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DjangoAPI

Download

Download zip file

or

• git clone https://github.com/phi-grib/RDTExtractorWeb

Install

Install and activate the environment

```
cd RDTextractorWeb
```

conda env create -f environment.yml

```
source activate RDTExtractorWeb
```

You will need to put the data files we will provide in the API/static/data/ folder. These files are distributed separately, please contact us to get them.

Run

In the root folder execute:

```
python manage.py runserver
```

and then navigate to http://127.0.0.1:8000.

Introduction

This tool is designed to extract data from the *in vivo* repeat-dose toxicity (RDT) studies' database generated within the context of the eTOX project. These data are expanded using an histopathological observation and an anatomical entity ontologies. The histopathological ontology is obtained from Novartis and can be used under the Apache License 2.0. The anatomical entities ontology is extracted from the following paper:

Hayamizu TF, Mangan M, Corradi JP, Kadin JA, Ringwald M. Genome Biol. 2005; 6(3): R29

In order to be able to aggregate the data by parent compound, some pre-processing has to be done to data as they exist in the database. Each substance is standardised according to the following protocol:

- From this repository use the *process_smiles.std* method to standardize, discard mixtures, discard compound with metal ions, and remove all salts. Also use the *neutralise.run* method to neutralise all charges when possible.
- Using molVS, get the canonical tautomer.

This project is an extension of the work published in the following paper:

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 López-Massaguer O, Pinto-Gil K, Sanz F, Amberg A, Anger LT, Stolte M, Ravagli C, Marc P, Pastor M. Toxicol Sci. 2018 Mar; 162(1): 287–300.

Manual

Exract studies' findings based on the given filtering and the organs' and morphological changes' ontologies-based expansions of these findings.

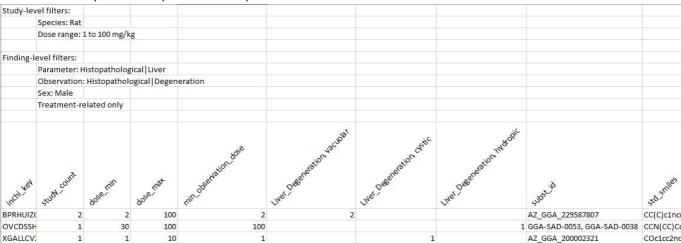
Output example

On clicking the 'Extract' button, two output files are generated, one with quantitative and the other with qualitative data. Both have a caption summarising the filtering criteria applied. After this caption, they both have a table with the data aggergated by parent compound. The table contains several fixed columns, namely 6 at the beginning:

- inchi_key: Parent compound's InChlKey.
- study_count: Number of relevant studies (according to the current filtering scheme) in which the compound appears.
- dose_min: Minimum dose at which the compound has been tested among the relevant studies.
- dose_max: Maximum dose at which the compound has been tested among the relevant studies.
- min_observation_dose: Minimum dose for which a relevant finding (according to the current filtering scheme) has been reported for the compound.
- is_active: Boolean indicating whether the substance has been found to have any toxicity according to the current finding-related filtering criteria. And two at the end:
- subst_id: All substance IDs corresponding to the parent compound.
- std_smiles: Smiles string corresponding to the standardised parent compound.

Between these two groups, there is a column for each relevant finding. In these columns a value is provided if the finding is reported for the given substance, and it is empty otherwise. The value will be the number of studies that report the finding in the qualitative file, and the minimum dose at which the finding is reported in the quantitative file.

This is an example of the qualitative output:



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This is an example of the quantitative output:

Study-lev	el filters:								
	Species: Rat								
	Dose range: 1	l to 100 mg/kg	3						
Finding-le	evel filters:								
Parameter: Histopathologi			ical Liver						
	Observation: Histopathological Degen			eration					
	Sex: Male		THE STATE OF THE S						
	Treatment-related only								
	·			nin observation	Notes Set	Liver Deserver	Liver Deserver	tion hydropic	
inchi yet	Study Count	dose min	dose mat	min olsen	Liver Degen	Liver Degen	They Dege	subst.id	std smiles
inchi ket BPRHUIZO	Study Count	dose min	dose mat	min observ	I'Met Degel	Iner Degen	The Dese.	substid AZ GGA 229	std smiles
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