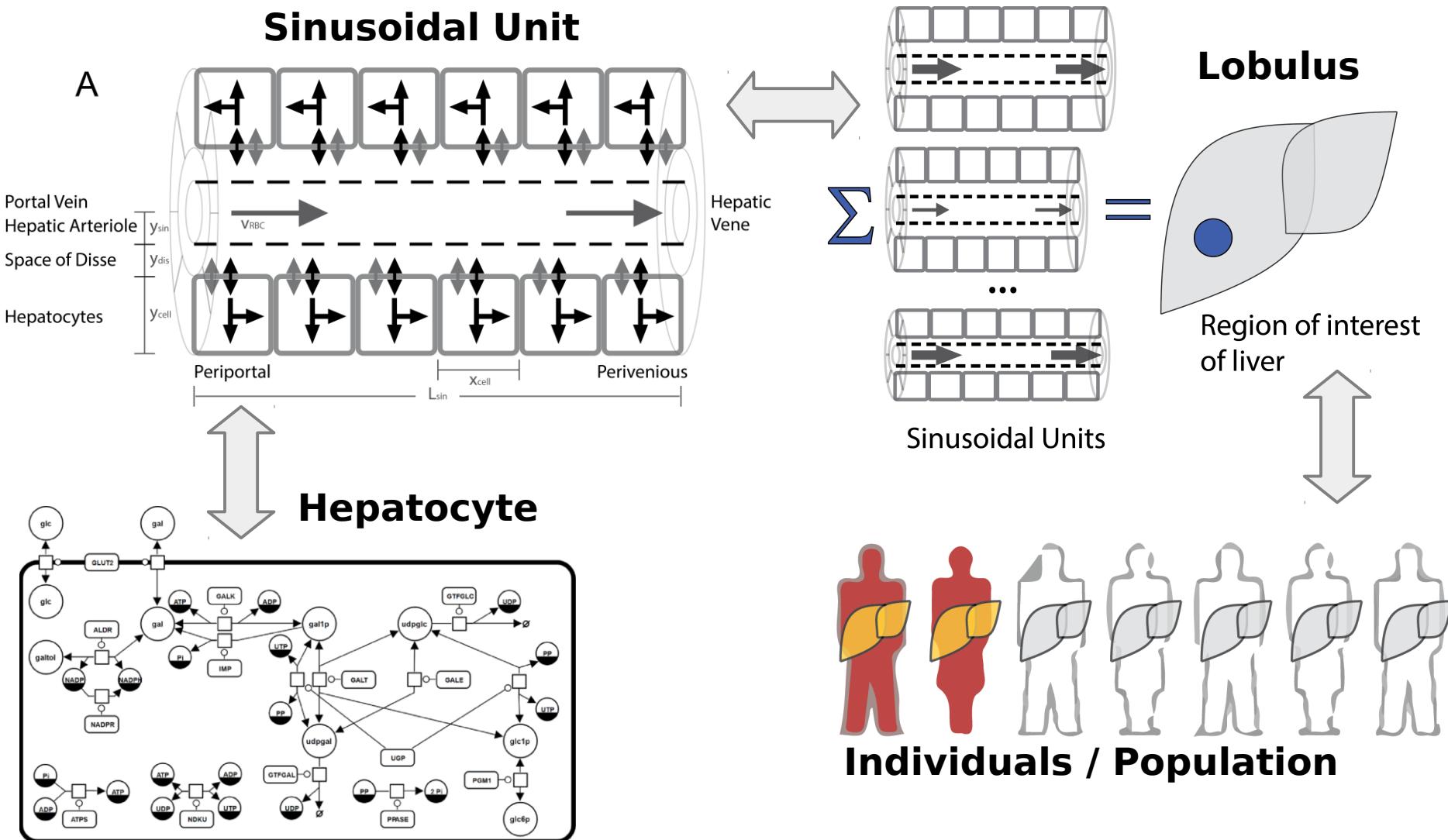


Population Variability in Liver Function

König M. & Holzhütter HG.



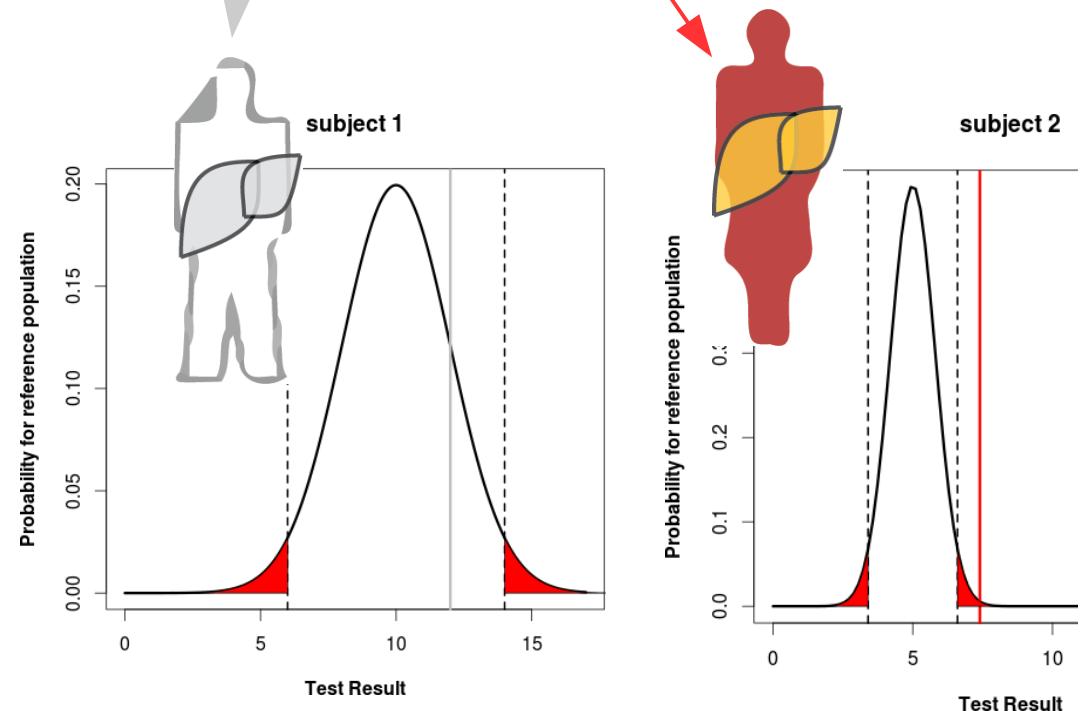
A high-angle photograph of a massive crowd of people filling the frame. The individuals are packed closely together, creating a complex, textured pattern of heads and shoulders. The crowd is diverse in age and appearance, with many young adults visible. The colors of their clothing range from dark blues and blacks to lighter whites and yellows, though the overall tone is somewhat muted due to the density and lighting.

We are all different ...

... so why no individualized predictions?

Individualized Liver Function Tests

- Large population variability in liver volume & blood flow
 - Age, gender, bodyweight, height, ethnicity, ...
- What are the effects of this variability on model prediction?
- Improved evaluation of liver function in given anthropomorphic context
 - Individualized liver function test
 - Reference range for sample of comparable individuals
 - Values outside range → further investigation & treatment



Variability & Sensitivity

- **Variability**

- The focus of variability analysis is to evaluate the range of values that a parameter expected to be present in individuals may have in a population and the impact of that variability on the simulation prediction.

