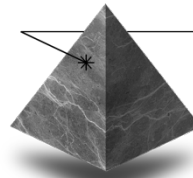


Application of a Multiplexed Flow Cytometric Assay and Machine Learning to Provide Genotoxic Mode of Action Information



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**Litron
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****This presentation contains unpublished data – please do not distribute****



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Today's Theme

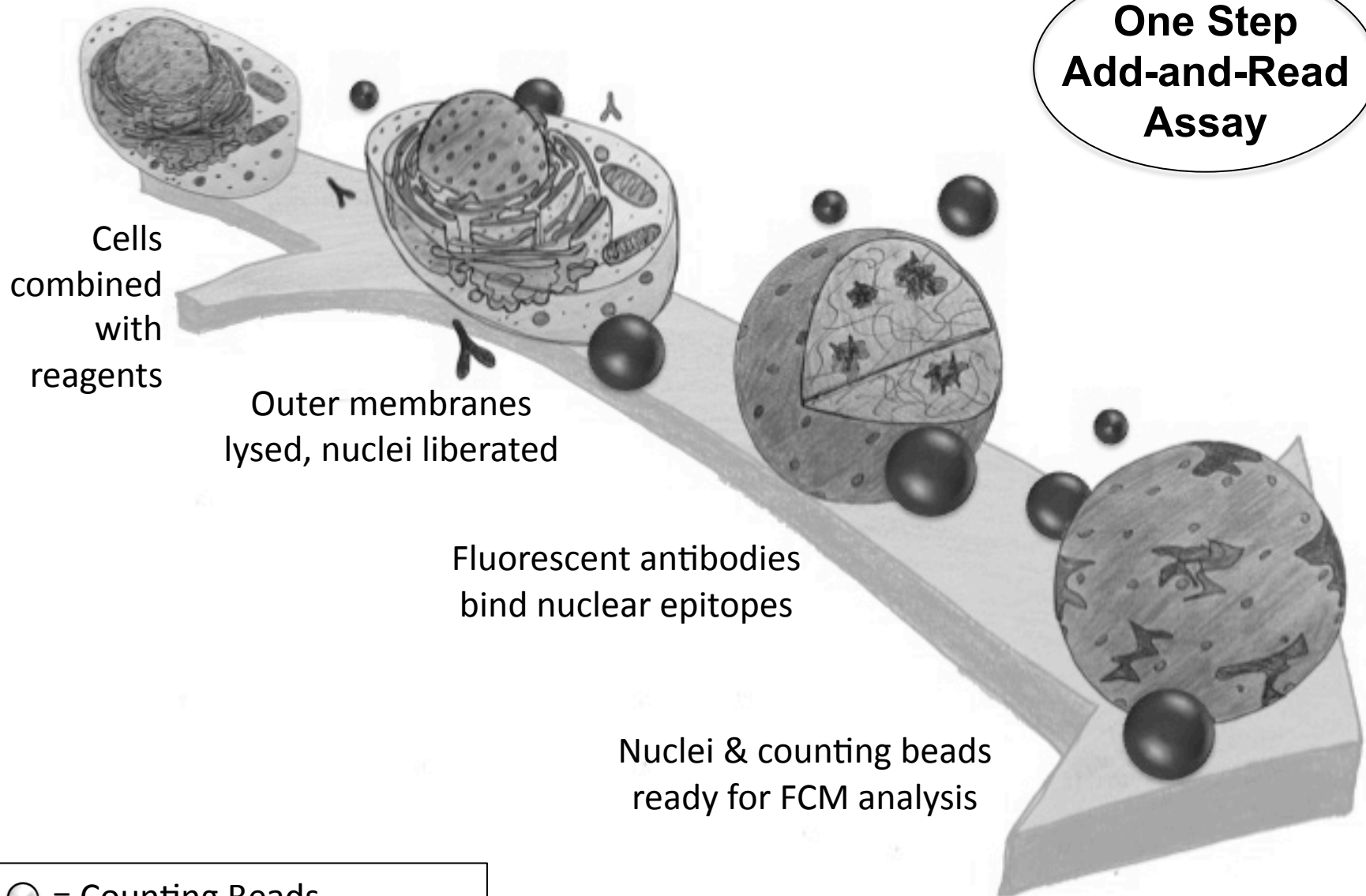
- More efficient, higher information content assays...
 - improve compound screening
 - complement regulatory-required assays
 - expand range of information

Coming soon!



MultiFlow™ Method

**One Step
Add-and-Read
Assay**



○ = Counting Beads

MultiFlow™ Reagents

REAGENT	PURPOSE
Detergent	Liberates nuclei for analysis
Nucleic acid dye + RNase	Cell cycle and polyploidization
Latex microspheres at a known density	Facilitates absolute nuclei counts, a measure of cytotoxicity
Anti-phospho-H3-PE	Metaphase cell marker
Anti- γ H2AX-Alexa 647	DNA double strand break marker
Anti-p53-FITC (human only)	Genotoxicity marker, specific to nuclear translocated target
Anti-Cleaved PARP	Apoptosis

*

* combination kits

single Ab kits ←

Case Study:

**H2AX, Phospho-Histone H3, p53 kit -
to determine genotoxic Mode of Action**

Bryce SM, Bernacki DT, Bemis JC, Dertinger SD. Genotoxic mode of action predictions from a multiplexed flow cytometric assay and a machine learning approach. *Environ Mol Mutagen.* 2016 Apr;57(3):171-89.

