

Applying a PBTK model for IVIVE

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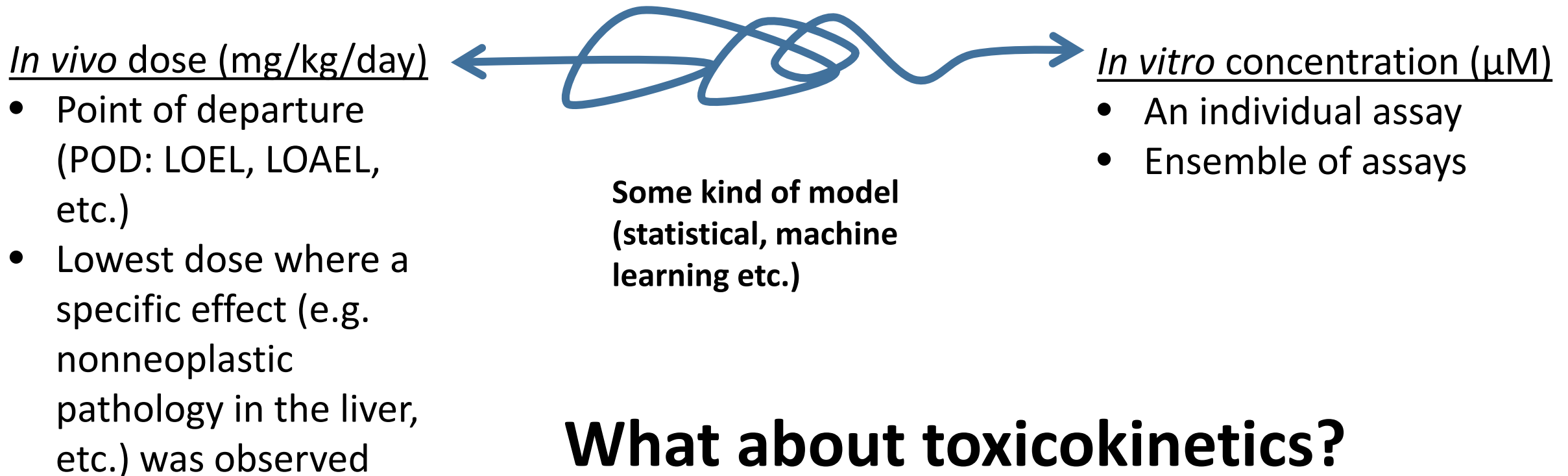
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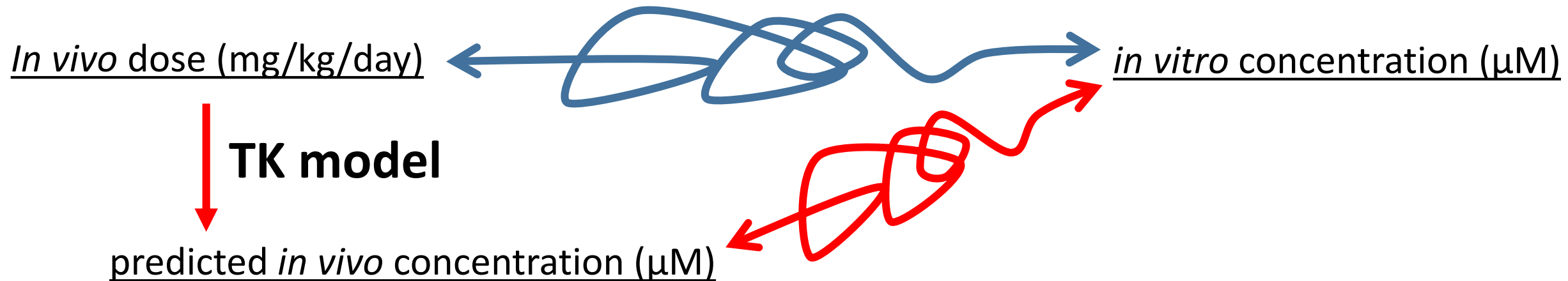
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In Vitro to In Vivo Extrapolation (IVIVE)

Enables use of high throughput toxicity assays as an alternative to animal testing



In Vitro to In Vivo Extrapolation (IVIVE)

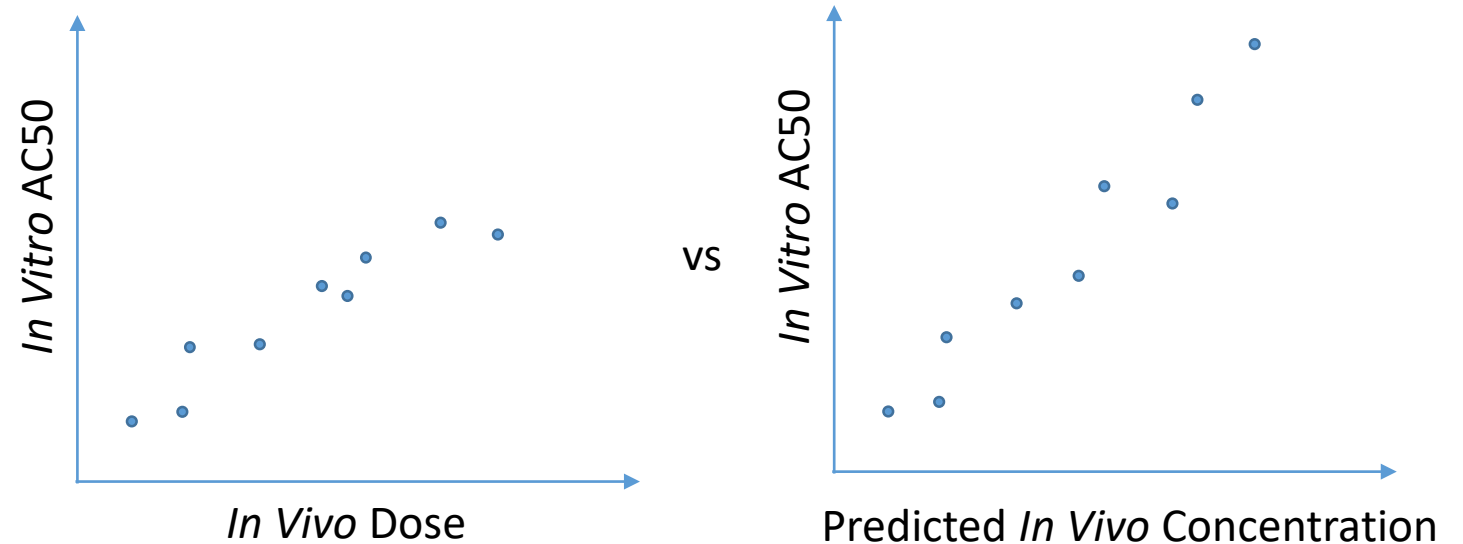


Does incorporating toxicokinetics improve the correlation between *in vitro* and *in vivo* toxicity data?

What are the effects of the assumptions in the application of TK?

