



# **IGRH model: a mechanistic model Integrating the relationship between average Glucose levels ( $C_{g,avg}$ ), RBCs & HbA1c in a mixed population of healthy volunteers and diabetic subjects**

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# Background

## HbA1c ?

- Chronic glycemia biomarker (2-3 months)

Standard biomarker for:

- Diagnosis (ADA 2010)
- Adequacy of glycemic management

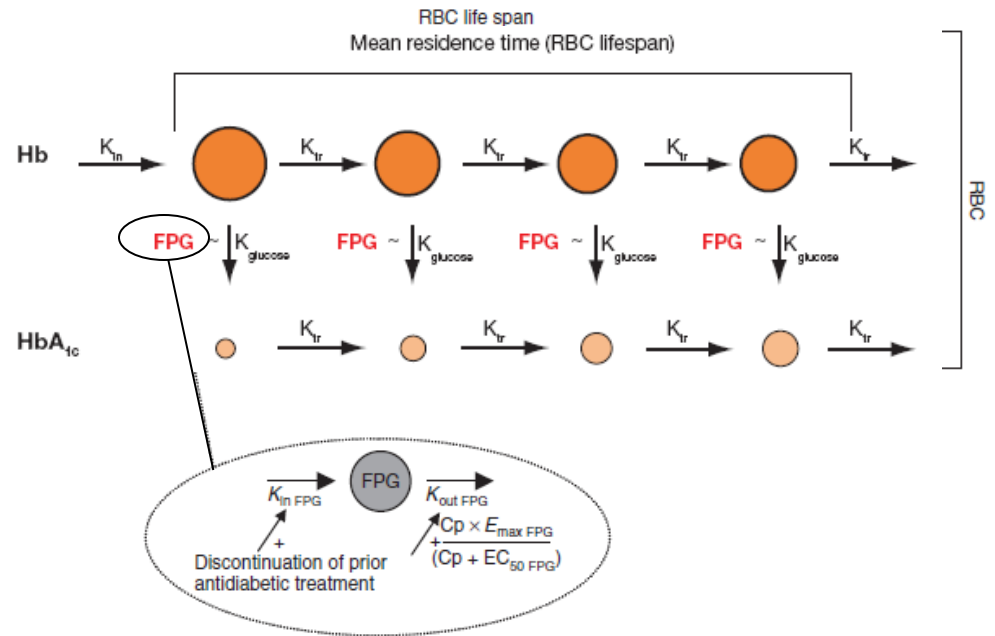




# Background

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## FPG – HbA1c Model Hamrén et al. 2008



- Mechanistic PKPD model:

RBC ageing and glycosilation to HbA1c

- Gaps:

FPG ~~X~~ chronic glycemia → to relate to HbA1c

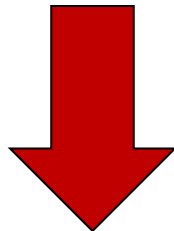
HbA1c depends on RBC LS → the LS model from a small group of Tesaglitazar therapy



# Rationale - Why?

- We wanted to explore more and fill up these gaps!
- To use average glucose concentration ( $C_{g,avg}$ )  
better descriptor of chronic glycemia
- Better RBC LS description
- Empirical models exist that relate:

$$C_{g,avg} \sim HbA1c$$



Lacking mechanism-based model



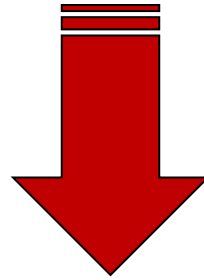
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# Aims

To derive a dynamic mechanism-based model for describing the underlying relationship between  $C_{g,avg}$ - HbA1c **using information from literature**. Including sources of variability (i.e: IIV RBC life-span,...)



How to build the model when  
you have different sources of  
data?



Integrating the data  
formal analysis Nonmem

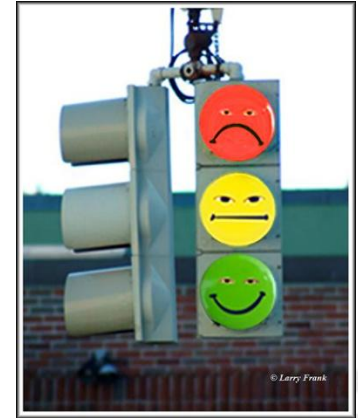


# Overview – Integrating different sources

Digitized data Nathan et al. 2008 :  $C_{g,avg} \sim HbA1c$  at steady state

Mechanistic re-inforcement by **literature priors** in structural & variability components (i.e. LS, IIV-LS, KG,...)

