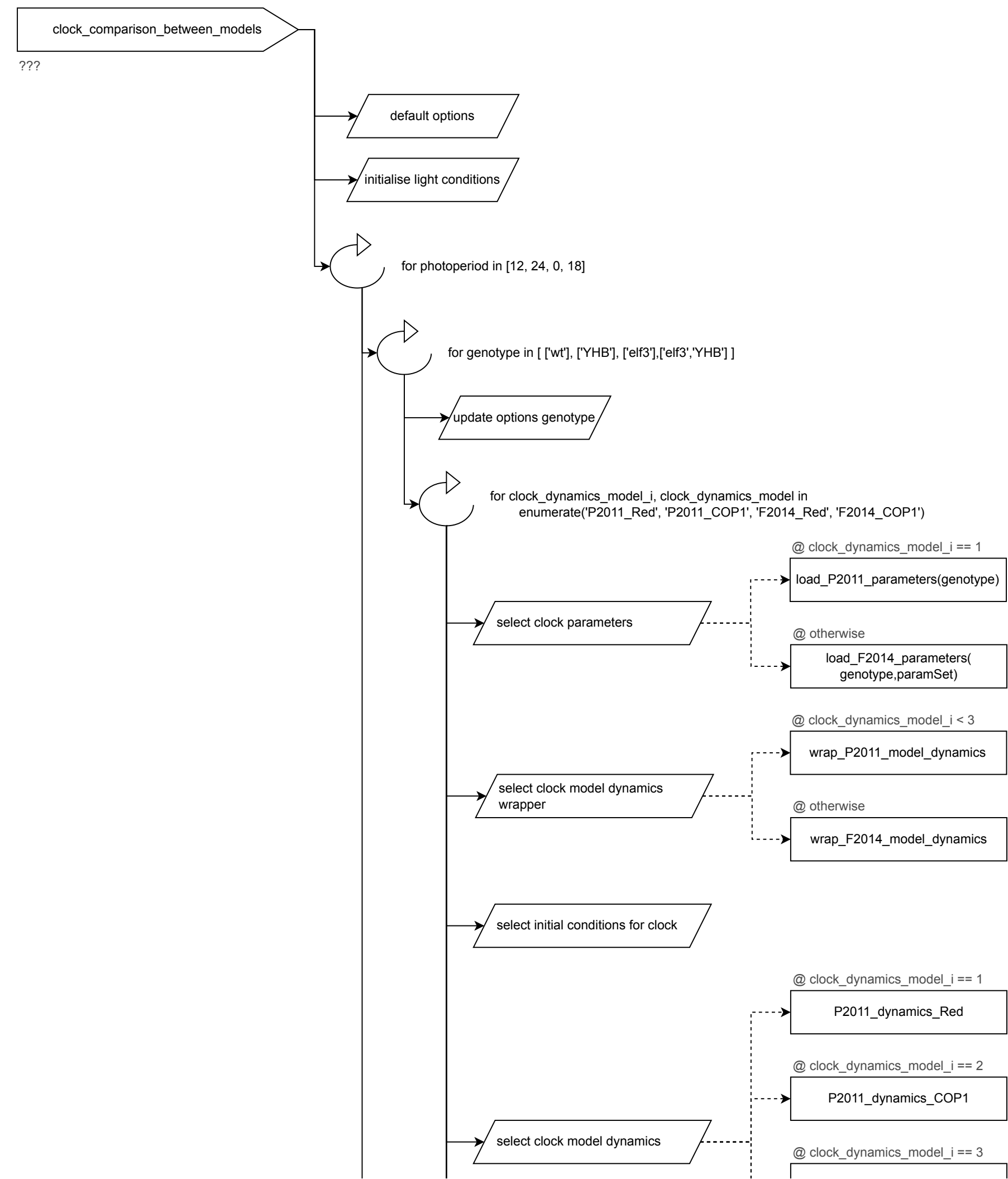