Definitions

- **Clastogens**

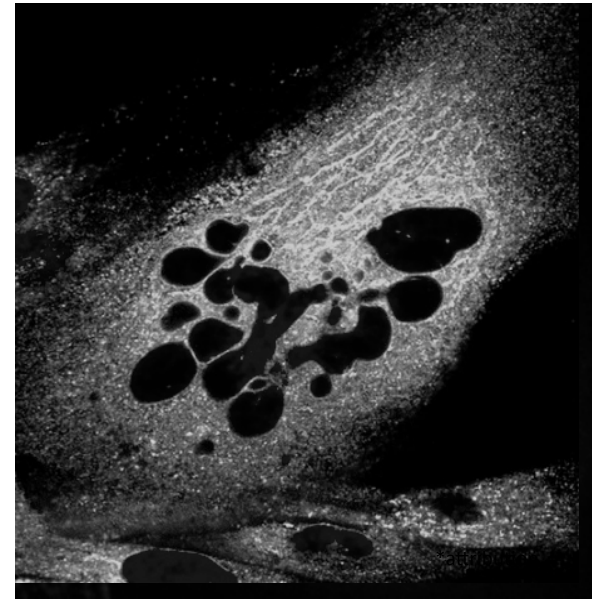
- DNA double strand breaks

- **Aneugens**

- malsegregation of chromosome(s)
- polyploidy

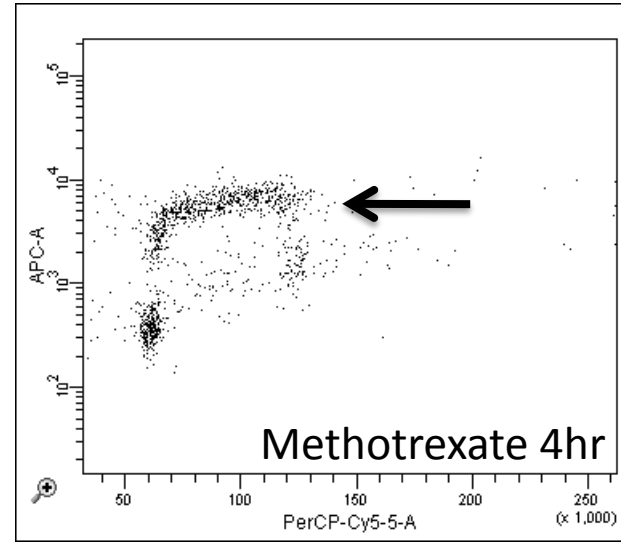
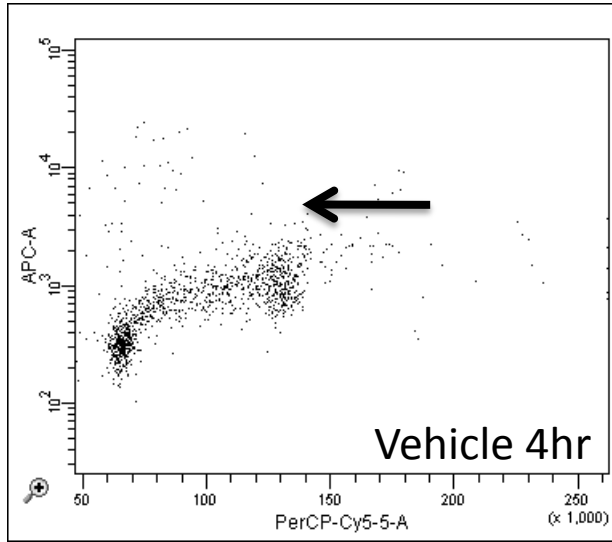
- **Non-genotoxicants**

- may be cytotoxic
- may damage DNA through secondary pathways (apoptosis, damage to lysosomes, etc)

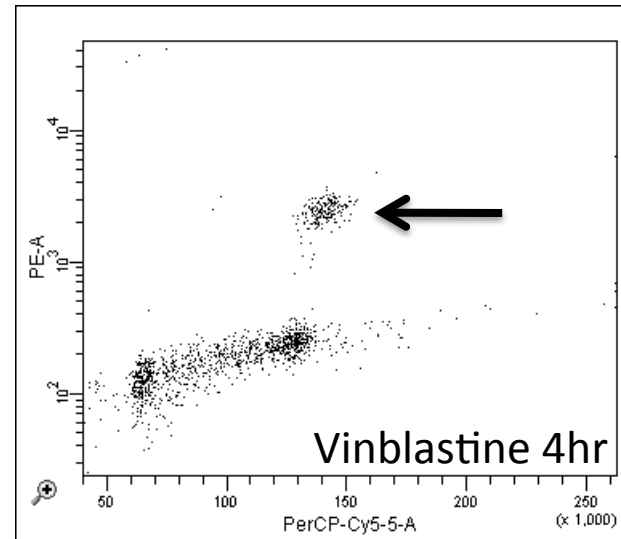
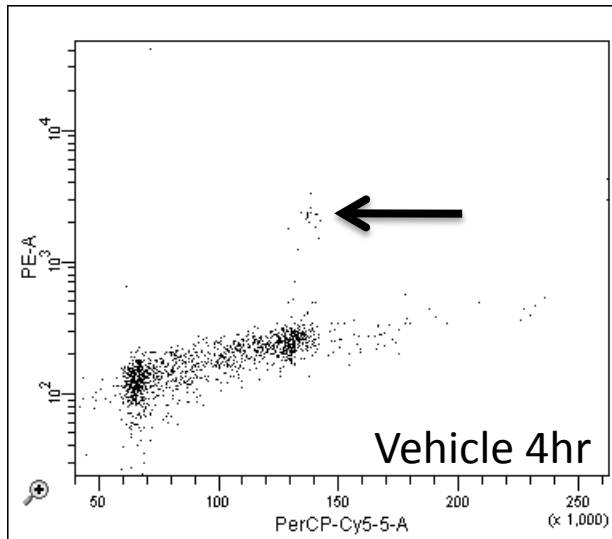