**Digitized data & clinical data as external validation :** hypothesis testing of specific mechanisms with high impact



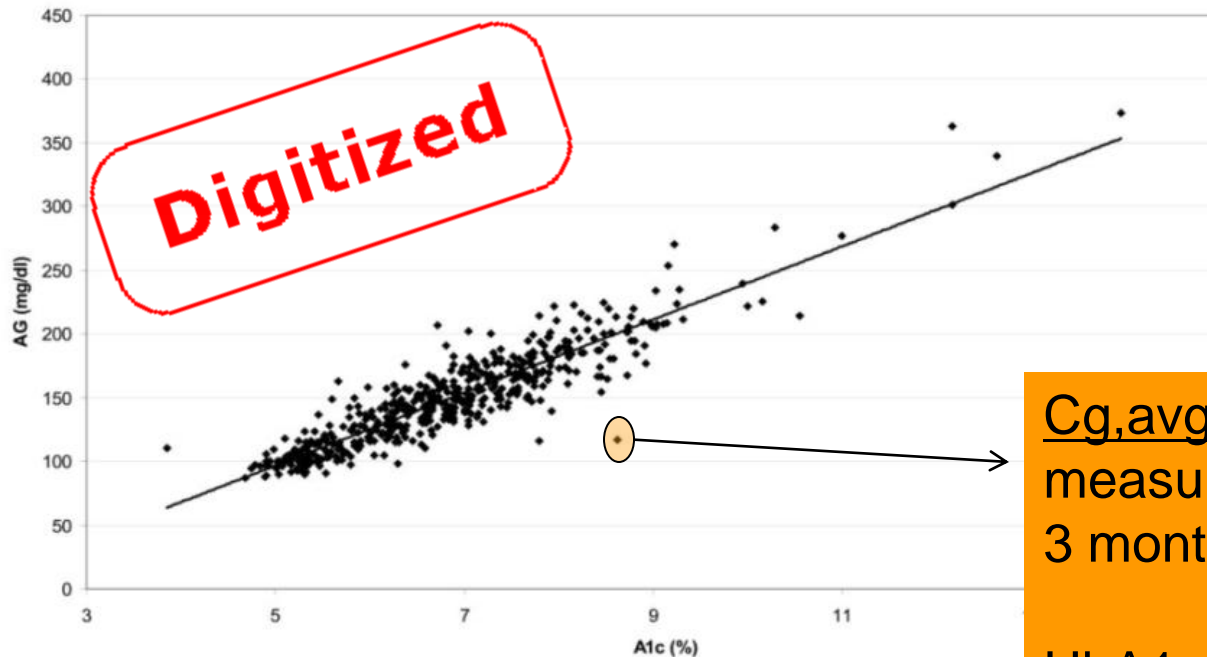
Mechanistic model in NONMEM: IGRH model



# Methods – Integrating literature data

**Main analysis:** ADAG study - Nathan et al. 2008

Cg,avg ~ HbA1c relationship  
(N=507 ; Diabetic & Non-diab.)



Cg,avg ( CGM ~2500  
measurements / subject in a  
3 month period)

HbA1c (monthly measured)

Figure 1—Linear regression of A1C at the end of month 3 and calculated AG during the preceding 3 months. Calculated  $AG_{mmol} = 1.59 \times A1C - 2.59$  ( $R^2 = 0.84$ ,  $P < 0.0001$ ).





# Methods - Structural model

## HbA1c formation:

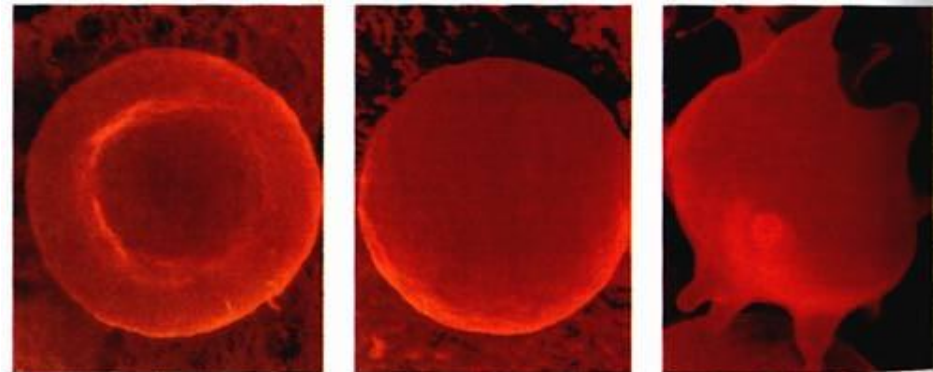
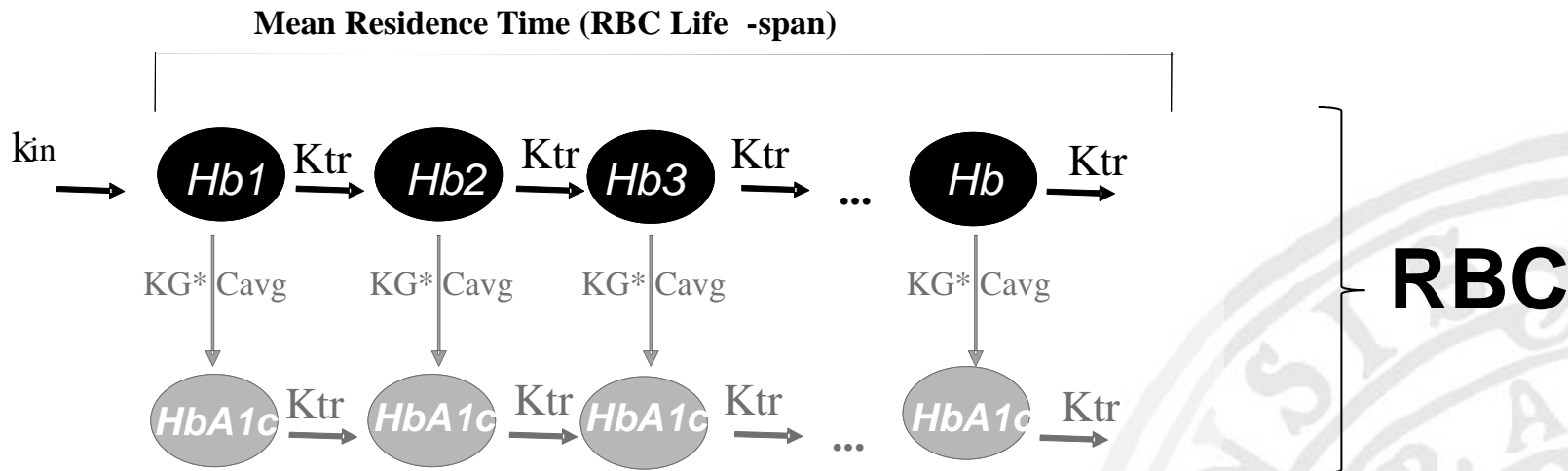
- i) RBC life-span and life-span distribution
- ii) Synthesis rate of HbA1c:  $f(C_{g,avg})$
- iii) HbA1c contribution in RBC precursors
- iv) Impact of HbA1c and  $C_{g,avg}$  measurement imprecision
- v) The fractional nature of HbA1c



