

Construction of a Machine Learning tool based on Differential Equations integrated into a dynamic tree allowing biological data assimilation

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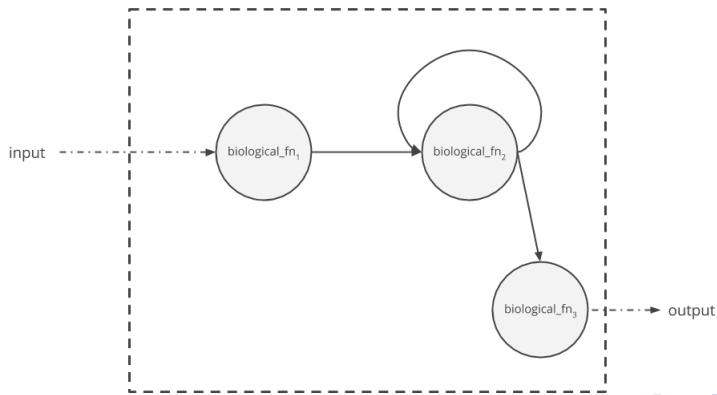
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- Modeling of the digestive behaviour
- The goal: build Machine Learning algorithms based on the discretization of Differential Equations in order to predict biological responses and assimilate biological data.

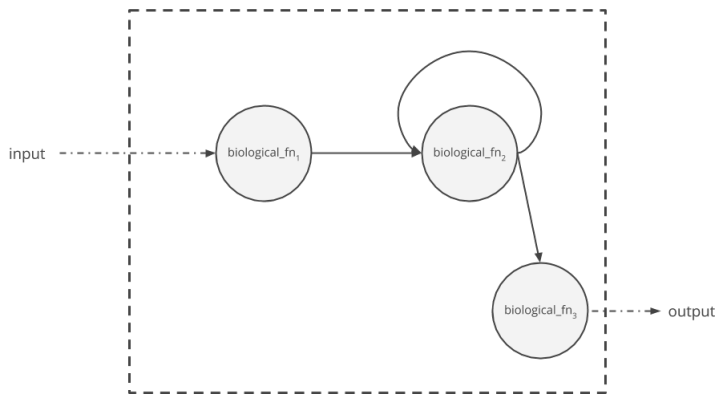
Graph-oriented Model

- The main idea: Assimilate an animal to a graph in which nodes are virtual organs and edges represent fluxes within the organism.



Graph-oriented Model

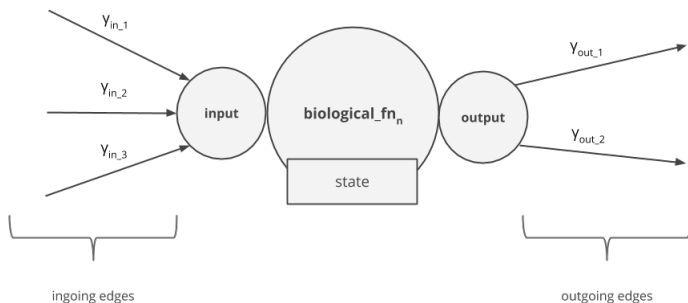
- Each virtual organ carry a biological function, which aimed to reproduce at best its actual behaviour.



Virtual animal

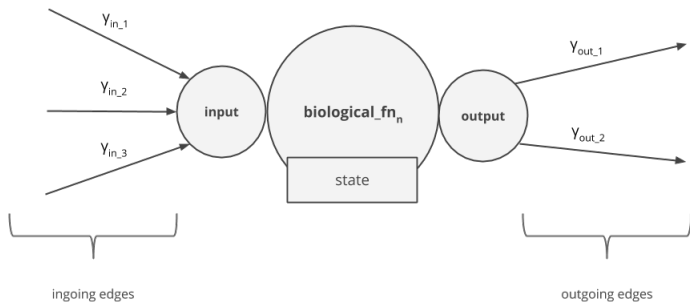
Graph-oriented Model

- Each link is weighted with a parameter γ_i , on which indicates how inputs and outputs are actually dispatched, and to give more or less significance on links between organs.