Example flow plots

γ H2AX



Phospho
-histone
H3

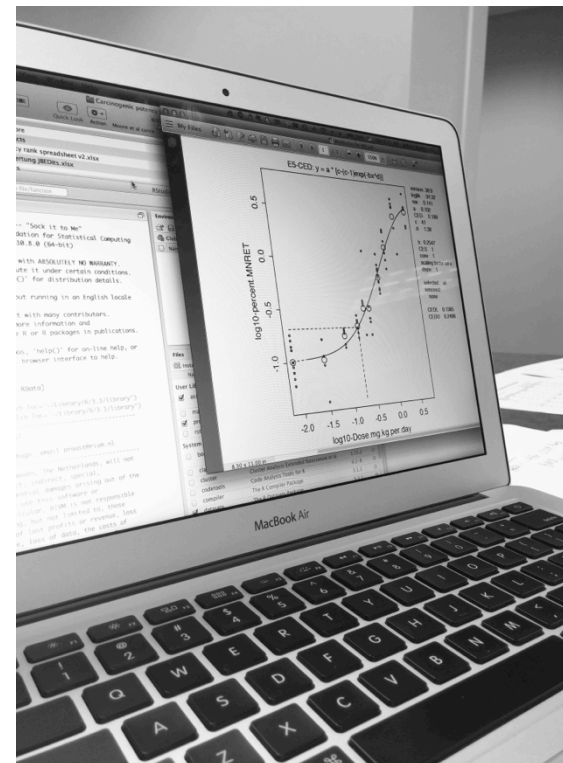
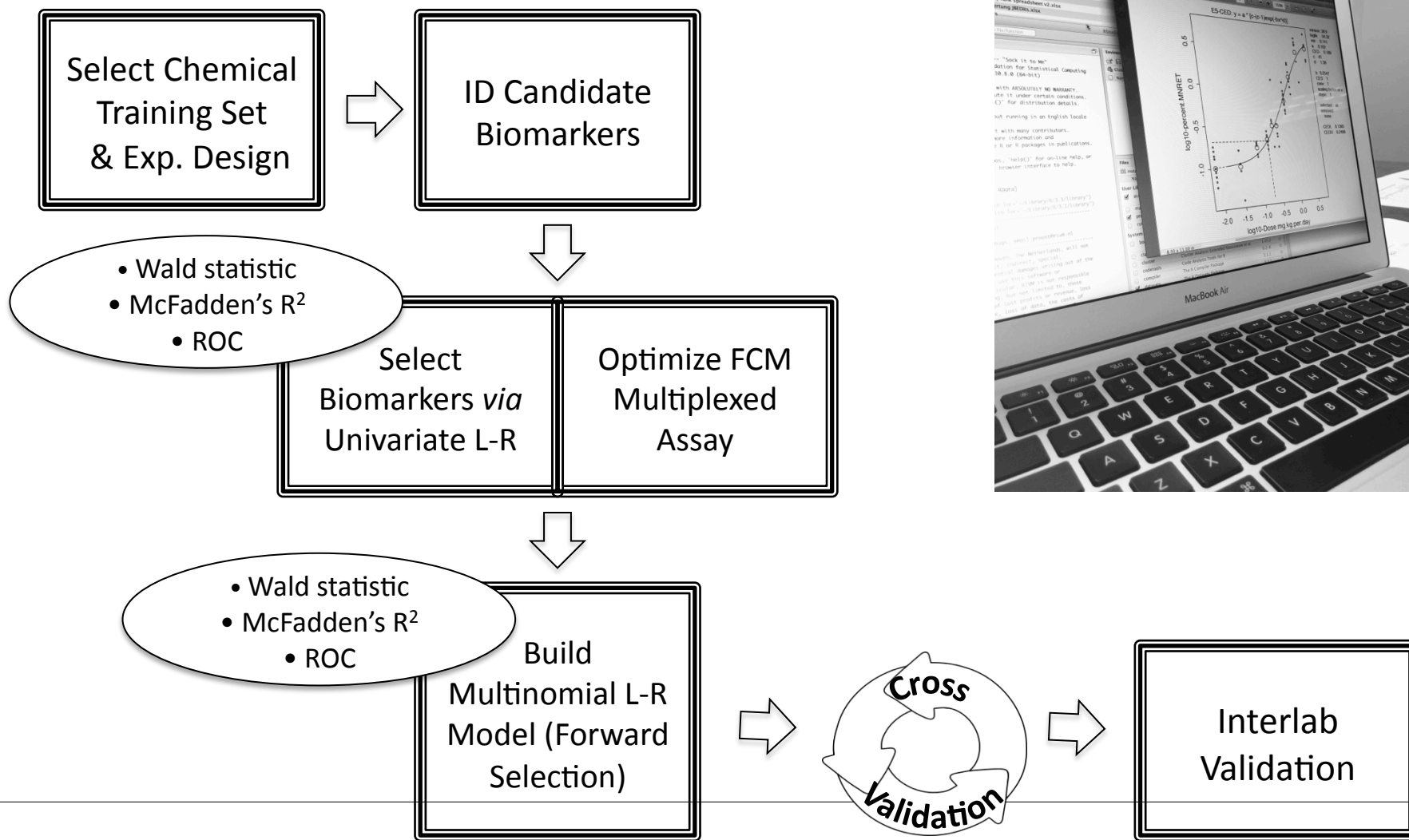


Experimental Design

- 79 chemicals from ECVAM published list and expert consensus
 - 14 aneugens, 27 clastogens, and 38 non-genotoxicants
- Treated TK6 cells in 96 well plates; solvent usually DMSO; top concentration 1 mM
 - extreme cytotoxicity or precipitation dictated some top conc.
- Up to 20 closely spaced concentrations (Sqr2), no replicate wells
- Concurrent vehicle and positive controls
- 24 hrs continuous exposure
- Study an early and late time point: 4 & 24 hr

