

Improving the External Validity of Conjoint Analysis: The Essential Role of Profile Distribution

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Conjoint Analysis and Average Marginal Component Effect

- Conjoint analysis: experiments to study multidimensional preferences
- Evaluate two profiles that have randomized characteristics
- Examples: political candidates with varying gender, party, policy, ...
- Main causal estimand: **Average Marginal Component Effect (AMCE)**
 - causal effect of a factor while marginalizing the other factors
 - effect of being female while marginalizing over party, policy...
- AMCE = weighted average of causal effects across **different profiles**
(not just average across respondents)
- **Problem:** Choice of marginalizing distribution has been ignored
- More than 90% of the papers use uniform randomization
 - ~~ equal weights given to all possible profiles
 - ~~ undermine the external validity of the AMCE

Population AMCE (pAMCE)

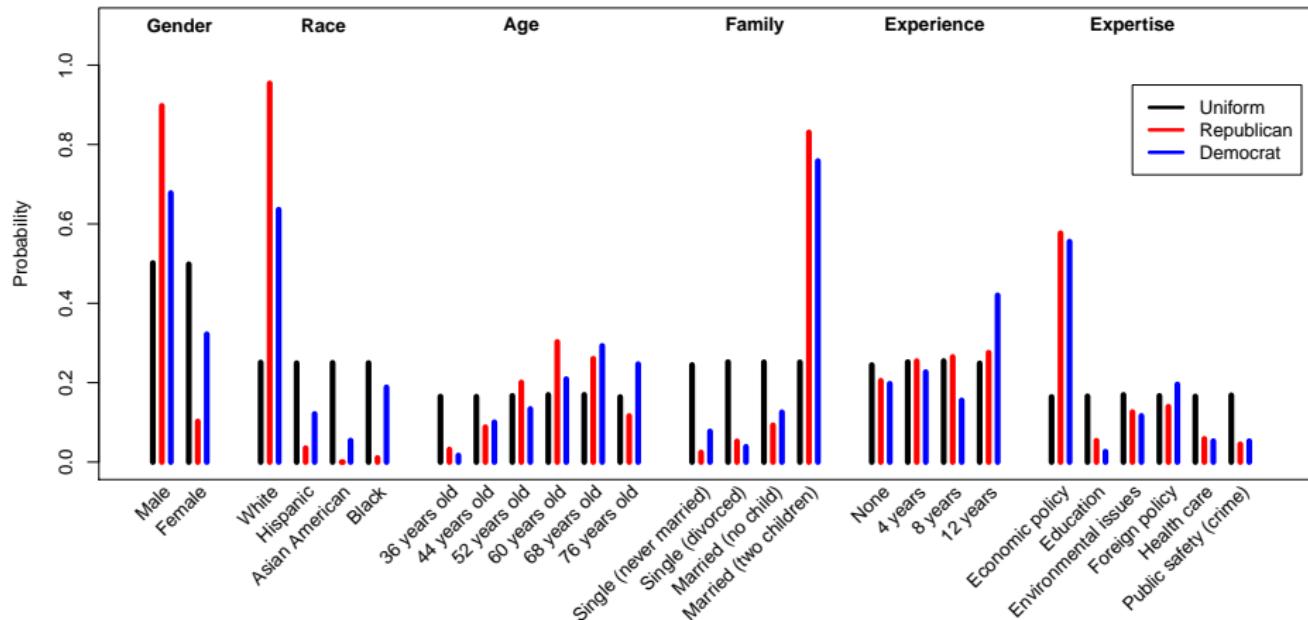
- Use population distribution when marginalizing over the other factors
- Distribution of actual candidates' attributes (party, policy,...)
- Effects in the real world \rightsquigarrow greater external validity

Three Contributions:

- ① Compare the pAMCE with the AMCE based on uniform distribution
- ② Design-based Confirmatory Analysis
 - incorporate population distribution at the design stage
 - new experimental design and estimator
- ③ Model-based Exploratory Analysis
 - explore the pAMCEs using existing conjoint data
 - flexible two-way interaction model

The Effect of Gender on Voter Choice (Ono and Burden, 2018)

- Voters choose hypothetical candidates with varying 13 factors
- Original: all factors are randomized with uniform
- Population distribution based on Republican & Democratic legislators



