



# ***Case Study on the Use of an Integrated Approach to Testing and Assessment for Identifying Estrogen Receptor Active Chemicals***

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U.S. Environmental Protection Agency

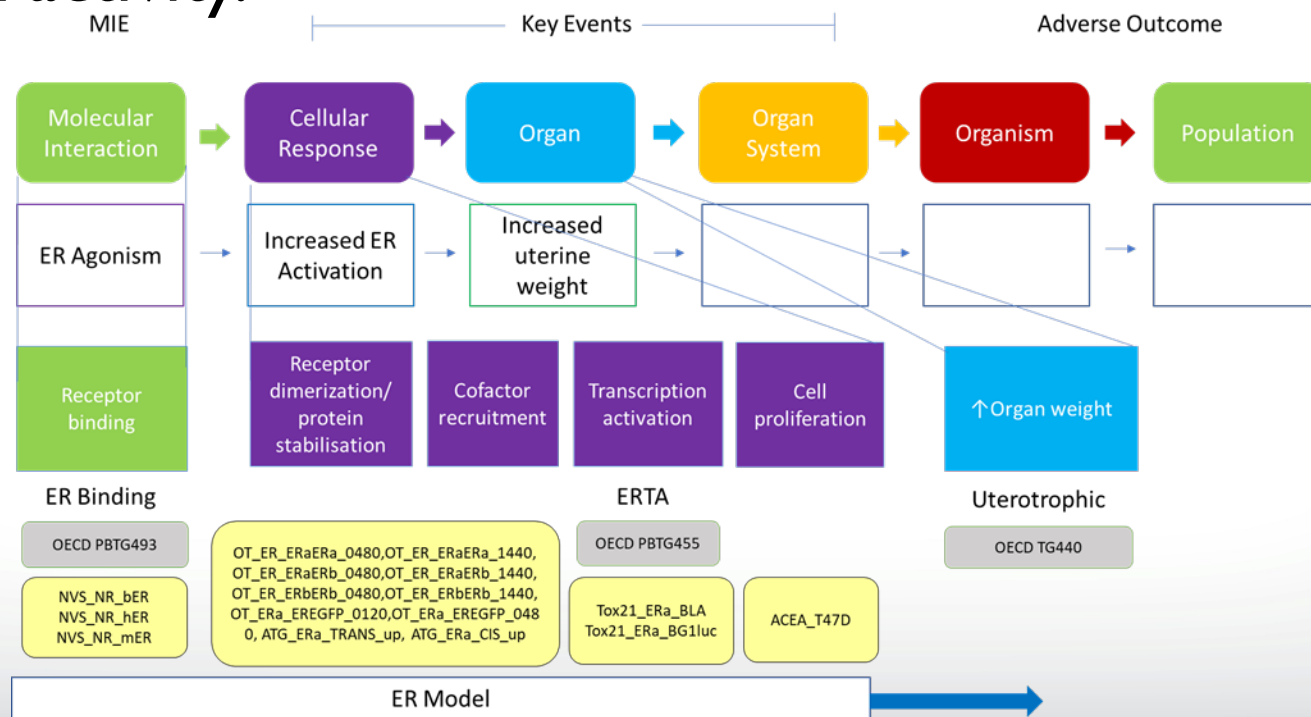
Office of Research and Development

Washington, DC

*The views expressed in this presentation are that of the presenter and do not represent the views and/or policies of the US Environmental Protection Agency.*

- The intended application of this IATA is for
  - screening of environmental chemicals based on their ER agonist activity
  - determining whether further evaluation of endocrine-related activity in higher tier in vivo tests (e.g., female pubertal assay, two generation reproductive toxicity study) is needed

- To use a combination of 4 to 16 in vitro high throughput screening (HTS) assays and a computational model for estrogen receptor (ER) agonist activity, as an alternative to low and medium throughput in vitro and in vivo tests for ER activity.







# IATA Process

## Run Assays

(Assays 1 - n,  
cytotoxicity)

- Choose at least 4 in vitro HTS assays that fit the criteria as described. Briefly, the assays chosen must probe diverse points in the ER pathway and use diverse assay reporting technologies and cell types.
- The possible combinations of 4 assays that can be used together are listed in Annex III. More than 4 assays can be chosen, in which case the results of all chosen assays must be modelled and the results reported. Concurrently evaluate a phenotypic response, cellular proliferation.