Graph-oriented Model

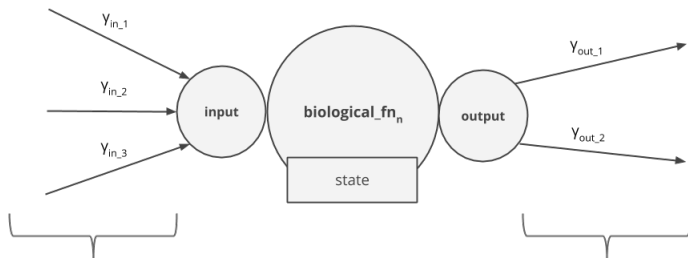
- Each node (organ) is given some memory, which are states. They can thus store and use several values, which can be part of their function definition.



Graph-oriented Model

The process of such function is as follows:

- 1) Aggregate all the ingoing fluxes coming from other functions
- 2) Given those inputs, compute the function output for this time step.
- 3) Propagate the computed output to all outgoing edges.



Global Optimization

Optimize the whole animal in order to find the parameters values that would make it fit at best the real data.

2 approaches:

- 1) Analytic approach: Analytically resolve the functions, by fixing some parameters each by each.

Global Optimization

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2 approaches:

- 1) Analytic approach: Analytically resolve the functions, by fixing some parameters each by each.
- 2) Machine Learning approach: Optimization (Genetic optimization methods, Least square method...)
⇒ Fast to implement.

Applied case: Ross Chicken growth



ROSS 708

<http://en.aviagen.com/brands/ross/products/ross-708>

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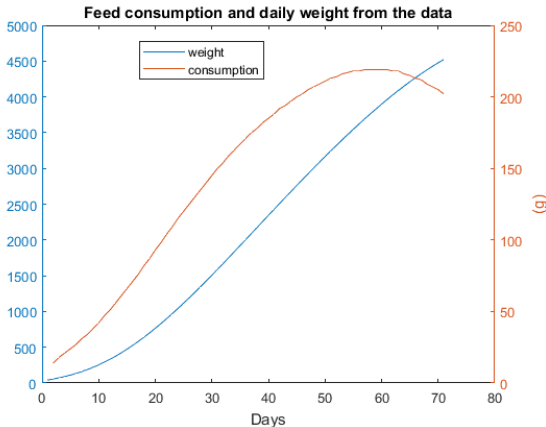
Some information on those animals are available, for each sex of the strains:

- Daily consumption of food.
- Daily growth (Weight).



ROSS 708

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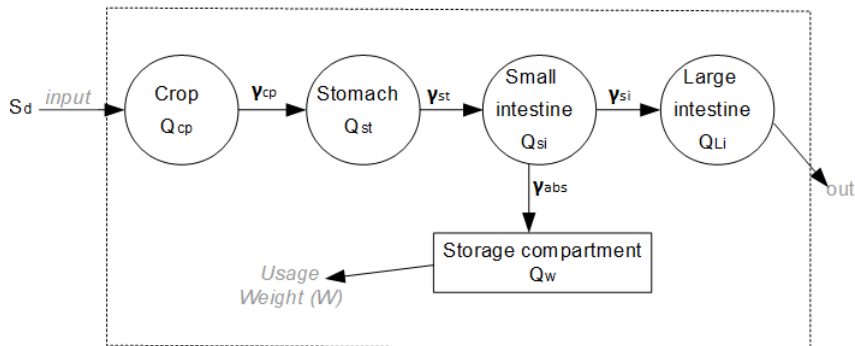


- At first, we consider the growth(mass gained each day) .
- Flux to consider is the food consumption which becomes later a nutrients flux.

