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Tutorial: Introduction to Markov modelling

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Coin toss

Assume a coin has been tossed n times

$P_{\text{tails}} = 1/2$ (if perfect coin)

Q: What is the probability of "tails" at the next toss

$P_{n+1}(\text{"tails"}) = P_{\text{tails}}$

Outcome is dependent on P_{tails} only and not dependent on outcome of previous tosses



Coin toss

Assume a coin has been tossed n times

$P_{\text{tails}} = 1/2$ (perfect coin)

**Q: What is the total number of "tails" (Σ_{n+1} ("tails"))
after the next toss**

$$P_{n+1}(\Sigma_{n+1}(\text{"tails"}) < \Sigma_n(\text{"tails"})) = 0$$

$$P_{n+1}(\Sigma_{n+1}(\text{"tails"}) = \Sigma_n(\text{"tails"})) = P_{\text{tails}}$$

$$P_{n+1}(\Sigma_{n+1}(\text{"tails"}) = \Sigma_n(\text{"tails"}) + 1) = P_{\text{tails}}$$

$$P_{n+1}(\Sigma_{n+1}(\text{"tails"}) > \Sigma_n(\text{"tails"}) + 2) = 0$$

Outcome is dependent on P_{tails} and outcomes of previous
tosses

$\Sigma_n(\text{"tails"})$ contains all necessary information about prior
history



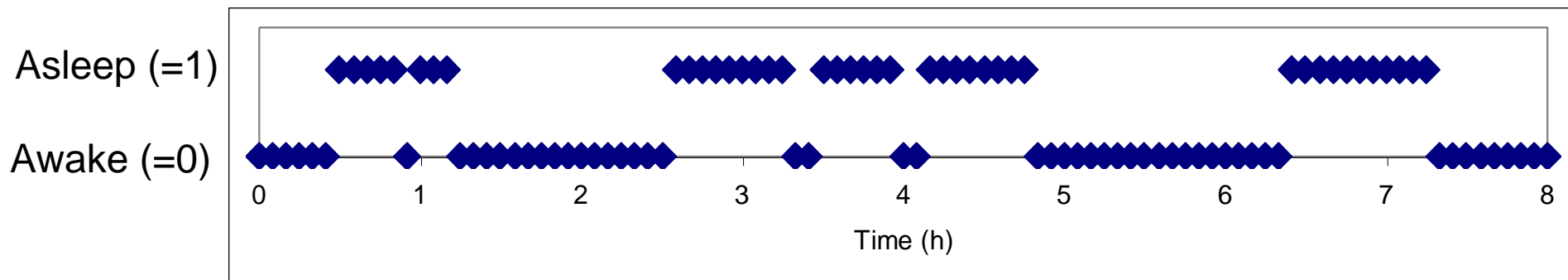
Definition Markov property

The Markov property, proposed by A.A. Markov (1856-1922), asserts that **the distribution of future outcomes depend only on the current state and not on the whole history**



Modeling of sleep/wake state

- Example: Nighttime observations of awake or sleep every 5th minute in an insomniac patient





Logistic model – NMTRAN

\$PROB Logistic model for sleep and awake

\$DATA data

\$INPUT DV

\$PRED

P1 = THETA(1)

IF(DV.EQ.1) Y=P1

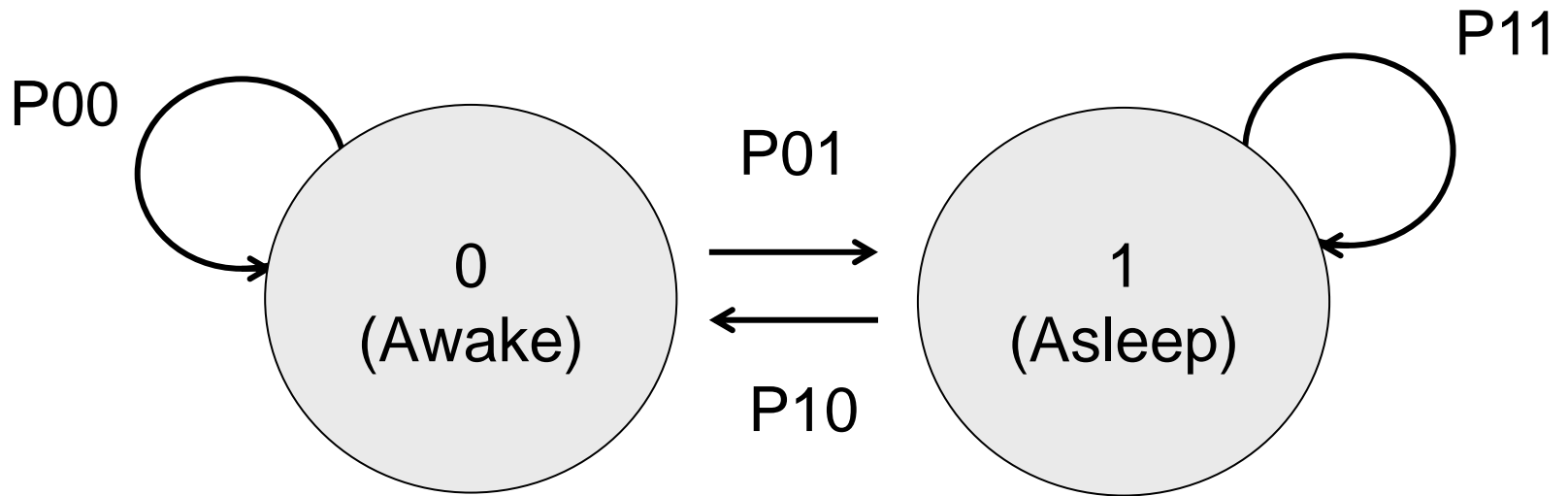
IF(DV.EQ.0) Y=1-P1

\$THETA (0,.5,1) ; PROBABILITY OF BEING ASLEEP

\$ESTIM LIKE



Two-state Markov model



$$P_{00} = 1 - P_{01}$$

$$P_{11} = 1 - P_{10}$$



Markov model – NMTRAN code

\$PROB Transition probabilities between sleep and awake

\$DATA data

\$INPUT DV PDV ;PDV=Previous DV

;PDV = Value of immediately preceding observation

\$PRED

P10 = THETA(1)

P01 = THETA(2)

IF(PDV.EQ.0.AND.DV.EQ.1) Y=P01

IF(PDV.EQ.0.AND.DV.EQ.0) Y=1-P01

IF(PDV.EQ.1.AND.DV.EQ.0) Y=P10

IF(PDV.EQ.1.AND.DV.EQ.1) Y=1-P10

\$THETA (0,.1,1) ; PROB AWAKE GIVEN ASLEEP

\$THETA (0,.1,1) ; PROB ASLEEP GIVEN AWAKE

\$ESTIM LIKE



Same model – MLXTRAN code

DESCRIPTION:

Categorical data model with Markovian dependence,
Binomial distribution

INPUT:

parameter = {p01, p11}

OBSERVATION:

```
Y = {  
    type = categorical  
    categories = {0,1}  
    dependence = Markov  
    P(Y=1 | Yp=0) = p01  
    P(Y=1 | Yp=1) = p11  
}
```