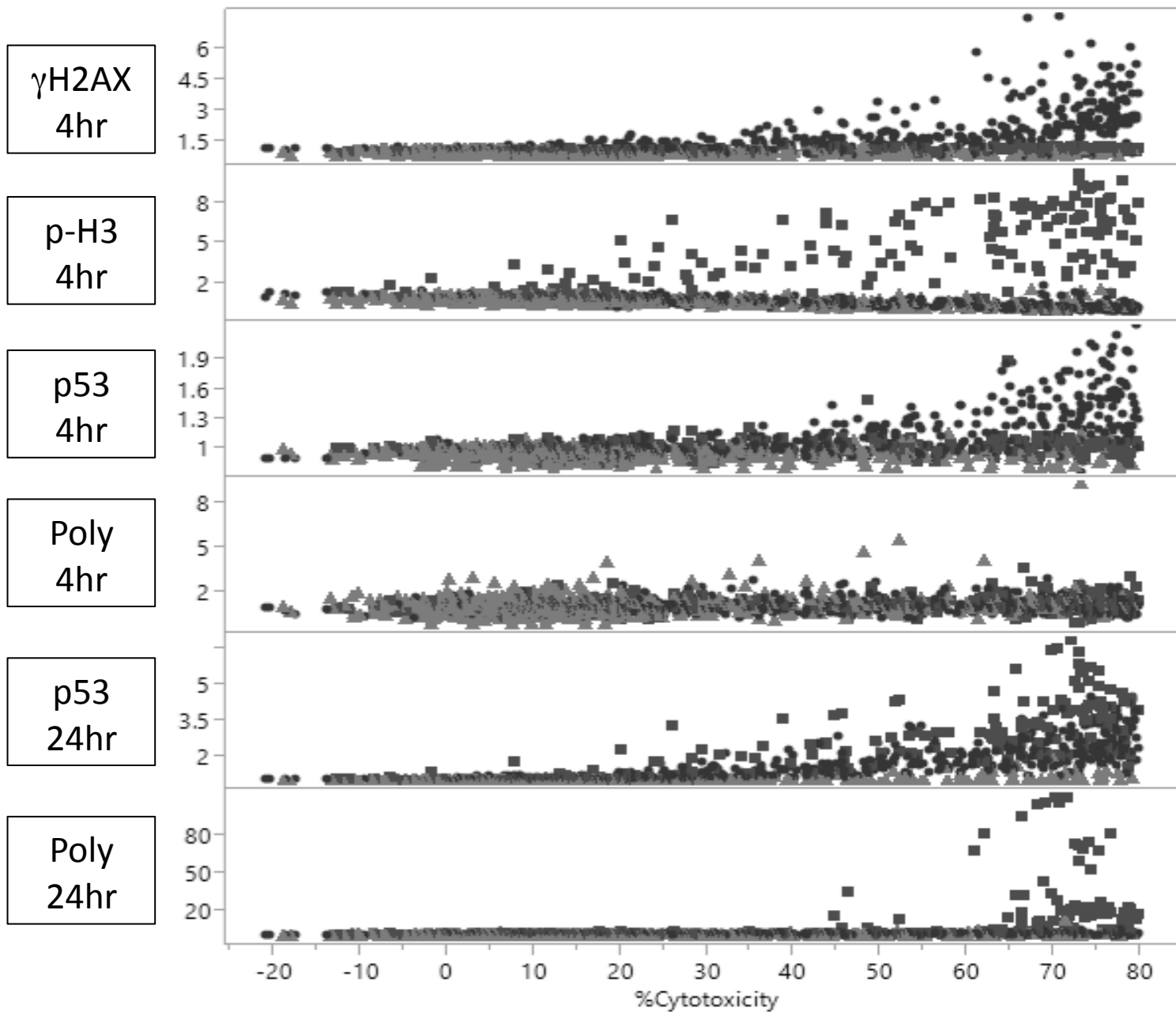


Overview



○ Clastogens □ Aneugens △ Non-Genotoxicants

Valuable
for Class
Discrimination?



YES

YES

YES/NO

NO

YES

YES

Univariate L-R Analyses

Based on Wald statistic, McFadden's R^2 and ROC values

- 4 and 24 hr γ H2AX are predictive of class
- 4 and 24 hr p-H3 are predictive of class
- 24 hr p53 is predictive of class
- 24 hr polyploidy is predictive of class
- Note: why provisionally 4 hrs when both 4 & 24 hr are predictive of class? There may be theoretical advantages to studying early effects that are not confounded by secondary processes

