



UPPSALA
UNIVERSITET

A strategy for residual error modeling incorporating both scedasticity of variance and distribution shape

Anne-Gaëlle Dosne,

Ron J. Keizer, Martin Bergstrand, Mats O. Karlsson

Pharmacometrics Research Group

Department of Pharmaceutical Biosciences

Uppsala University

Sweden



Traditional error modeling

Models and assumptions

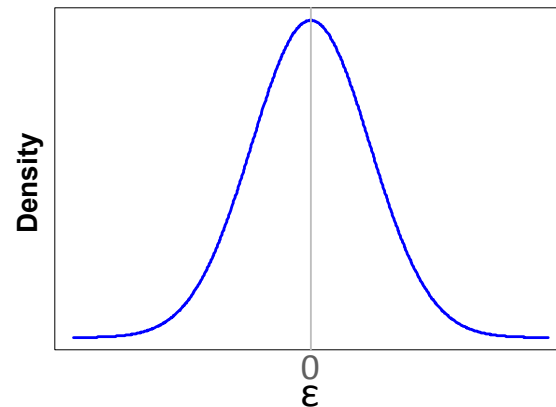
- Models**

$$y = f(\theta, x, Z) + \varepsilon$$

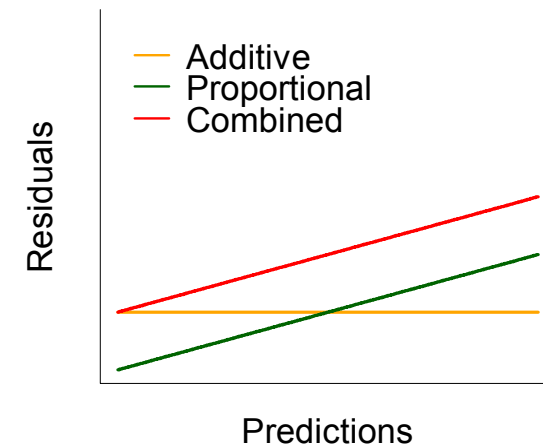
$$\varepsilon \sim N(0, \text{var}(\varepsilon))$$

$\text{var}(\varepsilon)$
σ_{add}^2
$\sigma_{\text{prop}}^2 \times f(\theta, x, Z)^2$
$\sigma_{\text{add}}^2 + \sigma_{\text{prop}}^2 \times f(\theta, x, Z)^2$

- Assumptions**



Normal distribution, mean 0

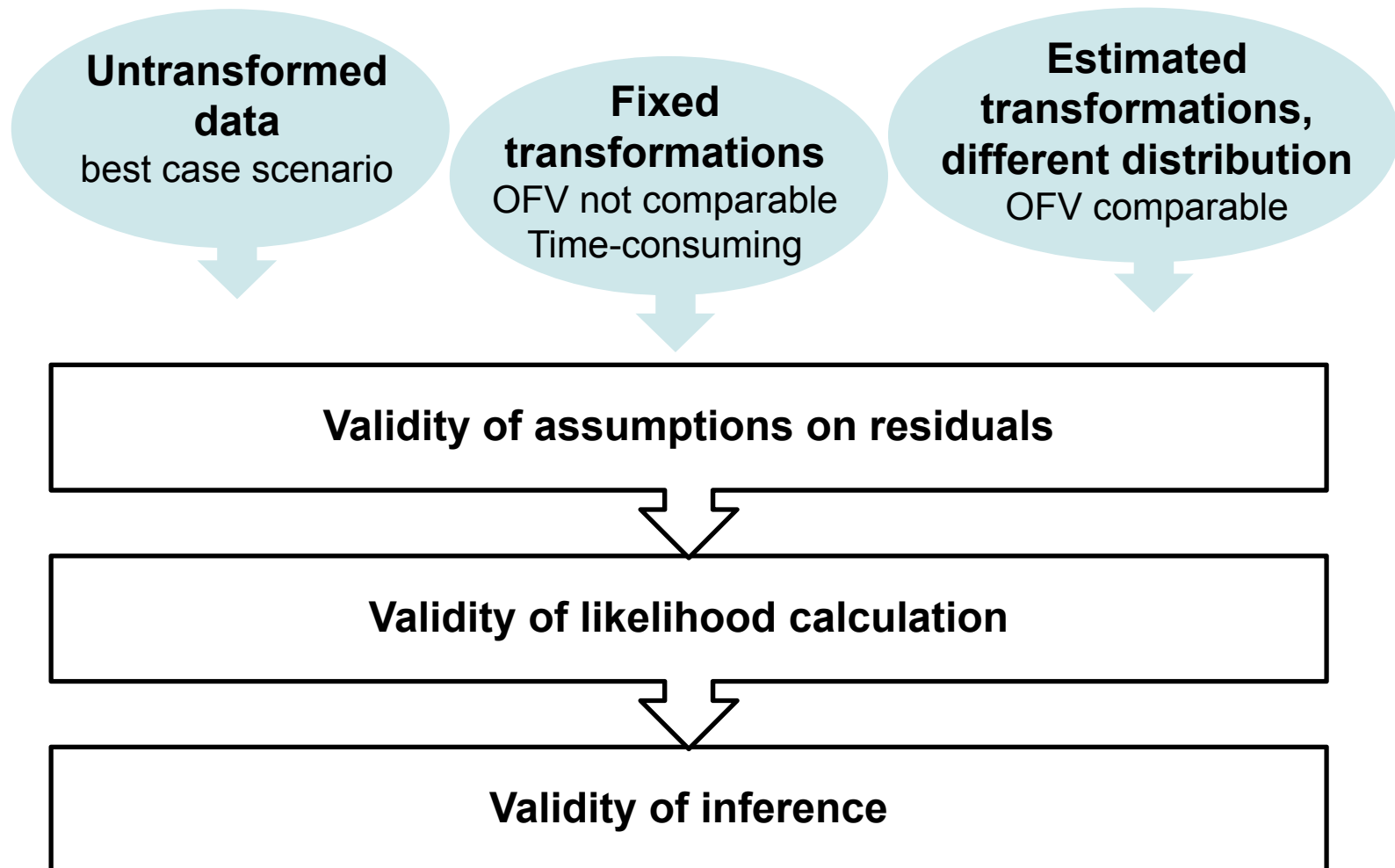


Fixed residual –
prediction relationship



What to do if assumptions do not hold?

