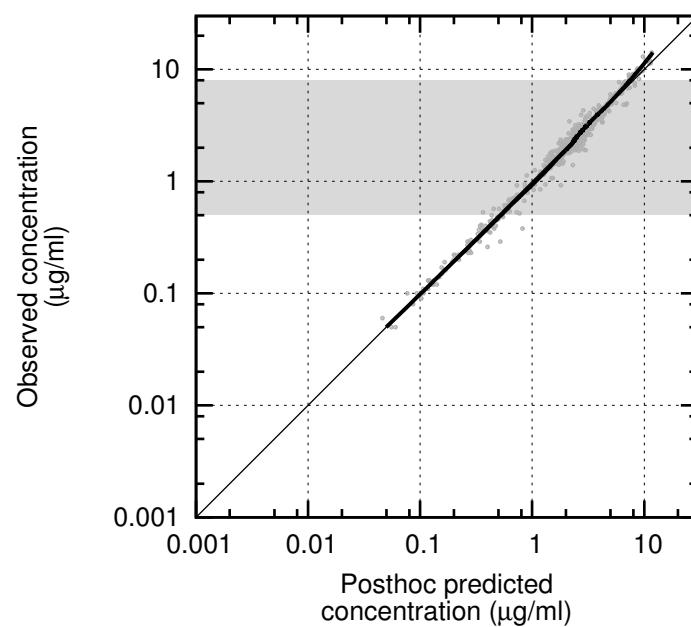
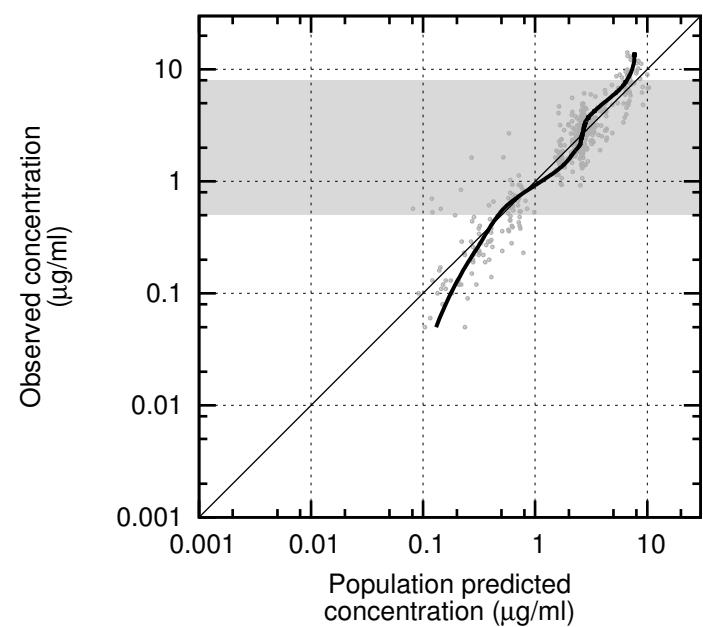
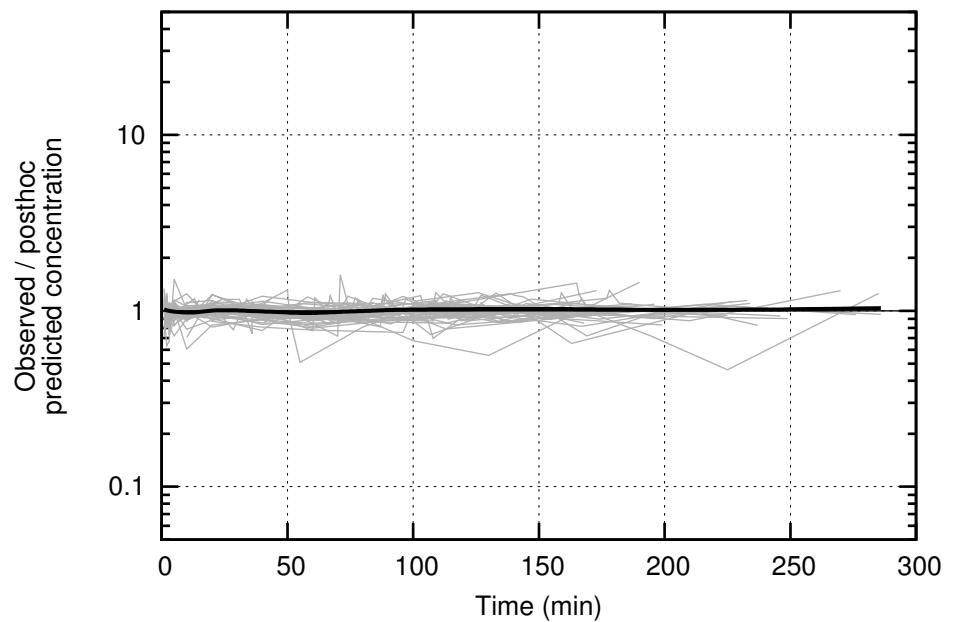
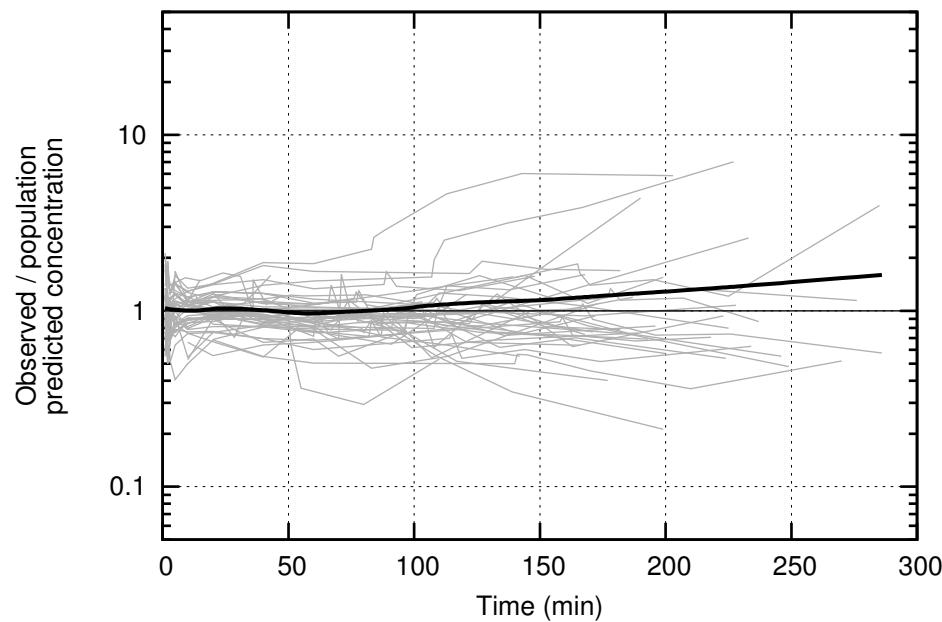
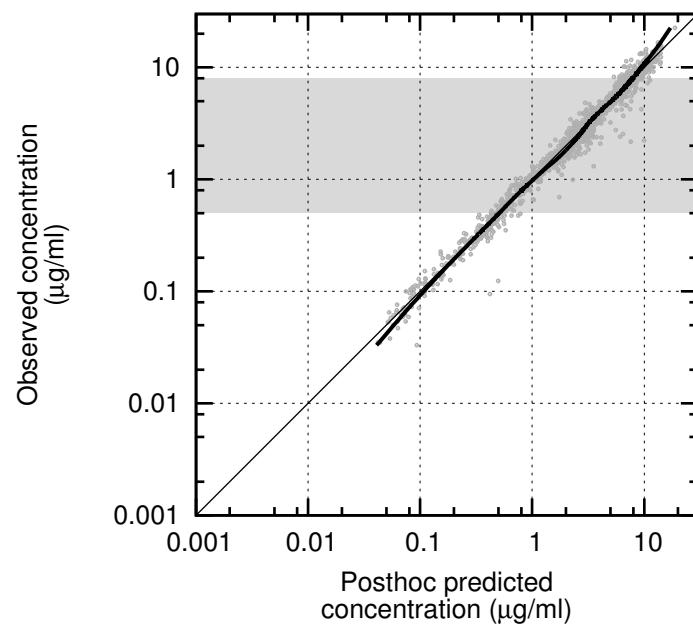
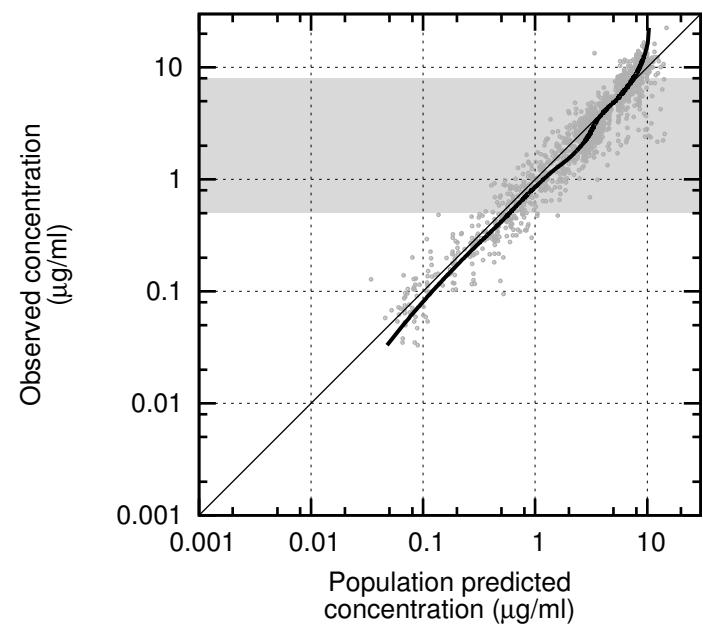
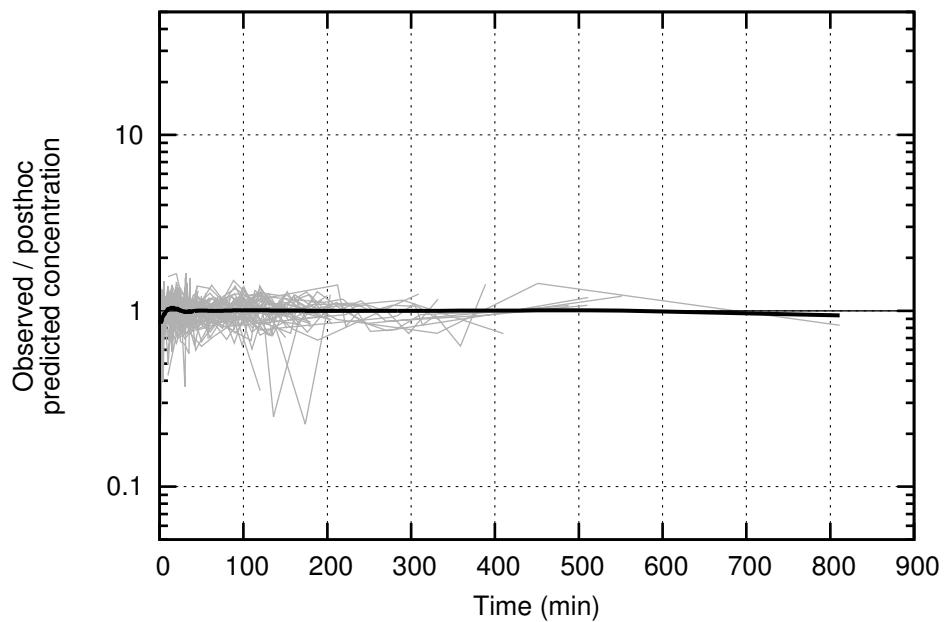
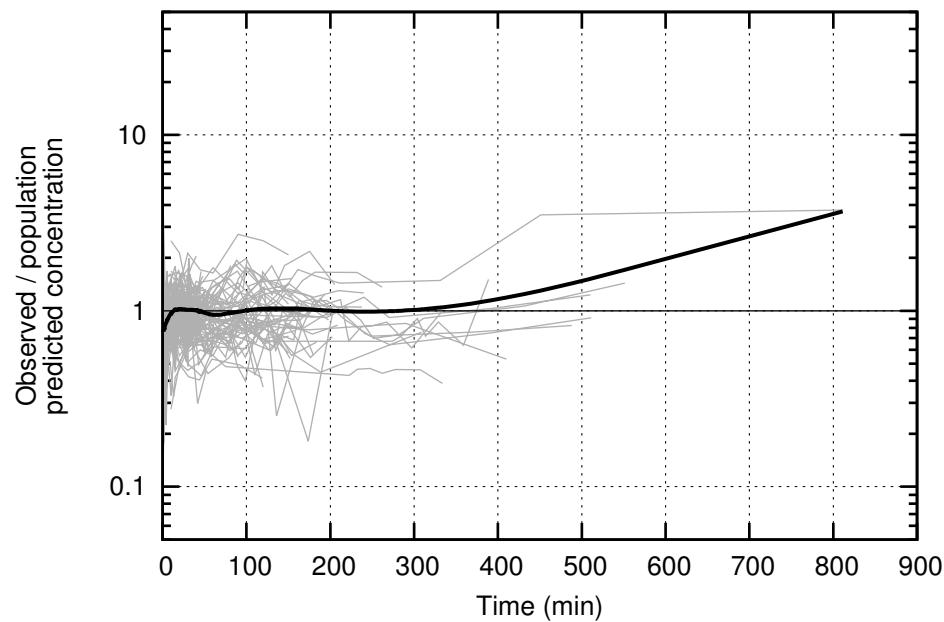


