# Model\_with\_VC&T\_correction

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## Description

-This file is about the building of the PESTIPOND model.

#### Set the file and call libraries

#### Read the data

```
t1 <- Sys.time()
data 1 <- read.csv2("data 1.csv", header = FALSE, skip=0, nrows = 101)
                                                                                     #read the CSV file of d
names(data_1)=data_1[1,]
                                                                                     #rename the columns of
data 1 \leftarrow data 1[-1,1:11]
                                                                                     #delete the first line
row.names(data_1)= data_1[,2]
                                                                                     #rename the lines of da
                  <- as.Date(data_1$Day , format = "%d/%m/%Y")
                                                                                     #convert day (d/m/y) to
data_1$Date
data 1
                  <- data_1[,-1]
                                                                                     #remove the old date co
                  <- as.data.frame(data 1)</pre>
                                                                                     #convert to a data fram
data 1
                  <- data_1[,c(11,1:10)]
data_1
                                                                                     #reorder columns
for (i in 2:11){
 data_1[,i] <- as.numeric(data_1[,i])</pre>
                                                                                     #convert to numeric
}
data_1 <- as.data.frame(data_1)</pre>
          <- 5269 #m2
Α
          <- 0.6*11e6/100 #g
Mveg
```

Build the input functions We don't need to calculate hw with a water balance, we can deduce it from hw = H - ds; with H=1m, and ds is deducted from Msed

```
Qin <- approxfun(data_1$`Step i`, data_1$`Qin (1/s)`, method = 'constant', rule = 2)

Qout <- approxfun(data_1$`Step i`, data_1$`Qout (1/s)`,method = 'constant', rule = 2)

TSS_in <- approxfun(data_1$`Step i`, data_1$`TSS_in (g/l)`,method = 'constant', rule = 2)

TSS_out <- approxfun(data_1$`Step i`, data_1$`TSS_out (g/l)`,method = 'constant', rule = 2)

Cin <- approxfun(data_1$`Step i`, data_1$`Cin (ug/l)`,method = 'constant', rule = 2)

Cp_in <- approxfun(data_1$`Step i`, data_1$`Cp_in(ug/g)`,method = 'constant', rule = 2)

TC <- approxfun(data_1$`Step i`, data_1$`T (°C)`,method = 'constant', rule = 2)

inputs <- NULL
```

## Write down the differential equations

- -The pesticides fate is described with a mass balance equation.
- -In this test case we will consider all of the compartments of the pond:
  - Dissolved pesticides
  - Particulate pesticides
  - Sediments
  - Vegetation

-We'll consider all the processes corrected according to vegetative cover (VC) and temperature (teta).

Now we will write down this model equation in R:

#### Parameters

Time We want the concentration C (ug/l & ug/g) to be calculated at every t = 0.01 day

```
times \leftarrow seq(1, 100, by = 0.01)
```

# Initial values

For this case we choose Isoda solver:

#### **ODE** solving

```
out <- as.data.frame(ode(yin, times, model, parms, method = 'lsoda', verbose = T, atol=1e-12, rtol=1e-1
#saveRDS(out, pasteO(getwd(), '/out.RDS'))
print(summary(out))
t2 <- Sys.time()
difftime(t2,t1)
#out <- as.data.frame(readRDS('C:/aya_local/Model/R_model_new/SA_OUTPUTS_15/16997.RDS')[[2]])</pre>
```

## Plot output

```
par(mfrow=c(2,2))
plot(times,out$Cdiss, col='blue',type='1',lwd=2, main='Water',ylab='ug/l', xlab = 'days')
plot(times,out$Cp, col='brown',type='1',lwd=2, main='Suspened solids',ylab='ug/g', xlab='days')
plot(times,out$Cs, col='red',type='1',lwd=2, main='Sediments',ylab='ug/g', xlab='days')
plot(times,out$Cv, col='green',type='1',lwd=2, main='Vegetation',ylab='ug/g', xlab='days')
```