Common answers, limitations, needs





Alternative strategies

dTBS and Student's t-distribution

1
dTBS

Transform data and prediction so ε has a symmetric density

- Box-Cox (skewness) + Power term (scedasticity)

2
t-distribution

Change distribution assumption

- Student's t-distribution (heavy-tailed)



UPPSALA
UNIVERSITET

Residual error modeling with dynamic Transform Both Sides (dTBS)



Dynamic Transform Both Sides

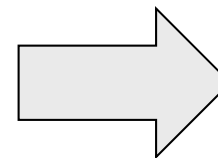
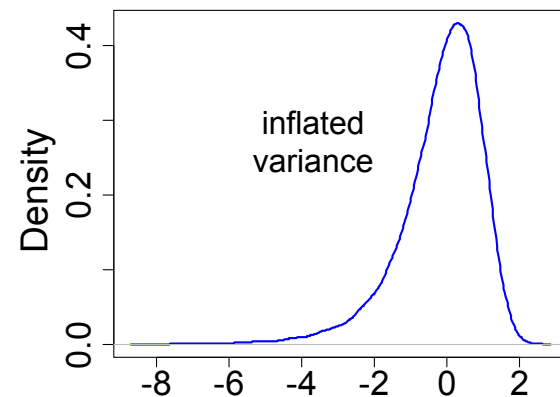
Box-Cox transformation in NLMEM

$$h(y) = \begin{cases} \lambda \neq 0 & \frac{y^\lambda - 1}{\lambda} \\ \lambda = 0 & \log(y) \end{cases}$$

$\lambda = 1$	normal
$\lambda = 0$	log
$\lambda > 1$	left skewed
$\lambda < 1$	right skewed

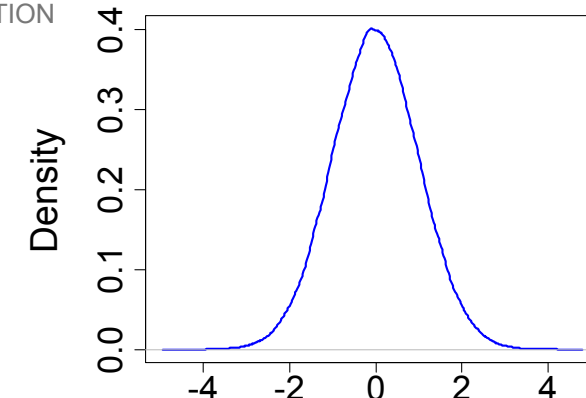
In NLMEM $h(DV) = h(IPRED) + \varepsilon$

Untransformed scale:
skewed residuals



TRANSFORMATION

Transformed scale:
normal residuals





Dynamic Transform Both Sides

Likelihood calculation given the transformation

- Likelihood of *untransformed* data– **allows OFV comparison**^{1,2}

$$p(y) = \int p(y|b)p(b)db$$

$$p(y|b) = \underbrace{\prod \frac{1}{\sqrt{2\pi\sigma^2}} \times e^{-\frac{(h(y)-h(\hat{y}))^2}{2\sigma^2}}}_{\text{Likelihood of transformed data}} \times \underbrace{\frac{d h(y)}{dy}}_{\text{First derivative of Box Cox: } y^{\lambda-1}}$$

¹ Carroll & Ruppert. *Transformation and Weighting in Regression*. Chapman and Hall. 1998

² Oberg & Davidian. *Estimating Data Transformations in Nonlinear Mixed Effects Models*. Biometrics. 2000



Dynamic Transform Both Sides

Implementation in NONMEM

- **Dynamic** estimation of λ^1
- Additional files^{1,2}
 - data transformed during estimation
 - redefinition of likelihood

$$\underbrace{-2LL'}_{\text{Likelihood of untransformed data}} = \underbrace{-2LL}_{\text{Likelihood of transformed data}} - \underbrace{2(\lambda - 1)\log(y)}_{\text{Contribution of transformation}}$$

¹ Frame B. *Within Subject Random Effect Transformations with NONMEM VI*. Wolverine Pharmacometrics Corporation. 2009.

² Robert Bauer, adaptation courtesy for NM7



Dynamic Transform Both Sides

Power term for dynamic heteroscedasticity

- **Power dTBS:** $\frac{DV^{\lambda-1}}{\lambda} = \frac{IPRED^{\lambda-1}}{\lambda} + IPRED^{\zeta} * \varepsilon$
 $\varepsilon \sim N(0, \sigma^2)$
- Dynamic heteroscedasticity
- Parameterization: $\zeta = \lambda + \delta$

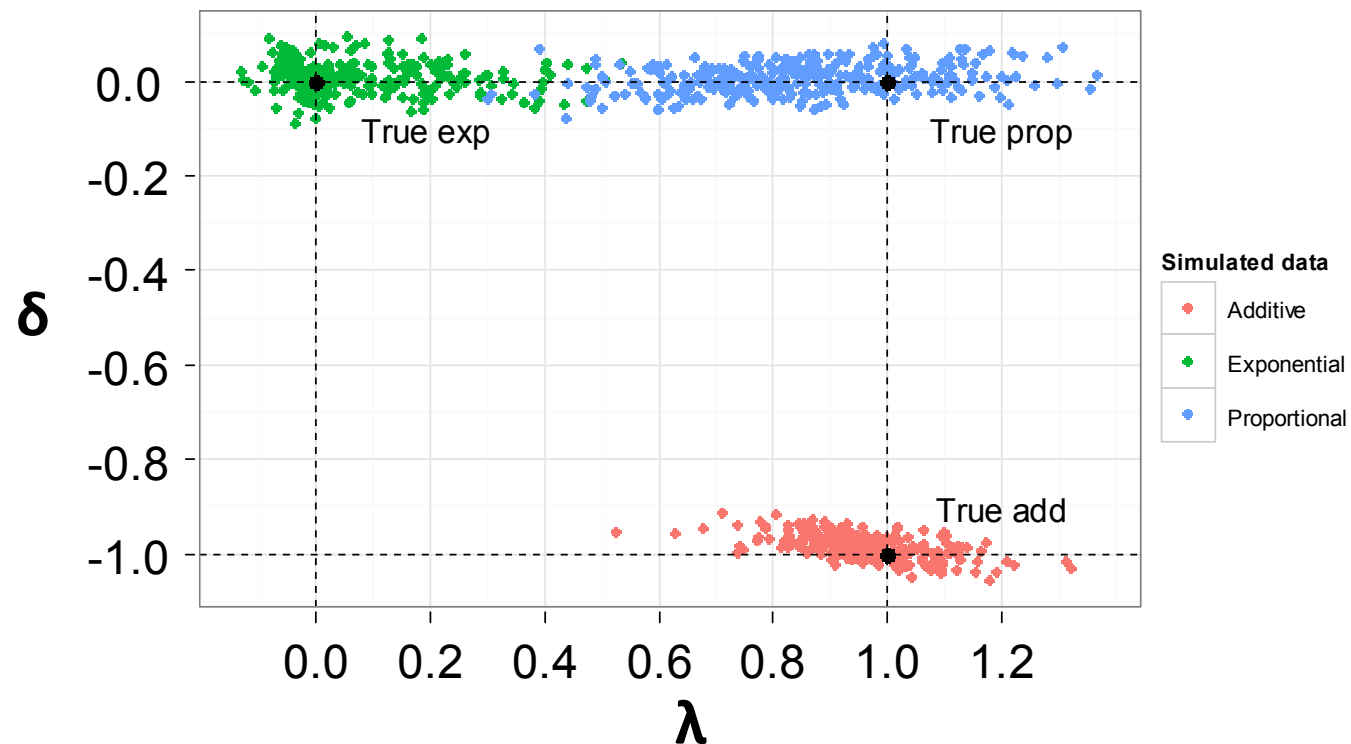
Model	Equation	$\widetilde{sd}(\varepsilon)$ untransformed scale	sd (ε) transformed scale
Power	$DV = IPRED + IPRED^{\zeta} * \varepsilon$	$\sigma \times IPRED^{\zeta}$	NA
Box-Cox	$h(DV) = h(IPRED) + \varepsilon$	$\sigma \times IPRED^{1-\lambda}$	σ
dTBS	$h(DV) = h(IPRED) + IPRED^{\zeta} * \varepsilon$	$\sigma \times IPRED^{1-\lambda+\zeta}$	$\sigma \times IPRED^{\zeta}$



Dynamic Transform Both Sides

Simulation : estimation of λ and δ

- Simulation **additive, proportional and exponential** error models
- Estimation dTBS + true model