Results

Logistic model

– $P1 = 0.43 \pm 0.05$

OFV 132.7

Markov model

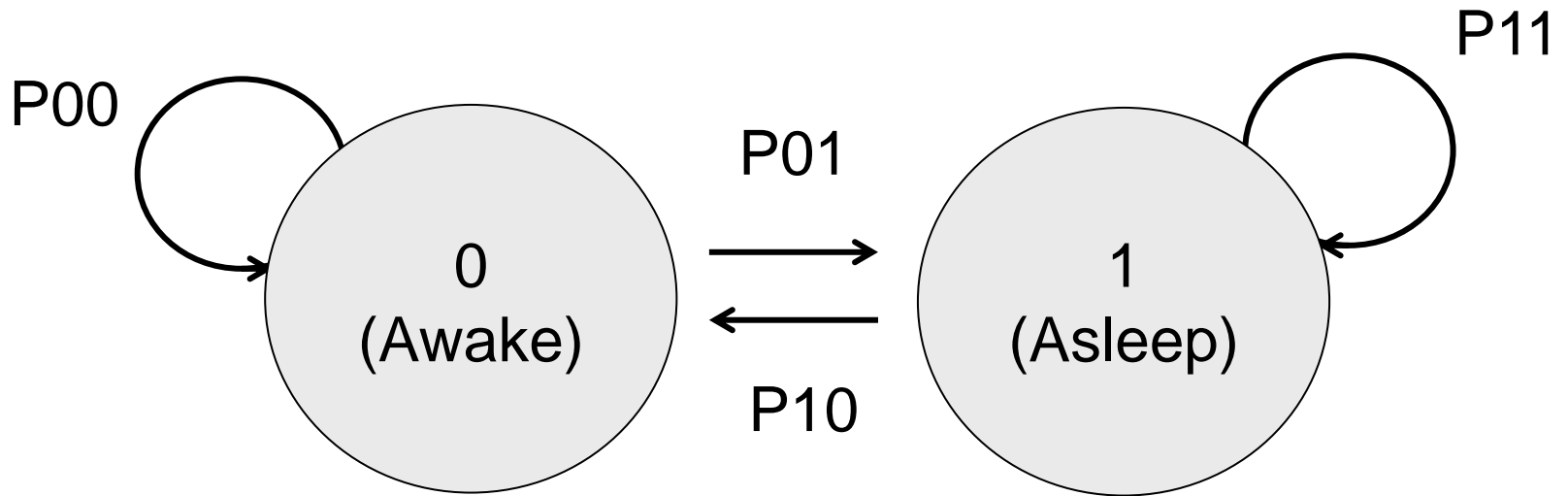
– $P01 = 0.11 \pm 0.04$

OFV 72.4

– $P10 = 0.14 \pm 0.05$



Two-state Markov model



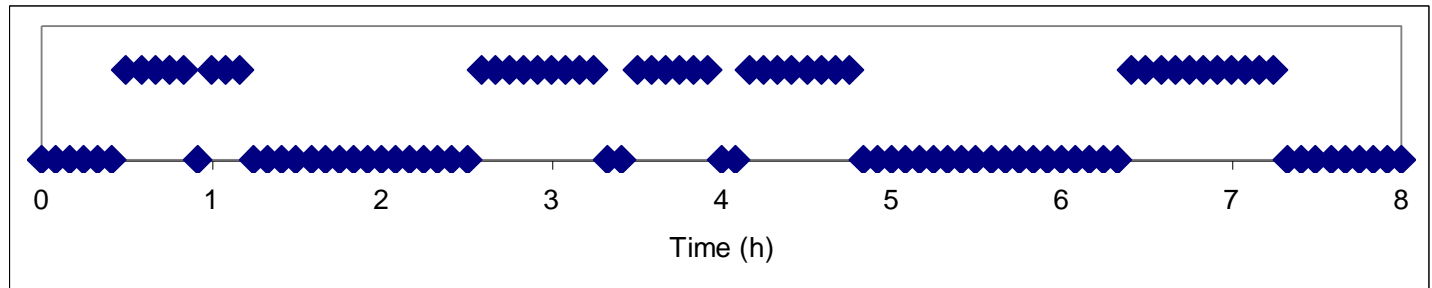
Transition Matrix

	PDV=0	PDV=1
DV=0	0.89	0.14
DV=1	0.11	0.86

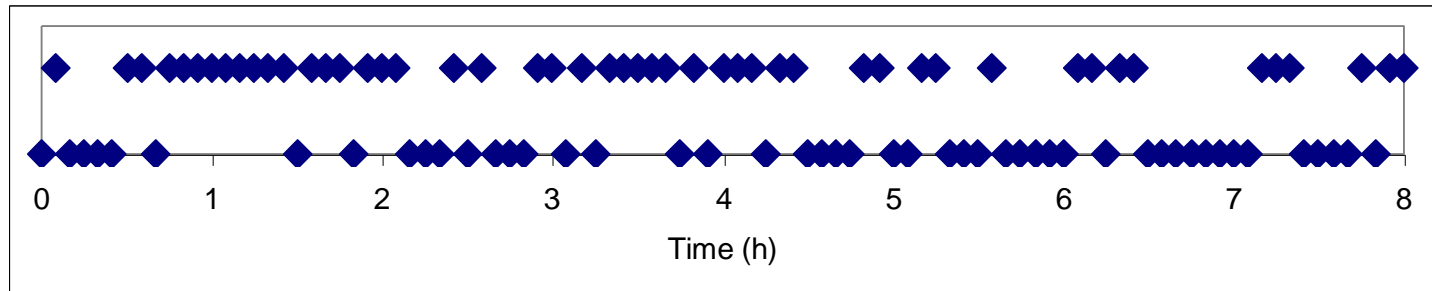


Simulations

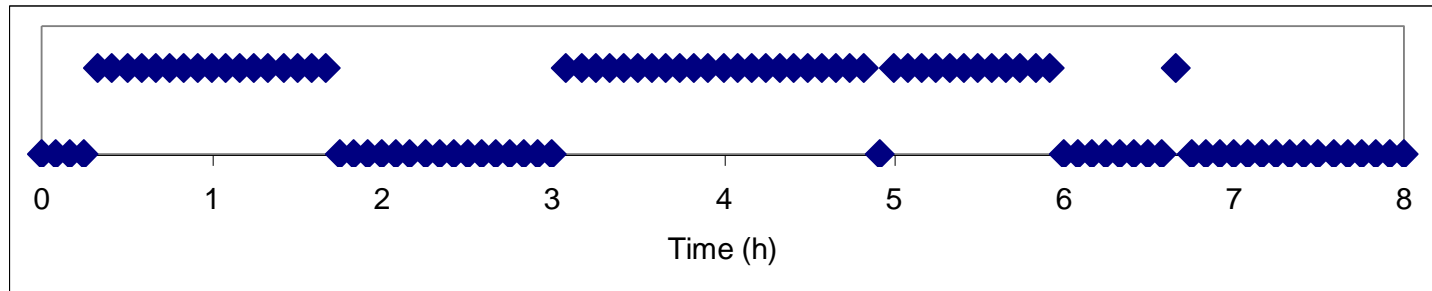
Original data



Simulation from
logistic model

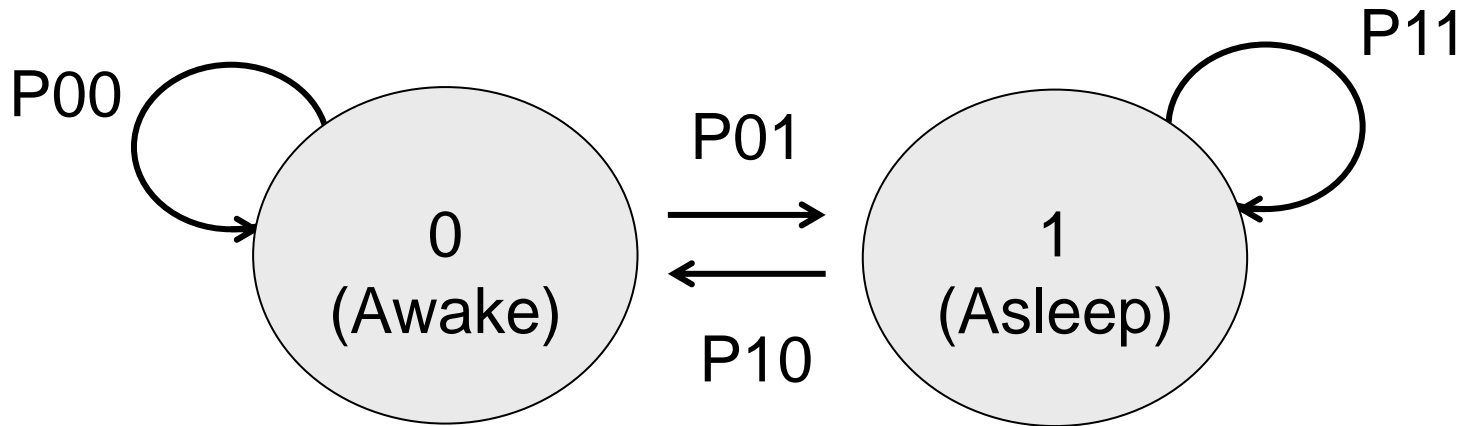


Simulation from
Markov model

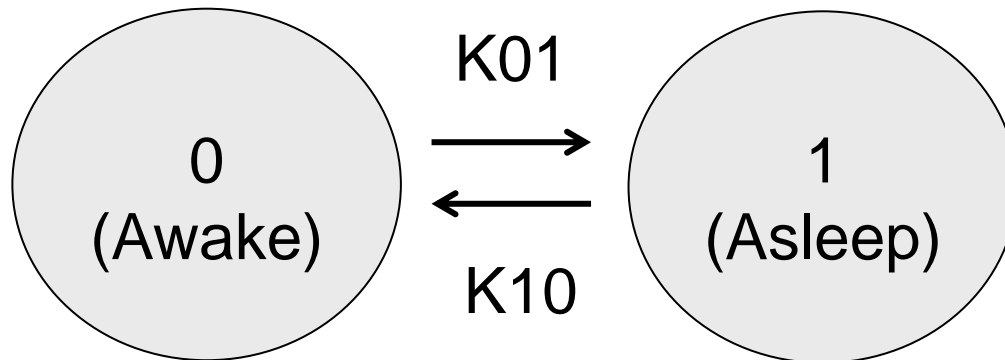




Discrete-time Markov model



Continuous-time Markov model





Continuous-time Markov model

\$PROB Continuous-time two-state Markov model for awake/asleep

\$DATA data

\$INPUT DV PDV DELTAT

\$PRED

K01 = THETA(1)

K10 = THETA(2)

P11 = (EXP(-(K10+K01)*DELTAT)*K10+K01)/(K01+K10)

P00 = (EXP(-(K10+K01)*DELTAT)*K01+K10)/(K01+K10)

IF(PDV.EQ.1.AND.DV.EQ.1) Y=P11

IF(PDV.EQ.1.AND.DV.EQ.0) Y=1-P11

IF(PDV.EQ.0.AND.DV.EQ.1) Y=P00

IF(PDV.EQ.0.AND.DV.EQ.0) Y=1-P00

\$THETA (0,1) ; K ASLEEP GIVEN AWAKE

\$THETA (0,1) ; K AWAKE GIVEN ASLEEP

\$ESTIM LIKE



Continuous–time Markov model - differential eqn parametrization

\$PROB Transition probability between awake and asleep

\$DATA data

\$INPUT DV TIME AMT CMT EVID

; "Complex" data set to initiate and empty compartments

; to reset compartment amounts after each observation

\$MODEL COMP=PRWAKE COMP=PRSLP

\$PK

K10 = THETA(1)

K01 = THETA(2)

\$DES

DADT(1) = - K01*A(1) + K10*A(2) ;Represents Probability of 0 - awake

DADT(2) = K01*A(1) - K10*A(2) ;Represents Probability of 1 - asleep

\$ERROR

IF(DV.EQ.0) Y = A(1)

IF(DV.EQ.1) Y = A(2)

\$THETA (0,1) ; K ASLEEP GIVEN AWAKE

\$THETA (0,1) ; K AWAKE GIVEN ASLEEP

\$ESTIM LIKE



Data set for differential eqn solution to Markov model

\$PROB Two-state Markov model

; Entire system updated after each observation

\$DATA data

\$INPUT DV TIME AMT CMT EVID

; . 0 1 1 1 ;awake at start - initialization

; 0 1 0 . 0 ;obs awake at T=1

; . 1 1 1 4 ;set probabilities to P(awake)=1

; 1 2 0 . 0 ;obs asleep at T=2

; . 2 1 2 4 ;set probabilities to P(asleep)=1



Data set for differential eqn solution to Markov model

\$PROB Two-state Markov model

;Data set structure with updating compartment by compartment

;May be needed when not entire system is to be updated (e.g. PKPD model)

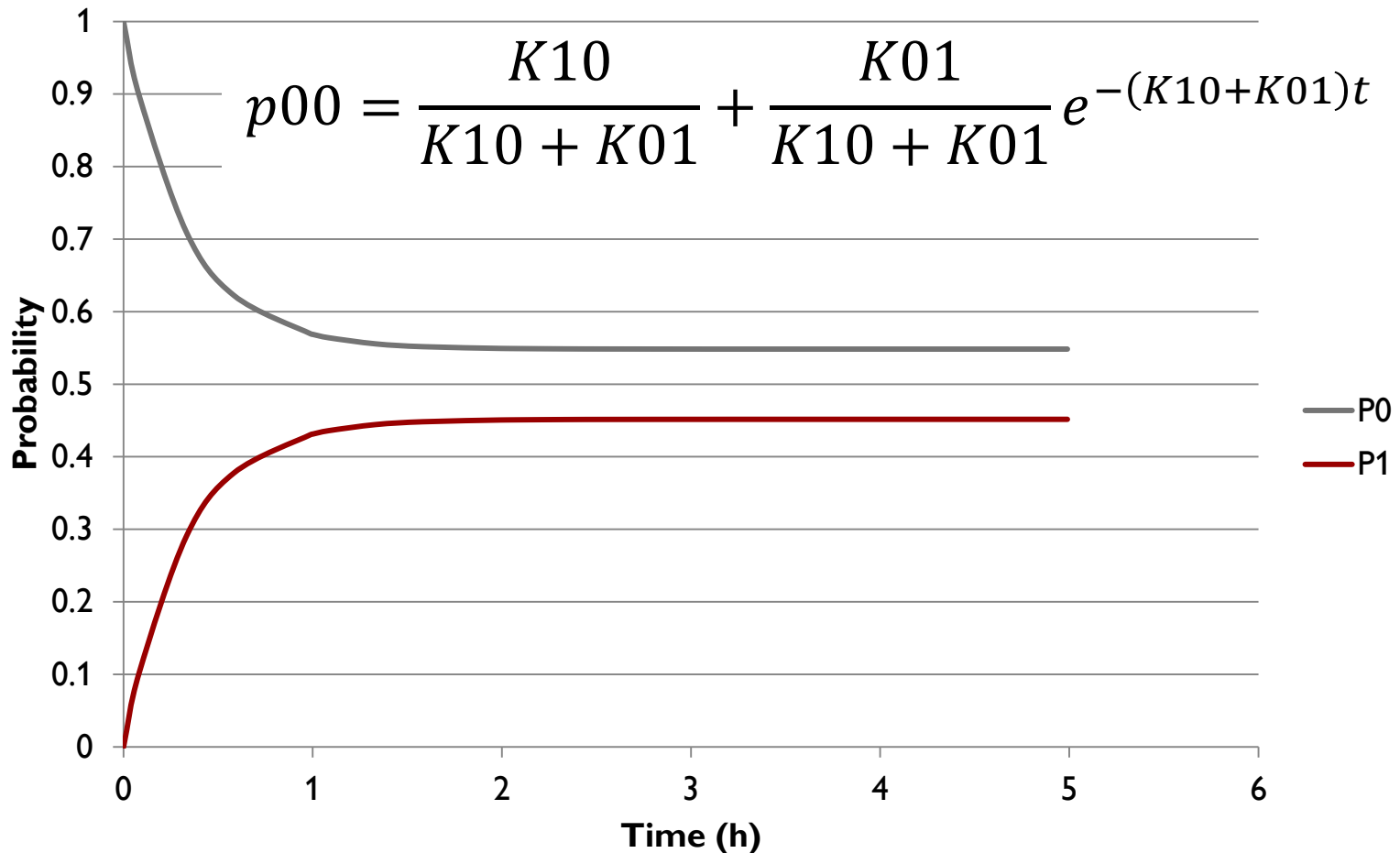
\$DATA data

\$INPUT DV TIME AMT CMT EVID

;	.	0	1	1	1	;awake at start - initialization
;	0	1	0	.	0	;awake at T=1
;	.	1	0	-1	2	; empty compartment 1
;	.	1	0	-2	2	; empty compartment 2
;	.	1	1	1	1	; reinitialize comp 1 to 1
;	.	1	0	2	2	; restart comp 2 with 0
;	1	2	0	.	0	;asleep at T=2
;	.	2	0	-1	2	; empty compartment 1
;	.	2	0	-2	2	; empty compartment 2
;	.	2	1	2	1	; reinitialize comp 2 to 1
;	.	2	0	1	2	; restart comp 1 with 0

