

!
 ! Magnolia CSL model file created on 2019-04-18T10:21:02.815
 ! PBPK Gas Uptake Model for 2,2-DCPe converted from original acslx model by EMK
 ! Units of this model are liters (L), hours (h), milligrams (mg), BW in kilograms (kg)

model DCP22

initial ! The INITIAL section contains statements evaluated 1x @ begin simulation

!*****! Physiological Parameters*****}

CONSTANT QPC = 13.2 ! Alveolar Ventilation Rate(l/hr-kg), Clewell et al., 2014, Table 6.1, p. 264
 CONSTANT QCC = 13.2 ! Cardiac Output(l/hr-kg), Clewell et al., 2014, Table 6.1, p. 264
 CONSTANT QLC = 0.242 ! Fract. Blood Flow to Liver, Table 25, p. 439, Brown et al., 1997
 CONSTANT QFC = 0.082 ! Fract. Blood Flow to Fat, Table 25, p. 439, Brown et al., 1997
 CONSTANT QSC = 0.257 ! Fract. Blood Flow to Poorly Perfused T., , Table 25, p. 439, Brown et al., 1997 (muscle + skin)
 QRC = 1.0 - QLC - QFC - QSC ! Fract. Blood Flow to Richly Perfused T

CONSTANT BW = 0.222 ! Body Weight(kg), experiment-specific, mean & SD all rats 0.222+/-0.0184
 CONSTANT VLC = 0.036 ! Volume Fraction Liver, Brown et al., 1997, Table 8, p. 418
 CONSTANT VFC = 0.076 ! Volume Fraction Fat, Brown et al., 1997, equation, p. 422 using avg BW
 CONSTANT VSC = 0.7 ! Volume Fraction Poorly Perfused T., muscle + skin, Table 5, p. 416, Brown et al., 1997
 VRC = 1.0 - VLC - VFC - VSC ! Volume Fraction Richly Perfused T.

!*****! Chemical Specific Parameters for DCPs*****}

! Partition Coeffs Estimated Based on method of Beliveau et al., (2003)}

CONSTANT PL = 1.81 ! Liver/Blood
 CONSTANT PF = 22.03 ! Fat/Blood
 CONSTANT PS = 0.84 ! Poorly/Blood (muscle is representative tissue)
 CONSTANT PR = 1.81 ! Richly/Blood (liver is representative tissue)
 CONSTANT PB = 3.75 ! Blood/Air
 CONSTANT MW = 112.99 ! Molecular Wt (g/mol)

!VmaxC & Km starting values - final values are estimated from set of vapor uptake curves

!CONSTANT VmaxC = 4.46 ! mg/hr-kg

!CONSTANT Km = 3.77 ! mg/L

CONSTANT KFC = 1.18 ! L/hr-kg, use if assuming first order metabolism

! Chamber/Rat/Exposure Parameters}

CONSTANT CONC = 500. ! Init. Inhaled Conc (ppm), experiment-specific

CONSTANT RATS = 1 ! Number of rats in chamber, always 1

CONSTANT VCHC = 3.8 ! Closed chamber vol (l)

CONSTANT KLOS = 0.04 ! Closed chamber loss, hr-1, experiment-specific

NETVOL = VCHC - (RATS * BW) ! Net Chamber Vol (l), i.e. volume of chamber not taken up by rats

AIO = (CONC * NETVOL * MW)/24450 ! Initial Conc of chemical in Chamber (mg/l)

CONSTANT TSTOP = 6.00 ! Length of experiment (hrs)

CONSTANT points = 100. ! No. of points in plot

!*****Scaled Parameters*****}

QC = QCC * (BW**0.75) ! Cardiac Output (l/hr)

QP = QPC * (BW**0.75) ! Alveolar Ventilation Rate (l/hr)

QL = QLC * QC ! Flow Liver Compartment (l/hr)

QF = QFC * QC ! Flow Fat Compartment (l/hr)

QS = QSC * QC ! Flow Slowly Perf. Tis. Cmpt. (l/hr)

QR = QRC * QC ! Flow Richly Perf. Tis. Cmpt. (l/hr)

VL = VLC * BW ! Volume Liver Compartment, Total

VF = VFC * BW ! Volume Fat Compartment

VS = VSC * BW ! Volume Slowly Perfused Tis. Cmpt.

VR = VRC * BW ! Volume Richly Perfused Tis. Cmpt.