Potential Outcomes for Conjoint Analysis

- Respondent $i \in \{1, \dots, N\}$ evaluates K choice tasks ($K = 10$)
- Within each task, choose one of J profiles ($J = 2$)
- Profile is characterized by L factors ($L = 13$) (e.g., gender and race)
- Within factor ℓ , the number of levels is D_ℓ (e.g., $D_\ell = 2$ for gender)
- Profile : vector \mathbf{T}_{ijk} of length L
- ℓ th factor of profile: \mathbf{T}_{ijkl} (e.g., Gender = female)
- Potential outcomes: $Y_{ijk}(\mathbf{t}_{ijk}, \mathbf{t}_{i,-j,k})$ (e.g, choose j th candidate)
 - \mathbf{t}_{ijk} : characteristics of candidate j
 - $\mathbf{t}_{i,-j,k}$: characteristics of the other candidates
- Under No Profile-Order Effects: $Y_{ijk}(\mathbf{t}_{ijk}, \mathbf{t}_{i,-j,k}) = Y_{ik}(\mathbf{t}_{ijk}, \mathbf{t}_{i,-j,k})$

Average Marginal Component Effect (AMCE; HHY 2014)

- AMCE: the average causal effect of changing levels within factor ℓ while marginalizing the other factors

$$\tau_\ell(\mathbf{t}_1, \mathbf{t}_0; \Pr(\mathbf{t}_{ijk,-\ell}, \mathbf{t}_{i,-j,k})) = \sum_{\substack{\mathbf{t}_{ijk,-\ell}, \\ \mathbf{t}_{i,-j,k}}} \underbrace{\mathbb{E}[Y_{ik}(\mathbf{t}_1, \mathbf{t}_{ijk,-\ell}, \mathbf{t}_{i,-j,k}) - Y_{ik}(\mathbf{t}_0, \mathbf{t}_{ijk,-\ell}, \mathbf{t}_{i,-j,k})]}_{\text{Causal Effect of factor } \ell \text{ for profile } (\mathbf{t}_{ijk,-\ell}, \mathbf{t}_{i,-j,k})} \underbrace{\Pr(\mathbf{t}_{ijk,-\ell}, \mathbf{t}_{i,-j,k})}_{\text{Marginalizing distribution}}$$

- Effect of being female relative to male while marginalizing Party and Abortion stance
- AMCE = an average of
 - DC: Effect of gender for candidate {Democrat, pro-choice}
 - DL: Effect of gender for candidate {Democrat, pro-life}
 - RC: Effect of gender for candidate {Republican, pro-choice}
 - RL: Effect of gender for candidate {Republican, pro-life}
- Weights for each effect \leftarrow marginalizing distribution

Uniform AMCE and Population AMCE

- Uniform randomization + Difference-in-means \rightsquigarrow uniform AMCE

$$u\text{AMCE} = \tau_\ell(t_1, t_0; \Pr^U(\mathbf{t}_{ijk, -\ell}, \mathbf{t}_{i, -j, k})) \quad (1)$$

- $u\text{AMCE} = 0.25\text{DC} + 0.25\text{DL} + 0.25\text{RC} + 0.25\text{RL}$
- **Problem:** equal weights to all profiles regardless of realism
 \rightsquigarrow External validity of the uAMCE is low
- New causal quantity of interest: population AMCE

$$p\text{AMCE} = \tau_\ell(t_1, t_0; \underbrace{\Pr^*(\mathbf{t}_{ijk, -\ell}, \mathbf{t}_{i, -j, k})}_{\text{Target population distribution}}) \quad (2)$$

- $p\text{AMCE} = 0.45\text{DC} + 0.01\text{DL} + 0.01\text{RC} + 0.53\text{RL}$
- Bias of the uAMCE relative to the pAMCE
 - (1) difference between uniform and population distributions
 - (2) causal interaction between the primary factor and the other factors

Design-based Confirmatory Analysis

- Goal: test several pre-specified hypotheses about the pAMCEs
- Idea: incorporate target population distribution at the design stage
- Main factors \mathbf{T}^M (e.g., gender and race) \leftarrow estimate the pAMCEs
- Control factors \mathbf{T}^C (e.g., age and party) \leftarrow control variables
- **Mixed Randomization Design:**
 - (1) Randomize main factors \mathbf{T}^M with uniform distributions
 - (2) Randomize control factors \mathbf{T}^C with their population distributions
- More main factors \rightarrow standard error for each pAMCE is larger

Weighted Difference-in-Means

- Weighted difference-in-means estimator

$$\frac{\sum_{i=1}^N \sum_{j=1}^J \sum_{k=1}^K \mathbf{1}\{T_{ijkl} = t_1\} w_{ijkl} Y_{ijk}}{\sum_{i=1}^N \sum_{j=1}^J \sum_{k=1}^K \mathbf{1}\{T_{ijkl} = t_1\} w_{ijkl}} - \frac{\sum_{i=1}^N \sum_{j=1}^J \sum_{k=1}^K \mathbf{1}\{T_{ijkl} = t_0\} w_{ijkl} Y_{ijk}}{\sum_{i=1}^N \sum_{j=1}^J \sum_{k=1}^K \mathbf{1}\{T_{ijkl} = t_0\} w_{ijkl}},$$

where the weights are

$$w_{ijkl} = \underbrace{\frac{1}{\Pr^R(T_{ijkl} | \mathbf{T}_{ijk,-\ell}, \mathbf{T}_{i,-j,k})}}_{\text{Inverse probability of receiving } T_{ijkl}} \times \underbrace{\frac{\Pr^*(\mathbf{T}_{ijk,-\ell}, \mathbf{T}_{i,-j,k})}{\Pr^R(\mathbf{T}_{ijk,-\ell}, \mathbf{T}_{i,-j,k})}}_{\text{Differences in distributions of the other factors}}$$