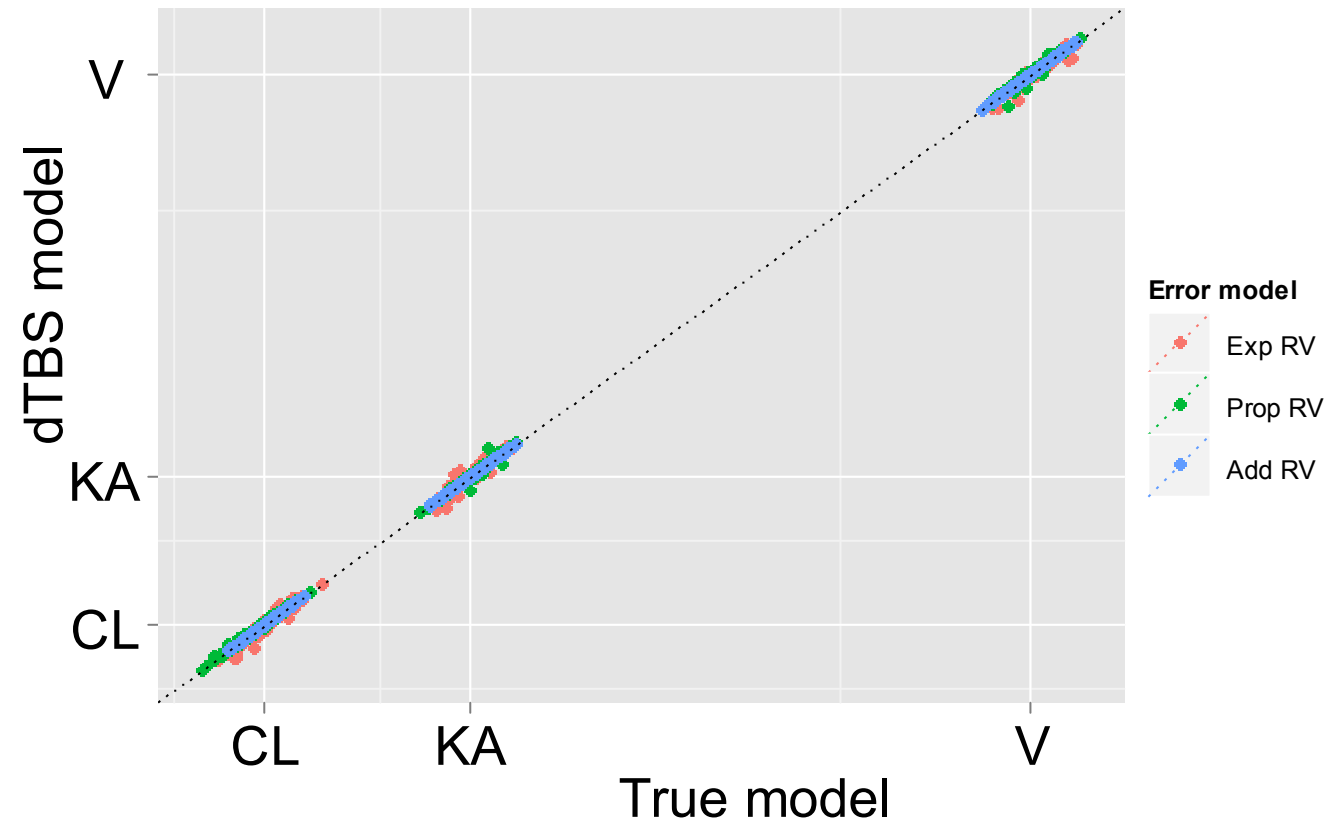




UPPSALA
UNIVERSITET

Dynamic Transform Both Sides

Simulation : satisfactory estimation of model parameters





Dynamic Transform Both Sides

Real data examples

Compound	Data	Model	RV model	Fixed transfo.	Obs.
ACTH & Cortisol	PD	turnover	combined	-	364
Cladribine	PK	IV 3 CMT	combined	-	488
Cyclophosphamide & metabolite	PK	4 CMT, CL induction	combined	-	383
Ethambutol	PK	2 CMT, transit	combined	-	1869
Moxonidine	PK	Oral 1 CMT	additive	Log	1021
Moxonidine	PD	E _{max}	additive	Log	1364
Paclitaxel	PD	Neutrophil	additive	Box-Cox	530
Pefloxacin	PK	IV 1 CMT	proportional	-	337
Phenobarbital	PK	IV 1 CMT	proportional	-	155
Prazosin	PK	Oral 1 CMT	proportional	-	887



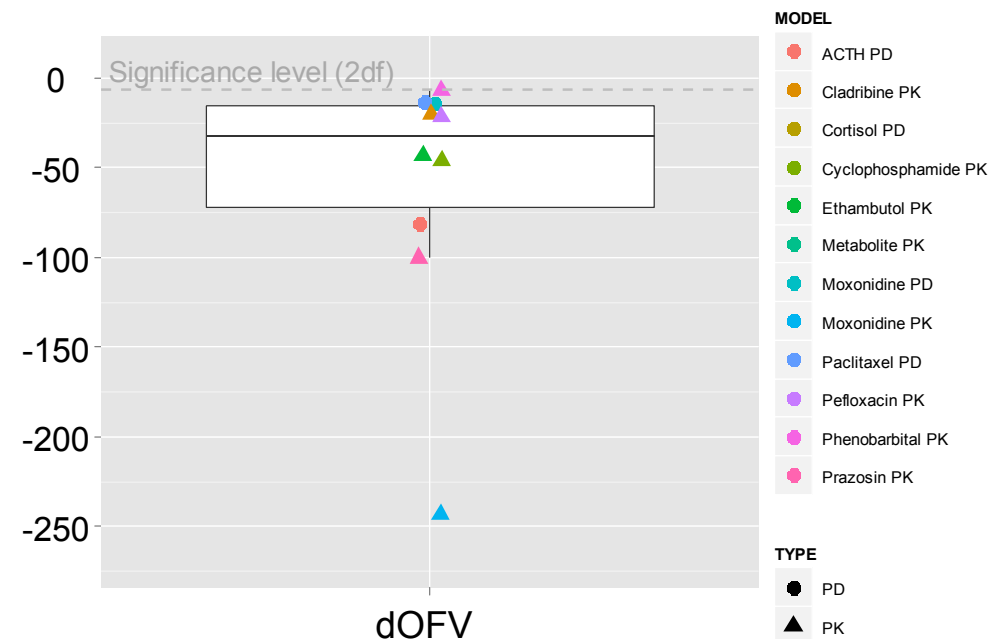
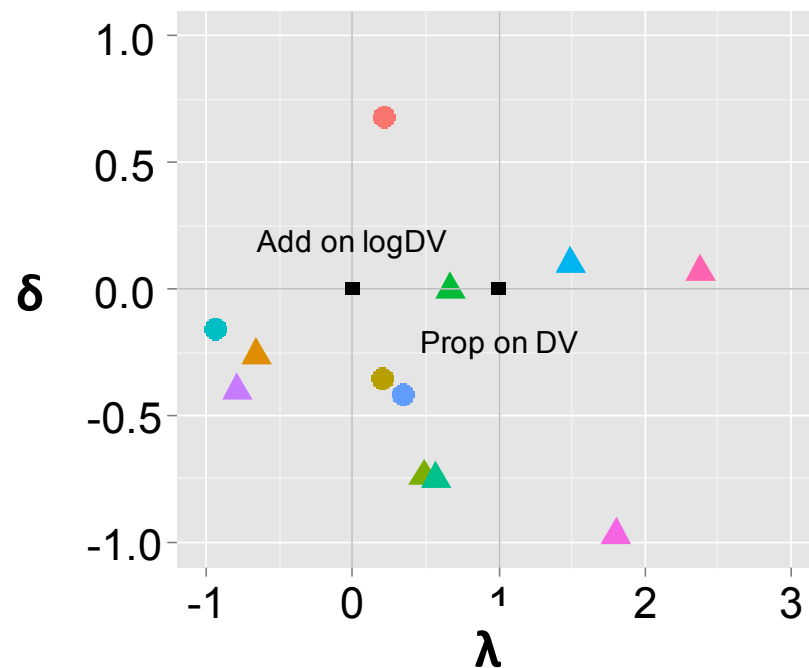
UPPSALA
UNIVERSITET

