

Introduction




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Welcome!

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- The goal of this workshop is not to give a comprehensive overview of all possible methods to analyze exposure to mixtures
- This is a very **hot** topic right now in the EH community
- New methods are being developed and adapted from other fields daily
- Instead, our goal is to give some **examples** of different approaches
 - That are used to answer **different research questions**
- Go over some tools in  to apply these methods
- Importantly, discuss what *types* of methods are appropriate for which research questions

What is a mixture?

- Actually, there is no strict definition
- According to NIEHS “a mixture must have at least three independent chemicals or chemical groups”
- Generally, exposure to a mixture indicates exposure to **multiple** “stressors” simultaneously
 - Chemical
 - Non-chemical (SES, diet, etc)

Why care about mixtures?

- We are exposed to hundreds (thousands?) of chemicals at any single time point
- Traditionally, epi studies have focused on single-chemical analyses
- This does not represent reality
- The combination of exposures to multiple chemicals likely induces different responses
 - Compared to exposure to each chemical independently

Million dollar question

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How can we represent the complexity of reality in a (single) statistical model?

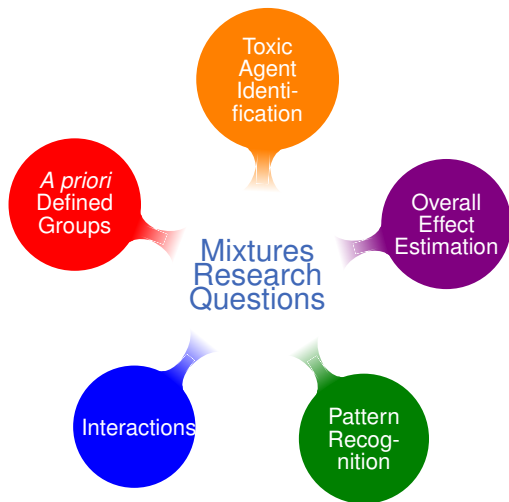
How do we deal with exposure to mixtures?

- This is still a very open question
- Existing methods have limitations
- There have been several workshops held by EPA and NIEHS to address this issue
- The most recent NIEHS workshop (2015) concluded that
 - 1 Although some methods performed better than others the presented estimated associations were still quite variable and not in agreement
 - 2 The choice of method should depend on the research question

- In fact, after the 2015 workshop NIEHS had an RFA for robust methods development
- Powering Research Through Innovative Methods for Mixtures in Epidemiology (PRIME)
- Grants were awarded starting January 2018
- Six grantees with very different proposed methods
- One of the requirements was that **all software** developed under PRIME will be publicly available
- More here: https://www.niehs.nih.gov/research/supported/exposure/mixtures/prime_program/index.cfm

Potential questions in mixtures analyses

For mixtures analyses the selected method depends on the primary research question



Bird's-eye (over)view of existing mixtures methods

*Not an exhaustive list of methods!!

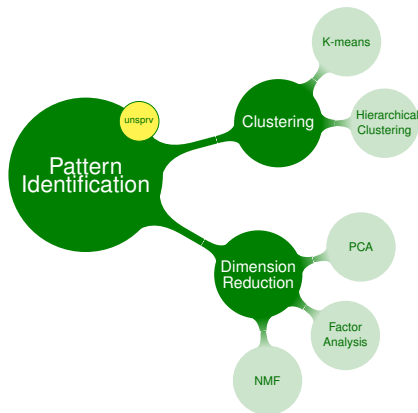


Some considerations

- 1 No single method outperforms all others for all potential questions
- 2 Interpretability
- 3 Robustness (stable solutions)
- 4 Computational scalability – as the dimensionality of our dataset increases (either N or p) some methods might start to fail
- 5 Exploration vs. hypothesis testing
- 6 Usually not a good idea to “blindly” use methods from other fields – may need to adjust them first

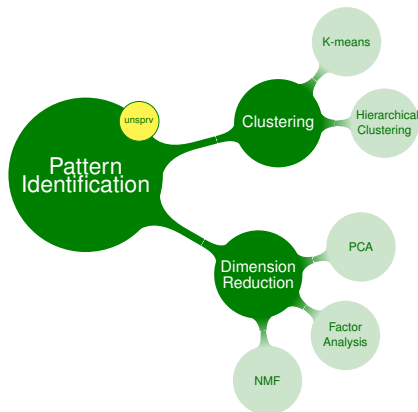
Exposure pattern recognition

- Why should we care about identifying **exposure patterns** to chemicals in a population?
 - Sources
 - Behaviors
- If we link these patterns to (multiple) adverse health outcomes
 - Efficient regulations
 - Targeted interventions



Exposure pattern recognition (cont'd)

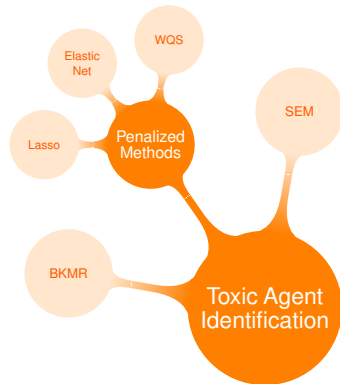
- Class of **unsupervised** methods
 - Solution independent of any outcome
- But supervised extensions exist (for most)
- Which version to use depends on the research question
 - Inform regulatory action, interventions
 - Better understand biological pathways