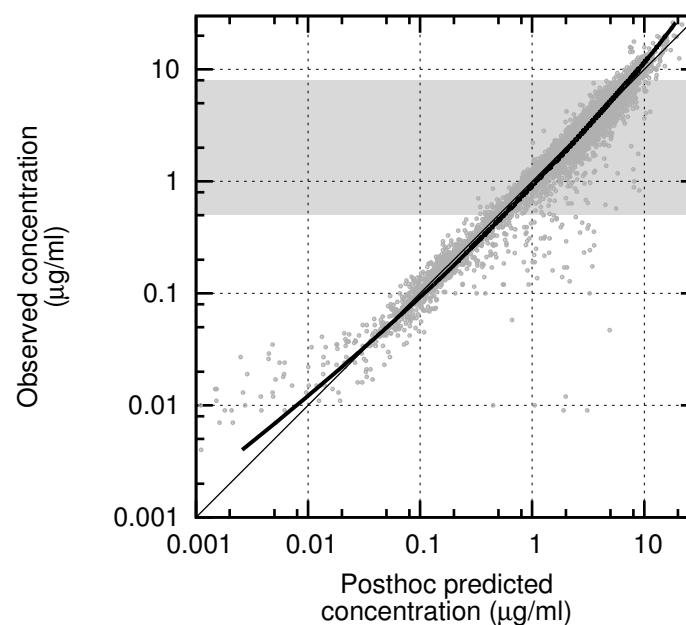
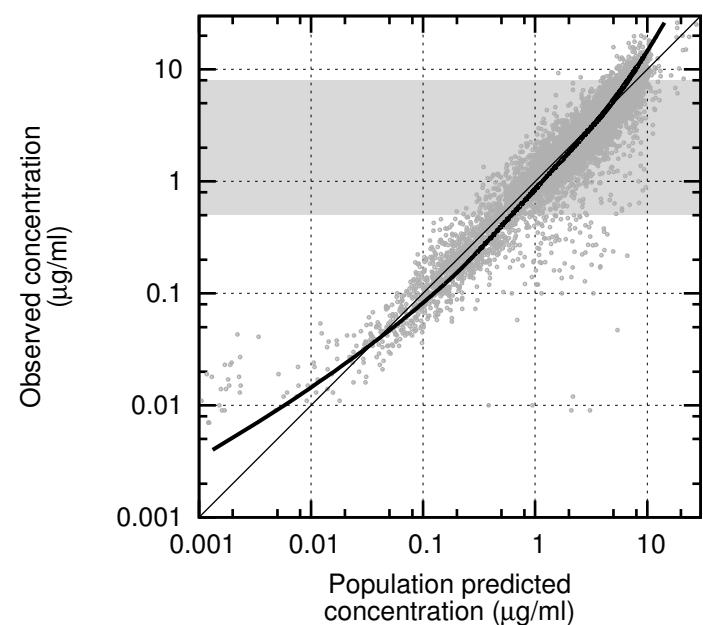
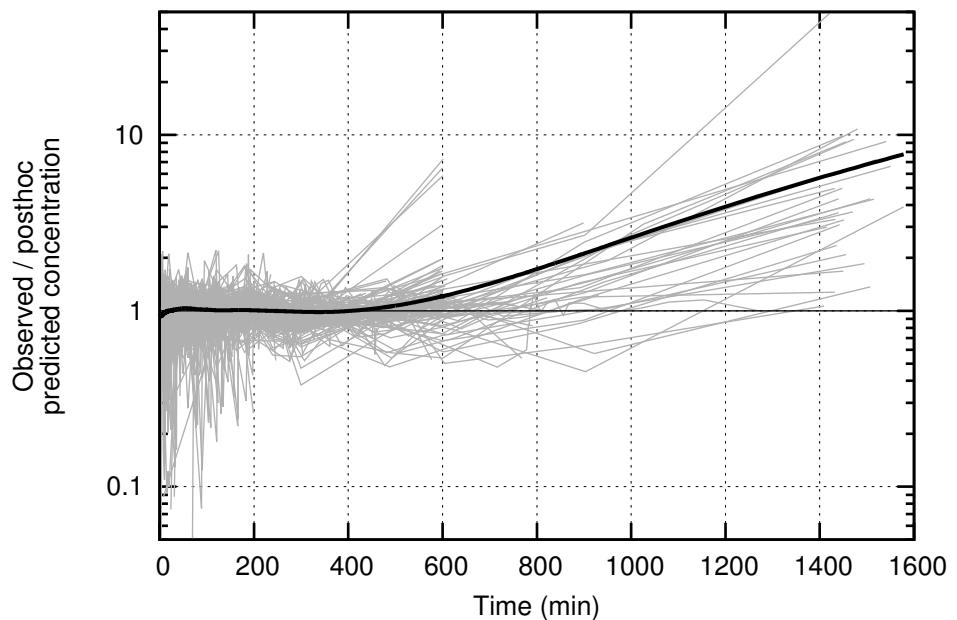
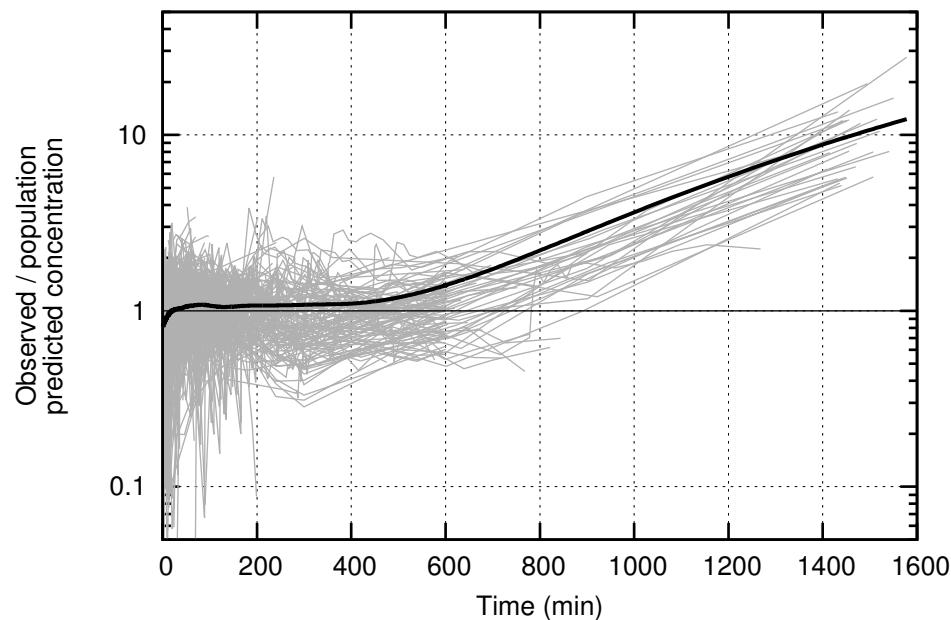
## Young children (age < 3 years)