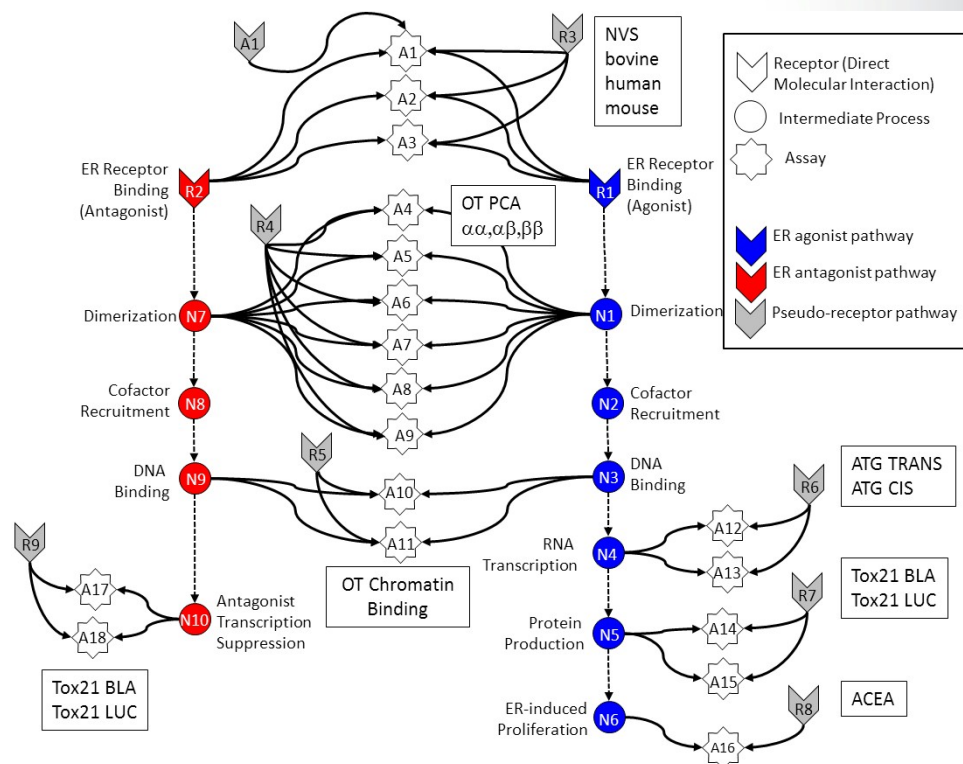
## Import Data and Run Model

- The next step is to collect data from the assays into a format that the computational model's R-code can import and analyse. Ideally, this process will be performed automatically (electronically) to reduce the chance of user-input error.
- Although useful, advanced computational expertise is not necessary to run the model. For example, the formula used for the subset model analysis could be implemented in a spreadsheet so limited computational expertise required.

## Generate Report

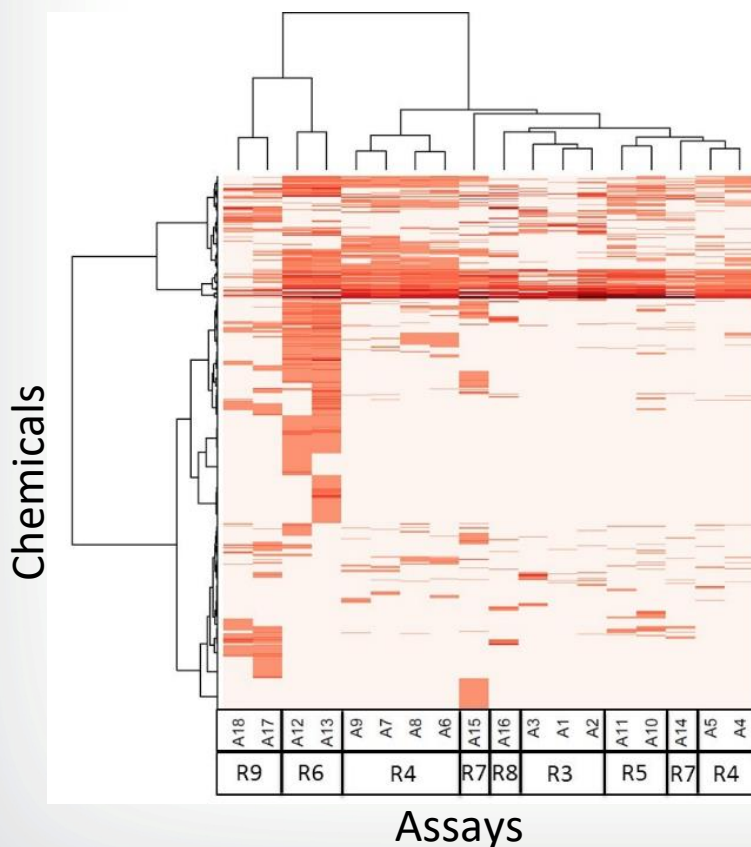
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- Any departure from the methodology of the ER pathway model as presented in this document must be thoroughly described along with the reason for the departure and the proposed impact on the screening results.

- **No *in vitro* assay is perfect**
  - Assay Interference
  - Noise
- **Use multiple assays per pathway**
  - Different technologies
  - Different points in pathway
- **Use model to integrate assays**
- **Evaluate model against reference chemicals**