Probabilities following an awake observation at time=0

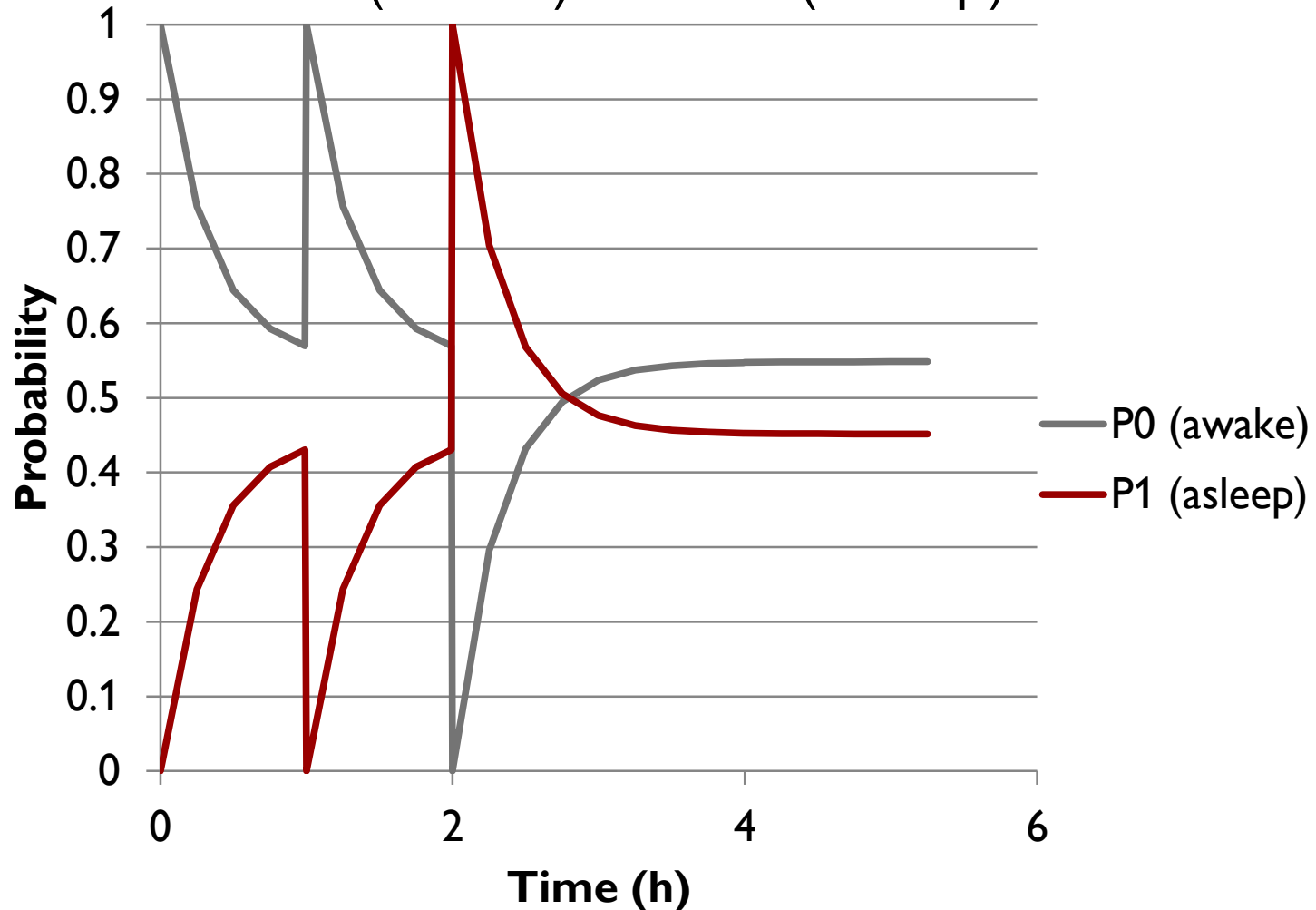
No additional observations made





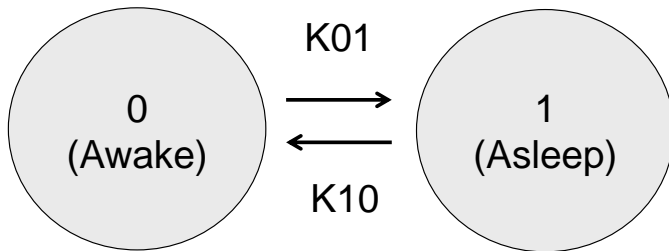
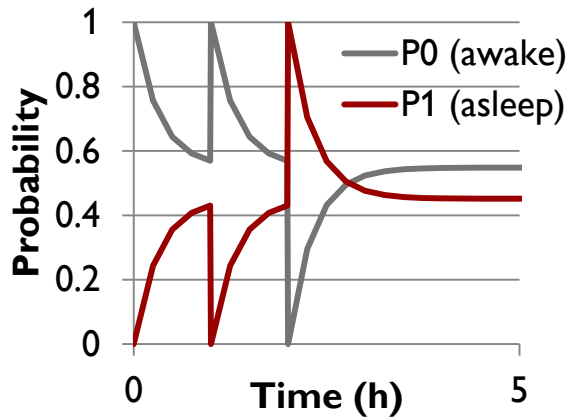
Probabilities following an awake observation at time=0

Additional observations made at
1 h (awake) and 2 h (asleep)

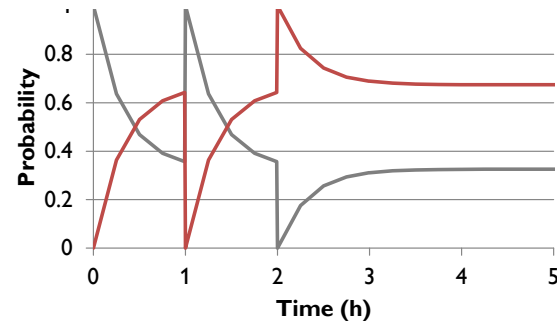
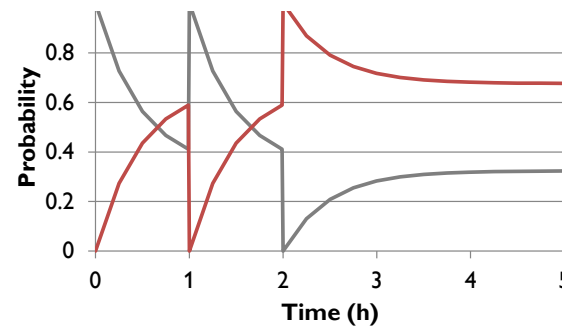
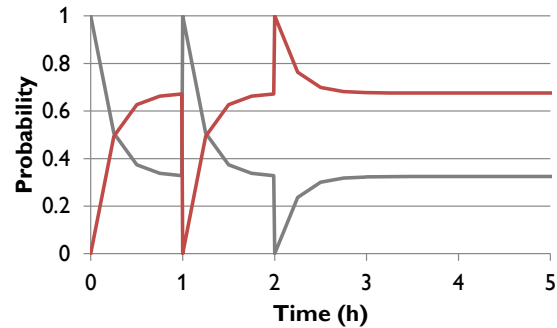


Introducing a treatment effect

Without treatment



With treatment



Promote falling asleep

K01 ↑
 K10 ↔
 K01+K10 ↑

Inhibit waking up

K01 ↔
 K10 ↓
 K01+K10 ↓

Both effects

K01 ↑
 K10 ↓
 K01+K10 ↔



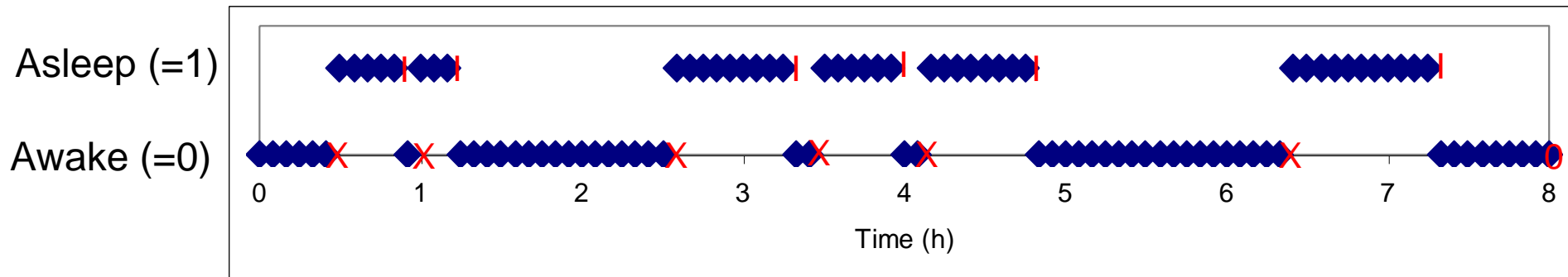
Clues about mechanism in data

Mechanism	Total sleep time	Average time of sleep episodes	Average time of awake episodes	# of arousals	# of times falling asleep	Total # of State transitions
Promote falling asleep	↑	↔	↓	↑	↑	↑
Inhibit waking up	↑	↑	↔	↓	↓	↓
Both effects K10+K01 ↔	↑	↑	↓	↔	↔	↔



Repeated time to event analysis an alternative to Markov model

- Assume transitions as events
- Assume no unobserved transitions
- Separate (constant) hazards for waking up and falling asleep



X – event during awake period (falling asleep) | – event during sleeping period (arousal)
0 – censoring during awake period