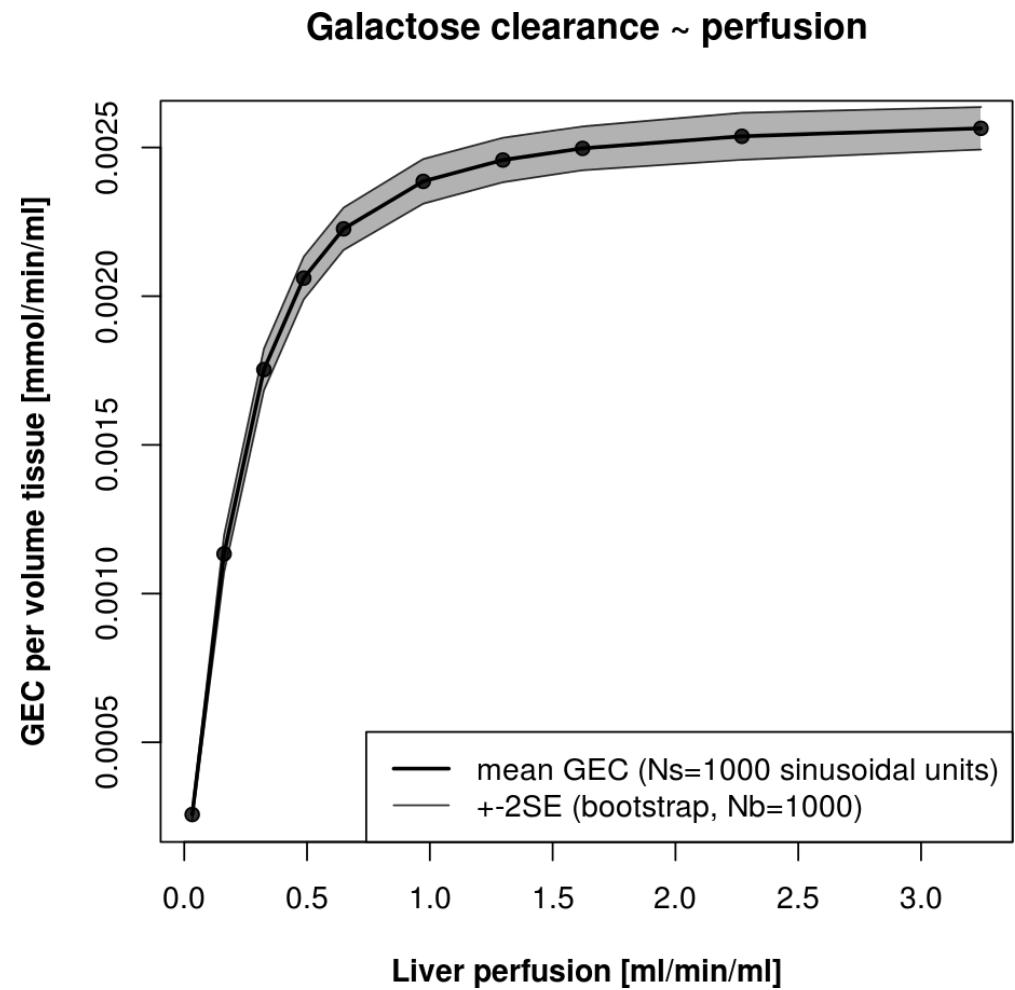
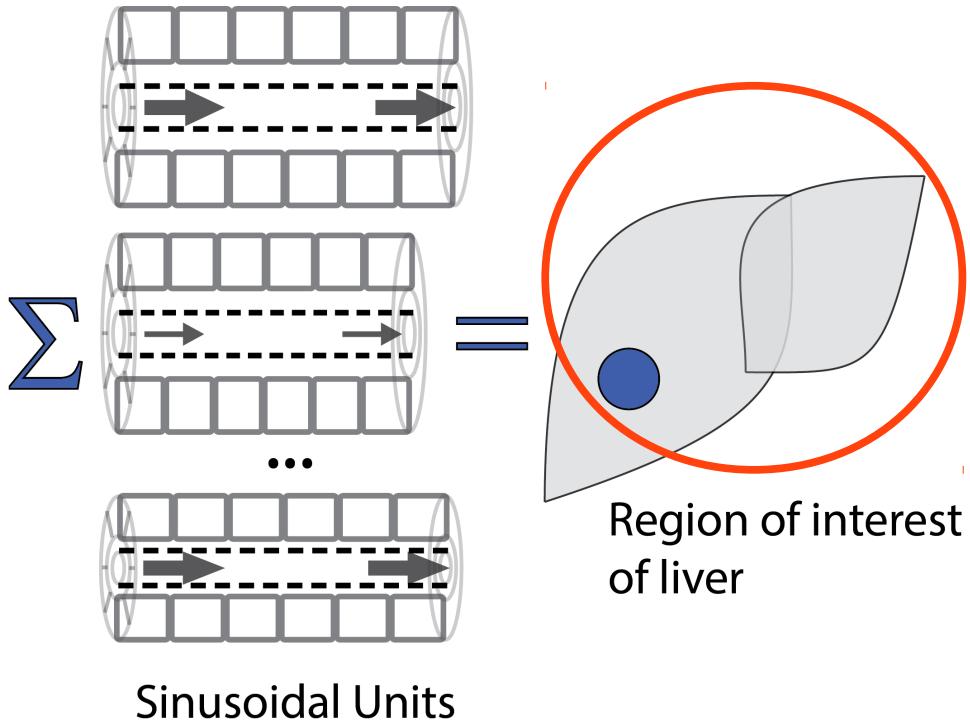


- **Sensitivity**

- Sensitivity analysis provides a quantitative evaluation of how parameters influence the simulation prediction of a model. Sensitivity analysis compares the magnitude in change in output for a defined change in each input parameter.

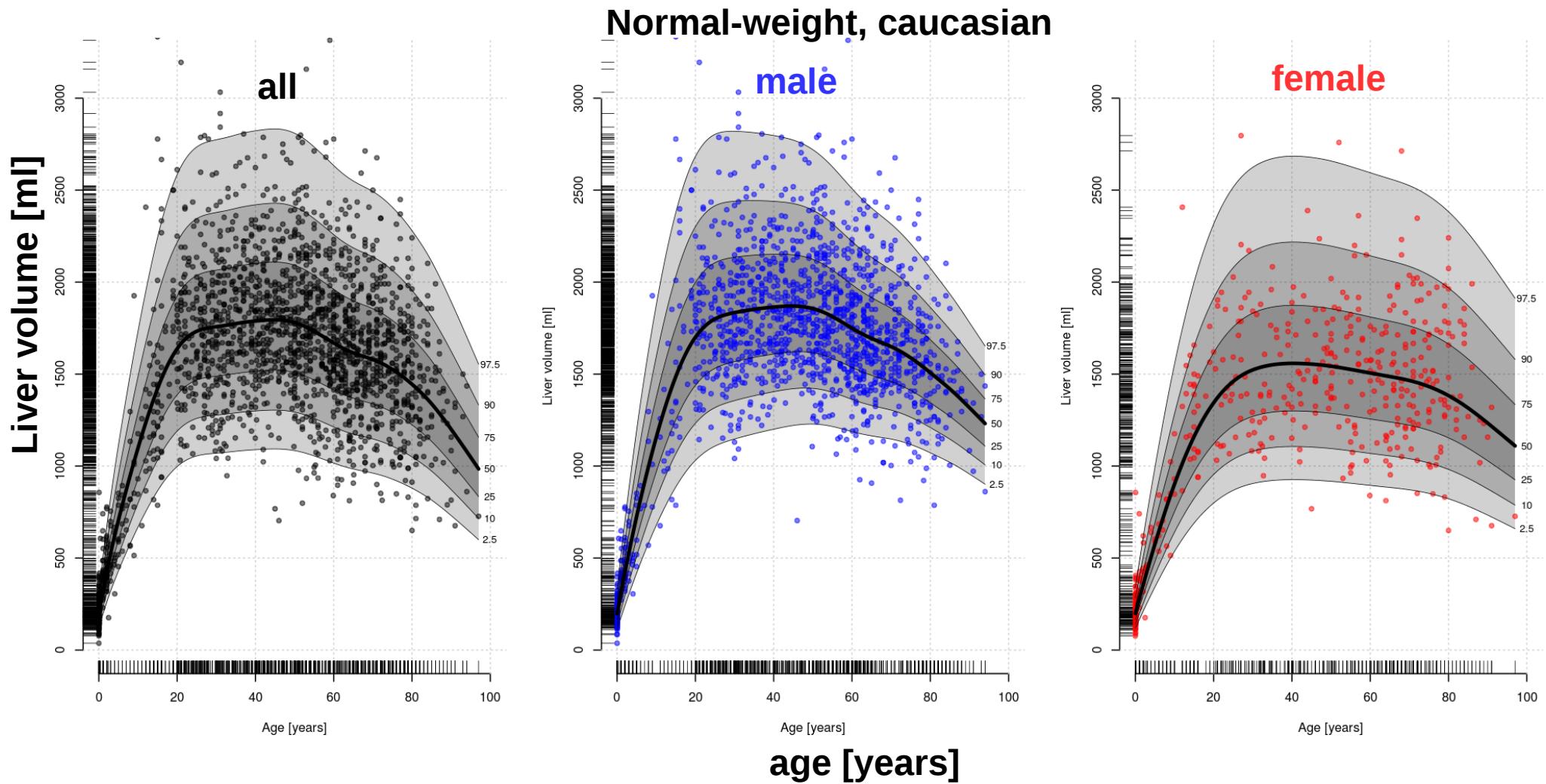
GEC - Liver Function Test

- Evaluation of liver function via galactose clearance capacity (GEC)
- Model prediction for galactose clearance per tissue volume for given perfusion in the region (**regional GEC**)
- Use individual **liver volume and perfusion** for scaling (**total GEC**)
- Evaluation of individual & population variability in GEC



Liver volume = f(sex, age, ...)