Identifying toxic agents

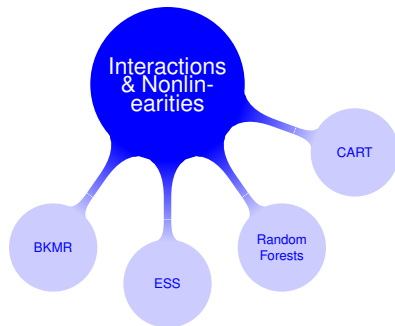
aka “bad actors”

- Which chemical(s) in my mixture are related to the outcome?
- Estimate chemical-specific *independent* effects
- While accommodating the (potentially very) high correlations among mixture members



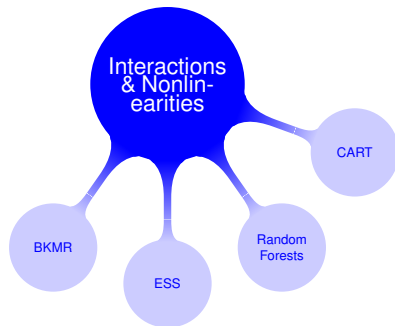
Interactions & non-linear relationships

- These are actually two different classifications of potential research questions
 - 1 Interactions among mixture members?
 - 2 Non-linear exposure – response curves?
- But methods tend to do both



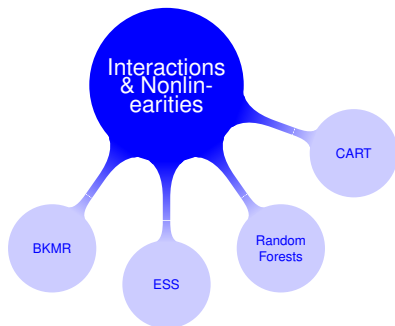
Non-linear relationships

- Because linearity is just an assumption . . .



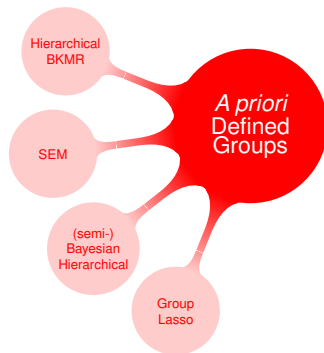
Interactions among mixture members

- If there is reason to believe that the combined health effect is greater (or less) than the sum of independent effects
 - Potential synergism
 - Most methods can accommodate *a priori* defined interactions
 - Need to hard code
 - Dimensionality ...
- Semi- or non-parametric methods preferred



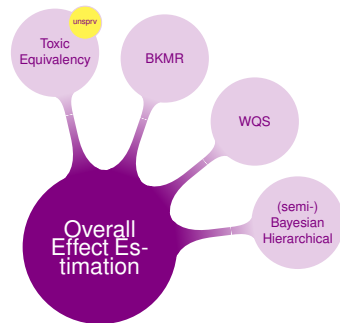
A priori defined groups

- We might have some prior knowledge or hypothesis on
 - How chemicals group naturally in the environment
 - Might share pathway to toxicity
- Methods exist to allow estimation both of group and within-group effects



Overall Mixture Effect

- Finally, we may want to estimate the overall mixture effect
- As chemical concentrations in the mixture increase, do we observe corresponding changes in the outcome?



Comparing results across methods

- Generally a good practice
 - Especially if complementary methods
 - Sensitivity analyses to assess robustness of results
- Even if different methods address different questions, consistency in findings is always welcome
- If/when differences across methods are detected → keep in mind what the aim of each method is!
- Trying different methods and choosing the answer we like the best should *a/ways* be avoided
 - I.e. no cherry-picking!

In summary...

- During this workshop we will present information on a few different methods
 - That are used to answer different research questions
 - By no means a comprehensive list
 - The goal is to have an open **Discussion** about how to quantify health effects of exposure to mixtures
 - ... With some extra technical details ;)
 - This is a very heterogeneous group in terms of experience with such methods
- Please share your previous experience!

The example we'll be using during this workshop

- We wanted to use a real-life application for all the labs
 - Instead of simulated datasets
- We decided on the Mitro et al paper:
 - ① High-dimensional exposure matrix
 - ② Publicly available data (NHANES)
- A big **Thank you!** to Dr. Ami Zota (GW)
 - Provided the datasets and code to get the final dataset as it is exactly on the paper
 - For consistency

The example we'll be using (cont'd)

For consistency with the Mitro et al. paper and across our labs

- We kept all parameters as in the paper
 - Log-transformed outcome and exposures
 - Same list of confounders included in the models
- Only included variables with at least 60% > LOD
- Since it is not the purpose of the workshop to discuss these choices, please refrain from asking such questions during the labs
- Happy to discuss these during the breaks
 - Although we might not be able to explain the choices the authors made

The platform we'll be using in the labs

<https://rstudio.cloud/>



- Looks like R studio but runs on the cloud
- Might take a bit to load – please load at the break before the first lab
 - or now
- We have shared instructions, but if you have any questions please ask at the break before the first lab!
- Most files are .rmd instead of .r → the code is in the gray chunks
- You also have the option to follow using the .html files
 - choose option “view in web browser”

Thank you!

Questions?

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