Biogeochemical Model Database bgc_md2

















Markus Müller, Holger Metzler, Verónika Ceballos Núñez, Kostiantyn Viatkin, Thomas Lotze, Jon Wells, Yu Zhou, Cuijuan Liao, Aneesh Chandel, Feng Tao, Yuanyuan Huang, Alison Bennett, Chenyu Bian, Lifen Jiang, Song Wang, Chengcheng Gang, Carlos Sierra, Yiqi Luo

Why would **you** want a model data base?









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Why would you want a model data base?

Find (rather than reinvent) the right model for a specific task?







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- **Implement** a **new** model but start from a similar one?







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Find and reduce sources of uncertainty in carbon predictions









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- **...** ???











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How can you build one?

We need:

collections







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How can you build one?

We need:

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 - of many models: (from A to Z):

Arora2005GCB-1 .CARDAMOM

Zelenev2000MicrobialEcology



(a) (b) NAU (b) (b) (c) (c) (c) (d)

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Biogeochemical Model Database bgc_md2 (a) (b) NAU (b) (b) (c) (c) (c) (d) (d)













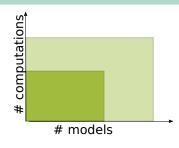


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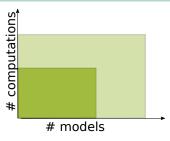


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- 'Copy and paste'?
 - → bilinear increase of code
 - \rightarrow unmaintainable, untestable,
 - errorprone, hard to change ...









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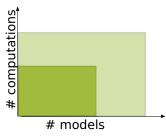
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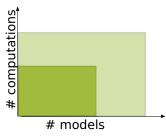
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Reproducible Carbon Cycle Models Biogeochemical Model Database bgc_md2 (a) (b) NAU (b) (b) (c) (c) (c) (d)

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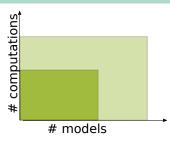
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- of many computable properties and diagnostics to compare them by
- Ways of **organizing** both collections



- 'Copy and paste'?
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 - \rightarrow unmaintainable, untestable, errorprone, hard to change ...
- better use the **dry** principle Don't repeat yourself











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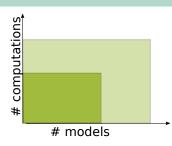
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- Ways of **organizing** both collections
 - building blocks for models



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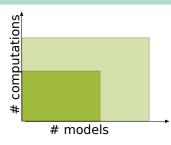
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- of many computable properties and diagnostics to compare them by
- Ways of **organizing** both collections
 - building blocks for models
 - functions of building blocks



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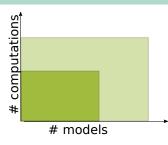
How can you build one?

We need:

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Zelenev2000MicrobialEcology of many computable

- properties and diagnostics to compare them by
- Ways of **organizing** both collections
 - building blocks for models
 - functions of building blocks
 - computational graph for **different** / **evolving** building blocks



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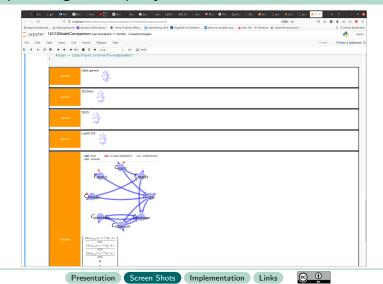
Reproducible Carbon Cycle Models

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Reproducible Carbon Cycle Models

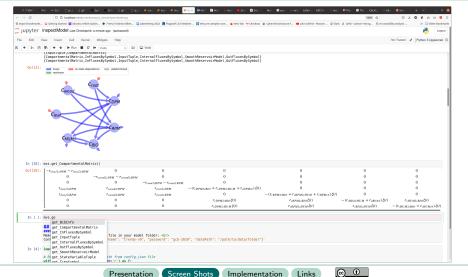
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Example widget for query result



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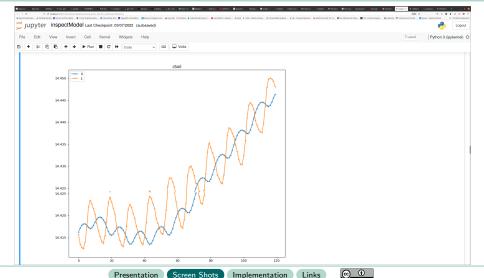
Analysis with symbolic tools (sympy) ...