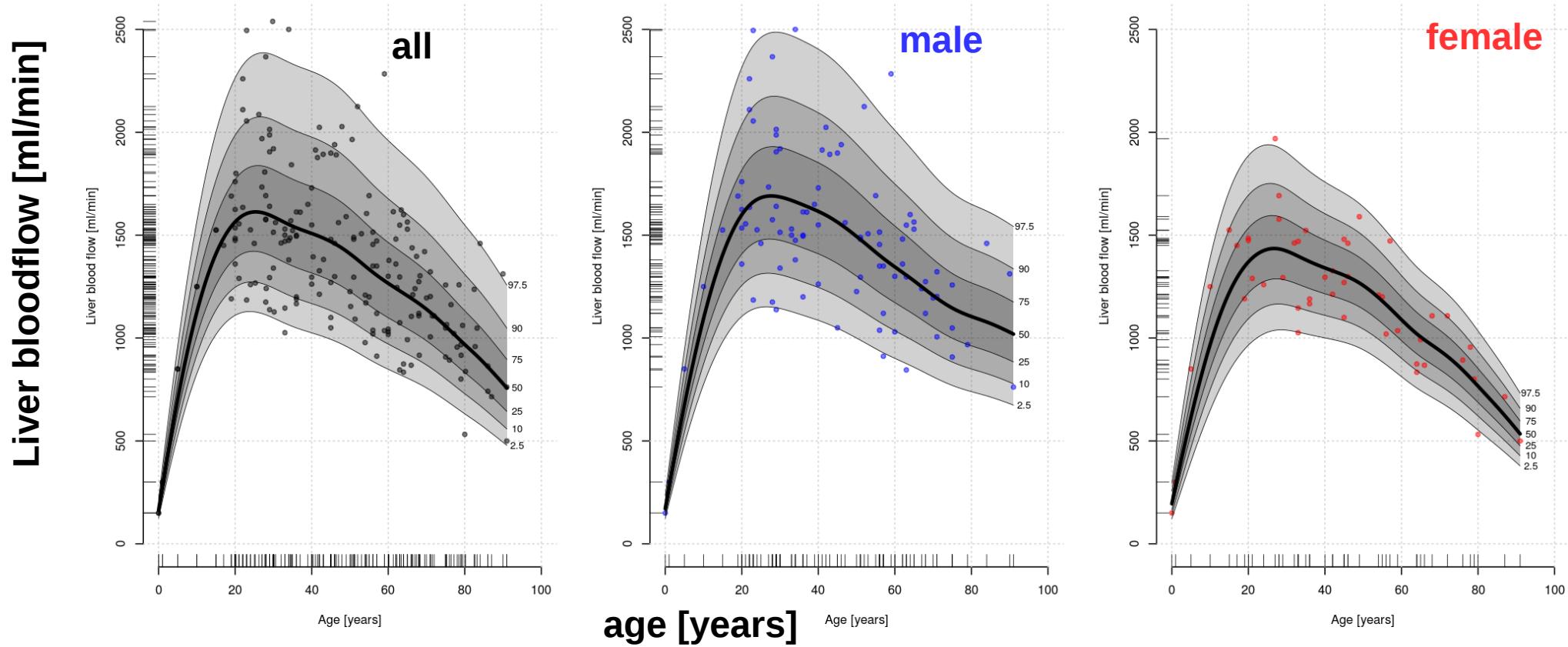
- Density estimation via generalized additive models (GAM) combined with centile estimation
- Combined dataset from multiple sources & experimental methods
- Analog $\text{volLiver} \sim \text{bodyweight}$, $\text{volLiver} \sim \text{height}$, $\text{volLiver} \sim \text{BSA}$, ...



Age & gender dependence blood flow

- Analog for bloodflow with additional $\text{flowLiver} \sim \text{volLiver}$ correlation
- Results:** probability densities for liver volume & liver bloodflow

Normal-weight, caucasian

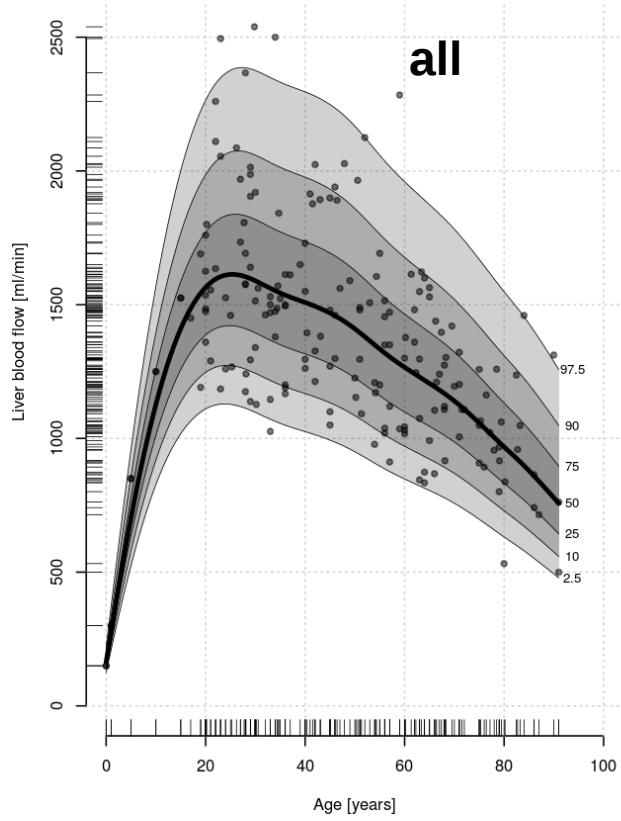


age=80 [y], sex=male

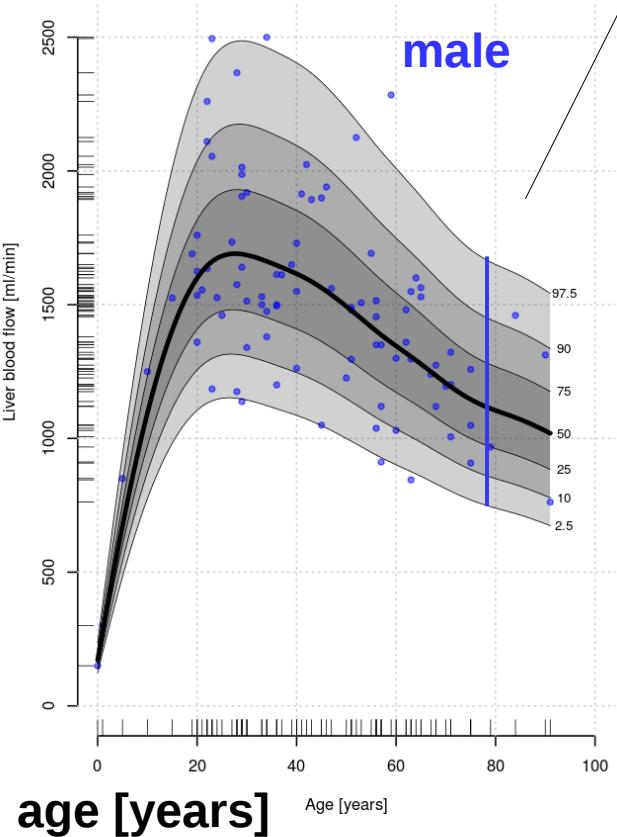
Probability distribution of liver blood flow

- normal-weight (BMI<24.9)
- caucasian
- male
- 80 years old

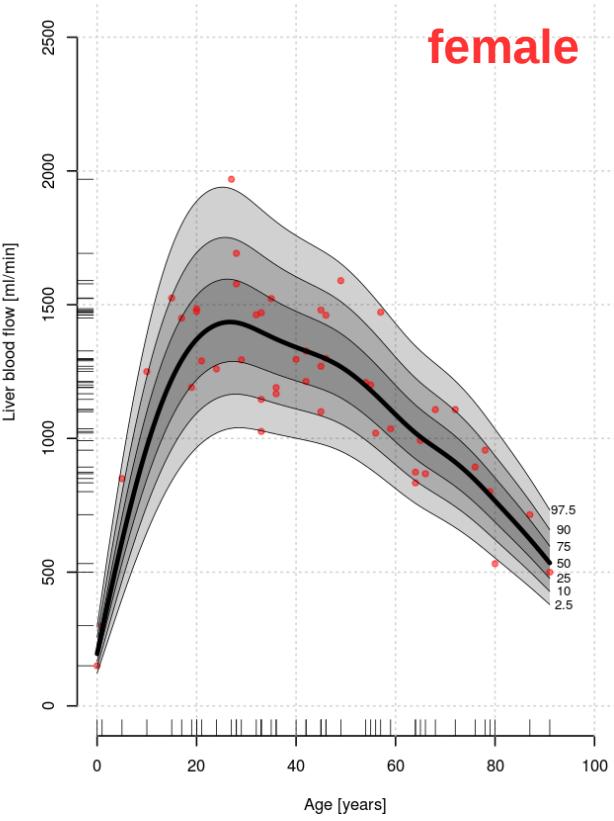
Liver bloodflow [ml/min]



