

Building a process-based model, from scratch

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Outline

- PART 1

- What is a process-based model?
- Why do we want process-based models?
- Building a model from scratch
- Modelling lake metabolism rates

- PART 2

- Step-by-step model implementation using R, GitHub and Binder

What is a process-based model?

- A representation of the way the world works
- Reproduces a level of realism
- However, models are necessary simplifications and do not explain all the variance in observational data
- A process-based model is a model that takes an object-oriented perspective, identifying:
 - the relevant entities (included properties)
 - their state
 - and their behaviours.

Why process-based models?

- Can be used for scenarios that are out of sample
- To challenge our level of understanding of the way our world works
 - From system A to system B
- Accommodates a variety of data, even if sampled irregularly
- Used to infill missing data

Building a model from scratch

- What do I want to model?
 - State variables
- What do I know about my state variables?
 - State
 - Behaviour = processes (= equations)
 - Boundary conditions
- What data (observations) do I need?
- How do I want to build my model?
 - Numerical scheme
 - Programming language
 - Numerical platform
- Let's go!

Application




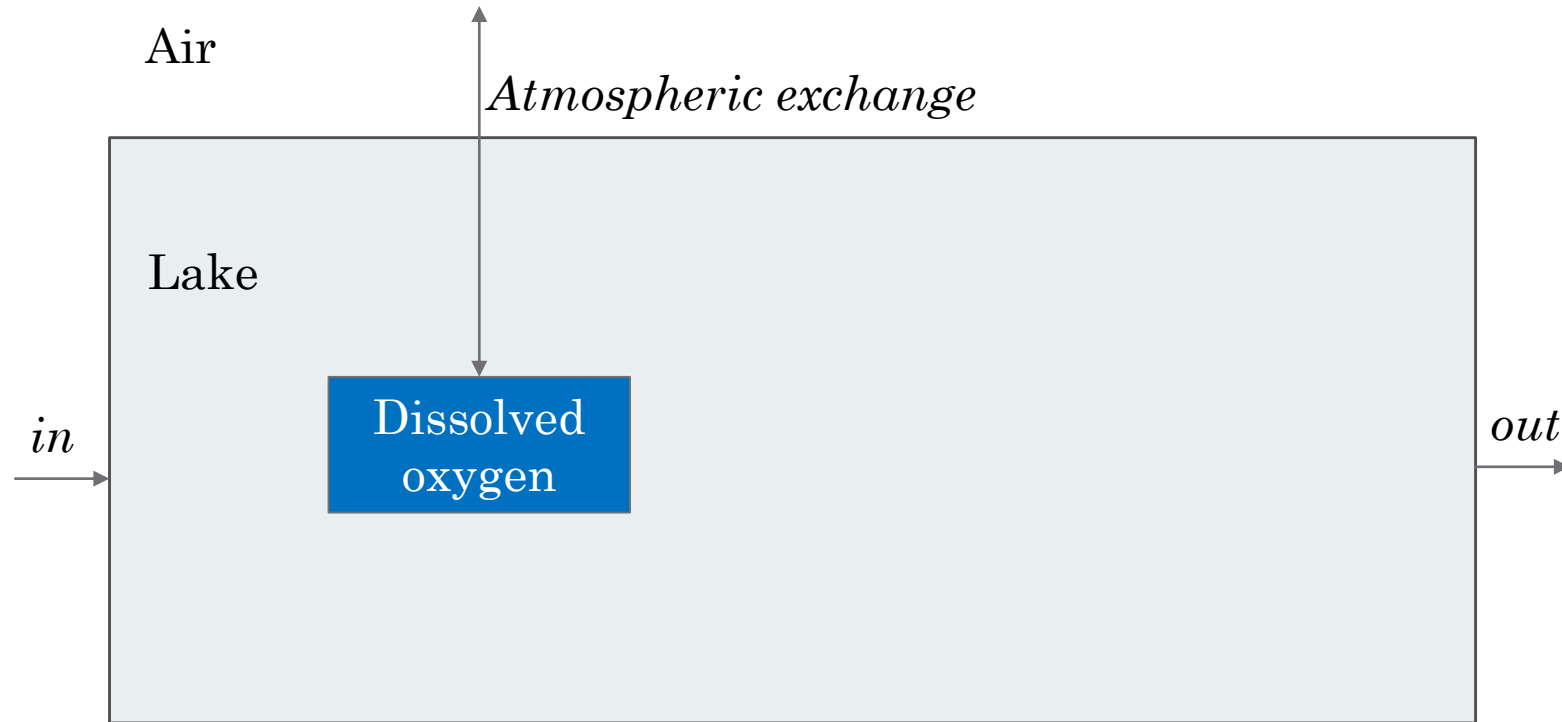
Modelling lake metabolism rates

Air

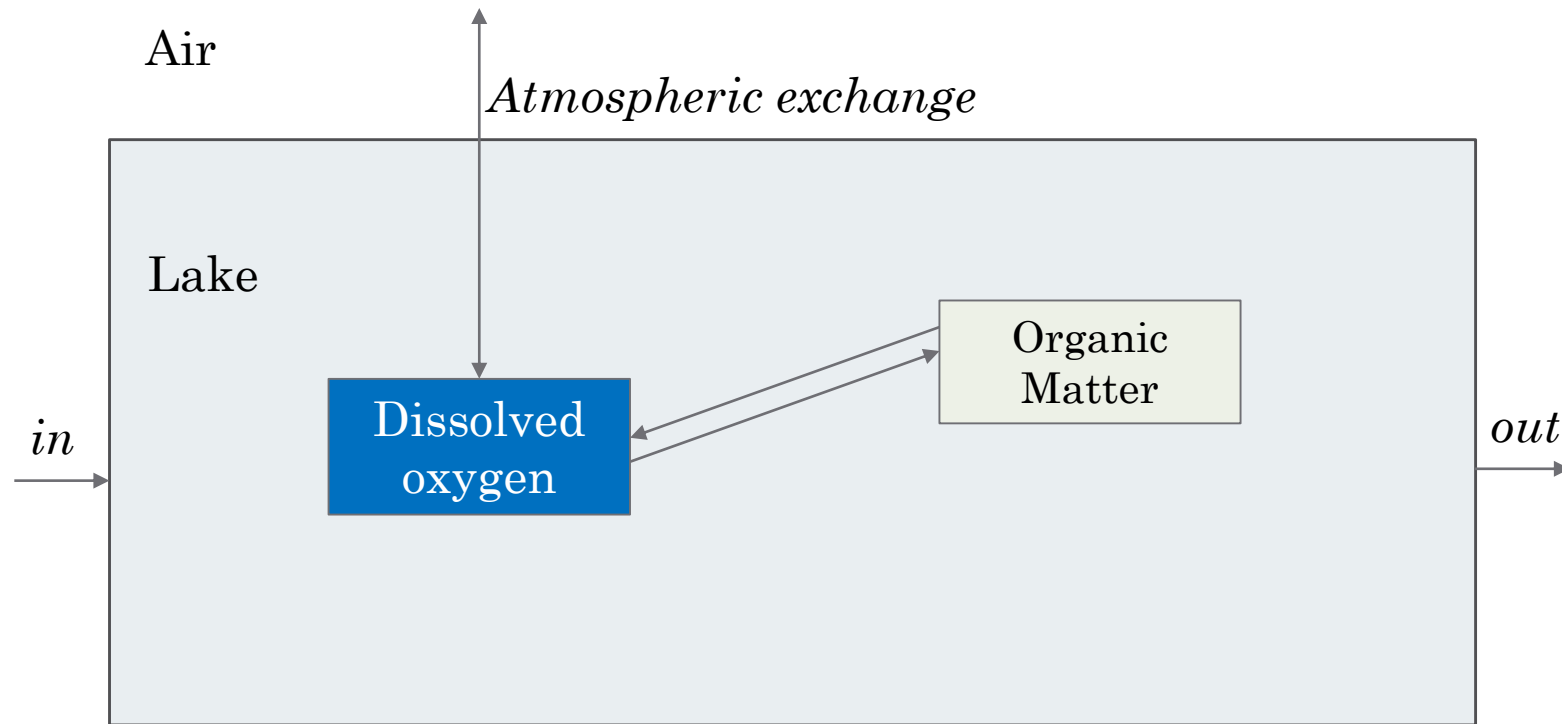
Lake

Dissolved
oxygen

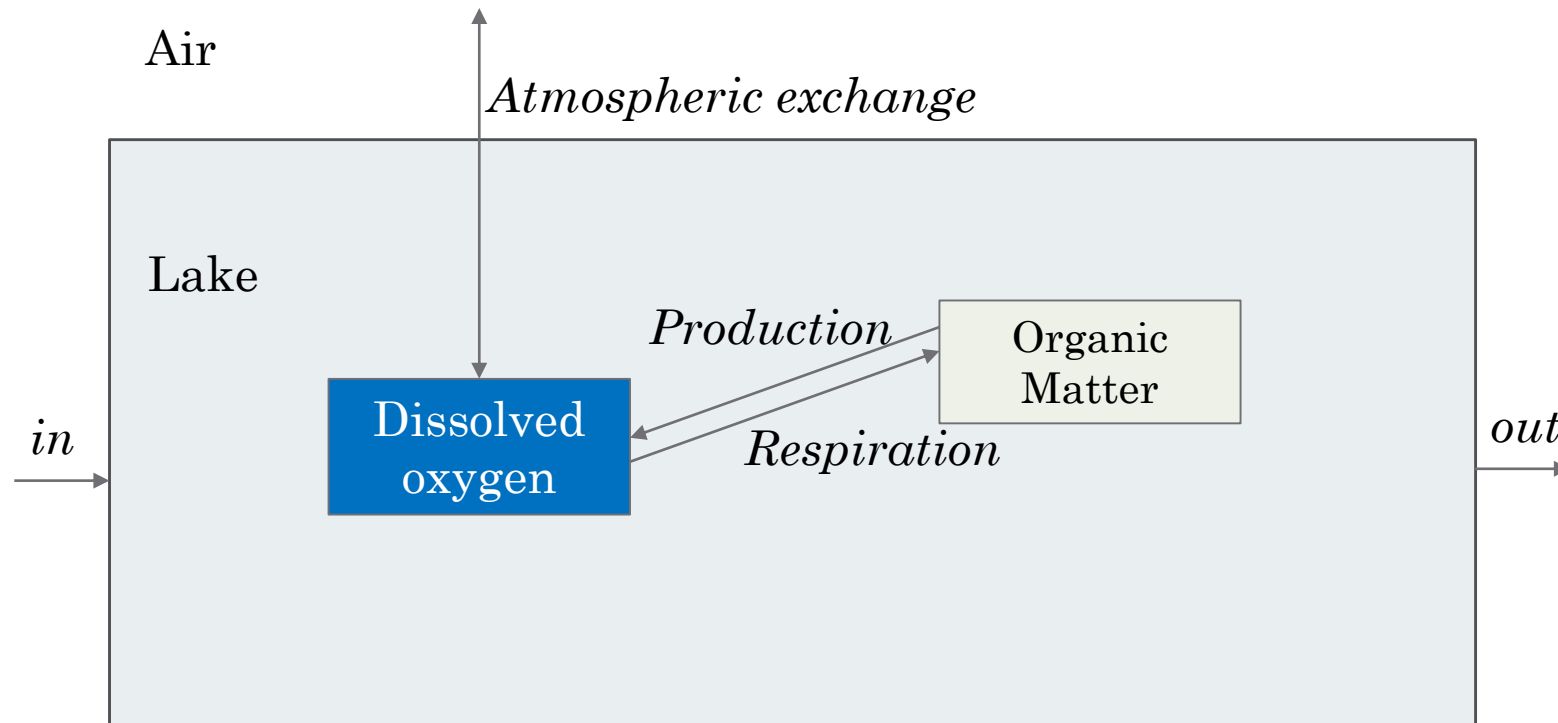
A diagram illustrating the concept of dissolved oxygen in a lake. It features a large, light blue rectangular area representing the lake. Inside this area, on the left side, is a smaller, solid blue rectangle. Within this blue rectangle, the words "Dissolved" and "oxygen" are written in white, stacked vertically. The word "Lake" is written in black text to the left of the large blue area, and the word "Air" is written in black text above the top-left corner of the large blue area.



