

Package ‘I2B2API’

May 28, 2015

Title API to i2b2 that is exposed to the R IDE

Version 1.0

Description

This documentation describes the available functions that can be used in the R-IDE web interface.

Depends R (>= 3.0.2)

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LazyData true

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getConcepts	<i>Get the prefixes of all concepts that start with anything from the concepts list (all concepts by default)</i>
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Description

Get the prefixes of all concepts that start with anything from the concepts list (all concepts by default)

Arguments

- concepts list of concepts to be compared to. Defaults to empty, meaning any prefix.
- level Aggregates ICD codes, e.g. Level 3 = ICD:M54*, Level 1: ICD:M*

Value

list of distinct concept_cd prefixes

Examples

```
getConcepts()  
getConcepts(c('K'), 3)
```

getObservations	<i>Get all observations with specified properties</i>
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Description

Get all observations with specified properties

Arguments

- interval specifies the time frame where the observations should come from
- concepts specifies a subset where the observations should come from. Default is any concept.
- level specifies ICD level to aggregate, e.g. Level 3 = ICD:M54*, Level 1: ICD:M*
- patient_set id of the desired patient set, defaults to -1

Value

list of observations with attributes patient_num, concept_cd_sub

Examples

```
getObservations(list(start=i2b2DateToPOSIXlt("01/01/2009"), end=i2b2DateToPOSIXlt("01/01/2010")), concepts=c("K"))  
getObservations(list(start=i2b2DateToPOSIXlt("01/01/2009"), end=i2b2DateToPOSIXlt("01/01/2010")), concepts=c("K"))
```

`GetObservationsForConcept`*Get all observations for a specified concept with specified properties*

Description

Get all observations for a specified concept with specified properties

Arguments

<code>interval</code>	specifies the time frame where the observations should come from
<code>concept.path</code>	specifies the path for the concept where the observations should come from.
<code>patient_set</code>	id of the desired patient set, defaults to -1

Value

list of observations with attributes `patient_num`, `concept_cd_sub`, counts

Examples

```
getObservationsForConcept(list(start=i2b2DateToPOSIXlt("01/01/2009"), end=i2b2DateToPOSIXlt("01/01/2010")), cor
```

`getObservationsWithLimit`*Get observations with a limit to speed up development*

Description

Get observations with a limit to speed up development

Arguments

<code>interval</code>	specifies start and end of the interval in a list
<code>concepts</code>	list of concepts for the observations
<code>level</code>	specifies the aggregation level of concepts
<code>patients_limit</code>	limit of observations to be returned

Value

list of observations with `patient_num`, `concept_cd>sub`, counts

Examples

```
getObservationsWithLimit(start=i2b2DateToPOSIXlt("01/01/2009"), end=i2b2DateToPOSIXlt("01/01/2010")), concepts=
```

`getPatients`*Get all patients of a patient set*

Description

Get all patients of a patient set

Arguments

`patient_set` Id of the desired patient set

Value

list of patients with `patient_num`, `sex_cd`, `birth_date`

Examples

```
getPatients(patient_set=-1)
```

`GetPatientSetDescription`*Get the description of a patient set*

Description

Get the description of a patient set

Arguments

`patient_set` Id of the desired patient set

Value

string with description of the patient set

Examples

```
getPatientSetDescription(patient_set=42)
```

`getPatientsWithLimit` *Get patients of a patient set with a limit to speed up development*

Description

Get patients of a patient set with a limit to speed up development

Arguments

<code>patient_set</code>	Id of the desired patient set
<code>limit</code>	limit of patients to be returned

Value

list of patients with `patient_num`, `sex_cd`, `birth_date`

Examples

```
getPatients(patient_set=-1, limit=100)
```

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