Recall of some growth functions

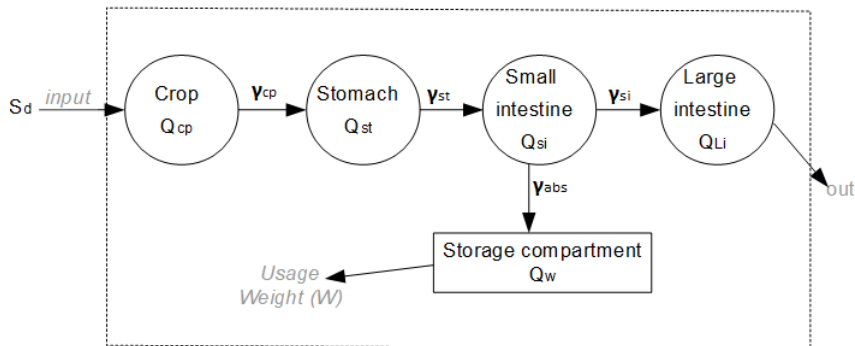
- Logistic: $W' = \mu_0 W \left(1 - \frac{W}{W_f}\right)$
- Gompertz: $W' = DW \ln \left(\frac{W_f}{W}\right)$
- Richards: $W' = DW \left(\frac{W_f^n - W^n}{nW_f^n}\right)$

Assumptions



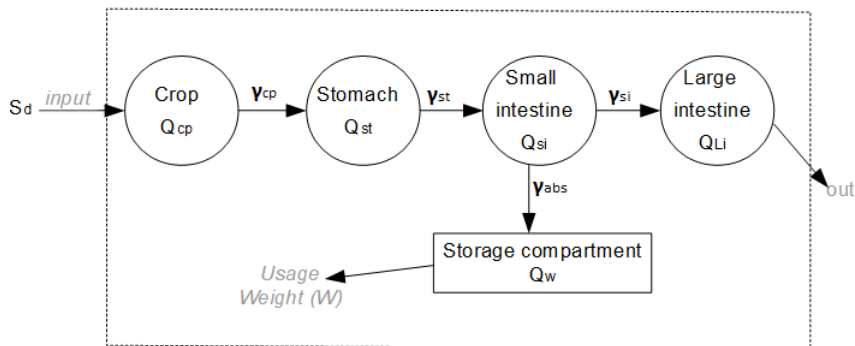
- A1 Ingested food is stored in the crop before moving into the stomach.

Assumptions



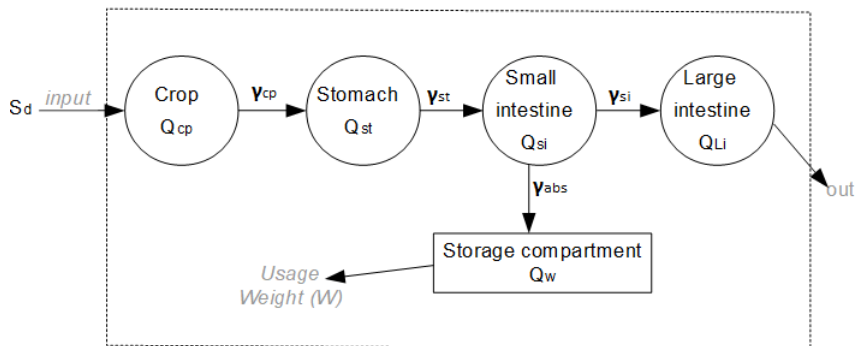
- A1 Ingested food is stored in the crop before moving into the stomach.
- A2 Some fluxes rates are constant between compartments.

Assumptions



A3 Existence of transmission delays in some compartments.