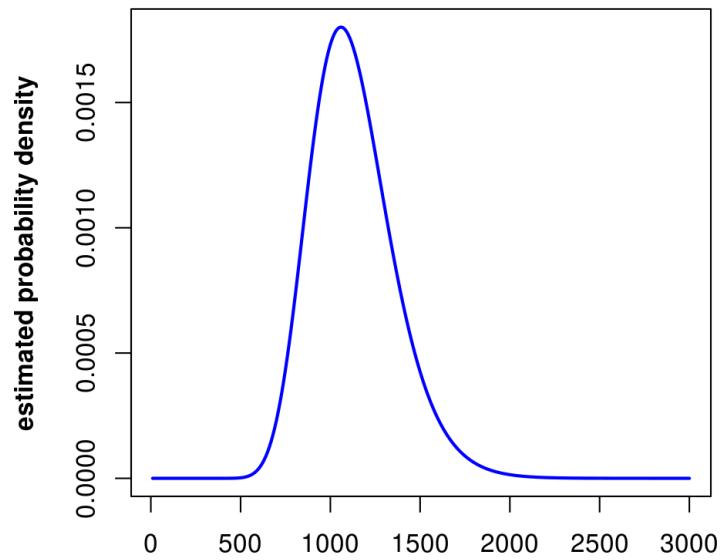
all



male



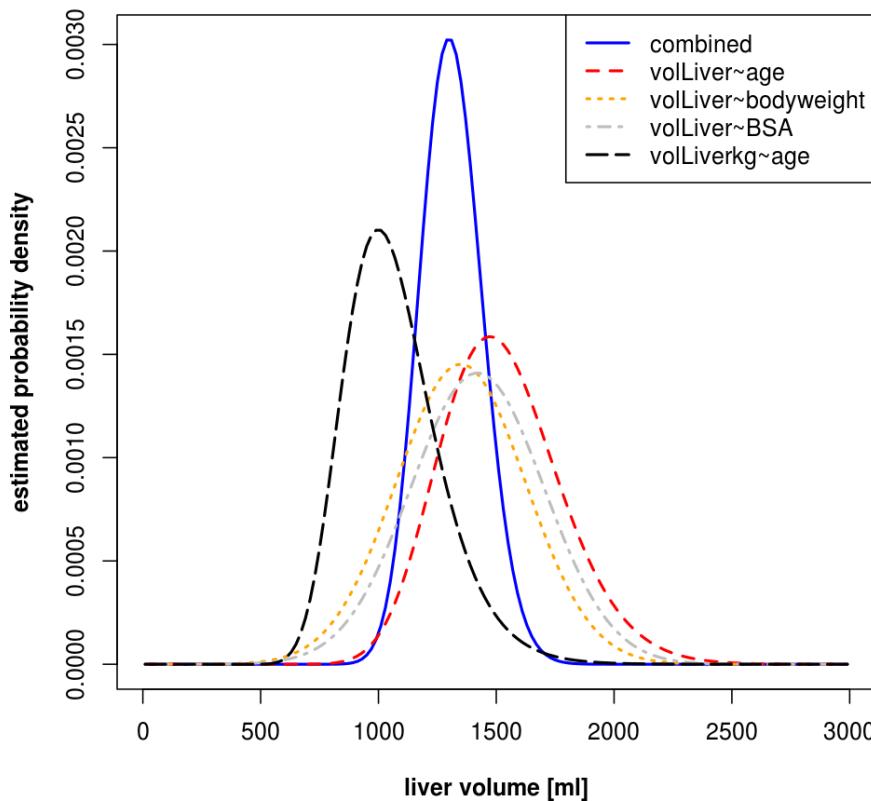
female



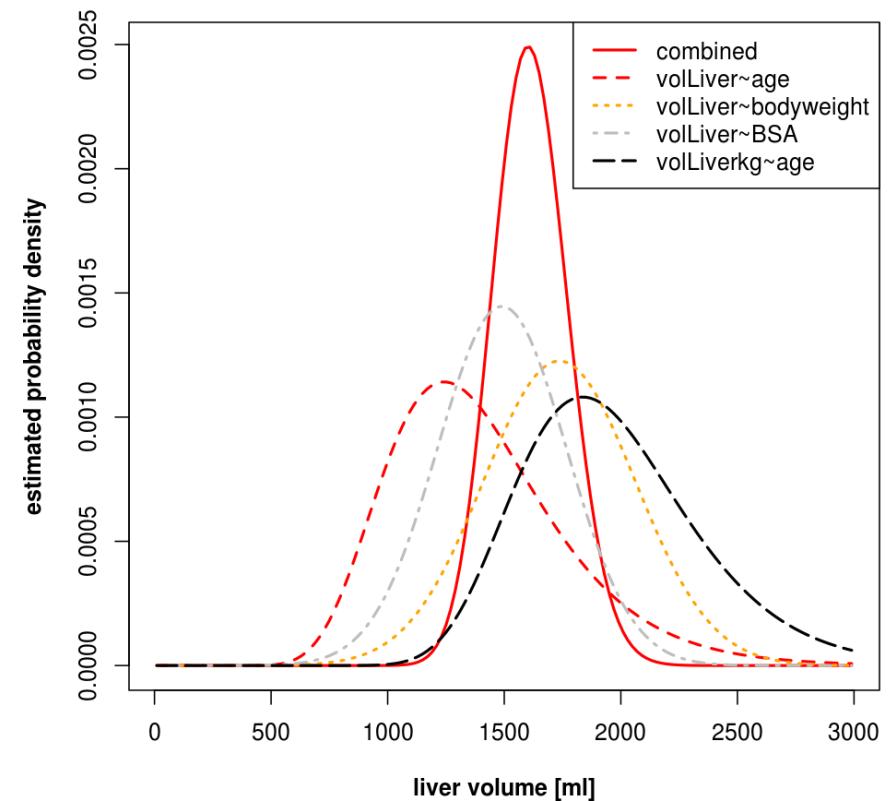
Combining the information

- Set of probability distributions for pair-wise correlations
- Important to keep the correlation structure between different data pairs
- Combination of individual probability distributions for best estimate
(use all the available knowledge)
- **Result: individual probability distributions** for liver volume and bloodflow based on normal reference population

age=80 [y], sex=male, bodyweight=55 [kg], BSA=1.6 [m^2]

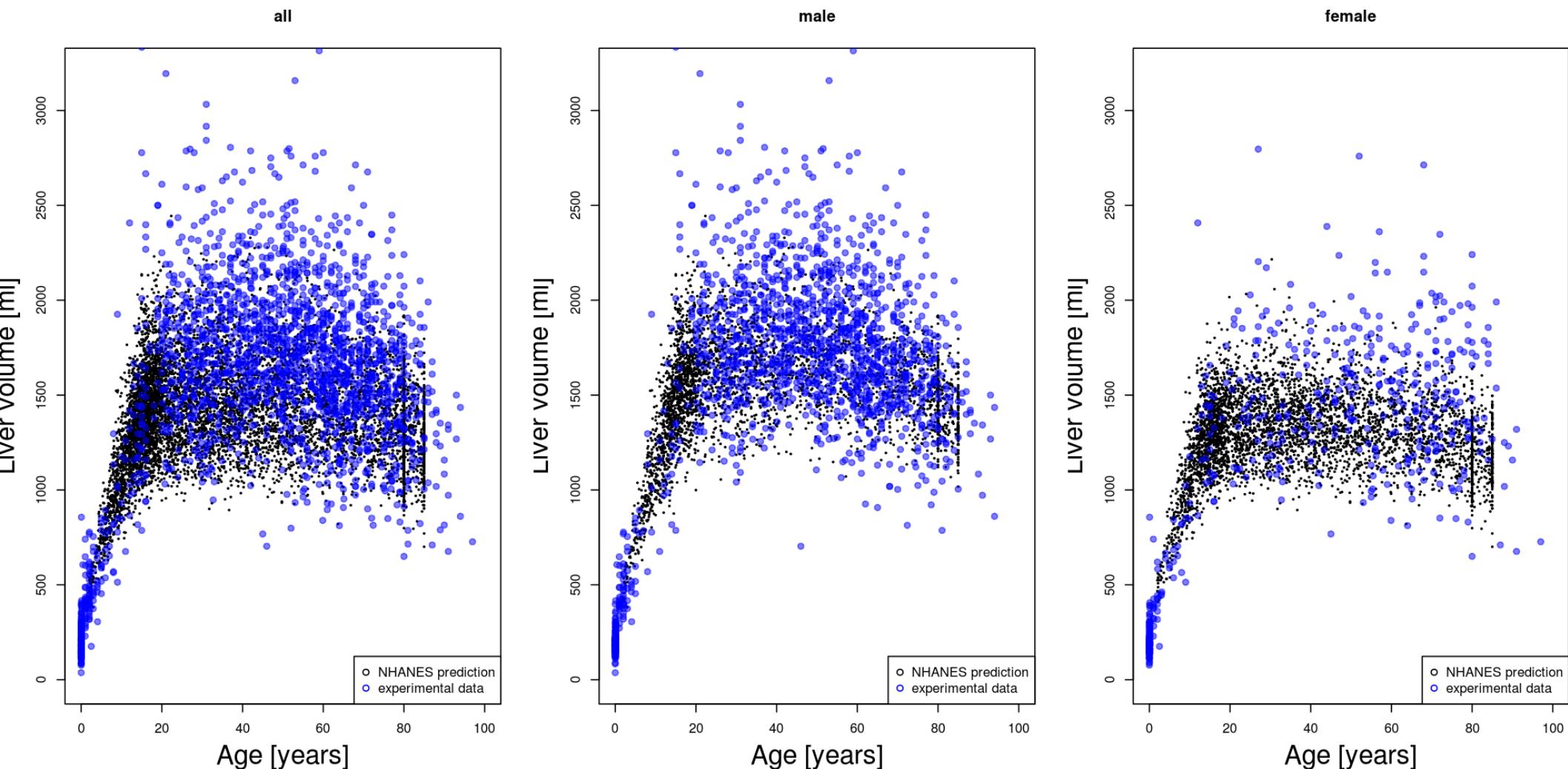


age=20 [y], sex=female, bodyweight=80 [kg], BSA=1.7 [m^2]