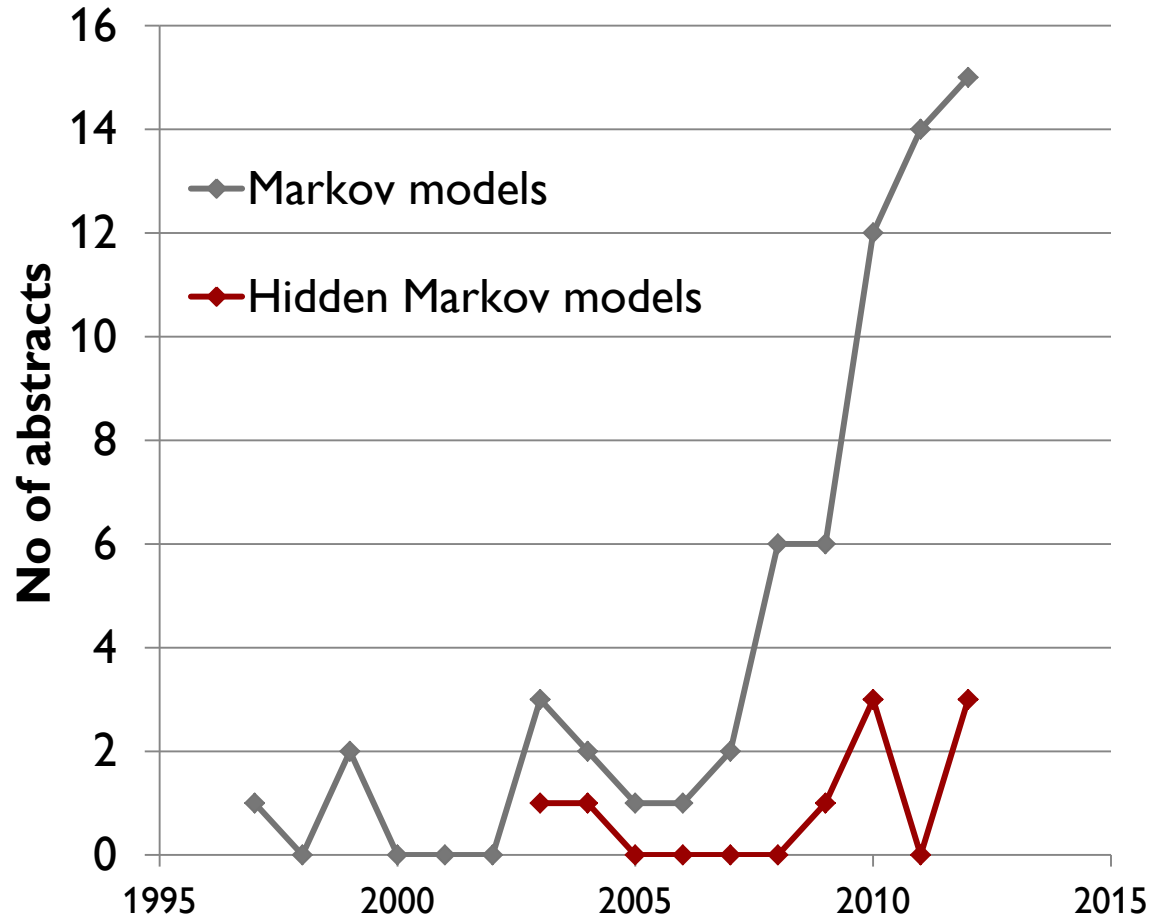


RTTE model – NMTRAN

```
$PROB Repeated Time To Event data - constant hazard
$INPUT ID DV TYPE DVID
$DATA data
$PRED
  IF(TYPE.EQ.0) HAZ= THETA(1) ;hazard for waking up
  IF(TYPE.EQ.1) HAZ= THETA(2) ;hazard for falling asleep
  CUMHAZ  = HAZ*DV           ;cumulative hazard
  SUR     = EXP(-CUMHAZ)    ;survival probability
  IF(DVID.EQ.1) Y = HAZ*SUR ;event
  IF(DVID.EQ.2) Y = SUR     ;censoring
$THETA (0, .146) ; HAZ_WK (/5min)
$THETA (0, .109) ; HAZ_SL (/5min)
$ESTIM LIKE
```



Markov models @ PAGE





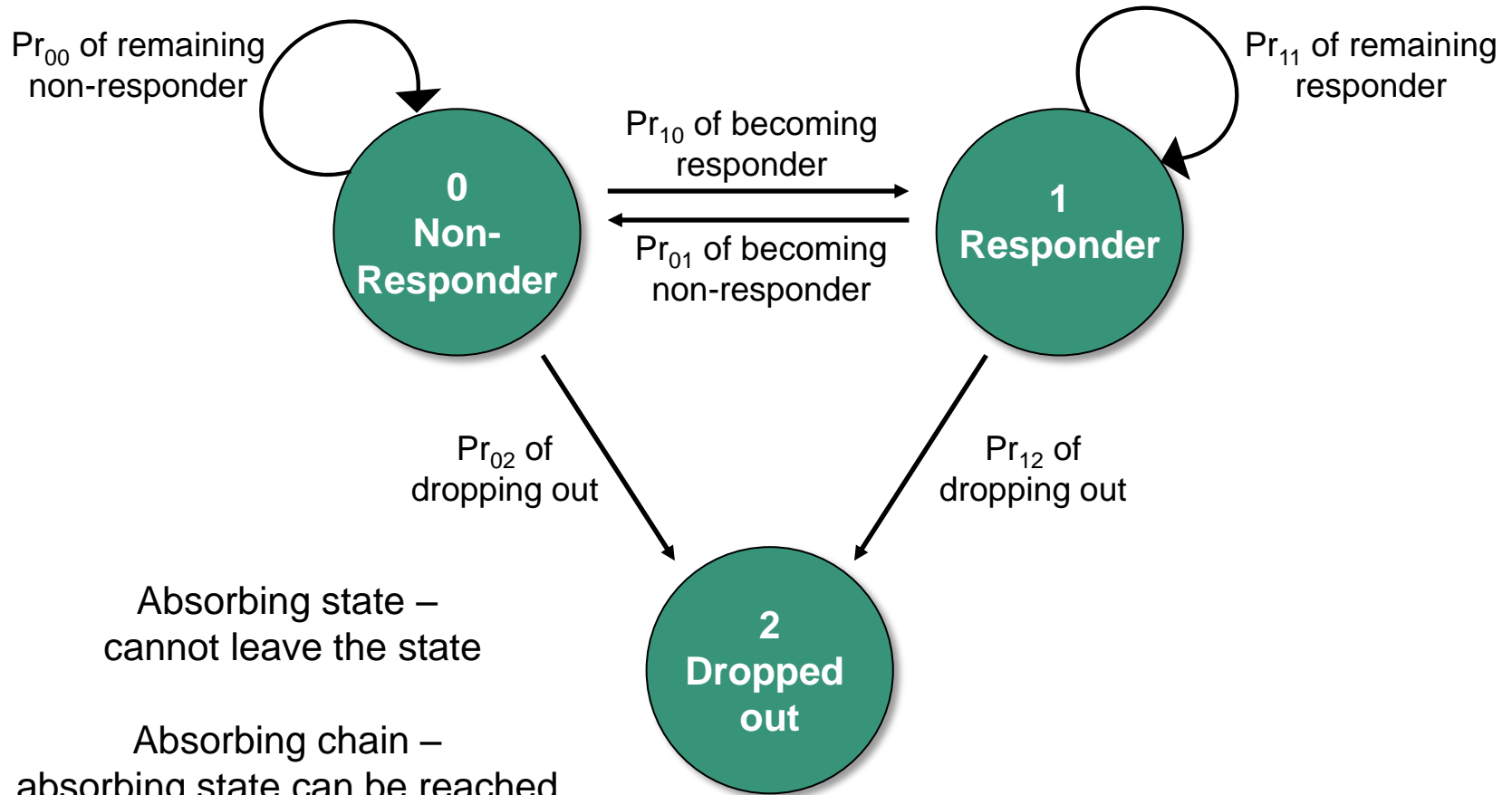
Pharmacometric Markov models

Bergstrand	GI movement	Non-ordered categorical	Compartmental chain	2009	CPT
Bizzotto	Sleep	Non-ordered categorical	1st order + stagetime	2008	PAGE
Girard	Adherence	3-state categorical	Discrete-time	1998	Stat Med
Henin	Hand-and-foot syndrome	Ordered categorical	1st order	2008	CPT
Ito	Dizziness	Ordered categorical	1st order	2008	CPT
Karlsson	Sleep	Non-ordered categorical	1st order + stagetime	2000	CPT
Karlsson	Sedation	Ordered categorical	1st order +stagetime	2002	Measurement and kinetics of in vivo drug effects
Kjellsson	Sleep	Non-ordered categorical	1st order + stagetime	2006	PAGE
Lacroix	ACR20	Binary + dropout	1st order	2009	CPT
Maas	Migraine	Ordered categorical	Hidden Markov	2006	Cephalalgia
Plan	None	Count	1st order	2009	JPKPD
Snoeck	Seizures	Count	1st order	2007	PAGE
Troconiz	Seizures	Count	1st order	2007	PAGE
Zandvliet	Follicules	Multinomial count	Compartmental chain	2008	PAGE
Zingmark	Side-effect	Ordered categorical	1st order	2005	JPKPD



Markov model for responder, non-responder and dropout

Ex, ACR20 score in Rheumatoid Arthritis



Absorbing state –
cannot leave the state

Absorbing chain –
absorbing state can be reached
from all other states



NMTRAN code

Responder, non-responder and dropout

\$PRED

;----transition from being a responder to non-responder---

LGT01=LOG(THETA(1)/(1-THETA(1))) + ETA(1)

P01=EXP(LGT01)/(1+EXP(LGT01))

;----transition from responder to dropout---

LGT21=LOG(THETA(2)/(1-THETA(2)))

P21=EXP(LGT21)/(1+EXP(LGT21))

;-- transitions from being a non-responder to a responder---

LGT10=LOG(THETA(3)/(1-THETA(3))) + ETA(2)

P10=EXP(LGT10)/(1+EXP(LGT10))

;---transition from non-responder to dropout---

LGT20=LOG(THETA(2)/(1-THETA(2)))

P20=EXP(LGT20)/(1+EXP(LGT20))

;----- transition's probabilities -----

IF (PREV.EQ.1.AND.DV.EQ.0) Y=P01

IF (PREV.EQ.1.AND.DV.EQ.2) Y=P21*(1-P01)

IF (PREV.EQ.1.AND.DV.EQ.1) Y=1-P01-P21*(1-P01)