#### Mass calculation

```
## Mass error ####
          <- parms[[1]]
VS
vr
          <- parms[[2]]
kbio_w <- parms[[3]]</pre>
kbio_TSS <- parms[[4]]</pre>
kbio_sed <- parms[[5]]</pre>
kbio_veg <- parms[[6]]</pre>
kp
         <- parms[[7]]
          <- parms[[8]]</pre>
kads_sed <- parms[[9]]</pre>
kads_TSS <- parms[[10]]</pre>
kads_veg <- parms[[11]]</pre>
        <- parms[[12]]
         <- parms[[13]]
teta
phi
         <- parms[[14]]
         <- parms[[15]]
rhob
                 <- NULL
Mass
Mass$'time'
                 <- times
Mass$'TC'
                 <- TC(times)
Mass
                 <- as.data.frame(Mass)</pre>
Mw
                 <- NULL
                 <- NULL
ďμ
Ms
                 <- NULL
Мτ
                 <- NUI.I.
```

```
MT
               <- NULL
Mdiss in
               <- NULL
Mdiss_out
               <- NULL
               <- NULL
Mp in
Mp_out
               <- NULL
MT_in
               <- NULL
               <- NULL
MT_out
for ( i in 1:length(times)) {
  Mw[i] <- out$Cdiss[i] * A * (1-out$Msed[i]/(rhob*A*10^6)) * phi * 1000 #ug #we calculate the net mas
 Mp[i] <- out$Cp[i] * out$MTSS[i] * phi #uq</pre>
 Ms[i] <- out$Cs[i] * out$Msed[i] * phi</pre>
                                            #uq
 Mv[i] <- out$Cv[i] * Mveg</pre>
 MT[i] \leftarrow Mw[i] + Mp[i] + Ms[i] + Mv[i]
}
Mass$'Mw' <- Mw
Mass$'Mp' <- Mp
Mass$'Ms' <- Ms
Mass$'Mv' <- Mv
Mass$'MT' <- MT
for (i in 1:length(times)){
  Mdiss_in[i] <- Cin(times[i]) * Qin(times[i]) * 24 * 3600 /99 #uq/0.01day
  #we set Mdiss_in(t=0)=0 to respect MB, that's why we divide the inlet mass of 10uq/L per 99 to obtain
  Mdiss_out[i] <- out$Cdiss[i] * Qout(times[i]) * 24 * 3600 * 0.01
  #we divide the equivalent mass of 150ug/g per 100 to obtain an instantaneous mass (per 0.01day)
               <- Cp_in(times[i]) * TSS_in(times[i]) * Qin(times[i]) * 24 * 3600/99 #ug</pre>
  Mp_in[i]
  #same goes for the particulate inlet/outlet mass
  Mp_out[i]
              <- out$Cp[i] * TSS_out(times[i]) * Qout(times[i]) * 24 *3600 * 0.01</pre>
  MT_in[i]
              <- Mdiss_in[i] + Mp_in[i] #ug
  MT_out[i]
               <- Mdiss_out[i] + Mp_out[i] #ug
}
#even the initial conditions should respect the MB (MO [in the pond]=M-1+MinO-MoutO-Mdeq0), if C(t=0)=0
Mdiss_in[1] \leftarrow 0
Mp_in[1]
          <- 0
```

```
MT_in[1] <- 0
Mass$'Min'
                <- Mdiss_in
Mass$'Mout'
                <- Mdiss_out
Mass$'Mp_in'
                <- Mp_in
Mass$'Mp_out'
                <- Mp_out
Mass$'MT_in' <- MT_in
Mass$'MT_out' <- MT_out</pre>
Mdeg
        <- NULL
Mdegw
        <- NULL
Mdegp
        <- NULL
        <- NULL
Mdegs
        <- NULL
Mdegv
Mads_sed <- NULL</pre>
Mads_TSS <- NULL</pre>
Mads_veg <- NULL</pre>
Mdes_sed <- NULL</pre>
Mdes_TSS <- NULL
Mdes_veg <- NULL</pre>
M_sett <- NULL
M_res <- NULL
TE <- Mass$'TC'
for (i in 1:length(times)){
  Mdegw[i]
             <- (kbio_w + kp + kv) * teta^(TE[i]-20) * Mw[i] * 0.01 #we correct degradation according</pre>
            <- kbio_TSS * teta^(TE[i]-20) * Mp[i] * 0.01</pre>
  Mdegp[i]
  Mdegs[i]
            <- kbio_sed * teta^(TE[i]-20) * Ms[i] * 0.01</pre>
  Mdegv[i]
           <- kbio_veg * teta^(TE[i]-20) * Mv[i] * 0.01</pre>
  Mdeg[i]
             <- Mdegw[i] + Mdegp[i] + Mdegs[i] + Mdegv[i] #ug #mdeg = dm/dt
  Mads_sed[i] <- kads_sed * Mw[i] * 0.01 #adsorbed during 0.01 day</pre>
  Mads_TSS[i] <- kads_TSS * Mw[i] * 0.01</pre>
  Mads_veg[i] <- kads_veg * Mw[i] * 0.01
  Mdes_TSS[i] <- kads_TSS * A * (1-out$Msed[i]/(rhob*A*10^6))/(out$MTSS[i]*Kd*1e-6) * Mp[i] * 0.01
  Mdes_veg[i] <- kads_veg * A * (1-out$Msed[i]/(rhob*A*10^6))/(Mveg*Kd*1e-6) * Mv[i] * 0.01
  M_{\text{sett}[i]} <- vs * 1/(1-out\$Msed[i]/(rhob*A*10^6)) * Mp[i] * 0.01
```

```
M_res[i] <- vr * 1/(1-out$Msed[i]/(rhob*A*10^6)) * Ms[i] * 0.01</pre>
}
Mass$'Mdeg' <- Mdeg
Mass$'Mdegw' <- Mdegw
Mass$'Mdegp' <- Mdegp
Mass$'Mdegs' <- Mdegs
Mass$'Mdegv' <- Mdegv
Mass$'Mads_sed' <- Mads_sed
Mass$'Mads_TSS' <- Mads_TSS</pre>
Mass$'Mads_veg' <- Mads_veg
Mass$'Mdes_sed' <- Mdes_sed
Mass$'Mdes_TSS' <- Mdes_TSS</pre>
Mass$'Mdes_veg' <- Mdes_veg
Mass$'M_sett' <- M_sett</pre>
Mass$'M_res' <- M_res
Mass <- round(Mass,4)</pre>
write_xlsx(Mass,"C:\\aya_local\\Model\\Resultats\\Mass_JDD.xlsx")
```