# All In vitro assays have false positives and negatives

Assays cluster by technology, suggesting technology-specific non-ER bioactivity

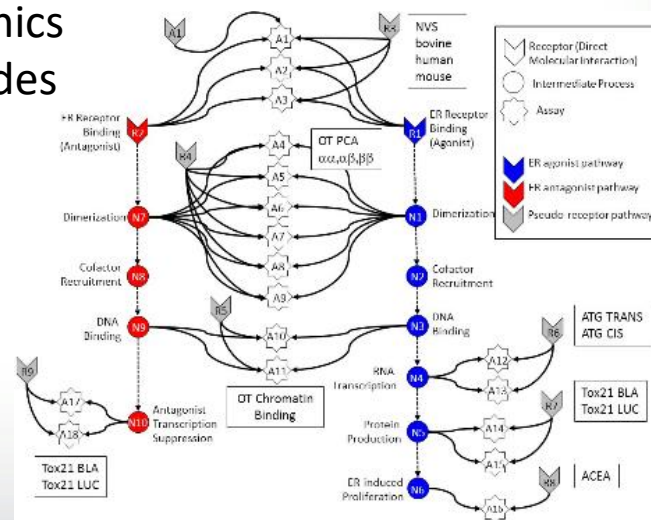


Much of this “noise” is reproducible

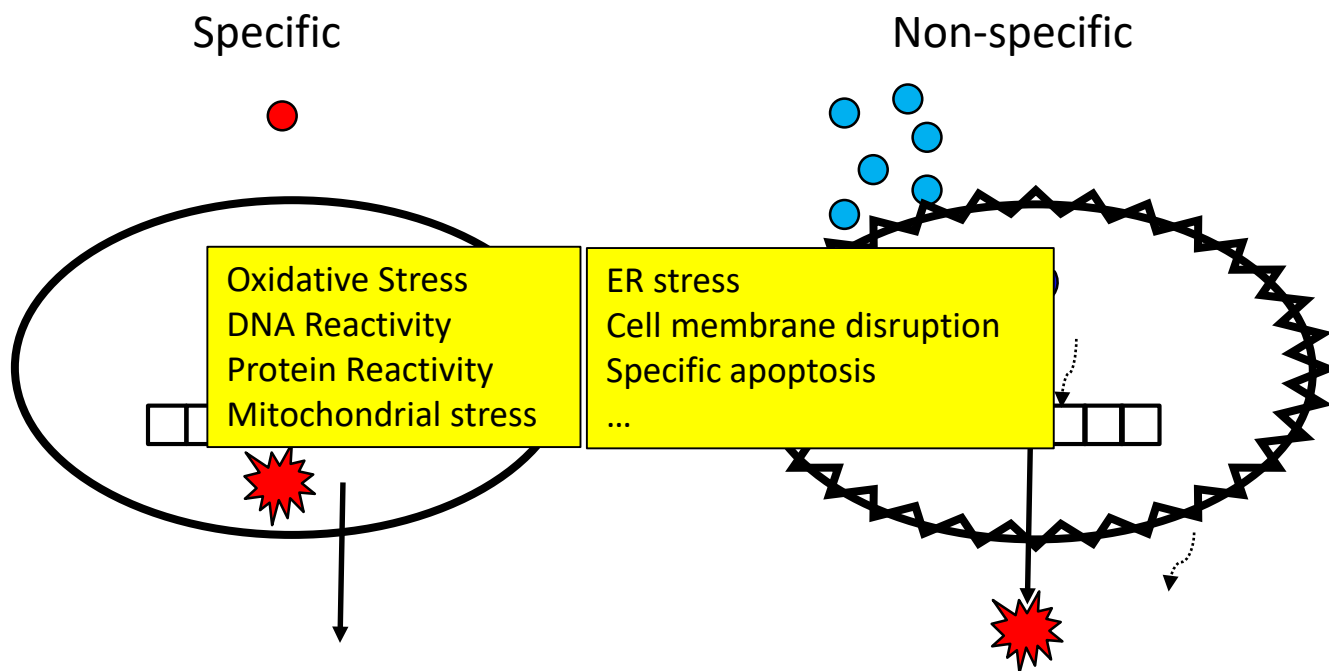
- “assay interference”
- Result of interaction of chemical with complex biology in the assay

Chemical universe is structurally diverse

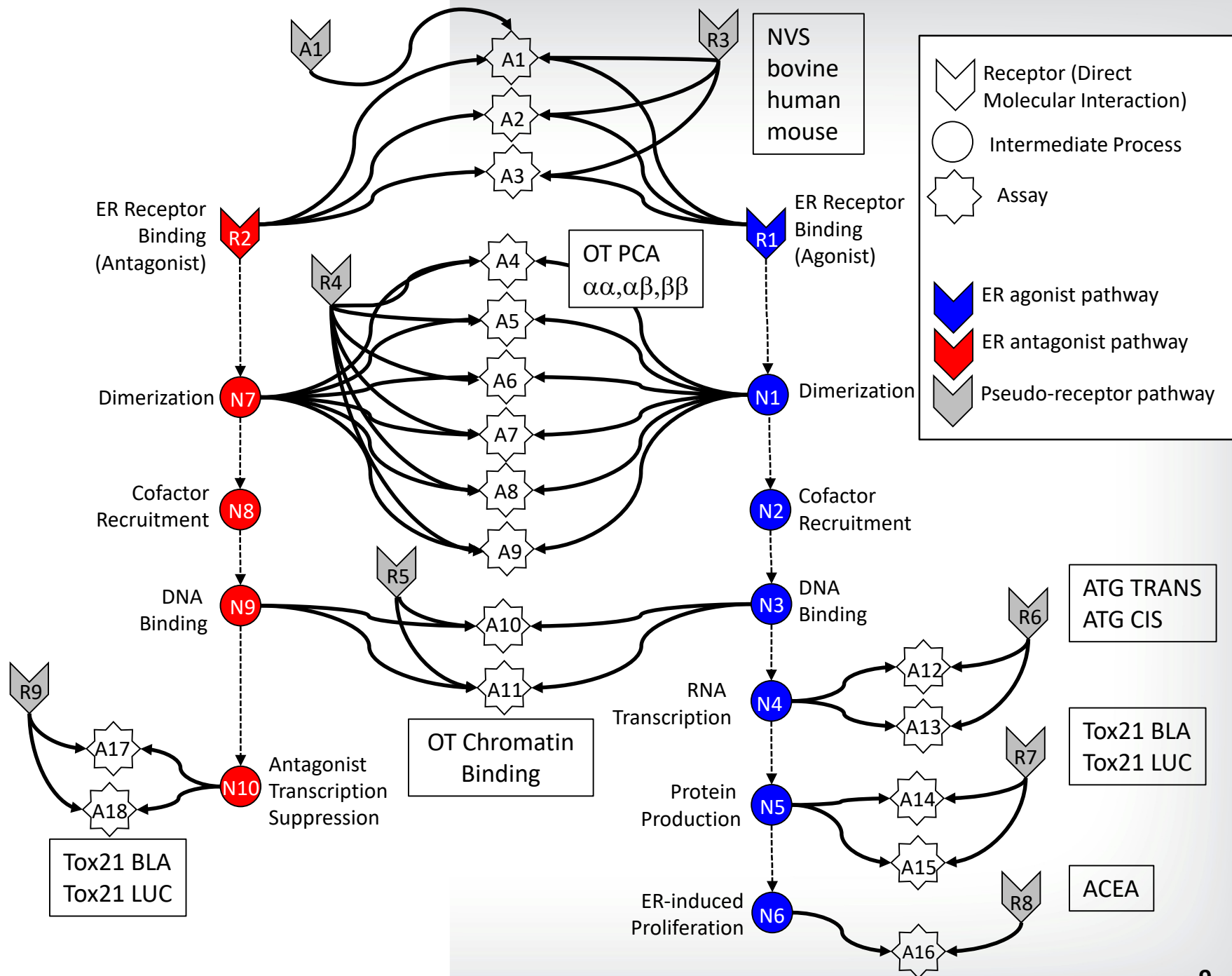
- Solvents
- Surfactants
- Intentionally cytotoxic compounds
- Metals
- Inorganics
- Pesticides
- Drugs