Dynamic Transform Both Sides

Real data examples: dTBS estimates & OFV drop

λ	δ
$[-0.94; 2.38]$	$[-0.97; 0.68]$

- 100% dOFV > 5.99
- **Mean drop -60**

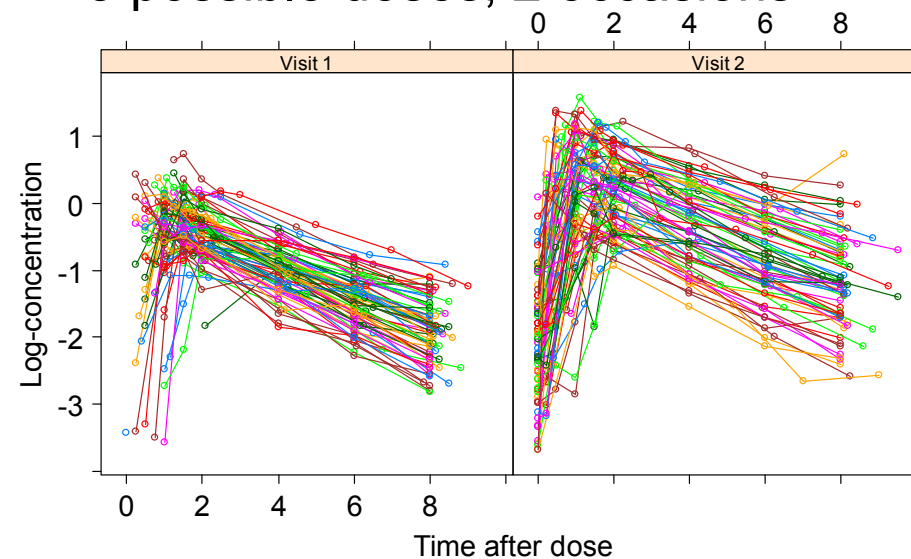




Dynamic Transform Both Sides

Moxonidine PK example: data and model

- **Data:** rich
 - 74 patients, 1021 observations
 - 3 possible doses, 2 occasions



- **Model:**
 - Additive error on log-transformed data
 - 1 CMT, first-order absorption, lag time



Dynamic Transform Both Sides

Moxonidine example: OFV and parameter estimates

Parameter	Fixed log		dTBS	
	Value	RSE(%)	Value	RSE(%)
OFV	-2173	-	-2416	-
dOFV	0	-	- 243	-
λ (-)	0	-	1.5	6.3
ζ (-)	0	-	1.6	-
$\Delta (\zeta - \lambda)$	0	-	0.1	39
CL (L.h ⁻¹)	27	1.2	26	3.4
V (L)	110	2.6	108	3.3
KA (h ⁻¹)	4.5	9	4.9	18
LAG (h)	0.24	1.3	0.23	2.3
RV (na)	0.33	1.3	0.25	4.1
IIV CL (%)	27	10	25	11
Cor IIV CL-V (-)	0.74	-	0.70	-
IIV V (%)	24	13	24	10
IIV KA (%)	168	12	136	14
IOV CL (%)	12	18	17	11
IOV KA (%)	71	18	88	16

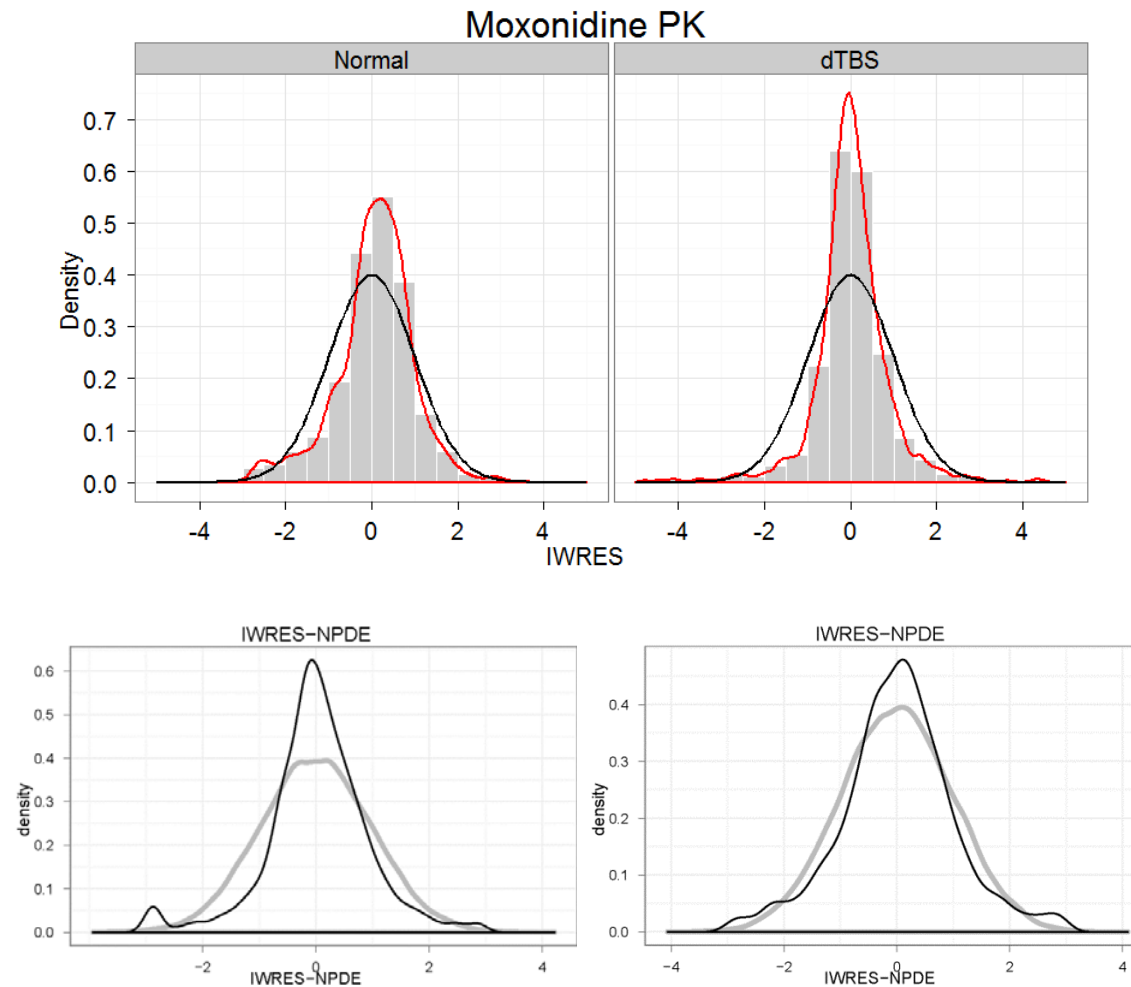
- Important OFV drop
- $\lambda \approx \zeta \approx 1.5$: left skewness
- Parameter estimates changed (RV on different scales)