In this section we estimate the mass balance error err(%):

## Total mass balance error

```
e <- NULL
N <- length(Mass$time)
e[1] <- 0
#instantaneous MB error of the total mass of pesticides in the pond
for (i in 2:N){
    #no need to estimate the error at the initial step because we already know the outputs there (i.e., i
    e[i] <- 100*(Mass$MT[i]-Mass$MT[1]-sum(Mass$MT_in[2:i])+sum(Mass$MT_out[2:i])+sum(Mass$Mdeg[2:i]))/sum
}
Mass$'err' <- e
summary(abs(Mass$'err'))
#total MB error of the total mass of pesticides in the pond
MBE <- 100*(Mass$MT[N]-Mass$MT[1]-sum(Mass$MT_in[2:N])+sum(Mass$MT_out[2:N])+sum(Mass$Mdeg[2:N]))/sum(Mprint(paste('MBE (%) = ',MBE))</pre>
```

Solids mass balance error

```
MTSS_in
              <- NULL
              <- NULL
MTSS_out
              <- NULL
Msett
Mres
              <- NULL
NS <- length(times)
for (i in 1:NS){
 MTSS_in[i]
               <- TSS_in(times[i]) * Qin(times[i]) * 24 * 3600 * 0.01
                                                                                           #g
 MTSS_out[i] <- TSS_out(times[i]) * Qout(times[i]) * 24 * 3600 * 0.01
for (i in 1:NS){
  Msett[i]
               <- vs * 1/(1-out$Msed[i]/(rhob*A*10^6)) * out$MTSS[i] * 0.01</pre>
 Mres[i]
               <- vr * 1/(1-out$Msed[i]/(rhob*A*10^6)) * out$Msed[i] * 0.01</pre>
}
#the solving scheme is forward Mi+1 = Mi + Mini - Mouti
Msolids
           <- out$MTSS + out$Msed
MBE_solids <- 100*(Msolids[NS] - Msolids[1] - sum(MTSS_in[1:NS-1]) + sum(MTSS_out[1:NS-1]))/sum(MTSS_in[
print(paste('MBE_solids (%) = ', MBE_solids ))
```

## MBE for water compartment

```
 \begin{aligned} & sum(Mass\$Mout[2:N]) + sum(Mass\$Mdegw[2:N]) + sum(Mass\$Mads\_sed[2:N]) + sum(Mass\$Mads\_TSS[2:N]) + sum(Mass\$Min[2:N]) + sum(Mass\$Mdes\_sed[2:N]) + sum(Mass\$Mdes\_veg[2:N]) \end{aligned}
```

# MBE for TSS compartment

## MBE for sediments compartment

```
e_s <-NULL
e_s[i] <- 0
#instantaneous MB error of the mass of pesticides in sediments

for (i in 2:N){

    e_s[i] <- 100*(Mass$Ms[i]-Mass$Ms[1]-sum(Mass$Mads_sed[2:i])+sum(Mass$Mdegs[2:i])+sum(Mass$Mdes_sed[2:i])+sum(Mass$M_sett[2:i]))}

    sum(Mass$M_sett[2:i])+sum(Mass$M_res[2:i]))/(sum(Mass$Mads_sed[2:i])+sum(Mass$M_sett[2:i]))}

Mass$'err_s' <- e_s
summary(abs(Mass$'err_s'))

#total MB error of the mass of pesticides in sediments

MBE_s <- 100*(Mass$Ms[N]-Mass$Ms[1]-sum(Mass$Mads_sed[2:N])+sum(Mass$Mdegs[2:N])+sum(Mass$Mdes_sed[2:N])</pre>
```

# MBE for vegetation compartment

```
e_v <- NULL
e_v[1] <- 0

#instantaneous MB error of the mass of pesticides in vegetation

for (i in 2:N){

    e_v[i] <- 100*(Mass$Mv[i]-Mass$Mv[1]-sum(Mass$Mads_veg[2:i])+sum(Mass$Mdegv[2:i])+sum(Mass$Mdes_veg[2])

Mass$'err_v' <- e_v
summary(abs(Mass$'err_v'))

#total MB error of the mass of pesticides in vegetation

MBE_v <- 100*(Mass$Mv[N]-Mass$Mv[1]-sum(Mass$Mads_veg[2:N])+sum(Mass$Mdegv[2:N])+sum(Mass$Mdes_veg[2:N])</pre>
```

# Total MBE in each compartment

```
cat('MBE_solids (%) = ', MBE_solids, '\n',
    'MBE_w (%) = ', MBE_w, '\n',
    'MBE_p (%) = ', MBE_p, '\n',
    'MBE_s (%) = ', MBE_s, '\n',
    'MBE_v (%) = ', MBE_v, '\n',
    'MBE (%) = ', MBE, '\n')
```