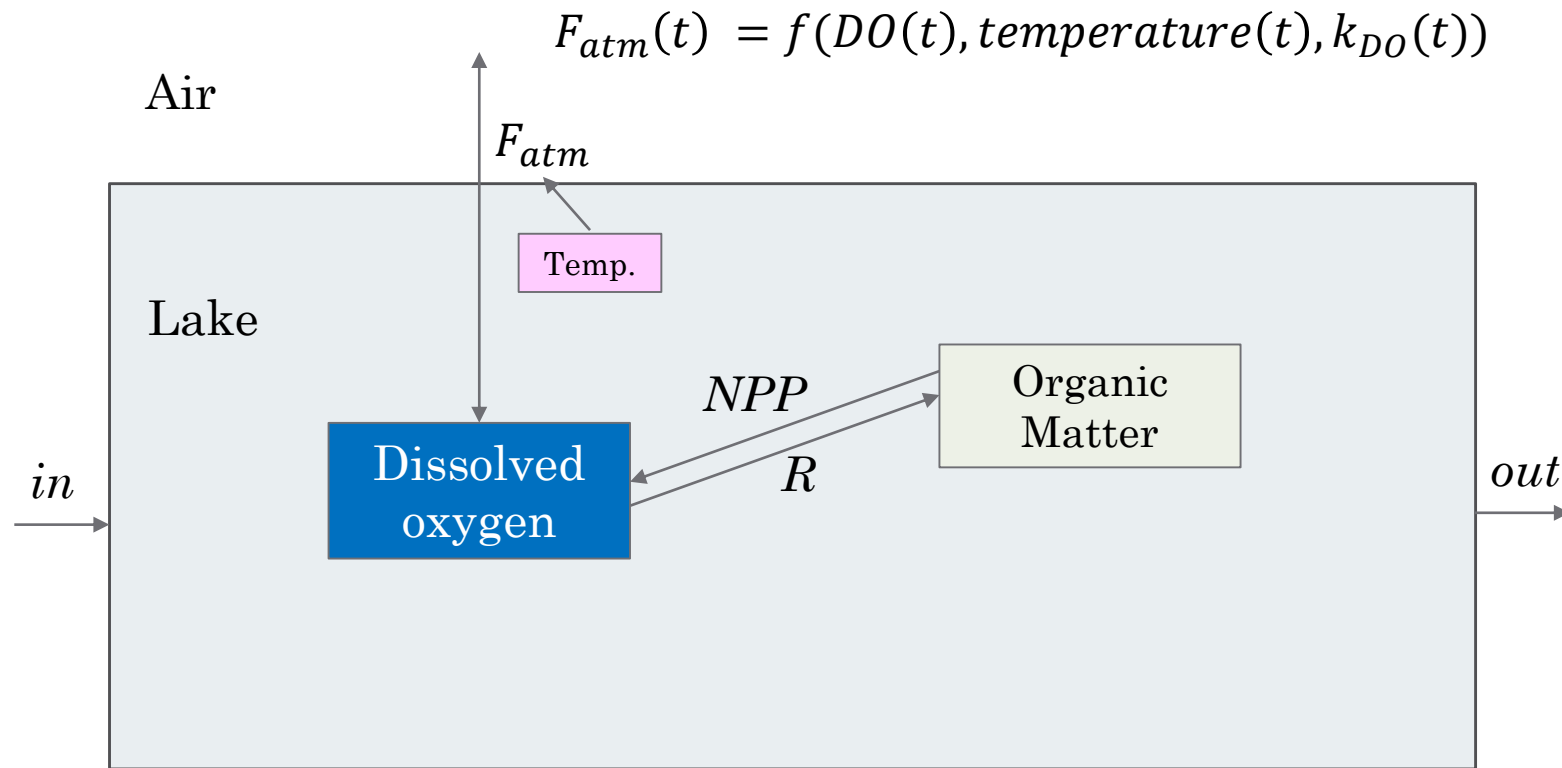
$$\frac{dDO(t)}{dt} = f(\text{Physical}_{fluxes}(t))$$