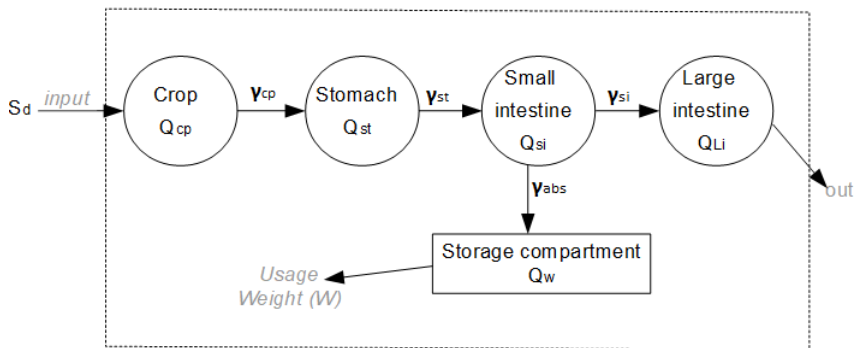
Assumptions



A3 Existence of transmission delays in some compartments.

A4 There exists a time τ for which the SI emptying dynamic starts.

Equations

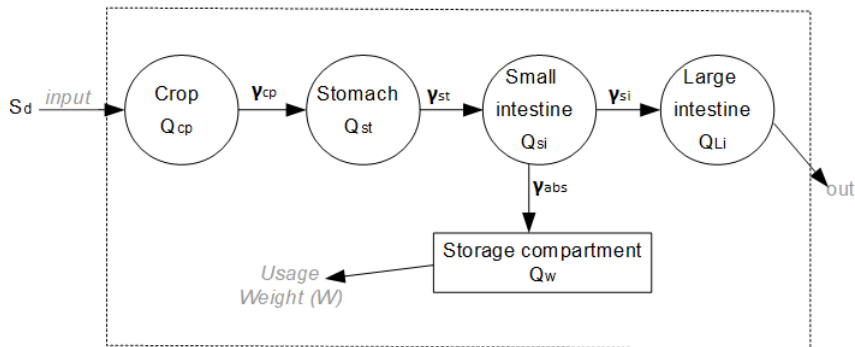


Crop

$$Q'_{cp,d} = -\gamma_{cp} Q_{cp,d}$$

$$Q_{cd,d}(0) = S_d$$

Equations

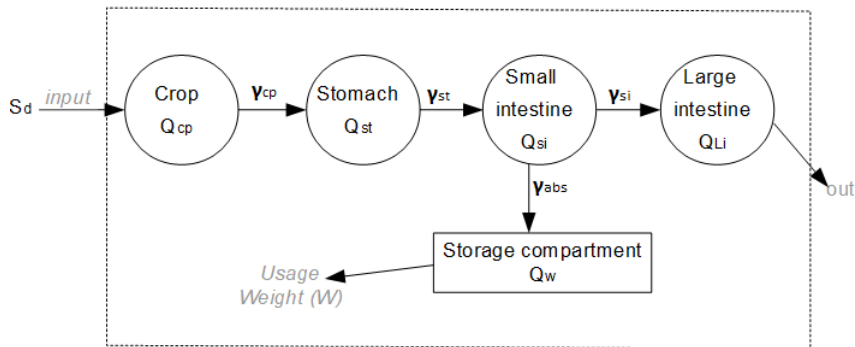


Stomach

$$Q'_{st,d} = \gamma_{cp} (1 - \exp(-\beta_1 t)) Q_{cp,d} - \gamma_{st} Q_{st,d}$$

$$Q_{st,d}(0) = 0.$$

Equations

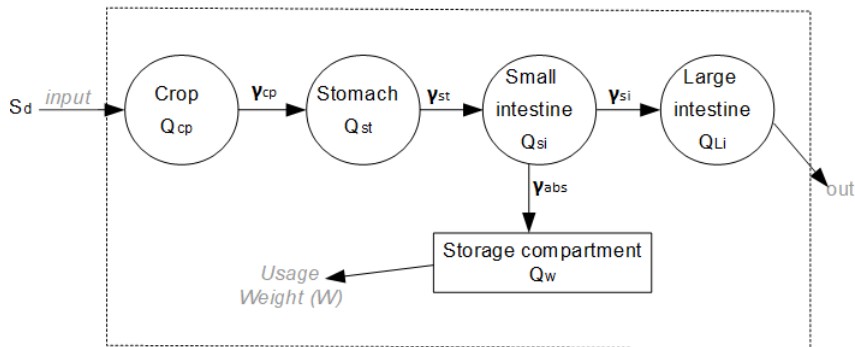


Small intestine

$$Q'_{si,d} = \begin{cases} \gamma_{st} Q_{st,d}, & t \leq \tau \\ \gamma_{st} Q_{st,d} - (\gamma_{abs} + \gamma_{si}) Q_{si,d}, & t > \tau \end{cases}$$

