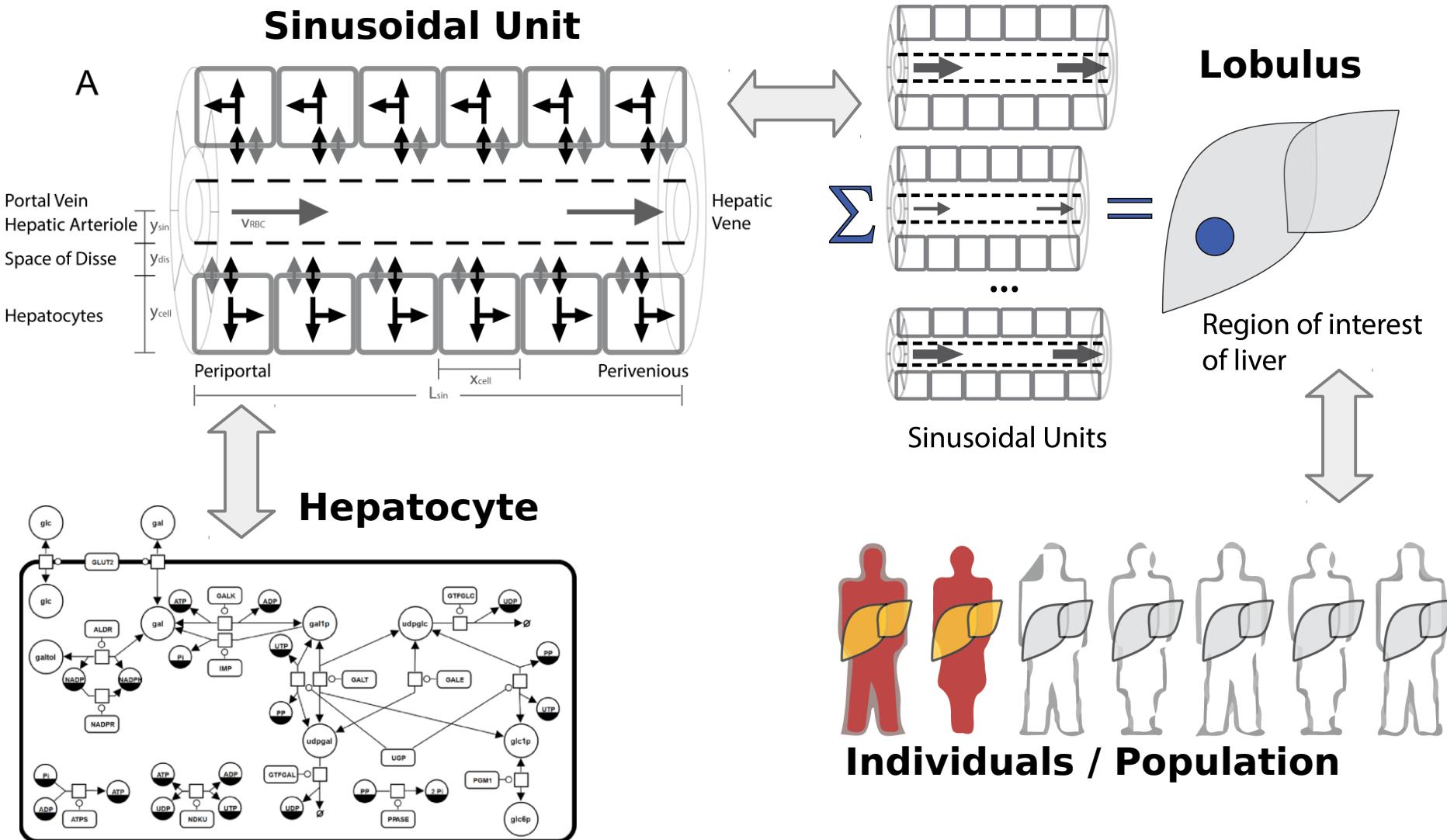
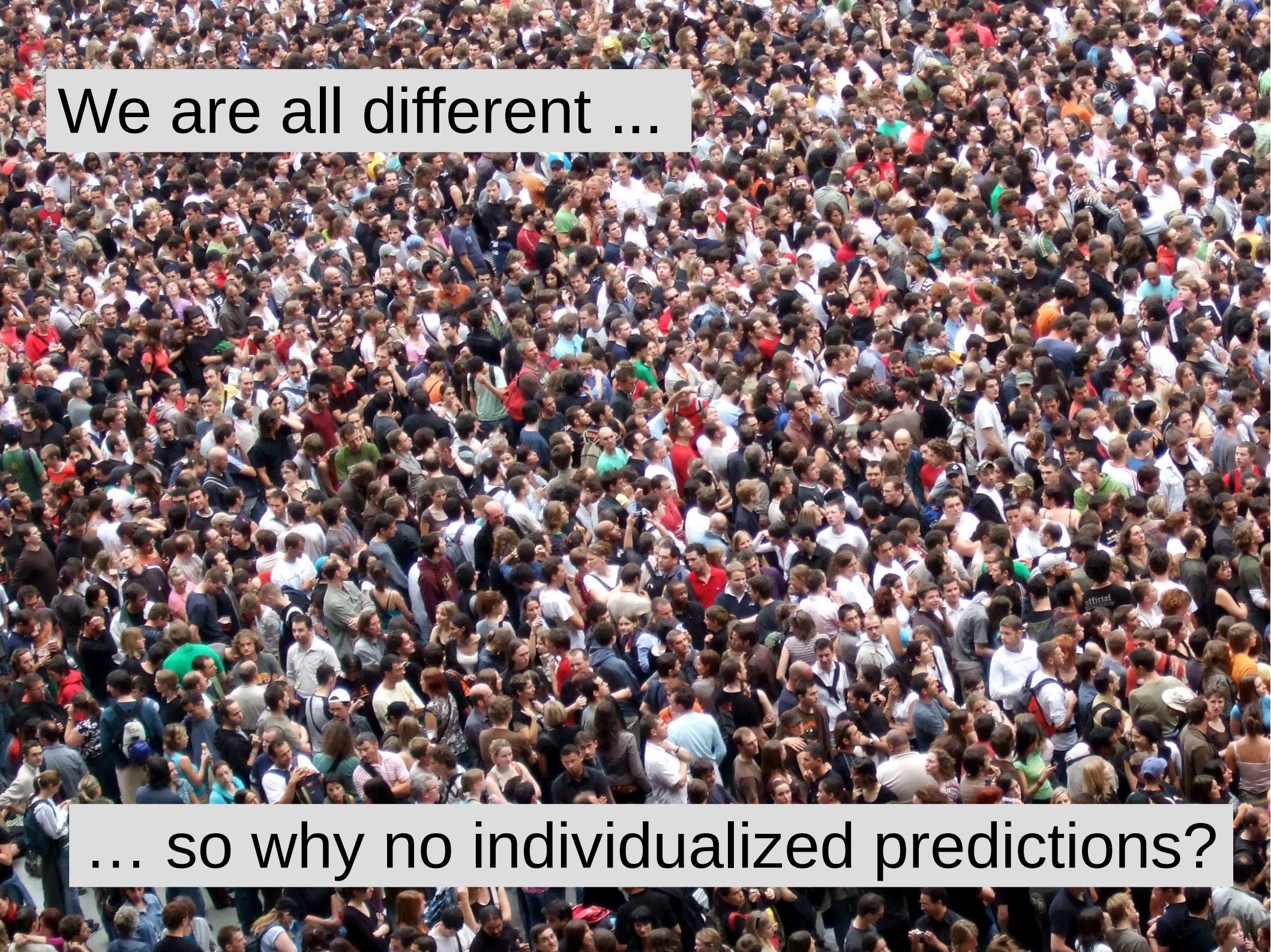


Population Variability of Liver Function

König M. & Holzhütter HG.