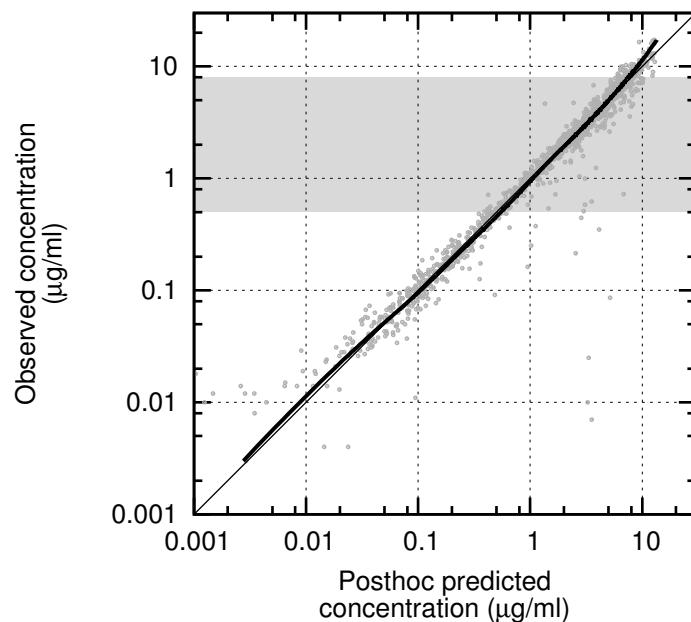
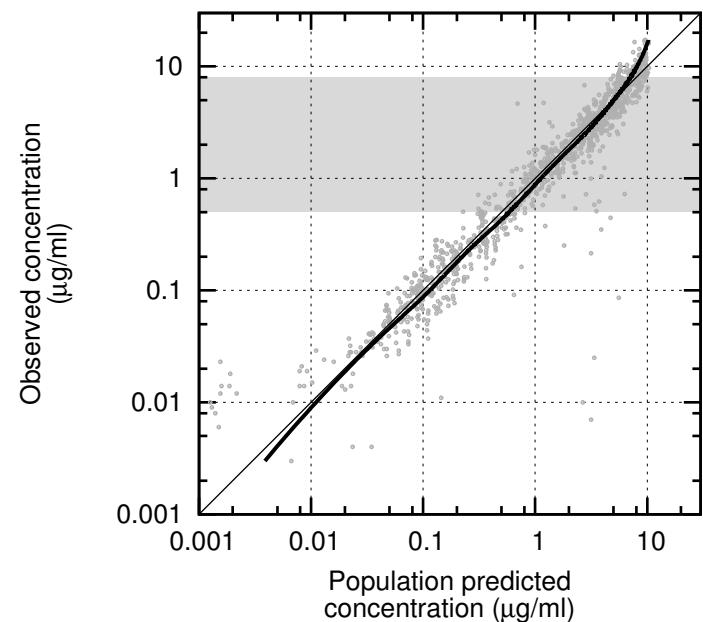
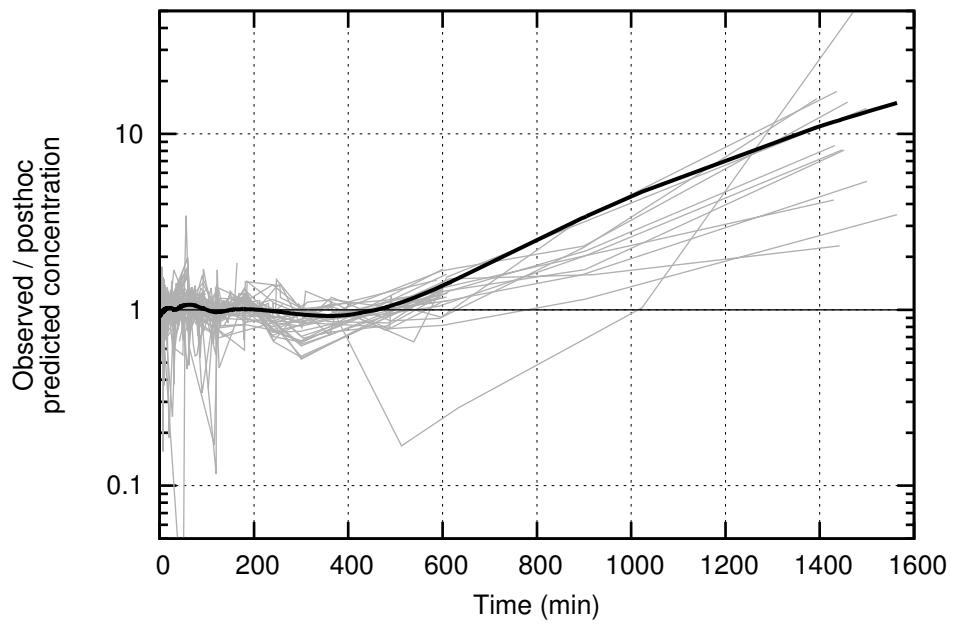
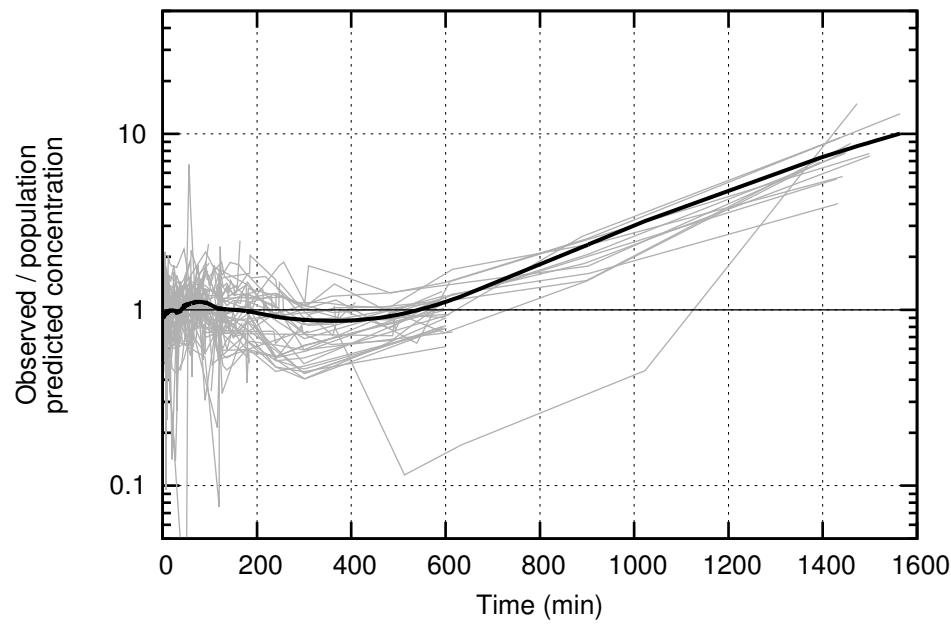
## Children (3 ≤ age < 18 years)



