

Clustering computer mouse tracking data via informed priors

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- 👉 Identifying molecular subtypes of cancer for personalized treatment.
- 👉 Multi-view modeling may try to identify group patients with similar genetic and clinical profiles.

- Integrating Multi-Omics Data in Precision Medicine
 - **Genomics:** DNA sequencing data
 - **Transcriptomics:** RNA expression levels
 - **Proteomics:** Protein concentration data
 - **Metabolomics:** Metabolic markers in blood
- 👉 Predicting drug response by combining different biological levels of information.
- 👉 Multi-view models improve patient stratification for targeted therapies.

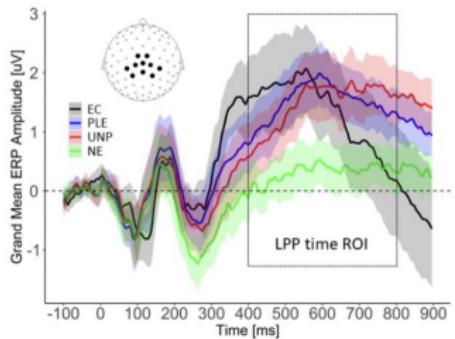
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- 👉 Multi-view models improve patient stratification for targeted therapies.
- Many other examples: **Public Health Surveillance** using hospital admissions, social media data, environmental data, etc.

Ultimate Goal: During a **smoking-cessation intervention**, we want to learn if an individual is at risk of **relapse**.

Neuroscientists may collect many data, including **EEG data**

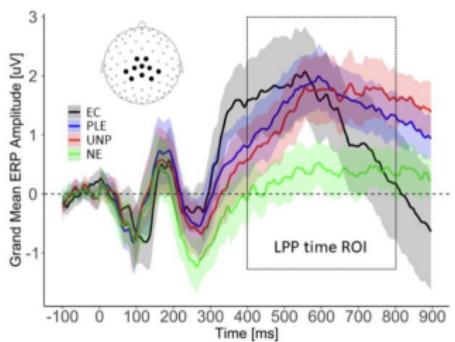
- They may also collect **behavioral data** where smokers are asked to make decisions in front of cigarette-related cues and other cues in experiments.
- 👉 Thus, to understand decision-makings behind smoker's behaviors, neuroscientists end up collecting data across multiple modalities and views.

- EEG experiment data
event-related potentials
(ERPs, a direct measure of brain activity)

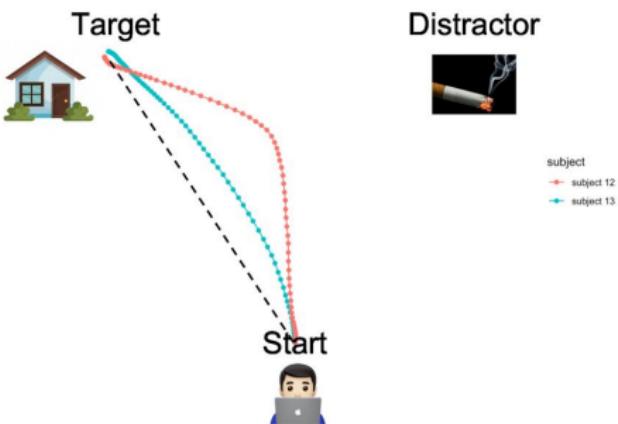


EEG data and computer-mouse tracking (CMT)

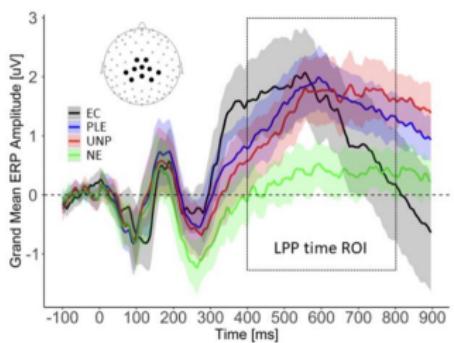
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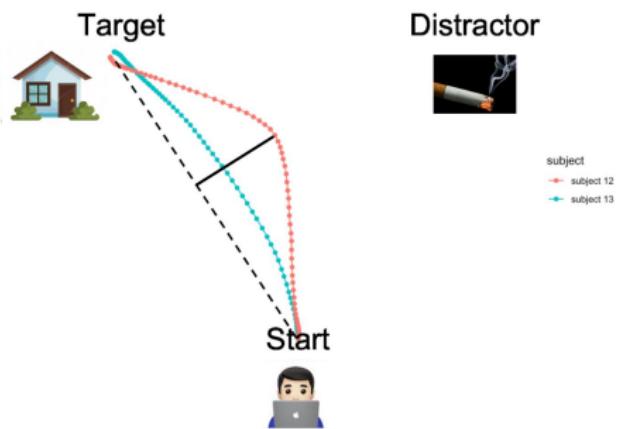
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(simpler and cheaper)