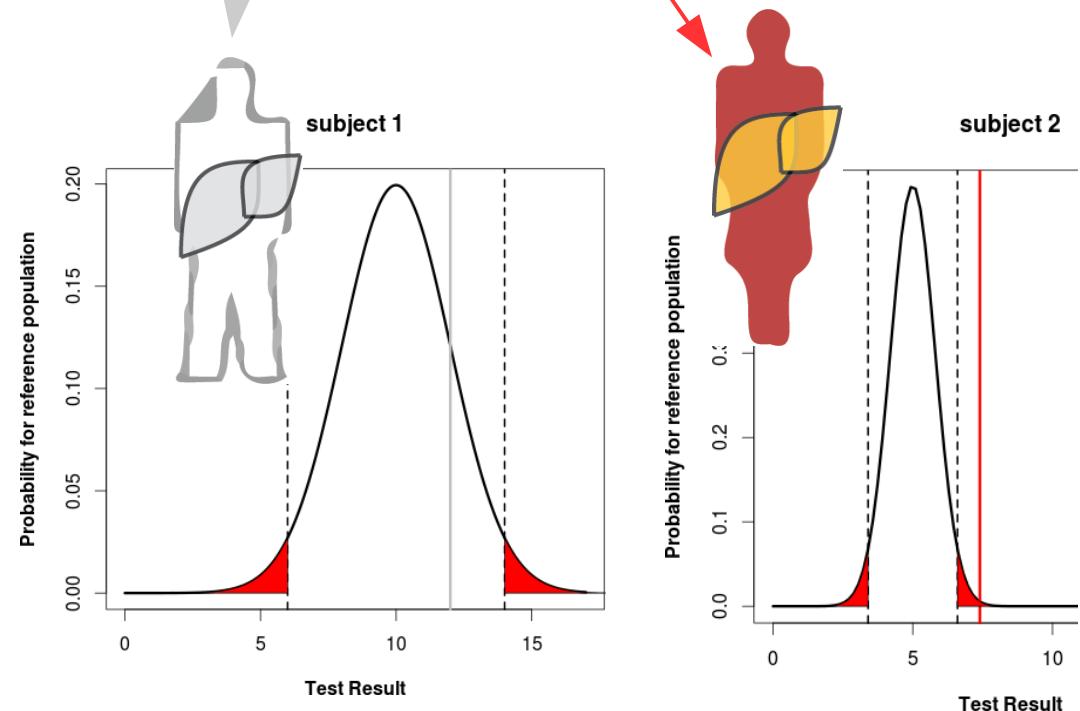
A high-angle photograph of a massive crowd of people filling the frame. The individuals are packed closely together, creating a sea of humanity. The crowd is diverse in age and appearance, with many young adults visible. They are dressed in a variety of casual clothing, including t-shirts, hoodies, and jackets. Some people are looking upwards or towards the camera, while others are looking down or to the sides. The overall impression is one of a large-scale public gathering or event.

We are all different ...

... so why no individualized predictions?

Individualized Liver Function Tests

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



Variability & Sensitivity of Parameters

- **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.**

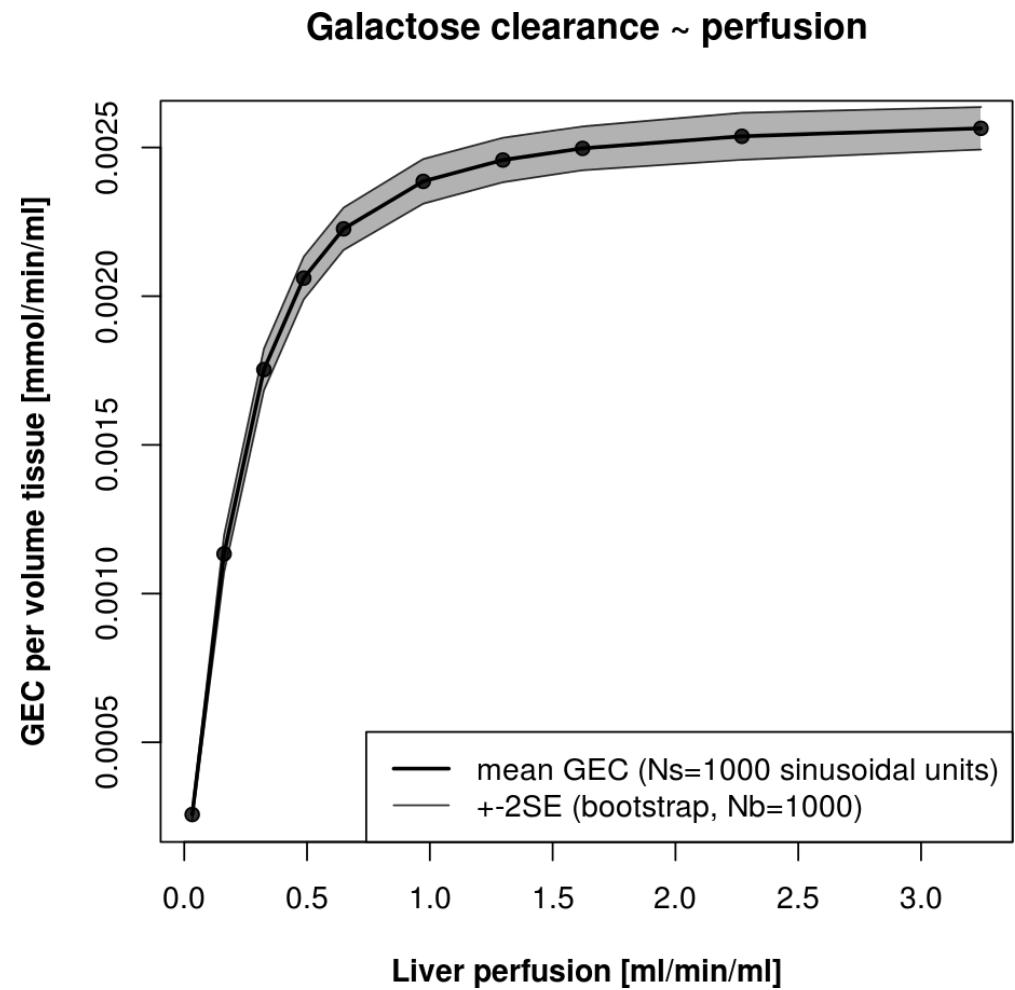
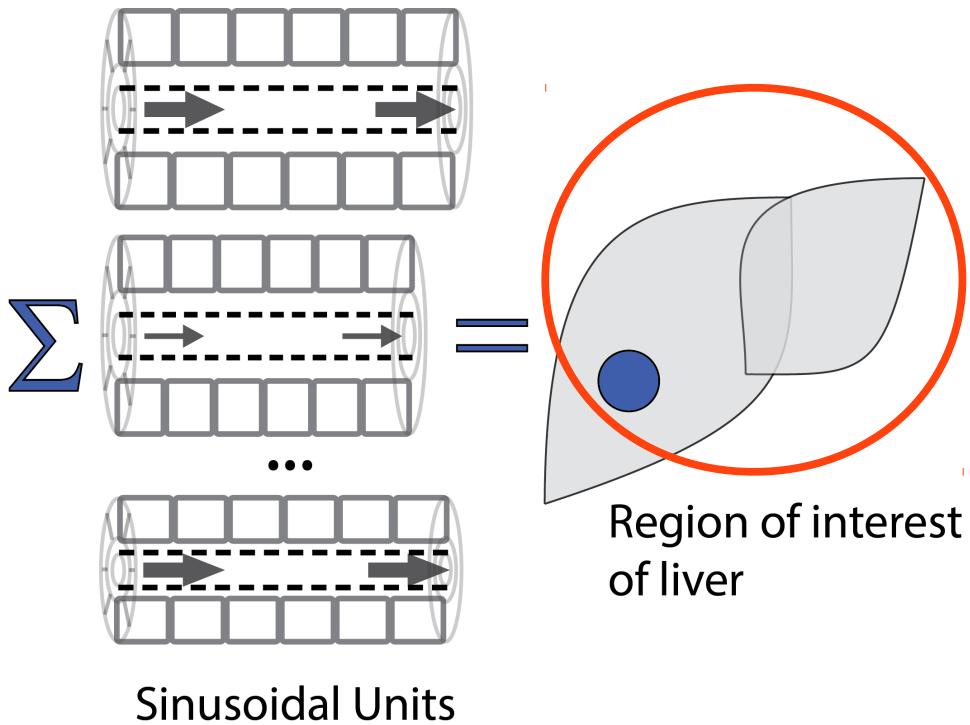


- **Sensitivity**

- Sensitivity analysis provides a quantitative evaluation of **how parameters influence the simulation prediction.**

