```
1 # install.packages("magrittr") # package installations are only needed the
  first time you use it
 2 # install.packages("dplyr") # alternative installation of the %>%
 3 library(magrittr) # needs to be run every time you start R and want to use
 4 library(dplyr) # alternatively, this also loads %>%
 5 # Part 6 - PPM scheme advection
 7 \```{r setup, include=FALSE}
 8 knitr::opts_chunk$set(echo = TRUE)
 9 library(ggplot2)
10 library(tidyverse)
11 \``
12
13 \```{r}
14 # Create the grid and initial conditions
                              # number of grid points in x-direction
15 \mid imax = 1000
16 | delx = 100.
                             # horizontal grid spacing (m)
17 \text{ delt} = 10.
                             # time increment (s)
                              # horizontal wind speed (m/s)
18 u = 5.
19|```
20
21 ```{r}
22 # Create initial concentration anomaly distribution in the x-direction
23 conc <- rep(0.0, imax) # initial concentration of background is zero
24 \text{ cmax} = 10.0
                                         # max initial concentration
25 \text{ conc}[100:150] \leftarrow \text{seq}(0., \text{cmax}, \text{len} = 51)
                                                          # insert left side of
   triangle
26 conc[150:200] <- seq(cmax, 0., len = 51)  # insert right side of
   triangle
27 conc[20:40] \leftarrow seq(0., -0.5*cmax, len = 21) # insert left side of triangle conc[40:60] \leftarrow seq(-0.5*cmax, 0., len = 21) # insert right side of
   triangle
29 conc_orig <- conc
30 ``
31
32
33 \```{r}
34 # create ideal solution
35 cideal <- rep(0.0, imax) # initial concentration of ideal background is
36 cideal[800:850] <- seq(0., cmax, len = 51) # insert left side of triangle
37 cideal[850:900] <- seq(cmax, 0., len = 51) # insert right side of triangle
38 cideal[720:740] <- seq(0., -0.5*cmax, len = 21) # insert left side of
39 cideal[740:760] <- seq(-0.5*cmax, 0., len = 21) # insert right side of
   triangle
40
41
42
43 ```{r}
44 | \text{nsteps} = (\text{imax} - 300) / (\text{u} * \text{delt} / \text{delx}) |
45 \times 1000 xvals = seg(1,1000)
46 \``
47
48
49
50 ## Plot 1
51 This has only the original concentration and the ideal solution
52
```

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```
53 \```{r}
54 plot(xvals, cideal, col = 'red', type = "l")
55 lines(xvals, conc, col = 'blue')
56 legend(400, 10, legend=c("cideal", "conc orig"),
          col=c("red", "blue"), lty=1, cex=0.8)
57
58 # plot <- ggplot() +
       geom\_line(aes(x = xvals, y = conc), color = "blue") +
59 #
       geom line(aes(x = xvals, y = cideal), color ="red") +
60 #
61 #
       theme bw() +
62 #
       xlab("grid index (i)") +
63 #
       ylab("quantity")+
       ggtitle("PPM plot")
64 #
65 ```
66 ## PPM scheme code
68 Here I use the code that was provided for the homework.
70 \```{r}
71 #### USING THE CODE PROVIDED
73 # 6) Use the HPPM method from CMAQ
74 # CW refers to the paper by Colella and Woodward.
75 # 1-D domain covers grid points i = 1 to imax. But 1 and imax are boundary-
76 # condition cells. The main interior computation is for i = 2 to (imax-1).
77 # Pre-calculate some constants
78 \text{ sixth} = 1.0/6.0
79 two3rds = 2.0/3.0
80 oneoverdelx = 1.0 / delx
81 # Allocate the vectors
82 dc = numeric(imax)
                            # nominal difference in concentration across a cell
83 clfirst = numeric(imax) # first guess of conc at left edge of cell i
                            # conc at right edge of cell i
84 cr = numeric(imax)
                            # conc at left edge of cell i
85 cl = numeric(imax)
86 | c6 = numeric(imax)
                            # this corresponds to parabola parameter a6 of CW
   eq.(1.4)
87 | FL = numeric(imax)
                            # pollutant flux into the left side of a grid cell
88 FR = numeric(imax)
                            # pollutant flux into the right side of a grid cell
89 # Iterate forward in time
90 for (n in 1:nsteps) {
                                           # for each time step n
91
92
       # To quarantee that solution is monotonic, check that the left edge of
93
   cell i
94
       #
            (which is between cells i and i-1) should not have a concentration
    lower
95
            or higher than the concentrations in those two neighboring cells
            Namely, is clfirst between c[i] and c[i-1]. If not, then fix.