# Methods: Structural model

## i) RBC life-span and life-span distribution

**Prior:** Kalicki et al. (PAGE meeting 2009)



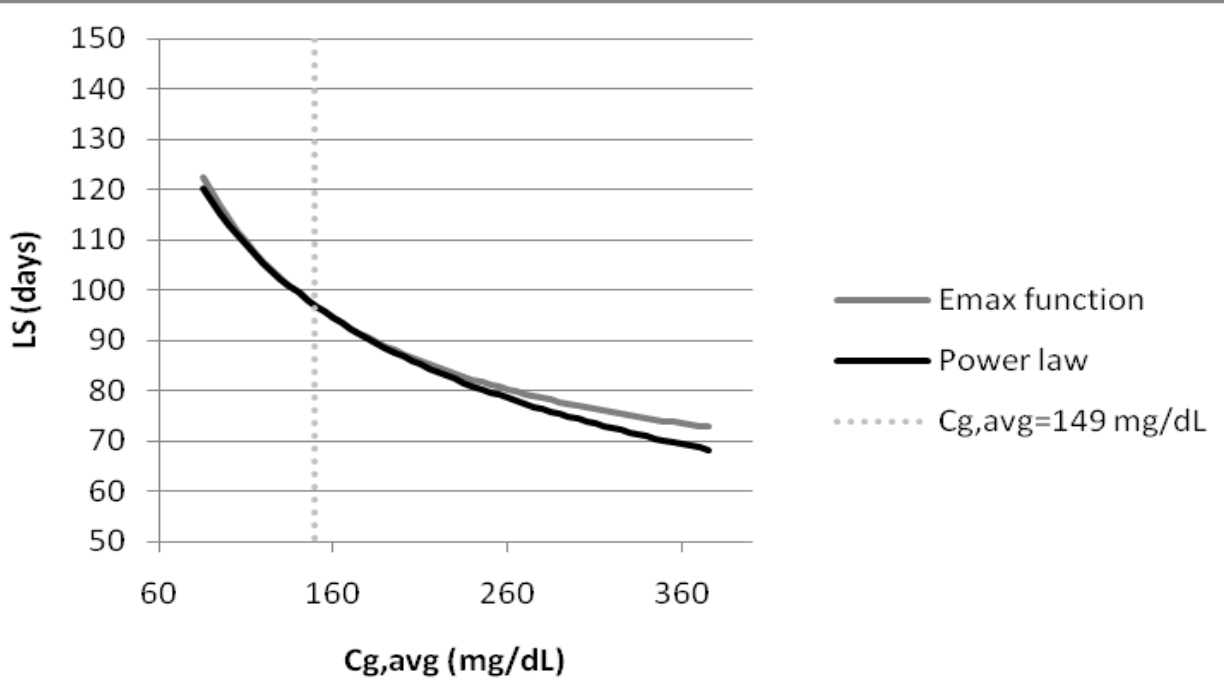


# Methods: Structural model

## i) RBC life-span and life-span distribution

### Influence of $C_{g,avg}$ on RBC life-span

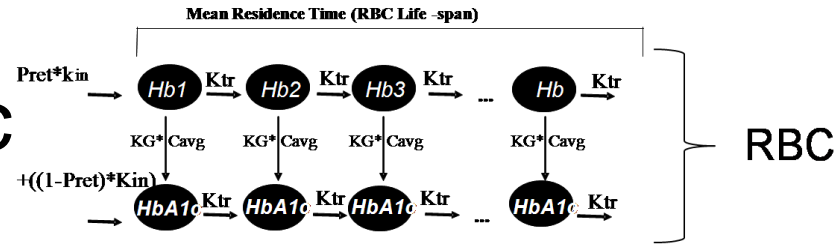
Power  $LS = TVLS \cdot \left( \frac{149}{C_{g,avg}} \right)^\delta \cdot \exp^\eta$





# Methods: Structural model

## ii) Synthesis rate of HbA1c



Linear  $Glycosilationrate = KG \cdot C_{g, avg} \cdot Hb$

Non-Linear  $Glycosilationrate = KG \cdot \left( \frac{149}{C_{g, avg}} \right)^\delta \cdot Hb$

**PRIORS from literature**

- Beach et al. 1979
- Higgins et al. 1981
- Mortensen-Volund et al. 1984
- Ladyzynski et al. 2008

