



UPPSALA
UNIVERSITET

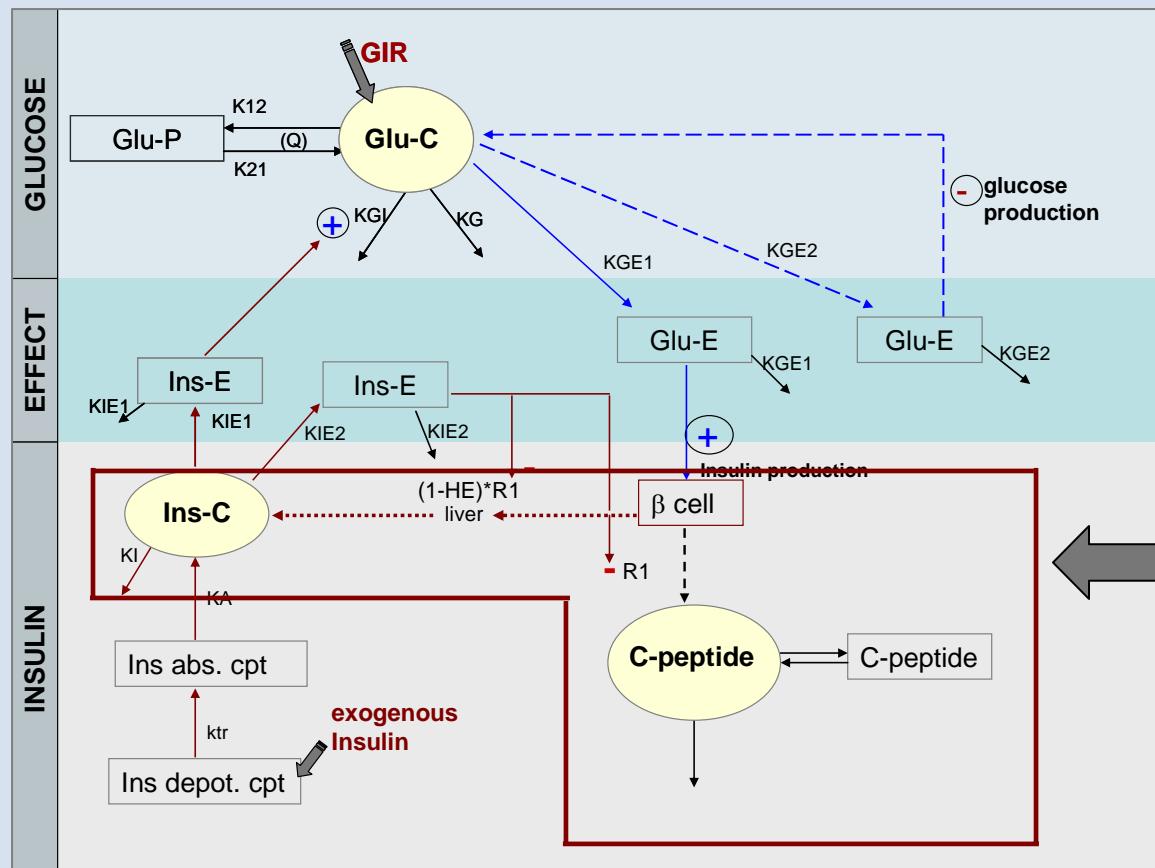
Insulin Secretion and Hepatic Extraction during Euglycemic Clamp Study: Modelling of Insulin and C-peptide data

Chantaratsamon Dansirikul
Mats O Karlsson

Division of Pharmacokinetics and Drug Therapy
Department of Pharmaceutical Biosciences
Uppsala University

Goal & Aim

Goal: "To develop an integrated glucose-insulin model"



Aim: "to characterize endogenous insulin secretion profile and to estimate hepatic extraction of insulin"

Introduction

Insulin Secretion & C-peptide

Insulin Secretion

- Secretion: basal & prandial
- Direct assessment of insulin secretion from blood concentration is not so easy!
- ~50% (40-85%)¹⁻³ is lost during single pass through liver

C-peptide

- Co-secreted with insulin from beta-cell on an equimolar basis
- A negligible amount is lost during single pass through the liver⁴
- A measure of pre-hepatic insulin secretion