In Vitro to In Vivo Extrapolation (IVIVE)

Ideally, we would know of an *in vitro* assay that is related to a specific *in vivo* effect (pathology); use this to evaluate the effect of incorporating TK



In the absence of this information, we:

1. Evaluate all *in vitro* assay endpoints against all *in vivo* effects (ToxRef)
2. Evaluate all *in vitro* assay endpoints against points of departure (ToxVal POD; determined across all *in vivo* effects)

Prior Work

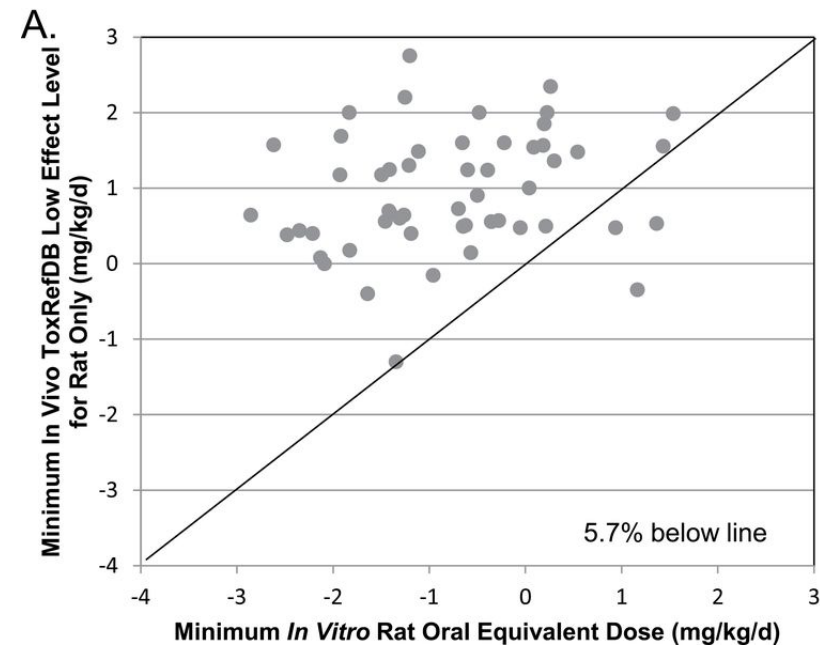
Measured rat-specific intrinsic clearance (Cl_{int}) and fraction of unbound chemical in plasma (f_{up}) for **56 chemicals** with rat ToxRefDB data.

Used **steady-state** PK model for *in-vitro* to *in-vivo* extrapolation.

$$\text{Rat } C_{ss} = \frac{ko}{(GFR \times F_{ub}) + \left(\frac{Q_l \times F_{ub} \times Cl_{int}}{Q_l + F_{ub} \times Cl_{int}} \right)}$$

- Using TK for IVIVE may help define exposure heuristics
- Incorporating TK did not otherwise improve predictive performance
 - Assumptions in the application of TK may have influenced this result

Comparison of the *in vitro* assay with the lowest oral equivalent dose with the *in vivo* response with the lowest LEL for each chemical.



This Work

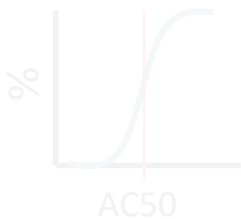
- PBTK model
- Evaluate assumptions:
 - Clearance
 - PBTK concentration selection (mean vs. max, etc.)
 - Accounting for partitioning in cell based assays
- 104 chemicals w/ rat specific *in vitro* measured values for (Cypotex)

To prepare this analysis:

- 1) Measured values for f_{up} and Cl_{int}
- 2) Select doses and examine scope of the data (i.e. what assays and *in vivo* effects can we

look at?)

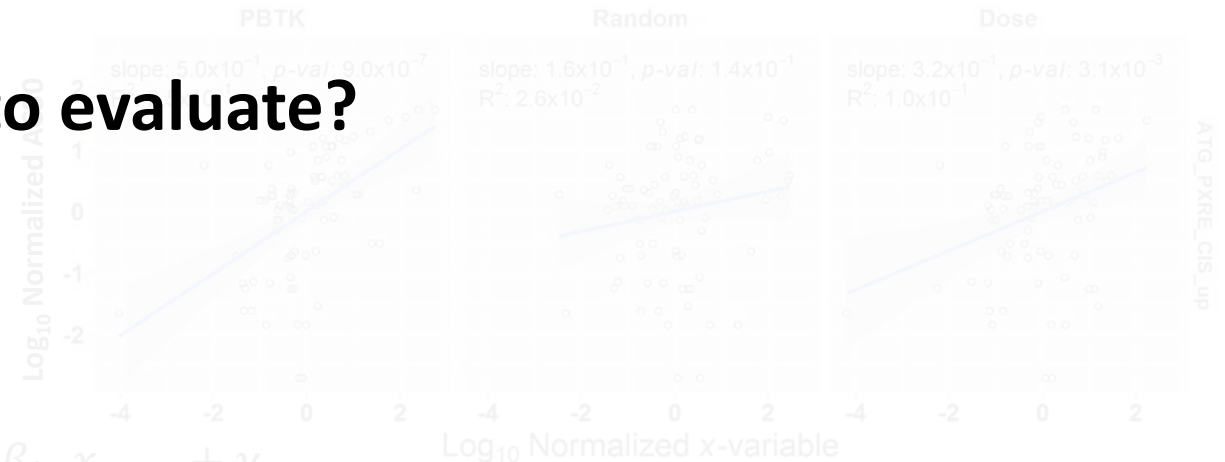
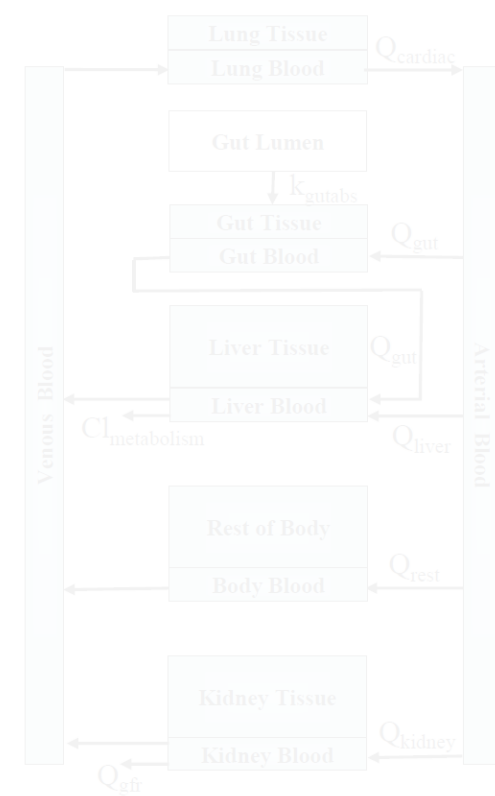
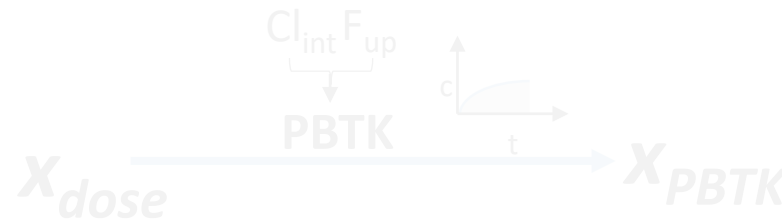
- 3) What assumptions to evaluate?