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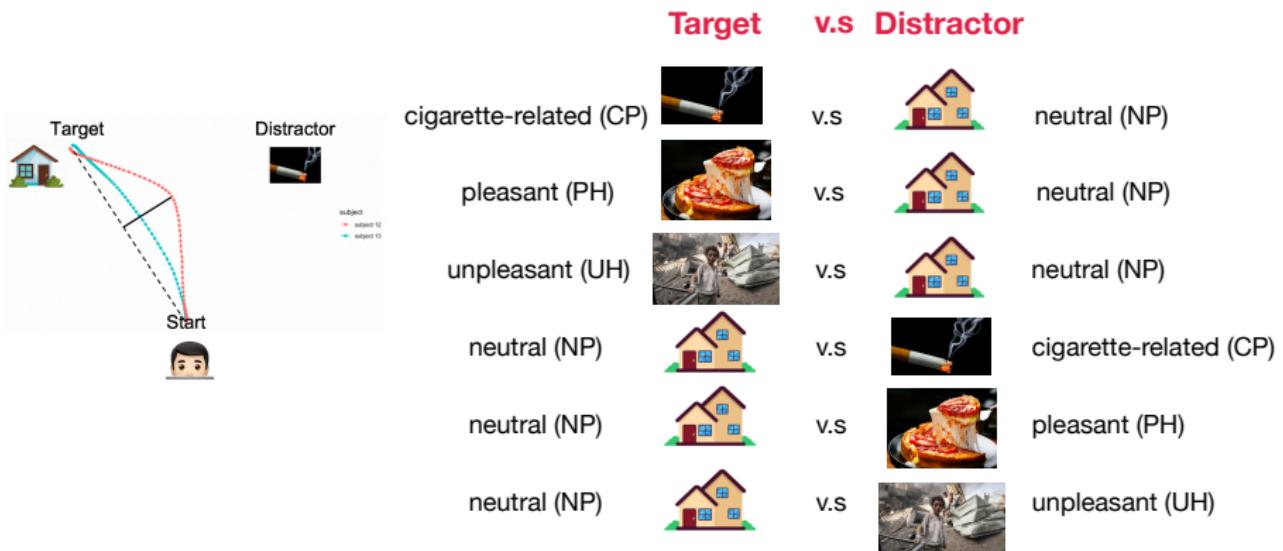


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Trajectories are often summarized using the maximum absolute deviation (MAD), which captures the greatest point of deviation from the idealized straight line

Multiple-experiment conditions

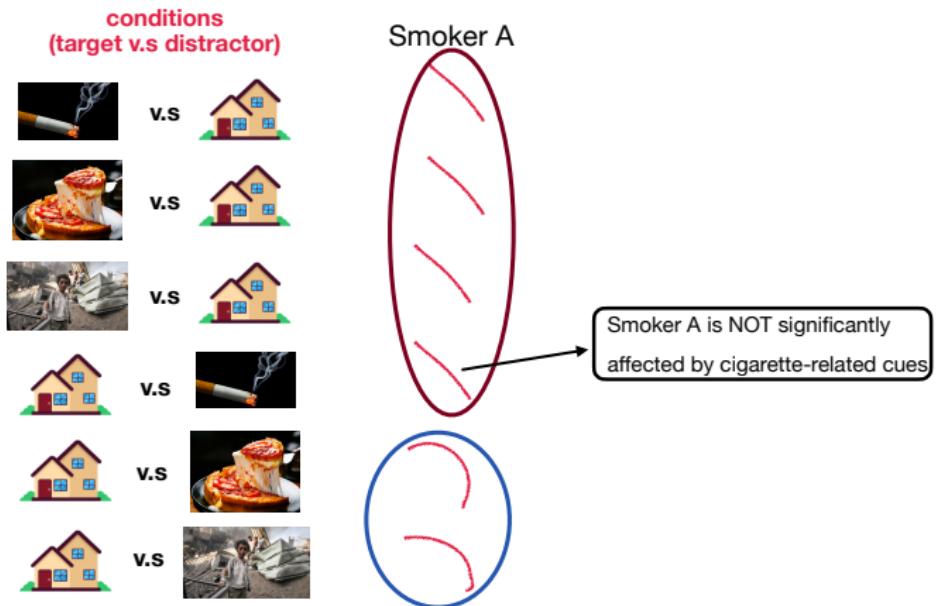


- Analyzing a smoker's trajectories under different conditions helps us understand the individual's degree of response competition to alternatives and the individual's decision-making processes.