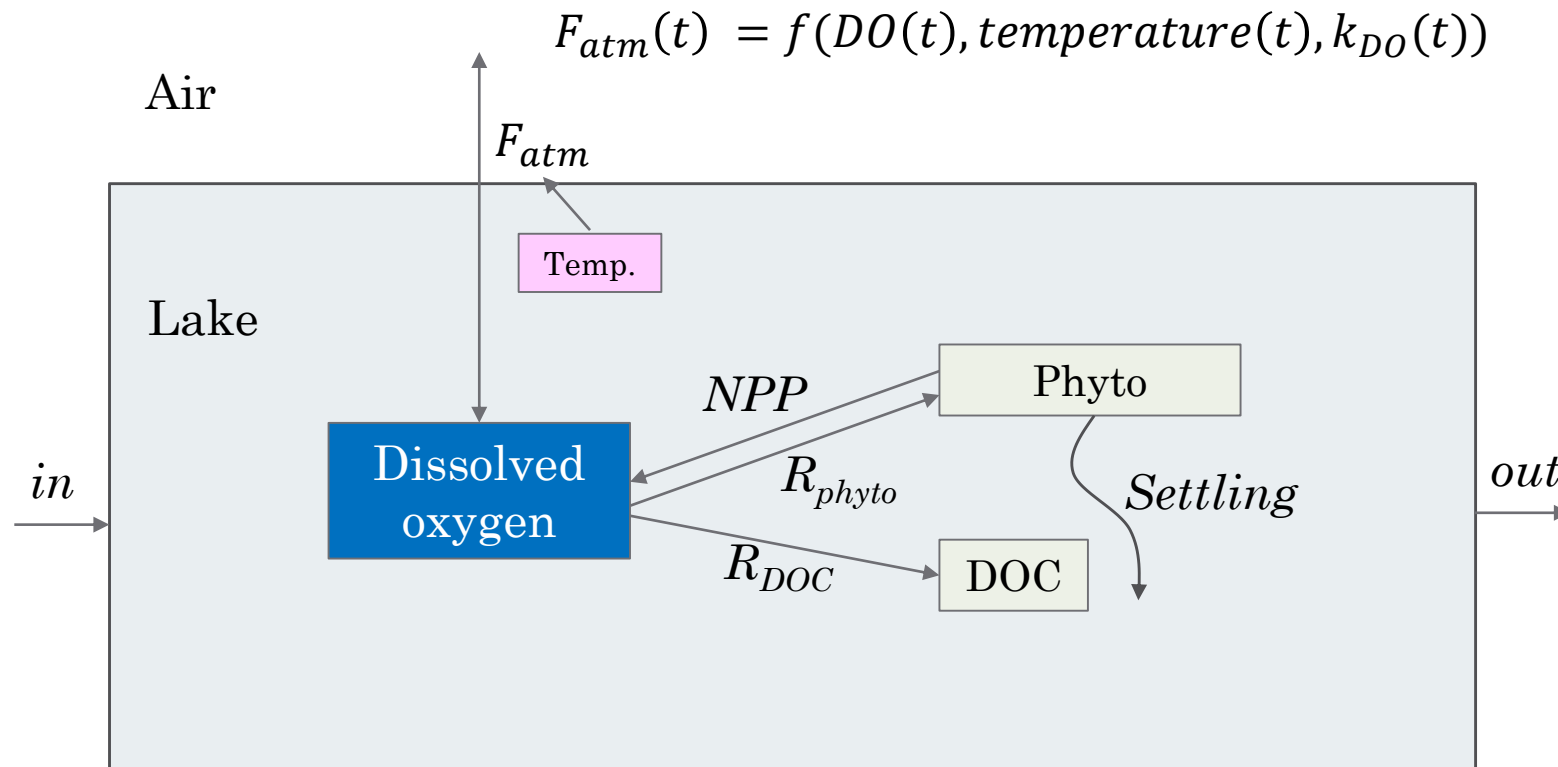
$$\frac{dDO(t)}{dt} = f(\text{Organic Matter}, \text{Physical}_{fluxes}(t))$$



$$\frac{dDO(t)}{dt} = Production(t) - Respiration(t) + Physical_{fluxes}(t)$$



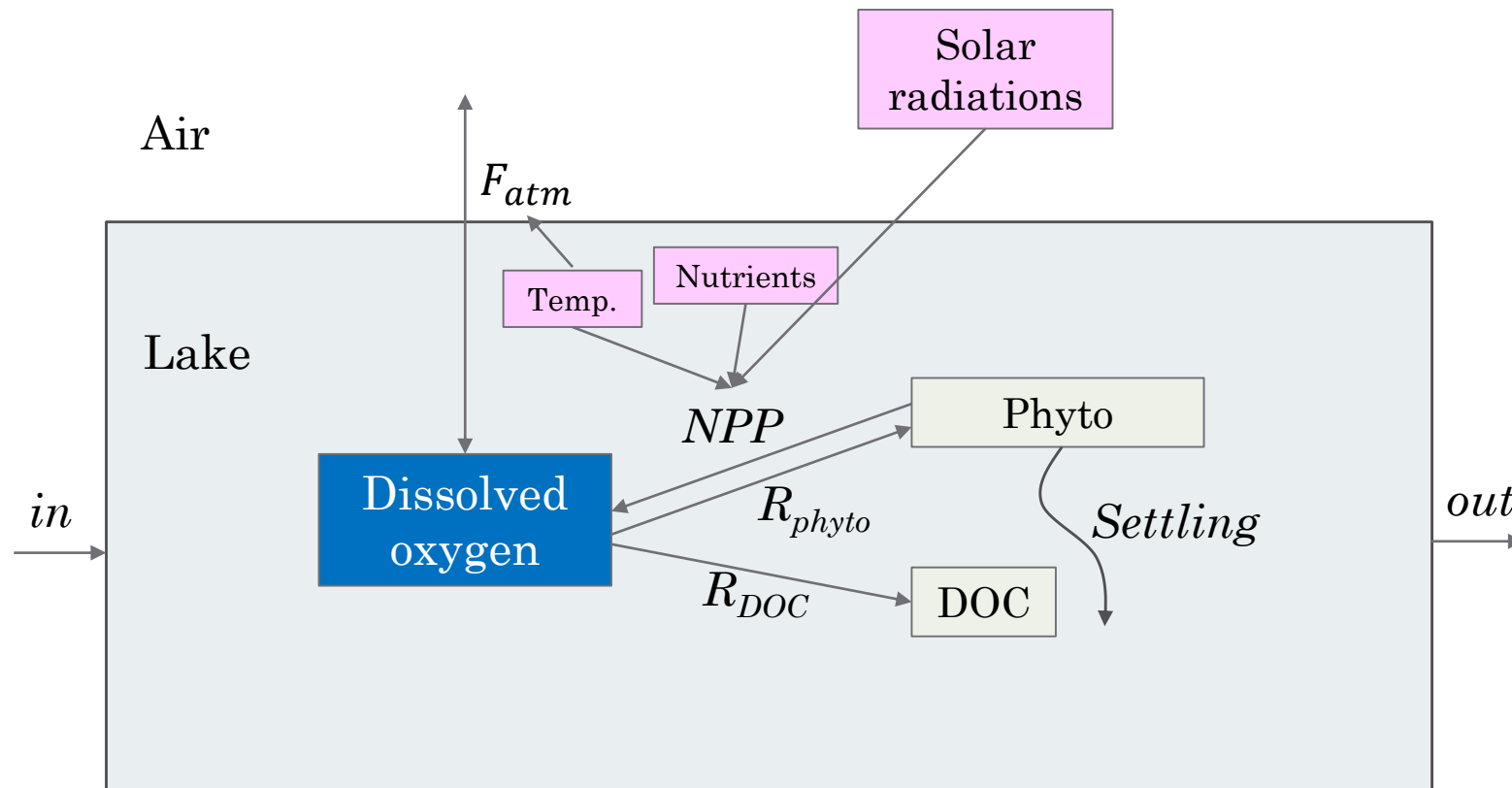
$$\frac{dDO(t)}{dt} = NPP(t) - R(t) + F_{atm}(t) + \varphi(t)$$