!Comment in/out eq below if running w/ michaelis-menten vs. first order}

!VMAX = VMAXC * BW**0.75 ! VMAX scaled

KF = KFC * BW**0.75 !

end ! initial

dynamic ! The DYNAMIC section statements evaluated at each output time point

derivative

$$\begin{aligned} \text{RAI} &= (\text{RATS} * \text{QP} * (\text{CX} - \text{CI})) - (\text{KLOS} * \text{AI}) && ! \text{ mg/hr} \\ \text{AI} &= \text{INTEG}(\text{RAI}, \text{AIO}) && ! \text{ mg} \\ \text{CI} &= \text{AI} / \text{NETVOL} && ! \text{ mg/l} \\ \text{CP} &= (\text{CI} * 24450 / \text{MW}) && ! \text{ chamber conc, ppm, data is in ppm} \end{aligned}$$

! chamber loss

$$\begin{aligned} \text{RLOSS} &= (\text{KLOS} * \text{AI}) && ! \text{ mg/hr} \\ \text{LOSS} &= \text{INTEG}(\text{RLOSS}, 0.0) \end{aligned}$$

$$\text{CA} = ((\text{QC} * \text{CV}) + (\text{QP} * \text{CI})) / (\text{QC} + (\text{QP} / \text{PB})) \quad ! \text{ arterial (mg/L)}$$

!exhaled breath

$$\begin{aligned} \text{RAX} &= \text{QP} * \text{CX} && ! \text{ mg/hr} \\ \text{AX} &= \text{INTEG}(\text{RAX}, 0.0) && ! \text{ mg} \\ \text{CX} &= \text{CA} / \text{PB} && ! \text{ Conc. DCPe in exhaled air (mg/l)} \\ \text{CXPPM} &= ((0.7 * \text{CX}) + (0.3 * \text{CI})) * (24.45 / \text{MW}) && ! \text{ Conc. DCPE in ppm exhaled breath} \\ \text{AXKG} &= \text{AX} / \text{BW} && ! \text{ mg exhaled/kg body weight} \end{aligned}$$

!slowly perfused tissue group

$$\begin{aligned} \text{RAS} &= \text{QS} * (\text{CA} - \text{CVS}) && ! \text{ mg/hr} \\ \text{AS} &= \text{INTEG}(\text{RAS}, 0.0) && ! \text{ mg} \\ \text{CVS} &= \text{AS} / (\text{VS} * \text{PS}) && ! \text{ Conc partition to slow per. tis. (mg/l)} \\ \text{CS} &= \text{AS} / \text{VS} && ! \text{ Conc in volume slow per. tis. (mg/l)} \end{aligned}$$

!rapidly perfused tissue group

$$\begin{aligned} \text{RAR} &= \text{QR} * (\text{CA} - \text{CVR}) && ! \text{ mg/hr} \\ \text{AR} &= \text{INTEG}(\text{RAR}, 0.0) && ! \text{ mg} \\ \text{CVR} &= \text{AR} / (\text{VR} * \text{PR}) && ! \text{ Conc partition to rap per. tis. (mg/l)} \\ \text{CR} &= \text{AR} / \text{VR} && ! \text{ Conc in volume rap per. tis. (mg/l)} \end{aligned}$$

!fat

```

RAF = QF * (CA - CVF)      ! mg/hr
AF = INTEG(RAF, 0.0)      ! mg
CVF = AF / (VF * PF)      ! Conc partition to fat(mg/l)
CF = AF / VF              ! Conc in fat volume(mg/l)

```

!liver

```

RAL = QL * (CA - CVL) - RAM    ! mg/hr
AL = INTEG(RAL, 0.0)         ! mg
CVL = AL / (VL * PL)         ! Conc partition to liver(mg/l)
CL = AL / VL                 ! Conc in liver volume(mg/l)

```

!Note: RAM rate of metabolism can be switched between being expressed as
!Michaelis-Menten terms (mg/hr) and Clearance terms (L/hr) by commenting
!in or out between the two equations below. Note that parameters need to
!be changed within the inital section also (2 places) VMAX vs. KF & VMAXC vs KFC}

```

! RAM = (VMAX * CVL) / (KM + CVL)
RAM = KF * CVL
AM = INTEG(RAM, 0.0)

```

```

CV = (QF*CVF + QL*CVL + QS*CVS + QR*CVR) / QC    ! venous blood conc (mg/l)

```

```

DOSE = AIO - AI                                ! amt dcp in rat (mg)

```

```

DMASS = (AF + AL + AS + AR + AM)*RATS

```

!Criterion - mass balance (MB) should be near zero, i.e. <= ~e10-8.}

```

MB = DOSE - DMASS - LOSS    ! mass balance (mg)