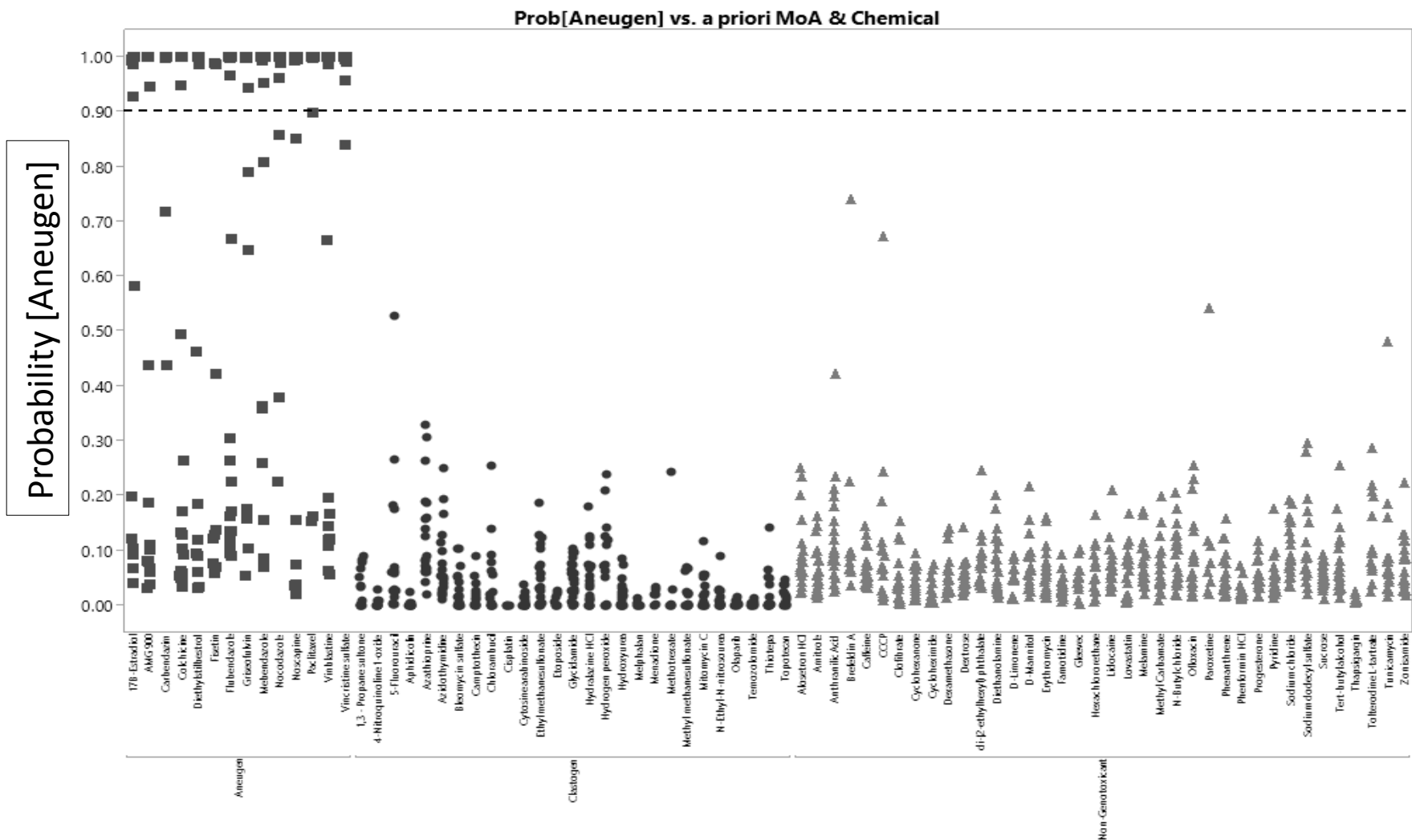
Summary Data

<u>Endpoints</u>	<u>Wald</u>	<u>Pseudo R²</u>	<u>ROC</u>
γ H2AX (4hr)	< 0.0001	0.2442	≥ 0.6464
p-H3 (4hr)	< 0.0001	0.2854	≥ 0.7572
p53 (24hr)	< 0.0001	0.3021	≥ 0.7181
Polyploidy (24hr)	< 0.0001	0.0960	≥ 0.5264
All 4 endpoints	≤ 0.0020	0.5502	≥ 0.8944
All 4 weighted on cytotoxicity	< 0.0001	0.7459	≥ 0.9596

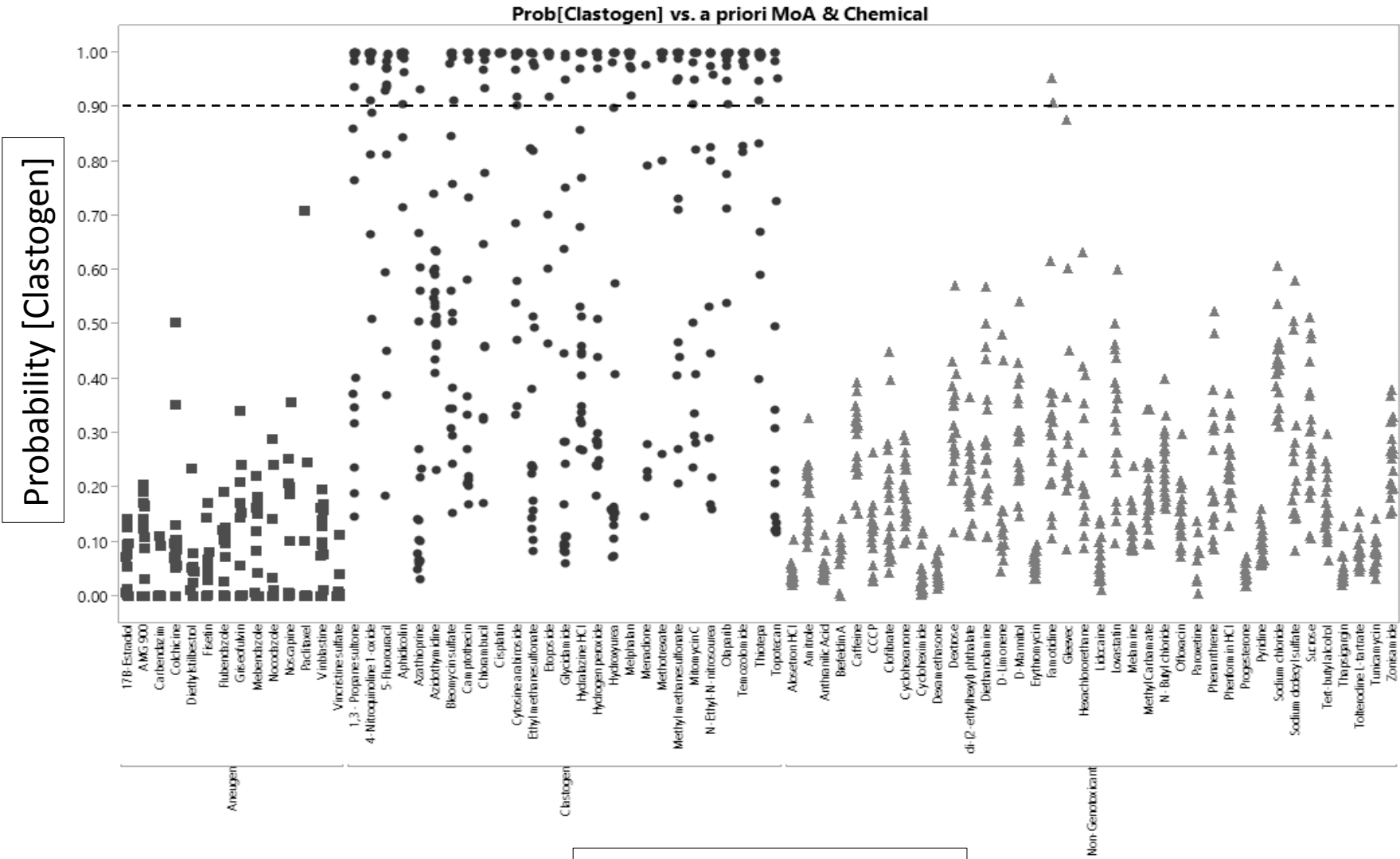
Synthesizing L-R Model Output

- Since L-R model is using data from every conc. the output consists of multiple probability scores—one for each conc.
- For a clastogen or aneugen call we require two consecutive concentrations to show $\geq 90\%$ probability
- Non-genotoxicant call = no concentration showing $\geq 90\%$ probability clastogen or aneugen
- Equivocal = only one concentration showing $\geq 90\%$ probability of clastogen or aneugen