- 👉 Clustering Computer Mouse tracking across subject using the known clustering of subjects based on EEG data as reference

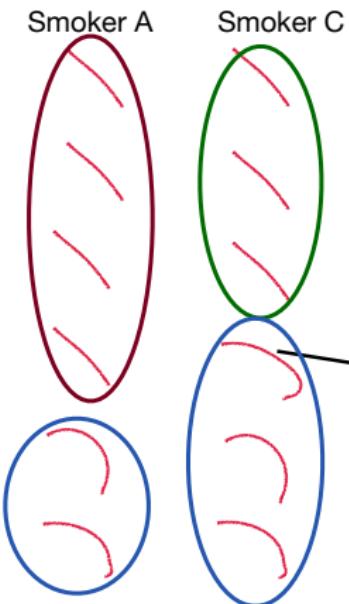
- 👉 Clustering Computer Mouse tracking across subject **using** the known clustering of subjects based on EEG data as **reference**
- 👉 The idea is then to try to inform the **probability of a partition** of the data based on available information on the partition of another set of data
- 👉👉 We want to work on the space of partition
 - !! we may not have other information on the reference data other than a point estimate of their partition (no experiment, no data)
- In addition, we need to take into consideration the several conditions of the computer-mouse tracking experiment

A smoker's behavioral patterns can be revealed by clustering the smoker's mouse tracking trajectories across experimental conditions.

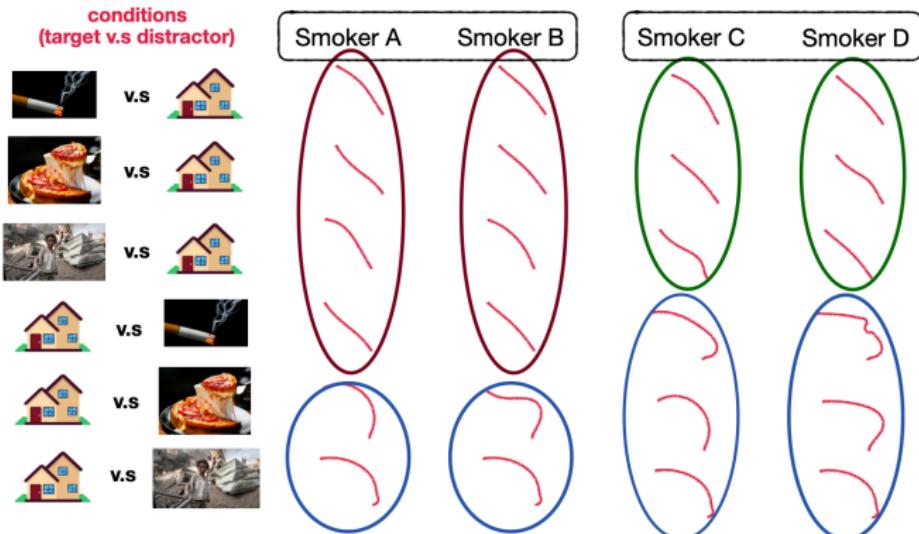


Computer Mouse Tracking data

A smoker's behavioral patterns can be revealed by clustering the smoker's mouse tracking trajectories across experimental conditions.



We are also interested in **clustering the smokers** to identify population subgroups with similar behavior patterns.



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👉 Random partition models

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👉 Random partition models

🤔 It is reasonable to expect that a group of smokers is identified by partitions of conditions that are just more similar (rather than identical) within a group than between groups.

- Goals:**
- 1 Reveal patterns of the smoker's decision-making behaviors via clustering individual responses by experimental conditions;
 - 2 Identify smokers' subgroups within the subgroups that exhibit similar behavioral patterns.
 - 3 incorporate prior domain knowledge on the partitioning of smokers and experiment conditions

Prior Behavioral Neuroscience knowledge and expectations about the partitioning of either smokers or experiment conditions

Computer Mouse Tracking

ERP — Brain Activity

Domain information on 43 smokers

- 43 smokers
 - Group 1: less sensitive to cigarettes
 - Group 2: more sensitive to cigarettes

Domain information on conditions



v.s



v.s



v.s



negligible deviations
from a straight line



v.s



moderate deviations



v.s



large deviations



v.s



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- In product partition models (PPMs), we consider a partition $\rho = \{S_1, S_2, \dots, S_k\}$, of a set of n units, where each S_j is a subset (cluster) of indices. The prior probability of the partition is given by

$$p(\rho) \propto \prod_{j=1}^k h(S_j),$$

where $h(S_j) \geq 0$ is the *cohesion function*.

The cohesion function quantifies how similar the elements within the subset S_j are considered a priori. In other words, it reflects our belief that items within the same cluster should be similar.