- Mixed design $\rightarrow w_{ijkl} = 1$ when the only main factor is ℓ
- Mixed randomization + Difference-in-means \rightsquigarrow the population AMCE
- Implement via a weighted linear regression
- Population distributions can be approximated by separate marginal distributions, e.g., $\Pr^*(\text{Gender, Age}) \approx \Pr^*(\text{Gender}) \Pr^*(\text{Age})$
- No approximation error unless strong three-way interaction effects

Model-based Exploratory Analysis

- Goal: explore the pAMCEs using previous conjoint experiments
- Idea: use models to adjust for differences in uniform and population
- Latent utility model (paired-choice):

$$\begin{aligned} \tilde{Y}_{ijk}(\mathbf{T}_{ijk}, \mathbf{T}_{ij'k}) = & \quad \tilde{\alpha} + \underbrace{\sum_{\ell=1}^L \mathbf{x}_{ijk\ell}^\top \tilde{\beta}_\ell + \sum_{\ell,\ell'} (\mathbf{x}_{ijk\ell} \times \mathbf{x}_{ijk\ell'})^\top \tilde{\gamma}_{\ell\ell'}}_{\text{Effect of factors of own profile}} \\ & - \underbrace{\sum_{\ell=1}^L \mathbf{x}_{ij'k\ell}^\top \tilde{\beta}_\ell - \sum_{\ell,\ell'} (\mathbf{x}_{ij'k\ell} \times \mathbf{x}_{ij'k\ell'})^\top \tilde{\gamma}_{\ell\ell'}}_{\text{Effect of factors of the opponent}} + \underbrace{\sum_{\ell=1}^L (\mathbf{x}_{ijk\ell} \times \mathbf{x}_{ij'k\ell})^\top \tilde{\delta}_{\ell\ell}}_{\text{Interaction across profiles}} + \tilde{\epsilon}_{ijk}, \end{aligned}$$

- Linear probability model (Egami and Imai, 2019)

$$\begin{aligned} \Pr(Y_{ik} = 1 \mid \mathbf{T}_{ijk}, \mathbf{T}_{ij'k}) = & \quad \alpha + \sum_{\ell} (\mathbf{x}_{ijk\ell} - \mathbf{x}_{ij'k\ell})^\top \underbrace{\beta_\ell}_{\text{Main effects}} \\ & + \sum_{\ell,\ell'} (\mathbf{x}_{ijk\ell} \times \mathbf{x}_{ijk\ell'} - \mathbf{x}_{ij'k\ell} \times \mathbf{x}_{ij'k\ell'})^\top \underbrace{\gamma_{\ell\ell'}}_{\text{within-profile Interaction}} + \sum_{\ell} (\mathbf{x}_{ijk\ell} \times \mathbf{x}_{ij'k\ell})^\top \underbrace{\delta_{\ell\ell}}_{\text{cross-profile Interaction}} \end{aligned}$$

Estimating the Population AMCE

- Estimating the pAMCE

$$\widehat{\tau}_\ell^*(t_1, t_0) = \underbrace{\widehat{\beta}_{\ell 1}}_{\text{Main effect}} + \sum_{\ell'=1}^L \sum_{d=1}^{D_{\ell'}-1} \underbrace{\widehat{\gamma}_{\ell 1 \ell' d}}_{\substack{\text{Interaction} \\ \text{effects}}} \times \underbrace{\Pr^*(T_{ijk\ell'} = d)}_{\substack{\text{Marginal distribution} \\ \text{of population}}}$$

- Bias of the uAMCE relative to the pAMCE (decomposition)

$$\widehat{\text{Bias}} = \sum_{\ell'=1}^L \widehat{\text{Bias}}_{\ell'} = \sum_{\ell'=1}^L \sum_{d=1}^{D_{\ell'}-1} \underbrace{\widehat{\gamma}_{\ell 1 \ell' d}}_{\substack{\text{Interaction} \\ \text{effects}}} \underbrace{\{\Pr^*(T_{ijk\ell'} = d) - \Pr^{\text{U}}(T_{ijk\ell'} = d)\}}_{\text{Difference in distributions}}$$