79 Chemical L-R Model (Aneugen Probability vs Chemical)



79 Chemical L-R Model (Clastogen Probability vs Chemical)



Performance of 4-Factor Model

- 79 chemicals, 75 correct, 2 equivocal
- 14/14 aneugens correctly classified
- 24/27 clastogens correctly classified
 - Misidentified AZT; azathioprine & menadione equivocal
- 37/38 non-genotoxicants correctly classified
 - Misidentified famotidine
- If we consider equivocal calls to be misclassifications, then concordance with *a priori* classification is 94.9%

Cross-Validation: Leave-One-Out

- Model generated with each chemical in turn left out of training set; make class prediction for hold-out
- 79 chemicals, 72 correct, 2 equivocal
- If we consider equivocal calls to be misclassifications, then concordance with a *priori* classification is 91.1%
- High concordance for L-O-O suggests model is not “overfitting” data and that it will be “generalizable” (meaning applicable to chemicals outside of training set)

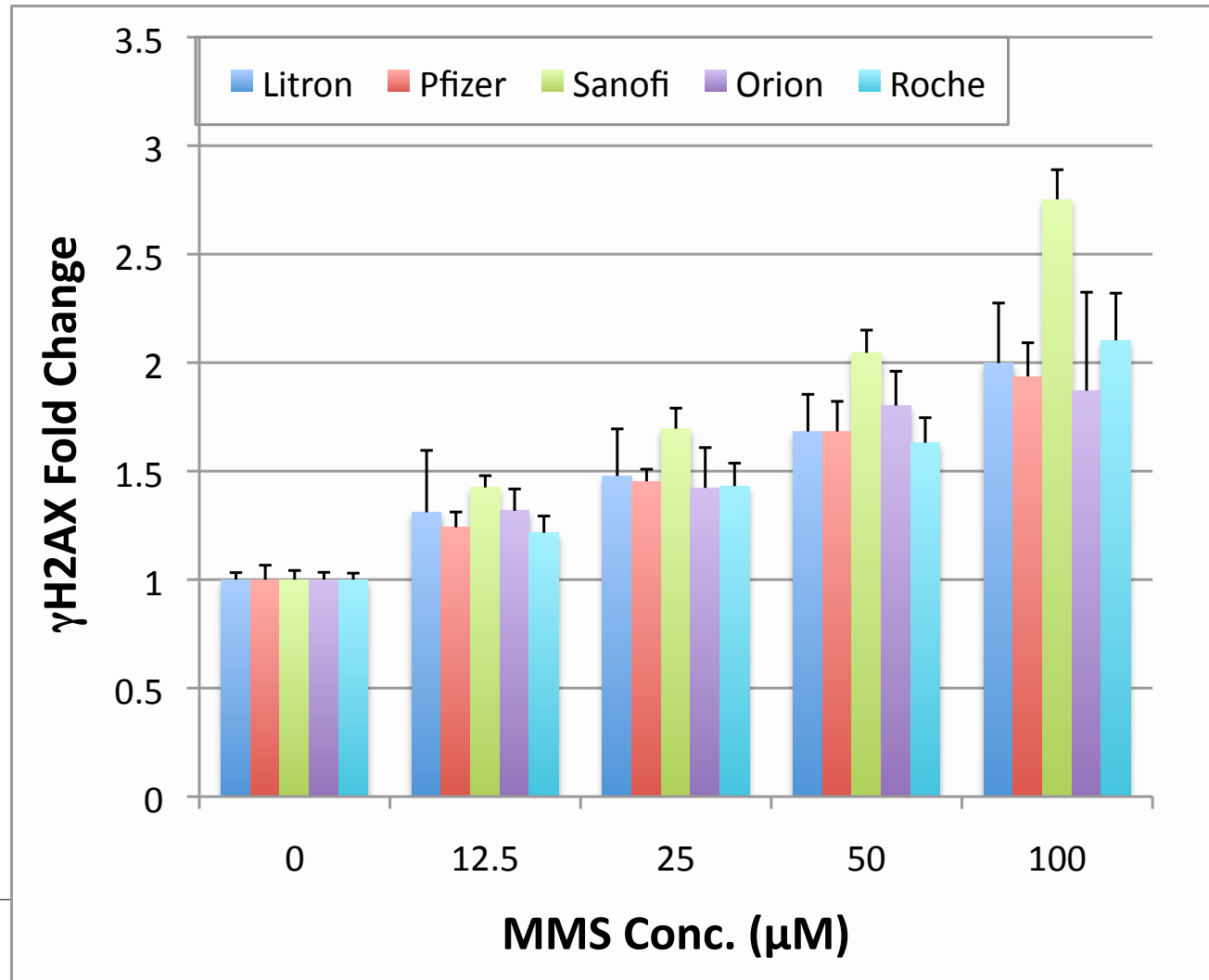
Interlaboratory Transferability

- Six external collaborating labs received kit-formatted reagents and instructions
- TK6 cells, continuous exposure
- First question—do reference agents elicit expected responses?