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- 📎 Park and Dunson (2010) and Müller, Quintana, and Rosner (2011) modified the cohesion functions in **product partition models** to allow for effects of covariates
- ➡ items closer in the covariate space have a higher probability of being grouped together a priori (nonexchangeable PPMs)

- Smith and Allenby (2020) introduce a **location-scale partition** (LSP) prior modifying the Polya-Urn scheme with similarity functions, incorporating the distance from a reference partition, say ρ_n

Possible Strategies: Shrinkage priors

- Smith and Allenby (2020) introduce a **location-scale partition** (LSP) prior modifying the Polya-Urn scheme with similarity functions, incorporating the distance from a reference partition, say ρ_n

$$\theta_i \mid \theta_{<i}, \rho_n, \tau \sim w_0(\rho_n, \tau) G_0(\theta_i) + \sum_{k=1}^{K^{(i)}} w_k(\rho_n, \tau) \delta_{\phi_k^*}(\theta_i)$$

Here $w_0(\cdot)$ and $w_k(\cdot)$ are positive similarity functions that satisfy $w_0(\cdot) + \sum_k w_k(\cdot) = 1$. More specifically, they build a model for auxiliary variables

$$w_0(s_i, \tau) \propto \int \text{Cat}(\xi_1, \dots, \xi_{C^{(i)}+1}) \text{Dir}(\tau_1, \dots, \tau_{C^{(i)}+1}) d\xi = \frac{\tau + 1}{\tau C^{(i)} + \tau + 1} \binom{s_i - C^{(i)} + 1}{\tau}$$

$$w_k(\{s_i, S_k\}, \tau) \propto \int \text{Cat}(\xi_1, \dots, \xi_{C^{(i)}+1}) \text{Dir}(\tau_1^*, \dots, \tau_{C^{(i)}+1}^*) d\xi = \frac{\tau + n_{S_k}^{s_i}}{\tau C^{(i)} + \tau + n_k}$$

with $n_{S_k}^c$ counting the number of elements in S_k equal to c .

Possible Strategies: Shrinkage priors

- 📎 Paganin et al (2021) propose a **centered partition process** which defines the prior on set partitions as proportional to a baseline EPPF multiplied by a penalization term of the type

$$p(\mathbf{c} \mid \mathbf{c}_0, \psi) \propto p_0(\mathbf{c}) e^{-\psi d(\mathbf{c}, \mathbf{c}_0)}$$

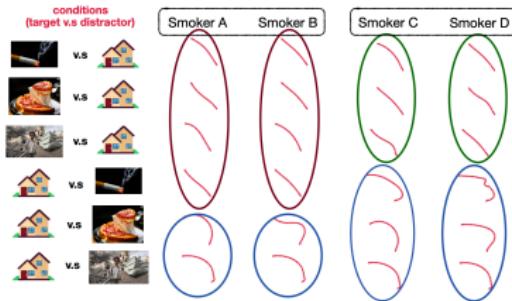
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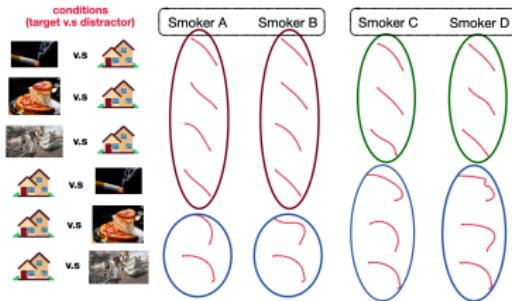
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- 📎 Dhal, Warr and Jensen (2022, [video](#)) and Dhal, Warr and Jensen (arXiv:2312) propose dependent random partitions by **shrinking toward an anchor partition**

- $(I \times J)$ data matrix of entries $Y_{i,j}$'s, from J smokers and I experiment conditions.
- Each $y_{i,j}$ is a MAD computed for the trajectory drawn by smoker j under condition i , for $i = 1, \dots, I$ and $j = 1, \dots, J$.



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- **condition partition** $\pi_j = \{ \pi_{1,j}, \dots, \pi_{I,j} \}$:
a partition of the I conditions MAD's for each smoker j
- **subject partition** $c = \{ c_1, \dots, c_J \}$:
a partition of the J smokers



- For any partition π , the probability function of a SP prior is

$$\pi \sim \text{SP}(\pi | \nu, \lambda, \delta, p_b)$$

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A base (anchor) partition suggested by **available expert information**

A nonnegative **shrinkage parameter** indicates how much we trust the base partition ν for inference

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The diagram illustrates the components of the SP prior formula. The formula is centered at the top, with four arrows pointing from below to its parameters. The top-right arrow points to p_b and is labeled "Permutation parameter". The bottom-left arrow points to ν and is labeled "A base (anchor) partition suggested by available expert information". The bottom-right arrow points to λ and is labeled "A nonnegative shrinkage parameter indicates how much we trust the base partition ν for inference".