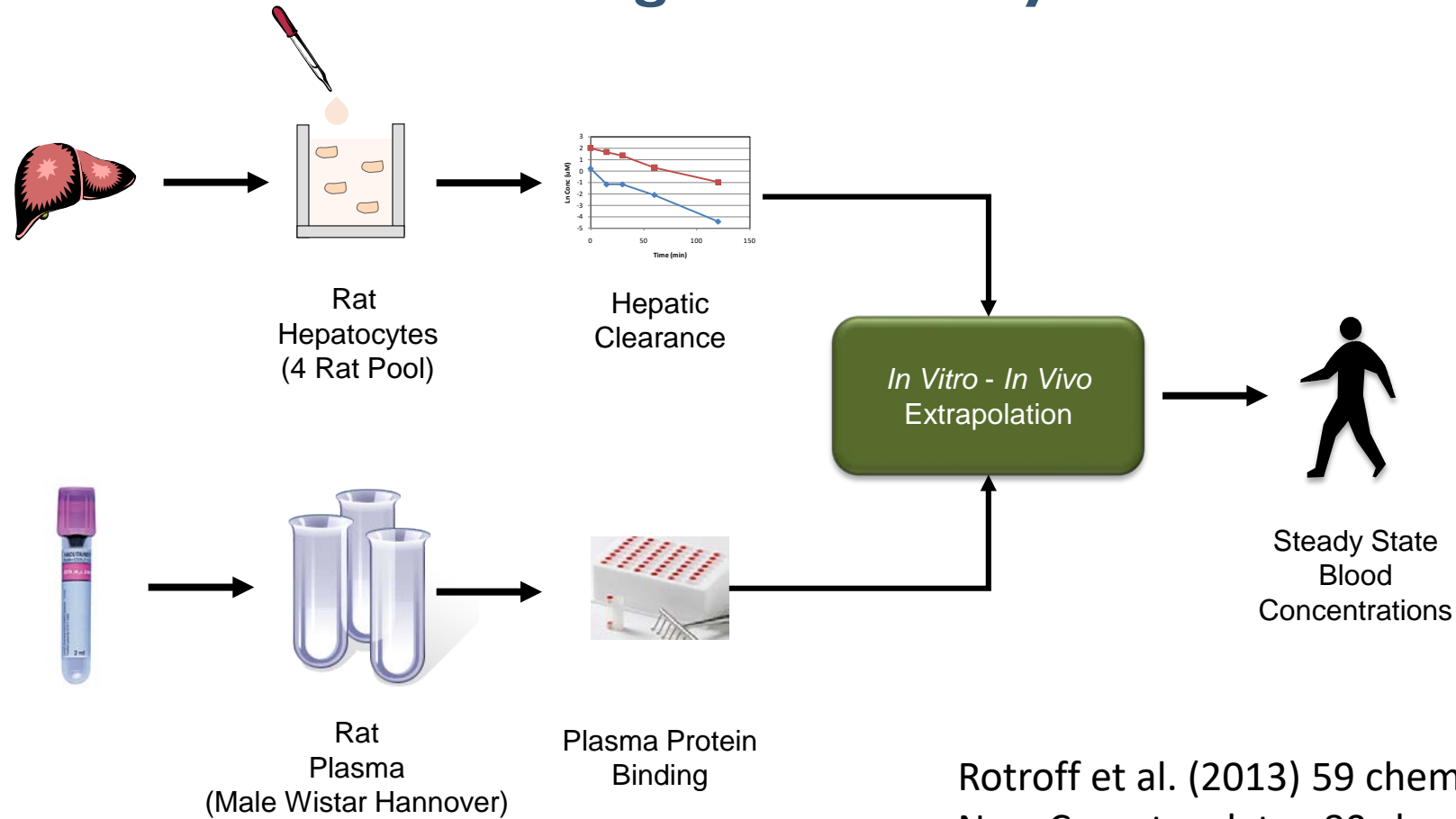
$$\begin{aligned} AC50 &\sim \beta_{i,1} x_{PBTK} + \gamma \\ AC50 &\sim \beta_{i,2} x_{rand} + \gamma \\ AC50 &\sim \beta_{i,3} x_{dose} + \gamma \end{aligned}$$

Multiple Regression

$$AC50 \sim \beta_{j,1} x_{PBTK} + \beta_{j,2} x_{rand} + \beta_{j,3} x_{dose} + \gamma$$



Characterizing Rat *In Vivo* Toxicokinetics Using *In Vitro* Assays



Rotroff et al. (2013) 59 chemicals
New Cyprotex data +80 chemicals

Slide from Barbara Wetmore

Scope of the Data – Selecting Dose

ToxRef Effect Level Dose (ELD)

- **ToxRef***: Detailed database of *in vivo* effect and dose
- **Effect level dose** defined as the **minimum dose** at which a **particular effect** (category-type-target, e.g. systemic-nonneoplastic pathology-liver) was observed for a given study and chemical, ignoring gender and strain; **a specific effect**

For a given chemical

- x_{dose} – doses for specific effect and study
- x_{PBTk} – concentration from transforming x_{dose} via PBTk
- x_{rand} – concentration from transforming x_{dose} via the randomly parameterized PBTk model

ToxVal POD Dose

- **ToxVal**: General database of *in vivo* POD
- Lowest observed effect level (**LOEL**) or lowest observed adverse effect level (**LOAEL**) for a given chemical and study; all usable rat studies; **across all effects**

For a given chemical

- x_{dose} – minimum POD across all studies
- x_{PBTk} – minimum concentration from transforming all POD via the PBTk model
- x_{rand} – minimum concentration from transforming all POD via the randomly parameterized PBTk model

Compare against every *in vitro* toxicity assay

Scope of the Data – An Example

ToxRef Effect: systemic-nonneoplastic pathology-liver

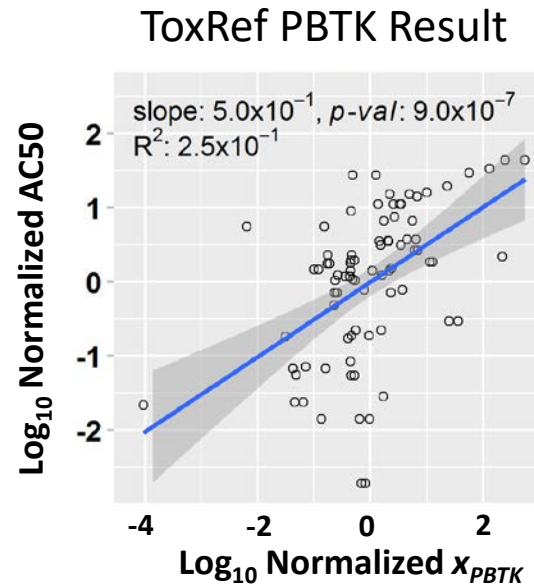
- Possibly multiple points for a given chemical