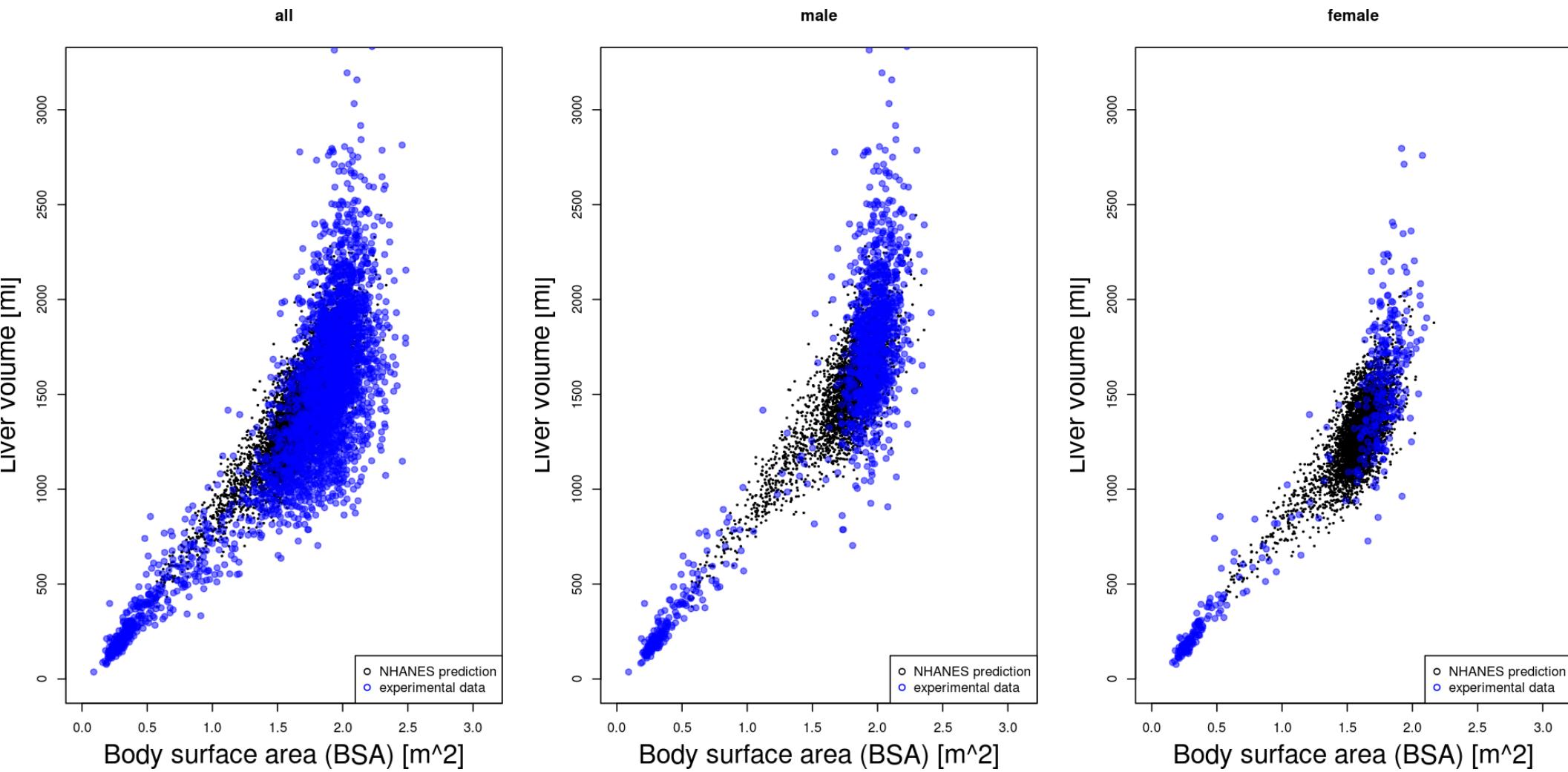


Prediction of population variability

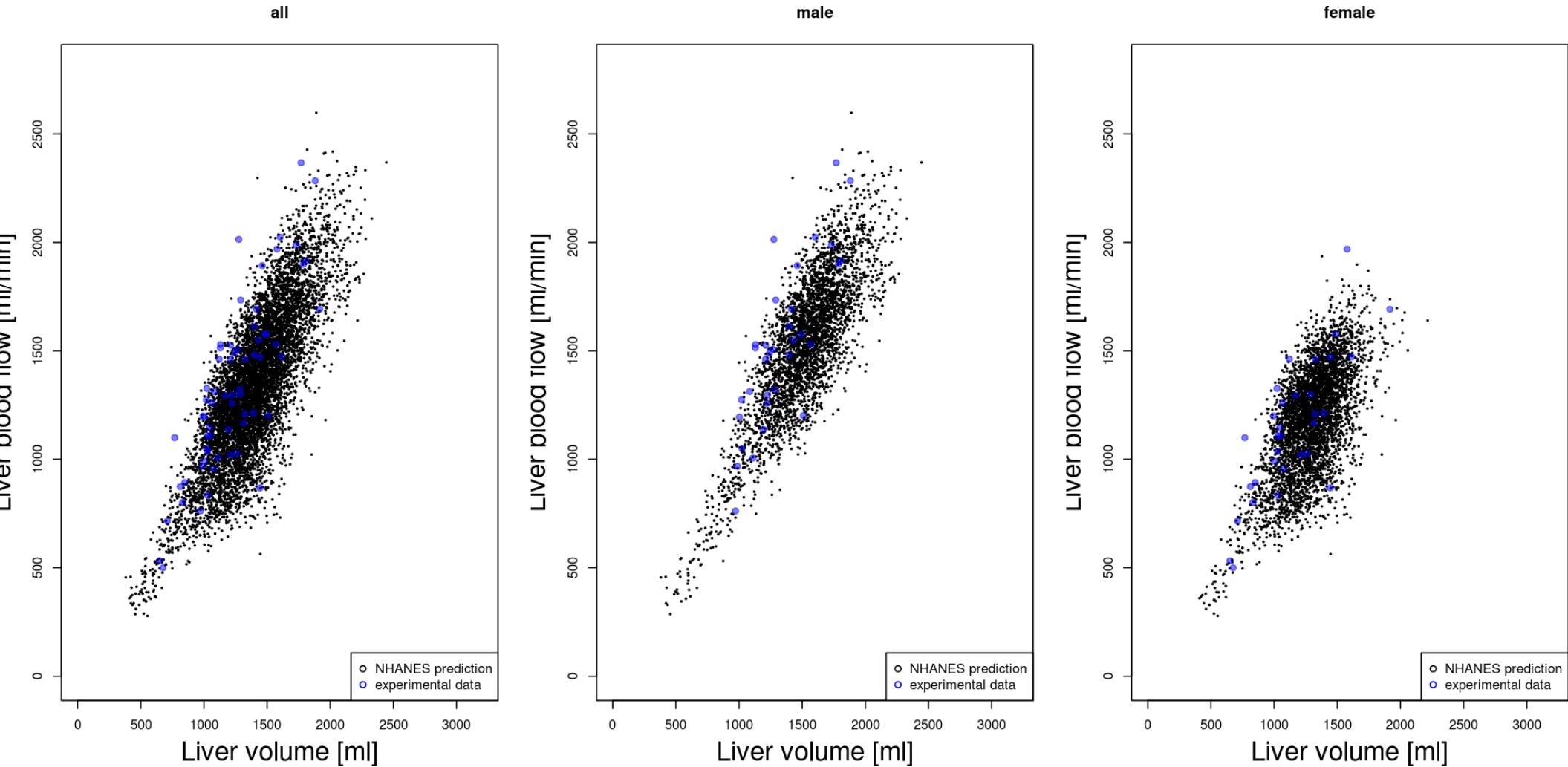
- NHANES prediction (~8000 normal-weight, caucasian)
 - Correlation structure of sex, age, bodyweight, height of US population
- Liver volume ~ age