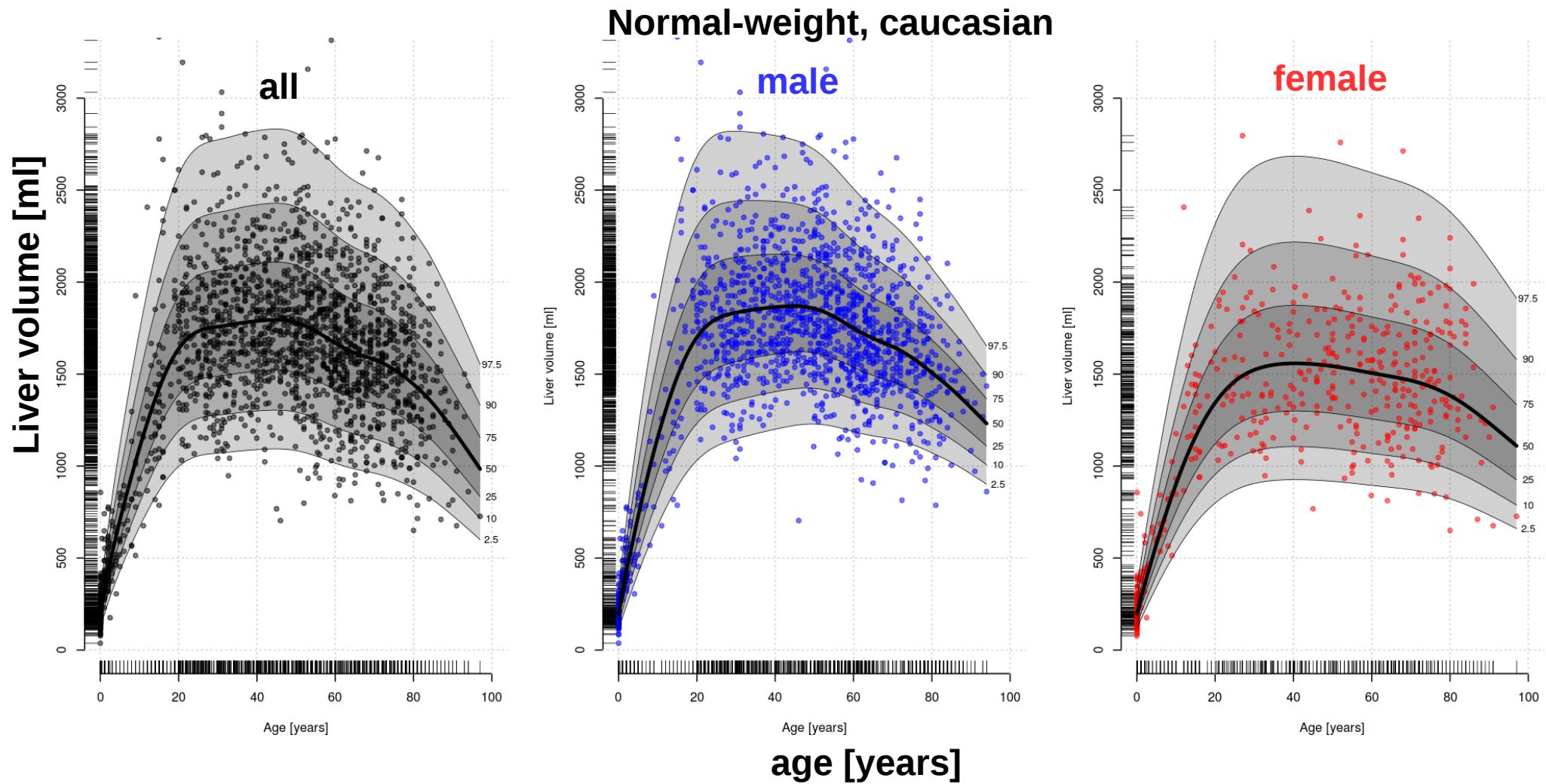
Liver Function Test - GEC

- Evaluation of liver function via galactose elimination capacity (GEC)
- Tissue-Model predicts the galactose clearance per tissue volume for given perfusion in the region (**regional GEC**)
- Scale with individual **liver volume and perfusion** (**total GEC**)
- Evaluation of individual & population variability in GEC



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

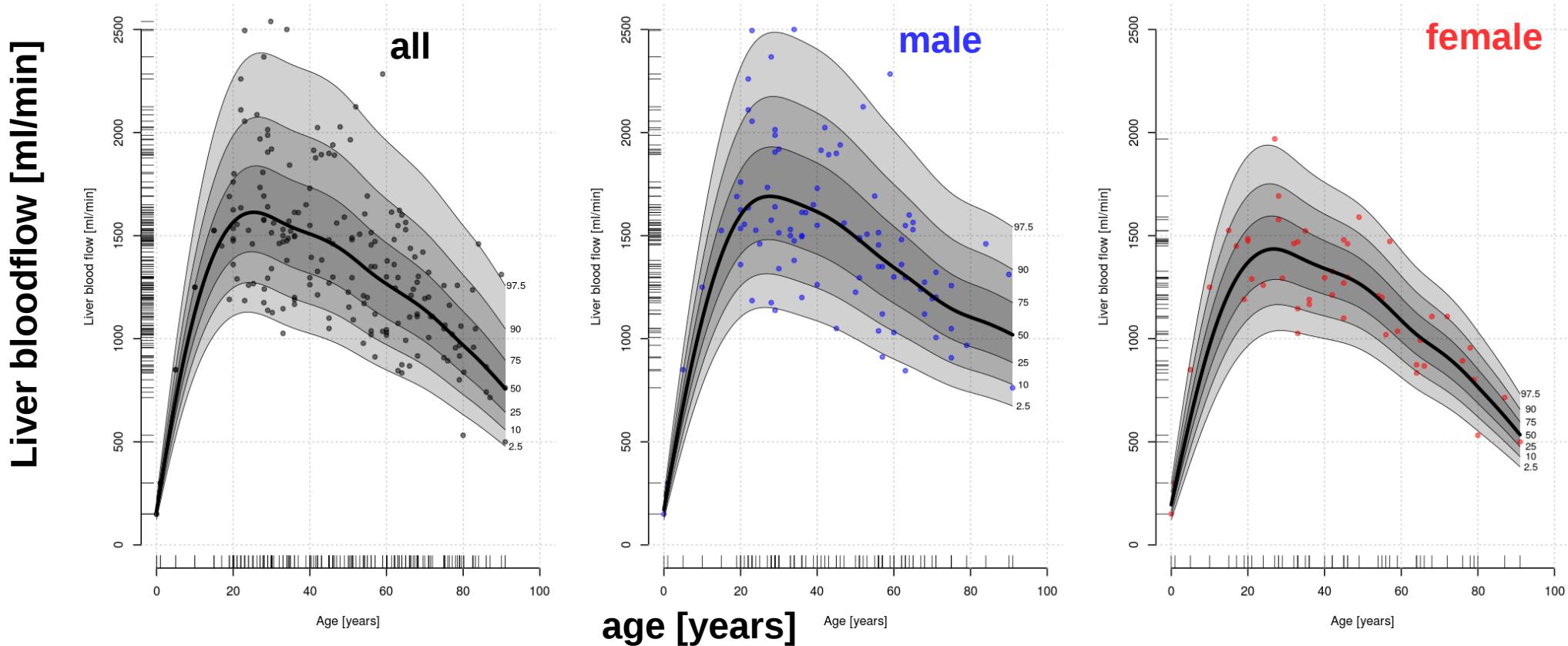
- Density estimation of liver **volume** ~ **age** via generalized additive models (GAM) combined with centile estimation
- Analog: volLiver ~ bodyweight, volLiver~height, volLiver~BSA, ...
- Combined dataset from multiple sources & experimental methods



Age & gender dependence blood flow

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

Normal-weight, caucasian

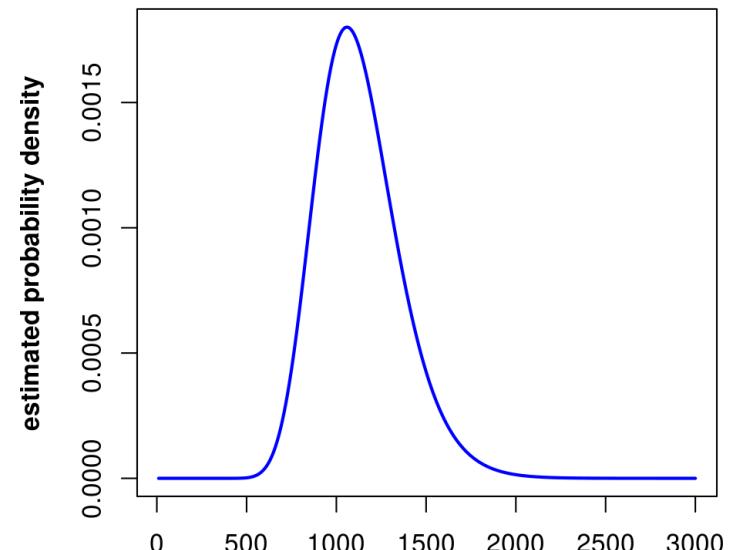
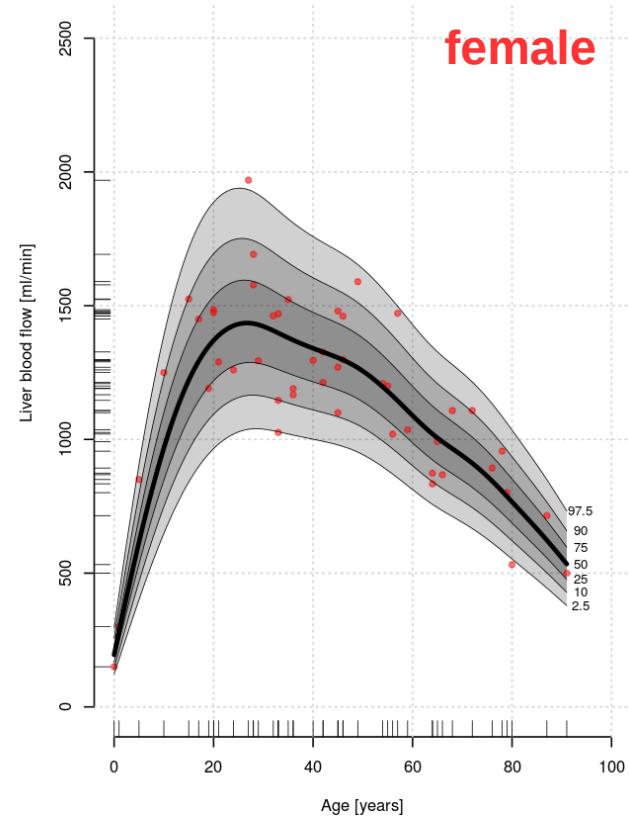
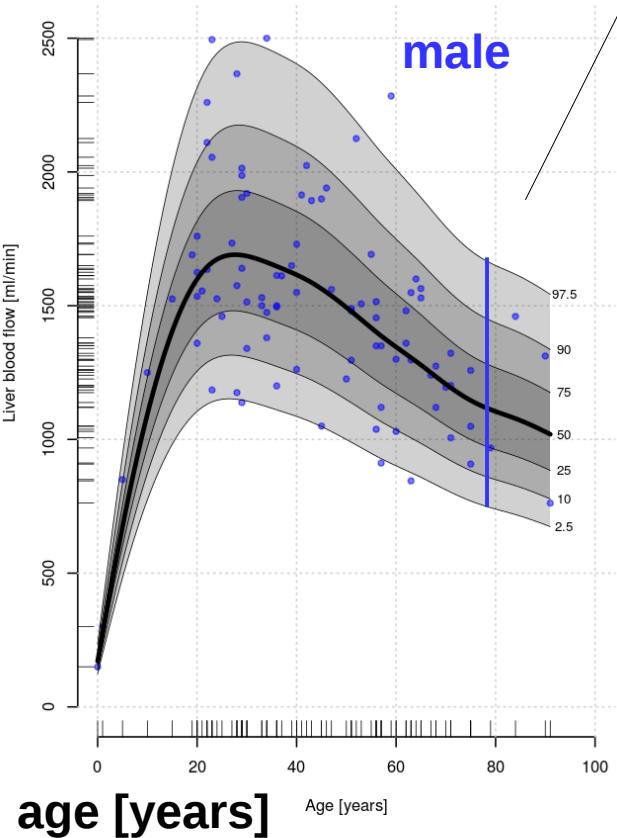
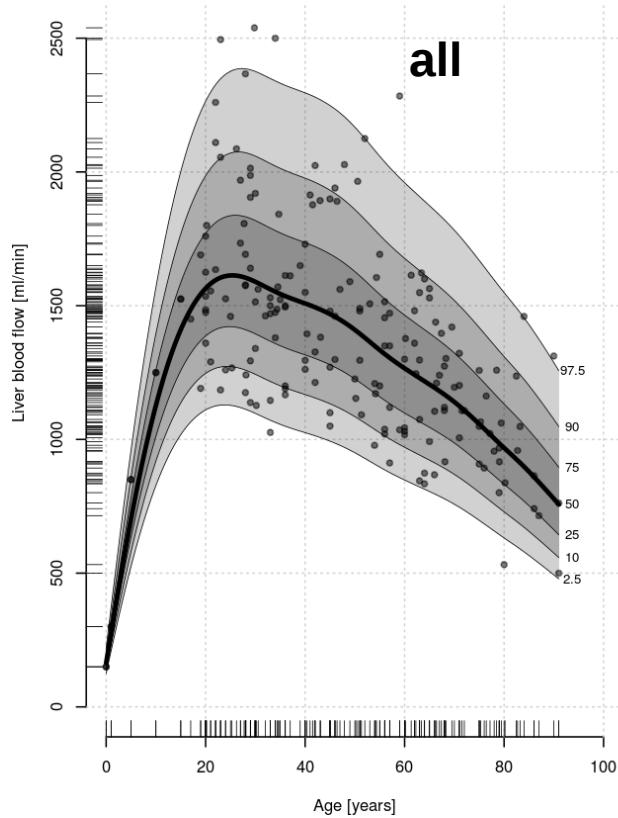