Assay endpoint: ATG_PXRE_CIS_up

- Single point for a given chemical

Number of points in the regression: 85

Number of chemicals in the regression: 49



**1 of ~40,000 assay
and *in vivo* effect
pairs**

ToxVal POD: LOEL or LOAEL

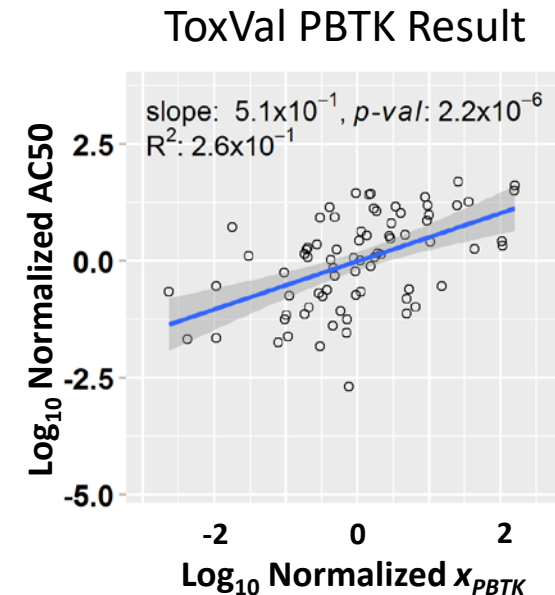
- Single point for a given chemical
(minimum taken across studies)

Assay endpoint: ATG_PXRE_CIS_up

- Single point for a given chemical

Number of points in the regression: 76

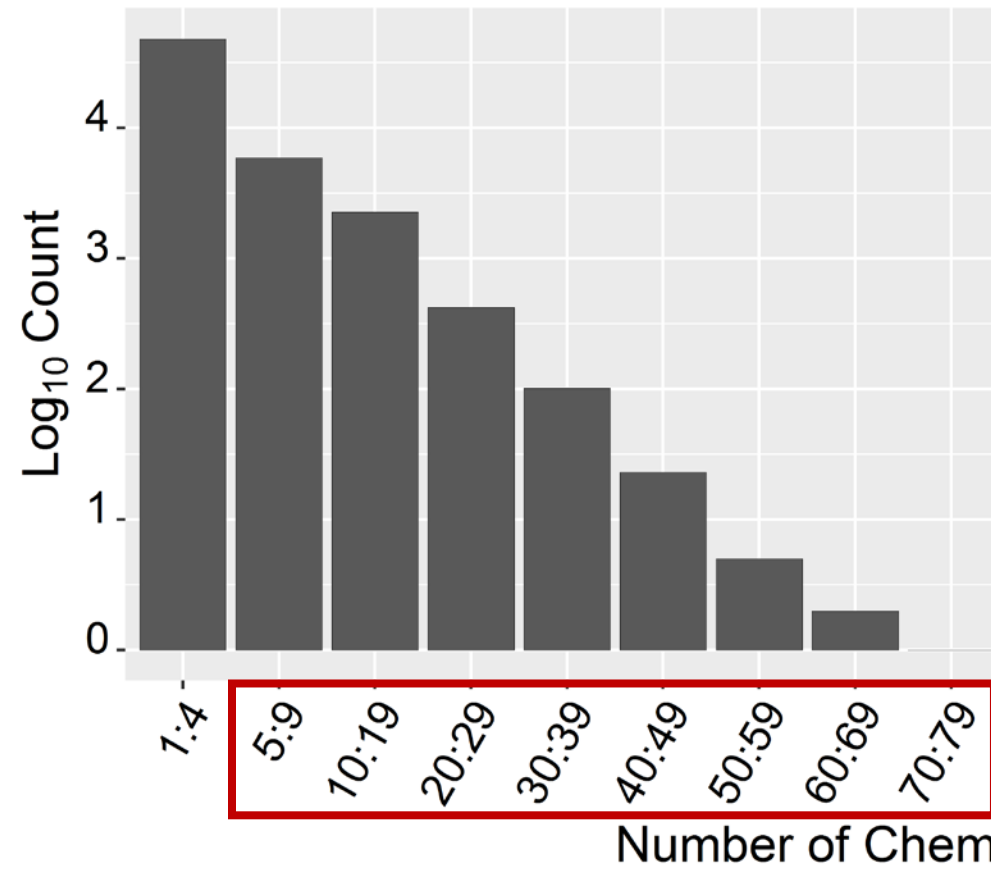
Number of chemicals in the regression: 76



**1 of ~1,000 assay
endpoints**

Scope of the Data

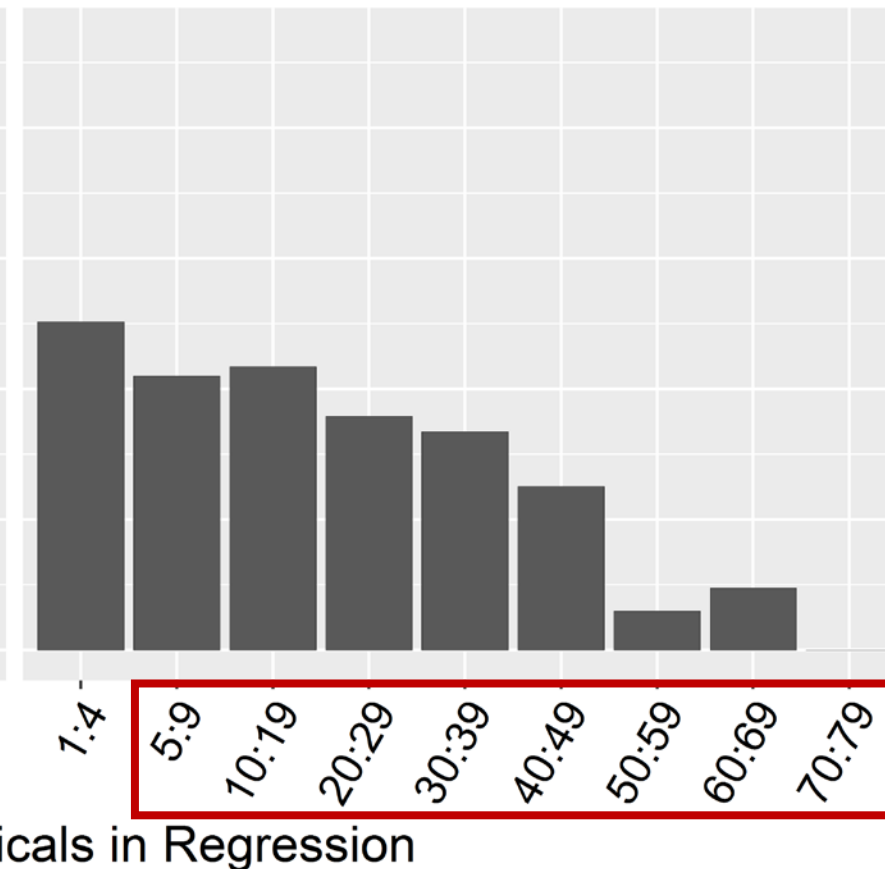
ToxRef Effect Level Dose



ToxRef Analysis:

- Total: 56,973 *in vitro* assay-*in vivo* effect pairs
- ≥ 5 chemicals: 8,731 *in vitro* assay-*in vivo* effect pairs
- ≥ 20 chemicals: 552 *in vitro* assay-*in vivo* effect pairs

ToxVal POD (1 POD per chem. across all effects)



ToxVal Analysis:

- Total: 734 *in vitro* assays
- ≥ 5 chemicals: 405 *in vitro* assays
- ≥ 20 chemicals: 133 *in vitro* assays

Evaluated Assumptions in Applying TK for IVIVE

Hepatic Clearance – Restrictive vs. Nonrestrictive

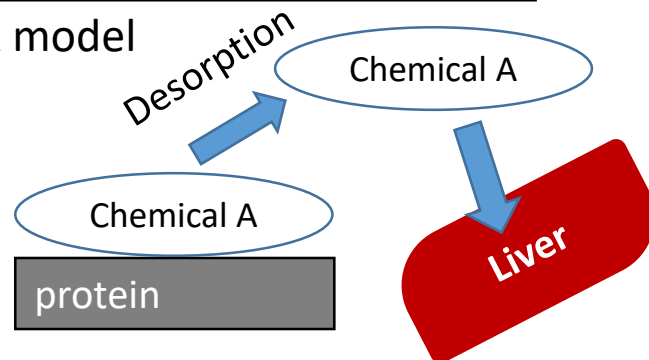
An assumption in the PBTK model

Restrictive Clearance

- **Slow** desorption rate
- Clearance $\propto f_{up}$

Non-Restrictive Clearance

- **Fast** desorption rate
- Clearance independent of f_{up}



Concentration Selection from the PBTK model

- C_{PBTK}
- Plasma vs tissue concentration (matched to cell type of cell based assay)
- Mean vs max concentration

Multiplication or Division by f_{up}

Multiplication: $C_{PBTK} * f_{up}$

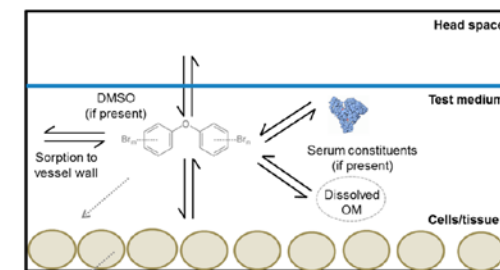
- Assume AC50 is closer to a free concentration

Division: $C_{PBTK} * f_{up}^{-1}$

- Assume AC50 is nominal concentration and convert to free concentration

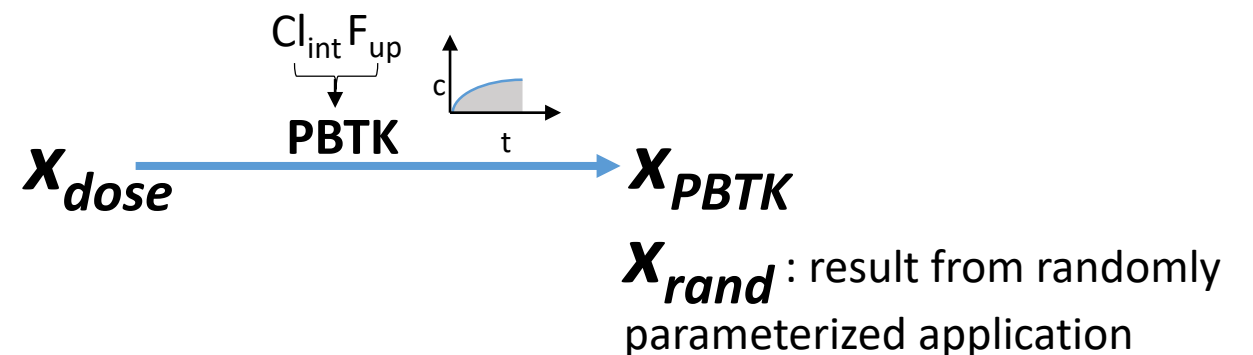
Partition model for cell based assays

- Nominal concentration (AC50) doesn't account for partitioning of chemical into serum and cells
- Determine free concentration from Armitage model
- Account for free concentration by defining a factor $\eta = \frac{C_{assay,free}}{C_{assay,nominal}}$ so that $x_{PBTK} = C_{PBTK}/\eta$



Armitage, J. M.; Wania, F.; Arnot, J. A. *Env. Sci. & Tech.* 2014, 49, 9770-9779.

Methods Summary



All assays:

- 8731 *in vitro* assay- *in vivo* effect pairs (analysis of ToxRef ELD)
- 405 *in vitro* assays (analysis of ToxVal POD)
- 24 combinations of assumptions to evaluate
- ~219,000 regressions (4 regressions per comparison)

Simple Regressions

$$AC50 \sim \beta_{i,1} x_{PBTk} + \gamma$$

$$AC50 \sim \beta_{i,2} x_{rand} + \gamma$$

$$AC50 \sim \beta_{i,3} x_{dose} + \gamma$$

Multiple Regression

$$AC50 \sim \beta_{j,1} x_{PBTk} + \beta_{j,2} x_{rand} + \beta_{j,3} x_{dose} + \gamma$$

Cell based assays: nominal vs free concentration (Armitage)

- 1531 *in vitro* assay- *in vivo* effect pairs(ToxRef ELD) with known FBS%
- 78 *in vitro* assays (ToxVal POD) with known FBS%
- an additional 24 combinations of assumptions to evaluate
- ~38,000 additional regressions

Clearance	f_{up} multiplier	Concentration	Concentration value
restrictive	none	plasma	mean
			max
		tissue	mean
			max
	f_{up}	plasma	mean
			max
non-restrictive	f_{up}	tissue	mean
			max
	f_{up}^{-1}	plasma	mean
			max
		tissue	mean
			max
non-restrictive	none	plasma	mean
			max
		tissue	mean
			max
	f_{up}	plasma	mean
			max
non-restrictive	f_{up}	tissue	mean
			max
	f_{up}^{-1}	plasma	mean
			max
		tissue	mean
			max

Example Results

Example Regressions – ToxRef ELD

in vivo Effect: systemic-nonneoplastic pathology-liver

Assumptions

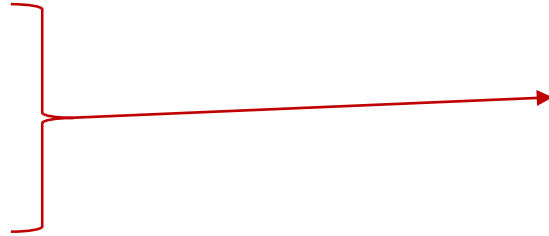
Clearance	f_{up} multiplier	Concentration	Concentration Value
nonrestrictive	none	plasma	mean