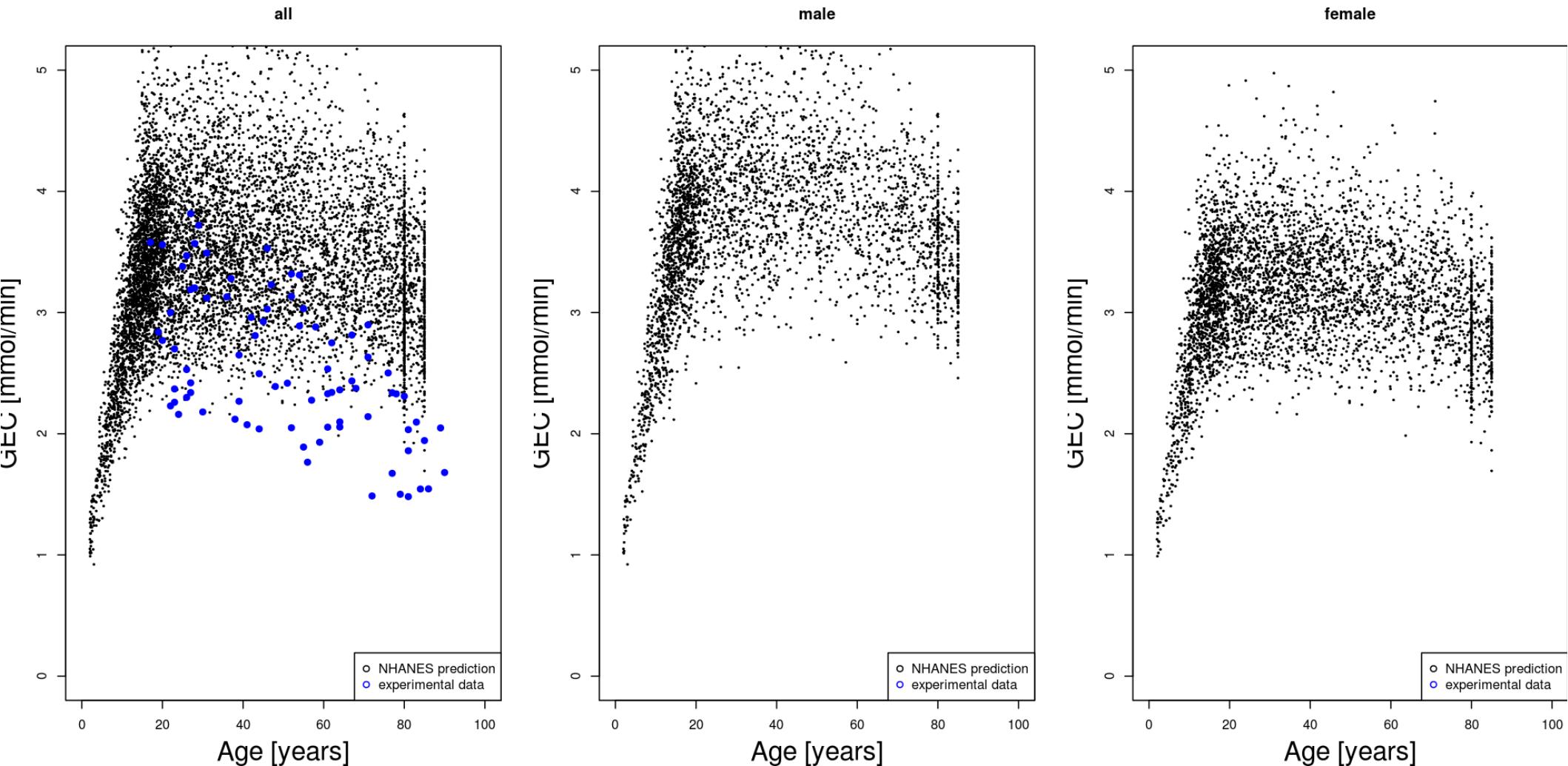
Liver volume ~ BSA (body surface area)



