## Oxygen mass transfer sensitivity in ABM

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```
library(ABM)
```

Default simulation

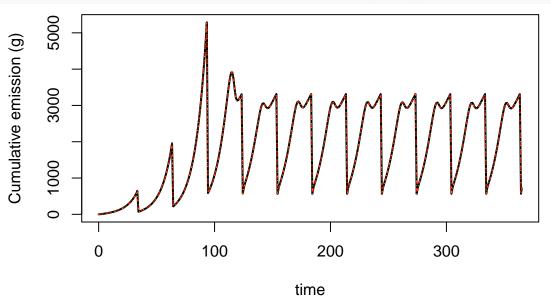
```
out1 <- abm()
```

Total cumulative emission (g) and emission rate (g/d) are plotted below.

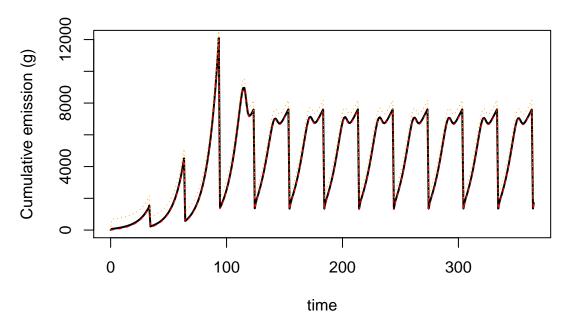
Completely shut off respiration, and also increase mass transfer coefficient by a factor of 10.

```
out2 <- abm(add_pars = list(kl.oxygen = 0))
out3 <- abm(add_pars = list(kl.oxygen = 5))

plot(CH4_emis_rate ~ time, data = out1, type = 'l', ylab = 'Cumulative emission (g)', lwd = 2)
lines(CH4_emis_rate ~ time, data = out2, col = 'red', lty = 2)
lines(CH4_emis_rate ~ time, data = out3, col = 'orange', lty = 3)</pre>
```

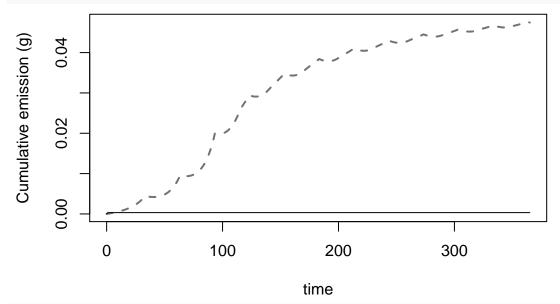


```
plot(CO2_emis_rate ~ time, data = out1, type = 'l', ylab = 'Cumulative emission (g)', lwd = 2)
lines(CO2_emis_rate ~ time, data = out2, col = 'red', lty = 2)
lines(CO2_emis_rate ~ time, data = out3, col = 'orange', lty = 3)
```



Conclude that respiration is minor sink for substrate, and uncertainty is not important. Compare CH4/respiration COD consumption:

```
plot(f_COD_CH4_cum ~ time, data = out3, type = 'l', col = 'gray45', lty = 2, ylab = 'Cumulative emission
lines(f_COD_respir_cum ~ time, data = out1, lwd = 1)
```



tail(out3[, c(1, which(grepl('^f\_COD', names(out3))))])

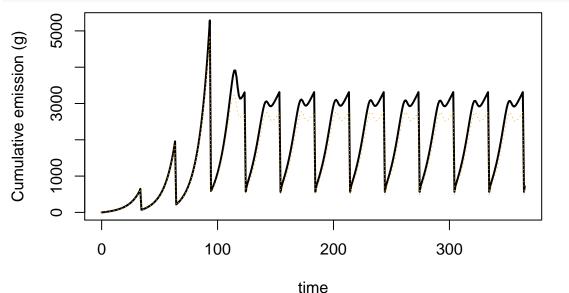
```
time f_COD_CH4_rate f_COD_CH4_cum f_COD_respir_cum f_COD_sr_cum
## 362 360.3300
                    0.07750056
                                   0.04732569
                                                    0.003268801
                    0.07905528
                                                    0.003268801
                                                                            0
   363 361.3300
                                   0.04741351
   364 362.3300
                    0.08062059
                                   0.04750516
                                                    0.003268801
                                                                            0
                                   0.04760058
                                                    0.003268801
                                                                            0
  365 363.3297
                    0.08218735
##
  366 364.3297
                    0.01407087
                                   0.04750855
                                                    0.003268801
                                                                            0
## 367 365.0000
                    0.01787083
                                   0.04745413
                                                    0.003268801
                                                                            0
```

Try increasing  $k_L 10x$  and decreasing the maximum depth. To do that we will increase surface area from 11 m2 to 333 m2, for a maximum depth of 1 cm.

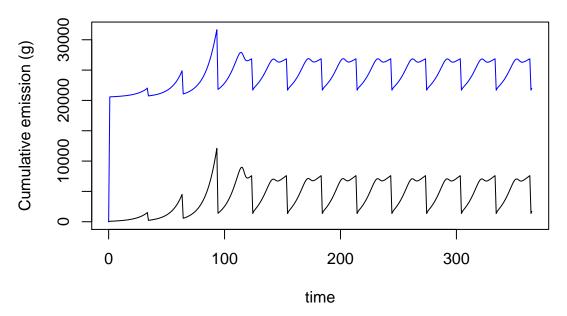
```
out4 <- abm(add_pars = list(kl.oxygen = 5, area = 333))</pre>
tail(out4[, c(1, which(grepl('^f_COD', names(out4))))])
##
           time f_COD_CH4_rate f_COD_CH4_cum f_COD_respir_cum f_COD_sr_cum
                    0.06536290
                                   0.04156057
                                                     0.09895553
## 362 360.3300
                                                                            0
## 363 361.3300
                    0.06652123
                                   0.04162965
                                                     0.09895553
                                                                            0
                    0.06770262
## 364 362.3300
                                   0.04170161
                                                     0.09895553
                                                                            0
## 365 363.3297
                    0.06889125
                                   0.04177642
                                                     0.09895553
                                                                            0
## 366 364.3297
                                                     0.09895553
                                                                            0
                    0.01216247
                                   0.04169514
## 367 365.0000
                    0.01553775
                                   0.04164710
                                                     0.09895553
                                                                            0
```

Now respiration becomes more important.

```
plot(CH4_emis_rate ~ time, data = out1, type = 'l', ylab = 'Cumulative emission (g)', lwd = 2)
lines(CH4_emis_rate ~ time, data = out4, col = 'orange', lty = 3)
```



```
plot(CO2_emis_rate ~ time, data = out4, type = 'l', ylab = 'Cumulative emission (g)', col = 'blue')
lines(CO2_emis_rate ~ time, data = out1)
```



But even so, for default conditions CH4 emission is not affected.