Interlab Transferability: γ H2AX Example

(only showing labs with $n \geq 4$)

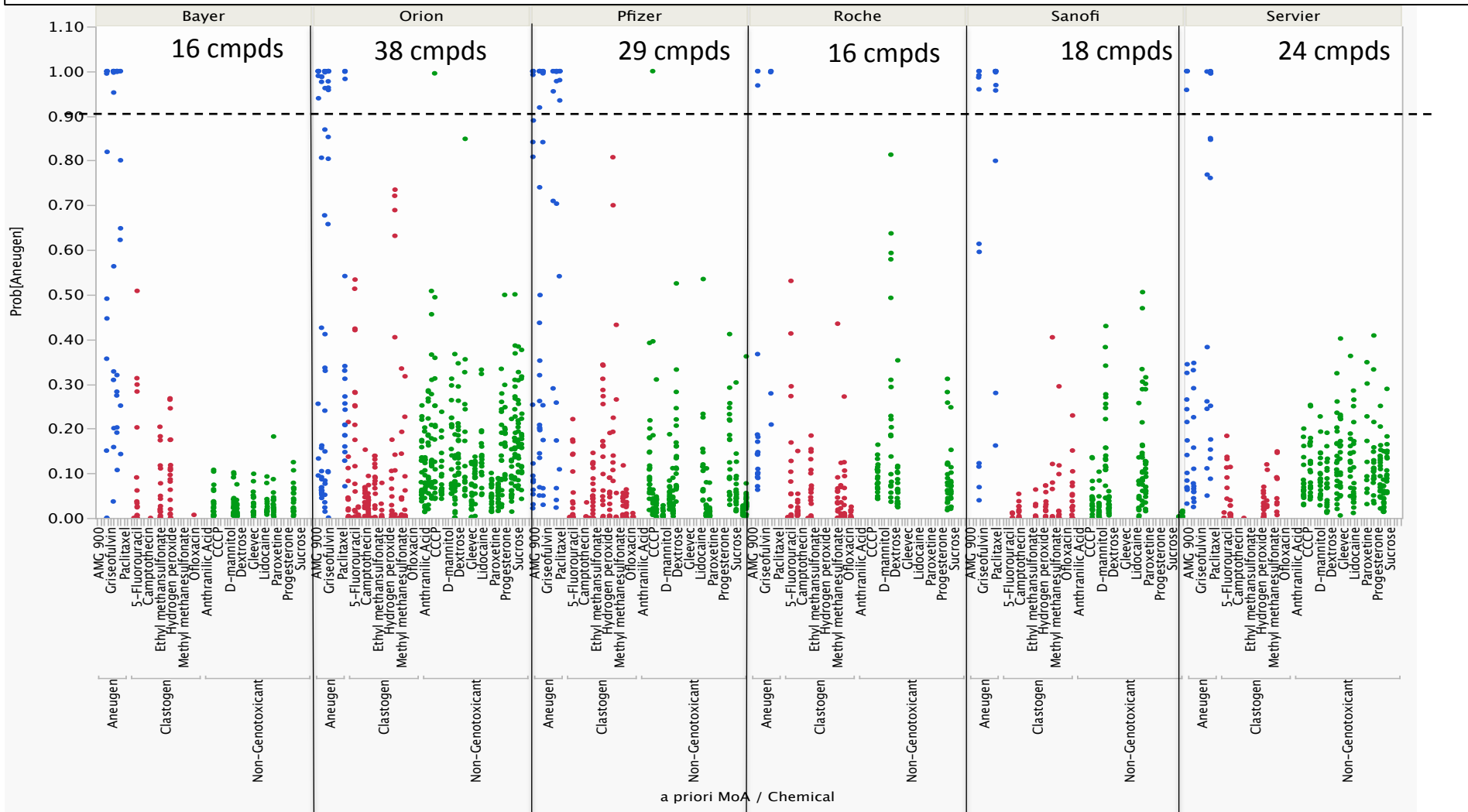


Interlaboratory Transferability

- Six external collaborating labs received kit-formatted reagents and instructions
- TK6 cells, continuous exposure
- First question—do reference agents elicit expected responses?
- Second question—do data from external labs work with reference lab model?
 - Collaborators chose from a list of compounds previously studied at Litron
 - Raw data returned to Litron and evaluated by MoA algorithm

