

# SuSiE: A Bayesian Method for Variable Selection in Regression

Method Paper Presentation

Rohan Dekate

Carlos Copana

Elaine Hu

# Variable Selection: Challenges & Solutions

- **Challenge:** Scientific conclusions depend on **which variables are selected**, yet selection is hard when predictors are **strongly correlated**.
- **Problem Statement:** How can we **quantify uncertainty** in variable selection under high correlation?
- **Goal:** Identify **causal variants** that truly affect traits — aiming for **scientific insight**, not just predictive accuracy.

## BVSR (Bayesian Variable Selection in Regression)

- Can assess uncertainty even with correlated variables.
- **Computationally intensive** and yields **complex posteriors**.

## SuSiE (Sum of Single Effects):

- Builds on BVSR but is **faster, simpler, and more interpretable**.
- Uses **Iterative Bayesian Stepwise Selection (IBSS)** and **variational approximation** for efficient model fitting.
- Provides **credible sets** — groups of variables that express uncertainty when multiple correlated candidates compete.

# Motivating Toy Example

Consider a linear model:

$$y = \mathbf{X}\boldsymbol{\beta} + \boldsymbol{\varepsilon}, \quad \boldsymbol{\varepsilon} \sim \mathcal{N}(0, \sigma^2 \mathbf{I})$$

with predictors  $x_1 = x_2$  and  $x_3 = x_4$  and  $\beta_1 \neq 0, \quad \beta_4 \neq 0, \quad \beta_2 = \beta_3 = 0$ .

- We wish to infer the statement:  $\beta_1 \neq 0$  **or**  $\beta_2 \neq 0$  **and**  $\beta_3 \neq 0$  **or**  $\beta_4 \neq 0$ .

Existing methods fall short:

- **LASSO / Elastic Net:** Picks one “best” model  $\Rightarrow$  ignores uncertainty.
- **BVSR: Sparse priors** on  $\boldsymbol{\beta}$  capture uncertainty; but **PIPs** lose detail.

**Single-Effect Regression (SER):** Only **one variable** has a **non-zero effect**.

$$\begin{aligned} y &= \mathbf{X}\mathbf{b} + \mathbf{e}, \quad \mathbf{e} \sim \mathcal{N}_n(\mathbf{0}, \sigma^2 \mathbf{I}_n), \quad \mathbf{b} = b\boldsymbol{\gamma}, \quad \boldsymbol{\gamma} \sim \text{Mult}(1, \boldsymbol{\pi}), \quad b \sim \mathcal{N}(0, \sigma_0^2) \\ \boldsymbol{\gamma} \mid \mathbf{X}, \mathbf{y} &\sim \text{Mult}(1, \boldsymbol{\alpha}), \quad b \mid \mathbf{X}, \mathbf{y}, \gamma_j = 1 \sim \mathcal{N}(\mu_j, \sigma_j^2) \end{aligned}$$

- $\boldsymbol{\pi} = (\pi_1, \dots, \pi_p)$  are the prior inclusion probabilities.
- $\boldsymbol{\gamma} \in \{0, 1\}^p$  is  $p$ -vector of indicator variables.
- The single-effect vector  $\mathbf{b}$  has exactly one non-zero element.

# Sum of Single-Effects Regression Model: SuSiE

- **Idea:** Extend the SER model to allow for **multiple effects**.

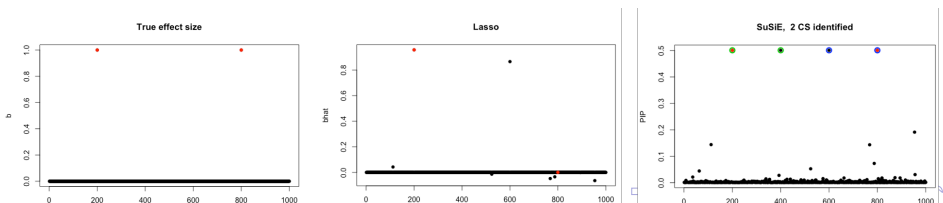
- **Model:**

$$\mathbf{y} = \mathbf{X}\boldsymbol{\beta} + \mathbf{e}, \quad \mathbf{e} \sim \mathcal{N}_n(\mathbf{0}, \sigma^2 \mathbf{I}_n).$$

- Represent the regression coefficient vector as a **sum of single effects**:

$$\boldsymbol{\beta} = \sum_{l=1}^L \mathbf{b}_l, \quad \mathbf{b}_l = b_l \boldsymbol{\gamma}_l, \quad b_l \sim \mathcal{N}(0, \sigma_{0l}^2), \quad \boldsymbol{\gamma}_l \sim \text{Mult}(1, \boldsymbol{\pi}).$$

- When  $L = 1$ , SuSiE reduces to SER; when  $L \ll p$ , it approximates BVSR.
- Each component  $\mathbf{b}_l$  corresponds to one **independent signal** in the data.