Simple Regressions

$$AC50 \sim \beta_{i,1} x_{PBTk} + \gamma$$

$$AC50 \sim \beta_{i,2} x_{rand} + \gamma$$

$$AC50 \sim \beta_{i,3} x_{dose} + \gamma$$

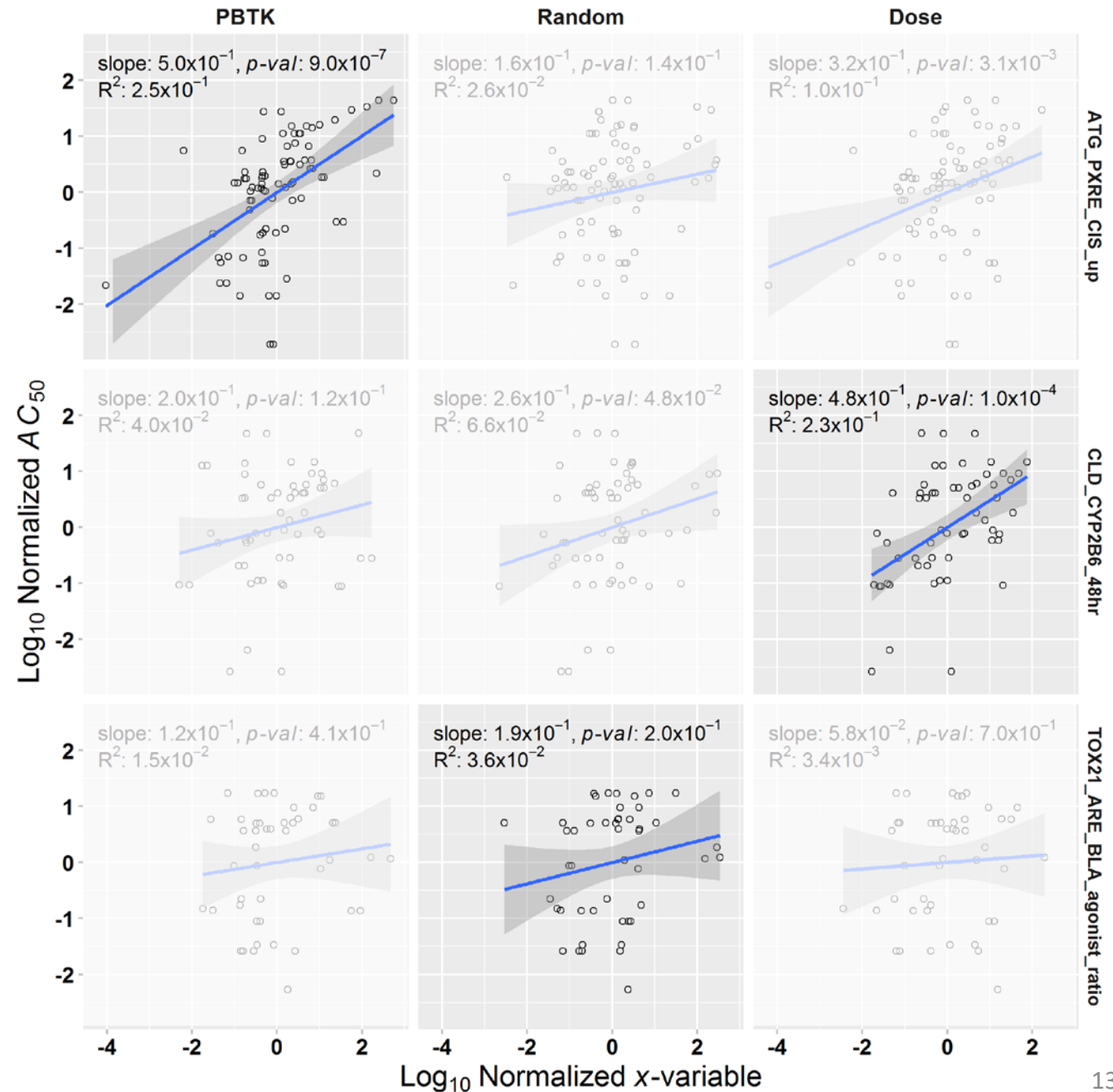


Multiple Regression:

$$AC50 \sim \beta_{j,1} x_{PBTk} + \beta_{j,2} x_{rand} + \beta_{j,3} x_{dose} + \gamma$$

Assay	Slopes			p-values		
	PBTk	Random	Dose	PBTk	Random	Dose
ATG_PXRE_CIS_up	0.50	-0.05	0.04	1.1E-04	6.8E-01	8.1E-01
CLD_CYP2B6_48_hr	0.06	-0.01	0.47	6.5E-01	9.7E-01	2.2E-03
TOX21_ARE_BLA_agonist_ratio	0.09	0.22	-0.10	5.9E-01	2.4E-01	5.8E-01

Filter based on result of multiple regression



Example Regressions – ToxVal POD

Clearance	f_{up} multiplier	Concentration	Concentration Value
nonrestrictive	none	plasma	mean