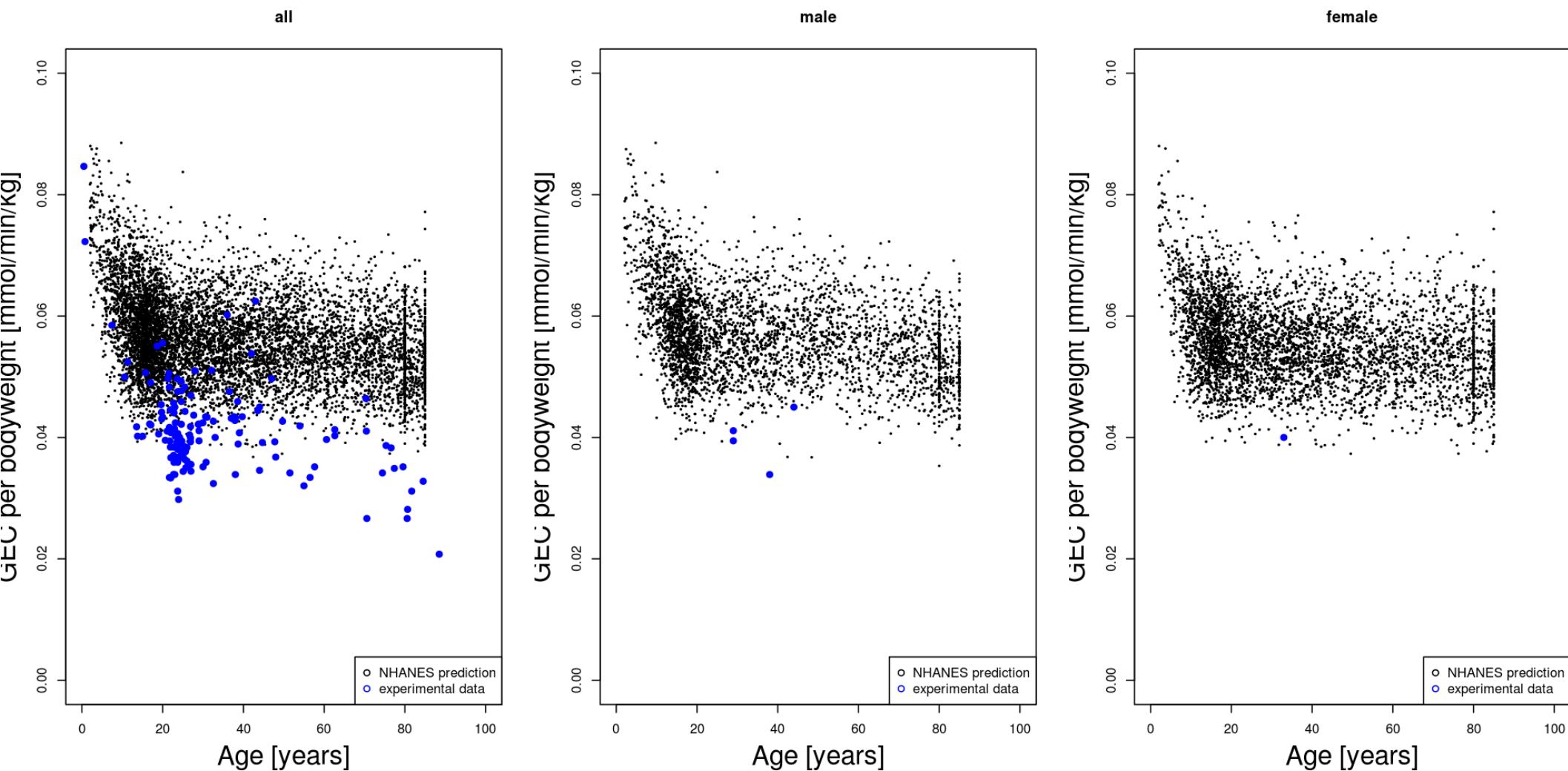
liver blood flow ~ liver volume



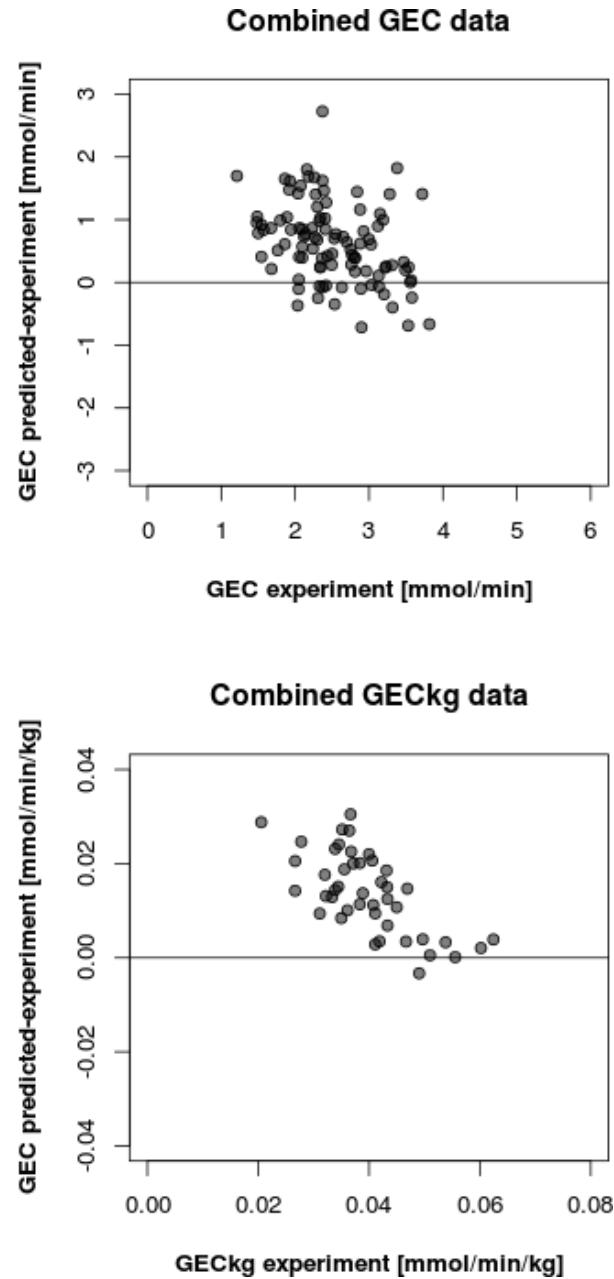
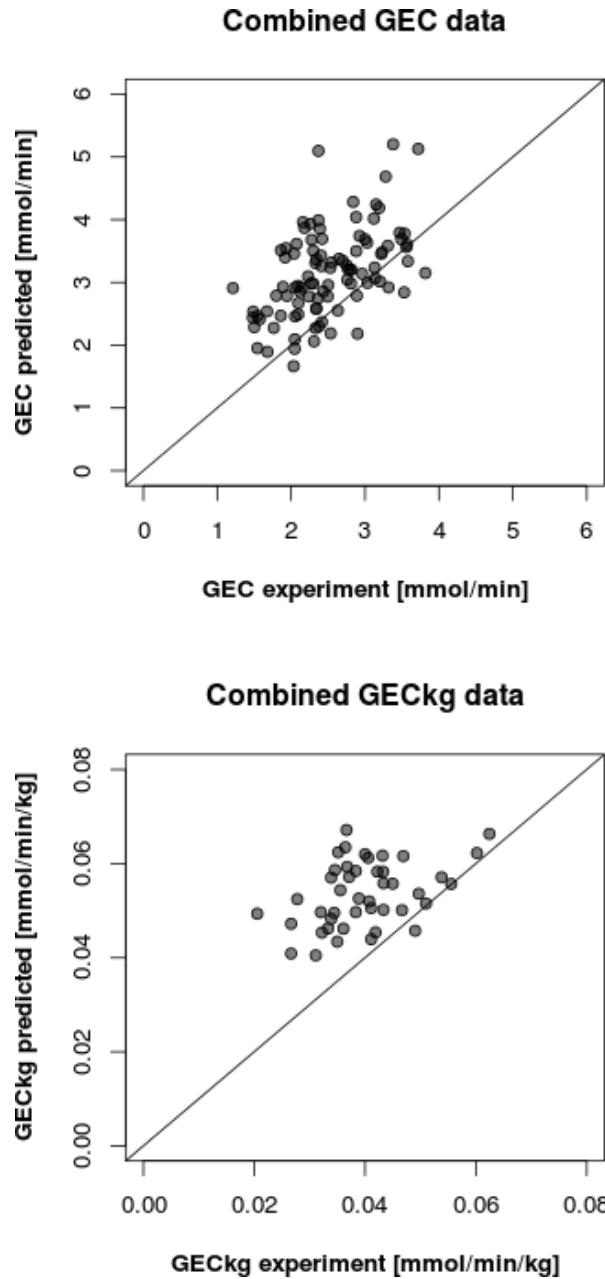
Prediction of GEC (galactose clearance capacity)



Prediction of GEC per bodyweight



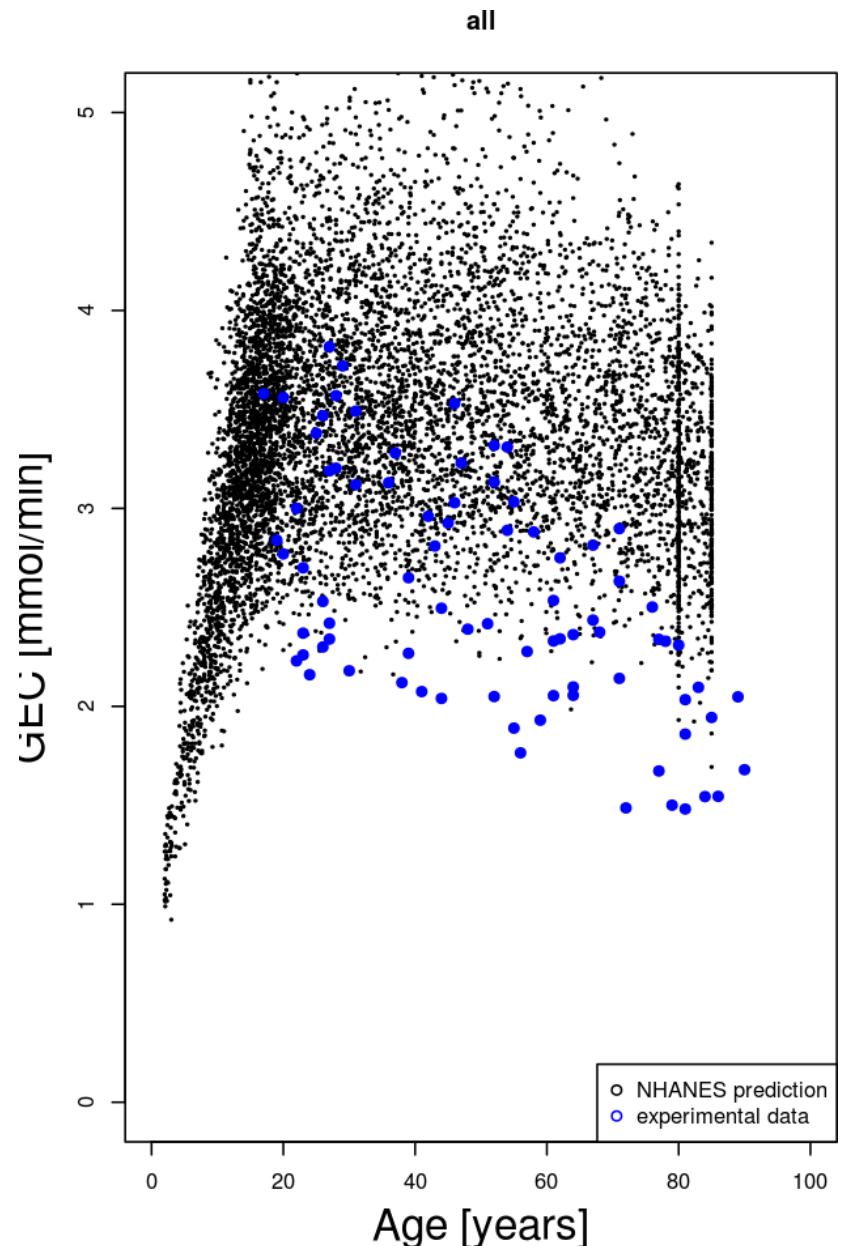
Individualized GEC prediction



First draft predictions,
no adaption of metabolic capacity

GEC variability

- Large part of GEC variability is explained by the population variability
 - Additional variability in metabolic (V_{max}) & morphological components in population (\rightarrow sensitivity analysis)
 - For instance age changes in structure (pseudo-capillarization) & decrease in metabolic capacity (?)
- Next steps
 - Adaption of galactose clearance on cellular level
 - Prediction of additional GEC datasets