Models can be used for prediction of individual parameter distributions.

Probability distribution of liver blood flow

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

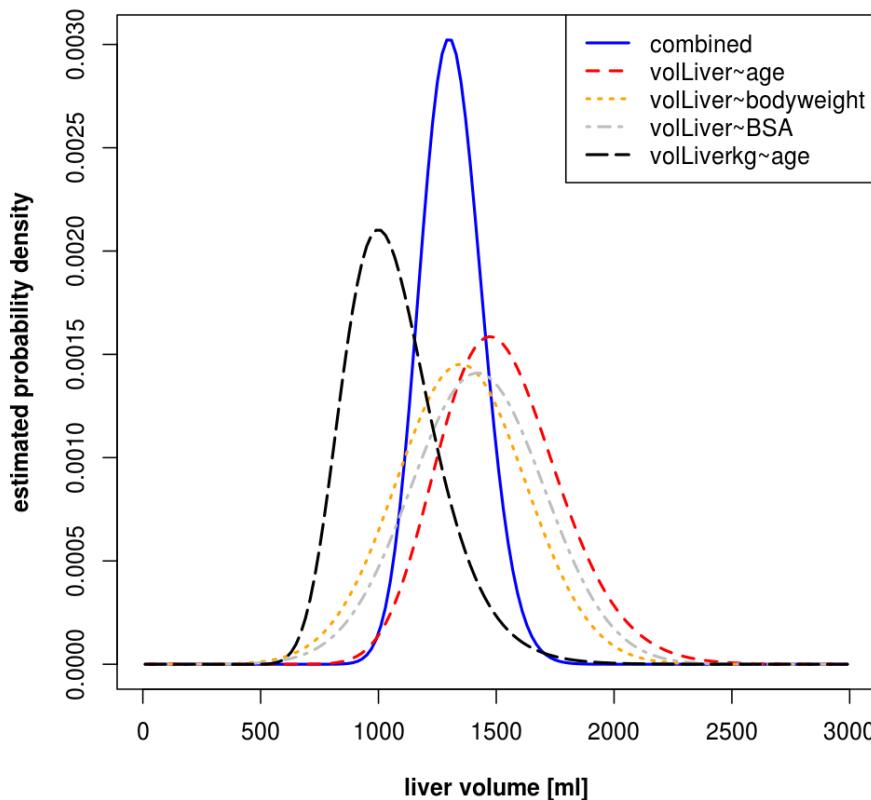
Liver bloodflow [ml/min]



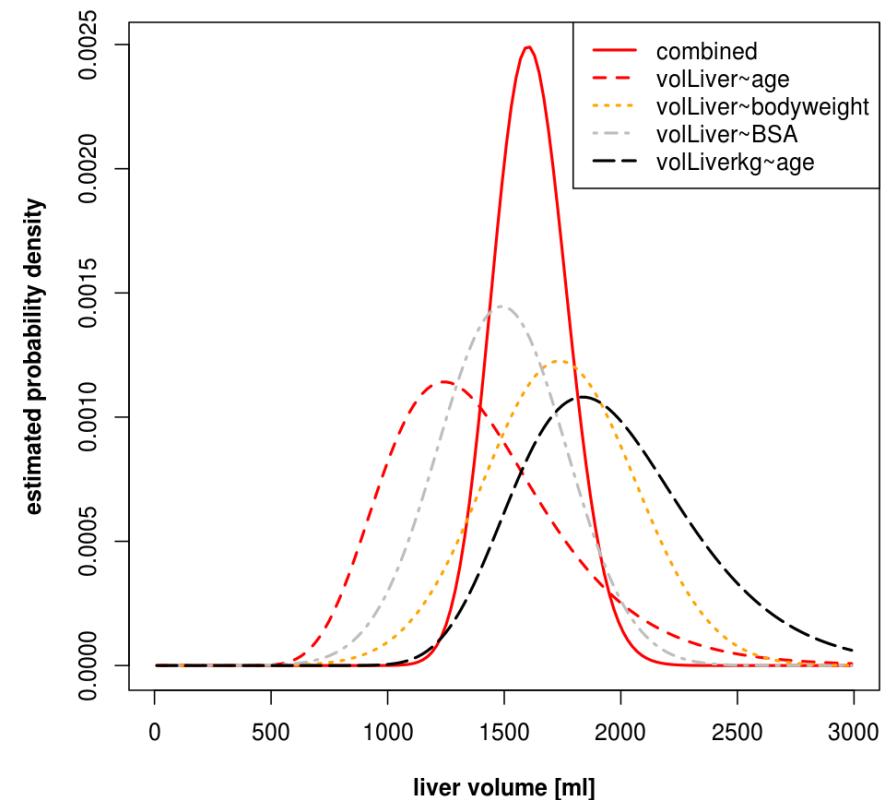
Combining the information

- Set of probability distributions for pair-wise correlations
- Important to represent the correlation structure between different pairs
- **Solution:** 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]