IF (PREV.EQ.0.AND.DV.EQ.1) Y=P10

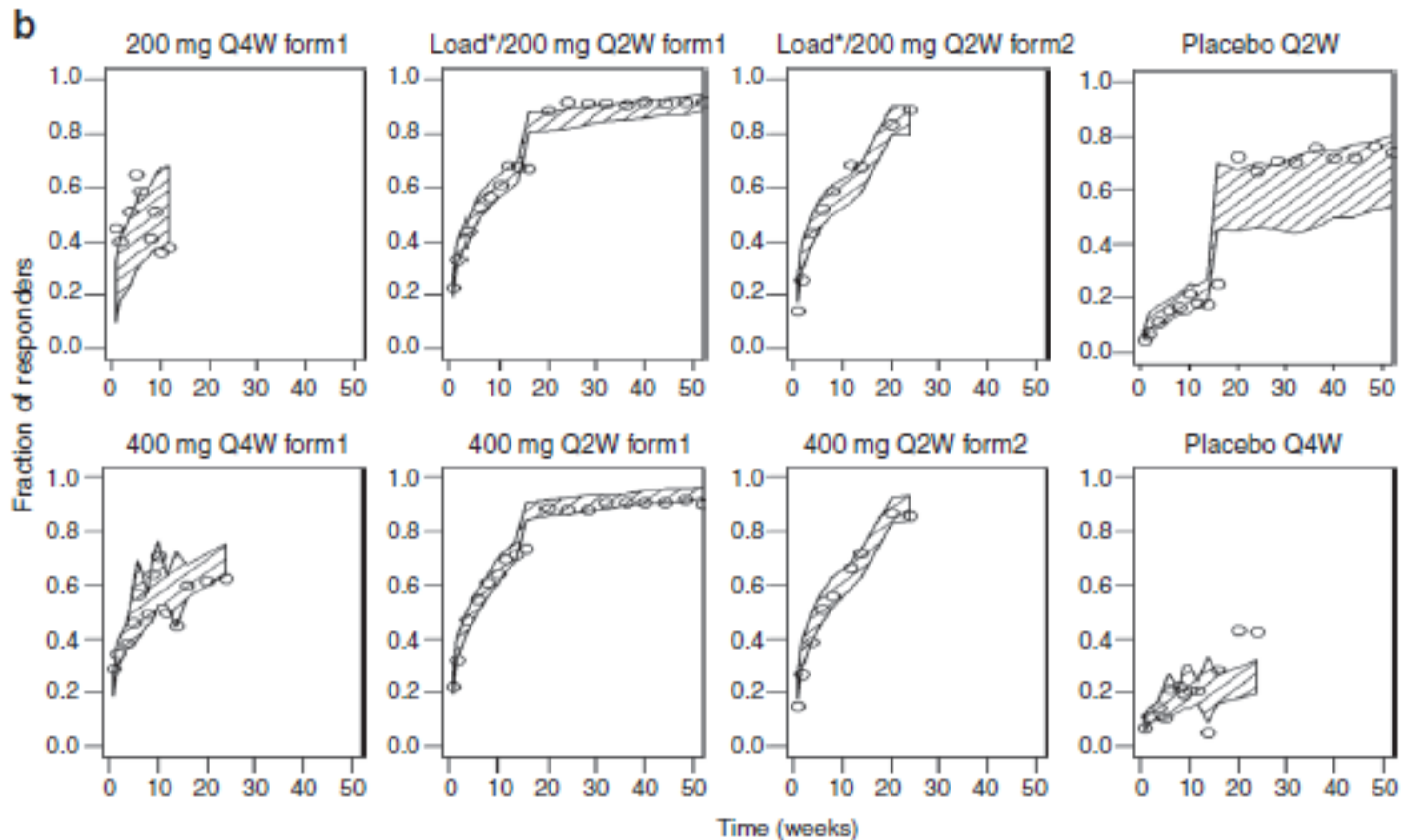
IF (PREV.EQ.0.AND.DV.EQ.2) Y=P20*(1-P10)

IF (PREV.EQ.0.AND.DV.EQ.0) Y=1-P10-P20*(1-P10)

\$EST METH=1 LAPLACE LIKE

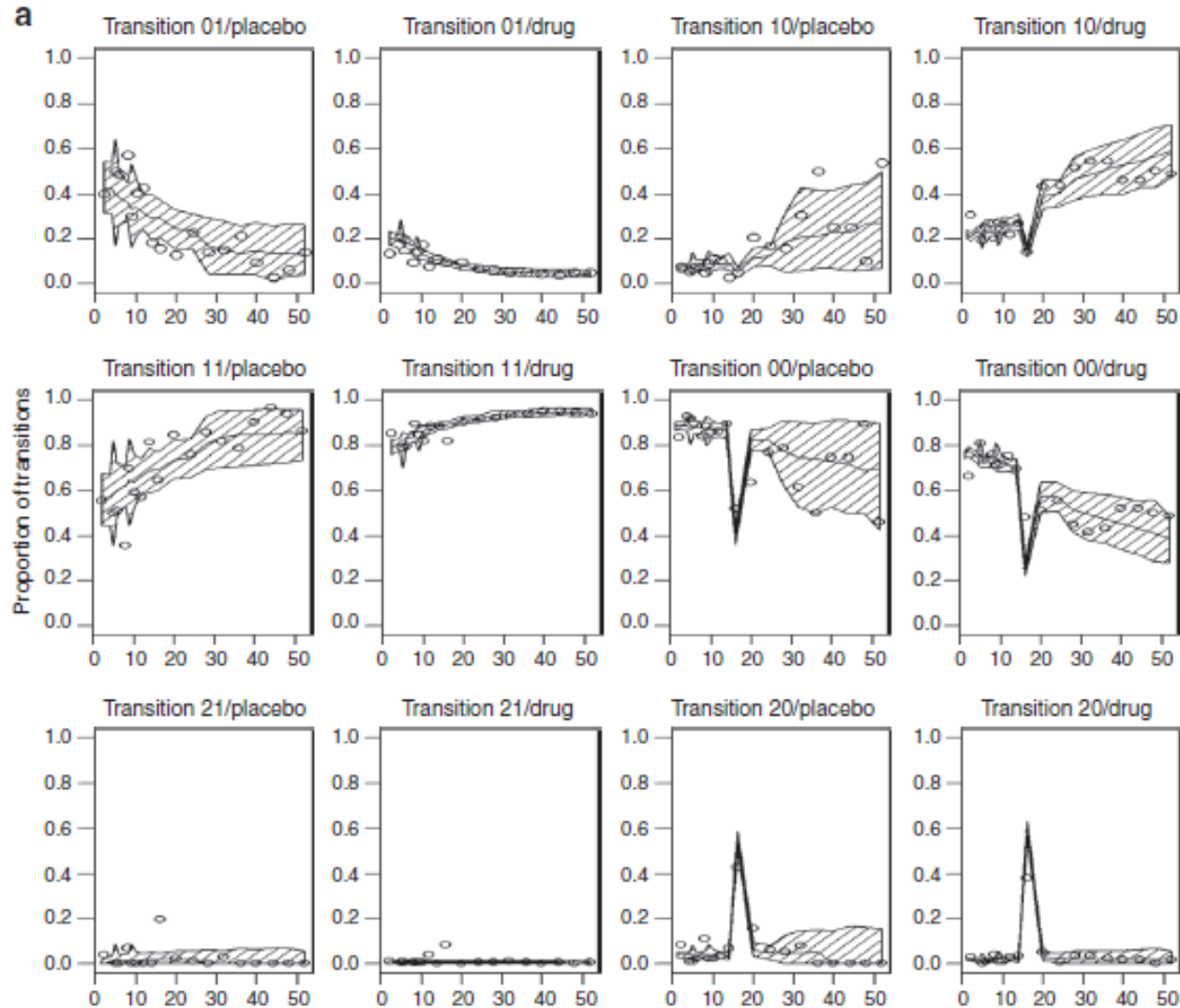


Diagnostics – ACR20 model - proportion responders



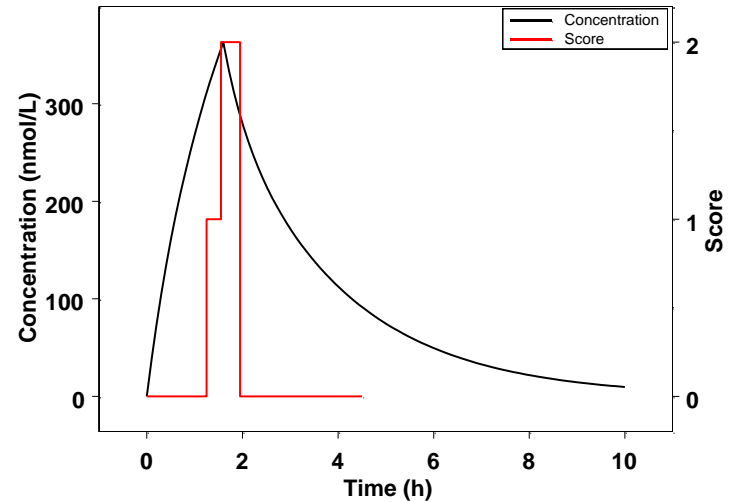
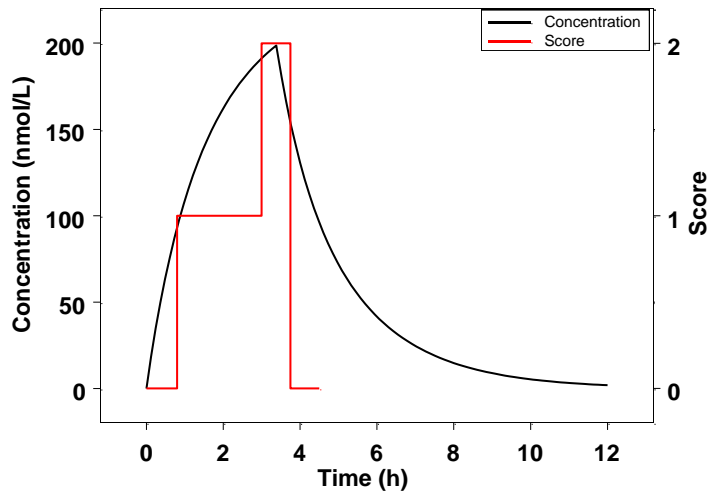
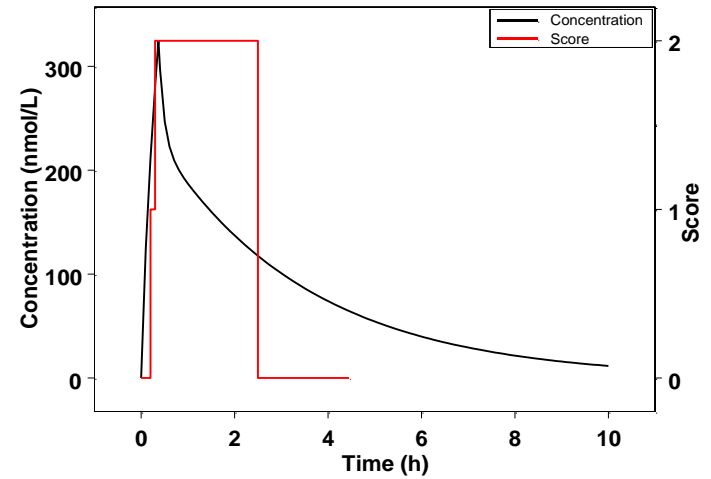
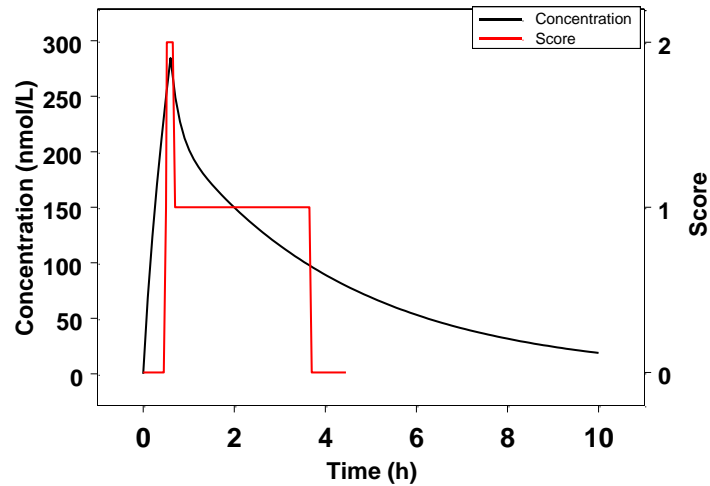