# Iterative Bayesian Stepwise Selection (IBSS)

**Goal:** Efficiently fit the SuSiE model using repeated SER updates.

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## Algorithm outline:

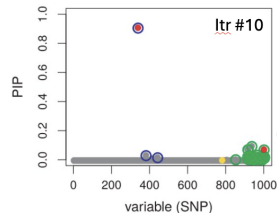
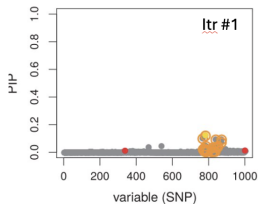
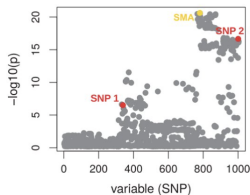
- ❶ Initialize posterior means  $\bar{\mathbf{b}}_l = 0$  for  $l = 1, \dots, L$ .
  - ❷ Repeat until convergence:
    - ❶ For each  $l$ :
      - Compute residuals excluding effect  $l$ :  $\mathbf{r}_l = \mathbf{y} - \mathbf{X} \sum_{l' \neq l} \bar{\mathbf{b}}_{l'}$ .
      - Fit SER to  $\mathbf{r}_l$  and obtain :  $(\alpha_l, \mu_l, \sigma_l^2) = \text{SER}(\mathbf{X}, \mathbf{r}_l; \sigma^2, \sigma_{0l}^2)$ .
      - Update effect estimate:  $\bar{\mathbf{b}}_l = \alpha_l \circ \mu_l$  (element-wise product).
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## Posterior Inference and Key features:

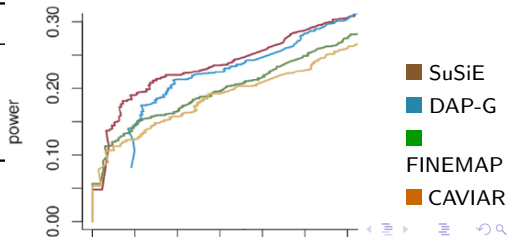
- Under SuSiE we obtain:  $\text{PIP}_j = \Pr(b^j \neq 0 \mid \mathbf{X}, \mathbf{y}) \approx 1 - \prod_{l \in \mathcal{L}} (1 - \alpha_{lj})$
- **Choice of L:** If L exceeds the number of detectable effects in the data, then many of the L credible sets are large.
- Identifiability and label switching  $p(\mathbf{b}_1, \dots, \mathbf{b}_L \mid \mathbf{y}) = p(\mathbf{b}_{v(1)}, \dots, \mathbf{b}_{v(L)} \mid \mathbf{y})$

# Simulation and Numerical Comparison

**Setup:** 6000 simulated datasets comparing SuSiE, DAP-G, FINEMAP, and CAVIAR on credible sets, runtime, and power–FDR trade-off.



Alg.	Time (s)	Credible Sets.
SuSiE	0.64	High power
DAP-G	2.87	Equal or less
FINEMAP	23.0	—
CAVIAR	2907	—



# Limitation and Change Point Detection

## Main Limitation:

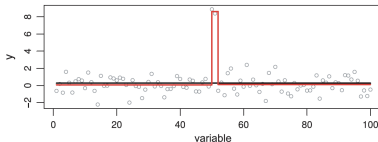
- IBSS uses coordinate-wise updates and can get stuck in **local optima**.
- Struggles to detect signals requiring **joint selection** of multiple correlated effects.

## Example – Change Point Detection:

$$y_t = \mu_t + \varepsilon_t, \quad \varepsilon_t \sim \mathcal{N}(0, \sigma^2), \quad \mathbf{y} = \mathbf{X}\mathbf{b} + \mathbf{e},$$

where  $\mathbf{X}$  encodes step functions marking potential change points.

- SuSiE reformulates this as regression, but IBSS may miss **adjacent change points** when signals cancel each other.



## Future Directions:

- *Better Initialization; Optimization; Model extension*

- G. Wang, A. Sarkar, P. Carbonetto, and M. Stephens (2020). *A simple new approach to variable selection in regression, with application to genetic fine mapping*. *Journal of the Royal Statistical Society, Series B*, **82**(5), 1273–1300. <https://doi.org/10.1111/rssb.12388>