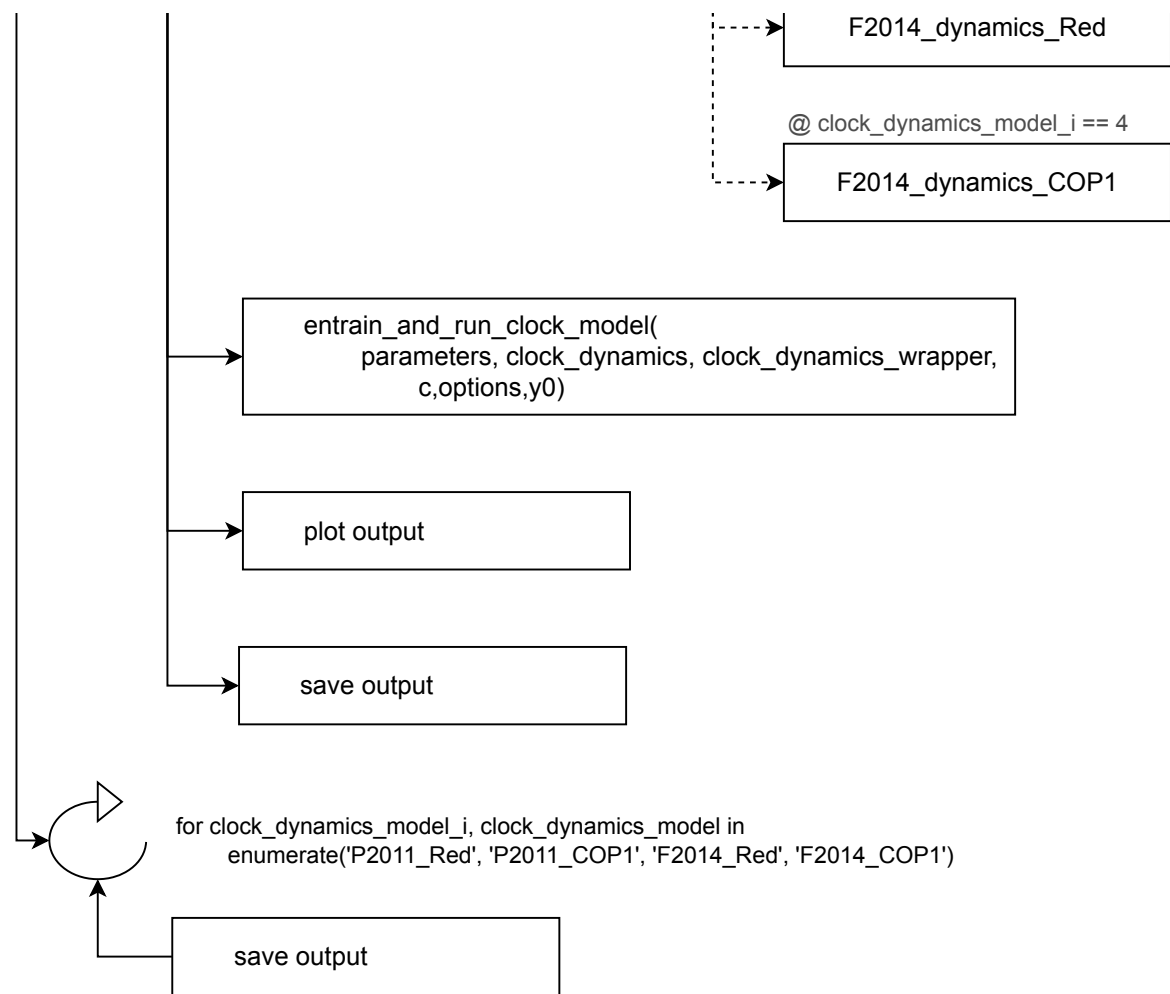