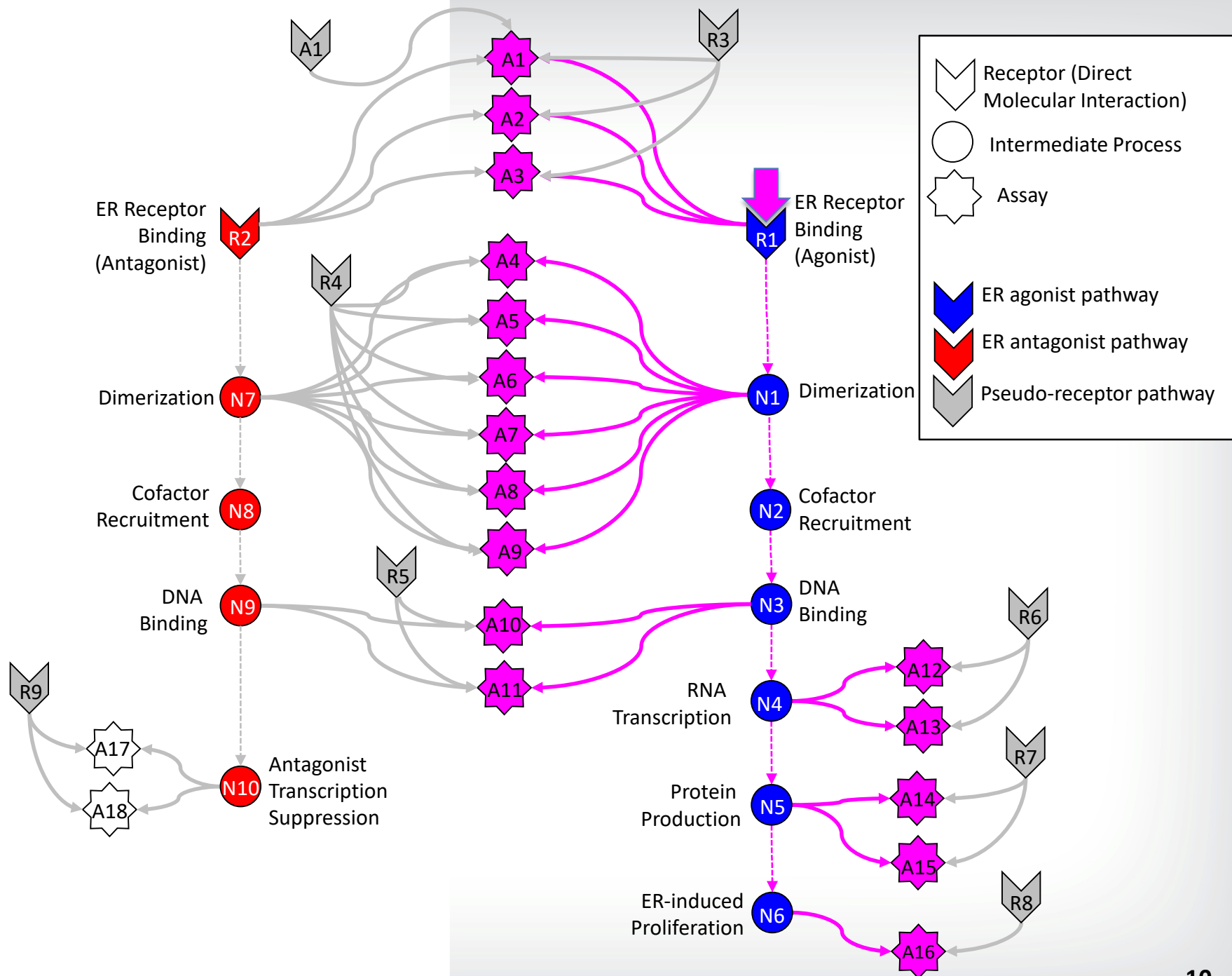


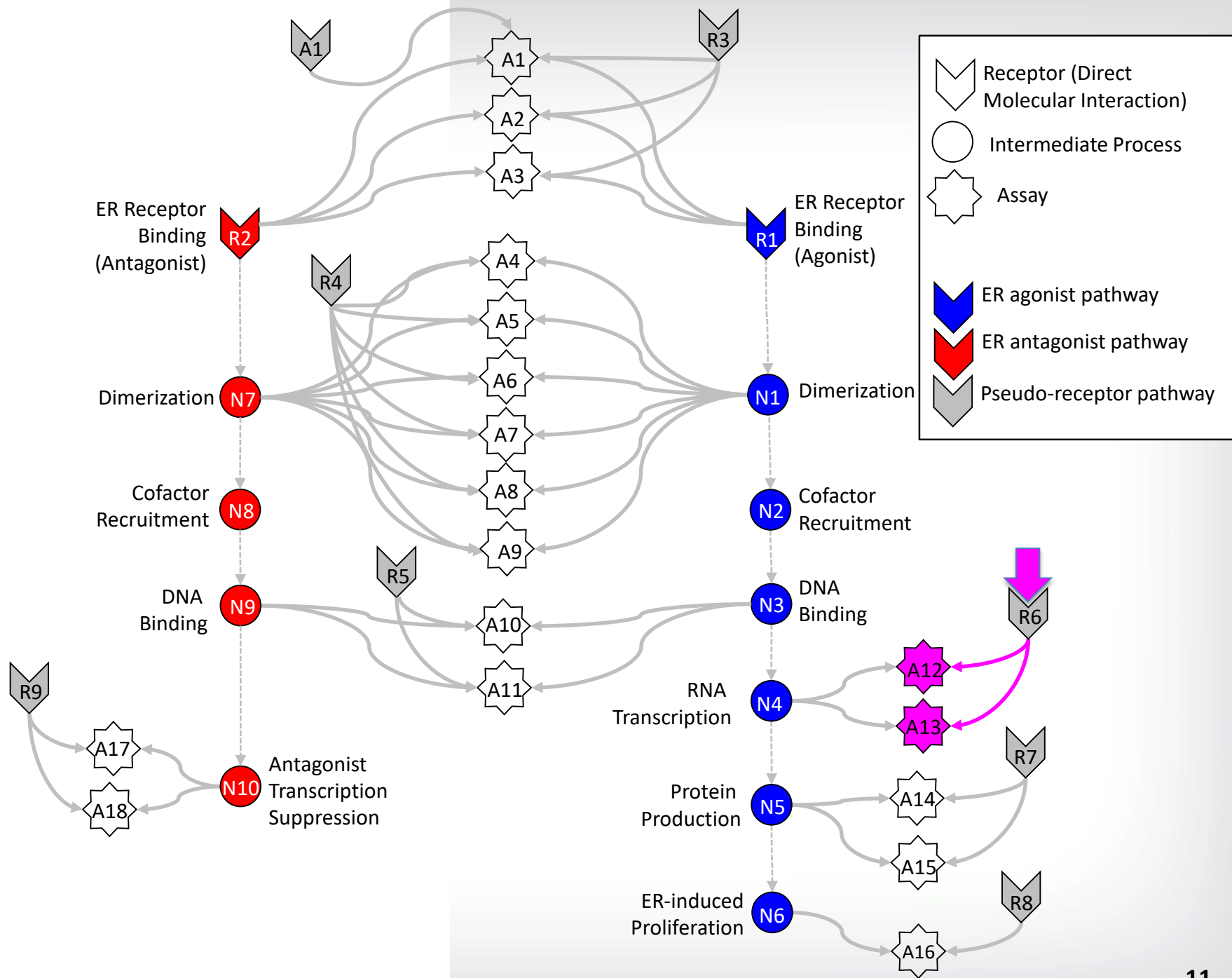
## Schematic explanation of non-specific activity