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Permutation parameter

A random partition prior distribution,
e.g., CRP or Pitman-Yor process

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- If shrinkage parameter $\lambda = 0$, the partition $\pi \sim p_b$ and π is **independent of the base partition ν** .

The SP prior assigns the joint probabilities $P_{SP}(\pi)$ through a sequential allocation scheme:

$$\Pr_{SP}(\pi_{\delta_t} = s \mid \pi_{\delta_1}, \dots, \pi_{\delta_{t-1}}, \nu_{\delta_1}, \dots, \nu_{\delta_t}, \lambda, p_b)$$

$$\propto \begin{cases} \Pr_b(\pi_{\delta_t} = s \mid \pi_{\delta_1}, \dots, \pi_{\delta_{t-1}}) \times \exp\left(\lambda \frac{\sum_{k=1}^{t-1} I\{\pi_{\delta_k} = s\} I\{\nu_{\delta_k} = \nu_{\delta_t}\}}{\sum_{k=1}^{t-1} I\{\nu_{\delta_k} = \nu_{\delta_t}\}}\right) & \text{if } s \in \{\pi_{\delta_1}, \dots, \pi_{\delta_{t-1}}\}, \\ \Pr_b(\pi_{\delta_t} = s \mid \pi_{\delta_1}, \dots, \pi_{\delta_{t-1}}) \times \exp(\lambda I\{\sum_{k=1}^{t-1} I\{\nu_{\delta_k} = \nu_{\delta_t}\} = 0\}) & \text{if } s \notin \{\pi_{\delta_1}, \dots, \pi_{\delta_{t-1}}\}, \end{cases}$$

$$t = 1, \dots, n.$$

Typically, $\lambda_t = \lambda$, $t = 1, \dots, n$.

For large λ , the SP distribution favors partitions drawn from the CRP that are very similar to the base partition. This behavior is induced by the exponential term, which encourages alignment with the base partition

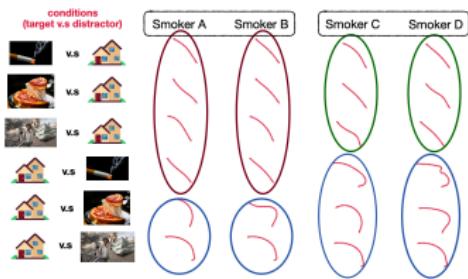
Hierarchical Shrinkage prior

smokers clustering

base partition of smokers given by ERPs results

shrinkage parameter

$$c \sim SP(c | c_0, \tau, \zeta, CRP(\alpha_0))$$



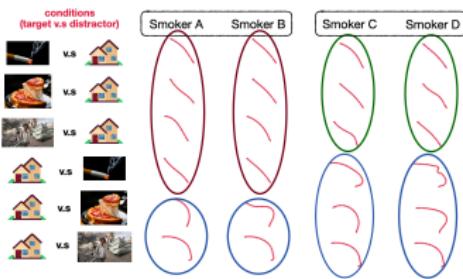
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experiment conditions
clustering for each
smoker

$\pi_j \sim SP\{\pi_j | \nu_j, \lambda, \delta_j, CRP(\beta)\}$

base condition partition for smoker $j = 1, \dots, J$

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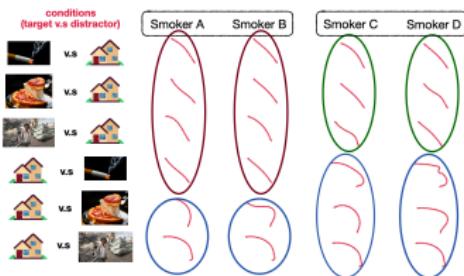
Smokers within a group share a common base partition of conditions ν_k^*

$$\nu_j = \nu_{j'} = \nu_k^* \text{ if } c_j = c_{j'}$$

experiment conditions clustering for each smoker

$$\pi_j \sim \text{SP} \left\{ \pi_j \mid \nu_j, \lambda, \delta_j, \text{CRP}(\beta) \right\}$$

base condition partition for smoker $j = 1, \dots, J$



If smoker j and j' are in the same group according to c , we anticipate they exhibit similar behaviors π_j and $\pi_{j'}$, so we let them share an identical base partition of conditions, i.e., $\nu_j = \nu_{j'} = \nu_k^*$

Hierarchical Shrinkage prior

smokers clustering

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$$c \sim SP(c | c_0, \tau, \zeta, CRP(\alpha_0))$$