Diagnostics – ACR20 model - transitions

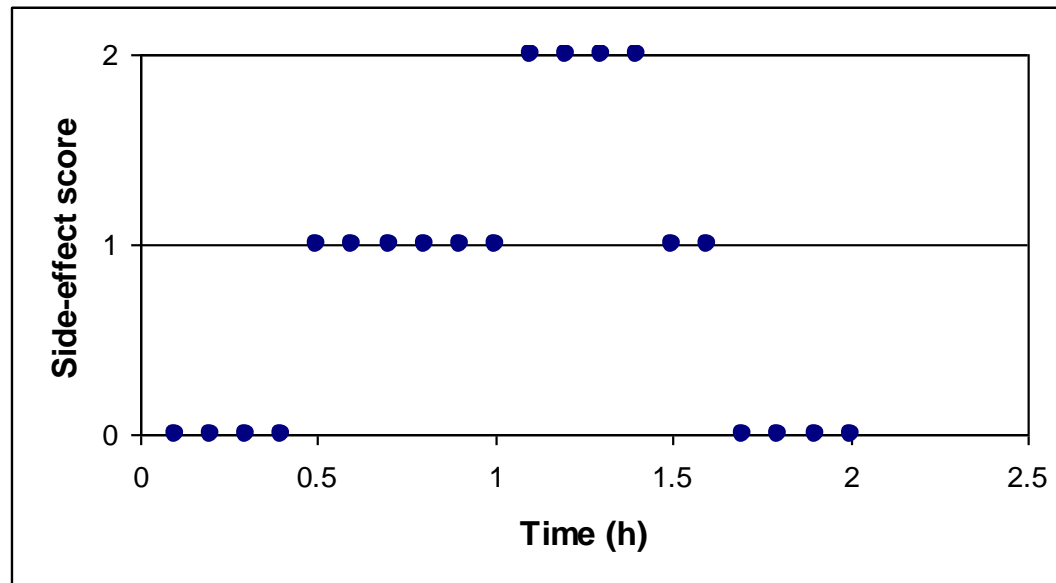
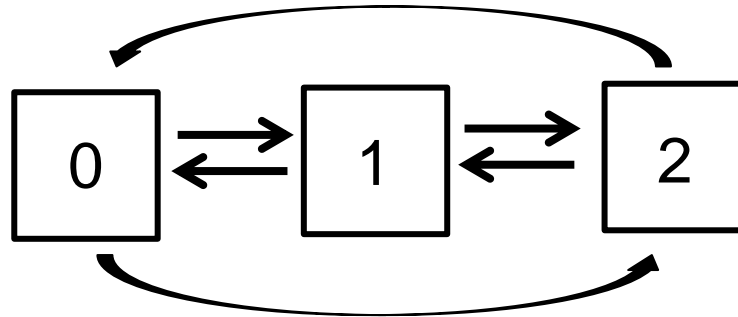




Spontaneous reporting of a side-effect



Alternative assumptions regarding nature of transitions





Parameterization of the model

$$\begin{aligned} [l_{S \geq 1} | pre = 0] &= b_1 + D \\ [l_{S=2} | pre = 0] &= b_1 + b_2 + D \\ [l_{S \geq 1} | pre = 1] &= b_3 + D \\ [l_{S=2} | pre = 1] &= b_3 + b_4 + D \\ [l_{S \geq 1} | pre = 2] &= b_5 + D \\ [l_{S=2} | pre = 2] &= b_5 + b_6 + D \end{aligned}$$

$$PC_x = \frac{e^{1x}}{1 + e^{1x}}$$

$$p_{S=0} = 1 - PC_{S \geq 1}$$

$$p_{S=1} = PC_{S \geq 1} - PC_{S=2}$$

$$p_{S=2} = PC_{S=2}$$

$$D = \frac{E_{\max} |_{pre=0,1,2} \cdot e^{\eta_i} \cdot Ce}{EC_{50} \cdot (1 + C_{tol} / TC_{50}) + Ce}$$



Spontaneous reporting of a side-effect - model simulations

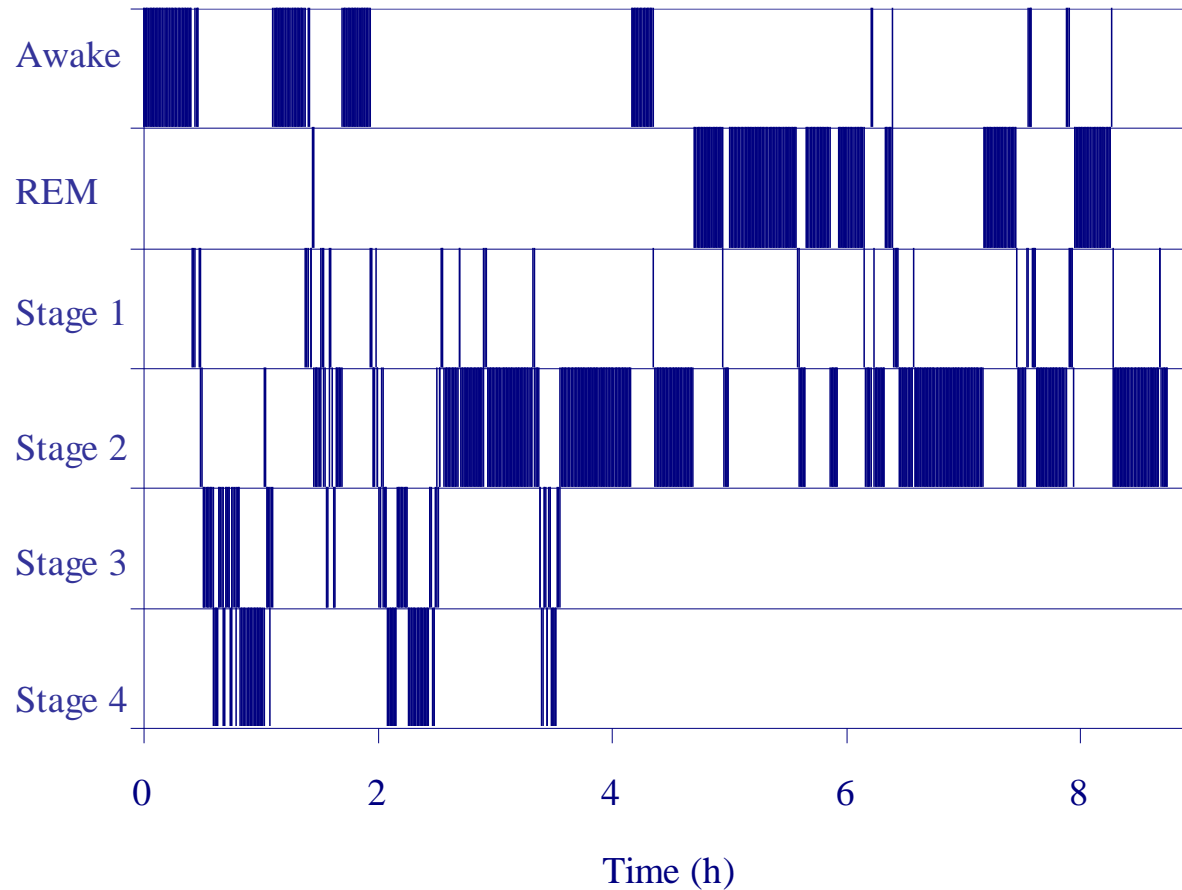
Number of transitions	Proportional odds model			Markov model			Observed data
	1 min	3 min	6 min	1 min	3 min	6 min	
0-2	289 (249; 335)	117 (102; 133)	72 (62; 82)	11 (9; 14)	11 (8; 15)	11 (8; 15)	11
1-2	178 (133; 223)	79 (64; 95)	48 (39; 59)	25 (21; 29)	25 (20; 29)	25 (21; 29)	23

The results shown are the average and (10th and 90th percentiles) from 100 simulated datasets

The performance of the proportional odds model, but not the Markov model, is dependent on the choice of observation frequency



Sleep scoring – non-ordered categorical data



(Karlsson et al., CPT 2000)

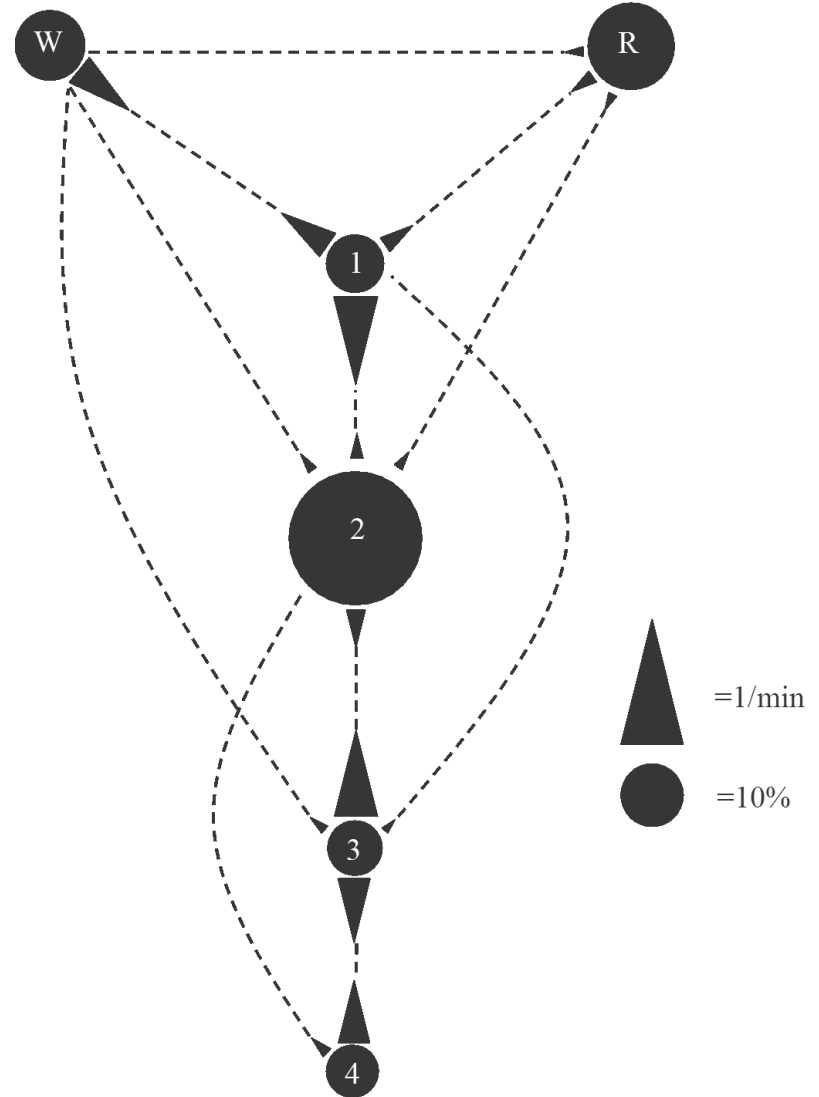


Choice of transitions to model

- To reduce the number of transitions to model, three criteria were defined to identify the transitions of interest representing:
 - (i) >1% of all observations in a stage,
 - (ii) >10% of all transitions from a stage
 - (iii) >10% of all transitions to a stage
- A transition was modeled, if at least one of these criteria was fulfilled