Simple Regressions

$$AC50 \sim \beta_{i,1} x_{PBTk} + \gamma$$

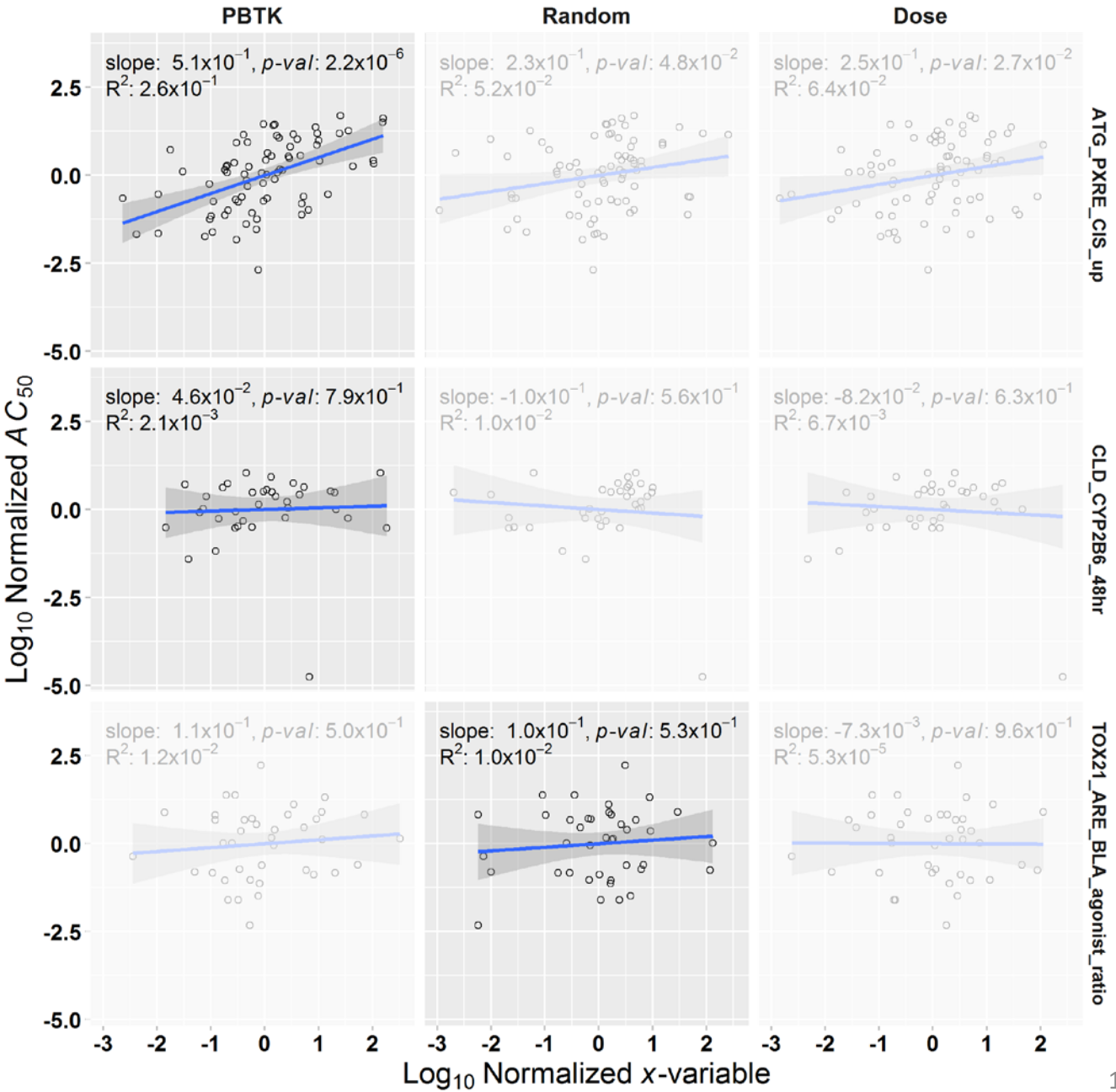
Summarize by looking at histograms of slope, p-value, and R² for every pair of **ToxRef Effect** and **assay endpoint** and every assay endpoint with corresponding **ToxVal POD**

Multiple Regression:

$$AC50 \sim \beta_{j,1} x_{PBTk} + \beta_{j,2} x_{rand} + \beta_{j,3} x_{dose} + \gamma$$

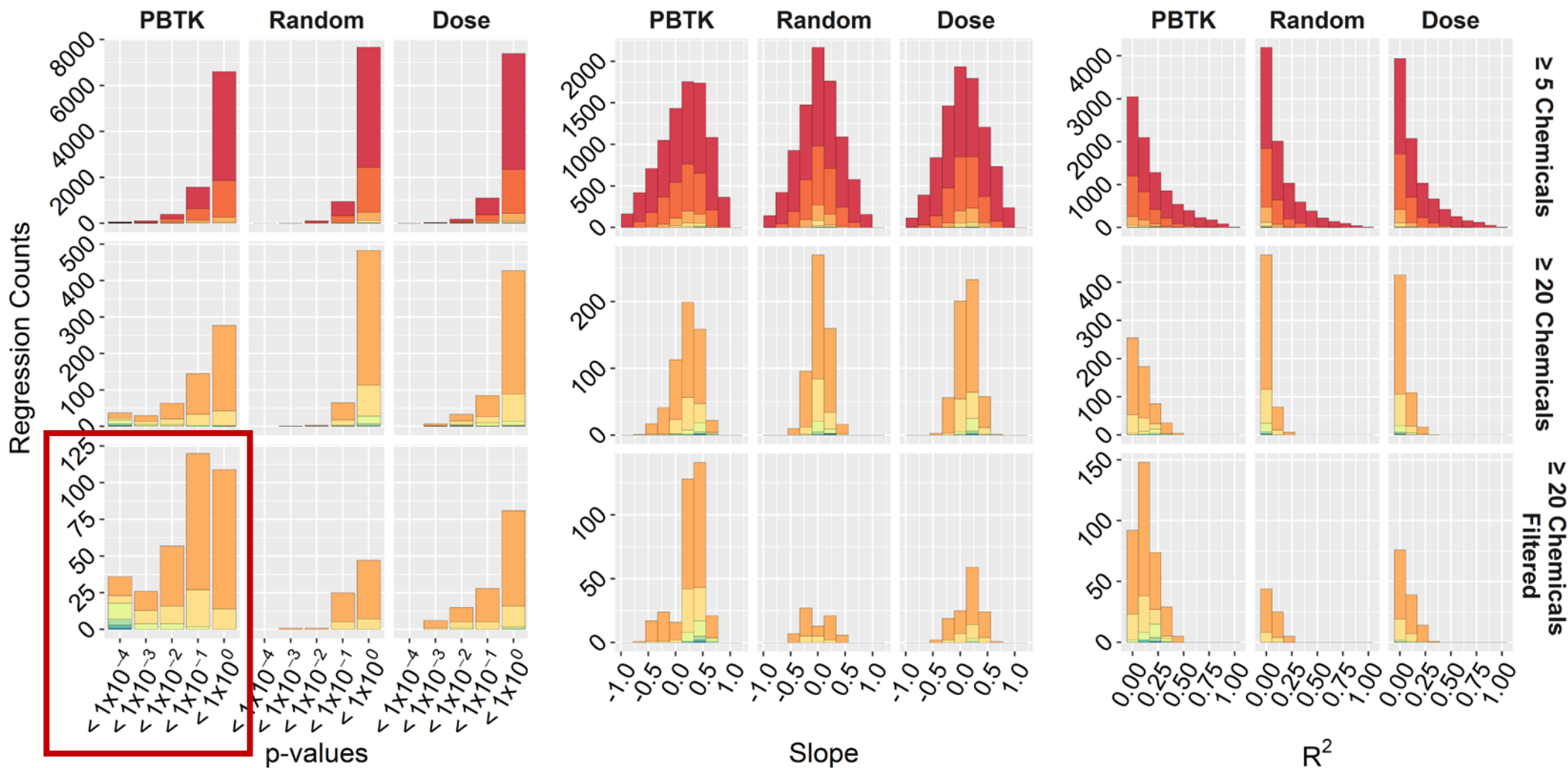
Assay	Slopes			p-values		
	PBTk	Random	Dose	PBTk	Random	Dose
ATG_PXRE_CIS_up	0.61	0.17	-0.24	1.3E-05	2.1E-01	1.3E-01
CLD_CYP2B6_48_hr	0.11	-0.05	-0.10	6.2E-01	8.1E-01	6.8E-01
TOX21_ARE_BLA_agonist_ratio	0.19	0.20	-0.25	3.5E-01	3.6E-01	3.2E-01

Filter based on result of multiple regression



ToxRef ELD Example Result

Clearance	f_{up} multiplier	Concentration	Concentration Value
nonrestrictive	none	plasma	mean