$$\frac{dDO(t)}{dt} = NPP(t) - R(t) + F_{atm}(t) + \varphi(t)$$

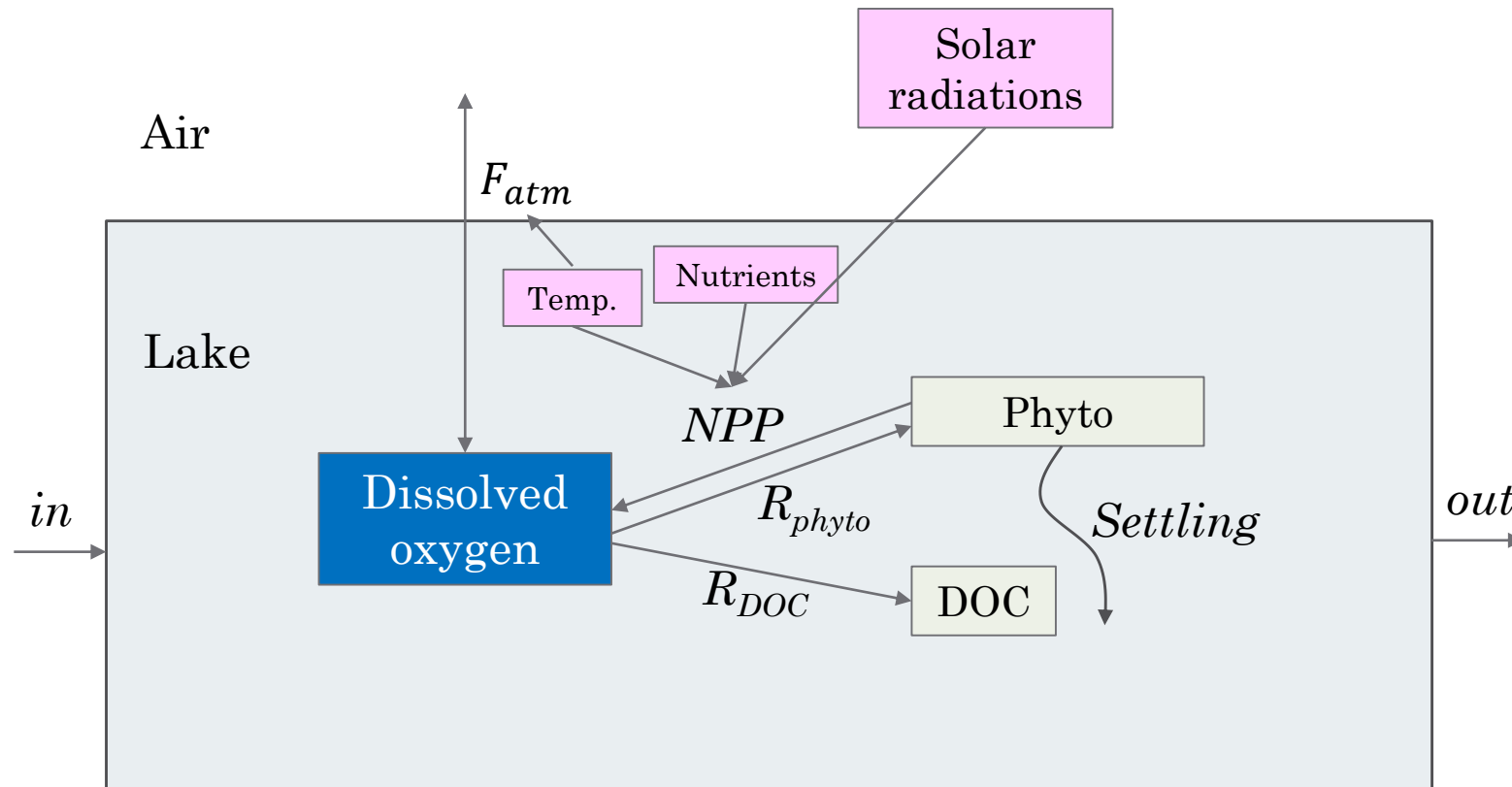
$$R(t) = R_{phyto}(t) + R_{DOC}(t)$$

$$\frac{dPhyto(t)}{dt} = NPP(t) - R_{phyto}(t) - Settling(t)$$



$$\frac{dDO(t)}{dt} = NPP(t) - (R_{phyto}(t) + R_{DOC}(t)) + F_{atm}(t) + \varphi(t)$$

