

Plasma urine example

Here are several examples of simultaneous modelling of plasma and urine data. The examples are adapted from a model for the analysis of Dextromethorphan data provided by Khaled Mohammed Abduljalil. The first involves data from the parent drug. The second involves both parent and metabolite data. In the control streams of these examples plasma and urine data are simulated. The same control streams could also be used for analysis of the data. MU_ modelling is used for the simulation to maintain consistency with subsequent analysis runs but MU_ modelling has no effect on simulation. If MU_ variables are renamed to avoid the character "_" (e.g., rename MU_1 as MU1), the same control streams may be run with NONMEM VI.

EXAMPLE 1 sim_parent_a2.ctl

The data dextroparent.dat contains data from only the parent drug. The parent drug is eliminated directly to urine, and also via conversion to metabolite, but there are no measurements of metabolite in plasma or urine. The fraction F0 of parent drug in the urine is computed as in Guide V (See (4.6) in Chapter 4.3.1 and (4.7) in Chapter 4.3.4.1).

The L2 data item is not required, but is used because multiple observations at the same TIME are in effect multivariate observations. Values of SIGMA corresponding to unused elements EPS(3) and EPS(4) are defined so that the EPS vector has the same length in both examples. See Remarks below.

Although ADVAN2 is used, the same control stream could be implemented with ADVAN5 or ADVAN6, with minor changes.

```
$PROBLEM Parent drug, using ADVAN2
$INPUT ID TIME AMT UVOL DV CMT MDV EVID L2
$DATA dextroparent.dat IGNORE=#
$SUBROUTINES ADVAN2 TRANS1
$PK
  K12=THETA(1)
  KA=K12
  MU_1=LOG(THETA(2))
  V2=EXP(MU_1+ETA(1))
  MU_2=LOG(THETA(3))
  CLP=EXP(MU_2+ETA(2)) ; RENAL CL FOR PARENT
  CLB=THETA(4) ; METABOLIC CL FOR METABOLIC
  K23=CLP/V2+CLB/V2
  K=K23
  F0=CLP/(CLP+CLB)
  S2=V2
  S3=UVOL
$ERROR (EVERY EVENT)
  ACMT=ABS(CMT) ; output compartment may have negative value of CMT
  IF(ACMT.EQ.2) Y=F*(1+EPS(1))
  IF(ACMT.EQ.3) Y=F*(1+EPS(2))
$THETA
  (0.01,0.8,6) ;KA
  (0.01,43,1000);V2
  (0.0001,20,190);CLP
  (0.01,15,90);CLB
$OMEGA
  0.05 0.05
$SIGMA
  .01 .01
  .01 .01 ; eps(3) and eps(4) for consistent EPS with sim_metab_a6 example
$SIM (111111) ONLYSIM
$TABLE ID TIME AMT UVOL DV SIMP=PRED CMT MDV EVID L2 NOAPPEND FILE=sim_parent_a2.tab
```

A fragment of the data file is as follows:

#	ID	TIME	AMT	UVOL	DV	CMT	MDV	EVID	L2
2.000		0.000	30000.000	0.000	0.000	1.000	1.000	1.000	1

2.000	0.171	0.000	0.000	0.000	3.000	1.000	2.000	1
2.000	2.000	0.000	0.000	0.000	2.000	0.000	0.000	1
2.000	2.000	0.000	93.100	0.000	-3.000	0.000	0.000	1
2.000	2.000	0.000	0.000	0.000	3.000	1.000	2.000	2
2.000	3.150	0.000	0.000	0.000	2.000	0.000	0.000	2
2.000	3.150	0.000	133.600	0.000	-3.000	0.000	0.000	2

There is a dose at TIME=0. At TIME=.171, the urine compartment (which is the default compartment for output) is turned on. At TIME=2, there are observations of both plasma and urine. The urine amount is reset to 0 and the compartment turned back on for the start of a new urine collection. At TIME 3.15 there are observations of plasma and urine.

In the model, different EPS variables are used for the two types of observations. In the table, values of PRED (SIMP) are computed with ETA and EPS equal to 0. The DV values are computed with simulated ETA and EPS, and would be used as the DV in subsequent analysis runs.

EXAMPLE 2 sim_metab_a6.cfl

The data dextrometab.dat includes measurements of metabolite in plasma and urine. The model also include predictions of the metabolite in both plasma (CMT=4) and urine (CMT=5; the default compartment for output). Different compartments are used for parent and metabolite in urine, but the same value of UVOL applies to both. The F0 parameter is not needed because the model for K23 now uses only CLP, the clearance of the parent drug. No analytic ADVAN model has two output compartments, so a general ADVAN (ADVAN6 or ADVAN5) must be used.