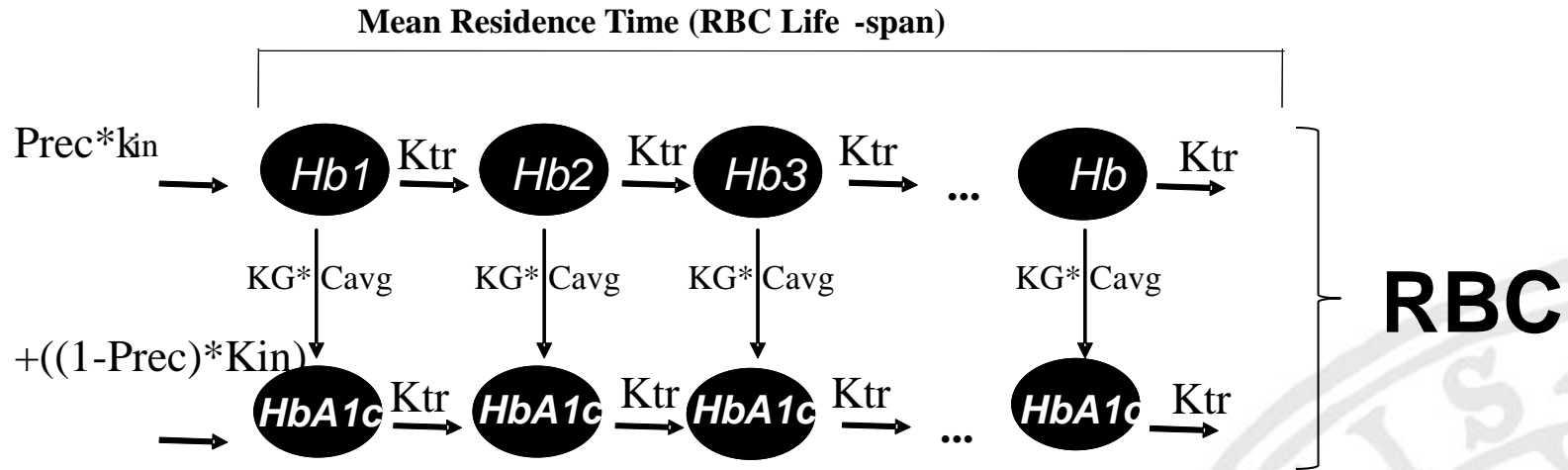
Non-Linear due to  
Glut1 saturation

$Glycosilationrate = KG \cdot \left( \frac{C_{g, avg} \cdot Km}{C_{g, avg} + Km} \right) \cdot Hb$



# Methods: Structural model

## iii) HbA1c contribution in RBC precursors



$$Prec = \exp(-KG * C_{g,avg} * LSP)$$

**PRIORS from literature**



## Hypothesis testing:

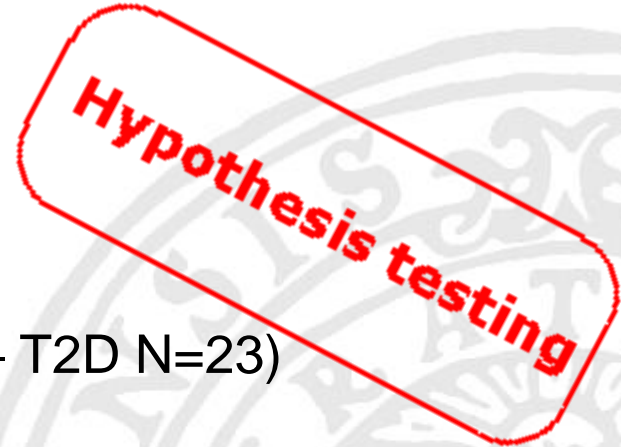
digitized data and clinical data as external validation for mechanisms

### Digitized literature data:

- Virtue et al. 2004 data (LS vs GHb – T2D N=23)

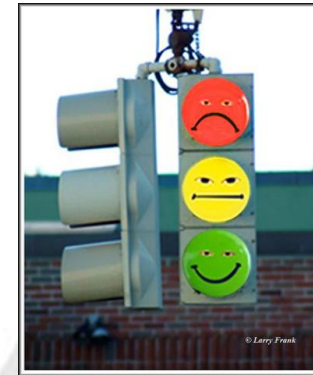
### Clinical data (shared by the authors)

- Nuttall et al. 2004 data (LS & GHb vs FPG - Non-diab. N=37)
- Ribbing et al. 2010 data (HbA1c vs FPG – T2D N=1460)





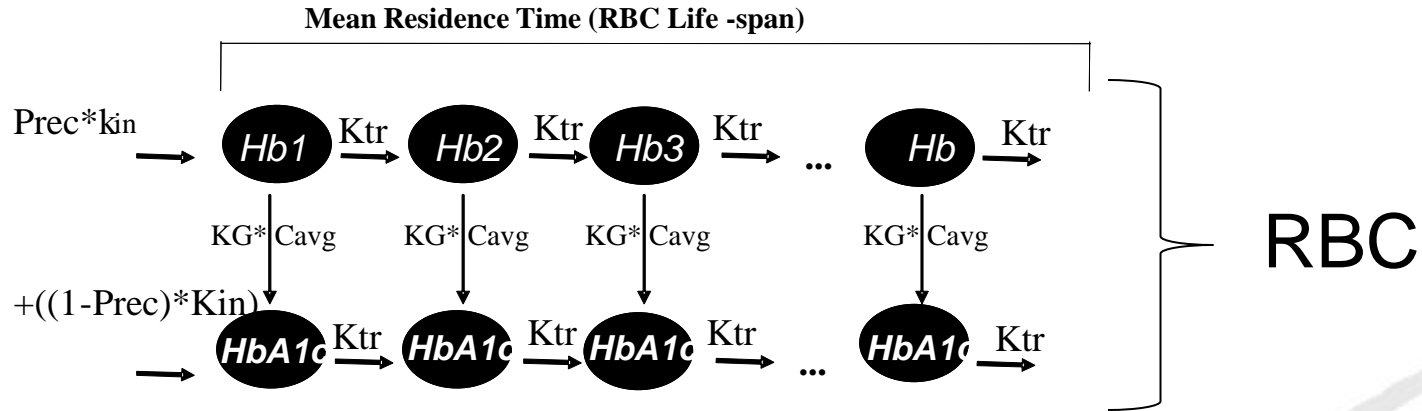
The integration of the  
data allowed to  
derived the