- Standard errors are large ← Need to estimate many interactions
- Regularization to collapse redundant levels ← Improve efficiency

The Effect of Gender on Voter Choice: Ono & Burden (2018)

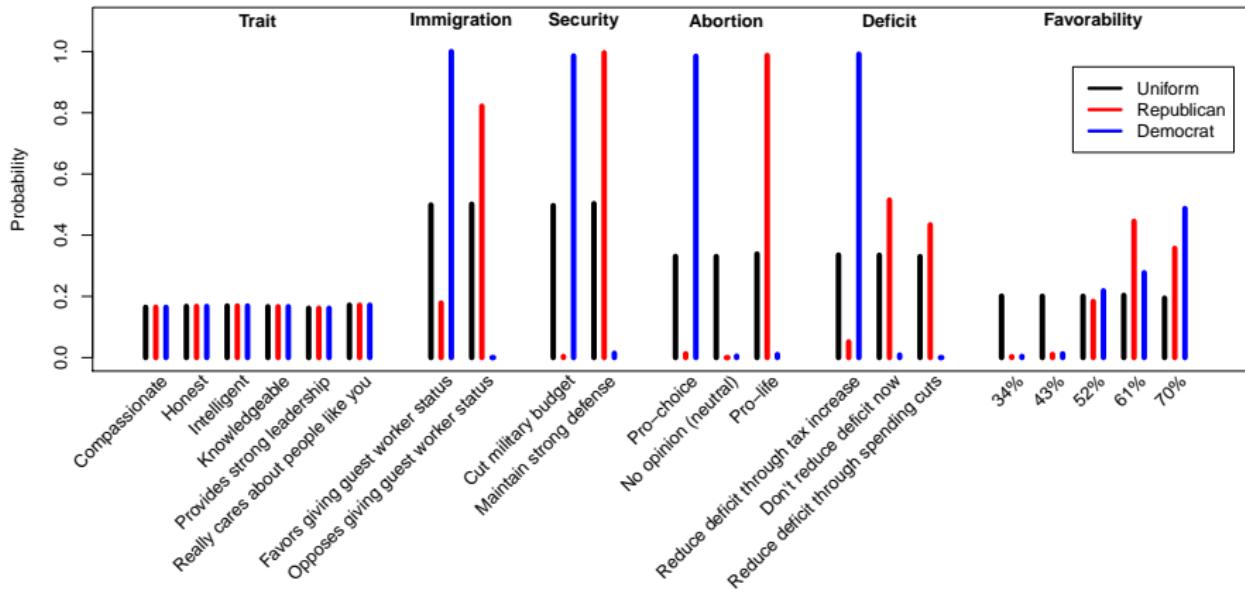
- Voters choose one of the two hypothetical candidates
- Candidates differ in 13 factors
Gender, Race, Age, Family, Experience, Expertise, Trait, Party, Office, Immigration Policy, Security Policy, Abortion Policy, Deficit Policy, Favorability Rating
- Each factor is randomized with uniform distribution
- Interested in the AMCE of being female relative to male

Two Main Findings from Original Analysis:

- (1) Find the uniform AMCE to be a small negative effect
→ Bias against female candidates exists but small
 - (2) The uniform AMCE is negative for presidential candidates
but no effect for congressional candidates
→ Bias against female candidates only for presidential candidates
- Revisit the findings by estimating the population AMCEs

Building Population Distributions

- Target Population: 115th U.S. Congress
- Data sources: (1) biographical dataset (2) electoral returns (3) issue-based scorecards (4) actual bills (5) Wikipedia (6) TheHill.com
- Population distributions separately for Democrats and Republican

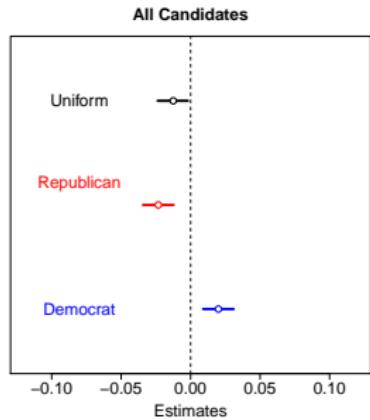


Design-based Confirmatory Analysis

- Goal: test pre-specified hypotheses about the effect of gender
- Mixed randomization design
 - ★ Uniform distribution for Gender — factor of interest
 - ★ Population distribution for the remaining 12 factors
- Original experiments are done with Uniform
- → Illustrate with simulations based on the original conjoint data
 - (1) Fit the two-way interaction model
 - (2) Take estimated coefficients as true data generating process
 - (3) Randomize profiles according to mixed design
 - (4) Estimate the pAMCEs with weighted diff-in-means estimator
- Three Estimands: all profiles, congressional, presidential
- If mixed design had been used, how would conclusions change?