$$Q_{si,d}(0) = 0.$$

Equations



Large intestine

$$Q'_{Li,d} = \gamma_{si} Q_{Li,d} - \gamma_{Li} Q_{Li,d}$$
$$Q_{Li,d}(0) = 0.$$

Equations

Storage and usage of nutrients

$$Q'_{w,d} = \gamma_{abs} Q_{si,d} - \alpha Q_{w,d}$$

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Formation of the daily weight

$$W'_d = \kappa(t) Q_{w,d} - \left(V_{max} \frac{Q_{w,d}}{K + Q_{w,d}} \right) \frac{1}{K} W_d$$

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$$\text{Final weight } W_f = W_0 + \sum_{d=1}^{N_d} W_d(t_f)$$

N_d : number of days.

Simulations

- Observed data
 - Daily feed intake.
 - Daily weight measurement.
- Discretization: FDM(Forwards Euler method);
- fitting: γ_{cp} and τ fixed.

Estimate $\beta_1, \gamma_{st}, \gamma_{abs}, \gamma_{si}, \gamma_{Li}, \kappa_0, \mu_{max}, \beta_2, V_{max}, K$

Simulations

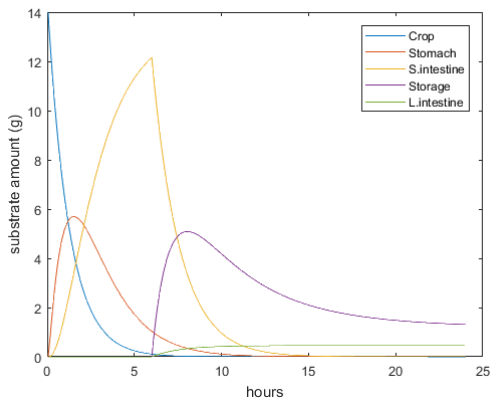


Figure: One day dynamic, $\beta_1 = 0.4$, $\gamma_{st} = 0.6$, $\gamma_{abs} = 0.6$, $\gamma_{si} = 0.023$, $\kappa_0 = 0.9$, $\mu_{max} = 0.01$, $\beta_2 = 0.01$, $V_{max} = 0.07$, $K = 26.8$

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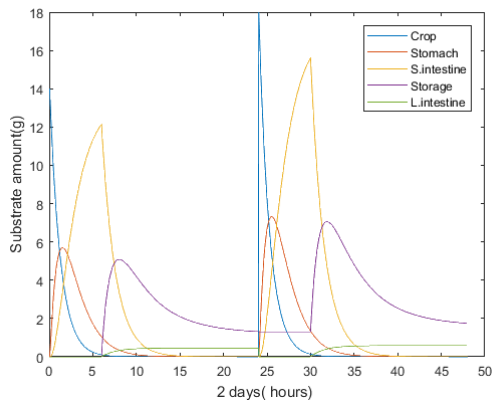


Figure: 2 days dynamic, $\beta_1 = 0.4$, $\gamma_{st} = 0.6$, $\gamma_{abs} = 0.6$, $\gamma_{si} = 0.023$, $\kappa_0 = 0.9$, $\mu_{max} = 0.01$, $\beta_2 = 0.01$, $V_{max} = 0.07$, $K = 26.8$