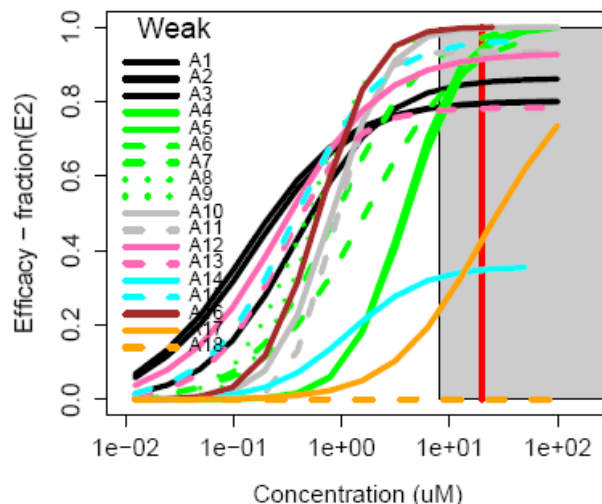




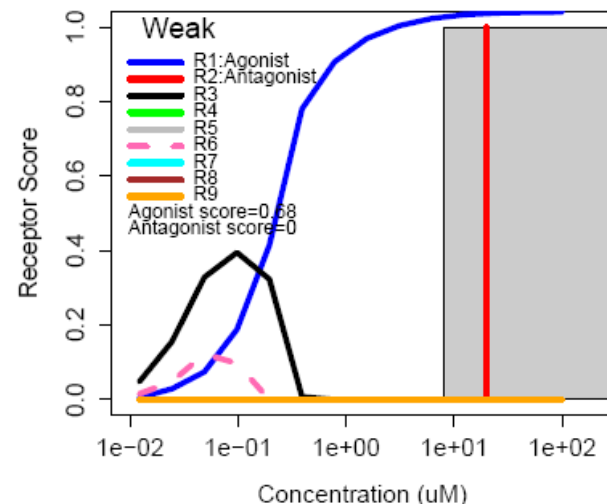
# Example chemicals: Observe quantitative uncertainty

True Agonist

80-05-7 : Bisphenol A

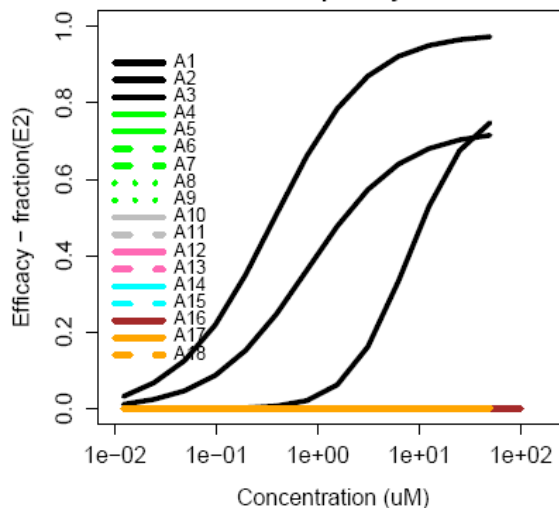


80-05-7 : Bisphenol A

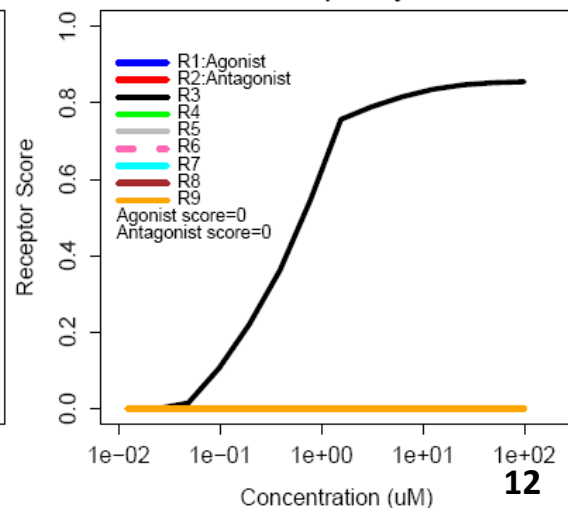


Assay Interference Example "R3"

10016-20-3 : alpha-Cyclodextrin



10016-20-3 : alpha-Cyclodextrin



- The IATA has been clarified further to demonstrate that:
  - a subset of as few as 4 assays as an alternative approach to the current guideline
  - the subset of assays can be any assays that fit into specific criteria as described
    - interrogates different points on the ER pathway
    - incorporate different technologies
    - Includes diverse cell types
- Included in this case study is an annex that describes the 9 subset models with 7 or fewer assays that achieve  $\geq 94\%$  balanced accuracy for all chemicals and the in vitro and in vivo reference chemical sets. (Annex III).
- Further articulated the benefits of using a subset of assays in the IATA, specifically the flexibility it gives users by allowing the use of any assays that fit the described criteria.