1. Cobelli C, et al. *Diabetes* **37**: 223-31 (1988)
2. Tura A, et al. *Am J Physiol Endocrinol Metab* **281**: E966-74 (2001)
3. Meier JJ, et al. *Diabetes* **54**: 1649-56 (2005)
4. Polonsky K, et al. *J Clin Invest* **72**:1114-23 (1983)

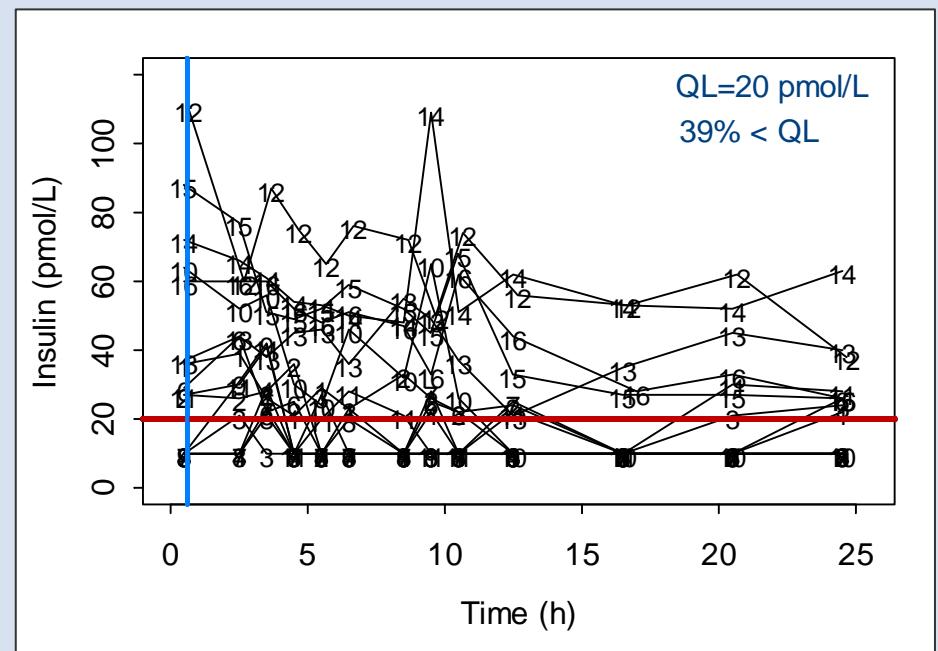
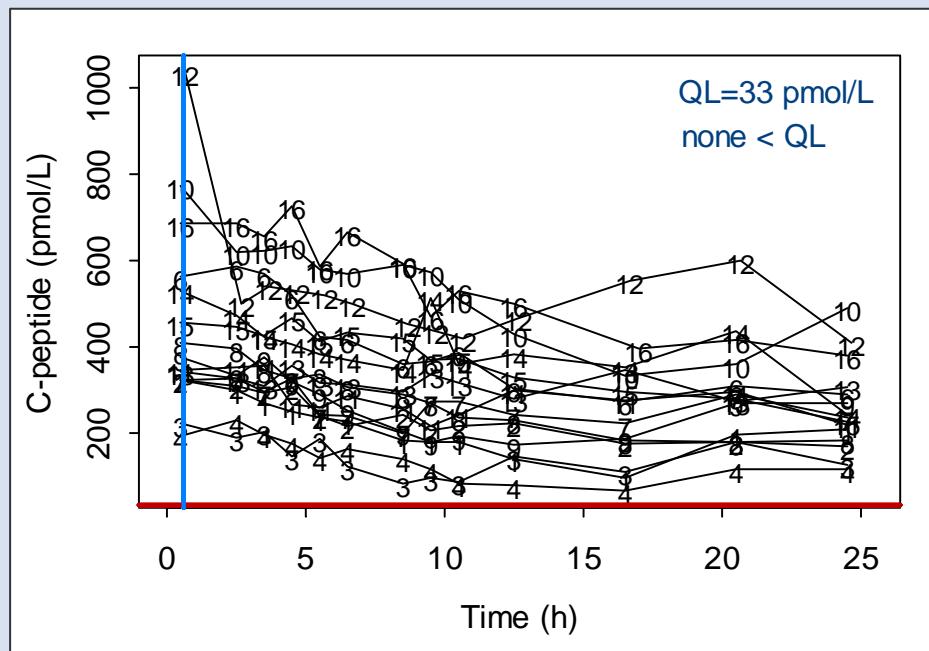