I<sub>ntegrated</sub> G<sub>lucose</sub> R<sub>BC</sub> H<sub>bA1c</sub> model



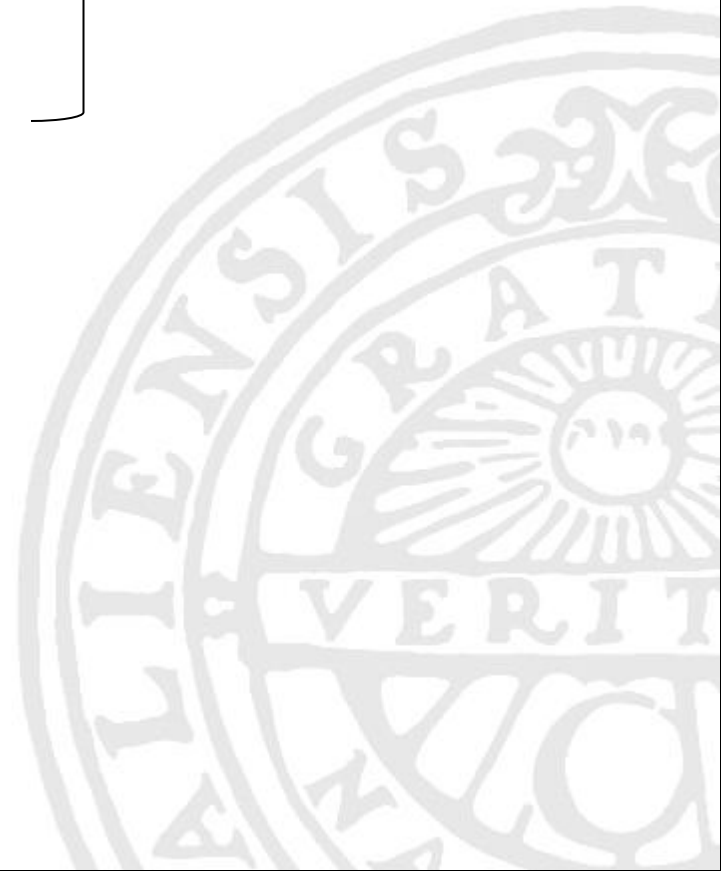
# Results – Final $I_{\text{ntegrated}}$ $G_{\text{lucose}}$ $R_{\text{BC}}$ $H_{\text{bA1c}}$ model



$$\text{Prec} = \exp(-KGP * C_{g,avg} * \text{LSP})$$

Power  $LS = TVLS \cdot \left( \frac{149}{C_{g,avg}} \right)^\delta \cdot \exp^\eta$

