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## Variability in organ-level effects in repeat dose animal studies

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Observations of liver, kidney, stomach, spleen, thyroid and

Building scientific confidence in the use of new approach methodologies (NAMs) in safety assessment may include performance comparison to *in vivo* study outcomes. This work defines the variability in organ-level effects and suggests qualitative and quantitative benchmarks for maximum NAM performance for prediction of organ-level effects in repeat dose studies of adult animals. Previous work suggests that the root residual mean square error (RMSE) for study-level lowest effect level (LEL) values (on a log<sub>10</sub>-mg/kg/day basis) approaches 0.5 log<sub>10</sub>-mg/kg/day.

## Table 1. Questions on animal study reproducibility

What is the range of possible

systemic effect values (mg/kg/day) in replicate studies? Residual root mean square error (RMSE) is an estimate of variance in the same units as the systemic The RMSE can also be used to define a minimum prediction interval, or estimate range, for a

What is the maximal accuracy of a model that attempts to predict a systemic effect values for an

 The mean square error (MSE) is used to approximate the unexplained variance (not explained by study descriptors).

 This unexplained variance limits the R-squared on a new model.

LOAEL

What is the probability that an effect in adult animals will be observed in replicate studies?

Understand the reproducibility of treatmentrelated changes in specific endpoint targets (e.g., any effect on liver).

Previous work suggests that the variance (estimated by RMSE) in study-level effect values from repeat dose studies in animals approaches 0.5 log10-mg/kg/day.

- Total variance in systemic toxicity effect values likely approaches 0.75-1 (units of  $(\log_{10}-mg/kg/day)^2$ )
- MSE (unexplained variance) is 0.2 0.4 (units of ( $\log_{10}$ mg/kg/day)<sup>2</sup>)
- RMSE is 0.45-0.60 log<sub>10</sub>-mg/kg/day
- RMSE is used to define a 95% minimum prediction interval (i.e., based on the standard deviation or spread of the

Calculating the minimum prediction interval width based on results of multi-linear regression modeling

lower bound on prediction =  $10^{qnorm(0.025*RMSE)}$ upper bound on prediction =  $10^{qnorm(0.075*RMSE)}$ 

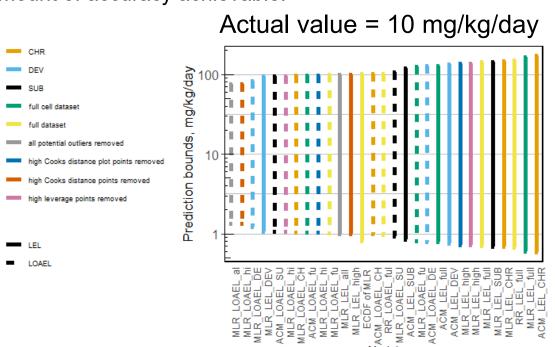
Based on Pham LL, Watford S, Pradeep P, Martin MT, Thomas RS Judson RS, Setzer RW, Paul Friedman K. Accepted. "Variability in in vivo studies: Defining the upper limit of performance for predictions of systemic effect levels." Computational Toxicology.