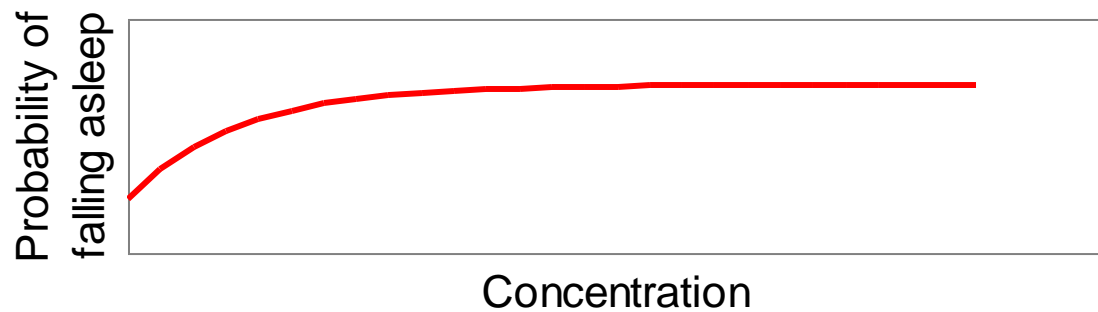
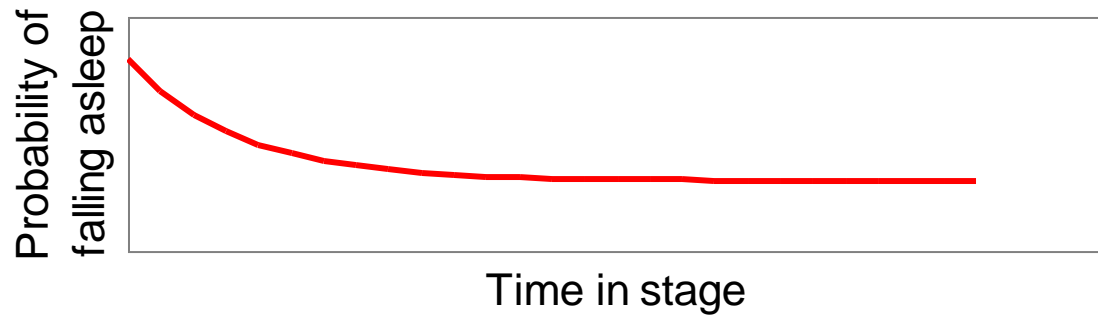
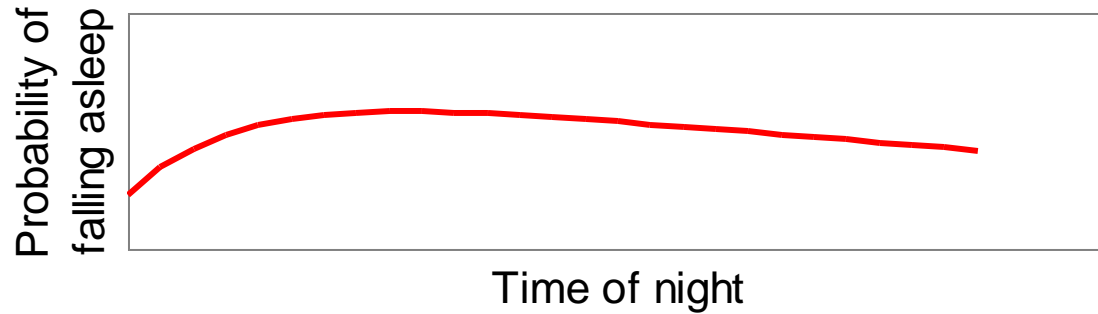


Overall sleep pattern





Transition dependences





What previous info to condition on?

- First-order Markov models often sufficient
- Sometimes biological processes dictate use of higher-order Markov elements
 - 2nd (3rd etc) – order elements
 - Time since entering present stage (stagetime)
 - Prior stage(s)

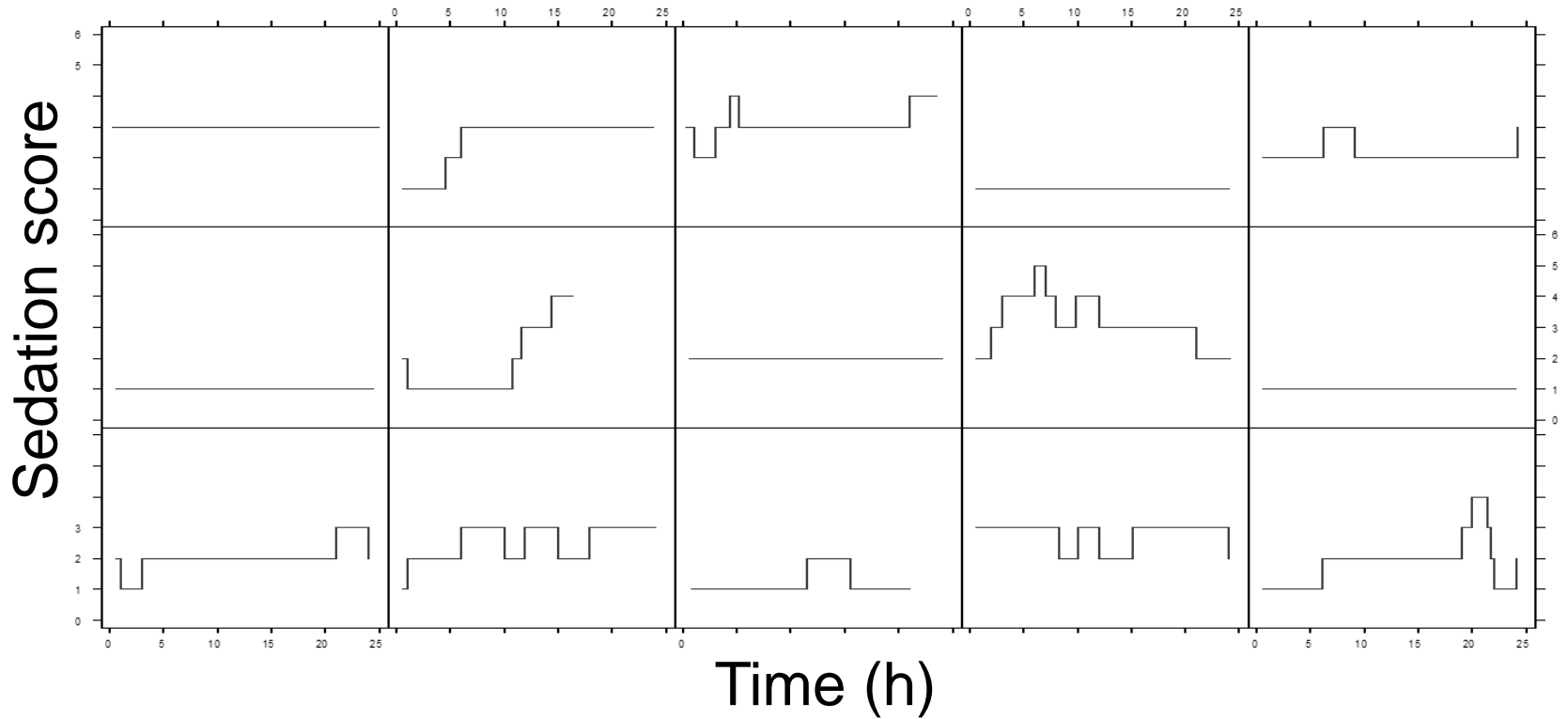


How to start a Markov chain?

- Sometimes obvious –
 - Sleep-wake data that starts at bed-time
 - Responder status for ACR20
- Sometimes screening data provide information about initial state
- Transform first observation(s) to a covariate for starting the chain