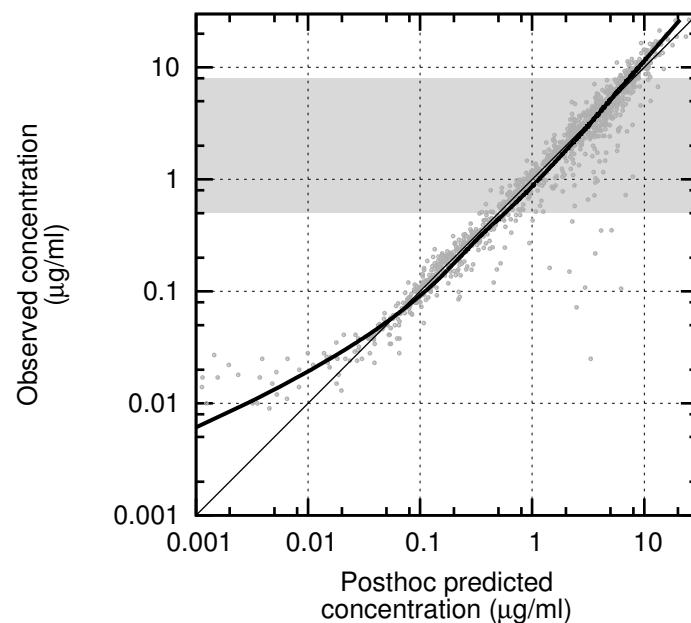
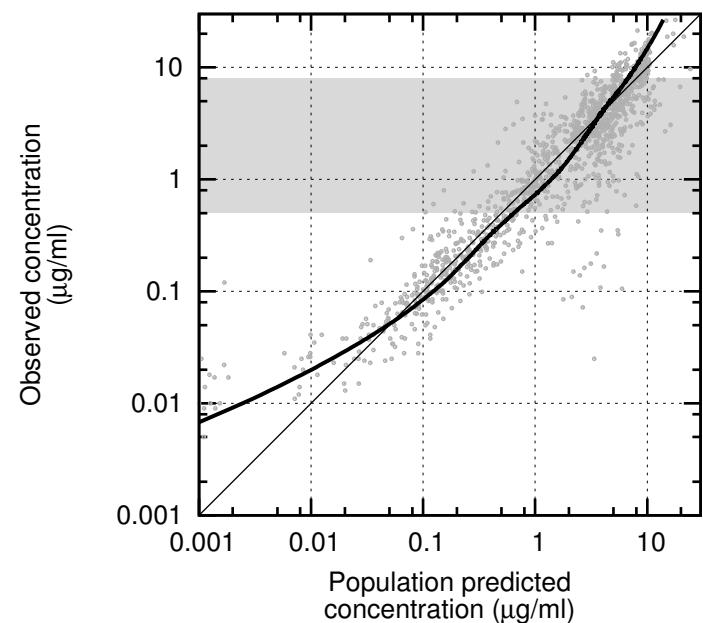
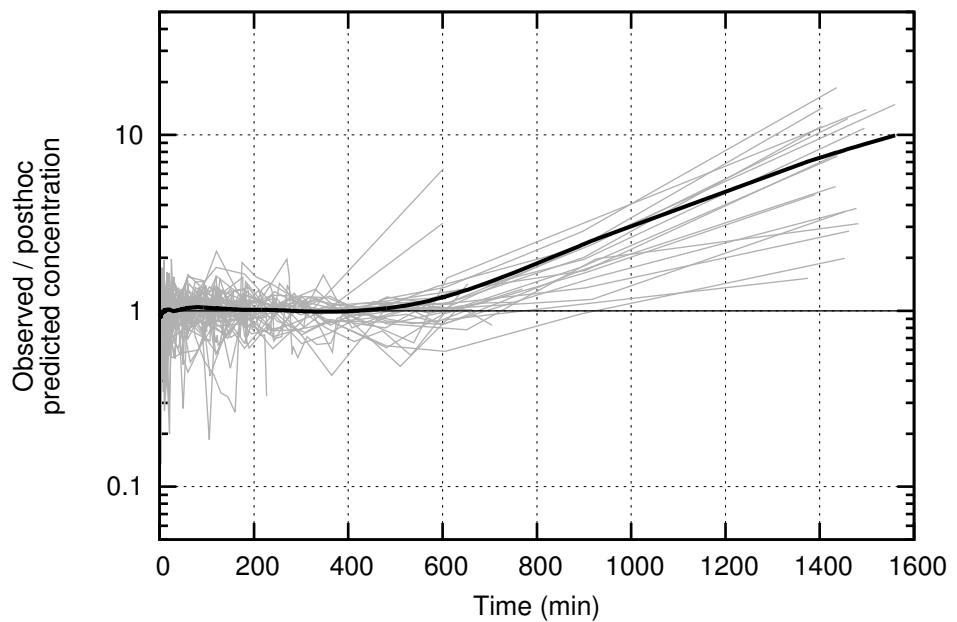
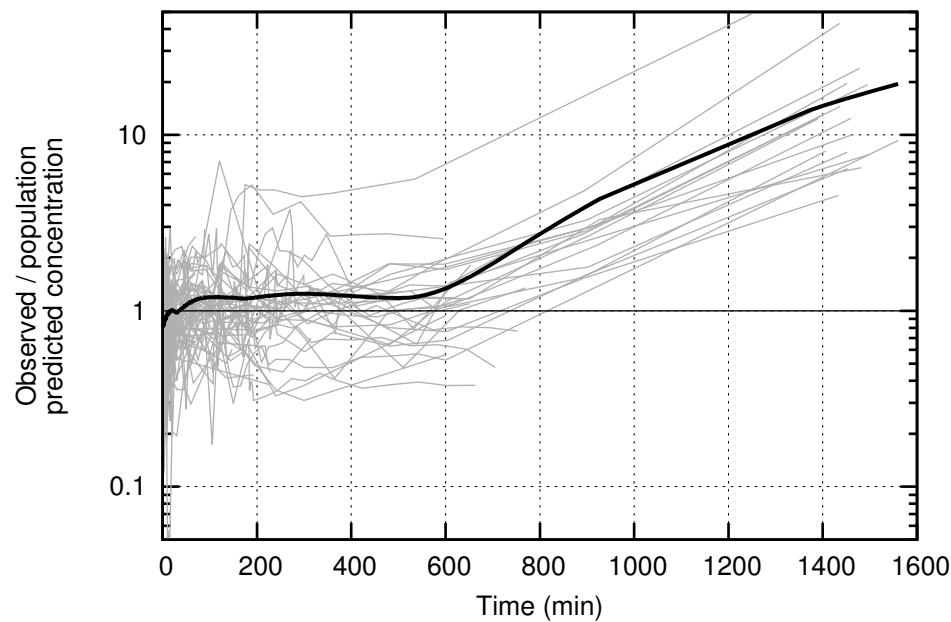
# Adults ( $18 \leq \text{age} < 70$ years, $\text{BMI} < 30$ )



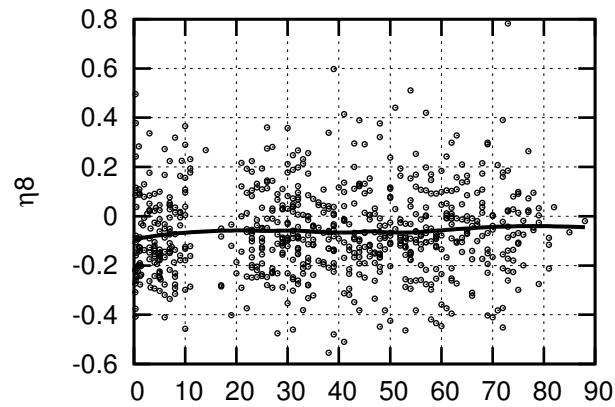
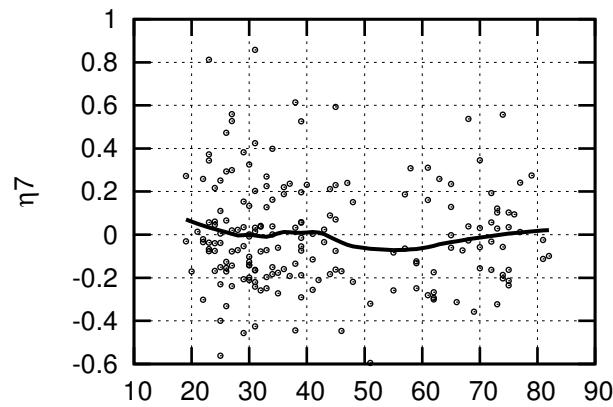
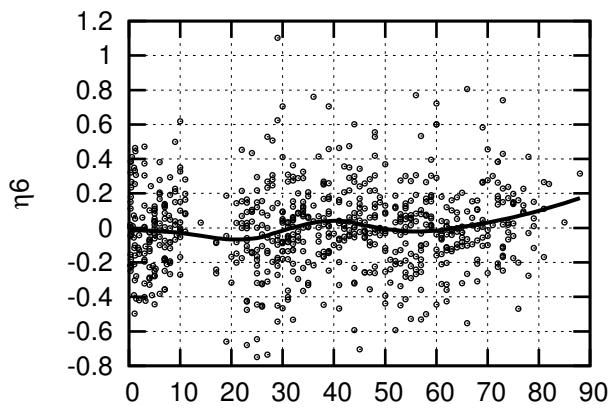
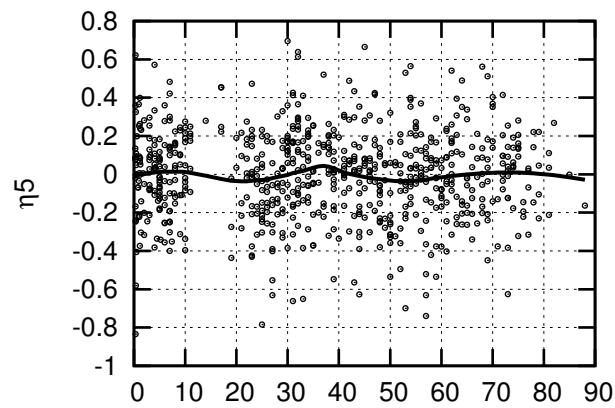
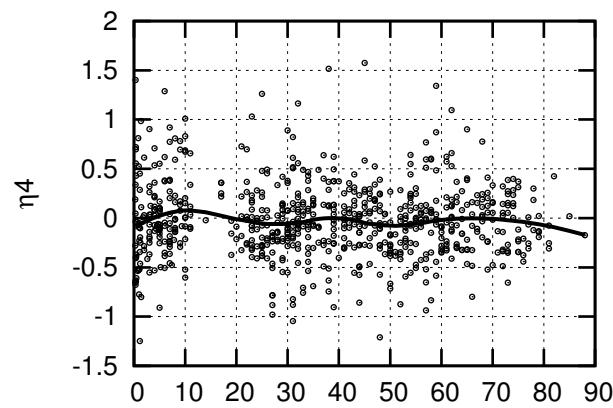
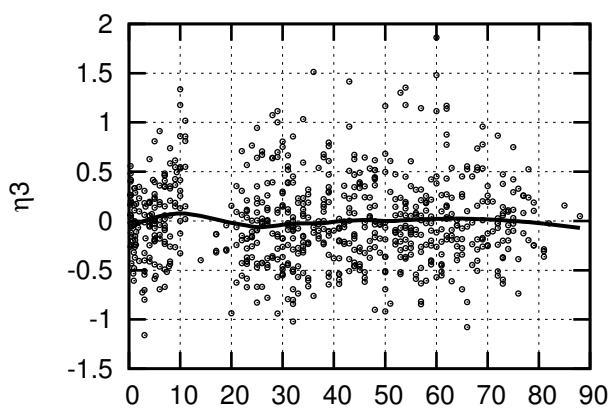
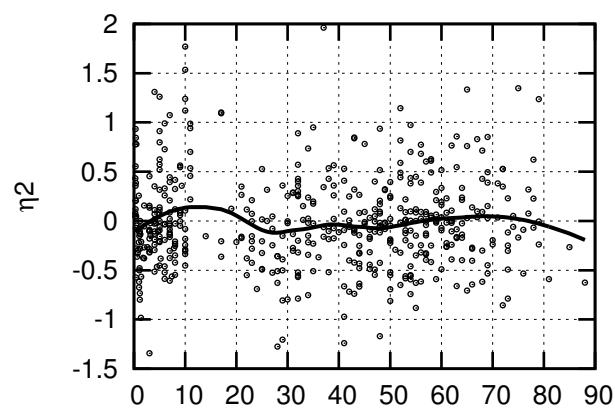
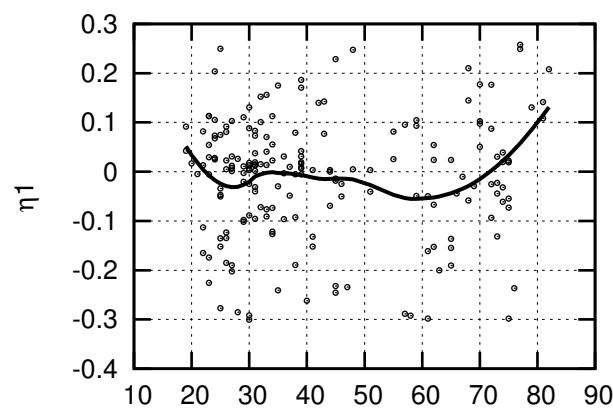
## Elderly (age $\geq 70$ years)