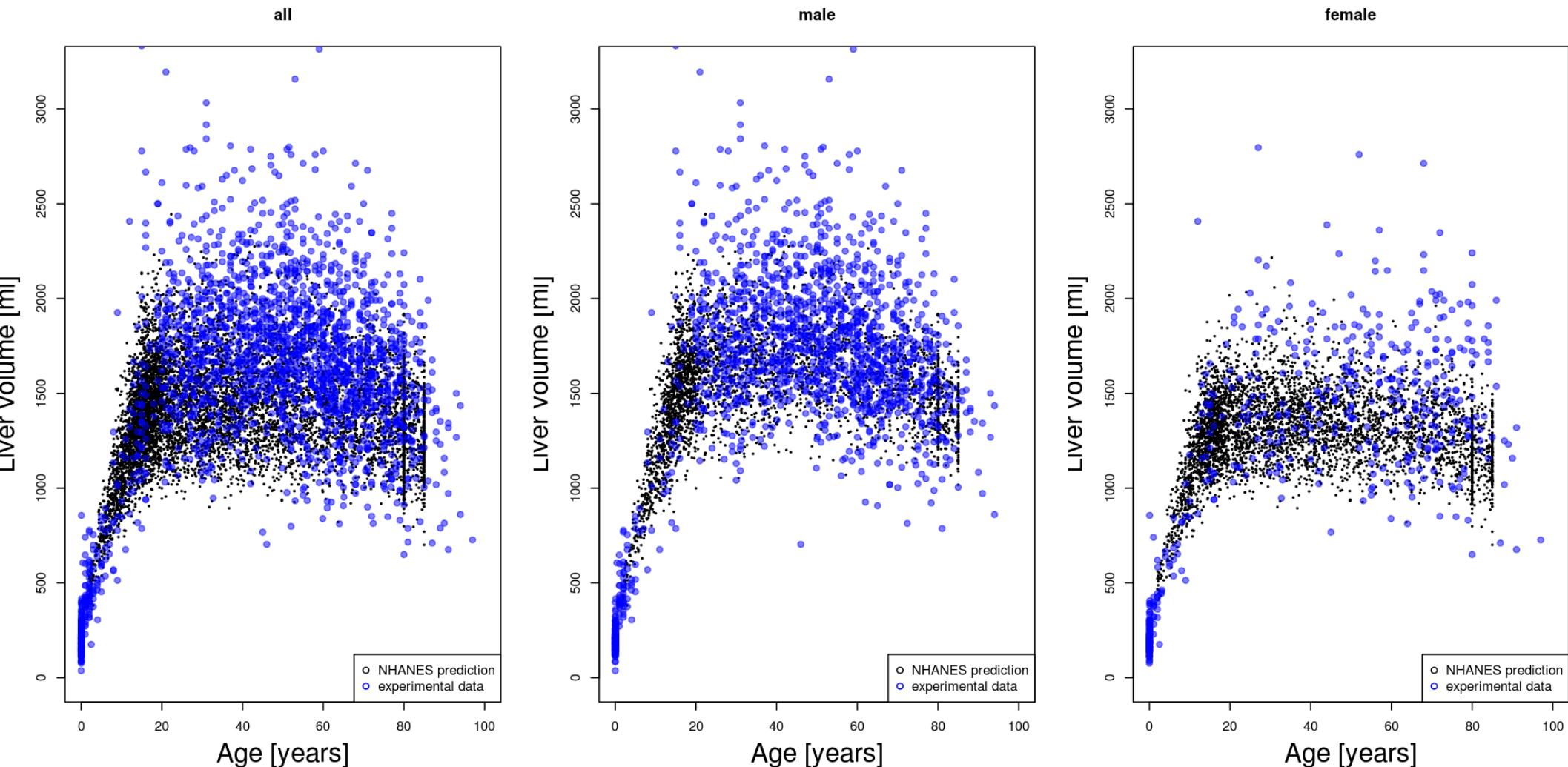


age=20 [y], sex=female, bodyweight=80 [kg], BSA=1.7 [m^2]

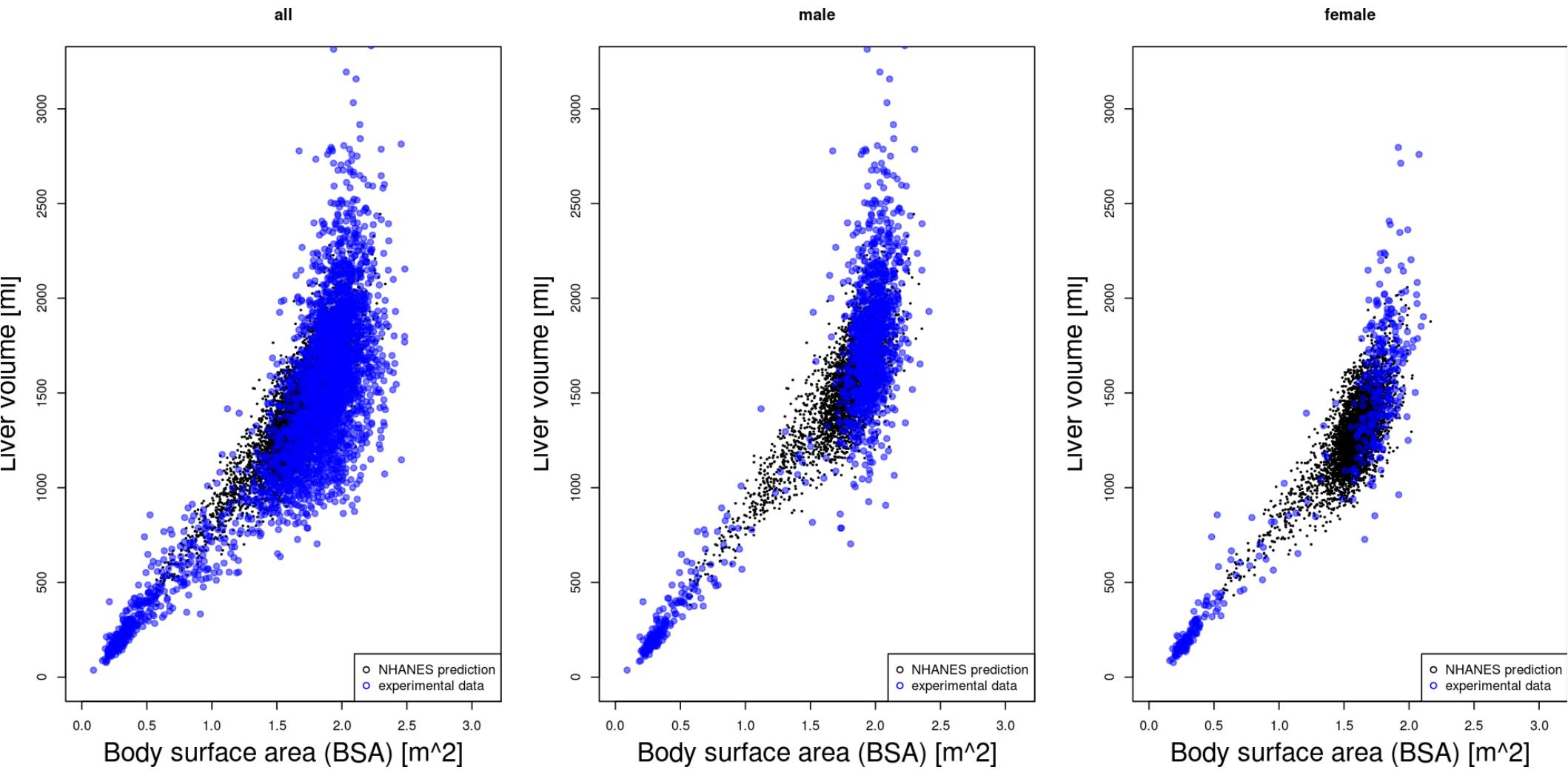


Prediction of population variability

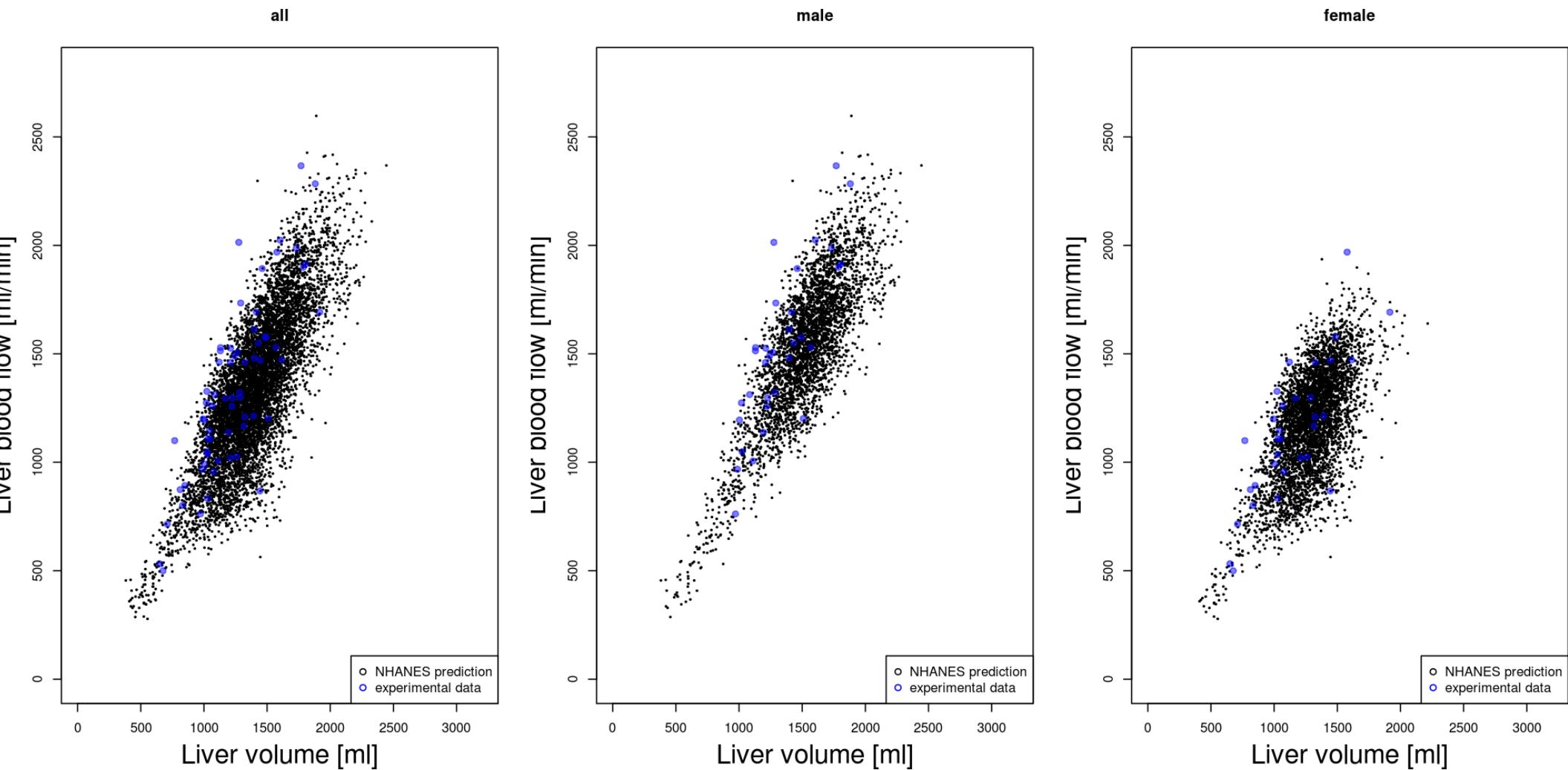
- NHANES prediction (~8000 normal-weight, caucasian)
 - Correlation structure of sex, age, bodyweight, height of US population
- **Liver volume ~ age**



