NOVEL COMPUTATIONAL TOOLS AND DATABASES

DOI: https://doi.org/10.18454/jbg.2018.4.9.1

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Received: 9.10.2018; Accepted: 18.10.2018; Published: 15.11.2018

CYTPOWER: IMPLEMENTATION OF CYTOTOXIC POWER CONCEPT IN PYTHON

Abstract

Currently existing methods and parameters are inadequate for representative and quantitative assessment of cytotoxic activity of chemical compounds. Recently, we have introduced a concept of cytotoxic power which provides an effective way to a reasonable estimation and comparative analysis of cytotoxic activity of chemical agents. The basis of this concept is an integrative parameter that relies on important toxicological principles: dose of compound related to a cell number and exposure-time-response relationship of its activity.

Herein, we propose an implementation of our concept of cytotoxic power as a package developed in Python programming language. The package is free and available on GitHub repository.

Keywords: cytotoxicity, apoptosis, necrosis, computational biology.

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CYTPower: Реализация концепции цитотоксической мощности на языке Python

Abstract

Currently existing methods and parameters are inadequate for representative and quantitative assessment of cytotoxic activity of chemical compounds. Recently, we have introduced a concept of cytotoxic power which provides an effective way to a reasonable estimation and comparative analysis of cytotoxic activity of chemical agents. The basis of this concept is an integrative parameter that relies on important toxicological principles: dose of compound related to a cell number and exposure-time-response relationship of its activity.

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Announcement

Cytotoxic power can be used to assess any type of cell death based on corresponding specific and/or non-specific cellular markers (as a rule, epigenetic in its nature, named phens). It is computed as a definite integral of a dose-response function divided by time delta. For instance, we can obtain cytotoxic power values for multiple cell indices: percentage of early apoptotic cells or late apoptotic and necrotic cells. These data may be further processed with basic and advanced statistical methods (such as ANOVA), multiple-criteria decision analysis (MCDA) or quantitative structure-activity relationship (QSAR) models.
\[ CP = \frac{\int_a^b f(x)dx}{\Delta t} \]

where \( x \) is a compound dose, \( f(x) \) is a dose-response function, \( \Delta t \) is incubation duration, \( a \) is minimal dose, and \( b \) is maximum dose. Thus, unit of cytotoxic power is mol % per number of cells per hour. This equation can be extended to a sum of integrals to calculate total cytotoxic power over multiple indices.

By default, spline is used to fit dose curves, but one can specify compound—function pairs to control the fitting process, e.g. sigmoid for compound 1, polynomial for compound 2. Bootstraping is applied to obtain multiple cytotoxic power values for multiple experimental results (number of iterations may be specified manually).

The algorithm of cytotoxic power calculation can be reduced to three steps:

1) Define the interval of effective doses (integration interval).
2) Choose an approximating function to fit dose-response relationship.
3) Compute the definite integral of a dose-response function and divide it by time delta. Optionally, sum definite integrals of functions to get the overall cytotoxic power based on cell death percentages or cellular markers (e.g. early apoptosis + late apoptosis and necrosis).

![Figure 1. Dose-response relationship of cytotoxic activity of compounds A1 and A2 plotted by cytpower package](image)

We have implemented the algorithm for calculating cytotoxic power as a package in Python programming language (https://www.python.org). The package provides two convinient functions to explore and plot results and compute cytotoxic power values: `calc()` and `plot()` methods. Cytpower is based on Numpy ndarrays that are optimized for fast and scalable computations and Pandas DataFrame objects to store, process, and export data to Excel, CSV, or any available format. The package is freely available on PyPi and GitHub (https://github.com/maximtrp/cytpower).

Typical procedure starts with (1) obtaining results and creating a dataset, (2) importing data with a Pandas read function, and choosing data columns according to cytpower API. Calling `plot()` method results in rendering publication-ready plots and can be used to explore data. Example plots are shown in Fig. 1. Final step is (3) calling `calc()` method to obtain cytotoxic power values.

We provide a simple example of cytpower use with Pandas:

```python
>>> import pandas as pd
>>> import cytpower as cp

# Step 1. Obtaining results and creating a dataset
          'dose': [1, 10, 50, 100, 500, 1, 10, 50, 100, 500],
          'apoptosis': [2, 10, 14, 35, 53, 1, 5, 17, 24, 30],
          'necrosis': [1, 4, 9, 28, 31, 0, 3, 9, 19, 21]}

# Step 2. Importing data
>>> df = pd.DataFrame.from_dict(data)

# Step 3. Calculating cytotoxic power
>>> result = cp.calc(cyt_cols=['apoptosis', 'necrosis'],
                   cmp_col='compound',
                   dose_col='dose',
                   df, iters=1)

# Table with results
>>> result.round(2)
          apoptosis  necrosis  compound  min  max
0  1683.24  1551.96          A    1  500
1   277.31   683.15          B    1  500
```

There are two compounds (“A” and “B”), the percentage values of apoptotic and necrotic cells, and dose values in this example dataset. Pandas `from_dict()` method will take Python dictionary keys as column names and values as indexed rows.
This table contains cytotoxic power values in “apoptosis” and “necrosis” columns. These results suggest that compound “A” is more powerful than compound “B”. Here, the results of a single experiment were processed with cytpower. But one can specify multiple values for each cell marker and dose. We also provide a complex cytotoxicity analysis workflow with basic statistics in a Jupyter Notebook on GitHub. 

Thus, cytpower package can be used to plot, calculate and compare cytotoxic power values of chemical agents analyzed with various cellular assays.

**Funding**

These studies were supported by State assignment AAAA-A18-118012290142-9.

**Conflict of Interest**

None declared.

**References**