https://doi.org/10.1016/j.comtox.2020.100126

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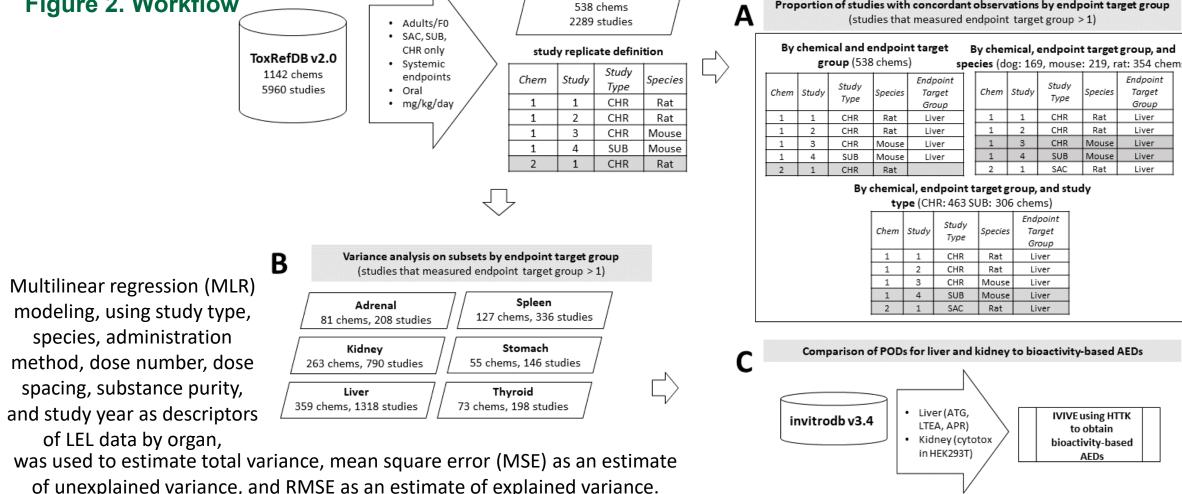
## Figure 1. Estimating minimum prediction intervals based on animal study variance (based on Pham et al., 2020).

If attempting to use a NAM-based predictive model for prediction of a reference systemic effect level value of 10 mg/kg/day, it is likely that given the variability in reference data of this kind, that a model prediction of somewhere between 1 and 100 mg/kg/day would be the greatest amount of accuracy achievable.



included in this analysis to understand the reproducibility of organ-level effects. Figure 2. Workflow 538 chems 2289 studies SAC, SUB, study replicate definition

adrenal gland from the Toxicity Reference database (v2.0) were



A: What is the qualitative reproducibility of organ-level effect observations in repeat dose studies of adult animals?

Table 2. Repeated concordance of organ-level chemical with positive finding in all studies + chemicals with negative finding in all studies

	Endpoint	%	Che				
	target group	Concord	m	+Pos	-Neg	Mixed	
	adrenal	60.2	538	8	316	214	
	Kidney	38.8	538	54	155	329	
	Liver	42.4	538	149	79	310	
	Spleen	56.5	538	17	287	234	
	Stomach	71.7	538	14	372	152	
	Thyroid	66.2	538	11	345	182	

	Stomach		7	<u> 1.7                                      </u>	538	<u> 14</u>	37	2	152
	Thyroid		6	<b>6.2</b> 538		3 11	345		182
۰	Endpoint								
	target	Stud	y	%					
	group	Тур	<u> </u>	Conc	ord	Chem	+Pos	-Neg	Mixe
	adrenal			67.	8	463	8	306	14
	kidney			49		463	58	169	23
	liver	CLIF	אווס	54.	6	463	160	93	21
	spleen	CHR	67.	8	463	16	298	14	
	stomach			79		463	22	344	97
	thyroid			70		463	10	314	13
	adrenal			73.	5	306	10	215	81
	kidney	SUB	<b>52</b> .	6	306	65	96	14	
	liver		66		306	143	59	10	
	spleen		68		306	24	184	98	
	stomach		85		306	10	250	46	
	thyroid			81		306	11	237	58
_									

% Concord = percent concordant chemicals; Chem = total # chemicals tested at the endpoint target group; +Pos = # chemicals with positive observations in all available studies; -Neg = # chemicals with negative observations in all available studies; Mixed = chemicals with at least 1 study that was not positive

Endpoint	%					
target group	Concord	Chem	+Pos	-Neg	Mixed	
	dog	84.6	169	8	135	26
adrenal	mouse	84	219	6	178	35
	rat	66.9	354	17	220	117
	dog	67.5	169	20	94	55
kidney	mouse	63.5	219	43	96	80
	rat	57.6	354	106	98	150
	dog	71	169	86	34	49
liver	mouse	67.1	219	96	51	72
	rat	61.3	354	157	60	137
	dog	78.1	169	9	123	37
spleen	mouse	74	219	16	146	57
	rat	65.5	354	31	201	122
	dog	87.6	169	2	146	21
stomach	mouse	80.4	219	7	169	43
	rat	79.9	354	11	272	71
	dog	78.7	169	8	125	36
thyroid	mouse	90.4	219	3	195	21
	rat	77.4	354	28	246	80

Replicate studies were defined by chemical only, chemical and species, and chemical and study type to estimate concordance in observed organ-level effects (repeated presence/absence of weight, gross or histopathological changes) for a chemical. Total concordance (% chemicals with positive or negative agreement across replicates), depending on the organ and replicate definition, ranged from 39 - 88%, with slightly greater average within-species concordance. Organs associated with more negative chemicals (stomach, thyroid, adrenal) had slightly higher rates of concordance in this range.