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...or numerically



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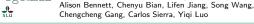












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Database records are python modules

```
from ComputabilityGraphs.ORTVS import ORTVS
from bgc_md2.helper import module_computers
from bgc_md2.models.BibInfo import BibInfo
              InFluxesBySymbol,
OutFluxesBySymbol,
InternalFluxesBySymbol,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         code*k+" = Function("[8]")".format(k)
import bac md2.resolve.computers as bac o
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       66 t=TimeSymbol("t")
67 beta_root = 1.0- (beta_leaf+beta_wood)
68 mms = OMTVS/
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C_wood_litter: r_C_wood_litter_rhfC_wood_litter*xi(t),
C_root_litter: r_C_root_litter_rhfC_root_litter*xi(t),
C_woil_fast: r_C_woil_fast_rhfC_woil_fast*xi(t),
C_woil_fast: r_C_woil_fast_rhfC_woil_fast*xi(t),
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       code=k+" = Function('(0)')".formet(k)
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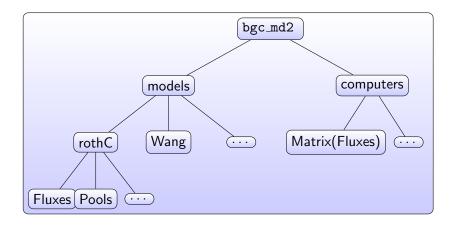


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Internal Structure of bgc_md2









The bgc_md library provides I:

- Datatypes defining building blocks of models e.g. CompartmentalMatrix, InternalFluxesBySymbol, ...
- Functions operating on those properties (forming the edges of the graph where the Datatypes are nodes)
- A user interface based on graph algorithms to
 - compute the set of computable properties (e.g. the comparable criteria for a set of models, database queries)
 - actually compute the desired properties by recursively connecting several function applications.
 - show what is missing to compute a desired property.



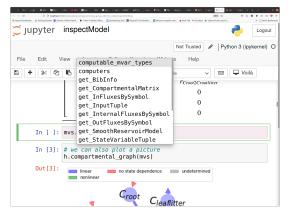






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Userinterface using computability graphs



Suggested methods automatically created by ComputabilityGraphs







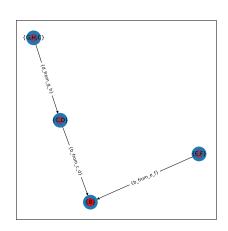




Biogeochemical Model Database bgc_md2

Finding what's missing in the model description

given a set of functions: a(i), b(c,d), b(e,f),c(b), d(b), d(g,h),e(b), f(b) and the target variable B e.g. CompartmentalMatrix, The algorithm computes all possible combinations and paths from which B can be computed.











The bgc_md library provides II:

- 30+ vegetation, soil or ecosystem models for carbon and nitrogen cycling as reusable python modules using the building blocks in a flexible way.
- An interface to many algorithms in CompartmentalSystems to compute diagnostic variables for many models in bgc_md2.



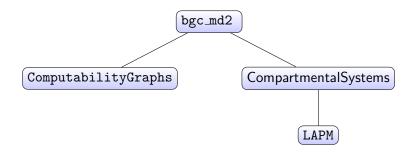




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Relation to other Python Packages









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Example computation via CompartmentalSystems

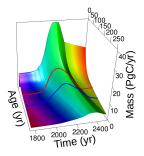


Figure: age distribution of a pool as function of time



Metzler, H., Müller, M., and Sierra, C. (2018).

Transit-time and age distributions for nonlinear time-dependent compartmental systems.











Links

- The README of the package on github (with installation instructions): https://github.com/MPIBGC-TEE/bgc_md2
- To explore some rudimentary tutorials without installation use https://mybinder.org/v2/gh/MPIBGC-TEE/bgc_md2/binder
 - Click on the link!
 - After jupyter lab has started go to
 /binder_notebooks/illustrativeExamples/
 - right click on createModel.py
 - choose Open With
 - choose Jupytext Notebook

This will open an example notebook exploring some of the concepts. More applied examples are coming.







