

Changes in liver volume II – a population-based approach to meta-analysis of paediatric, adult and geriatric populations - an update



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Objectives

- Liver volume (LV) is a critical parameter for both liver transplantation and as a scaling factor for predicting clearance (CL) in physiologically-based pharmacokinetic modelling (PBPK); therefore obtaining its accurate and precise estimation is essential.
- The objective of this study was to extend an existing meta-analysis for the estimation of LV to other race groups (e.g. Caucasian, Japanese, Chinese) and paediatric and geriatric populations using non-linear mixed effect modelling techniques.

Methods

The overall workflow (Fig 1) comprised the following steps:

- Development of a search string to ensure optimal retrieval of data.
- Extraction of data and simulation of missing variables to give a final dataset including both averaged study data as well as individual datum.
- Construction of a non-linear mixed effect model (NLME) of the data including covariates.
- Final covariate selection and model evaluation

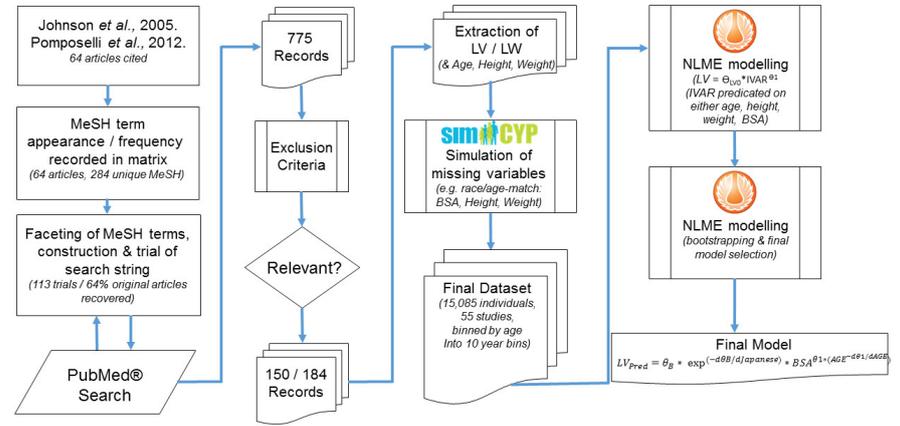


Figure 1. A workflow describing the meta-analysis and NLME modelling of the acquired LV data

Results

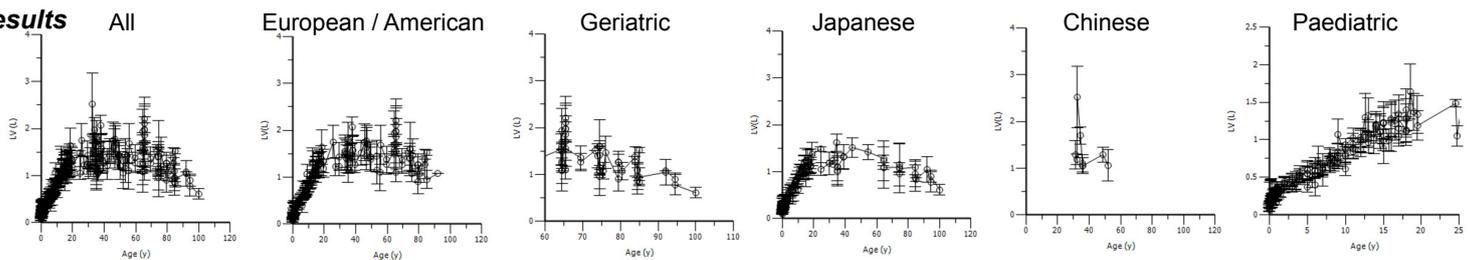


Figure 2. Observed LV (L) data by age (y) for different population groups

- Data collated from 55 studies included 2906 LV measurements from averaged data and 1787 from 'individuals'.
- In this context 'Individuals' are representative of the collated studies binned by age (i.e. 10y bands; Fig 2) and therefore IPRED was reflective of a study rather than individual prediction *per se*. Evaluation of covariate models revealed well performing models predicated using bodyweight and body surface area (BSA) as independent variables (IVAR) (see Table 1).
- The BSA predicated model was chosen (Final model), due to its parsimony when compared against the Urata model and it was therefore less complex for a similar 'goodness of fit'.
- Predicted population estimates of LV as a function of BSA (left and middle) and conditionally weighted residuals (CWRES) (right) were well determined against observed values (Fig 3).

Table 1. Comparison and evaluation of optimal models (predicated using different IVAR) NB. * additional constants in the Urata model are not considered within #Parms

Model*(IVAR)	Model #	Covariates retained	-2(LL)	AIC	BIC	#Parms	#Obs	#Subj	p-value
BSA simple run	1	dθ1/dAGE, dLV0/dJapanese	-631.03	-617.03	-591.51	7	283	55	3.80E-15
WT simple run	2	dLV0/dCaucasian, dθ1/dCaucasian, dLV0/dJapanese, dθ1/dJapanese, dLV0/dAGE	-633.61	-613.61	-577.15	10	283	55	5.20E-33
HT simple run	3	dLV0/dMETHODS, dθ1/dMETHODS, dθ1/dJapanese	-514.11	-496.11	-463.3	9	283	55	3.70E-76
Age simple run	4	dLV0/dMETHODS, dθ1/dMETHODS, dLV0/dJapanese	-338.7	-324.7	-299.19	7	283	55	1.10E-45
Yu simple run (WT & HT)	5	dθ1/dJapanese, dLV0/dChinese	-632.98	-612.98	-576.53	10	283	55	1.20E-08
Urata simple run (BSA)	6	dθ1/dMETHODS, dLV0/dJapanese	-631.11	-617.11	-591.59	7*	283	55	1.20E-15

$$\text{Final model: } LV_{Pred} = \theta_B \exp(-d\theta_B/d_{Japanese}) BSA^{\theta_1} (AGE^{-d\theta_1/dAGE})$$

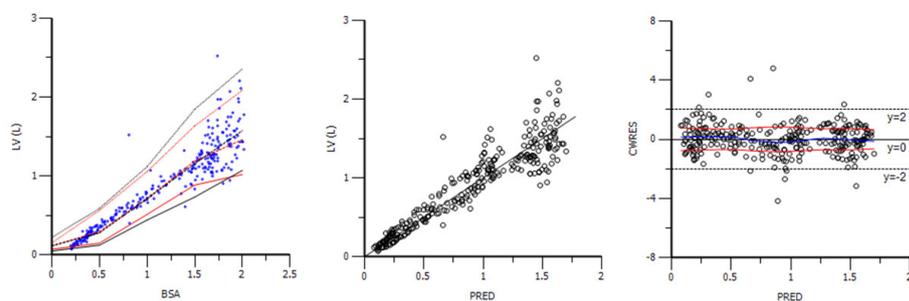


Figure 3. Bootstrapped predicted population estimates of liver volume as a function of BSA predicated model (left and middle panels) and CWRES (right panel) from this same model respectively. The bulk of observed quantiles (red) fall within the 5 – 95 % range of predicted quantiles (grey) for the prediction intervals

Conclusion

- Age and BSA are used here as important predictors of LV.
- Traceable model-based meta-analysis is important for arriving at accurate values for clinical parameters.
- The final model corrects some of the deficiencies of a previous model¹ and may be applied both clinically and to PBPK modelling.

References

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- Urata K et al. Hepatology 1995;21:1317-21.
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