Data

24-hour Euglycemic clamp study

C-peptide & Insulin Data

- 15 healthy volunteers
- 13 samples/subject



Methods

Modelling

- NONMEM VI
- Log-transformed data
- Estimation method
 - First Order Conditional Estimation (FOCE) method
 - Laplacian method (handling of data below quantification limit (BQL))



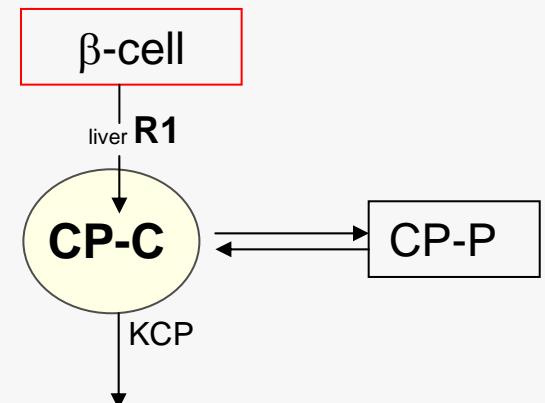
Methods

Model & Modelling Steps

1

C-peptide data:

- Population disposition parameters were fixed¹
- ✓ Insulin pre-hepatic secretion (R1) was characterised using a flexible staircase input model²

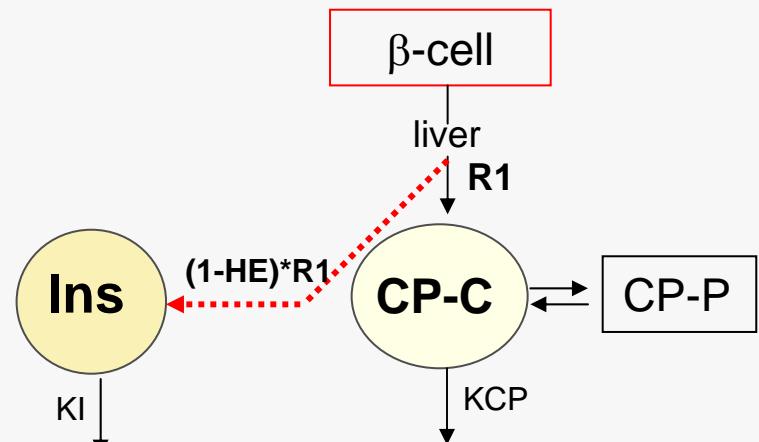


2

Sequential approach

C-peptide & Insulin data:

- Population disposition parameters of insulin were fixed³
- Pre-hepatic secretion (R1) estimated from C-peptide data was used
- ✓ Hepatic extraction (HE) of insulin



¹ Van Cauter E, et al. *Diabetes* **41**: 368-77 (1992); ² Lindberg-freij A et al. *Biopharmaceutics & Drug Disposition* **15**: 75-86 (1994);