- **Assume that the “full model” (16 agonist assays) provides acceptable prediction of ER agonist activity**
  - Model based on detailed biology
  - Validated against in vitro and in vivo reference chemicals
- **Build simple “subset models” using fewer assays**
  - Validate against the full model over 1811 chemicals and in vitro and in vivo reference chemicals
- **Input to the model are assay-chemical AUC values**



# Subset Model Input

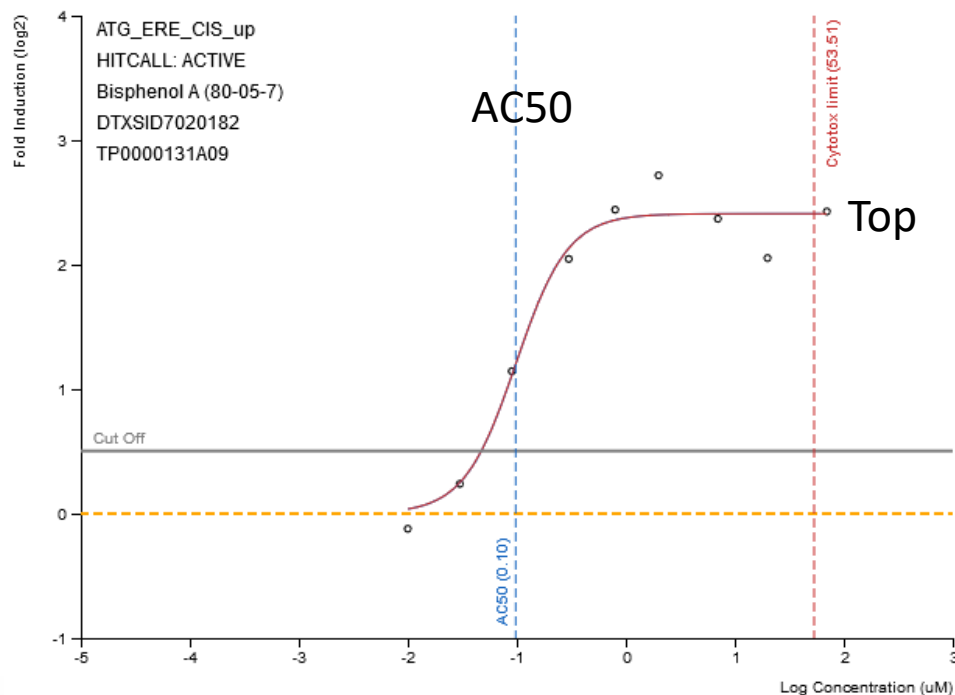
- Inputs to model are chemical-assay **AUC** (area under the curve) values
- Run assay in concentration-response mode
- Fit to model (e.g. Hill model)
- Calculate **AC50** and **Top**
- **AUC = -log(AC50) x Top**

Example curve for Bisphenol A

Data from:

EPA CompTox Chemicals Dashboard

<https://comptox.epa.gov>



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(Assays 1 - n,  
cytotoxicity)

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## Import Data and Run Model

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## Generate Report

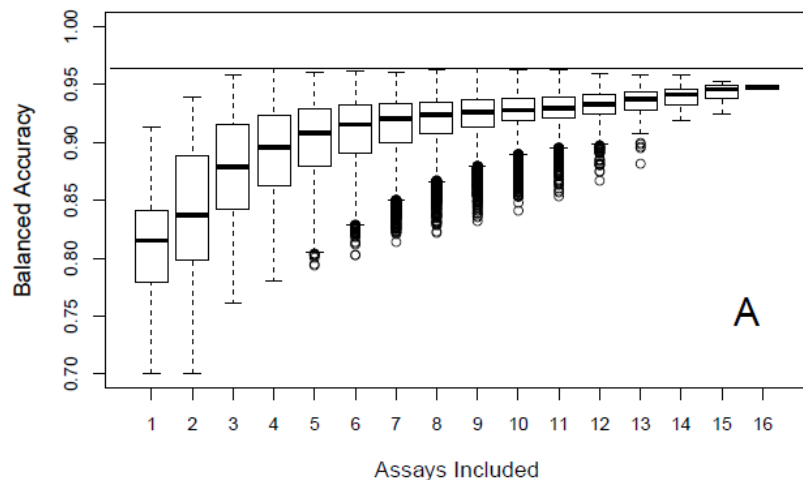
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- **Step 1: For each set of assays (all combinations from 1 to 16), calculate the pathway AUC using the following formula ...**
- $AUC_{subset}(chemical) = \sum_{i=1}^N c_i \times AUC(chemical, assay_i)$
- **Step 2: vary the weights ( $c_i$ ) to minimize the squared difference between the subset pathway AUC and the full model AUC across all 1811 chemicals**
  - This is a standard linear model inverse problem
- **Calculate several statistics on each subset pathway model**
  - Sensitivity, specificity, balanced accuracy for all chemicals
  - Sensitivity, specificity, balanced accuracy for in vitro reference chemicals
  - Sensitivity, specificity, balanced accuracy for in vivo reference chemicals
  - (balanced accuracy is average of sensitivity and specificity)
- **Allow a user to select any model (i.e. any subset of assays) that provides high enough sensitivity, specificity, balanced accuracy across the 3 chemical group**

# Statistical Results (I)

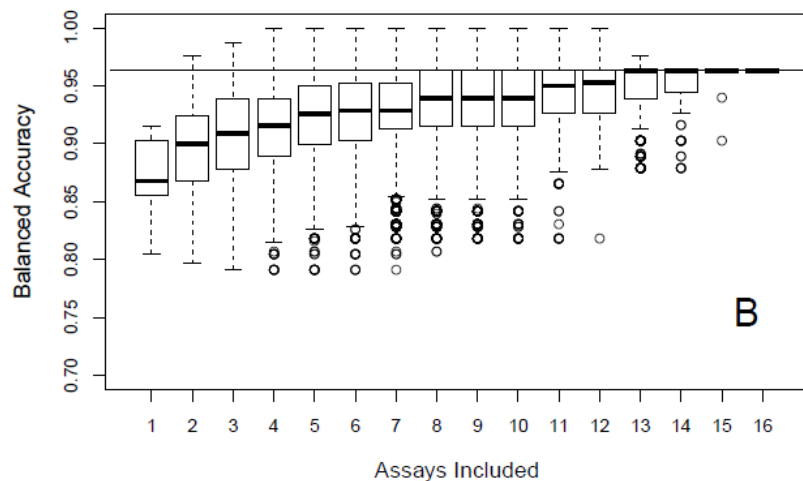
All Chemicals



Boxplots show balanced accuracy for all models with a specified number of assays

- Top (all chemicals)
- Bottom (all reference chemicals, combined in vitro and in vivo)