Linear  $Glycosilationrate = KG \cdot C_{g,avg} \cdot Hb$



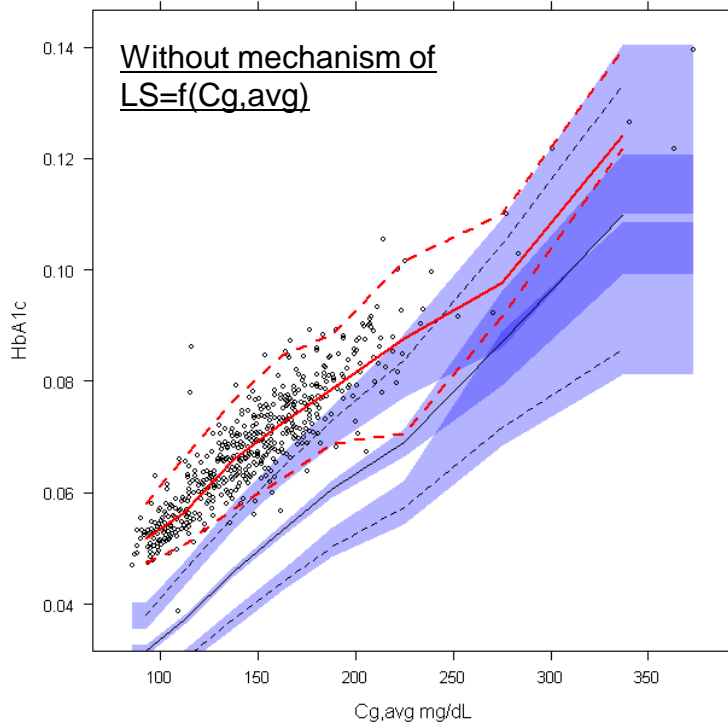




# Results – Final $I_{\text{ntegrated}} G_{\text{lucose}} R_{\text{BC}} H_{\text{bA1c}}$ model

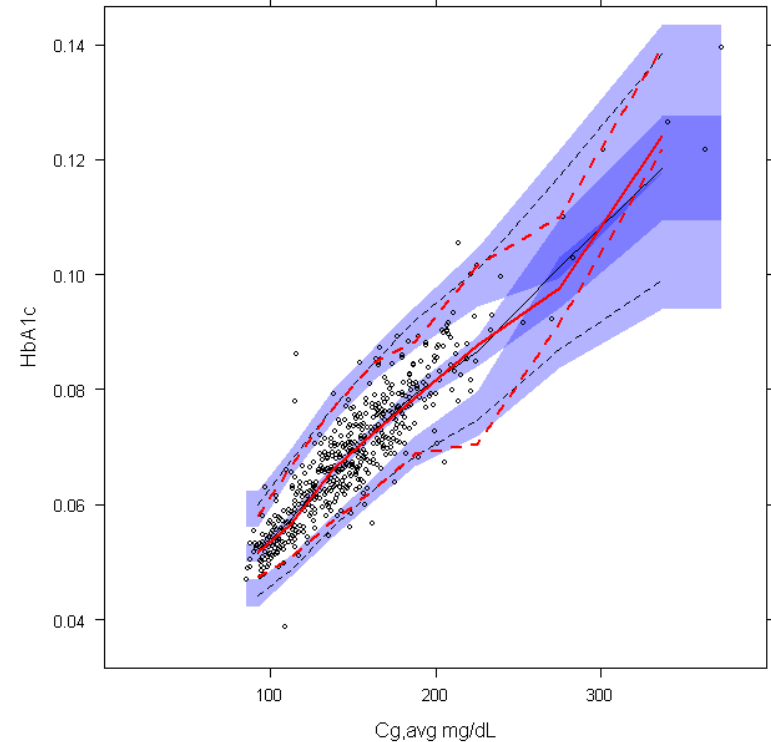
OFV=-4499.935

VPC run109



OFV=- 4888.172

VPC Final model (run86)



**Dots: Observations**

**Red lines: 5th, 50th & 95th percentiles of the observations**

**Black lines: 5th, 50th & 95th percentiles of the predictions (1000 simulations)**

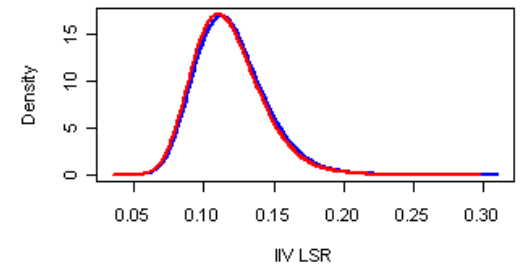
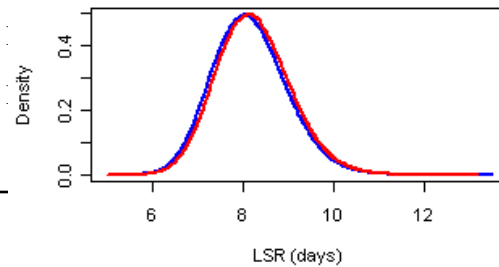
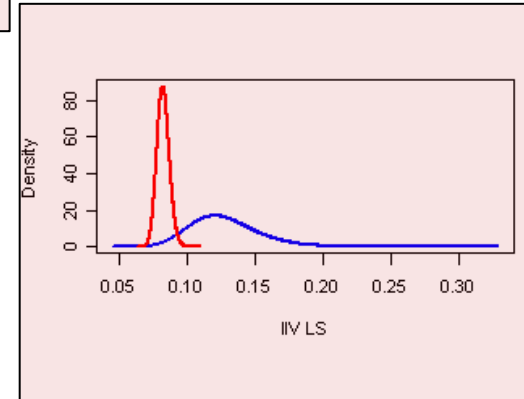
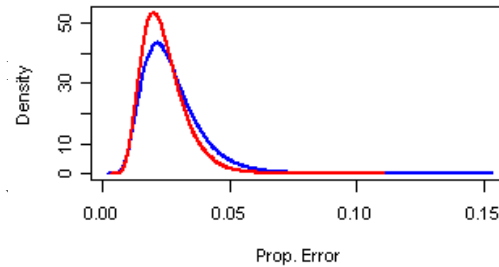
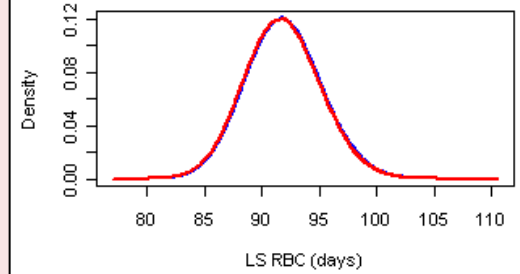
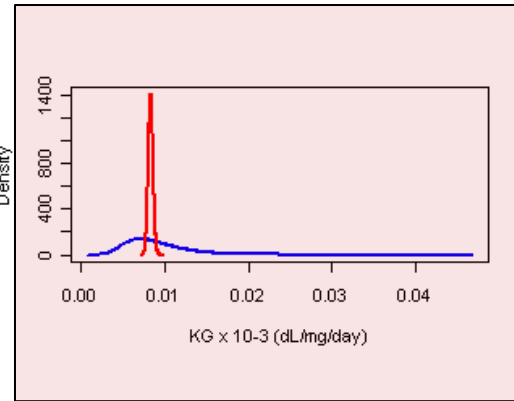
**Blue area: 90% CI of prediction intervals**



# Results – Final $I_{integrated}$ $G_{lucose}$ $R_{BC}$ $H_{bA1c}$ model

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Parameter	Estimated value	RSE%	Prior	RSE%
KG (dL/mg/day)	$8.37 \times 10^{-6}$	3.4%	$8.23 \times 10^{-6}$	27%
LS RBC (days)	91.7	3.6%	91.8	3.6%
IIV LS RBC (CV%)	8.22 %	5.5%	12.4%	19%
LS Prec. (days)	8.20	9.8%	8.11	10%
IIV LS Prec. (CV%)	11.5%	20%	12%	19%
Prop. RE	2.27 %	35%	2.5%	40%
$\delta$ (LS - Cg,avg)	0.381	4.5%		
OFV	-4888.172			