UPPSALA
UNIVERSITET

Dynamic Transform Both Sides

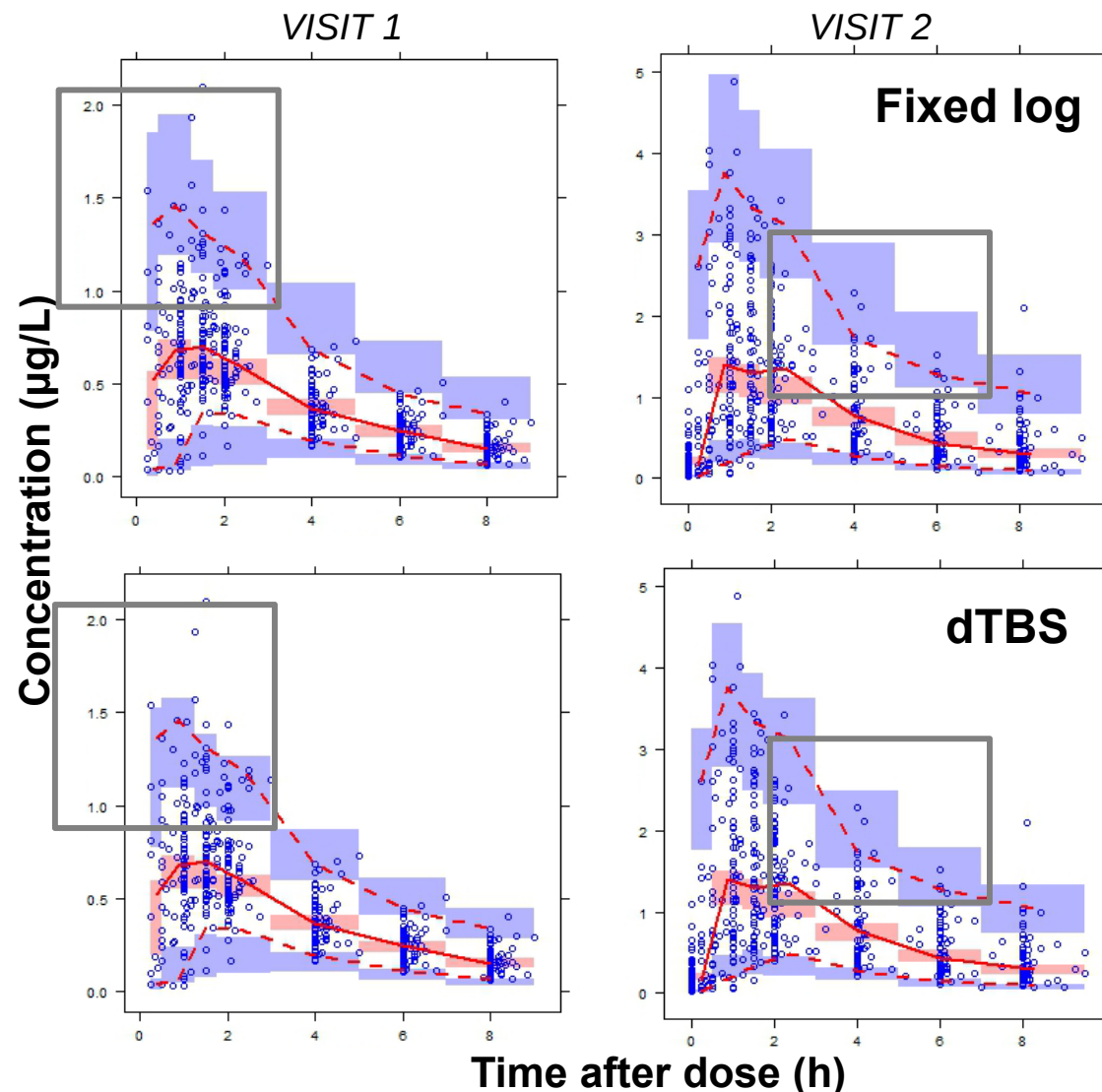
Moxonidine example: IWRES distribution





Dynamic Transform Both Sides

Moxonidine example: prediction properties



- **Narrower CI** with dTBS
- E.g. highest upper bound of CI_{95} 95th perc. = **2 vs 1.6** (log vs dTBS)
- **more precise predictions**



UPPSALA
UNIVERSITET

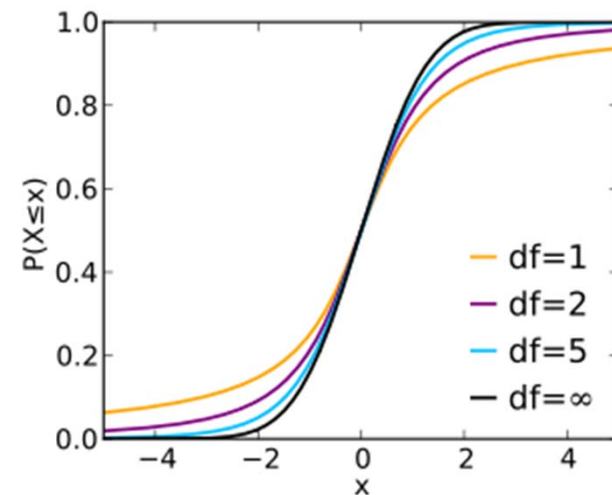
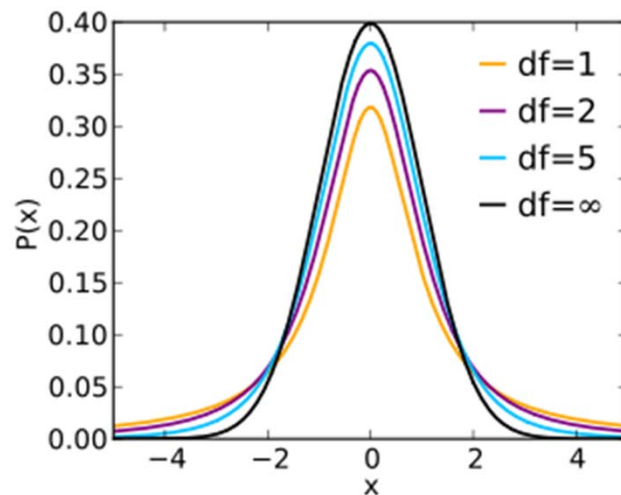
Residual error modeling with Student's t-distribution



Student's t-distribution

Function characteristics

- Parameter: **degrees of freedom ν (df)**
- Symmetric around 0, heavy-tailed
- Variance = $\frac{\nu}{\nu-2} > 1$ for $\nu > 2$
- Approaches normal distribution when $\nu \rightarrow \infty$





Student's t-distribution

Implementation: redefinition of likelihood

- **Estimation**

- Likelihood based on pdf of t-distribution¹

$$p(\mathbf{y}|\nu, \sigma, \mu) = \frac{\Gamma(\frac{\nu+1}{2})}{\Gamma(\frac{\nu}{2})\sqrt{\pi\nu\sigma^2}} \left(1 + \frac{1}{\nu} \frac{(\mathbf{x} - \mu)^2}{\sigma^2}\right)^{-\frac{\nu+1}{2}}$$