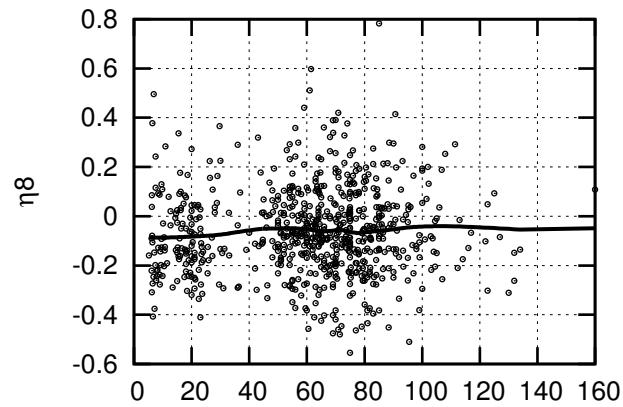
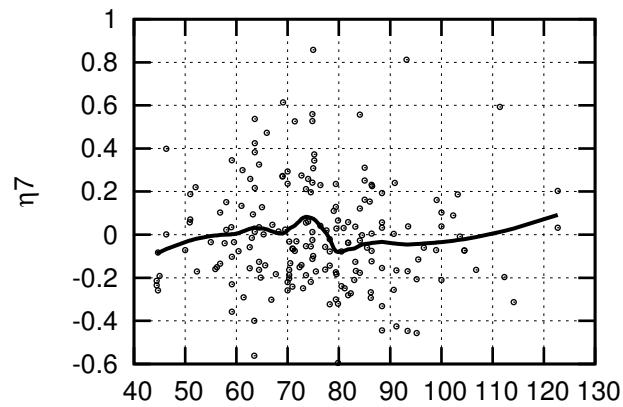
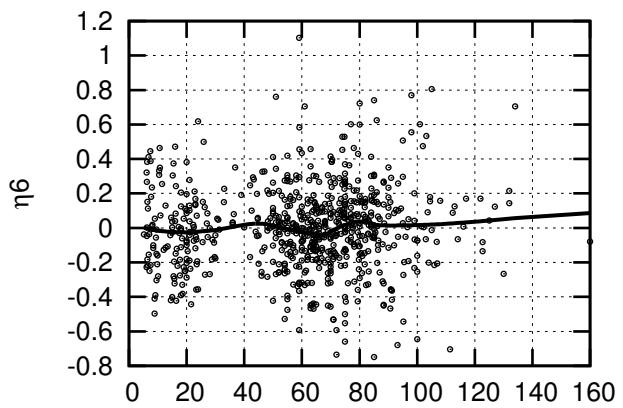
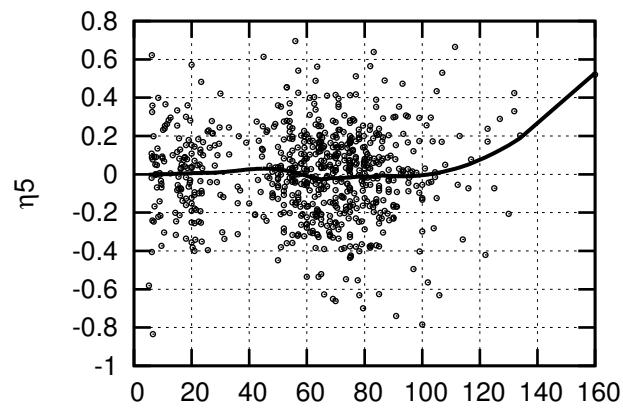
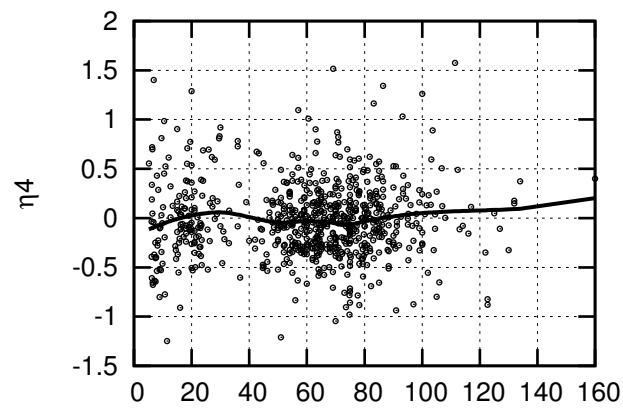
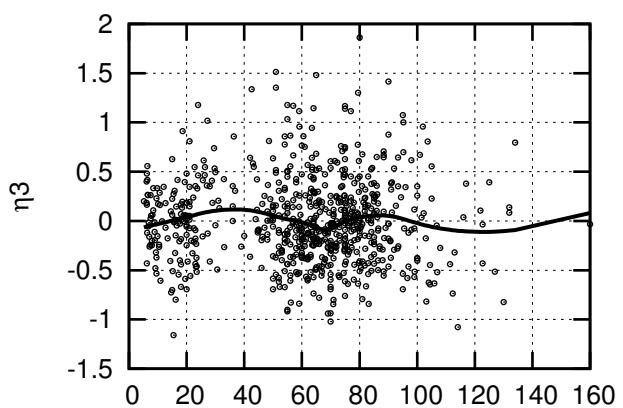
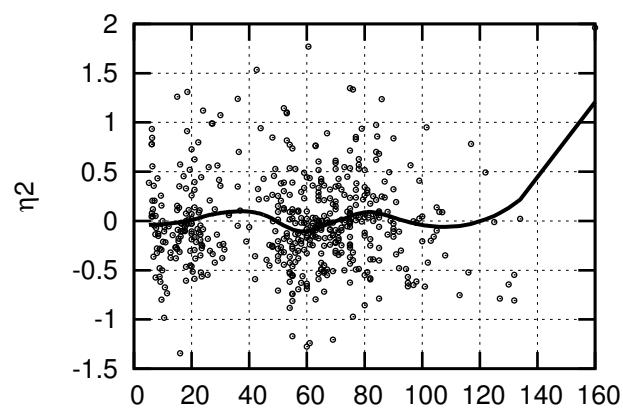
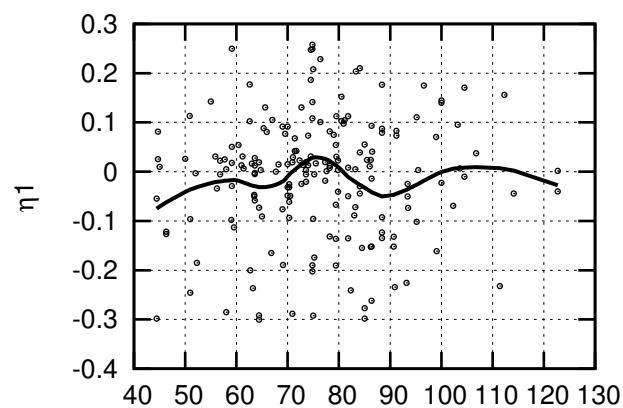
## High BMI (BMI $\geq 30$ )