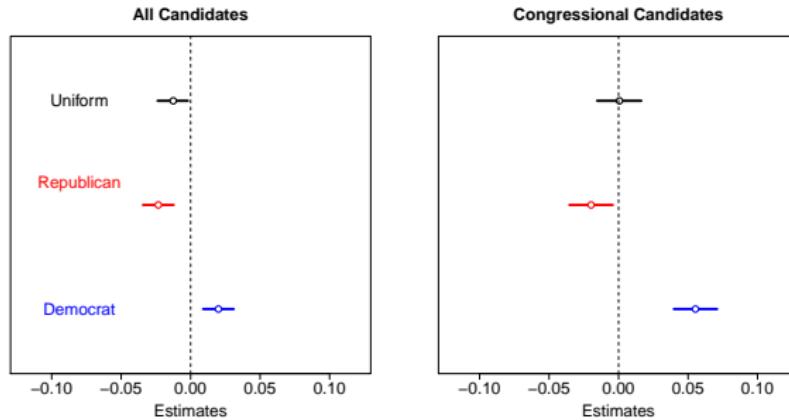
Real World Distribution Leads to Different Findings

Design-based Confirmatory Analysis



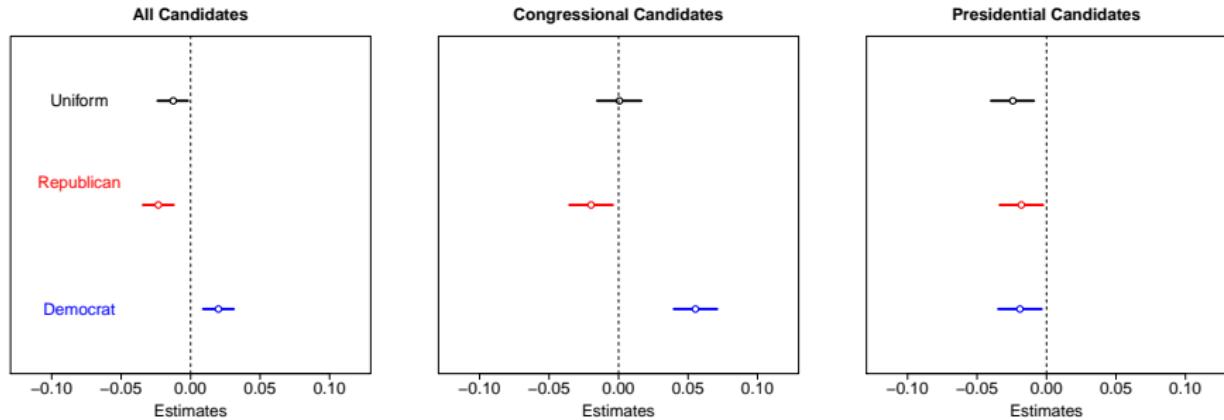
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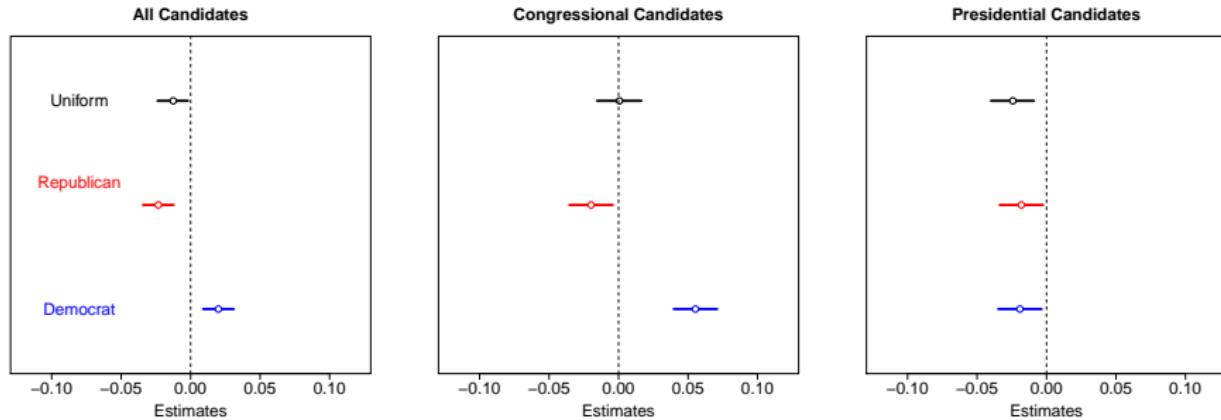


Model-based Exploratory Analysis

- Goal: explore pAMCEs using existing conjoint experiments
- Fit linear probability model with all the two-way interactions
- Regularization to collapse levels → improve efficiency
- Expect that standard errors are larger
- Unpacking the bias of the uAMCE relative to the pAMCE
 - (1) Decompose bias into each factor
 - (2) Is bias caused by interactions or difference in distributions?