$$NPP(t) = f(Radiations(t), [Nutrients](t), temperature(t))$$

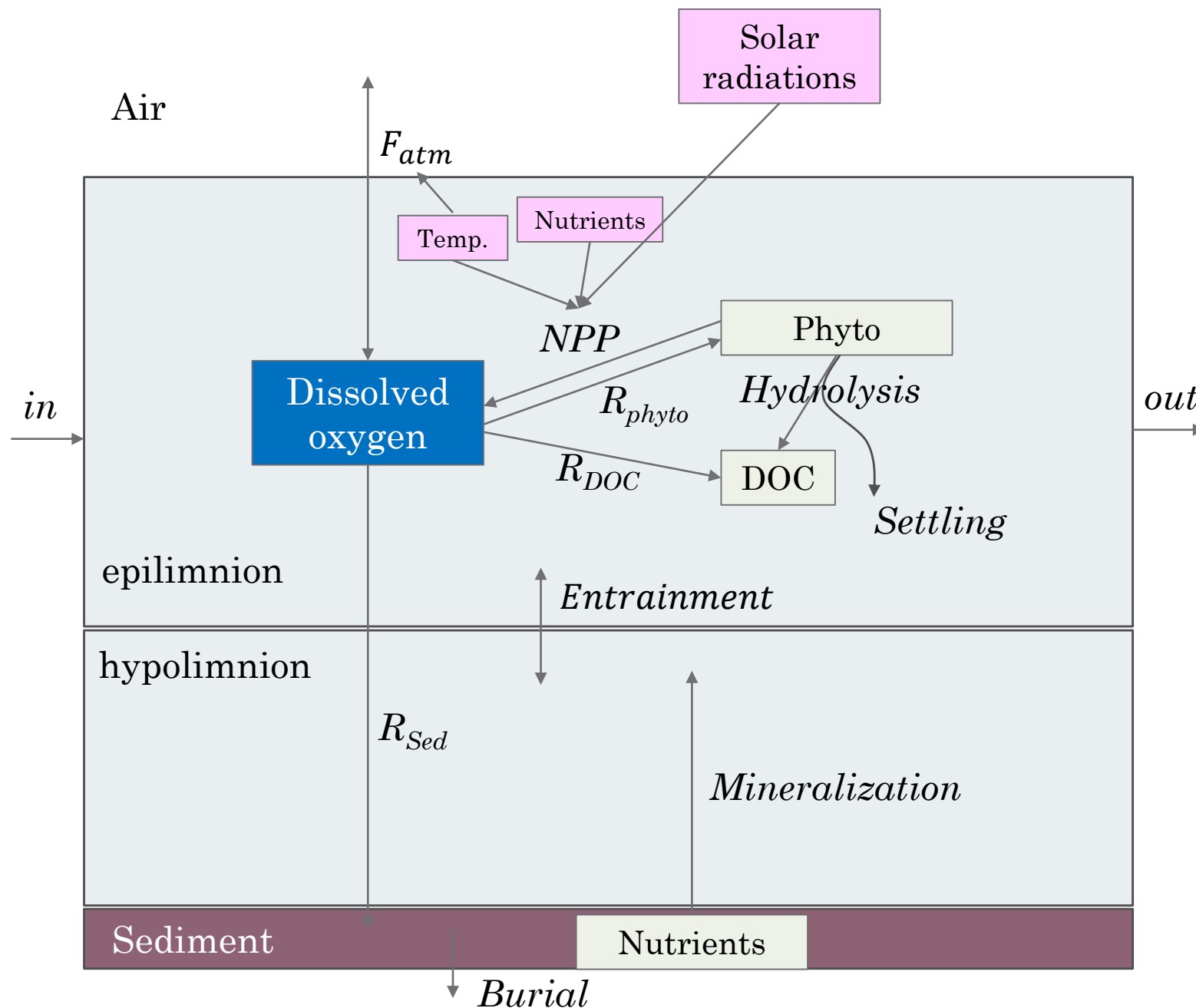


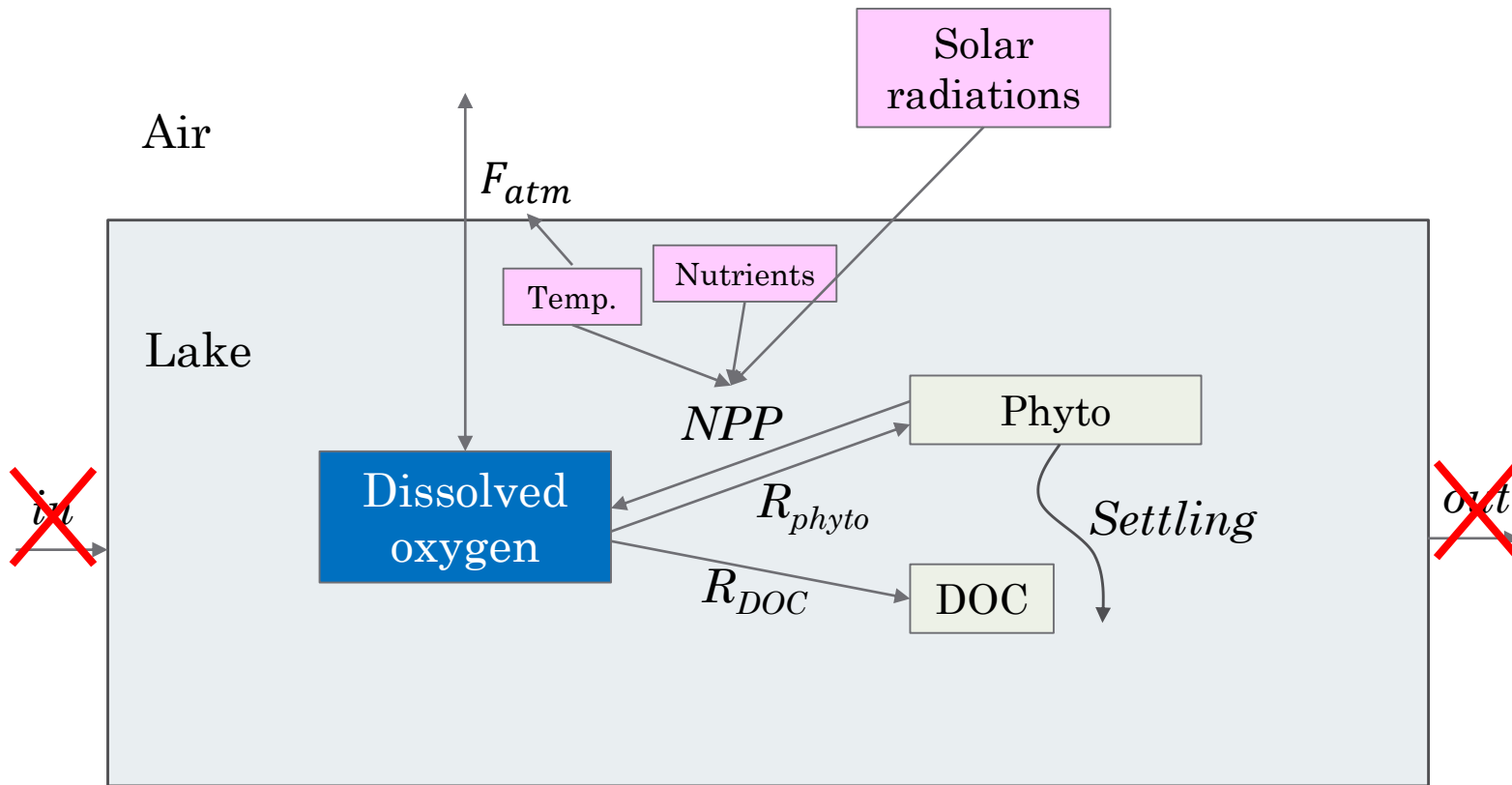
$$\frac{dDO(t)}{dt} = NPP(t) - (R_{phyto}(t) + R_{DOC}(t)) + F_{atm}(t) + \varphi(t)$$