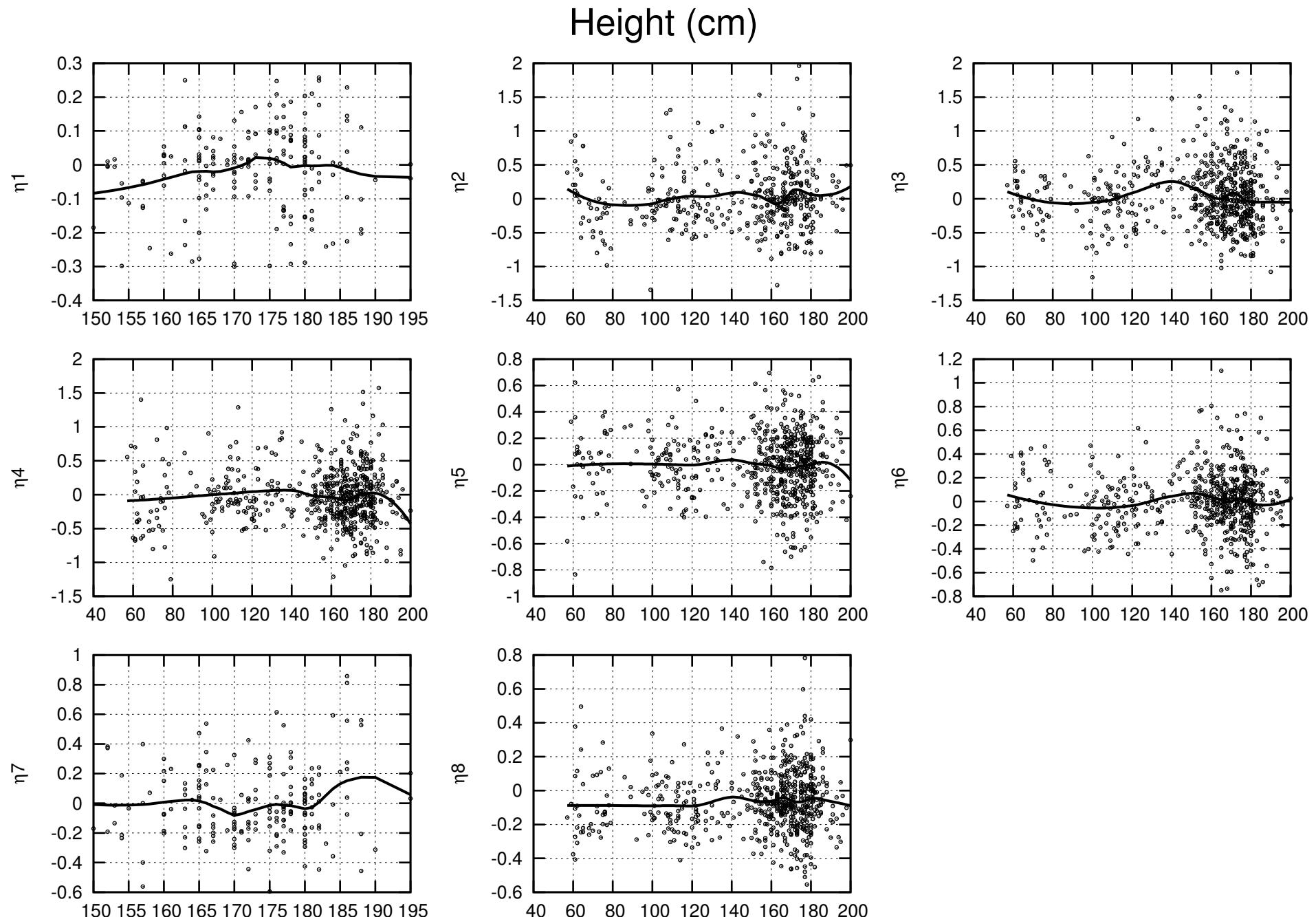


Age (years)

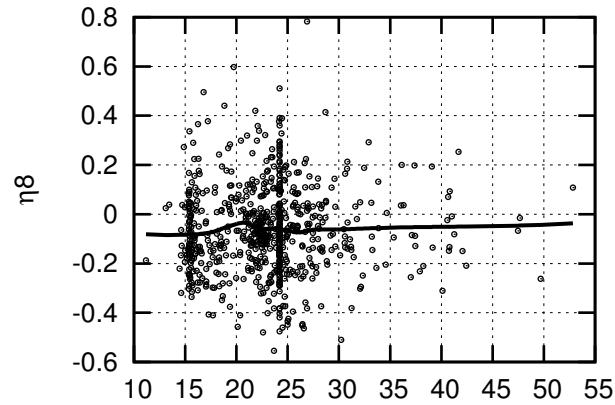
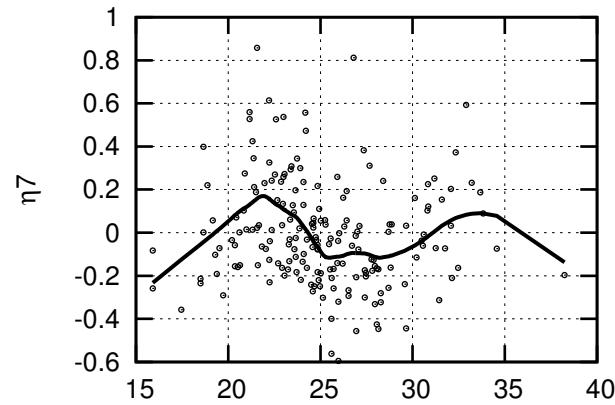
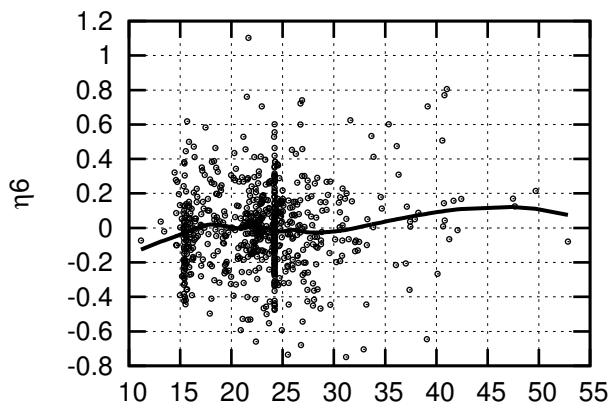
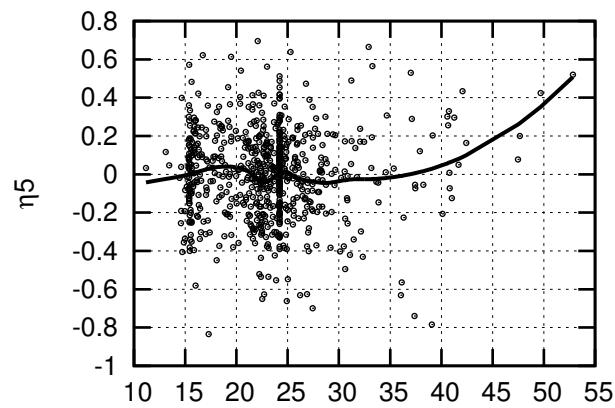
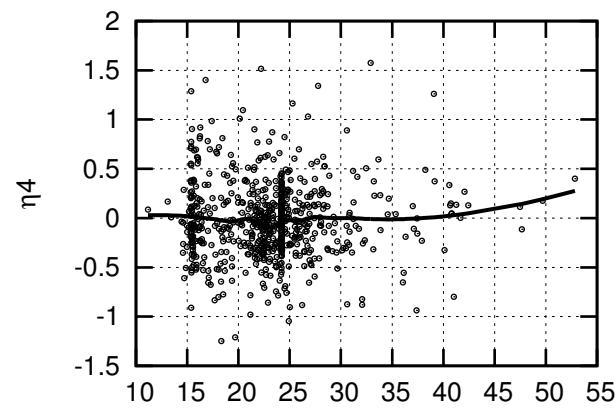
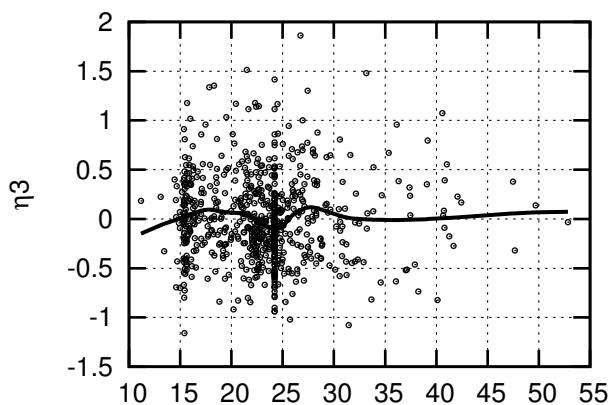
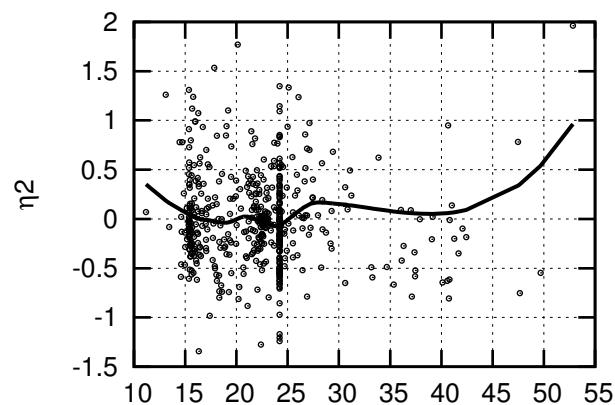
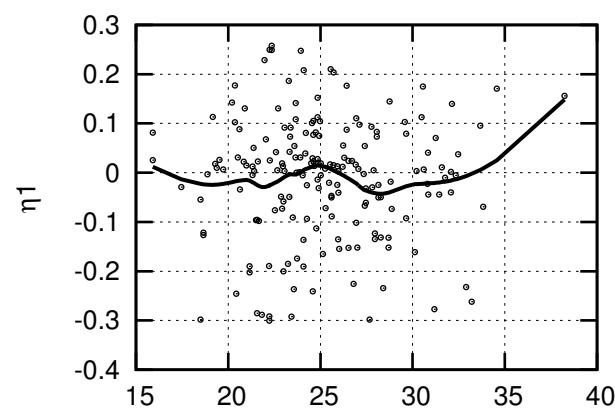