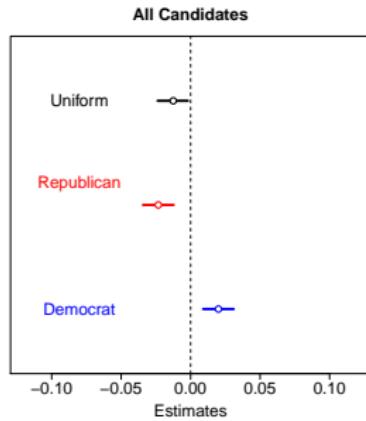
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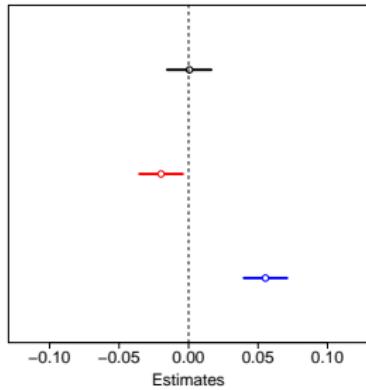


Real World Distribution Leads to Different Findings

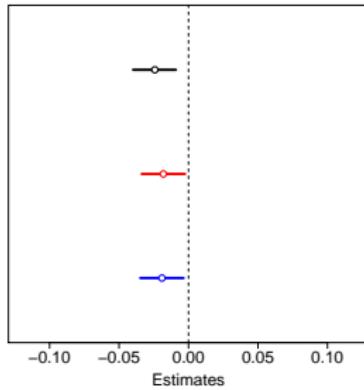
Design-based Confirmatory Analysis



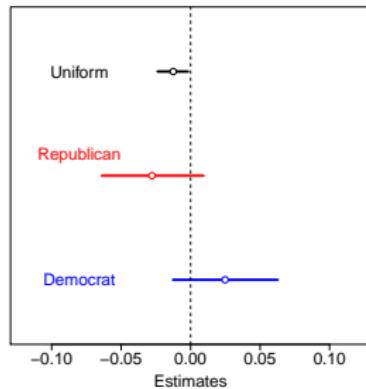
Congressional Candidates



Presidential Candidates

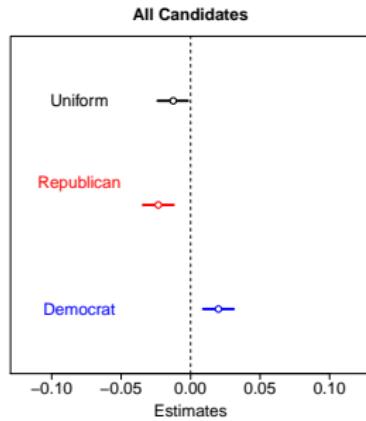


Model-based Exploratory Analysis

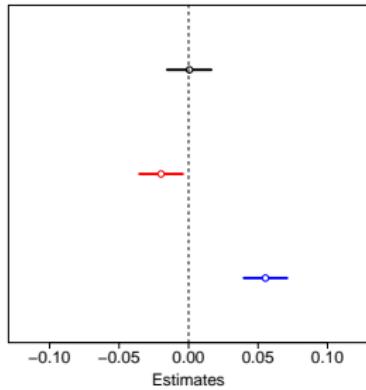


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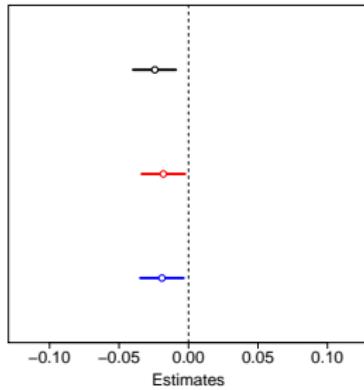
