



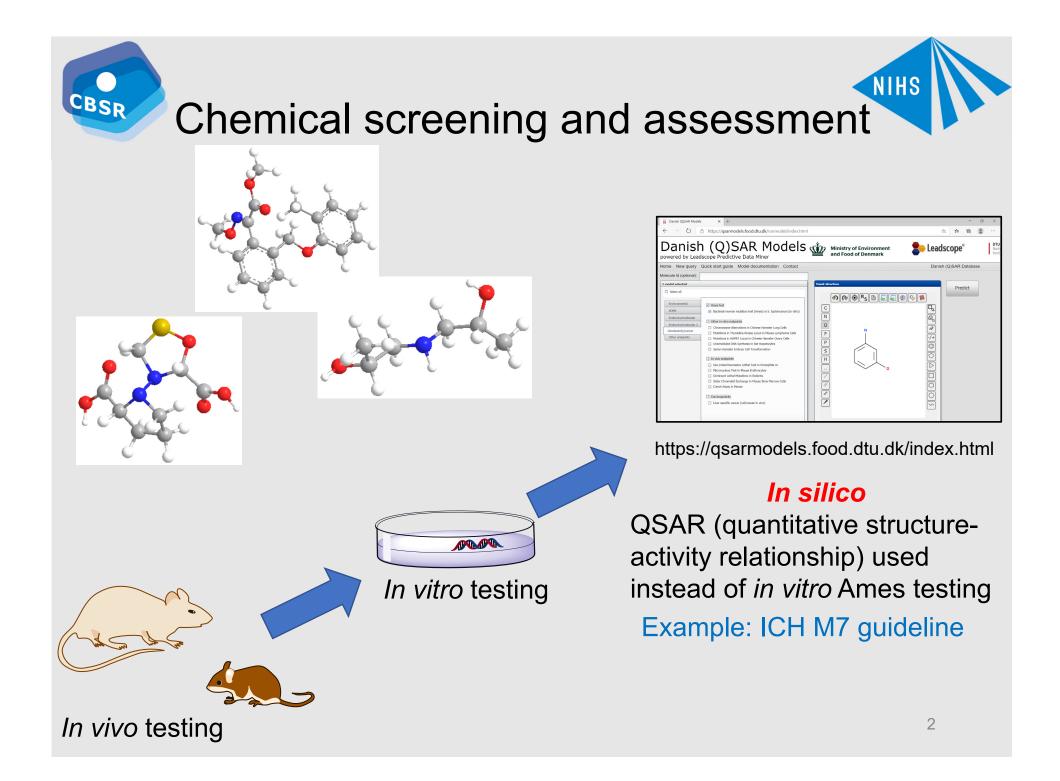
Bigdata Analysis: Outcome of the 2nd AMES/QSAR International Challenge Project

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The opinions in this presentation are my own and do not necessarily reflect the views and policies of NIHS and Ministry of Health, Labour and Welfare of Japan (MHLW) or else.

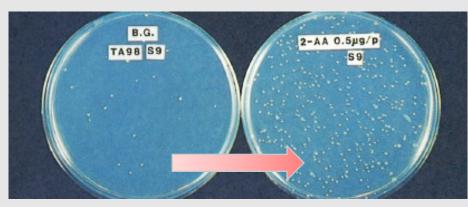




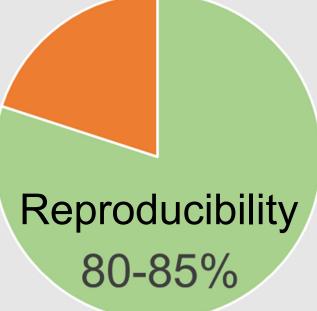
Ames test



- In vitro genotoxicity testing
- High reproducibility
- Sensitive to detect carcinogens and non-carcinogens compared with other tests.



Positive



 \rightarrow In silico Ames QSAR is effective for regulation.





- High sensitivity
- High negative predictivity
 - Minimum false-negative
- Wide coverage
 - Max chemical space

→ Need high quality data



Bigdata used in the 1st and 2nd Ames/QSAR International Challenge Projects



Proprietary dataset by DGM/NIHS

- 1. Ames database with >10,000 new chemicals.
- 2. The origin of the Ames test reports is ANEI-HOU, MHLW
- 3. The reports were originally undisclosed but the outcomes (positive or negative) were made available for validation, development and improvement of QSAR tools.