# Weight (kg)

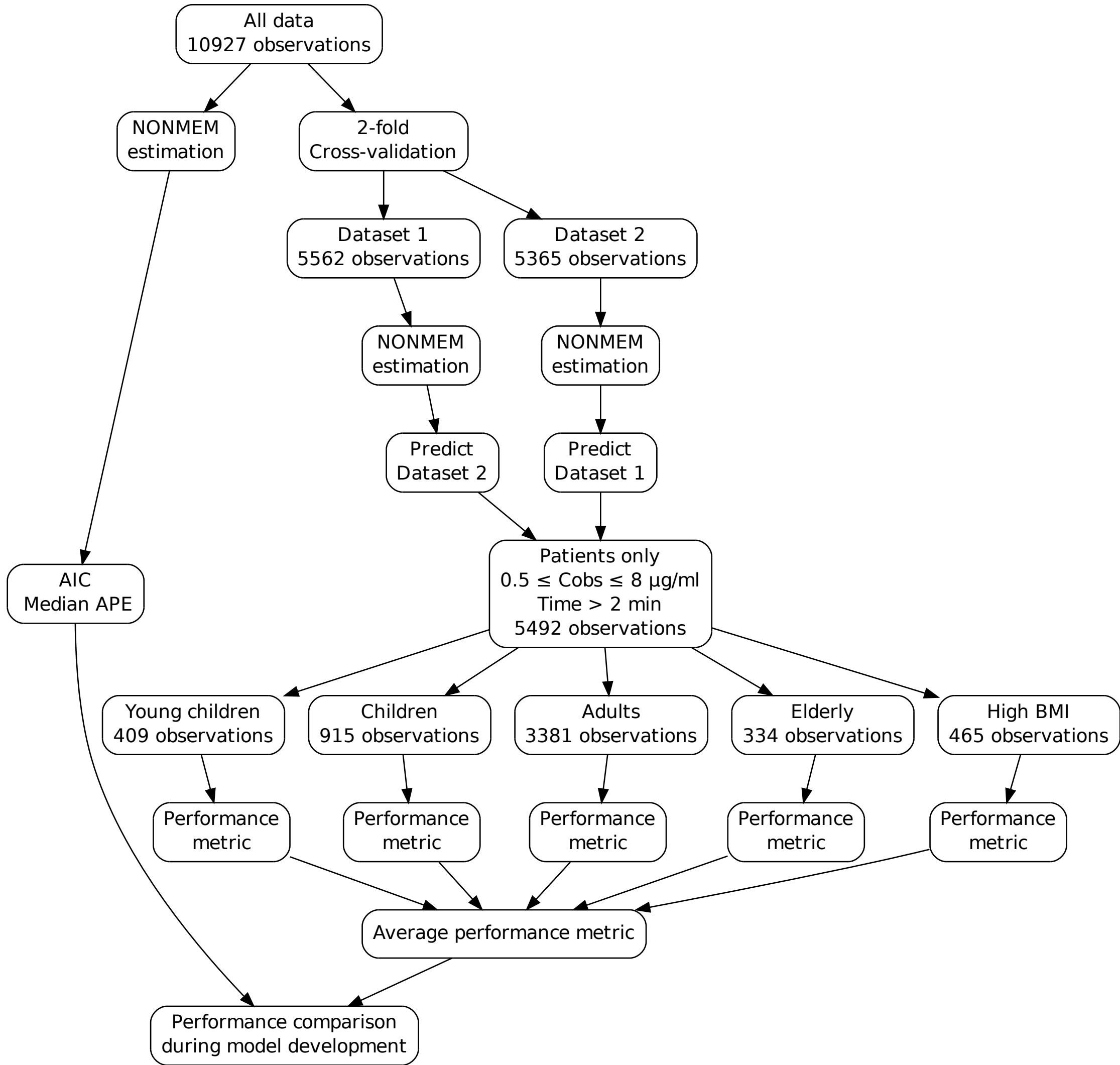




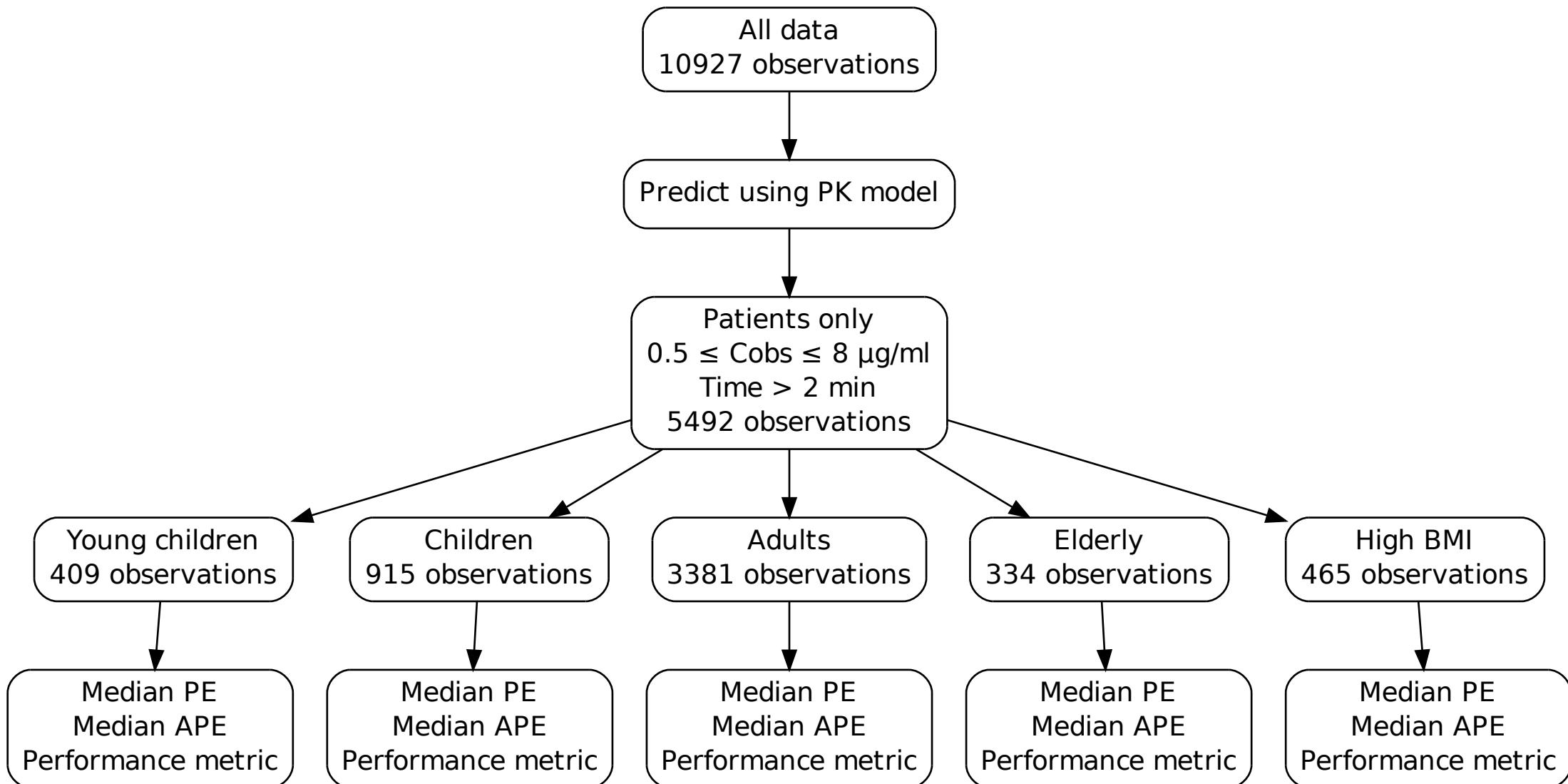
# BMI



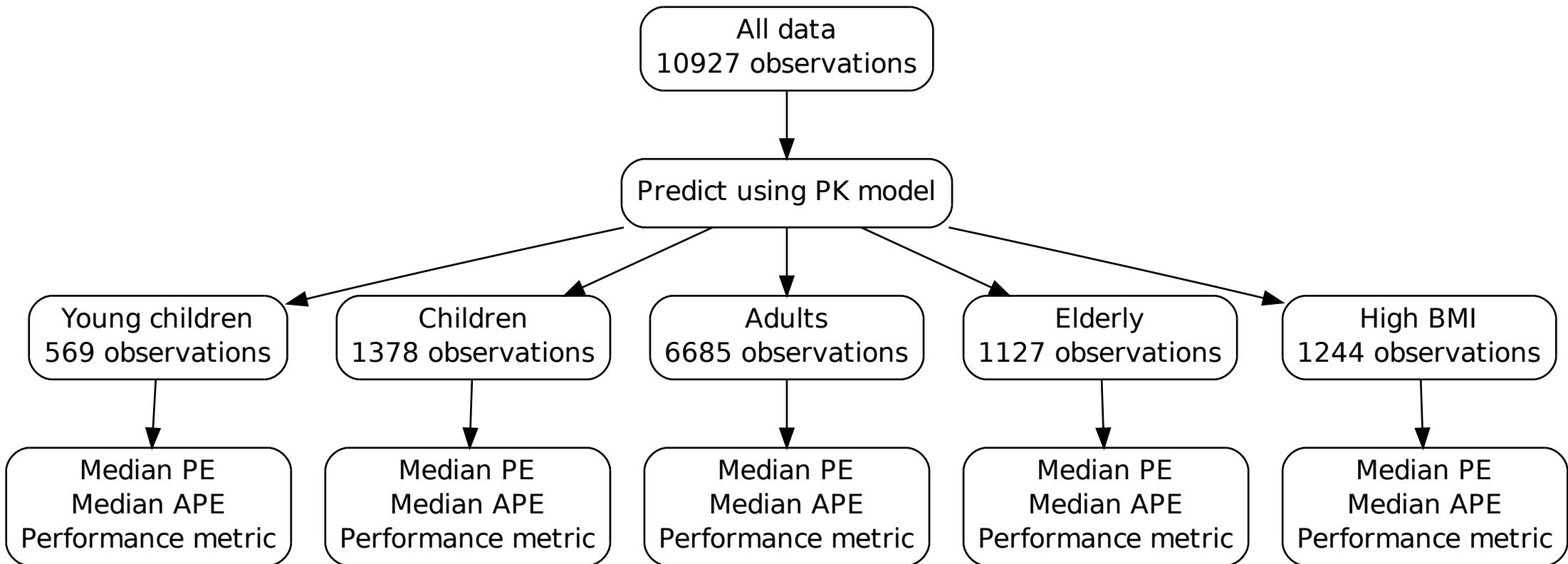
# Evaluating model performance during model development



## Evaluation of intra-operative predictive performance (Table 5)



## Evaluation of overall predictive performance (Table 6)



## Bootstrap resampling and Likelihood profiles

Uncertainty in the estimated population typical parameters was evaluated using bootstrap resampling and likelihood profiles. For bootstrap resampling 350 resamples were evaluated. The results of estimations were used regardless of NONMEM termination criteria. The red line is the NONMEM estimated population typical value. In the likelihood profiles the dark shaded area indicates the p<0.05 region and the light shaded area indicates the p<0.01 region.

$$\begin{aligned}
f_{sigmoid}(x, E50, \lambda) &= x^\lambda / (x^\lambda + E50^\lambda) \\
ADLT &= f_{sigmoid}(WGT, \Theta_{13}, \Theta_{14}) \\
f_{aging}(x) &= e^{-0.001 \cdot (AGE - 35) \cdot x} \\
CLAG &= 1 - f_{sigmoid}(PMA, \Theta_{16}, 8) \\
PMAL &= \begin{cases} f_{sigmoid}(PMA, \Theta_{15}, 8), & \text{male} \\ 0, & \text{female} \end{cases} \\
KGEN &= e^{\Theta_{23} \cdot (CLAG \cdot (1 - PMAL) + (1 - CLAG) \cdot PMAL) \cdot ADLT} \\
V1(l) &= \begin{cases} \Theta_1, & \text{healthy} \\ \Theta_7, & \text{patient} \end{cases} \cdot \frac{ADLT}{ADLT_{ref}} \cdot \begin{cases} 1, & \text{male} \\ \Theta_{22}, & \text{female} \end{cases} \cdot f_{aging}(\Theta_{18}) \cdot e^{\begin{cases} n1, & \text{healthy} \\ n2, & \text{patient} \end{cases}} \\
V2(l) &= \begin{cases} \Theta_2, & \text{healthy} \\ \Theta_8, & \text{patient} \end{cases} \cdot \left( \frac{WGT}{70} \right) \cdot f_{aging}(\Theta_{19}) \cdot e^{\eta_3} \\
V3(l) &= \begin{cases} \Theta_3, & \text{healthy} \\ \Theta_9, & \text{patient} \end{cases} \cdot \left( \frac{WGT}{70} \right)^{\Theta_{17}} \cdot \frac{ADLT}{ADLT_{ref}} \cdot e^{\eta_4} \\
CL(l/\text{min}) &= \begin{cases} \Theta_4, & \text{healthy} \\ \Theta_{10}, & \text{patient} \end{cases} \cdot \left( \frac{WGT}{70} \right)^{0.75} \cdot \frac{KGEN}{KGEN_{ref}} \cdot f_{aging}(\Theta_{18}) \cdot e^{\eta_5} \\
Q2(l/\text{min}) &= \begin{cases} \Theta_5, & \text{healthy} \\ \Theta_{11}, & \text{patient} \end{cases} \cdot \left( \frac{V2}{\Theta_8} \right)^{0.75} \cdot f_{aging}((1 - ADLT) \cdot \Theta_{20} + ADLT \cdot \Theta_{21}) \cdot e^{\eta_6} \\