$$NPP(t) = f(Radiations(t), [Nutrients](t), temperature(t))$$

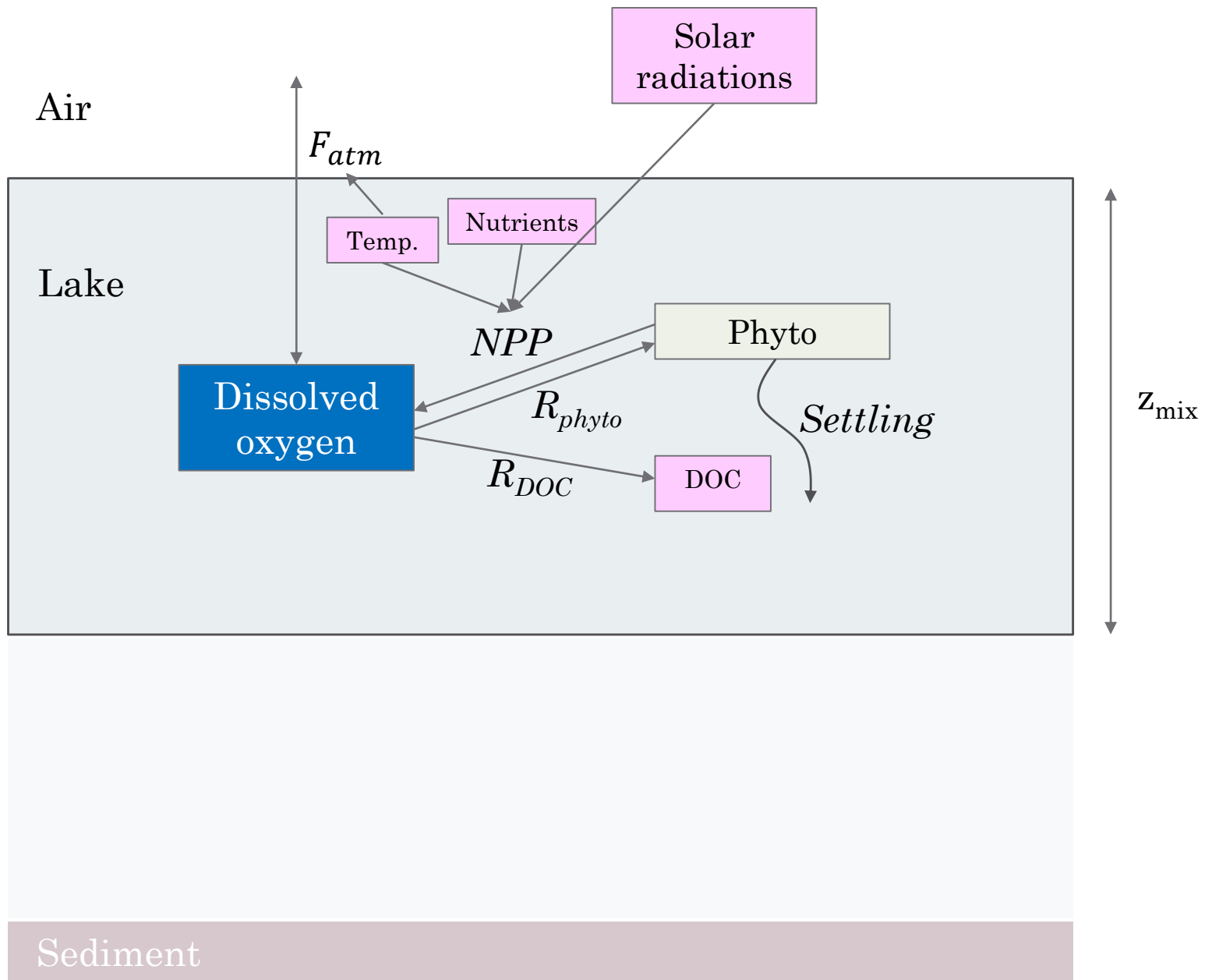
$$R_{DOC}(t) = f([DOC](t))$$

$$Settling(t) = f(z_{mix}(t), Phyto(t))$$





$$\frac{dDO(t)}{dt} = NPP(t) - (R_{phyto}(t) + R_{DOC}(t)) + F_{atm}(t) + \cancel{\varphi(t)}$$



In discrete time form

$$\frac{dDO(t)}{dt} = NPP(t) - (R_{phyto}(t) + R_{DOC}(t)) + F_{atm}(t)$$

$$DO(t) = DO(t-1) + NPP(t) \cdot k_{C \rightarrow O_2} - (R_{phyto}(t) + R_{DOC}(t)) \cdot k_{C \rightarrow O_2} + F_{atm}(t)$$

Oxygen at the current time step (t), is equal to what it was at the previous time step ($t-1$), plus the mass movement of oxygen through a set of processes : gross primary production, respiration, atmospheric exchange and physics.