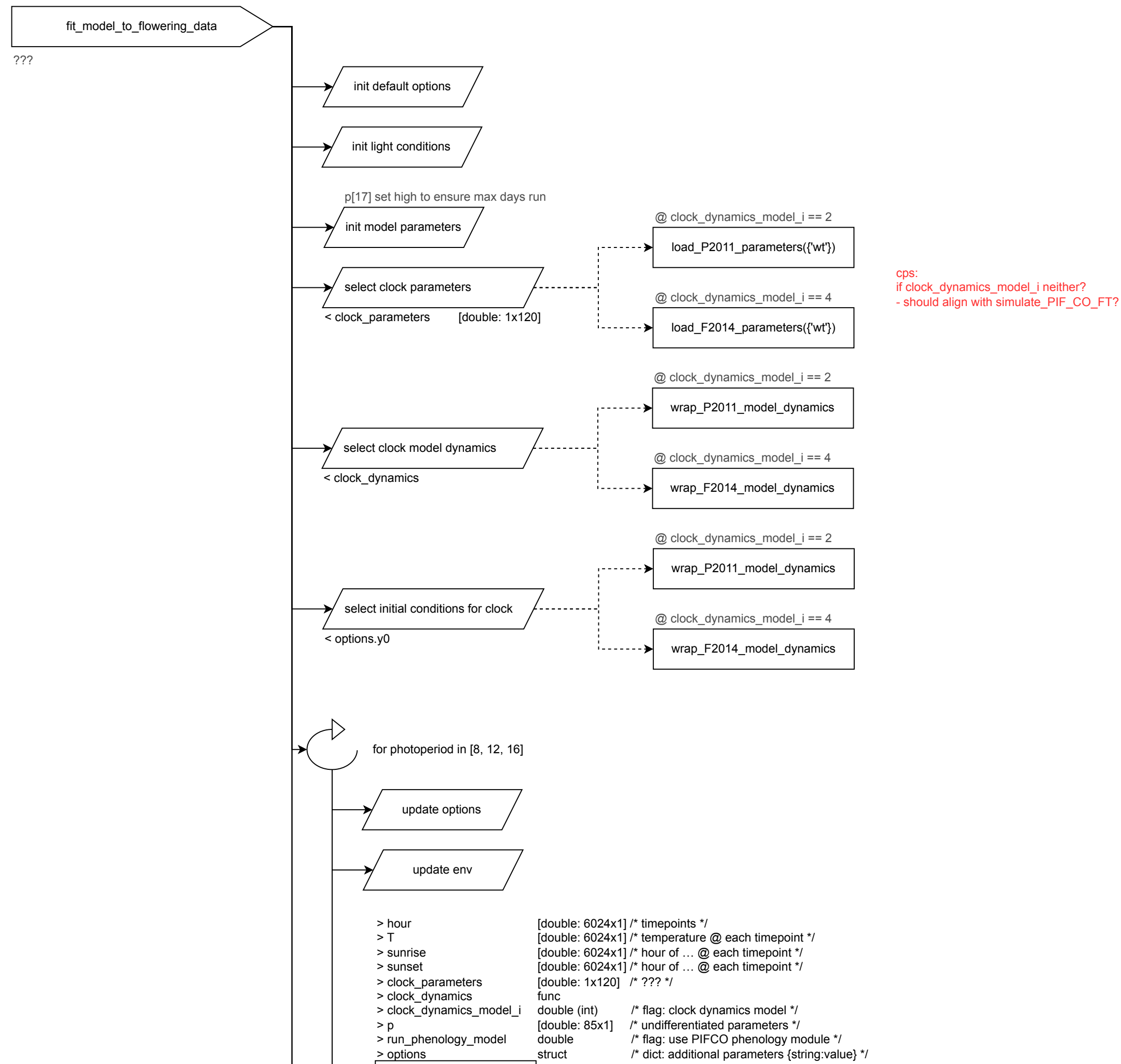
Functional Structure of Plant Model

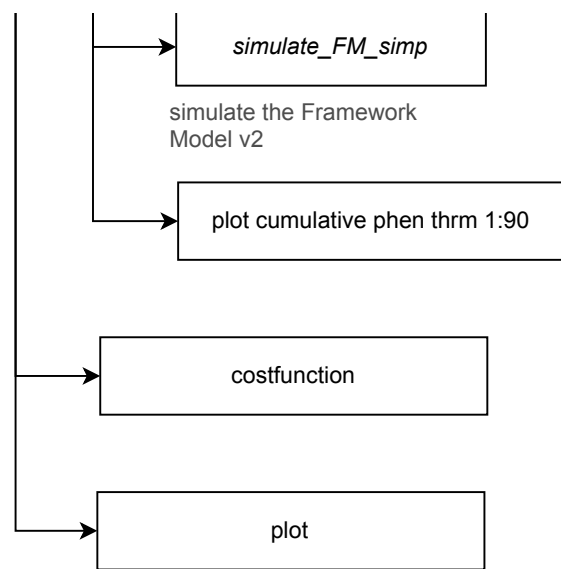
KEY

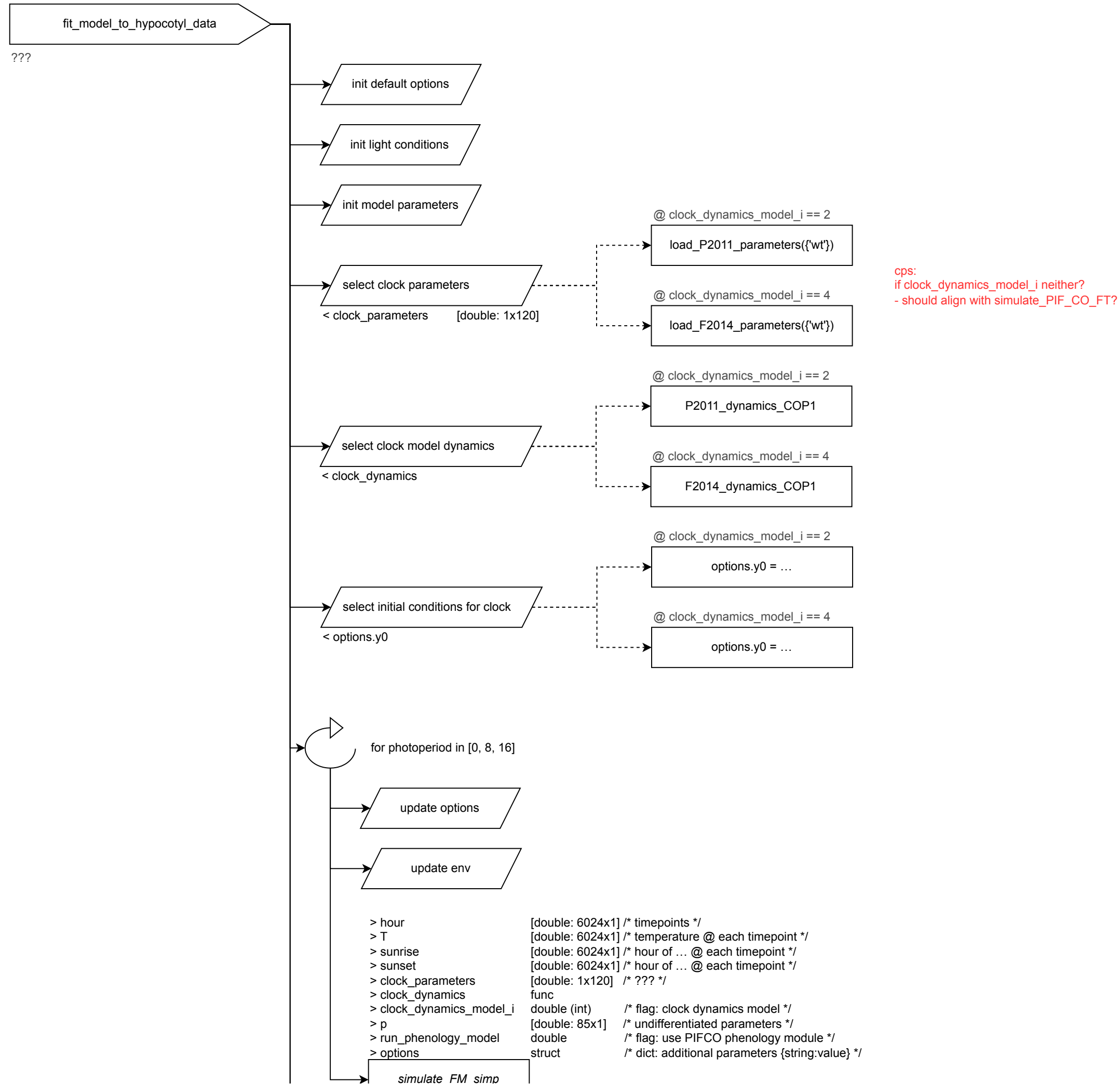
entry point	Entry point — module / function that calls rest of functions
module / function	module / function — functional component of model, may call other functions
<i>module / function reference</i>	reference — placeholder / reference for module / function unpacked elsewhere in document
code chunk	Chunk of code / logic — logical chunk of code