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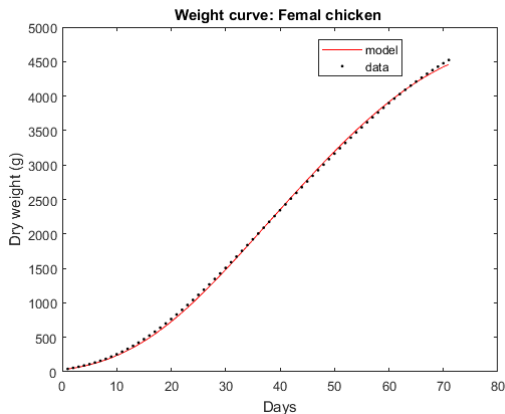


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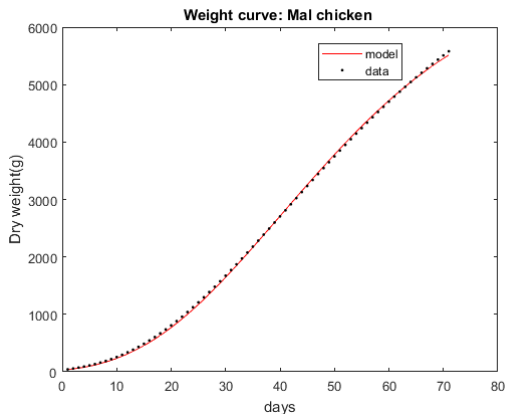
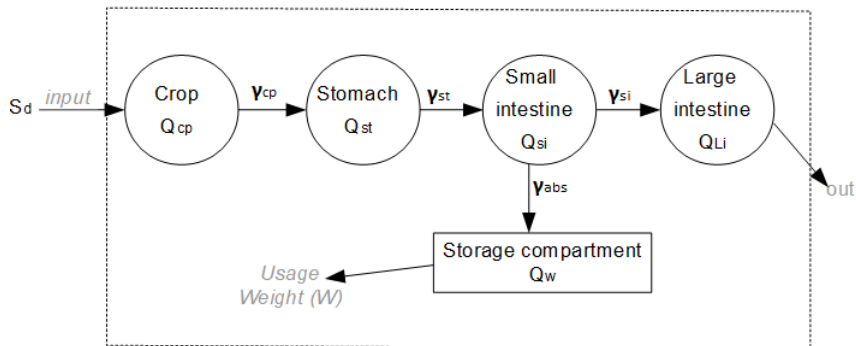
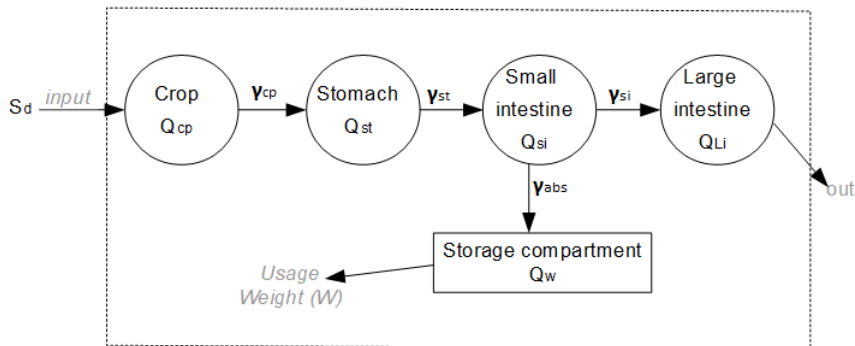
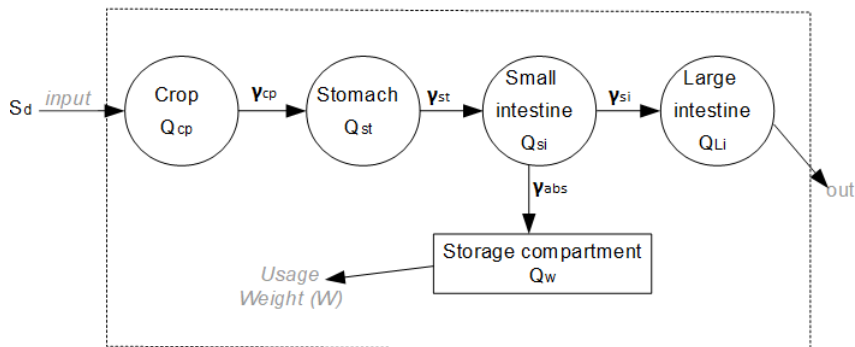