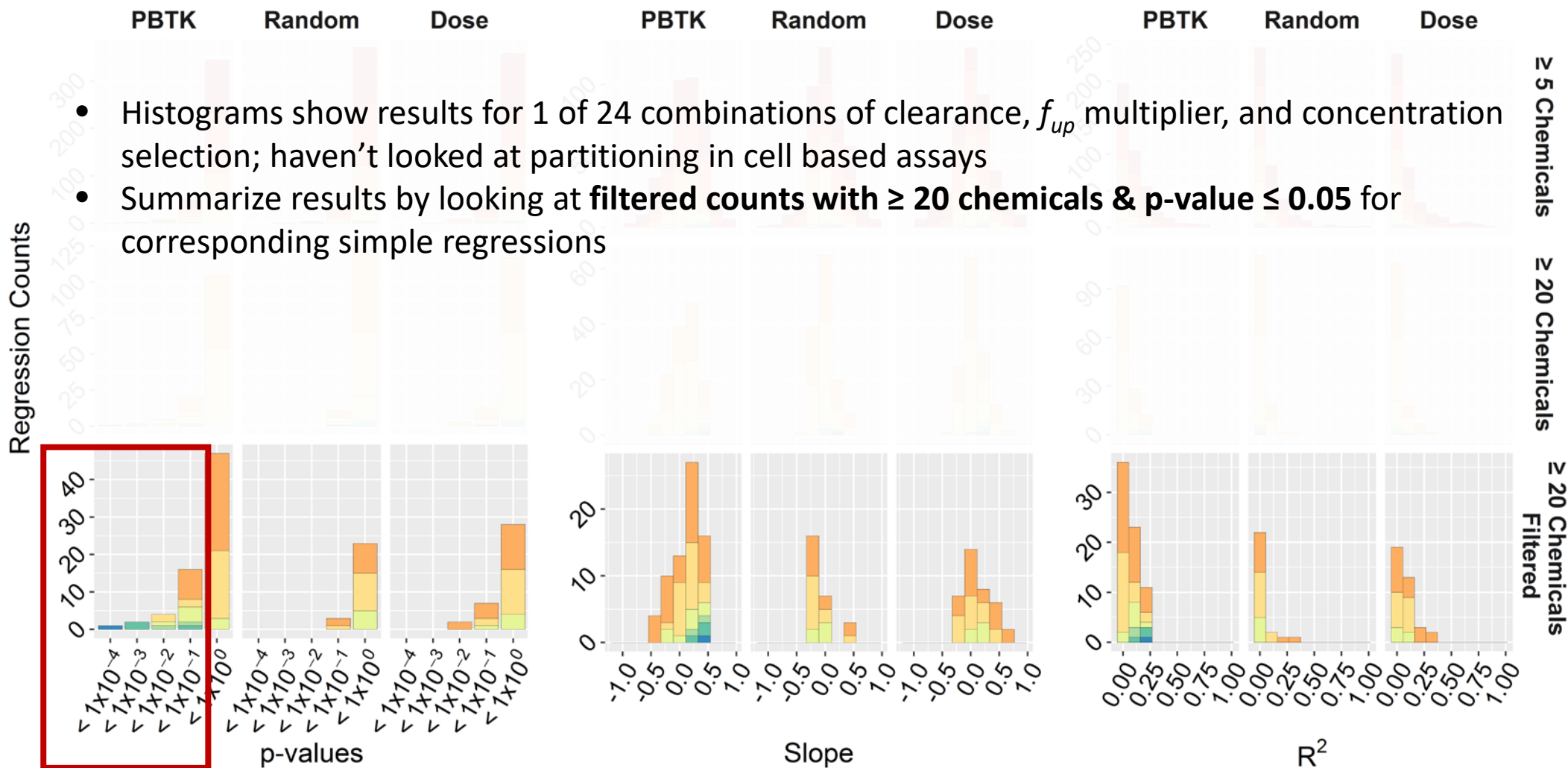
PBTK gives best result

Number of Chemicals

5:9	20:29	40:49	60:69
10:19	30:39	50:59	70:79

ToxVal POD Example Result

Clearance	f_{up} multiplier	Concentration	Concentration Value
nonrestrictive	none	plasma	mean



PBTK gives best result



Filtered counts with p-value ≤ 0.05 and # of chemicals ≥ 20

All Assays

ToxRef ELD

ToxVal POD

Clearance	f_{up} multiplier	Concentration	Concentration value	PBTK	Random	Dose	PBTK	Random	Dose
restrictive	none	plasma	mean	108	23	50	2	6	7
			max	101	20	39	4	5	4
		tissue	mean	94	19	59	3	3	7
			max	75	17	54	5	2	6
	f_{up}	plasma	mean	189	22	30	9	4	5
			max	183	20	28	13	2	5
		tissue	mean	145	24	48	5	5	8
			max	133	25	39	7	3	5
	f_{up}^{-1}	plasma	mean	40	17	77	1	5	9
			max	38	26	72	2	5	8
		tissue	mean	50	25	77	4	6	8
			max	35	25	76	2	5	8
non-restrictive	none	plasma	mean	214	19	34	15	2	6
			max	210	19	29	15	2	6
		tissue	mean	198	16	36	11	4	8
			max	206	14	29	11	2	8
	f_{up}	plasma	mean	213	12	34	19	2	6
			max	224	15	28	20	2	6
		tissue	mean	187	21	38	15	2	8
			max	192	20	34	14	2	8
	f_{up}^{-1}	plasma	mean	141	29	53	7	5	7
			max	124	25	56	5	4	7
		tissue	mean	124	24	51	6	6	7
			max	106	25	52	6	4	7

Cell Based Assays

nonrestrictive clearance, no f_{up} multiplier,
mean plasma concentration

Cell Assay Model	ToxRef			ToxVal		
	PBTK	Random	Dose	PBTK	Random	Dose
None	22	7	0	1	0	2
Armitage	30	2	1	3	1	0

- **Nonrestrictive** clearance appears to give better result
- Using a f_{up} multiplier has an effect
- **Plasma** slightly better than tissue concentration
- No significant difference between using mean or max concentration
- Using a cell assay partition model appears to improve results for cell based assays

ToxRef: total pairs ≥ 20 chemicals = 552

ToxVal: assays ≥ 20 chemicals = 133

Conclusions

- In general, **using the PBTK model improves the correlation between the evaluated *in vitro* and *in vivo* toxicity data**
- In some cases, untransformed dose remains a better predictor
- The assumptions in the application of the PBTK model for IVIVE matter
- **Nonrestrictive clearance** with **plasma concentration** gave the best result and should be used as a starting basis when applying PBTK for IVIVE, using the predicted *in vivo* unbound concentration (multiplying by f_{up}) may be beneficial
- Although some correlations were significant, they were not predictive on their own
- Toxicokinetics should be considered when building ensemble models to relate *in vitro* toxicity assay results to particular *in vivo* endpoints – *but only had data for 104 chemicals in this analysis*

Acknowledgements

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