Reference Chemicals



Observe that with as few as 4 assays, there are subset models where reference chemical balanced accuracy is perfect and other subset models where all-chemical balanced accuracy is as high as it gets (93%)

## Statistical Results (II)

Model that optimizes one parameter does not always optimize the other

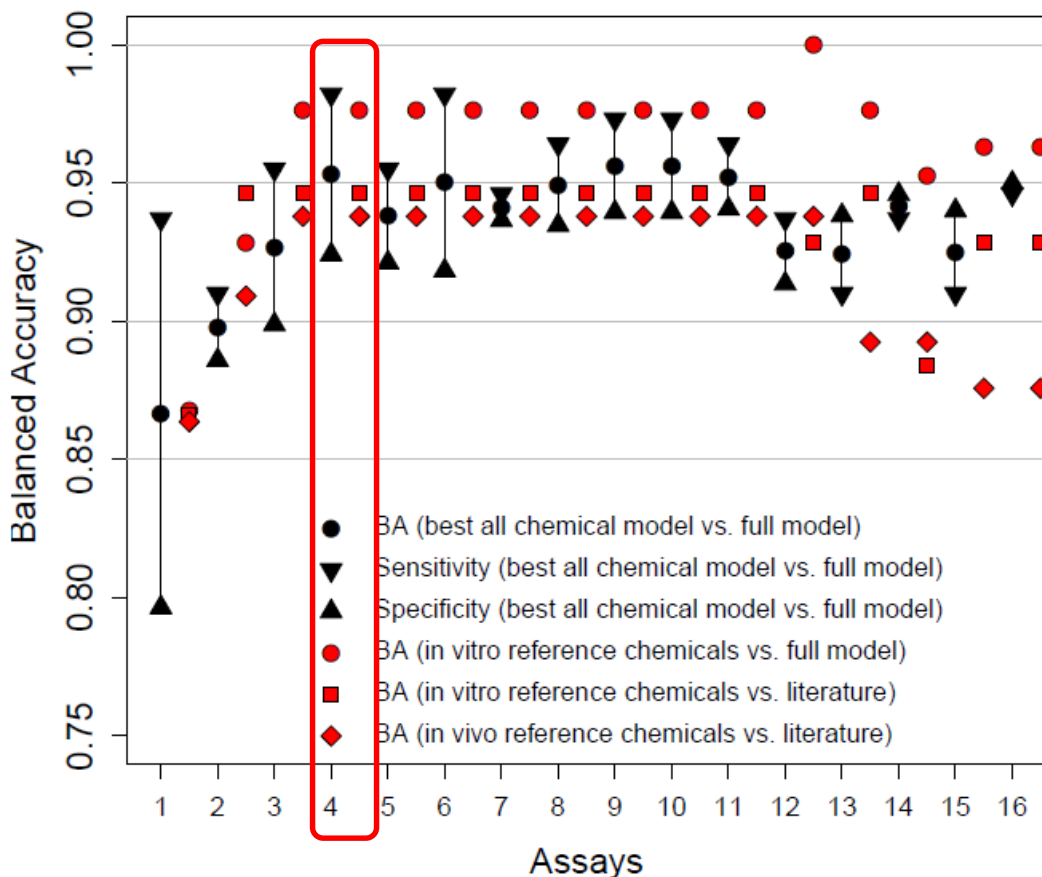
This plot shows the statistics for the best overall model for a given number of assays

Example: best overall 4-assay model gives balanced accuracy of

- All chemicals: 0.955
- In vitro reference chemicals: 0.945
- In vivo reference chemicals: 0.94

Sensitivity is higher than specificity (minimize false negatives)

Note that models with intermediate number of assays (4-11) perform better than those with more or fewer