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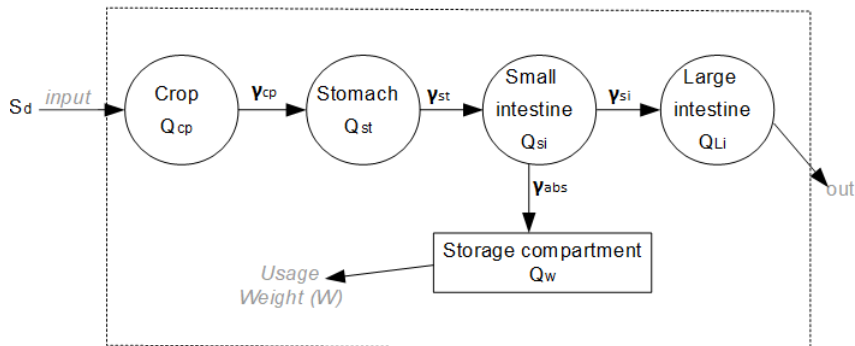


Python code: Virtual chicken.



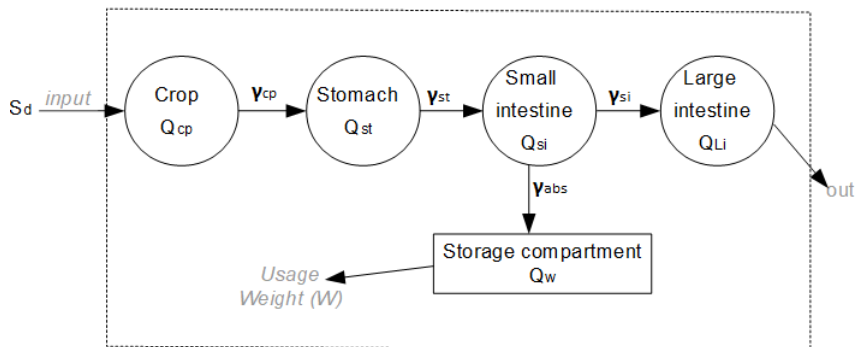
Python code: Virtual chicken.

- Daily food consumption and its composition.



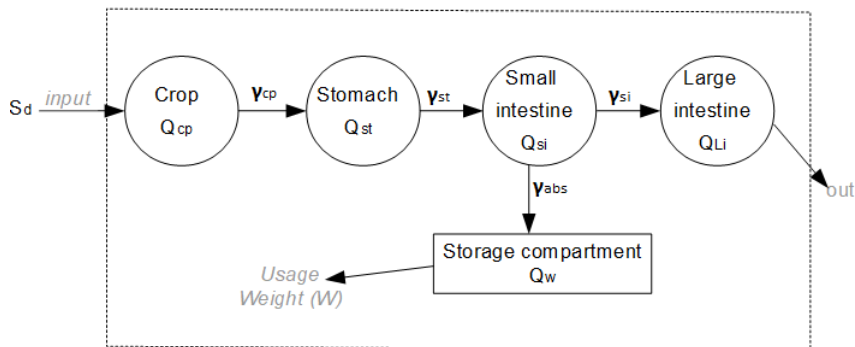
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Python code: Virtual chicken.

- Daily food consumption and its composition.
- Mathematical description of biological functions.
- Optimize different parameters.
- Predicts the daily growth.

Challenges and perspectives

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 - Estimation of different fluxes rates.
 - Important biological processes happening in different compartments.

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- Perspectives:

1. Find needed DATA in order to build a better descriptive model.
2. Build a virtual chicken that allows fitting parameters using Machine Learning approach.

Références

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THANK YOU