High quality Ames test data

- 1. ANEI-HOU test guideline, similar to OECD TG 471
- 2. Five strains with/without metabolic activation

MHLW: the Ministry of Health Labour and Welfare of Japan ANEI-HOU: Industrial Safety and Health Act; To secure safety and health in the workplace, new chemicals in >100 kg/year require Ames test. Honma et al., *Mutagenesis* 34, 2-16, 2019 modified.



The 1st project achievements



Mutagenesis Special Issue, **34** (2019)

Mutagenesis, 2019, 34, 3–16 doi:10.1093/mutage/gey031 Original Manuscript

OXFORD

Original Manuscript

Improvement of quantitative structure-activity relationship (QSAR) tools for predicting Ames mutagenicity: outcomes of the Ames/QSAR International Challenge Project

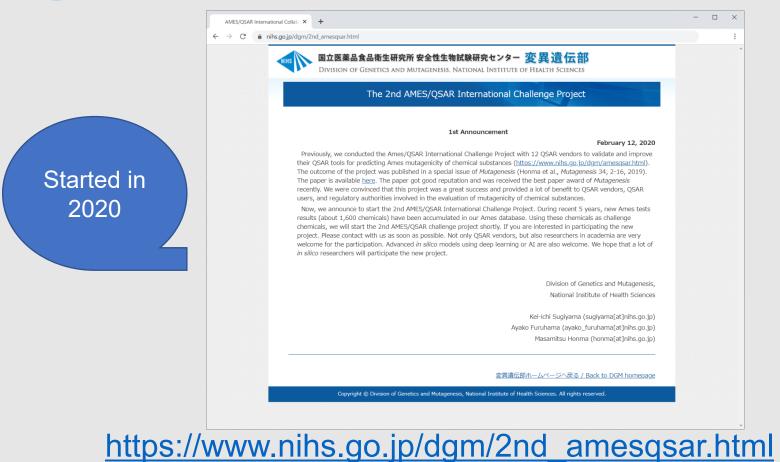
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This paper received best paper award of *Mutagenesis* in 2019



The 2nd Ames/QSAR project



Chemical Hazards Control Division, Industrial Safety and Health Department, MHLW of Japan for providing the ANEI-HOU Ames dataset and allowing us to use the data in the projects.

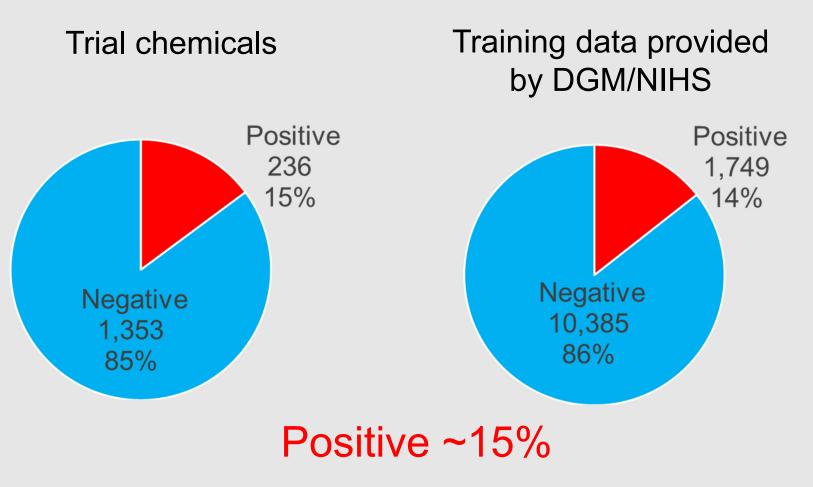




Where	1st project DGM/NIHS	2nd project DGM/NIHS			
When	2014 - 2017	2020			
Who participate	12 teams (7 countries) Mainly QSAR venders	19 teams (11 countries) Academia/non-commercials			
What kind predictions	Three trials: ~4,000 chemicals/trial	One trial: ~1,600 chemicals			
How many training data provided by DGM	Trial I: 0 data Trial II: ~4,000 data Trial III: ~8,000 data	>12,000 data used at the 1st project			
Why	QSAR tool improvements				
by using	Statistical, rule-based, its consensus models	Machine learning models and AI based systems			