- Function Γ : Nemes approximation²
- $\nu \geq 3$: stability and full distribution definition

¹ Jackman, Simon. *Bayesian Analysis for the Social Sciences*. Wiley. 2009

² Nemes, Gergő. *New asymptotic expansion for the Gamma function*. Archiv der Mathematik . 2010



Student's t-distribution

Real data examples

Compound	Data	Model	RV model	Fixed transfo.	Obs.
Moxonidine	PK	Oral 1 CMT	additive	Log	1021
Pefloxacin	PK	IV 1 CMT	proportional	-	337
Phenobarbital	PK	IV 1 CMT	proportional	-	155
Prazosin	PK	Oral 1 CMT	proportional	-	887

Compound	dOFV	ν
Moxonidine	- 400	3
Pefloxacin	- 16	4.7
Phenobarbital	0	∞
Prazosin	- 169	3



Student's t-distribution

Moxonidine example: OFV and parameter estimates

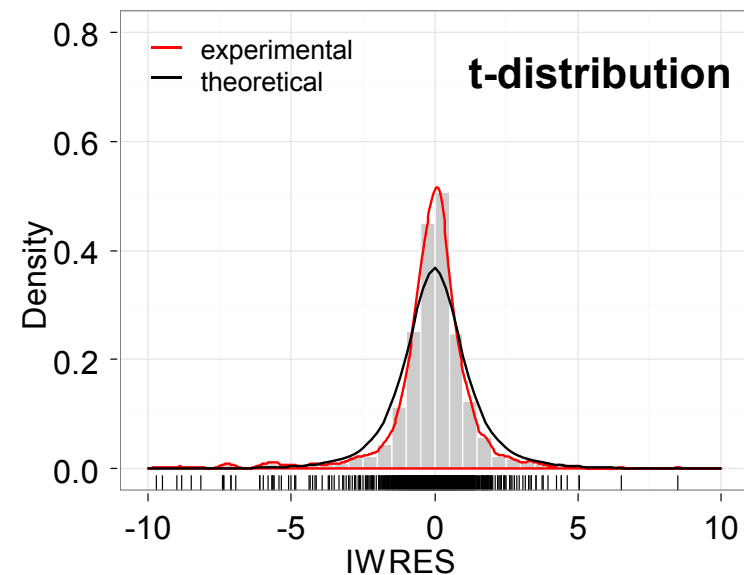
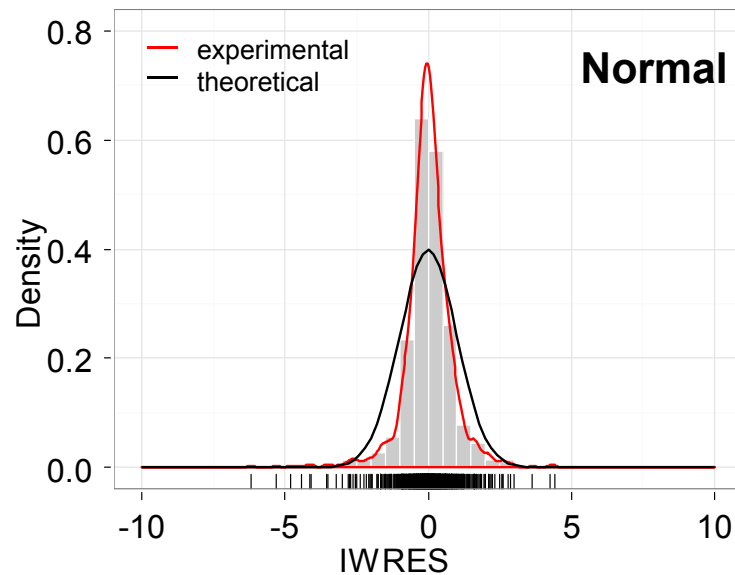
Parameter	Normal	t-distribution
OFV	1221	820
dOFV	0	- 400
DF	∞	3
CL (L.h ⁻¹)	26	26
V (L)	107	102
KA (h ⁻¹)	5.4	4.7
LAG (h)	0.24	0.24
RV (na)	0.33	0.17
IIV CL (%)	27%	28%
Cor IIV CL-V (-)	0.74	0.77
IIV V (%)	24%	23%
IIV KA (%)	162%	150%
IOV CL (%)	12%	12%
IOV KA (%)	63%	20%

- Important OFV drop
- DF at lower boundary
- Variance = 3
- Changes in absorption parameters



Student's t-distribution

Moxonidine PK: Residual distribution

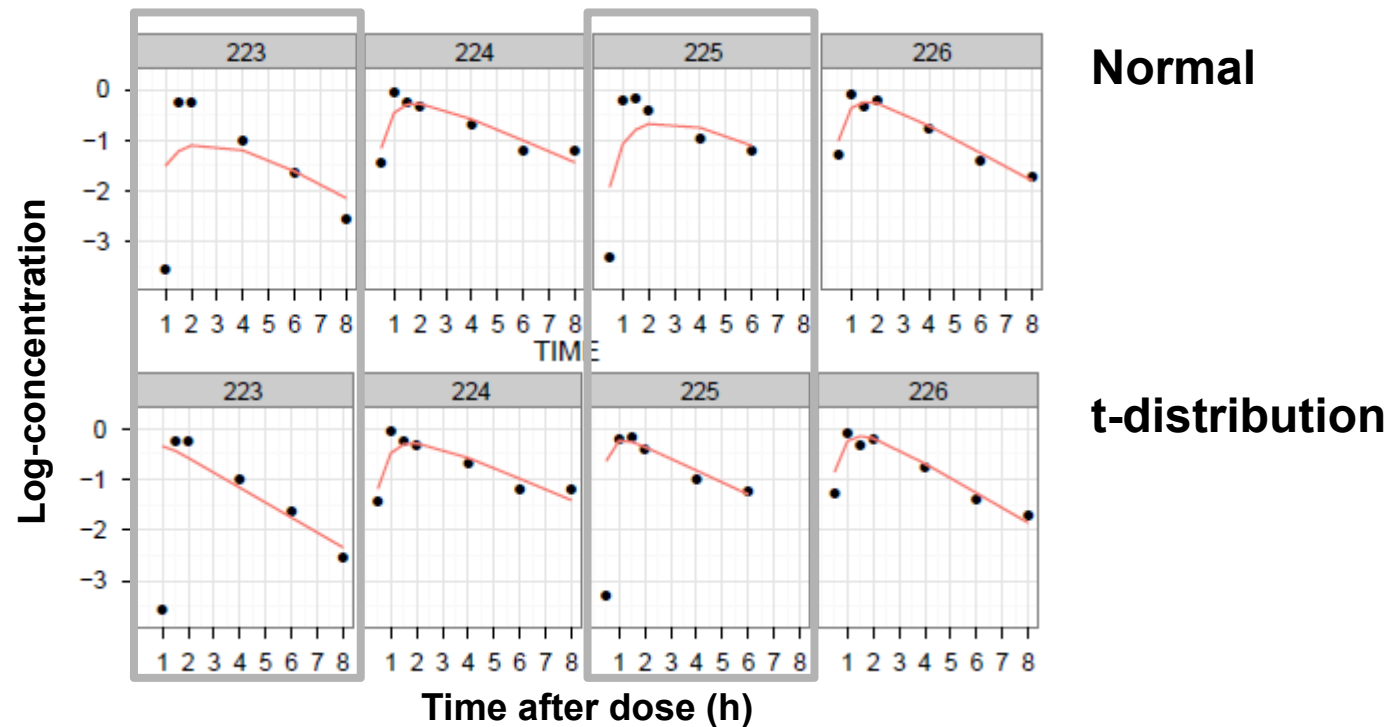


- Better agreement between experimental and theoretical IWRES distribution for t-distribution