# Results – Final I<sub>integrated</sub> G<sub>lucose</sub> R<sub>BC</sub> H<sub>bA1c</sub> model

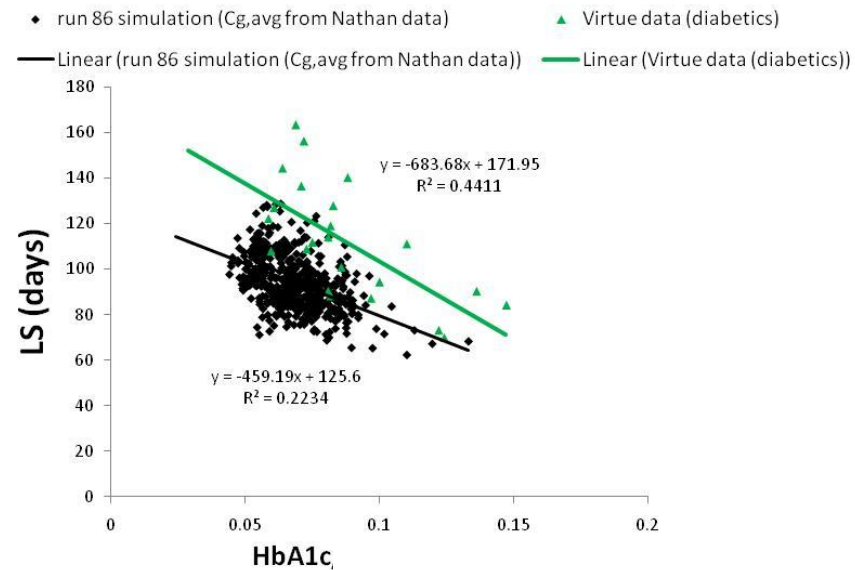
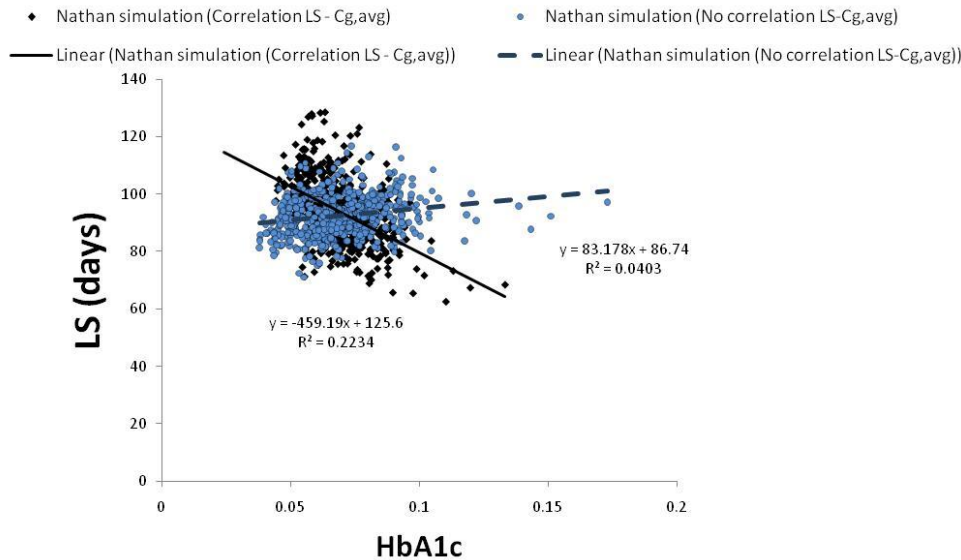
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## Data supporting this relationship between LS- Cg,avg:

Ribbing et al. (HbA1c vs FPG)  $\delta=0.48$  vs  $\delta$  IGRH model=0.38

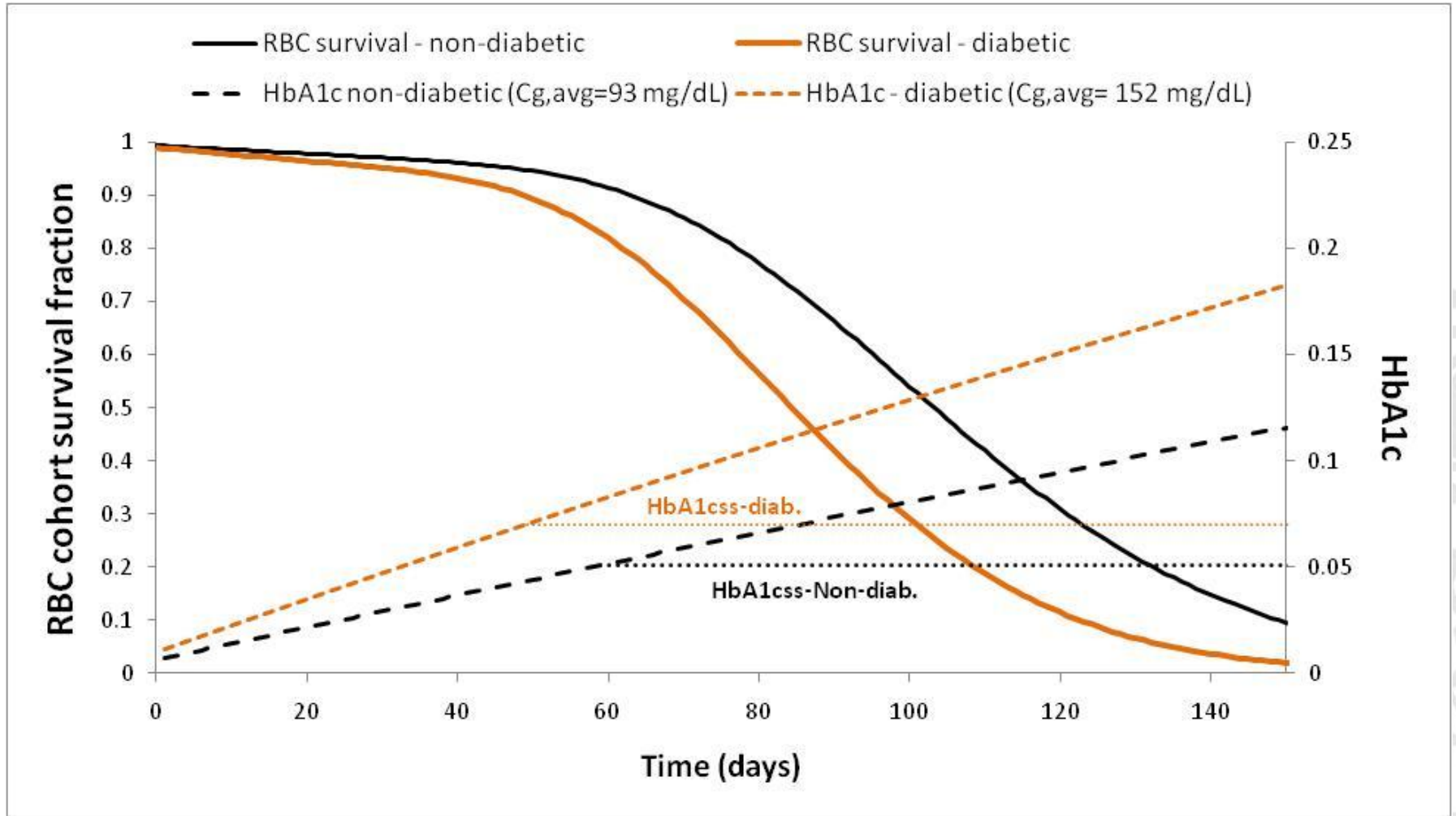
Nuttall et al. (LS vs FPG)  $\delta=0.49$  vs  $\delta$  IGRH model=0.38