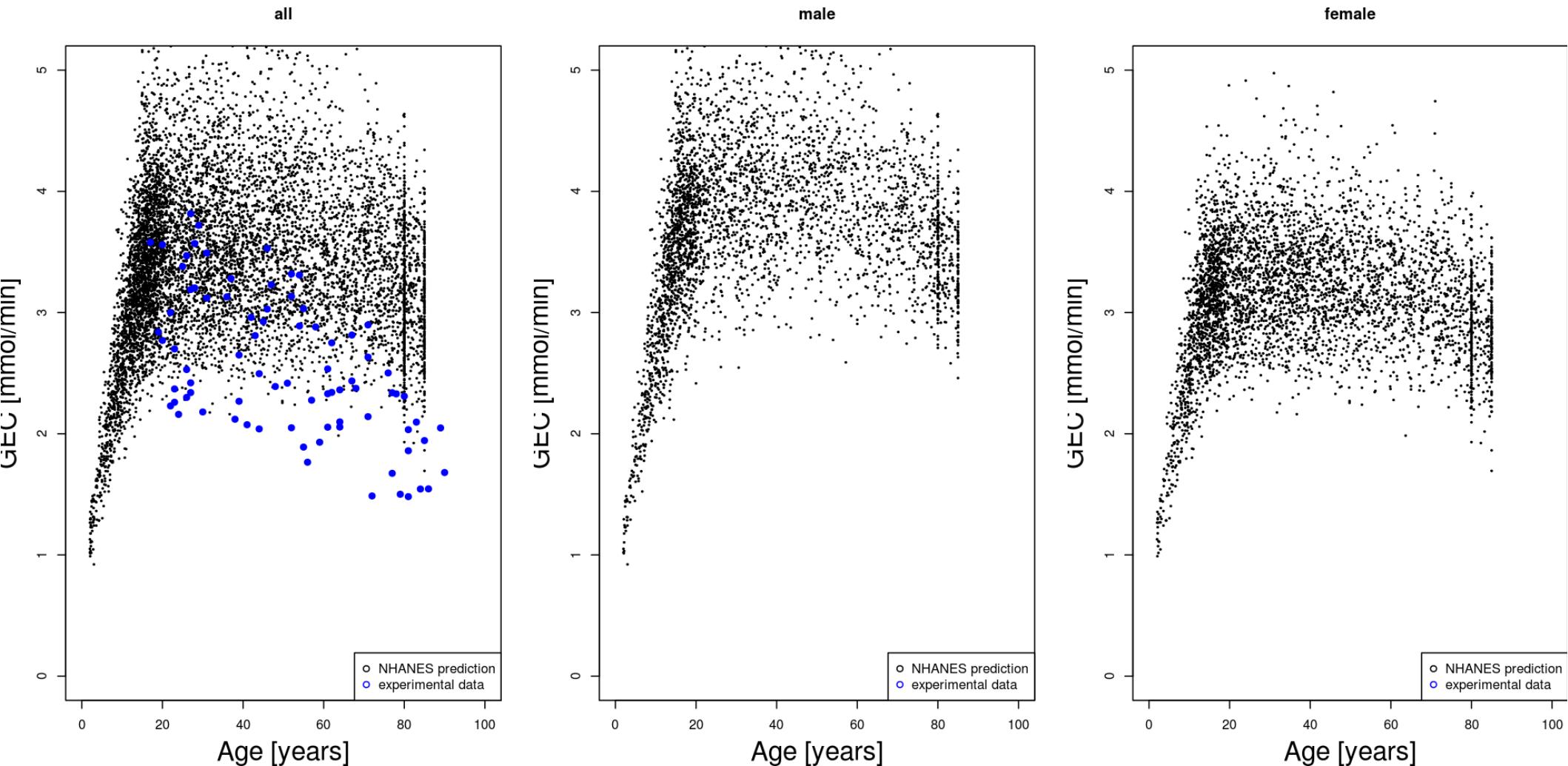
Prediction: Liver volume ~ BSA (body surface area)



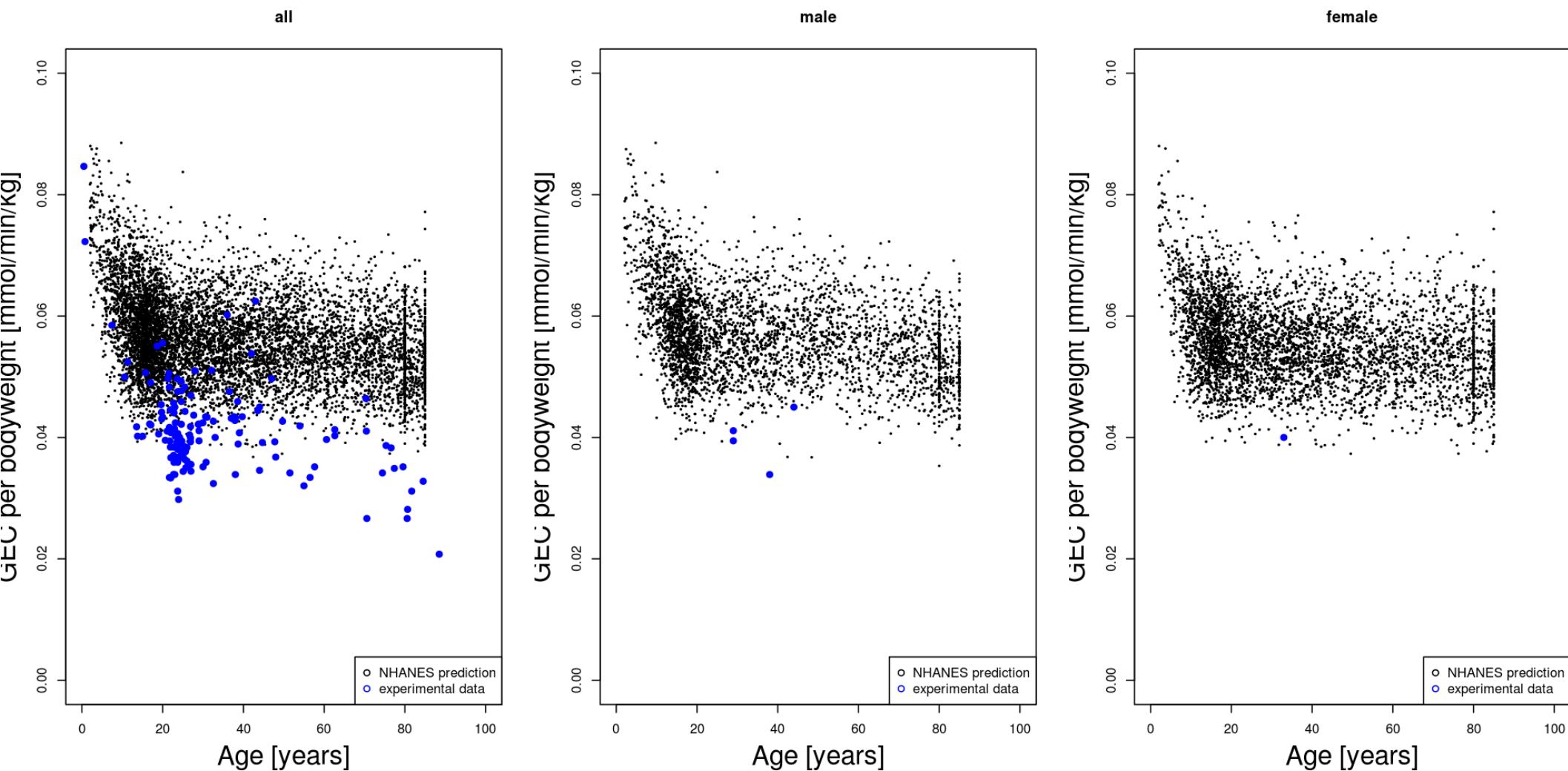
Prediction: liver blood flow ~ liver volume