In a R code:

```
doPredic[i] = doPredic[i-1] + npp[i]*CtoO2 - Rtot[i]*CtoO2 + fatm[i]
```

In discrete time form

$$\frac{dPhyto(t)}{dt} = NPP(t) - R_{phyto}(t) - Settling(t)$$

$$phyto(t) = phyto(t - 1) + NPP(t) - R_{phyto}(t) - Settling(i)$$

Phytoplankton biomass at the current time step (t), is equal to what it was at the previous time step ($t-1$), plus what is being produced through photosynthesis, minus what is lost by respiration and settling

In a R code:

```
phyto[i] = phyto[i-1] + npp[i] - Rphyto[i] - Settling[i]
```

Additional equations

$$F_{atm}(t) = dt \cdot k_{DO}(DO_{sat}(t) - DO(t - 1))/z_{mix}$$

$$DO_{sat}(t) = -6 \cdot 10^{-5} \cdot wTemp(t)^3 + 69 \cdot 10^{-4} \cdot wTemp(t)^2 - 0.3906 \cdot wTemp(t) + 14.578$$

$$Settling(t) = dt \cdot phyto(t - 1) \frac{settling_{phyto}}{z_{mix}}$$

$$NPP(t) = dt \cdot PAR(t) \cdot phosphorus \cdot pNPP \cdot \theta_{NPP}^{(wTemp(t)-20)}$$

$$R_{DOC}(t) = dt \cdot DOC \cdot docR$$

$$R_{phyto}(t) = dt \cdot phyto(t - 1) \cdot phytoR \cdot \theta_R^{(wTemp(t)-20)}$$

$$R(t) = R_{DOC}(t) + R_{phyto}(t)$$

These equations (in a R code)

```
fatm[i] = dt * kD0 * (dosat[i] - doPredic[i-1]) / zMix
```

```
dosat[i] = -0.00006 * wTemp[i]^3 + 0.0069 * wTemp[i]^2  
          - 0.3906 * wTemp[i] + 14.578
```

```
Settling[i] = dt * phyto[i-1] * settlingPhyto/zMix
```

```
npp[i] = dt * PAR[i] * phosphorus * pNPP *  
         thetaNPP^(wTemp[i] - 20)
```

```
RDOC[i] = dt * DOC * docR
```

```
Rphyto[i] = dt * phyto[i-1] * phytoR * thetaR^(wTemp[i] - 20)
```

```
Rtot[i] = RDOC[i] + Rphyto[i]
```

Constants

Constant name	value	units	Definition
$k_{C \rightarrow O_2}$	32/12	d.l.	Molar ratio between C and O ₂
k_{DO}	1	m/d	Gas exchange coefficient, usually computed from wind speed
z_{mix}	10	m	Depth of mixed layer (or epilimnion)
$settling_{phyto}$	1	m/d	Settling velocity of phytoplankton cells
$phosphorus$	0.050	µgP/L	Phosphorus concentration
$pNPP$	0.08	mgC/unitP/unitLight	Conversion factor
θ_{NPP}	1.2	d.l.	Arrhenius coefficient for temperature adjustment for NPP
DOC	5	mg/L	Dissolved organic C concentration
$docR$	0.02	d.l.	1 st order decay of DOC
$phytoR$	0.8	d.l.	1 st order decay of phytoplankton
θ_R	1.08	d.l.	Arrhenius coefficient for temperature adjustment for R

Demo

<https://github.com/camilleminaudo/oxygen-model-student>

Part 2

Model implementation

You have RStudio installed



<https://github.com/camilleminaudo/oxygen-model-students>

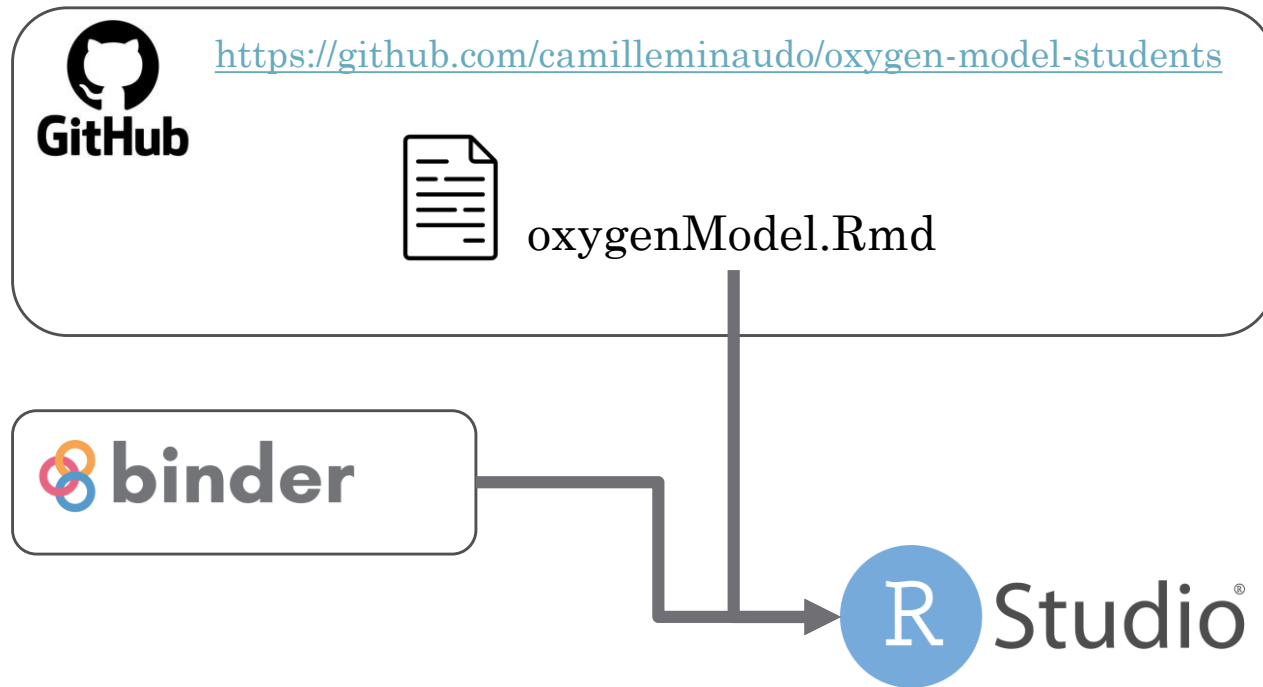
- Option A: download the entire repository
- Option B: clone to this repository



oxygenModel.Rmd



You don't have RStudio installed



The model, step by step

- (Create a GitHub account)
- Go to <https://github.com/camilleminaudo/oxygen-model-students>
- Click on “lauch binder” in the README.md file
- Wait for installation and Jupyter install
- Open a Rstudio notebook
- Open the file oxygenModel.Rmd
- Follow instructions and run the model