```

```

TERMT(T >= TSTOP)        ! Condition for terminating simulation

```

end ! derivative

end ! dynamic

end ! program

! Parameter estimation script for 2,2-DCP PBPK model
 ! Closed-chambered rat data, 2,2-dichloropropane

load 'DCP22.csl'

! Any or all but one individual line can be commented out to simulate a single expt.
 ! Each line below calls data file and species parameters that change with each experiment
 data @file='dcp2_60a.csv' ds1 t='t' cp='cp' tstop=4.17 bw=0.217 klos=0.0305 conc=49
 data @file='dcp2_60b.csv' ds2 t='t' cp='cp' tstop=4.17 bw=0.222 klos=0.0332 conc=50
 data @file='dcp2_60c.csv' ds3 t='t' cp='cp' tstop=5.00 bw=0.214 klos=0.0208 conc=57
 data @file='dcp2_60d.csv' ds4 t='t' cp='cp' tstop=5.00 bw=0.209 klos=0.0312 conc=60
 data @file='dcp2_200a.csv' ds5 t='t' cp='cp' tstop=4.33 bw=0.213 klos=0.045 conc=190
 data @file='dcp2_200b.csv' ds6 t='t' cp='cp' tstop=4.33 bw=0.226 klos=0.025 conc=183
 data @file='dcp2_230a.csv' ds7 t='t' cp='cp' tstop=4.17 bw=0.211 klos=0.043 conc=221
 data @file='dcp2_500a.csv' ds8 t='t' cp='cp' tstop=6.0 bw=0.231 klos=0.032 conc=454
 data @file='dcp2_500b.csv' ds9 t='t' cp='cp' tstop=6.1 bw=0.267 klos=0.036 conc=496
 data @file='dcp2_600a.csv' ds10 t='t' cp='cp' tstop=5.67 bw=0.219 klos=0.059 conc=586
 data @file='dcp2_600b.csv' ds11 t='t' cp='cp' tstop=6.17 bw=0.210 klos=0.065 conc=617
 data @file='dcp2_1150a.csv' ds12 t='t' cp='cp' tstop=5.92 bw=0.189 klos=0.0468 conc=1105
 data @file='dcp2_1150b.csv' ds13 t='t' cp='cp' tstop=6.1 bw=0.266 klos=0.063 conc=1141
 data @file='dcp2_1150c.csv' ds14 t='t' cp='cp' tstop=6.33 bw=0.248 klos=0.0035 conc=1242
 data @file='dcp2_1300a.csv' ds15 t='t' cp='cp' tstop=5.92 bw=0.197 klos=0.0518 conc=1288

! A procedure to plot all data sets
 ! Used to plot the fits before and after parameter estimation
 procedure plotall

```
set tstop=4.17 bw=0.217 klos=0.0305 conc=49
start
plot cp 'ds1:cp' @title='2,2-DCP 60a'
```

```
set tstop=4.17 bw=0.222 klos=0.0332 conc=50
start
plot cp 'ds2:cp' @title='2,2-DCP 60b'
```

```
set tstop=5.00 bw=0.214 klos=0.0208 conc=57  
start  
plot cp 'ds3:cp' @title='2,2-DCP 60c'
```

```
set tstop=5.00 bw=0.209 klos=0.0312 conc=60  
start  
plot cp 'ds4:cp' @title='2,2-DCP 60d'
```

```
set tstop=4.33 bw=0.213 klos=0.045 conc=190  
start  
plot cp 'ds5:cp' @title='2,2-DCP 200a'
```

```
set tstop=4.33 bw=0.226 klos=0.025 conc=183  
start  
plot cp 'ds6:cp' @title='2,2-DCP 200b'
```

```
set tstop=4.17 bw=0.211 klos=0.043 conc=221  
start  
plot cp 'ds7:cp' @title='2,2-DCP 230a'
```

```
set tstop=6.0 bw=0.231 klos=0.032 conc=454  
start  
plot cp 'ds8:cp' @title='2,2-DCP 500a'
```

```
set tstop=6.1 bw=0.267 klos=0.036 conc=496  
start  
plot cp 'ds9:cp' @title='2,2-DCP 500b'
```

```
set tstop=5.67 bw=0.219 klos=0.059 conc=586  
start  
plot cp 'ds10:cp' @title='2,2-DCP 600a'
```

```
set tstop=6.17 bw=0.210 klos=0.065 conc=617
```

```
start
plot cp 'ds11:cp' @title='2,2-DCP 600b'

set tstop=5.92 bw=0.189 klos=0.0468 conc=1105
start
plot cp 'ds12:cp' @title='2,2-DCP 1150a'

set tstop=6.1 bw=0.266 klos=0.063 conc=1141
start
plot cp 'ds13:cp' @title='2,2-DCP 1150b'

set tstop=6.33 bw=0.248 klos=0.0035 conc=1242
start
plot cp 'ds14:cp' @title='2,2-DCP 1150c'

set tstop=5.92 bw=0.197 klos=0.0518 conc=1288
start
plot cp 'ds15:cp' @title='2,2-DCP 1300a'

end

prepare @clear T CP

set kfc = 1.08 @min=0.1 @max=2.16

plotall

fit @method=ml @errormodel=mixed kfc

plotall
```