Q3(l/\text{min}) &= \begin{cases} \Theta_6, & \text{healthy} \\ \Theta_{12}, & \text{patient} \end{cases} \cdot \left( \frac{V3}{\Theta_9} \right)^{0.75} \cdot f_{aging}(\Theta_{21}) \cdot e^{\begin{cases} n7, & \text{healthy} \\ n8, & \text{patient} \end{cases}} \\
C_{observed} &= C_{predicted} (1 + RES^2 \varepsilon_1 \cdot e^{n8}) + \varepsilon_2
\end{aligned}$$

Where  $\Theta_{1-23}$  are estimated parameters from the model and  $\eta_1 - \eta_8$  represent random variable of variances denoted in Table 4.  $AGE$ ,  $WGT$  and  $PMA$  represent an individual's age in years, weight in kg and post-menstrual age in years ( $AGE + 40/52$ ), respectively.  $RES$  is the proportional residual error for each component dataset. Error variance  $\varepsilon_1$  was fixed to a variance of 1 and  $\varepsilon_2$  was estimated from the data. Constants with a subscript  $ref$  are calculated for the reference individual.

Table 1: Estimated model parameters compared to results from bootstrap resampling.

Parameter	Estimated value	350 Bootstrap resamples			
		Mean	Median	Percentile	
				25%	75%
$\Theta_1$	5.74	5.70	5.75	5.51	5.92
$\Theta_2$	11.8	11.8	11.8	10.6	12.9
$\Theta_3$	222	223	225	210	238
$\Theta_4$	1.83	1.84	1.84	1.79	1.88
$\Theta_5$	3.10	3.14	3.11	2.84	3.39
$\Theta_6$	1.08	1.08	1.08	0.99	1.17
$\Theta_7$	9.77	9.65	9.66	9.32	9.97
$\Theta_8$	29.0	29.1	29.2	27.6	30.6
$\Theta_9$	134	133	133	128	139
$\Theta_{11}$	1.53	1.55	1.54	1.51	1.57
$\Theta_{11}$	1.42	1.44	1.44	1.39	1.49
$\Theta_{12}$	0.61	0.60	0.60	0.58	0.63
$\Theta_{13}$	16.6	16.5	16.4	15.7	17.2
$\Theta_{14}$	2.75	2.79	2.78	2.63	2.93
$\Theta_{15}$	22.2	24.0	22.3	20.5	25.3
$\Theta_{16}$	69.2	68.8	68.6	66.7	71.0
$\Theta_{17}$	0.35	0.35	0.34	0.30	0.39
$\Theta_{18}$	3.55	3.45	3.46	3.09	3.82
$\Theta_{19}$	8.91	9.02	8.84	7.59	10.24
$\Theta_{20}$	27.6	27.1	27.2	24.7	29.3
$\Theta_{21}$	6.34	6.21	6.33	5.24	7.33
$\Theta_{22}$	0.830	0.837	0.836	0.809	0.864
$\Theta_{23}$	0.225	0.24	0.242	0.214	0.263
Variance					
$\eta_1$	0.032	0.037	0.029	0.017	0.049
$\eta_2$	0.323	0.326	0.330	0.295	0.352
$\eta_3$	0.294	0.296	0.294	0.267	0.322
$\eta_4$	0.318	0.318	0.316	0.289	0.344
$\eta_5$	0.068	0.068	0.068	0.064	0.072
$\eta_6$	0.130	0.125	0.126	0.103	0.147
$\eta_7$	0.091	0.088	0.087	0.070	0.105
$\eta_8$	0.044	0.042	0.041	0.037	0.046

### Bootstrap distribution of population parameter estimates (350 resamples)