Student's t-distribution

Moxonidine example: Individual fits



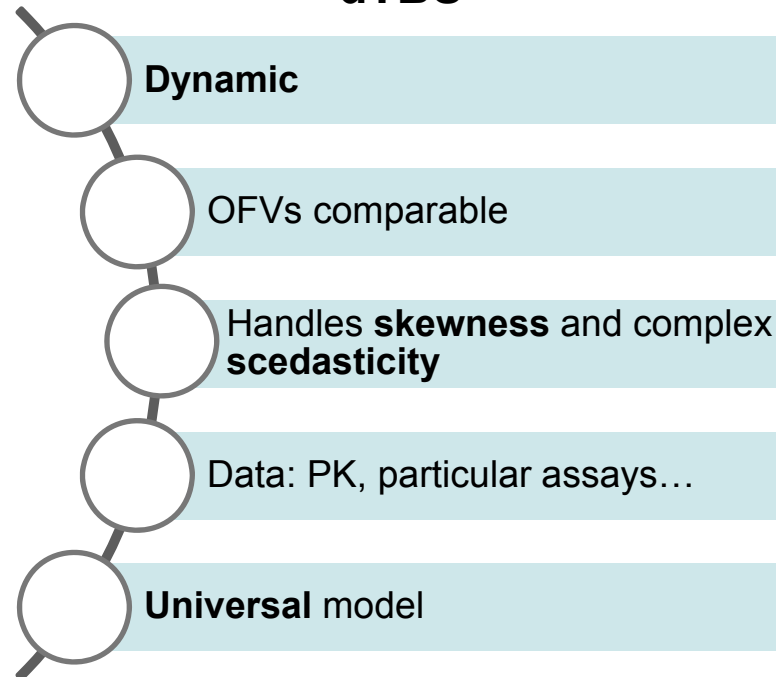
- Better global fit by allowing some predictions to be further away from observations



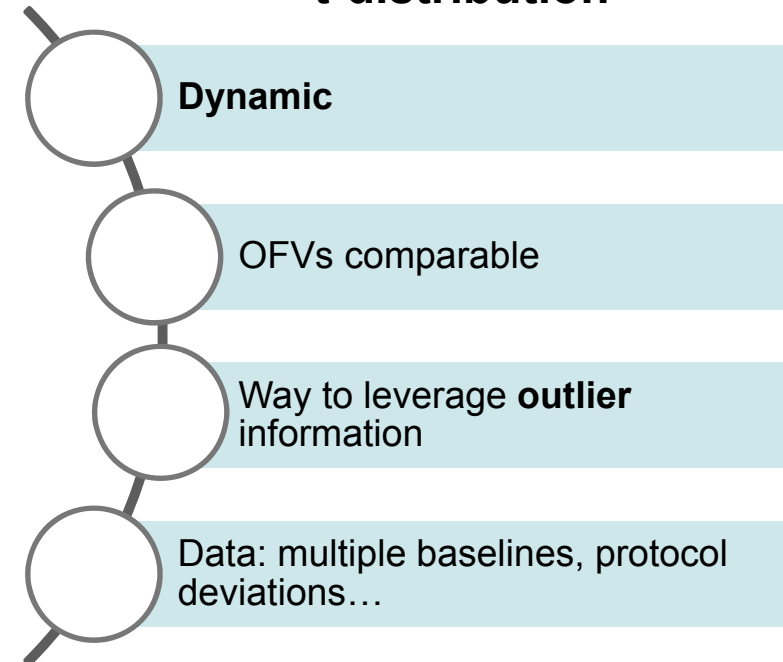
Conclusion

New possibilities for residual error modeling

dTBS



t-distribution



- Could be used jointly
- Can improve quality of parameter estimates and predictions



UPPSALA
UNIVERSITET

Acknowledgements

The research leading to these results has received support from the Innovative Medicines Initiative Joint Undertaking under grant agreement n° 115156, resources of which are composed of financial contributions from the European Union's Seventh Framework Programme (FP7/2007-2013) and EFPIA companies' in kind contribution. The DDMoRe project is also supported by financial contribution from Academic and SME partners. This work does not necessarily represent the view of all DDMoRe partners.

- Uppsala Pharmacometrics Research Group
- Novartis for financial support





UPPSALA
UNIVERSITET

Additional slides

- *dTBS ccontra and contr files*
- *dTBS model file code*
- *dTBS simulation code*
- *t-distribution simulation code*
- *Type I error investigation for dTBS and t-distribution*



Dynamic Transform Both Sides

ccontra and contr files

ccontra

```
subroutine ccontr (icall,c1,c2,c3,ier1,ier2)
  USE ROCM_REAL, ONLY: theta=>THETAC,y=>DV_ITM2
  USE NM_INTERFACE,ONLY: CELS
!   parameter (lth=40,lvr=30,no=50)
!   common /rocm0/ theta (lth)
!   common /rocm4/ y
!   double precision c1,c2,c3,theta,y,w,one,two
double precision c1,c2,c3,w,one,two
dimension c2(:),c3(:, :)
data one,two/1.,2./
if (icall.le.1) return
w=y(1)

if(theta(7).eq.0) y(1)=log(y(1))
if(theta(7).ne.0) y(1)=(y(1)**theta(7)-one)/theta(7)

call cels (c1,c2,c3,ier1,ier2)
y(1)=w
c1=c1-two*(theta(7)-one)*log(y(:l))

return
end
```

DV transformation

Redefinition of log-likelihood

contr

```
subroutine contr
(icall,cnt,ier1,ier2)
  double precision cnt
  call ncontr
(cnt,ier1,ier2,l2r)
  return
end
```



UPPSALA
UNIVERSITET

Dynamic Transform Both Sides

Model file example: Modification of \$ERROR

Power RUV

*Transform. of
IPRED*

\$ ERROR

```
IPR1 = A(2)/V
IF(IPR1.LE.0) IPR1 = 0.001
```

```
LAMBDA = THETA(7)
ZETA    = LAMBDA + THETA(6)
```

```
W      = THETA(5)*IPR1**ZETA
```

```
IPRED = IPR1
```

```
IPRTR=IPRED
```

```
IF (LAMBDA .NE. 0 .AND. IPRED .NE.0) THEN
  IPRTR=(IPRED**LAMBDA-1)/LAMBDA
```

```
ENDIF
```

```
IF (LAMBDA .EQ. 0 .AND. IPRED .NE.0) THEN
  IPRTR=LOG(IPRED)
```

```
ENDIF
```

```
IF (LAMBDA .NE. 0 .AND. IPRED .EQ.0) THEN
  IPRTR=-1/LAMBDA
```

```
ENDIF
```

```
IF (LAMBDA .EQ. 0 .AND. IPRED .EQ.0) THEN
  IPRTR=-10000000000
```

```
ENDIF
```

```
IPRED=IPRTR
```

```
>IRES = IPRED-DV
```

```
IWRES = IRES/W
```

```
IWRTR=IWRES
```