96
97
       for (i in 2:(imax - 1)) {
                                         # for each interior grid point i
           del_cl = conc[i] - conc[i-1] # concentration difference with cell
98
   at left
           del cr = conc[i+1] - conc[i] # concentration difference with cell
99
   at right
           dc[i] = 0.5*(del_cl + del_cr) # 1st guess of avg conc difference
100
   across cell i
           if ((del cl*del cr)>0.0) { # then revise average difference
101
   across cell i
               dc[i] = sign(dc[i]) * min(abs(dc[i]), 2*abs(del_cl),
102
   2*abs(del cr) )
103
           } else {dc[i]=0.0}
                                           # for the special case of constant
   conc across cell
```

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```
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                                                hw7_ppm.R
            }
                                                 # end of grid-point (i) loop
    104
    105
    106
            # First guess for concentration at left edge of each cell, using revised
        dc value
    107
            for (i in 2:(imax - 1)) {
                                                 # for each interior grid point i
                 clfirst[i] = 0.5*(conc[i]+conc[i-1]) - sixth*(dc[i]-dc[i-1])
    108
    109
                                                 # end of grid-point (i) loop
    110
            # find parameters for the piecewise-continuous parabola in cell i
    111
            for (i in 2:(imax - 1)) {
    112
                                                 # for each interior grid point i
    113
                 # conc at the right edge (cr) of cell i equals concen at left edge of
    114
        cell i+1
                 cr[i] = clfirst[i+1]
                                                 # concentration at right edge of cell
    115
                 cl[i] = clfirst[i]
                                                 # concentration at left edge of cell
    116
    117
    118
                 # Check whether cell i is an extremum (is a peak or valley in the
        conc plot)
    119
                 if (((cr[i]-conc[i]) * (conc[i] - cl[i])) > 0.0) { # then not
        extremum
    120
                     # Find the two coefficients of the parabola: dc and c6:
    121
                     dc[i] = cr[i] - cl[i]
                                                 # updated concen diff. between right
    122
        and left edges
                     c6[i] = 6*(conc[i] - 0.5*(cl[i]+cr[i]))
    123
    124
                     if ((dc[i]*c6[i]) > (dc[i]*dc[i])) { # then adjust for
    125
        overshoot at left edge
                         cl[i] = 3.0*conc[i] - 2.0*cr[i]
    126
                     } else if ((-dc[i]*dc[i]) > (dc[i]*c6[i])) { # then adjust for
    127
        overshoot at right
    128
                         cr[i] = 3.0*conc[i] - 2.0*cl[i]
                     }
                                                 # end of block of "not extremum"
    129
        calculations
    130
                 } else {
                                                 # For an extremum, don't use a
    131
        parabola.
    132
                     cl[i] = conc[i]
                                                 # Instead, assume concen is constant
        across the cell,
                                                 # Thus, left and right concentrations
    133
                     cr[i] = cl[i]
        equal average conc.
    134
                 }
                                                 # end of grid-point (i) loop
    135
                 # second guess of coefficients for the parabola, from CW eq. (1.5)
    136
                dc[i] = cr[i] - cl[i]
    137
    138
                 c6[i] = 6.0*(conc[i] - 0.5*(cl[i] + cr[i]))
    139
            }
                                                 # end of grid-point (i) loop
    140
    141
    142
            # Initialize to 0 the fluxes into the left and right sides of cell i
    143
            FL \leftarrow rep(0.0, imax)
    144
    145
             FR \leftarrow rep(0.0, imax)
    146
    147
    148
             # Next, use parabolic fits within each cell to calculate the fluxes
        betweeen cells
     149
```

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