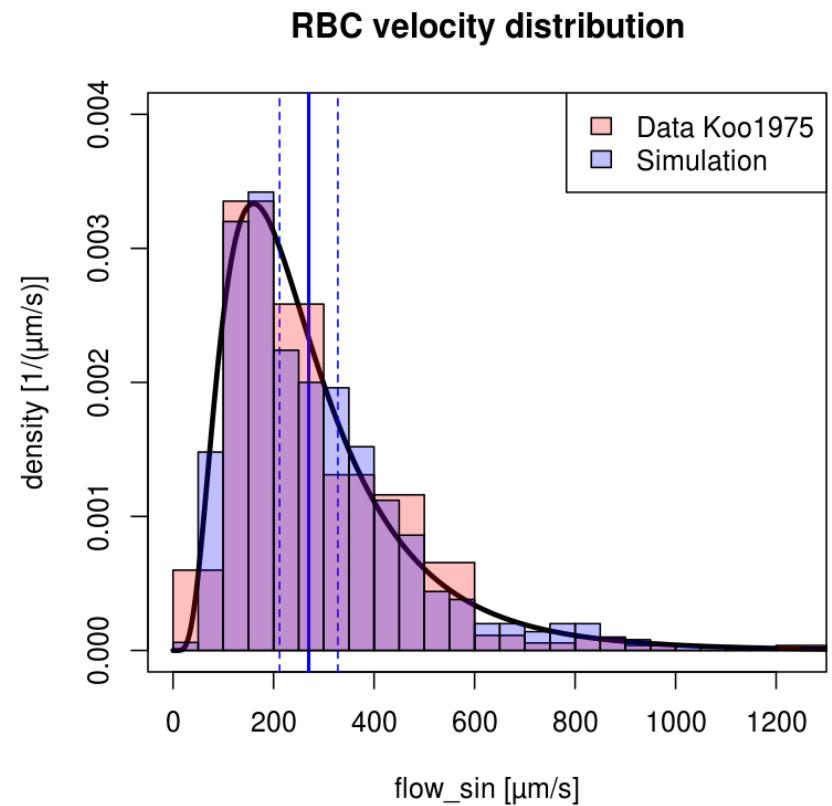
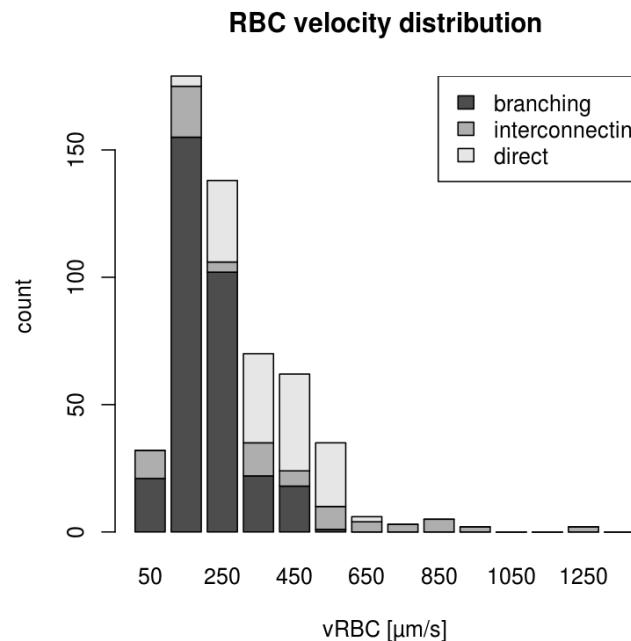
Acknowledgements

- **Concept & Modeling**
 - Prof. Holzhütter
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- **Funding**
 - Charité
 - Virtual Liver Network



Fitting Distributions

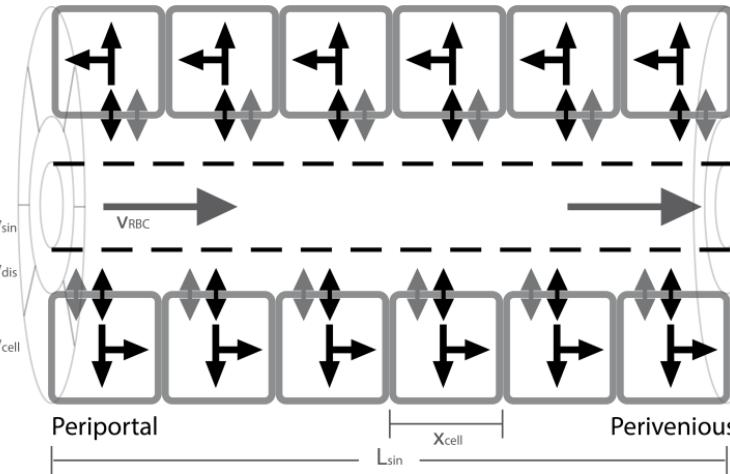
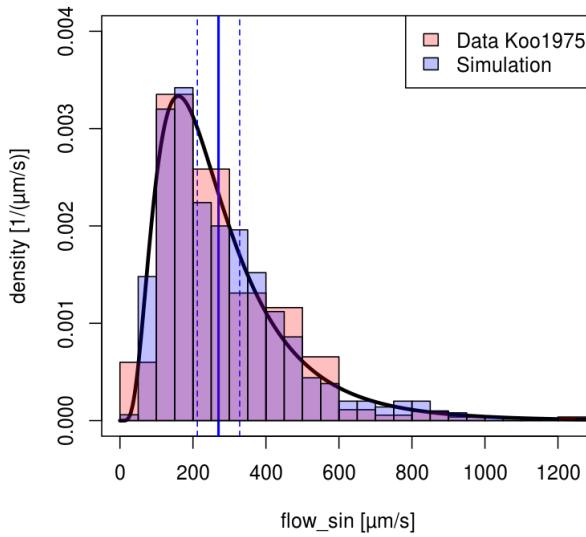
- maximum-likelihood method for univariate distributions
(log-normal)



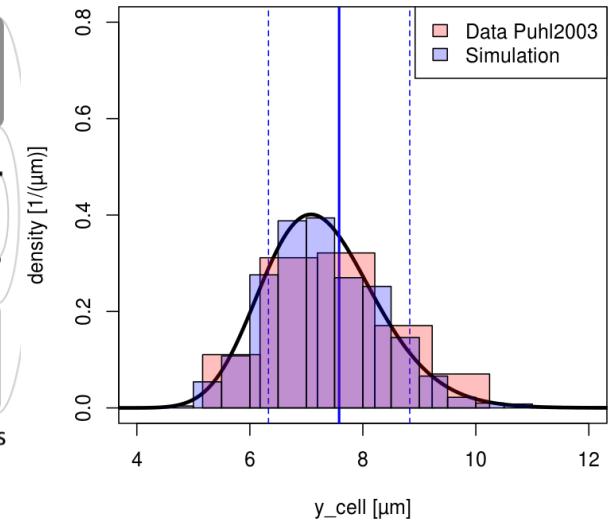
Parameter		meanlog	stdlog	mean μ (reported)	standard deviation (reported)	SD	references
Sinusoidal length	L_{\sin}	6.184	0.2462	500 μm	125 μm		based on distance between central veins 809 \pm 199 μm (SD, n=79, young rat, SEM of corrosion cast) (Warren, et al., 2008) scaled to human sinusoidal length
Sinusoidal radius	y_{\sin}	1.465 (± 0.010)	0.1017 (± 0.0073)	4.4 μm	0.45 μm		Based on distribution of sinusoidal diameter 8.8 \pm 0.9 μm (SD, n=440 in N=11 human, OPS) (Puhl, et al., 2003)
Width of Disse space	y_{dis}	0.1296	0.3246	1.2 μm	0.4 μm		0.4-1.5 μm (human, SEM, estimated from imaged) (Muto, et al., 1977) 0.5-1.2 μm (human, SEM, estimated from image) (Burwen, et al., 1982)
Hepatocyte sheet thickness	y_{cell}	1.977 (± 0.014)	0.1390 (± 0.0099)	7.58 μm	1.25 μm		7.58 μm Calculated from functional sinusoidal density FSD FSD 391 \pm 30 [1/cm] (SD, n=88, human, OPS) (Puhl, et al., 2003)

Tissue-scale Sinusoidal Unit

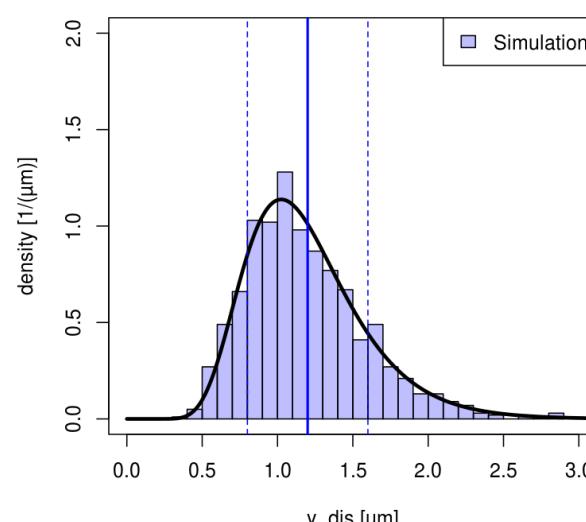
RBC velocity distribution



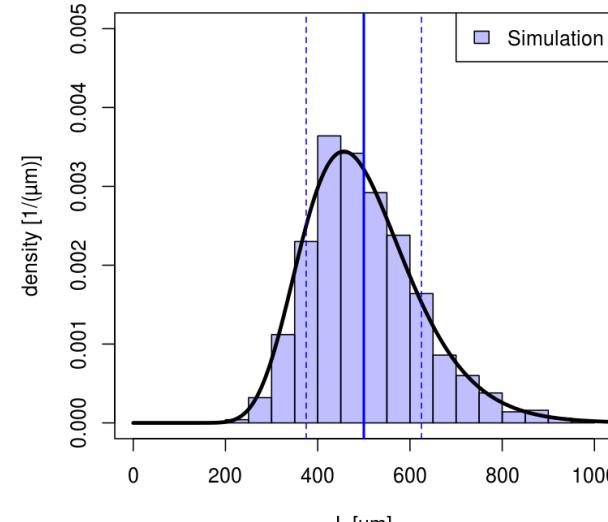
y_{cell} distribution



Width space of Disse



Sinusoidal length



Sinusoidal radius