B: What is the variance in organ-level effects in repeat dose studies, and is it smaller than study-level variance?

Table 3. Results of MLR to estimate unexplained and explained variance in organ LELs.

organL	$EL \sim b_0 + cnemical * b_1 + species * b_2$
+ study	$type * b_3 + administration method * b_4$
+ dose s	$spacing * b_5 + number of dose levels * b_6$
+ study	$year * b_7 + \%$ substance purity $* b_8$

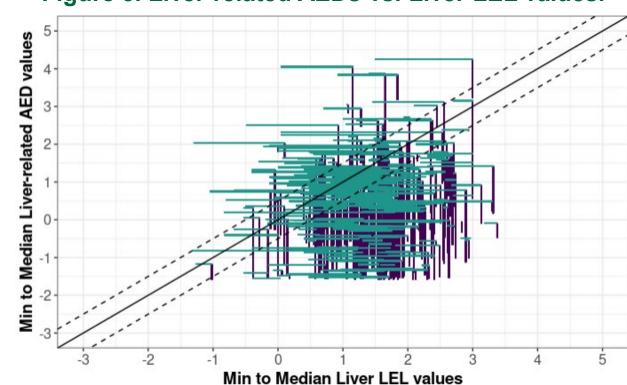
Endpoint						
Target						% var
Group	Chem	Ν	Var	MSE	RMSE	explained
adrenal	81	208	0.756	0.349	0.591	53.8
kidney	263	790	0.765	0.316	0.562	58.7
liver	359	1318	0.745	0.355	0.596	52.3
spleen	127	336	0.671	0.318	0.564	52.6
stomach	55	146	0.553	0.173	0.416	68.7
thyroid	73	198	0.721	0.378	0.615	47.6

Total variance at the organ level is generally less than or equal to total variance at the study-level. The RMSE at the organ level is similar to the study level RMSE in Pham et al. The % variance explained is similar to the lower estimate of % variance explained at the study level in Pham et al.

## C. How well do liver-related bioactivity-derived administered equivalent doses approximate liver-related LEL values?

A 3 compartment steady-state model (R library(httk)) was used in the in vitro to in vivo extrapolation of administered equivalent doses in order to include the largest number of chemicals in this comparison. The plasma steady state concentration for the median individual based on Monte Carlo simulation of human physiological parameters was used.

Figure 3. Liver-related AEDs vs. Liver LEL values.



The linear relationship between liver LELs and liverrelated AEDs is poor, but the AED values generally provide a conservative estimate of liver LEL.

- Wilcoxson tests of the LEL and AED values by DSSTox chemical identifier suggest that mean values are different (p<0.05) for 163/306 chemicals in this comparison (53% of the time).
- However, the median liver-related AED was less than the minimum liver LEL value for 175/306 chemicals (57%) and the minimum liver-related AED was less than the minimum liver LEL value for 267/306 (87%). 286/306 chemicals (93%) demonstrated a minimum liver-related AED less than the minimum liver LEL + 0.5 log10-,mg/kg/day.

With MSE used to indicate unexplained variance, results suggest study descriptors accounted for 52-69% of the total variance in organ-level LELs. A NAM would be unlikely to explain more than the variance explained by study level descriptors, or 70%, of the variance in these data. The RMSE for these organ-level statistical models ranged 0.4 – 0.6 log<sub>10</sub> mg/kg/day, suggesting organ-level variance in LEL values was similar to overall study LEL variance. Therefore, a good NAM might predict organ-level LELs within ± 1 log<sub>10</sub>-mg/kg/day. This work suggests thresholds on NAM accuracy for repeat dose, organ-level effects in adult animals.