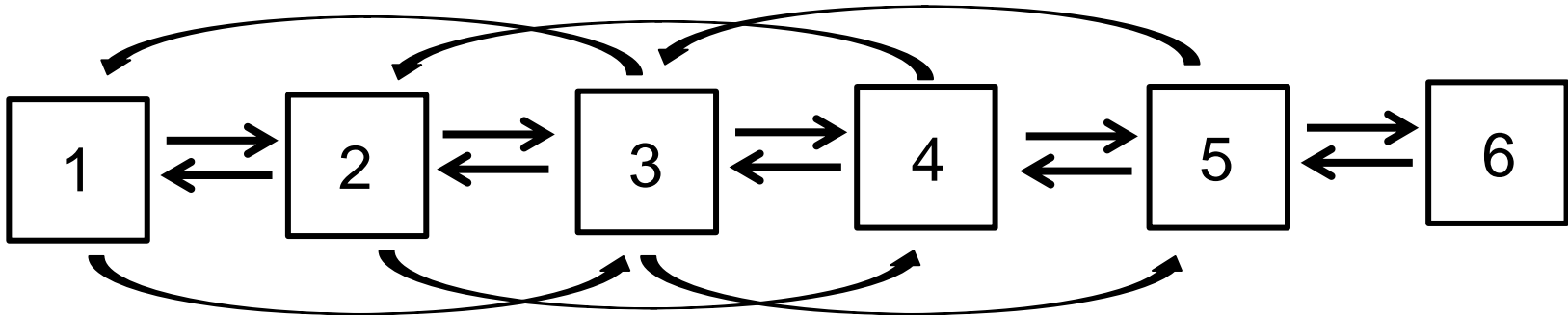


Sedation scores following stroke



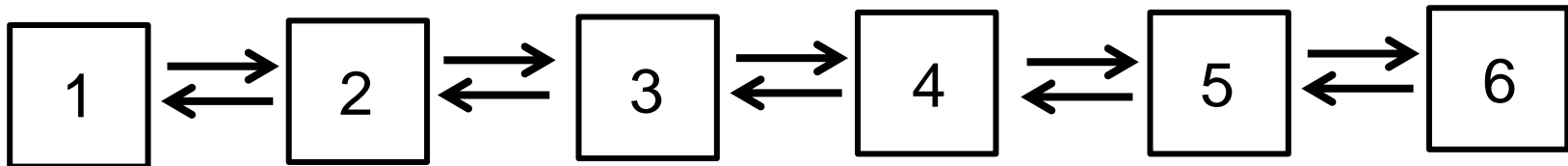


Alternative assumptions regarding nature of transitions



Many transitions to model

No assumption about intermediate transition states



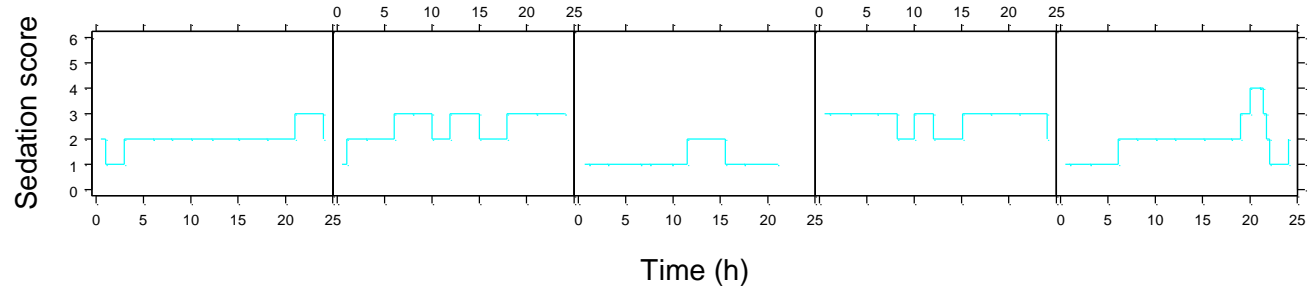
Fewer transitions to model

Assumption about intermediate transition states

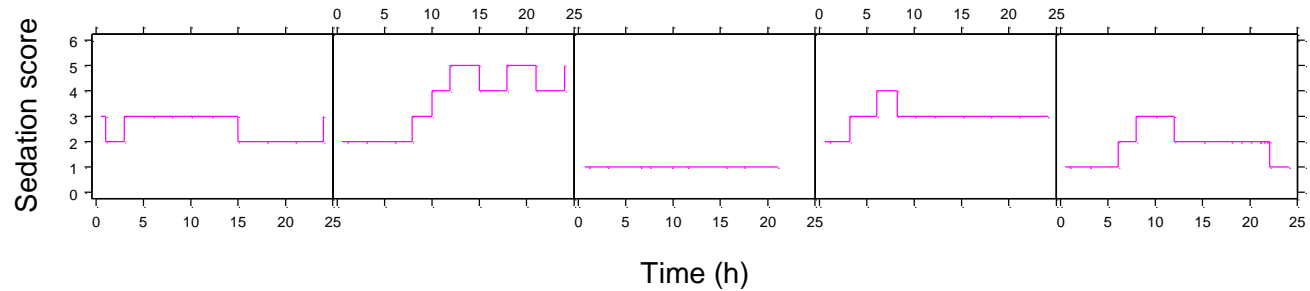


Individual time courses of sedation

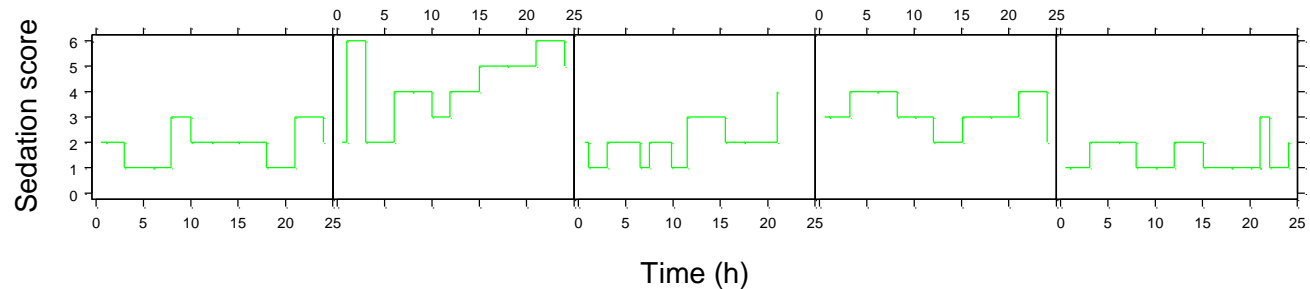
Observations



Markov model
simulations

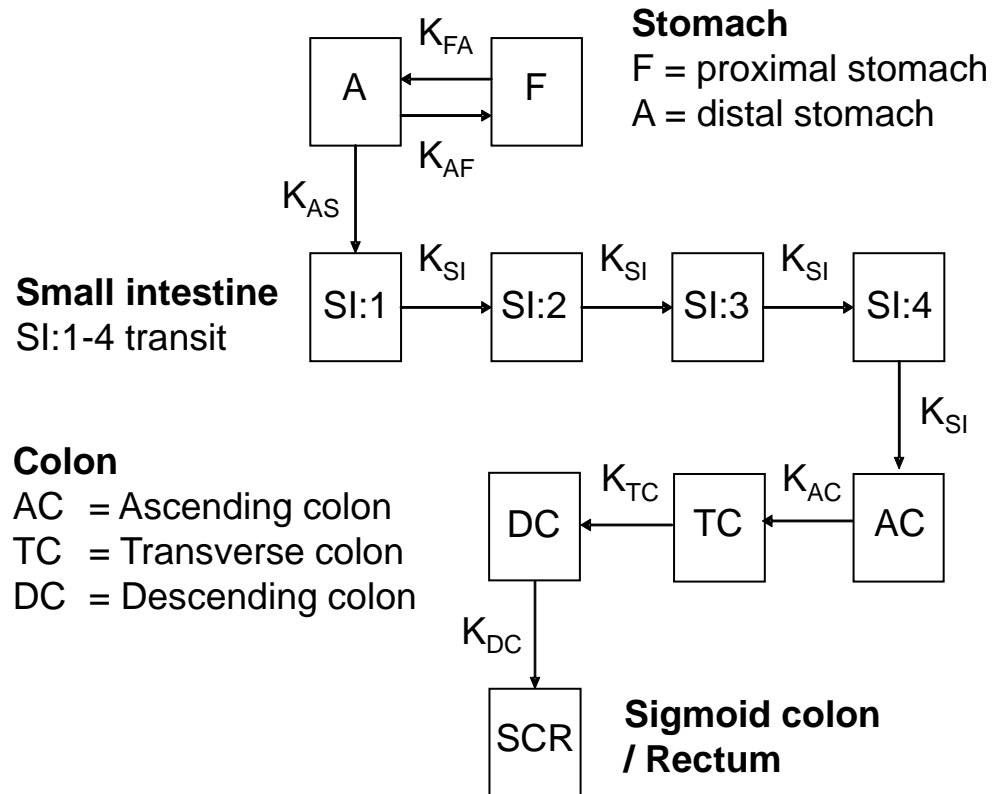


Ordered categorical
model simulations





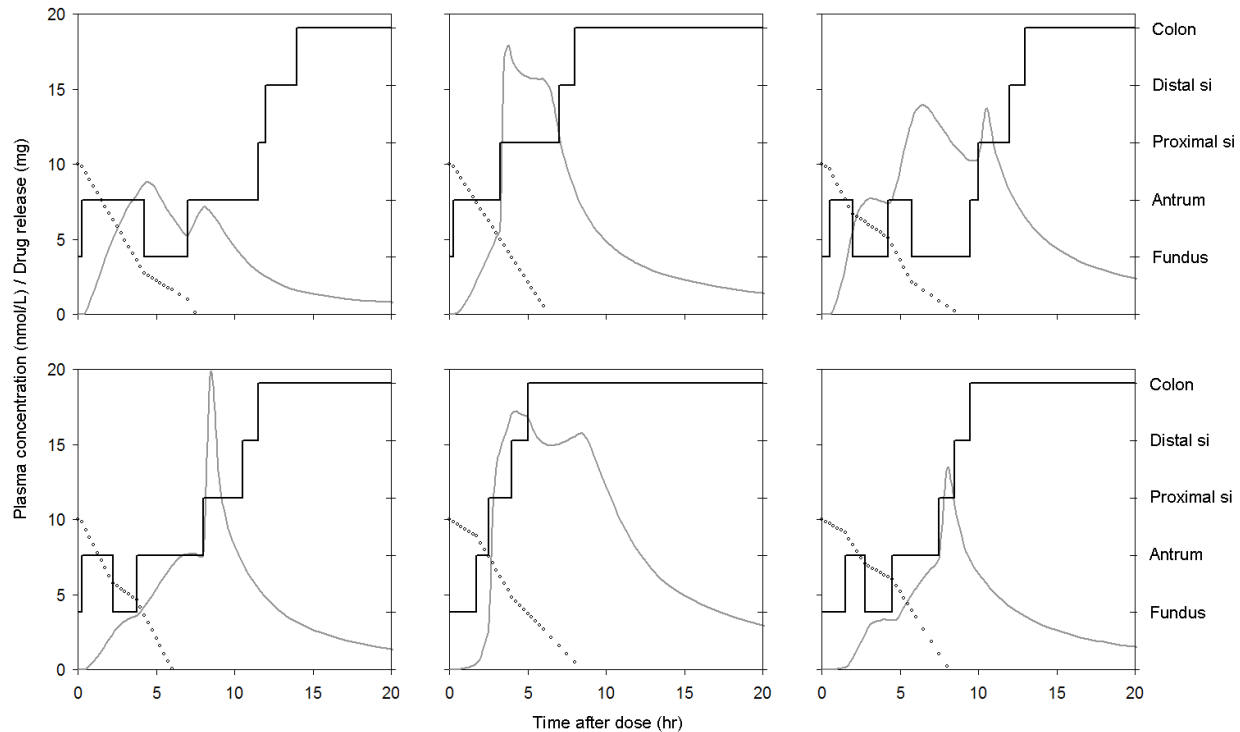
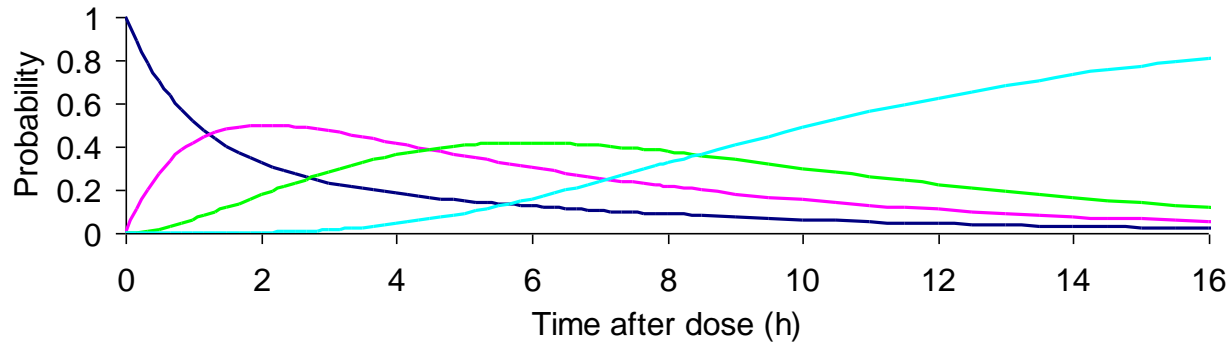
Model for GI transit



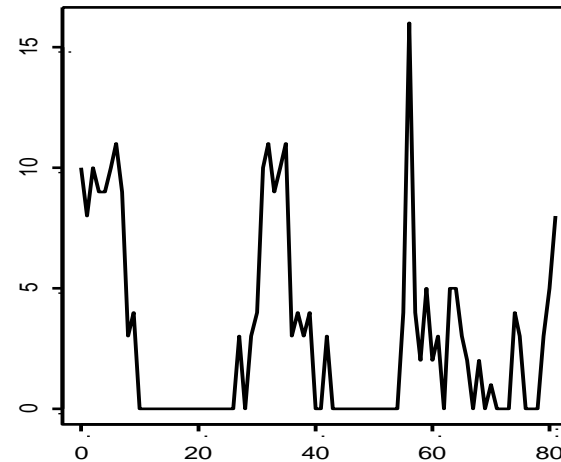
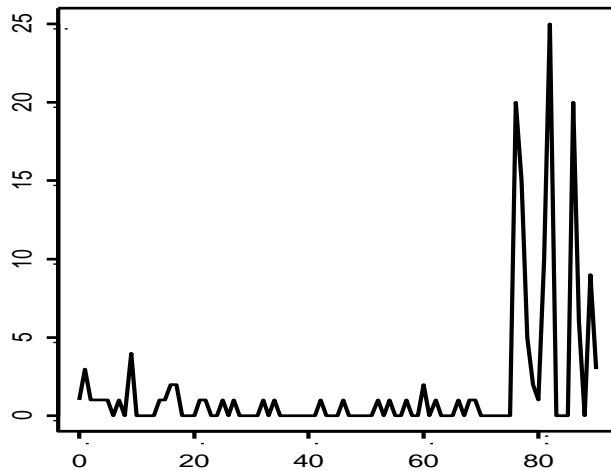


GI position

- Proximal stomach
- Distal stomach
- Small intestine
- Colon



Count data - daily seizure scores in epileptic patients



Markov model; Troconiz et al., JPKPD 36:461-77 (2009)

Hidden Markov model; Delattre et al., JPKPD 39:263-71 (2012)

Consequences of ignoring Markov properties - estimation

Information content in data overestimated

SEs underestimated

Hypothesis tests inappropriate

Interindividual variability overestimated

Potential structural model misspecification

No info on time-course of dependence

Consequences of ignoring Markov properties – simulation & design

Duration of state periods too low

Inflated number of transitions

Inflated number of extreme value occurrences

E.g. distribution of maximum severity score in population

Individualisation strategy suboptimal

Value of more frequent observations overrated

Positioning of observation times

Optimal design results inappropriate*



Markov model or not?

Start with Markov

Frequent observations

Many consecutive same-state observations

Many levels of response

Non-ordered categorical data

Start without Markov and diagnose

Check number of transitions

Check average duration of same-state periods



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Thank you!