Ames results in the 2nd project







Additional data provided to the participants of the 2nd project

CAS RN Chemical Name Structure			Result	Purity	ity Solvent	Without metabolic activation (- S9)				With metabolic activation (+ S9)					
CAS KN Chemical Name	Siluciule	Result	(%)	Solvent		TA1535	WP2 uvrA	TA98	TA1537	TA100	TA1535	WP2 uvrA	TA98	TA1537	
			++	>99	H2O	++	-	++	++	++	+	+	++	+	+
•			++	99.5	DMSO	++	+	+	+	-	++	+	+	+	-

- Purity
- Solvent
- Results of each strain

++: Strongly Positive +: Positive -: Negative



2nd Project participants



Name	Country
1 Shanghai Institute of Organic Chemistry, Chinese Academy of Sciences	China
2 Altox Ltd.	Brazil
3 MN-AM	Germany/USA
4 Leadscope, Inc.	USA
5 Istituto di Ricerche Farmacologiche Mario Negri - IRCCS	Italy
6 IdeaConsult Ltd.	Bulgaria
7 MultiCASE Incorporated	USA
8 Lhasa Limited	UK
9 Istituto Superiore di Sanità (ISS)	Italy
10 Gifu University	Japan
11 Massachusetts Institute of Technology	USA
12 Simulations Plus, Inc.	USA
13 Chemotargets	Spain
14 LMC - Bourgas University	Bulgaria
15 The University of Sydney	Australia
16 Meiji Pharmaceutical University	Japan
17 Liverpool John Moores University	UK
18 Evergreen AI, Inc.	Canada
19 Politecnico di Milano	Italy



The performance metrics of QSAR tools



1st project

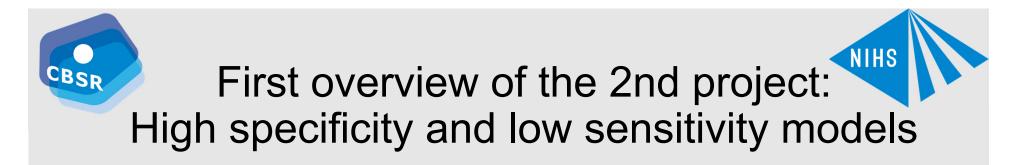
	Phase I	Phase II	Phase III
Sensitivity (%)	56.7	58.0	57.1
	(38.6–70.0)	(41.6-72.1)	(31.7–67.6)
Specificity (%)	77.7	84.2	79.9
	(62.5–91.5)	(64.9–92.8)	(60.7–93.0)
Accuracy (%)	74.7	80.3	76.7
	(63.6-83.9)	(65.8-87.7)	(68.0–87.3)

Honma et al., *Mutagenesis* 34, 2-16, 2019 modified.

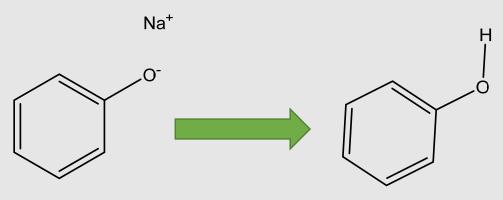
Sensitivity : the ability to detect mutagens Specificity : the ability to detect non-mutagens Accuracy: proportion of correct predictions

2nd project: under analysis

The models still show -Low sensitivity -High specificity



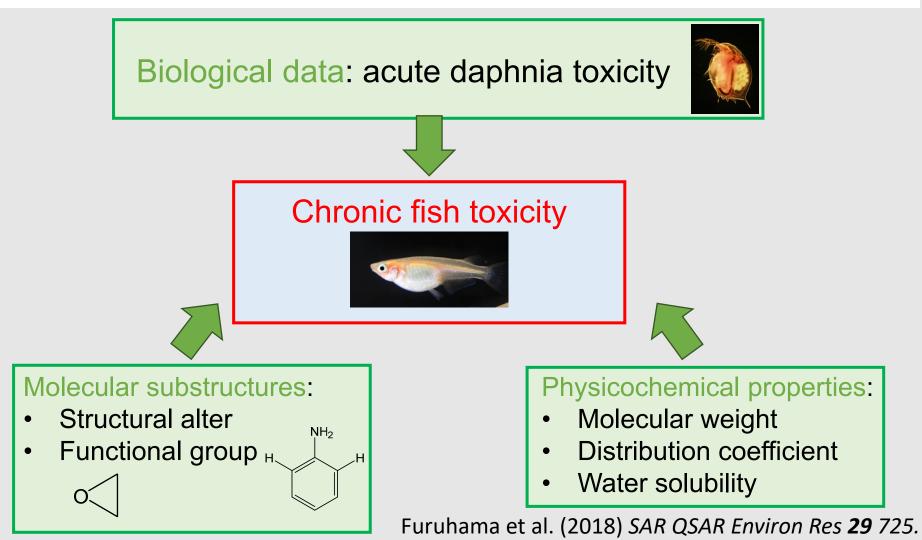
Lack of data curations may enhance the data vias
Valance of positive/negative data (ideal 1/1)
Chemical structures