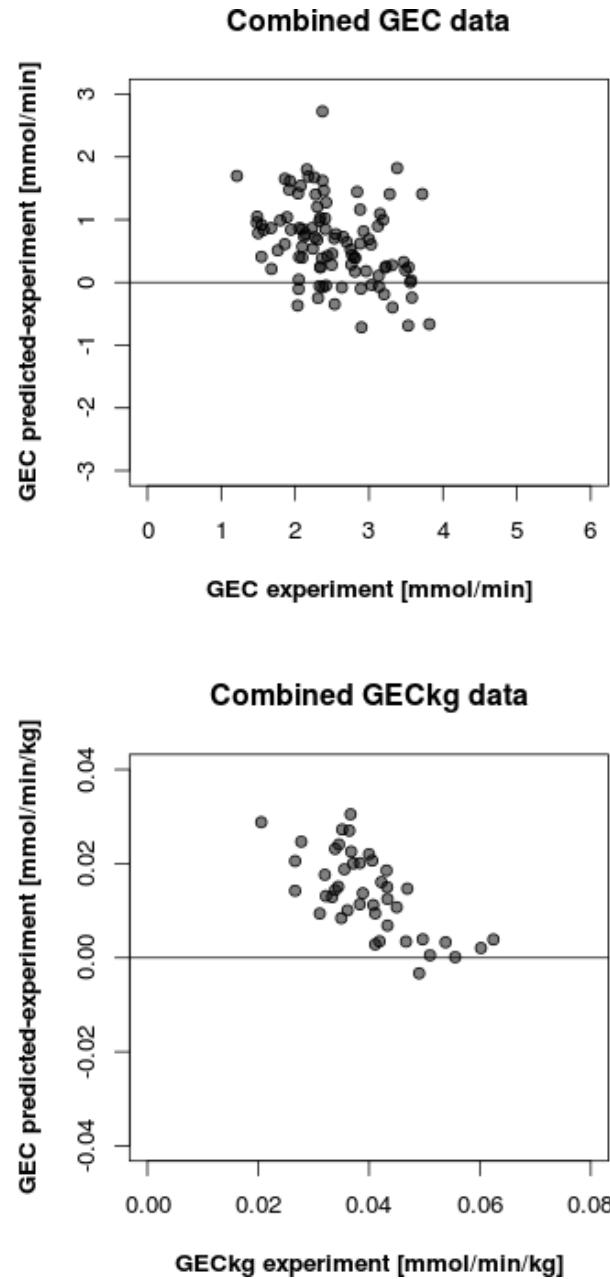
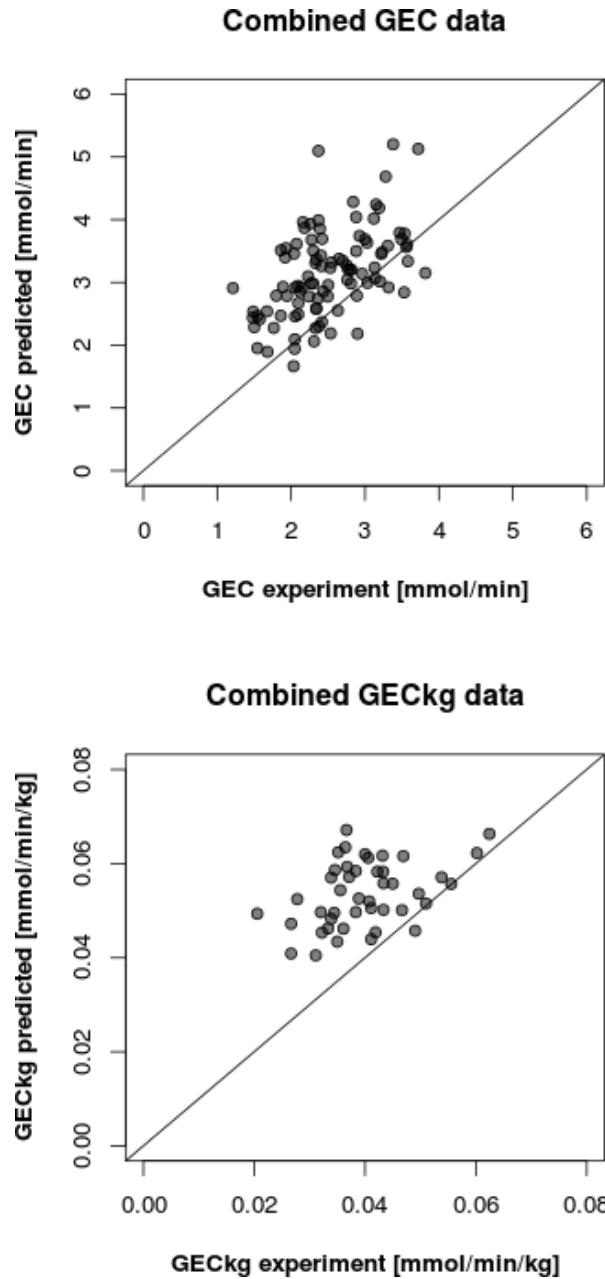
Prediction of GEC (galactose clearance capacity)



Prediction of GEC per bodyweight



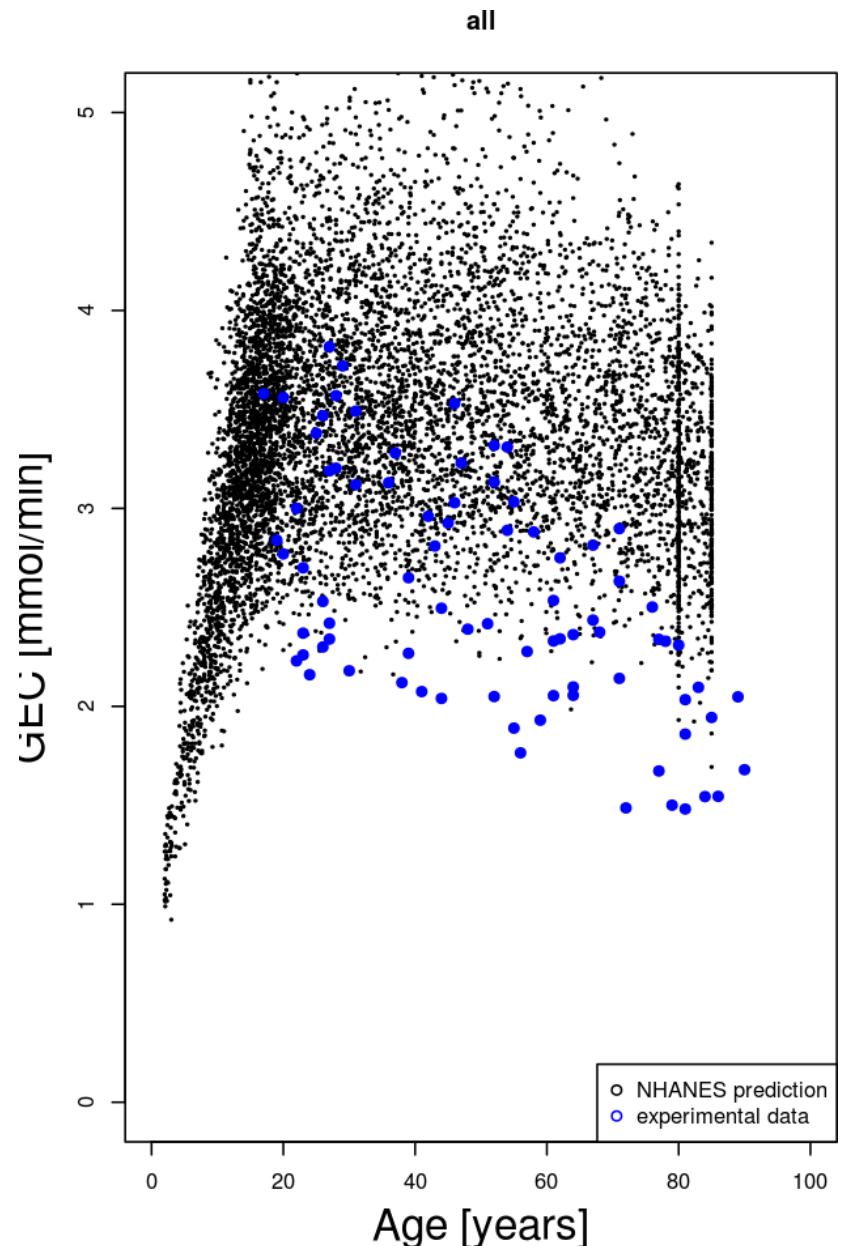
Individualized GEC prediction



- First draft predictions, no adaption of metabolic capacity
- No correction for large vessels

