULL,  
    code = ".xx",  
    description = NULL,  
    homepage = NULL,  
    uri_prefix = NULL,  
    license = NULL,  
    notes = NULL  
)
```

**Arguments**

name	<a href="#">character(1)</a> the path of the ontology.
version	<a href="#">character(1)</a> the version of the ontology.
path	<a href="#">character(1)</a> the path where the ontology shall be stored.
code	<a href="#">double(1)</a> format of a single code snippet that is concatenated for nested levels.
description	<a href="#">character(1)</a> a brief description of the new ontology.
homepage	<a href="#">character(1)</a> the URL to the homepage of the new ontology.
uri_prefix	<a href="#">character(1)</a> the basic URL to construct URIs for all concepts.
license	<a href="#">character(1)</a> any string describing the license under which this ontology can be (re)used.
notes	<a href="#">character(1)</a> any notes that might apply to this ontology.

**Value**

it returns the new, empty ontology and also stores that within the directory specified in path.

**Examples**

```
start_ontology(name = "crops", path = tempdir())
```

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