Note that Compartment 3 is no longer the default for output. Instead of being computed by PREDPP using mass balance, it is computed explicitly by the ADVAN routine itself (e.g., using a differential equation when ADVAN6 is used). If compartment 3 is defined in \$MODEL with default attributes, e.g.,

```
COMP=(DEXURIN)
```

there is an error message from PREDPP:

```
SPECIFIED COMPARTMENT MAY NOT BE TURNED OFF WITH AN OBSERVATION RECORD
```

Instead, the compartment may be defined as an output-type compartment:

```
COMP=(DEXURIN INITIALOFF NODOSE)
```

Now observations with CMT=-3 are permitted and no change to the original data records is needed.

```
$PROBLEM Parent drug and metabolite, using ADVAN6
```

```
$INPUT ID TIME AMT UVOL DV CMT MDV EVID L2
```

```
$DATA dextrometab.dat IGNORE=#
```

```
$SUBROUTINES ADVAN6 TRANS1 TOL=4
```

```
$MODEL
```

```
COMP=(DEPOT)
```

```
COMP=(PLASMA DEFOBS) ; PARENT IN PLASMA
```

```
COMP=(DEXURIN INITIALOFF NODOSE) ;PARENT IN URINE
```

```
COMP=METAB ;METABOLITE IN PLASMA
```

```
$PK
```

```
K12=THETA(1)
```

```
MU_1=LOG(THETA(2))
```

```
V2=EXP(MU_1+ETA(1))
```

```
MU_2=LOG(THETA(3))
```

```
CLP=EXP(MU_2+ETA(2)) ; RENAL CL FOR PARENT
```

```
CLB=THETA(4) ; METABOLIC CL FOR METABOLIC
```

```
CLMR=THETA(5) ; RENAL CL FOR METABOLITE
```

```
V4=1
```

```
K24=CLB/V2
```

```
K23=CLP/V2
```

```
; F0=CLP/(CLP+CLB) Omit F0 because parent and metab have different urine compts.
```

```
K45=CLMR/V4
```

```
S2=V2
```

```
S4=V4
```

```
S3=UVOL
```

```
S5=UVOL
```

```

$ERROR (EVERY EVENT)
  ACMT=ABS(CMT) ; output compartment may have negative value of CMT
  IF(ACMT.EQ.2) Y=F*(1+EPS(1))
  IF(ACMT.EQ.3) Y=F*(1+EPS(2))
  IF(ACMT.EQ.4) Y=F*(1+EPS(3))
  IF(ACMT.EQ.5) Y=F*(1+EPS(4))
$DES
  DADT(1)=-K12*A(1)
  DADT(2)=K12*A(1)-K23*A(2)-K24*A(2)
  DADT(3)=K23*A(2)
  DADT(4)=K24*A(2)-K45*A(4)
$THETA
  (0.01,0.8,6) ;KA
  (0.01,43,1000);V2
  (0.0001,20,190);CLP
  (0.01,15,90);CLB
  (0.0001,5,90);CLMR
$OMEGA
  0.05 0.05
$SIGMA
  .01 .01 .01 .01
$SIM (111111) ONLYSIM
$TABLE ID TIME AMT UVOL DV SIMP=PRED CMT MDV EVID L2 NOAPPEND FILE=sim_metab_a6.tab

```

A fragment of the data dextrometab.dat is:

#	ID	TIME	AMT	UVOL	DV	CMT	MDV	EVID	L2
2.000		0.000	30000.000	0.000	0.000	1.000	1.000	1.000	1
2.000		0.171	0.000	0.000	0.000	3.000	1.000	2.000	1
2.000		0.171	0.000	0.000	0.000	5.000	1.000	2.000	1
2.000		2.000	0.000	0.000	0.000	2.000	0.000	0.000	1
2.000		2.000	0.000	0.000	0.000	4.000	0.000	0.000	1
2.000		2.000	0.000	93.100	0.000	-3.000	0.000	0.000	1
2.000		2.000	0.000	93.100	0.000	-5.000	0.000	0.000	1
2.000		2.000	0.000	0.000	0.000	3.000	1.000	2.000	2
2.000		2.000	0.000	0.000	0.000	5.000	1.000	2.000	2
2.000		3.150	0.000	0.000	0.000	2.000	0.000	0.000	2
2.000		3.150	0.000	0.000	0.000	4.000	0.000	0.000	2
2.000		3.150	0.000	133.600	0.000	-3.000	0.000	0.000	2
2.000		3.150	0.000	133.600	0.000	-5.000	0.000	0.000	2

Observations of metabolite in plasma (CMT=4) and metabolite in urine (CMT=5) are present at the same values of TIME as observations of parent in plasma (CMT=2) and parent in urine (CMT=3), respectively, and have the same values of L2.

Remarks:

After the control streams are run, the predictions SIMP for compartments 2 and 3 are the same in the two table files sim_par-ent_a2.tab and sim_metab_a6.tab, as expected, because they are computed with EPS=0, i.e., they are not simulated values. Because of the use of L2 and the same-length EPS vector in both examples, the simulated DV's for compartments 2 and 3 are also the same. This is useful for illustrative and debugging purposes, but is not usually the case when the EPS structure is different, or the L2 data item is not used.

To summarize:

With L2, during simulation SIMEPS generates a new set of values for EPS only when L2 changes value.

Without L2, during simulation SIMEPS generates a new set of values for EPS with every record, even when MDV is 1 or the PREDPP item EVID is not 0.

(See **simeps**)

All control streams and data files are found in the NONMEM 7.4 examples directory.

dextroparent.dat (also displayed here)

dextro_sim_parent_a2.ctl (also displayed here)

dextro_sim_parent_a5.ctl

dextro_sim_parent_a6.ctl

dextrometab.dat (also displayed here)

dextro_sim_metab_a5.ctl

dextro_sim_metab_a6.ctl (also displayed here)

REFERENCES: None.