!

! Magnolia CMD script created on 2020-02-24T16:12:57.165

! Sensitivity Analysis 22-DCP Global Morris Method

load 'DCP22.csl'

set QPC = 13.2 @min=10 @max=16 ! Alveolar Ventilation Rate(l/hr-kg), Clewell et al., 2014, Table 6.1, p. 264

set QCC = 13.2 @min=10 @max=16 ! Cardiac Output(l/hr-kg), Clewell et al., 2014, Table 6.1, p. 264

set QLC = 0.242 @min=0.2 @max=0.28 !Fract. Blood Flow to Liver, Table 25, p. 439, Brown et al., 1997

set QFC = 0.082 @min=0.05 @max=0.11 ! Fract. Blood Flow to Fat, Table 25, p. 439, Brown et al., 1997

set QSC = 0.257 @min=0.23 @max=0.28 ! Fract. Blood Flow to Poorly Perfused T., , Table 25, p. 439, Brown et al., 1997 (muscle + skin)

set BW = 0.222 @min=0.19 @max=0.25 ! Body Weight(kg), experiment-specific, mean & SD all rats 0.222+/-0.0184

set VLC = 0.036 @min=0.028 @max=0.044 ! Volume Fraction Liver, Brown et al., 1997, Table 8, p. 418

set VFC = 0.076 @min=0.05 @max=0.09 ! Volume Fraction Fat, Brown et al., 1997, equation, p. 422 using avg BW

set VSC = 0.7 @min=0.65 @max=0.75 ! Volume Fraction Poorly Perfused T., muscle + skin, Table 5, p. 416, Brown et al., 1997

!*****! Chemical Specific Parameters for DCPs*****}

! Partition Coeffs Estimated Based on method of Beliveau et al., (2003)}

set PL = 1.81 @min=1.0 @max=2.6 ! Liver/Blood

set PF = 22.03 @min=18 @max=26 ! Fat/Blood

set PS = 0.84 @min=0.6 @max=1.2 ! Poorly/Blood (muscle is representative tissue)

set PR = 1.81 @min=1.0 @max=2.6 ! Richly/Blood (liver is representative tissue)

set PB = 3.75 @min=2.8 @max=4.7 ! Blood/Air

set MW = 112.99 ! Molecular Wt (g/mol)

!VmaxC & Km starting values - final values are estimated from set of vapor uptake curves

set KFC = 1.18 @min=0.1 @max=2.2 ! L/hr-kg, hepatic clearance

! Chamber/Rat/Exposure Parameters}

set CONC = 50. ! Init. Inhaled Conc (ppm), experiment-specific, use 50, 200, 500 and 1150 ppm

set RATS = 1 ! Number of rats in chamber, always 1

set VCHC = 3.8 ! Closed chamber vol (l)

set KLOS = 0.04 @min=0.02 @max=0.06 ! Closed chamber loss, hr-1, experiment-specific

```
set TSTOP = 6.00 ! Length of exposure
```

```
prepare t, cp, mb
```

```
output @clear t cp mb
```

```
start @nocallback
```

```
plot @title = 'Gas Uptake GSA Setup', @xlabel = 'Time (hrs)', @ylabel = 'CP (ppm)', cp
```

```
! Global sensitivity analysis, Morris method, response is cp
```

```
gsa @method=morris @nlevels=10 @nsamps=50 @response=cp @coeffile='cp_gsa.csv' QPC QCC QLC QFC QSC BW VLC VFC VSC...
```

```
PL PF PS PR PB KFC KLOS
```

```
line @xvar=t @file='cp_gsa.csv' QPC QCC QLC QFC QSC BW VLC VFC VSC...
```

```
PL PF PS PR PB KFC KLOS
```