Virtue et al. (GHb vs LS)





# Results – Final $I_{\text{integrated}} G_{\text{lucose}} R_{\text{BC}} H_{\text{bA1c}}$ model





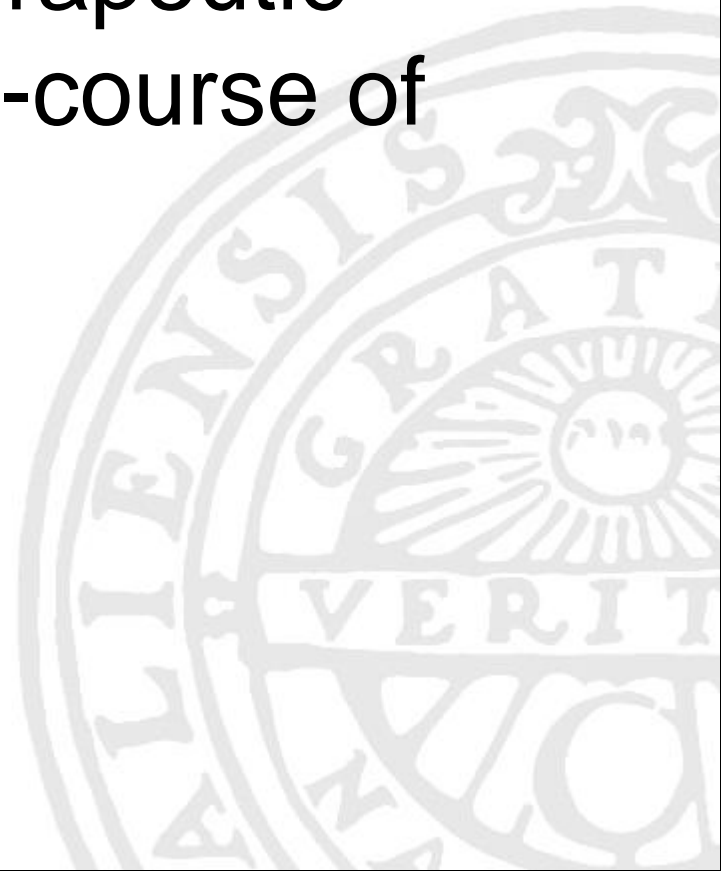
# Conclusions - I<sub>integrated</sub> G<sub>glucose</sub> R<sub>BC</sub> H<sub>bA1c</sub> model

- 1st quantitative description of the C<sub>g,avg</sub>-HbA1c relationship on mechanistic basis.
- The model describes well the relationship in both diabetic and non-diabetic patients



# Conclusions - I<sub>ntegrated</sub> G<sub>lucose</sub> R<sub>BC</sub> H<sub>bA1c</sub> model

- To predict the impact of changes in  $C_{g,avg}$  (due to diet or therapeutic interventions) on the time-course of HbA1c levels.





# Conclusions - I<sub>ntegrated</sub> G<sub>lucose</sub> R<sub>BC</sub> H<sub>bA1c</sub> model

- If any of the processes involved are subjected to change in an individual patient, the expected temporal and steady state change of HbA1c can also be predicted

(e.g. uremic patients (LS decreased))



# Conclusions - I<sub>ntegrated</sub> G<sub>lucose</sub> R<sub>BC</sub> H<sub>bA1c</sub> model

- Literature data can be used not only to support parameter estimates, but combined from different sources to test hypothesis and build structurally novel models!





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# Acknowledgements

- Prof. Frank Nuttall for sharing data

- *F. Hoffmann-La Roche* for financial support



Roche

The Roche logo, which consists of the word "Roche" in a blue, sans-serif font, enclosed within a blue hexagonal border. The logo is positioned to the left of the main text block for the second acknowledgment.