Transform. of IWRES

```
IF (LAMBDA.NE.0 .AND. DV.NE.0 .AND.
W.NE.0) THEN
```

```
  IWRTR=((DV**LAMBDA-1)/LAMBDA-IPRED)/W
```

```
ENDIF
```

```
IF (LAMBDA.EQ.0 .AND. DV.NE.0 .AND.
W.NE.0) THEN
```

```
  IWRTR=(LOG(DV)-IPRED)/W
```

```
ENDIF
```

```
IF (LAMBDA.NE.0 .AND. DV.EQ.0 .AND.
W.NE.0) THEN
```

```
  IWRTR=(-1/LAMBDA-IPRED)/W
```

```
ENDIF
```

```
IF (LAMBDA.EQ.0 .AND. DV.EQ.0 .AND.
W.NE.0) THEN
```

```
  IWRTR=(-10000000000-IPRED)/W
```

```
ENDIF
```

```
IWRES=IWRTR
```

```
Y=IPRED+EPS(1)*W
```



UPPSALA
UNIVERSITET

Dynamic Transform Both Sides

Model file : other changes and simulation code

```
$SUBROUTINE ADVAN2 TRANS1 CONTR=contr.txt CCONTR=ccontra_nm7.txt
```

Additional files needed

```
$THETA 0.0001 ;DELTA_ZETA  
$THETA 1 ;LAMBDA
```

```
$ESTIMATION METHOD=1 INTER MAXEVALS=9999
```

dTBS SIMULATION ON UNTRANSFORMED SCALE

```
IF (ICALL.EQ.4 .AND. LAMBDA.EQ.0) THEN  
  Y=EXP(Y)  
ENDIF  
IF (ICALL.EQ.4 .AND. LAMBDA.NE.0) THEN  
  Y=((Y*LAMBDA)+1)**(1/LAMBDA)  
ENDIF
```



UPPSALA
UNIVERSITET

Student's t-distribution

Model file example

\$ERROR

```
DF      = THETA(5)                ; degrees of freedom of Student distribution
W = THETA(4)*IPRED
SIG1    = W                        ; scaling factor for standard deviation of RUV
IWRES=(DV-IPRED)/SIG1
```

```
PHI=(DF+1)/2                      ; Approximation of gamma function
INN=PHI+1/(12*PHI-1/(10*PHI))
GAMMA=SQRT(2*3.14159265/PHI)*(INN/EXP(1))**PHI
```

```
PHI2=DF/2                         ; Approximation of gamma function
INN2=PHI2+1/(12*PHI2-1/(10*PHI2))
GAMMA2=SQRT(2*3.14159265/PHI2)*(INN2/EXP(1))**PHI2
```

```
COEFF=GAMMA/(GAMMA2*SQRT(DF*3.14159265))/SIG1
BASE=1+IWRES*IWRES/DF
IF(BASE.EQ.0) BASE=0.000001
POW=-(DF+1)/2
L=COEFF*BASE**POW
Y=-2*LOG(L)
```

```
$EST MAXEVAL=9999 -2LL METH=1 LAPLACE
```



UPPSALA
UNIVERSITET

Student's t-distribution

Simulation code example

\$ERROR

```
IPRED = ((DOSE/V)*(KA/C))*(A-B)
```

```
W = THETA(4)*IPRED
```

```
IWRES=(DV-IPRED)/W
```

```
DF = THETA(5)
```

```
IF (ICALL.EQ.4)
```

```
Y = IPRED +
```

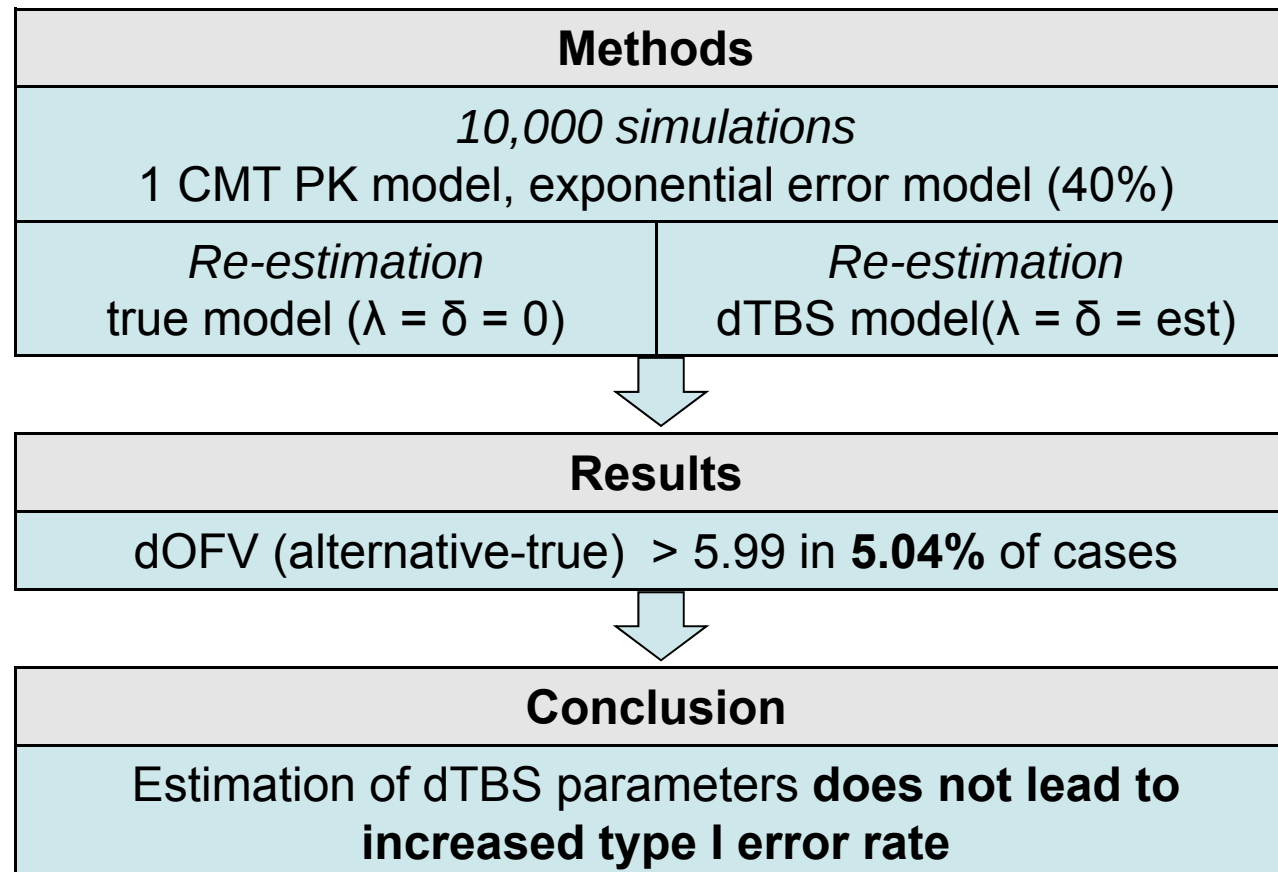
```
W*EPS(1)*(1+((EPS(1)**2+1)/(4*DF))+((5*EPS(1)**4+16*EPS(1)**2+3)/(96*DF**  
2))+((3*EPS(1)**6+19*EPS(1)**4+17*EPS(1)**2-15)/(384*DF**3)))
```

```
$SIGMA 1FIX
```



Dynamic Transform Both Sides

Type I error rate not increased





Student's t-distribution

Type I error rate not increased