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$$\nu_k^* \sim SP(\nu_k^* | \nu_0, \rho, e_k^*, CRP(\beta_0))$$

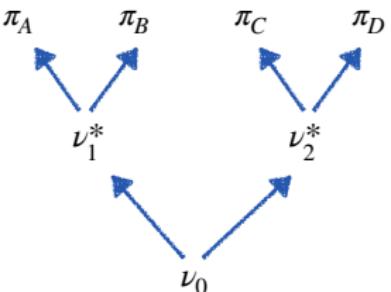
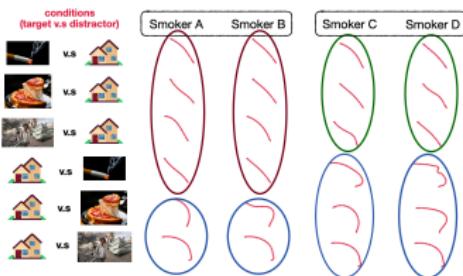
base condition partition given by collaborator on population level

how much we trust the base condition partition ν_0

experiment conditions clustering for each smoker

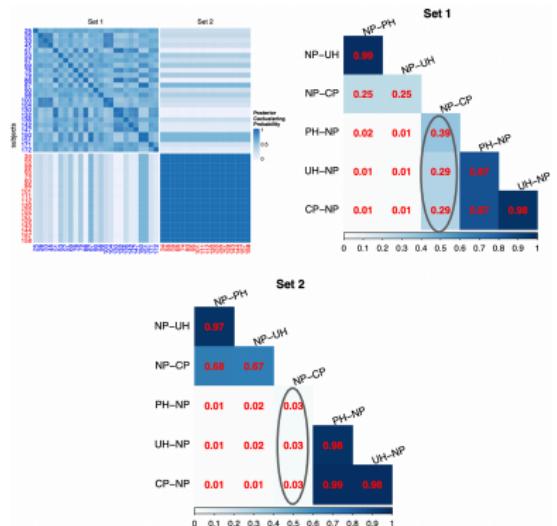
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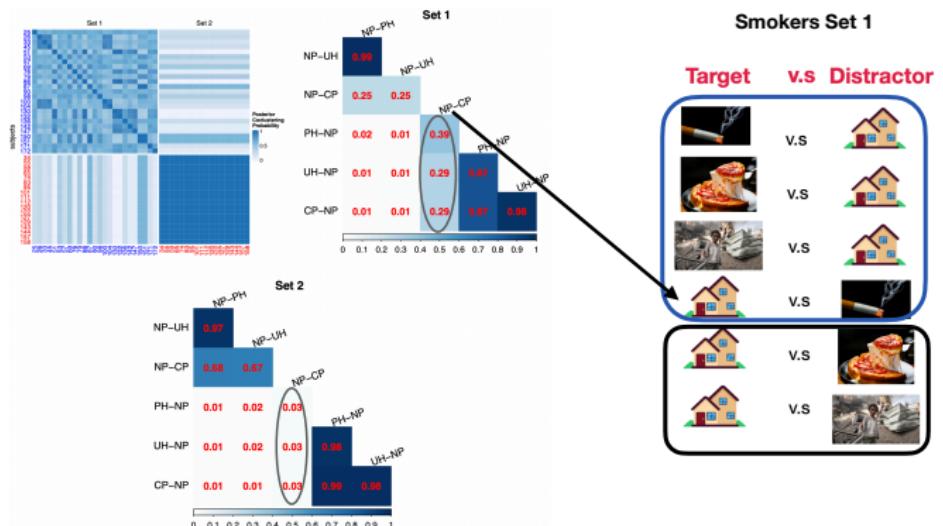
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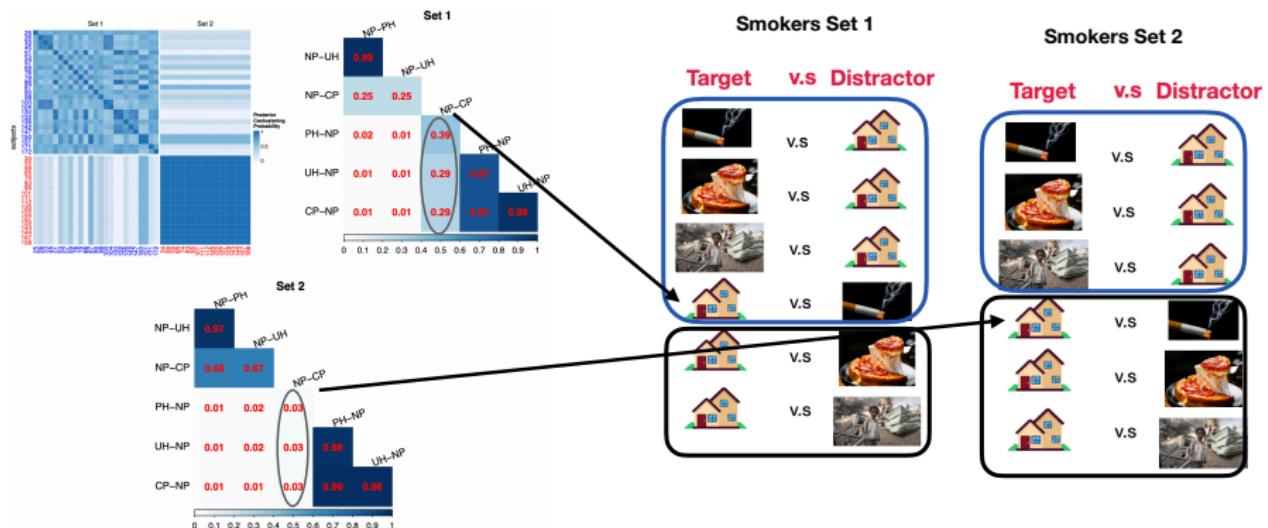
clustering of subjects and mouse-tracking experimental conditions