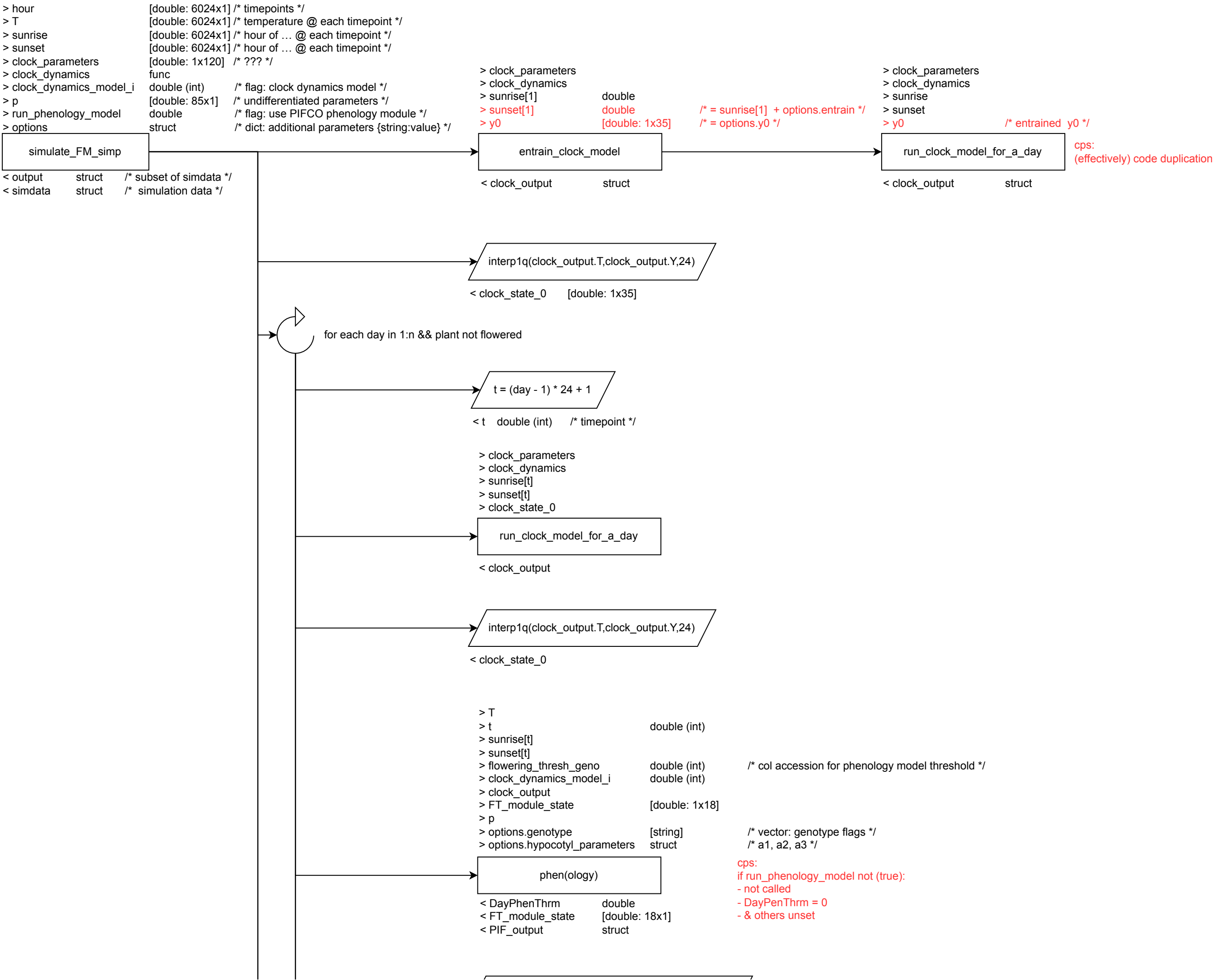


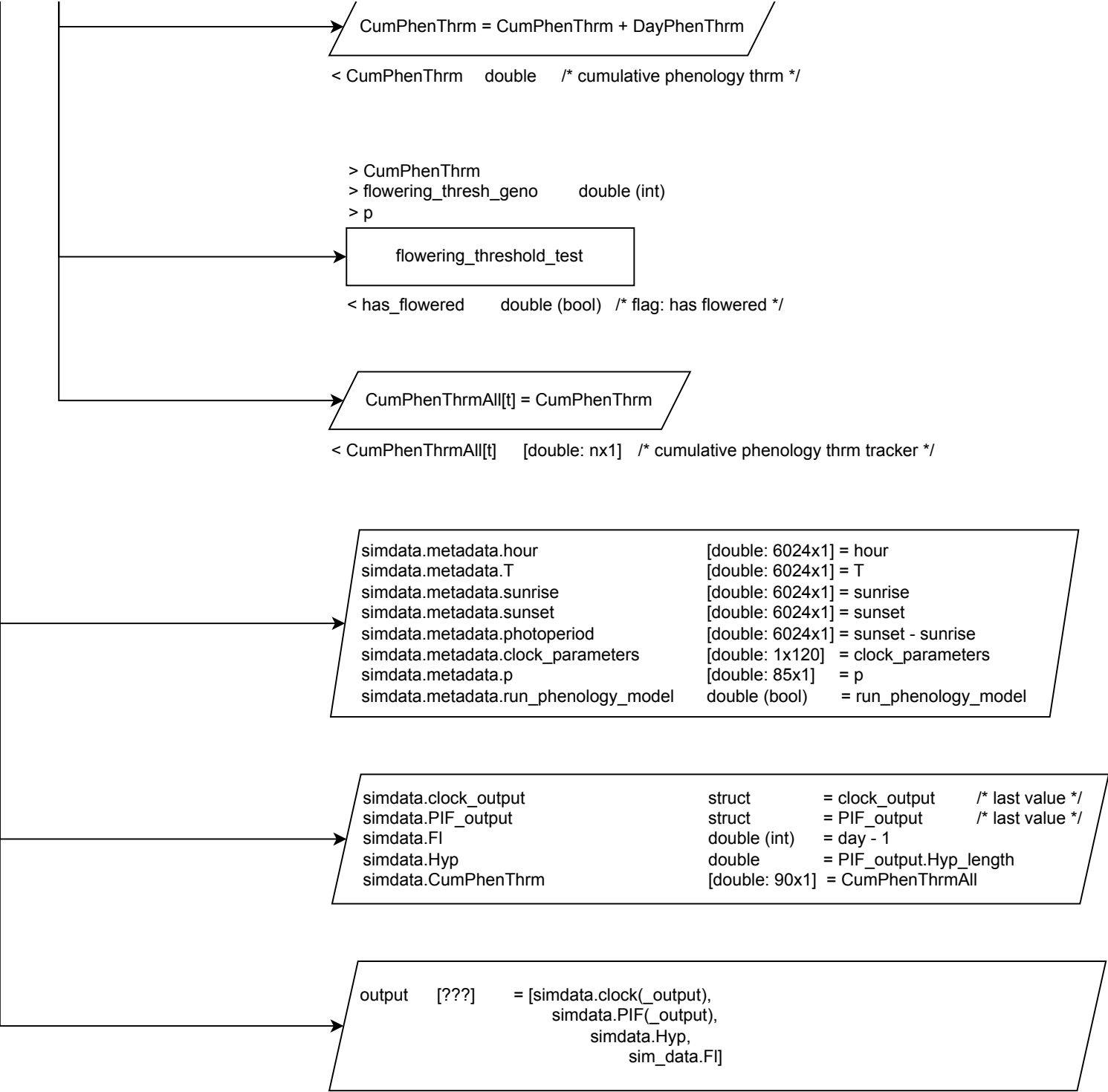
simulate the Framework

Model v2

costfunction

plot





> T

[double: 6024x1]

/* Temperature at time points */

> t

double (int)

/* timepoint */

> sunrise

double (int)

/* sunrise @ hours */

> sunset

double (int)

/* sunset @ hours */

> geno

double (int)

/* genotype index */

> clock_dynamics_model_i

double (int)

/* flag: clock dynamics model */

> clock_output

struct

/* clock model generated time series for parameters */

> FT_module_state

[double: 18x1]

/* state vector of FT model (conc. model components @ initial time point */

> p

[double: 85x1]

/* undifferentiated parameters */

> flowering_genotype

[string]

