Taxonomically Informed Metabolite Annotation
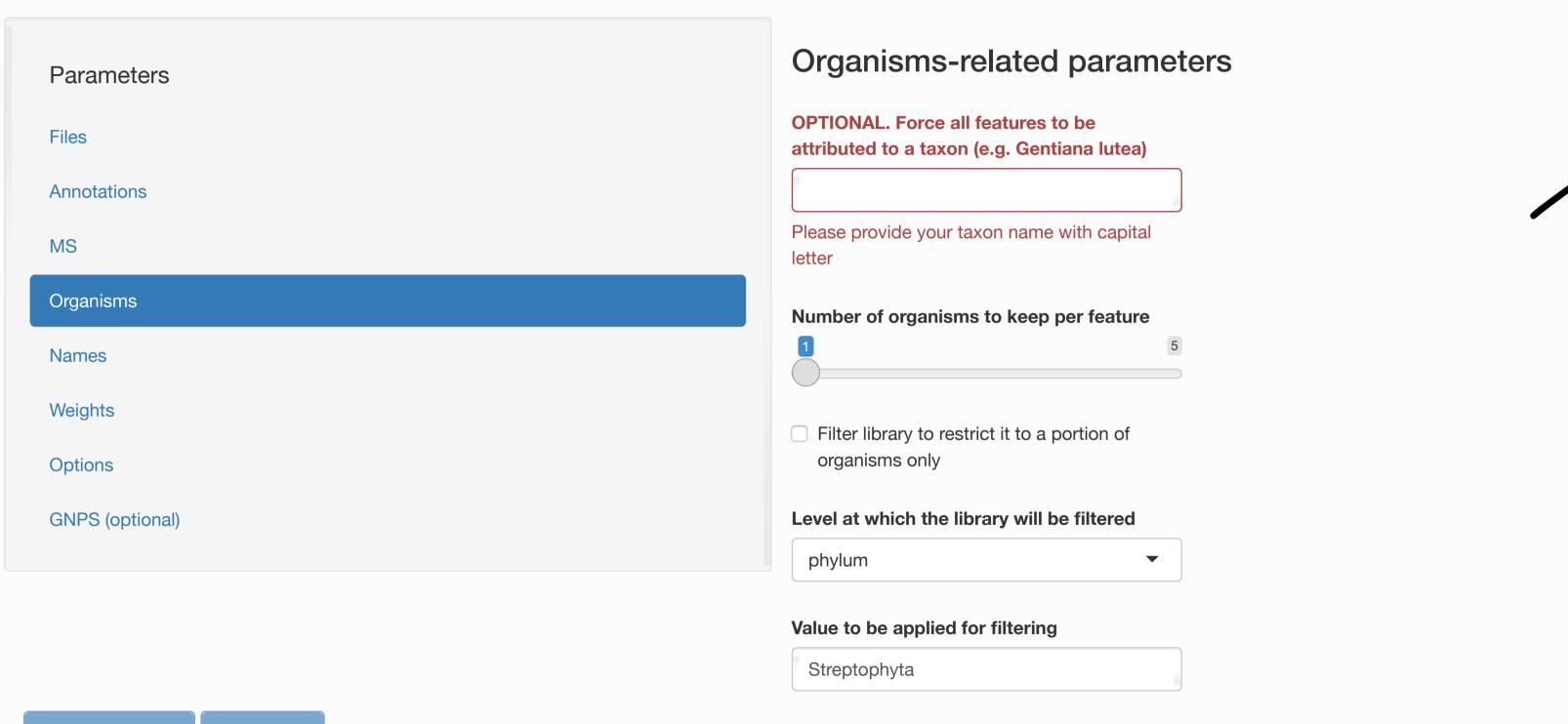
This app helps performing TIMA as described in the following documentation

Created by **Adriano Rutz** · **Main publication** · Code **on GitHub**

Parameters

Files

Annotations

MS

Organisms

Names

Weights

Options

GNPS (optional)

Organisms-related parameters

❓

OPTIONAL. Force all features to be 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

❓

Streptophyta

Save parameters   Launch job