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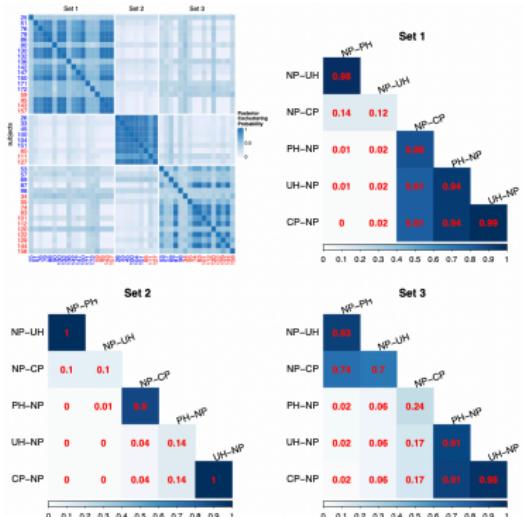


clustering of subjects and mouse-tracking experimental conditions

- **No ERP information** taken into account

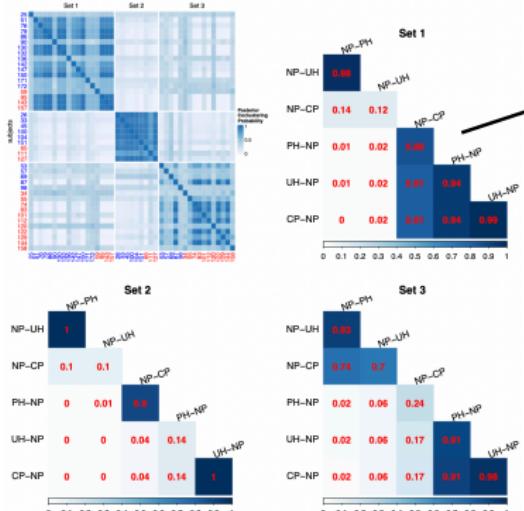
Exploratory Data Analysis

- No ERP information taken into account

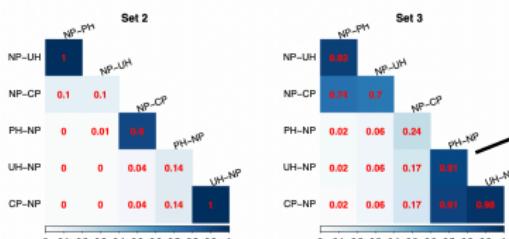
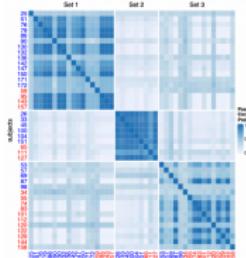


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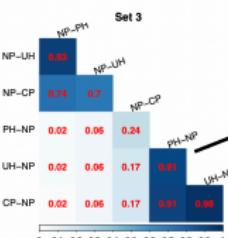
Smokers Set 1
Target v.s Distractor