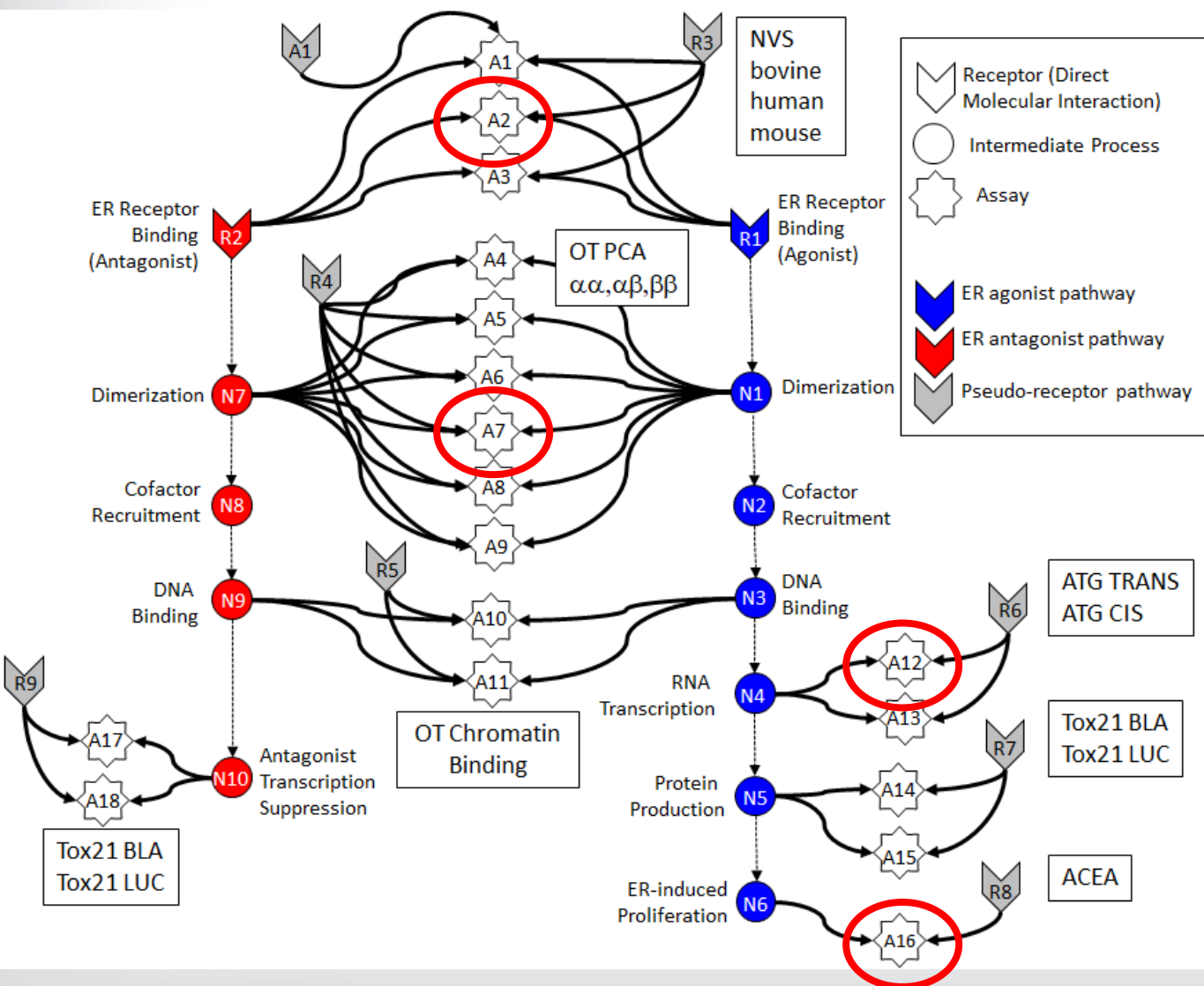




# What assays are used? (best assay sets)

Assays	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16
Sensitivity (all chemicals)	0.94	0.91	0.95	0.98	0.95	0.98	0.95	0.96	0.97	0.97	0.96	0.94	0.91	0.94	0.91	0.95
Specificity (all chemicals)	0.80	0.89	0.90	0.92	0.92	0.92	0.94	0.93	0.94	0.94	0.94	0.91	0.94	0.95	0.94	0.95
BA (all chemicals)	0.87	0.90	0.93	0.95	0.94	0.95	0.94	0.95	0.96	0.96	0.95	0.93	0.92	0.94	0.92	0.95
BA (in vitro reference chemicals)	0.87	0.95	0.95	0.95	0.95	0.95	0.95	0.95	0.95	0.95	0.95	0.93	0.95	0.88	0.93	0.93
BA (in vivo reference chemicals)	0.86	0.91	0.94	0.94	0.94	0.94	0.94	0.94	0.94	0.94	0.94	0.94	0.89	0.89	0.88	0.88
BA (minimum)	0.86	0.90	0.93	0.94	0.94	0.94	0.94	0.94	0.94	0.94	0.94	0.93	0.89	0.88	0.88	0.88
<b>Assay Selection</b>																
NVS NR bER	0	0	0	0	0	0	0	1	1	1	1	1	1	1	1	1
<b>NVS NR hER</b>	0	0	1	1	1	1	1	1	1	1	1	1	1	1	1	1
NVS NR mERa	0	0	0	0	1	1	0	1	1	1	1	1	1	1	1	1
OT ER ERaERa 0480	0	0	0	0	0	0	1	1	1	1	1	1	1	1	1	1
OT ER ERaERa 1440	0	0	0	0	1	0	1	1	1	0	0	1	1	1	1	1
OT ER ERaERb 0480	0	0	0	0	0	1	1	0	1	1	1	1	1	1	1	1
<b>OT ER ERaERb 1440</b>	0	0	1	1	0	1	0	0	0	1	1	1	1	1	1	1
OT ER ERbERb 0480	0	0	0	0	0	0	1	0	0	1	1	1	1	1	1	1
OT ER ERbERb 1440	0	1	0	0	0	0	0	0	0	0	1	1	1	1	1	1
OT ERa EREGFP 0120	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	1
OT ERa EREGFP 0480	0	0	0	0	0	0	0	1	1	1	1	0	0	0	1	1
<b>ATG ERa TRANS up</b>	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1
ATG ERE CIS up	0	0	0	0	0	0	0	0	0	0	0	0	1	1	1	1
TOX21 ERa BLA Agonist ratio	0	0	0	0	0	0	0	0	0	0	0	0	1	1	1	1
TOX21 ERa LUC BG1 Agonist	0	0	0	0	0	0	0	0	0	0	0	1	1	1	1	1
<b>ACEA T47D 80hr Positive</b>	0	0	0	1	1	1	1	1	1	1	1	1	0	1	0	1

# Where do these assays fall on the pathway?



Selected assays come from different points in the pathway and used different assay technologies

- **Many of these assays are not commercially available**
  - Novascreen (cell free binding assays) no longer offered by Novascreen, but other vendors can provide
  - Odyssey Thera (protein complementation assays). Company is out of business, no known commercial source of these assays
  - Attagene (RNA-based transactivation assay) Company currently offers these assays
  - ACEA (Cell proliferation, real time impedance measurement) Company currently offers these assays
  - Tox21 (protein-based transactivation assays) These are produced by a US Government lab, and are not available as a service
- **However, variants of these assays could be developed by independent labs**
  - Specific aspects of assays may be patent protected, but basic technology is not

- **Identify laboratories to develop or offer variants of assays in the following classes**
  - Cell-free ER binding
  - Protein complementation / transcription factor dimerization
  - Transactivation
  - Cell proliferation
- **Use different cell types and readout technologies**
- **Validate each assay against the OECD in vitro reference chemical set**
- **Build the 4-assay subset model against the in vitro reference chemical set to determine assay weights**
- **Validate subset model against further chemicals shown to be ER agonist positive and negative from the current full model**
- **Use the new subset model to evaluate new chemicals**

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