/* vector: genotype flags */

> hypocotyl_parameters

struct

/* a1, a2, a3 */

phen(ology)

< DayPhenThrm double

< FT_module_state [double: 18x1]

< PIF_output struct

set-up constants based on p

> sunrise double (int) /* sunrise @ hours */

> sunset double (int) /* sunset @ hours */

> clock_output struct /* clock model generated time series for parameters */

> clock_dynamics_model_i double (int) /* flag: clock dynamics model */

> FT_module_state [double: 1x18] /* state vector of FT model (conc. model components @ initial time point */

> temperatures [double: 6024x1] /* Temperature at time points */

> flowering_genotype [string] /* vector: genotype flags */

> hp struct /* = hypocotyl_parameters: a1, a2, a3 */

simulate_PIF_CO_FT_model

< dailyFTarea

< FT_module_state

< PIF_output

select clock parameters

< clock_parameters [double: 1x120]

@ clock_dynamics_model_i < 3

load_P2011_parameters

(flowering_genotype)

@ otherwise

load_F2014_parameters

(flowering_genotype)

@ clock_dynamics_model_i < 3

wrap_P2011_model_dynamics

@ otherwise

wrap_F2014_model_dynamics

select clock model dynamics

wrapper

< func_dynamics_wrapper

u = func_dynamics_wrapper(

clock_output.T,clock_output.Y,clock_parameters)

< u struct

> flowering_genotype [string] /* vector: genotype flags */

> temperature [double: 6024x1] /* Temperature at time points */

load_PIF_CO_FT_parameters.m

< parameters.PIF_CO_FT

cps:

tweaks parameters if temperature @ 27 °C