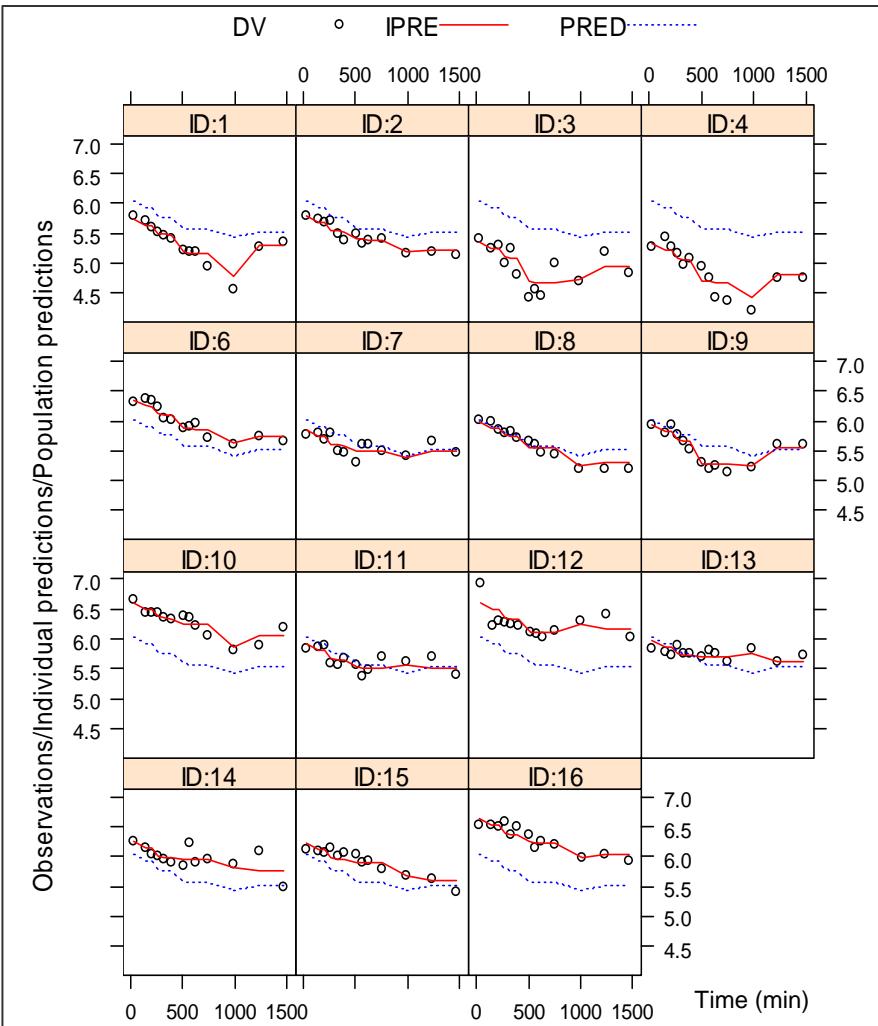
³ Silber H, et al. *J Clin Pharmacol* (in press)



Results

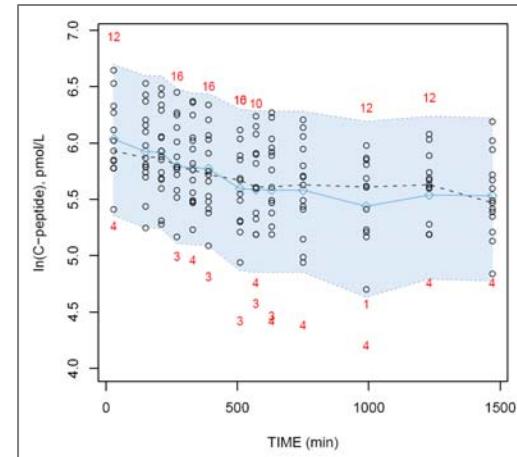
C-peptide secretion (Pre-hepatic secretion of Insulin)