Interlab Transferability: Aneugenicity Predictions

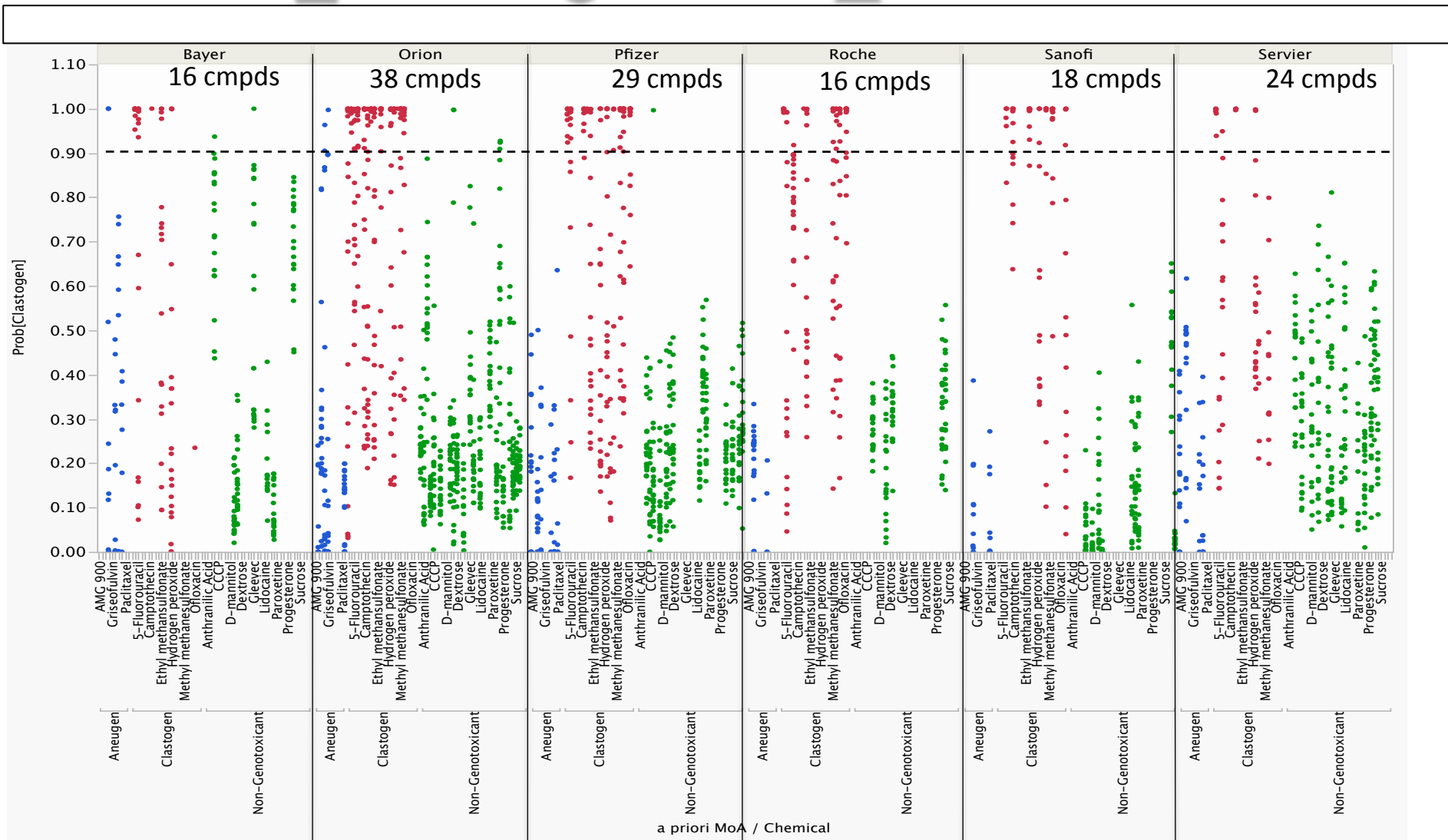
Aneugens
 Clastogens
 Non-Genotoxicants



Bayer	Orion	Pfizer	Roche	Sanofi	Servier
4/4 Aneugens	5/5 Aneugens	6/6 Aneugens	2/2 Aneugens	2/2 Aneugens	3/4 Aneugens

Interlab Transferability: Clastogenicity Predictions

Aneugens
 Clastogens
 Non-Genotoxicants



Bayer	Orion	Pfizer	Roche	Sanofi	Servier
5/5 Clastogens	13/13 Clastogens	12/12 Clastogens	8/9 Clastogens	7/7 Clastogens	5/7 Clastogens

Interlab Transferability: Summary

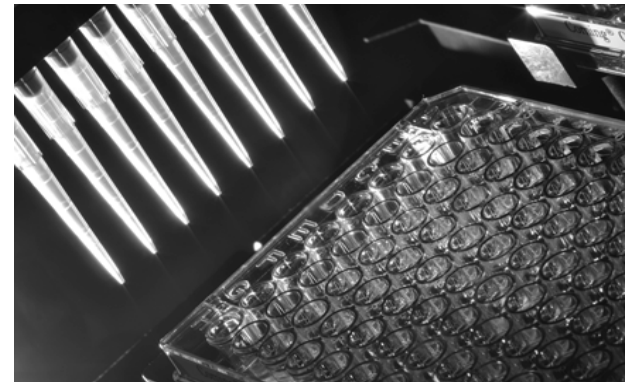
Lab	Total	Aneugens	Clastogens	Non-Genotoxicants	A Priori
Pfizer	29	6/6	12/12	9/11	93%
Sanofi	18	2/2	7/7	9/9	100%
Roche	16	2/2	8/9	5/5	93%
Orion	38	5/5	13/13	17/20	92%
Bayer	16	4/4	5/5	5/7	88%
Servier	24	3/4	5/7	13/13	88%

Final steps:

- collect and analyze all data
- explore methods of comparison
- publish/present outcomes

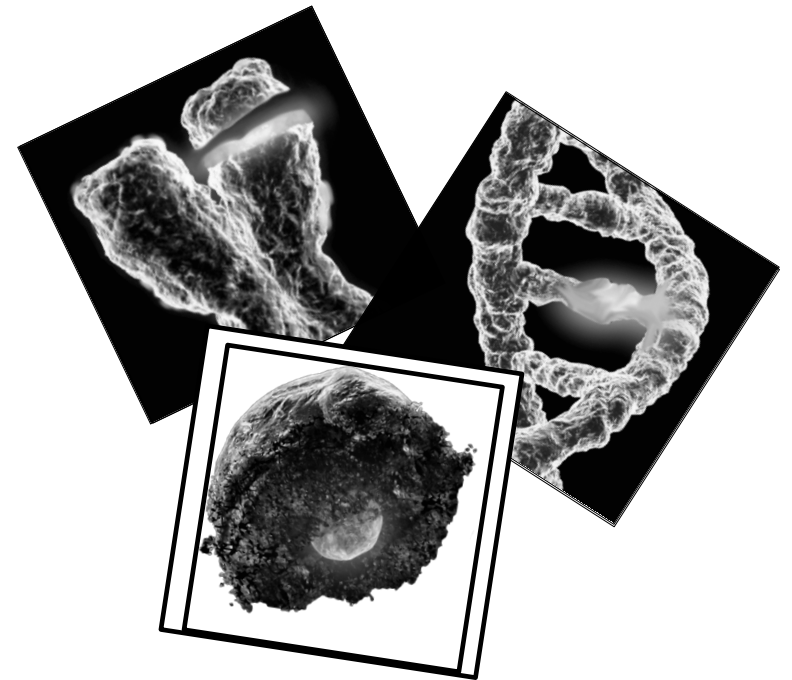
Conclusions

- Multiplexed genetox screening
 - reduced to a simple, one-step procedure
 - robust signals provides MoA classification
- Interlab transferability study
 - 6 external labs show excellent performance
- Future efforts
 - additional endpoints
 - expanded chemical sets
 - inter-lab studies



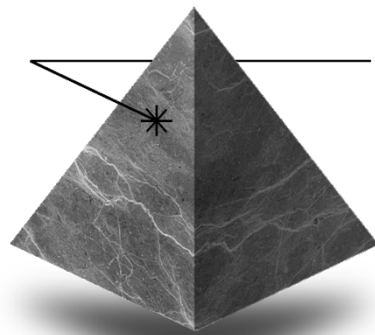
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 - SBIR grant # R44ES024039



Thank you!

- Questions – jbemis@litronlabs.com



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