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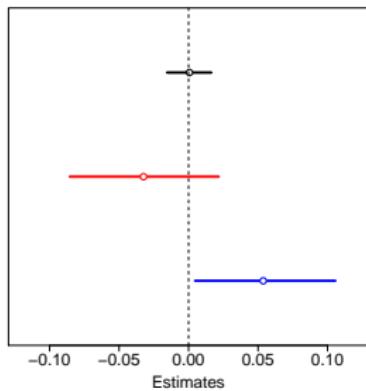
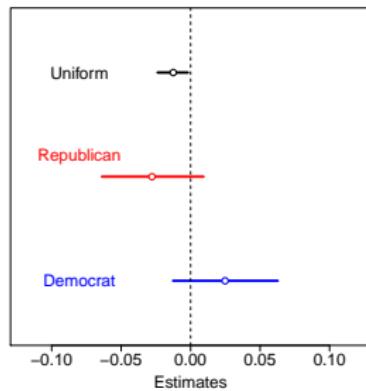
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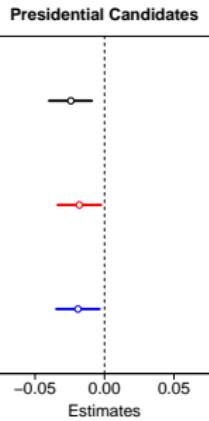
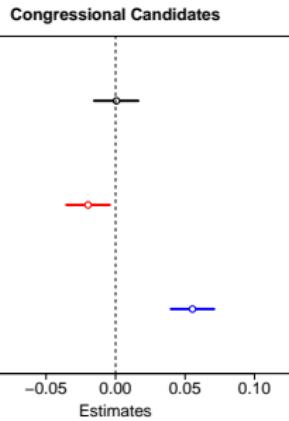
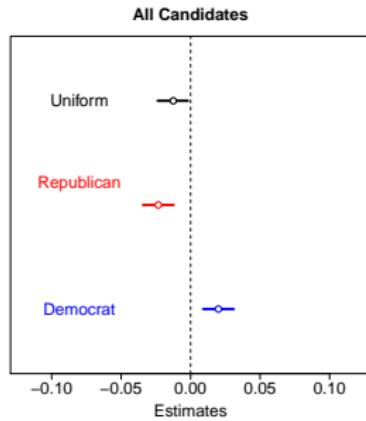


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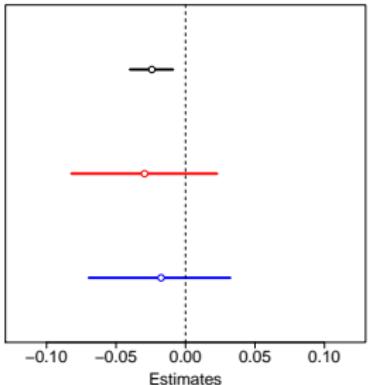
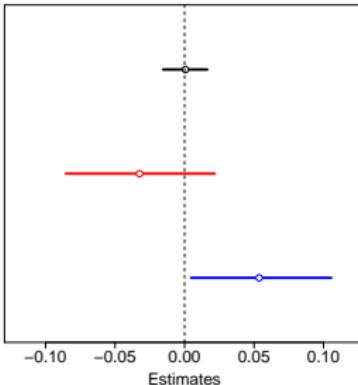
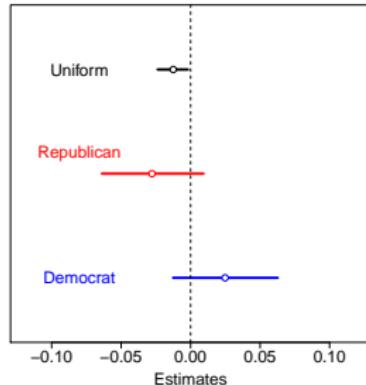


Real World Distribution Leads to Different Findings

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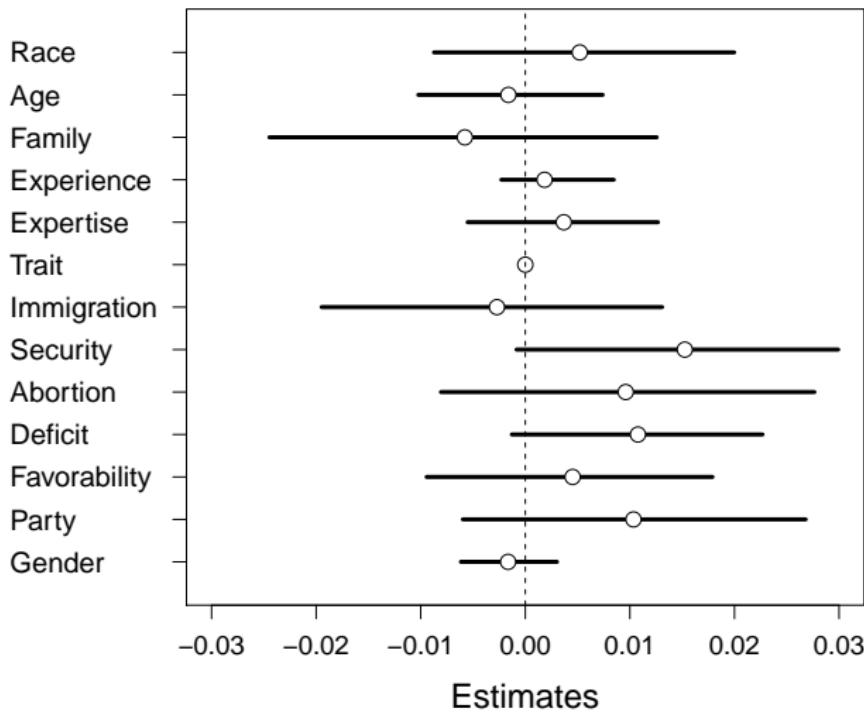