✓ describe data



✓ simulate new data

1. Visual Predictive Check e.g. 90% PI



2. Numerical Predictive Check e.g.

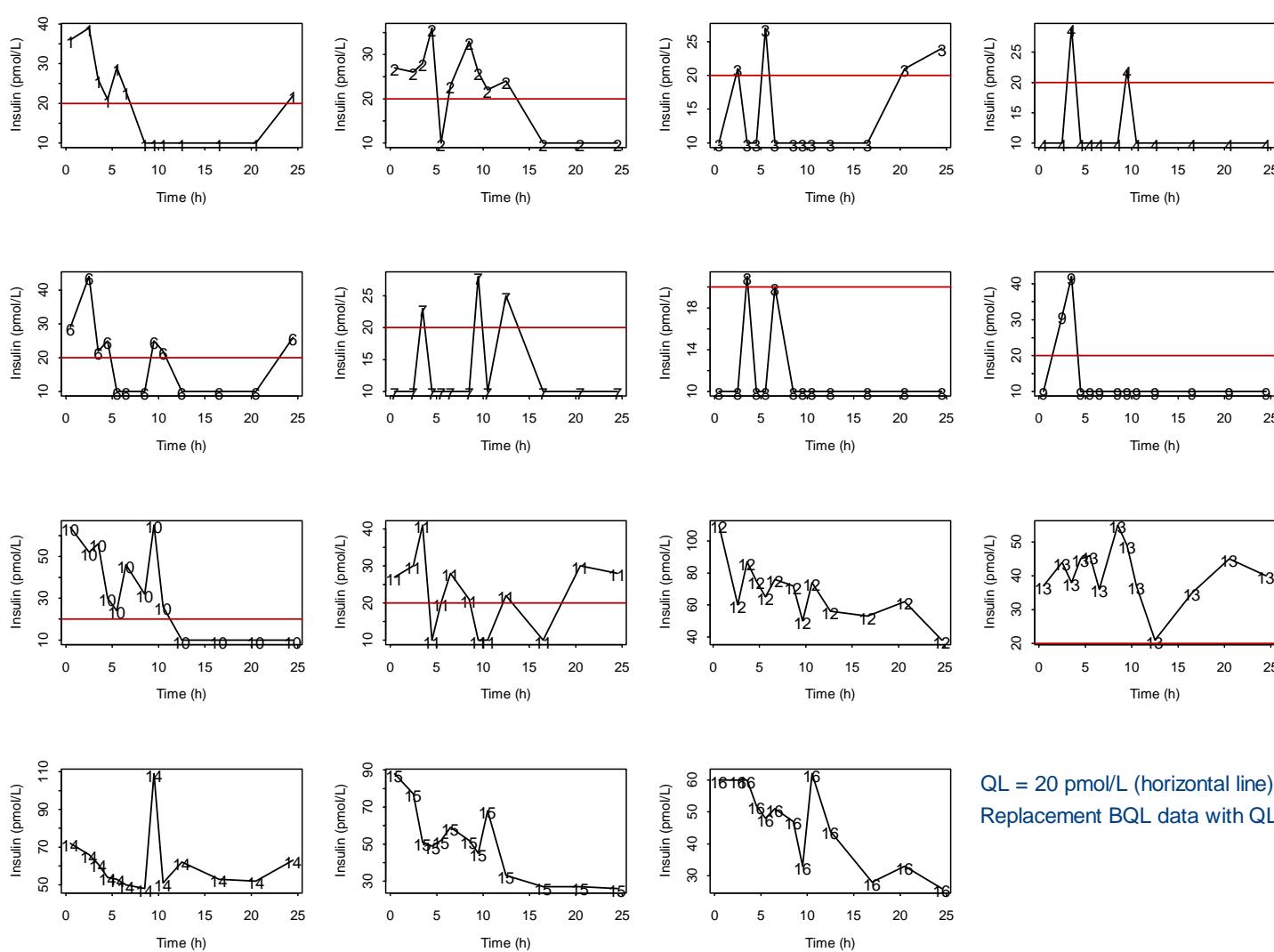
	%Observation (95%CI) ¹	Expectation
above simulated median	49.7 (28.2-71.8)	50
below simulated median	50.3 (28.2-71.8)	50
above simulated 90%PI	4.1 (0-16.9)	5
below simulated 90%PI	7.2 (0-15.4)	5
above simulated 50%PI	29.2 (8.21-44.6)	25
below simulated 50%PI	23.1 (8.21-45.1)	25

¹ obtained based on 1000 simulated datasets



Issue

39% of Insulin data are BQL



Method

How to handle BQL?

- Omission BQL data
- Replacement observation
 - QL/2 for all BQL data
 - QL/2 for the first BQL data, discard the rest
 - Replacing with 'zero'
- Likelihood approach: maximizing likelihood ...
 - For '**all data**' taken into account likelihood of a BQL observation being below QL
 - e.g. F_FLAG functionality in NONMEM VI
 - For '**data above QL**' conditional on the likelihood of being above QL
 - e.g. YLO functionality in NONMEM VI

Results

Hepatic Extraction (HE) of Insulin

- Slightly different parameter estimates

Method	HE		P-RV	
	Typical	95%CI	Typical	95%CI
1. Omission	0.474	(0.393-0.542)	0.236	(0.205-0.269)
2. QL/2	0.618	(0.50-0.699)	0.373	(0.313-0.430)
3. F_FLAG	0.561	(0.458-0.646)	0.275	(0.232-0.334)

HE= hepatic extraction, P-RV = proportional residual variability (on untransformed scale), 95%CI = bootstrap 95% confidence interval (1000 bootstrap samples)

- Neither the inclusion of additive residual error (fixed or estimated) nor ETA on EPS improved OFV value for each method

QL/2 or F_FLAG, which method should I go for?