Smokers Set 3
Target v.s Distractor

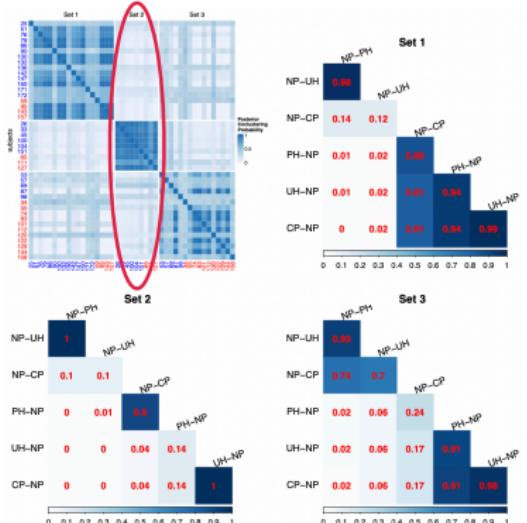


Set 3

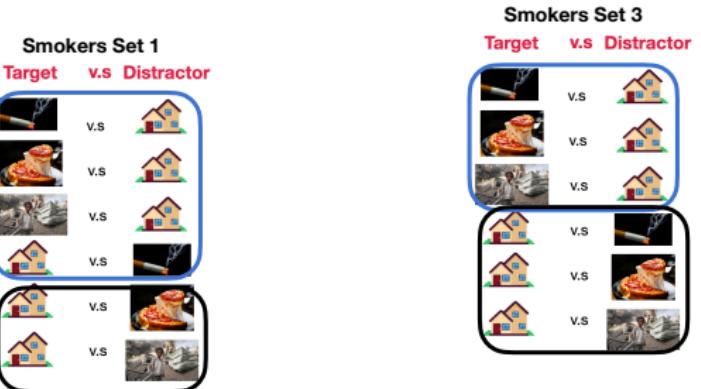


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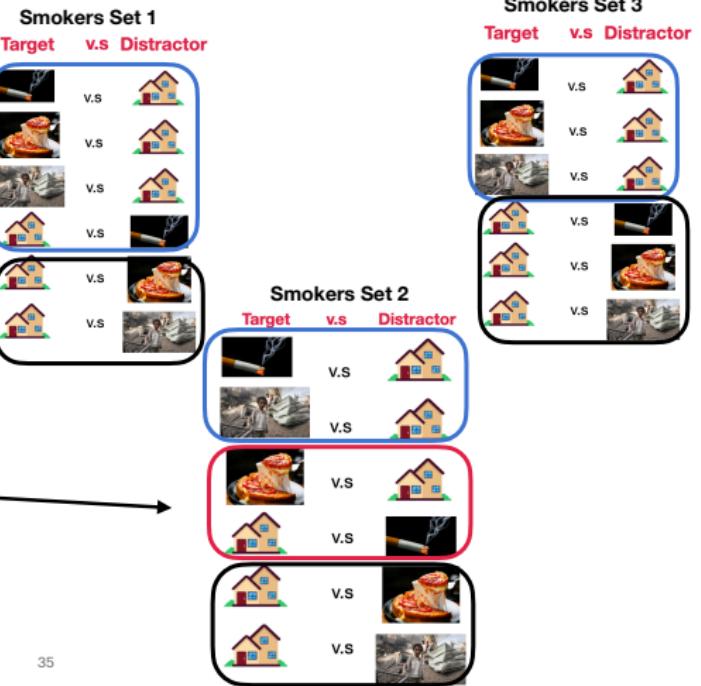
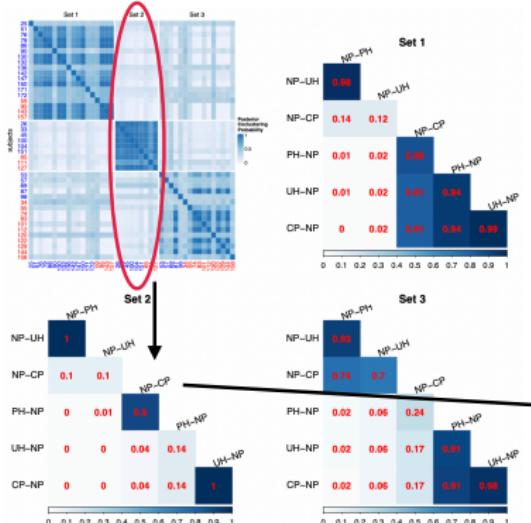


clustering of subjects and mouse-tracking experimental conditions



Exploratory Data Analysis

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clustering of subjects and mouse-tracking experimental conditions

- Incorporate valuable expert information through the base partitions for rows and columns.
- It is important to understand how much either base partition should be trusted.
 - E.g., in considering a base partition of rows, it is important to account for the degree of heterogeneity across all subjects in the original data.

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- Extensions:
 - Move beyond CRP for the baseline partition distributions;
 - Priors over the shrinkage parameters;
 - Directly model the curve trajectories to capture more complex patterns, such as velocity, acceleration, or curvature.
- Ziyi Song, Weining Shen, Marina Vannucci, Alexandria Baldizon, Paul M Cinciripini, Francesco Versace, Michele Guindani, (2024) Clustering computer mouse tracking data with informed hierarchical shrinkage partition priors, *Biometrics*