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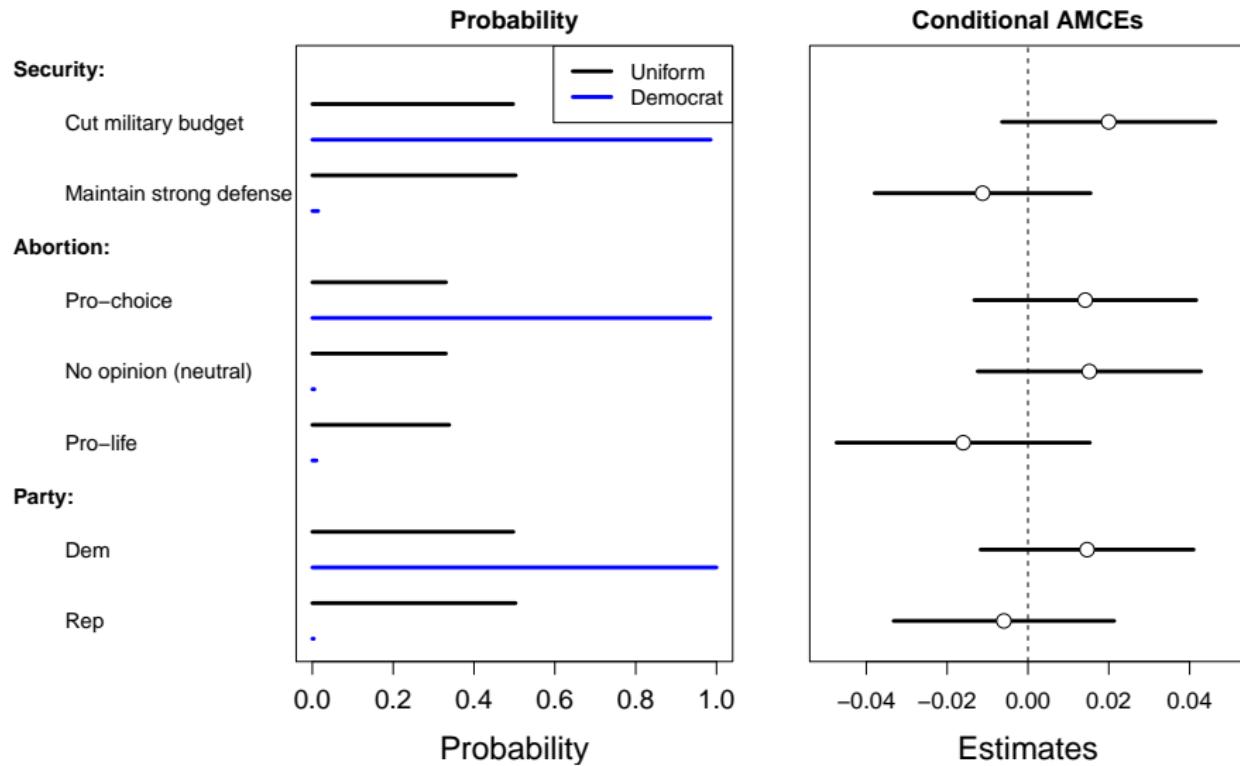


Bias Decomposition for Democratic Congressional pAMCE

Decomposition of Bias:
Democrat – Uniform



Bias in Uniform AMCE Caused By Distribution



Concluding Remarks

- AMCE critically depends on the marginalizing distribution
- **Problem:** Uniform distribution is used for convenience
- uAMCE has low external validity ← equal weights to all profiles
- **Population AMCE:** marginalize over population distribution
 - (1) the real-world distributions of actual politicians' attributes
 - (2) counterfactual distributions of information environments
- Two methods to estimate Population AMCE
 - Developing an open-source software R package
- External validity of treatments (profiles) are essential
 - Conjoint mimics real-world *outcome*
 - Population distribution essential to treatment realism

References

- de la Cuesta, Egami, and Imai. (2019). "Experimental Design and Statistical Inference for Conjoint Analysis: The Essential Role of Population Distribution."
- Egami and Imai. (2019). "Causal Interaction in Factorial Experiments: Application to Conjoint Analysis," *Journal of the American Statistical Association*.

Send comments and suggestions to us

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