Criteria	Relevance for comparison	Results	
		QL/2	F_FLAG
• Objective function value (OFV)	no	inapplicable	inapplicable
• Goodness of fit plots (GOF)	limited		
• Model estimation stability <ul style="list-style-type: none"> • Different initial estimates • Bootstrapping 	yes	stable 74% Cov step success	stable 72% Cov step success
• Bias & Imprecision (Simulation & re-estimation ¹)	yes	ME < 1% RMSE < 10%	ME < 1% RMSE < 10%
• Numerical Predictive Check	yes	✓	✓ (slightly better)
• Run time	yes	< 10 min	< 10 min

¹ 100 datasets was simulated using final estimates from each method. Parameters were then re-estimated. Mean error (ME) and root mean squared error (RMSE) were computed and presented as a percentage of the true value.



Results

Numerical Predictive Check

- % observation above 90% and 50% prediction interval (PI)

% observations above expectation		QL/2		F_FLAG	
		observed	95%CI ¹	observed	95%CI ¹
90%PI	5	12.3	0.51 - 11.8	10.8	0.513 - 13.3
50%PI	25	34.9	12.3 - 40.0	31.3	11.3 - 42.1

¹ The 95% confidence interval (CI) is obtained based on 1000 simulated datasets

- % BQL observation simulated from each method

method	%BQL observation	
	observed	95%CI ¹
QL/2	39	27 - 56
F_FLAG	39	15 - 46

¹ The 95% confidence interval (CI) is obtained based on 1000 simulated datasets

Conclusions

- Endogenous insulin secretion profile was well characterized using a flexible staircase input model
- In the presence of BQL data, F_FLAG method was chosen
 - Theoretically appealing
 - Slightly better with numerical predictive check

Discussion Issue: I

Selection among different methods of handling BQL

- **Original Question**

"How to select which method to be used to handle BQL when the OFV (and GOF) can not be used?"

- **Approach**

- Model estimation stability
- Bias & imprecision (simulation & re-estimation)
- Numerical Predictive Check
- Run time

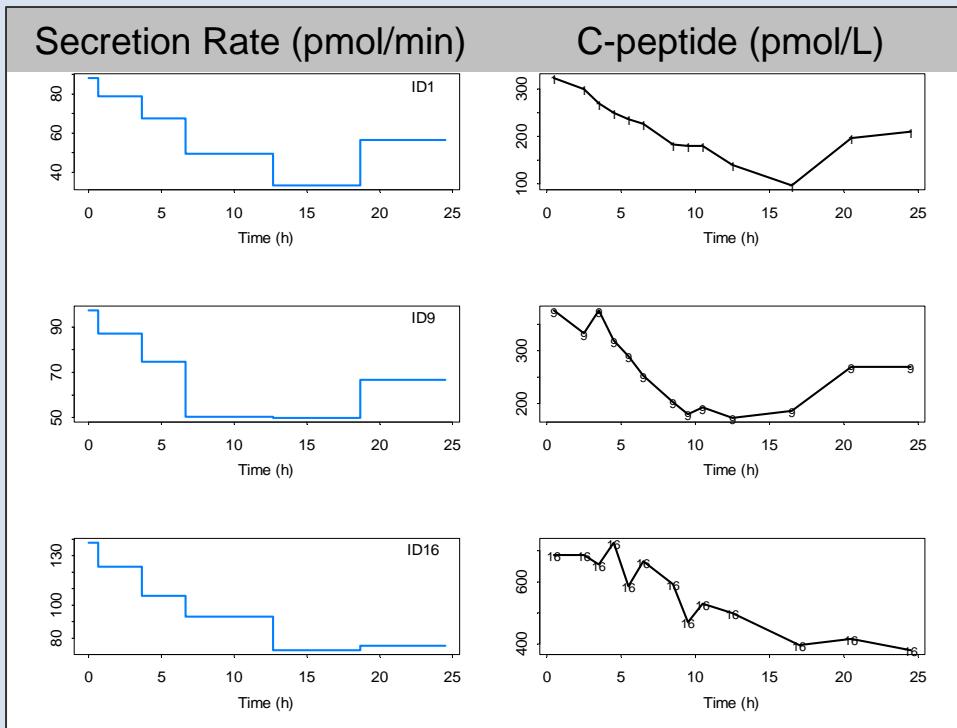
What else can be considered?

Discussion Issue: II

Flexible staircase input model & Endogenous secretion

Flexible Staircase Input Model

- Secretion rate is the same within a time interval
- Changes are characterised using staircase function
- Secretion at baseline and fraction of changes are estimated



Any comments?