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



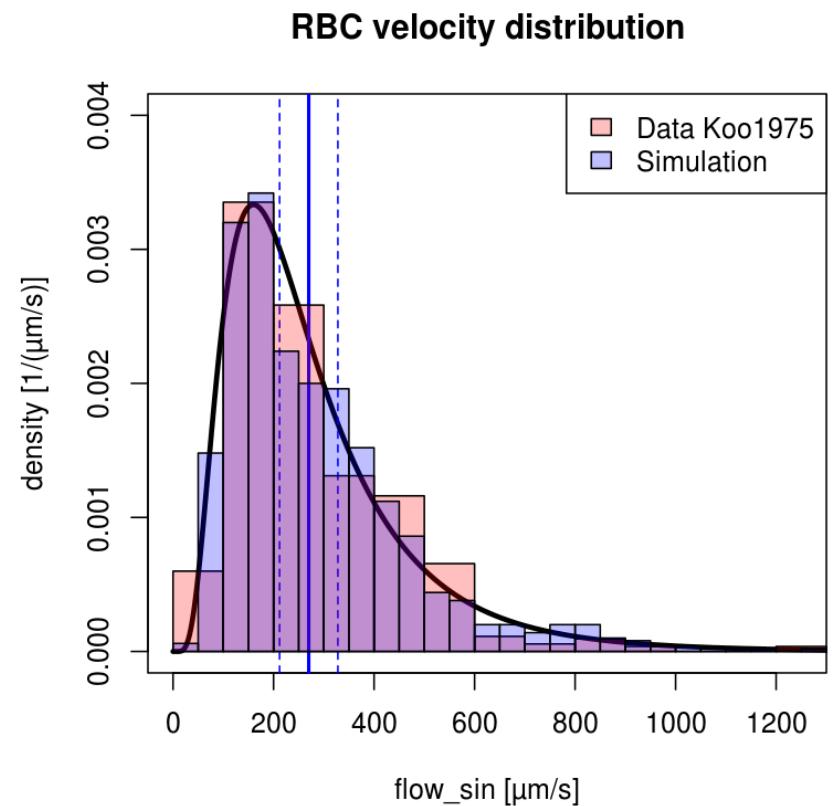
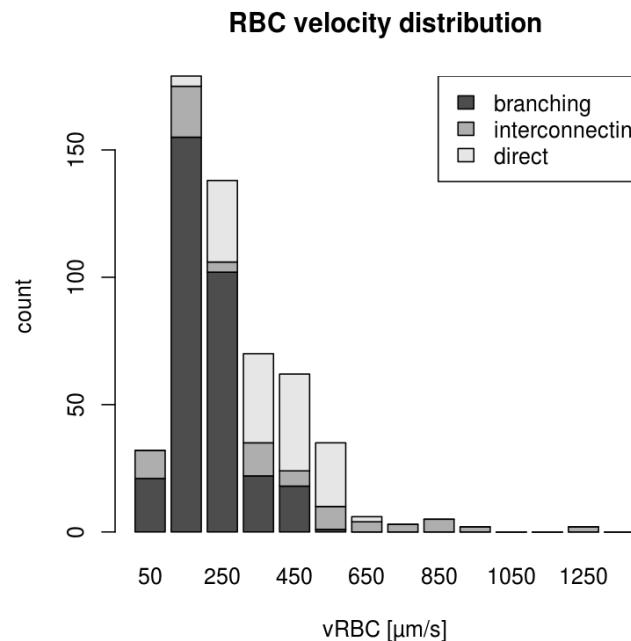
Acknowledgements

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 - Prof. Holzhütter
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 - Martin Golibiewski, Wolfgang Müller, Renate Wittig
- **Funding**
 - Charité
 - Virtual Liver Network



Fitting Distributions

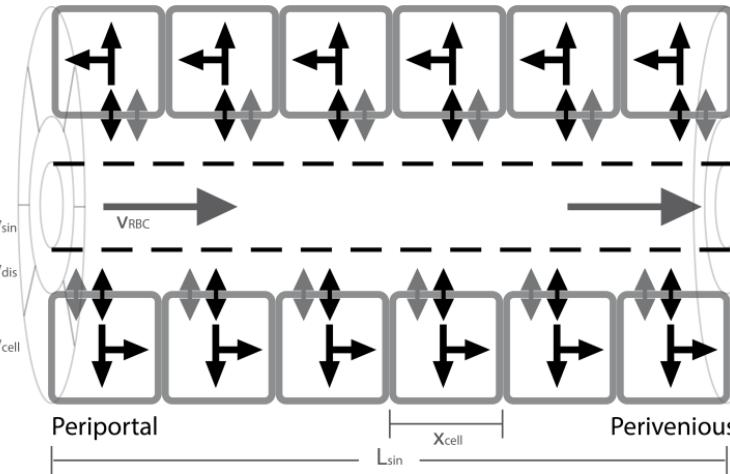
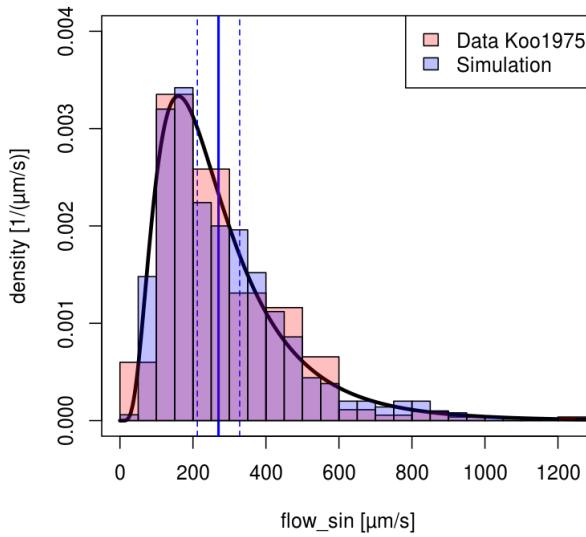
- maximum-likelihood method for univariate distributions
(log-normal)



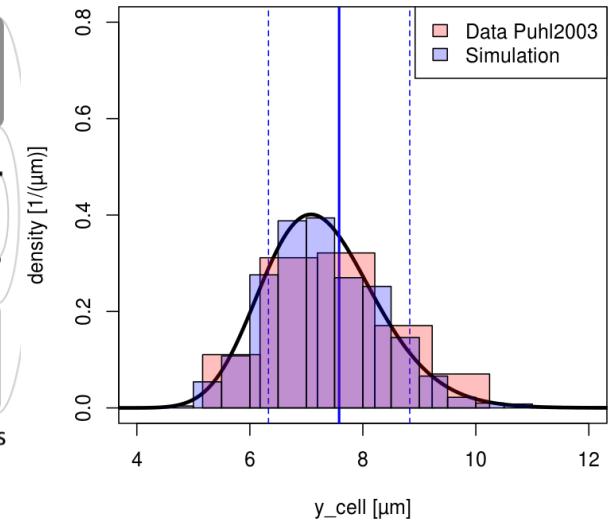
Parameter		meanlog	stdlog	mean μ (reported)	standard deviation (reported)	SD	references
Sinusoidal length	L_{\sin}	6.184	0.2462	500 μm	125 μm		based on distance between central veins 809 \pm 199 μm (SD, n=79, young rat, SEM of corrosion cast) (Warren, et al., 2008) scaled to human sinusoidal length
Sinusoidal 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

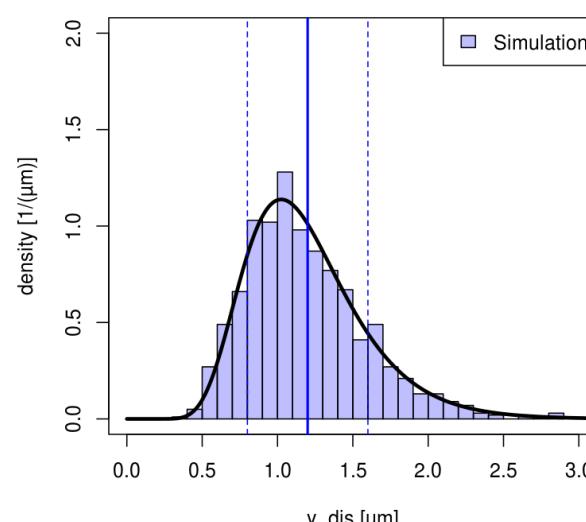
RBC velocity distribution



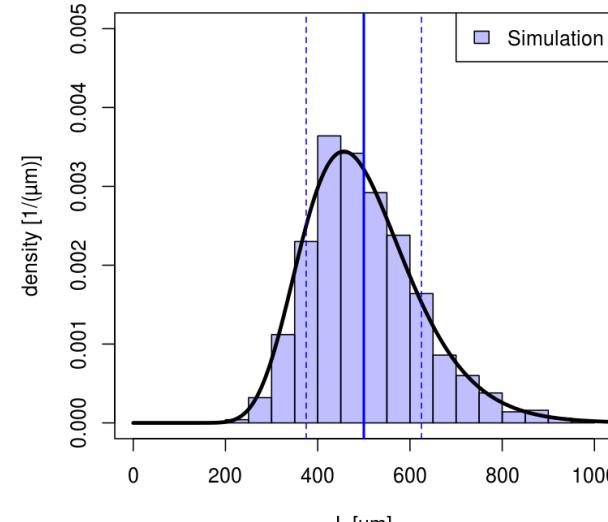
y_{cell} distribution



Width space of Disse



Sinusoidal length



Sinusoidal radius

