

Taxonomically Informed Metabolite Annotation

This app helps performing TIMA as described in the [following document](#)

Created by [Adriano Rutz](#) • [Main publication](#) • Code [on GitHub](#)

Parameters

Files

Annotations

MS

Organisms

Names

Weights

Options

GNPS (optional)

You have to input the scientific name of your taxon.

Will be used for the weighting of your annotations

Example: *Gentiana lutea*

All features will be attributed to this source.

For finer attribution, you need to provide a metadata file in the `Files` panel.

Okay

attributed to a taxon (e.g. *Gentiana lutea*)

Please provide your taxon name with capital letter

Number of organisms to keep per feature

1 5

☐ Filter library to restrict it to a portion of organisms only

Level at which the library will be filtered

phylum ▼

Value to be applied for filtering

Save parameters

Launch job