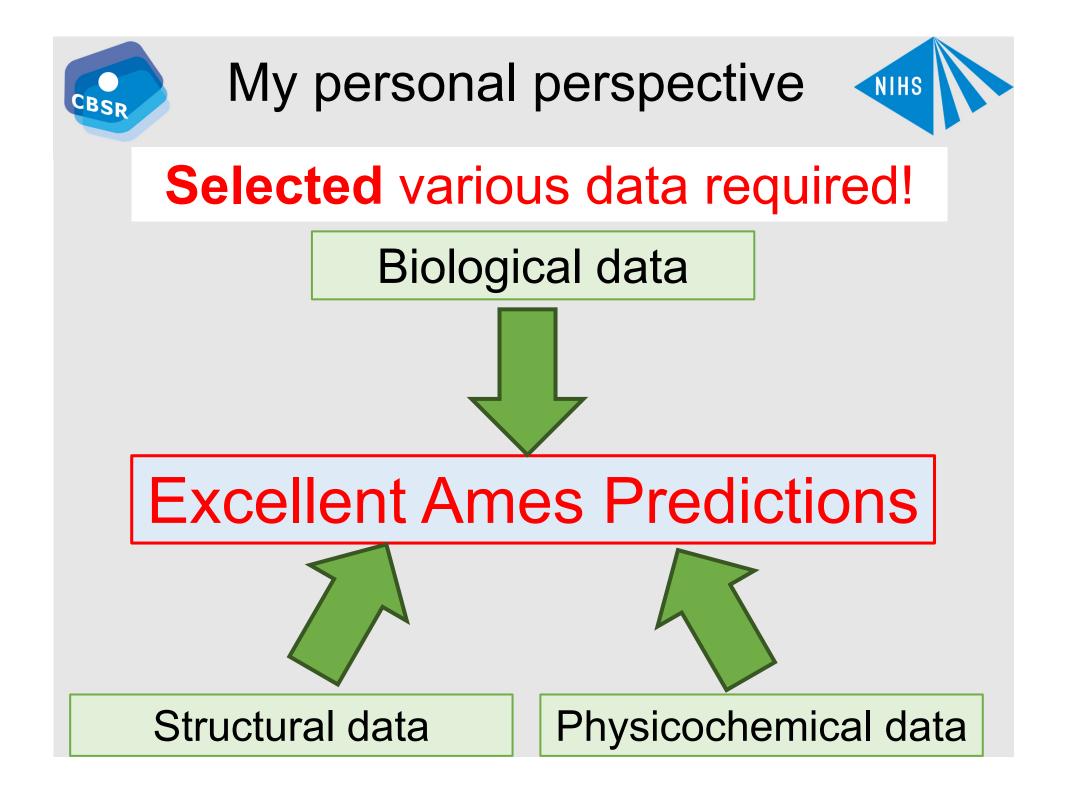


>10,000 data: small for the Ames prediction of new chemicals
Experiences may cover a small chemical space

Ayako's previous study: model development for chronic ecotoxicity prediction

Daphnia toxicity + selected structural and physicochemical data brought good fish chronic toxicity perdition models







Summary



- The high reproductivity and amount of Ames data allow us the use of QSAR.
- DGM/NIHS started the 2nd Ames/QSAR project providing with informative >10,000 Ames data.

→ First overview: Models show high specificity and low sensitivity. >10,000 Ames data are not enough.

• For improving predictivity, experiences and various type of selected data might be important as well as the number of Ames data.

 \rightarrow We need to investigate how to improve the reliable QSAR models and *in silico* predictions including machine learning and AI.



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