Clustering of spatial scRNA-seq data

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1 Ising model

1.1 Description

Let the expression y of cell i be determined by $y_i = \mu I(z_i = 1) + \epsilon_i$ with priors

$$\mu | \mu_0, \lambda_0 \sim \mathcal{N} \left(\mu_0, \lambda_0^{-1} \right)$$
$$\epsilon_i | \lambda \sim \mathcal{N} \left(0, \lambda^{-1} \right)$$
$$\lambda | \alpha, \beta \sim \text{Gamma} \left(\alpha, \beta \right),$$

and known hyperparameters

$$\mu_0 = \bar{y}$$

$$\lambda_0 = \frac{1}{100}$$

$$\alpha = 1$$

$$\beta = 0.01.$$

Then

$$y_i|z_i, \mu, \lambda \sim \mathcal{N}\left(\mu I(z_i=1), \lambda^{-1}\right).$$

The conditional posterior distributions are given by

$$\mu|\boldsymbol{y},\boldsymbol{z},\lambda \sim \mathcal{N}\left(\frac{\lambda_0\mu_0 + \lambda\sum_{i=1}^n y_i I(z_i=1)}{\lambda_0 + \lambda\sum_{i=1}^n I(z_i=1)}, \left(\lambda_0 + \lambda\sum_{i=1}^n I(z_i=1)\right)^{-1}\right)$$
(1)

$$\lambda | \boldsymbol{y}, \boldsymbol{z}, \mu \sim \text{Gamma}\left(\alpha + \frac{n}{2}, \beta + \frac{\sum_{i=1}^{n} [y_i - \mu I(z_i = 1)]^2}{2}\right).$$
 (2)

Let n be the total number of cells, arranged on a lattice. For every cell $i \in \{1, ..., n\}$, the cell has state $z_i \in \{-1, 1\}$. Using the Ising model, the energy of the lattice system is given by

$$H(oldsymbol{z}) = rac{\gamma}{|\langle i | j \rangle|} \sum_{\langle i | j
angle} z_i z_j,$$

and the probability of the system is given by

$$p(\boldsymbol{z}|\boldsymbol{y}) \propto \exp(H(\boldsymbol{z})) p(\boldsymbol{y}|\boldsymbol{z})$$

where $\langle i j \rangle$ denotes nearest neighbors on the lattice and γ controls the magnitude of the dependence between neighbors.

The Metropolis-Hastings algorithm can be used to explore the lattice space, updating the state for one cell at a time. For each new proposal z', the acceptance probability α is given by

$$\alpha(z', z) = \min \left\{ \frac{\exp(H(z')) p(y|z')}{\exp(H(z)) p(y|z)}, 1 \right\}.$$
(3)

The hyperparameters can be estimated using the following procedure:

- 1. Initialize $\mu = \bar{y}$, $\lambda = \frac{\alpha}{\beta} = 100$, $z_i = -1 \ \forall \ i \in \{1, ..., n\}$.
- 2. Given y, z, λ , sample μ from the conditional distribution given in (1).
- 3. Given y, z, μ , sample λ from the conditional distribution given in (2).
- 4. Given \mathbf{y} , $\{z_2, ..., z_n\}$, μ , λ , sample z_1 (transition from -1 to 1 or 1 to -1) with acceptance probability given by (3).
- 5. Repeat step 4 for all other indices of z.
- 6. Repeat steps 2-5 for N iterations.

1.2 Simulation

All cells on a 100 by 100 lattice are set to have state -1, except for a radius 10 circle and side length 10 square that have state 1. For this simulation $\mu = 5$ and $\lambda = 0.5$.

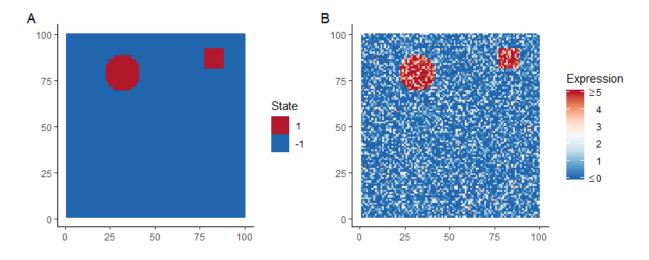


Figure 1: The ground truth (A) and simulated data (B).

One chain is generated for each of three values of γ : 2, 4, 6. 1000 iterations are generated, including a burn-in period of 100 iterations.

The bias is large when $\gamma = 2$, but negligible with a higher smoothing parameter.

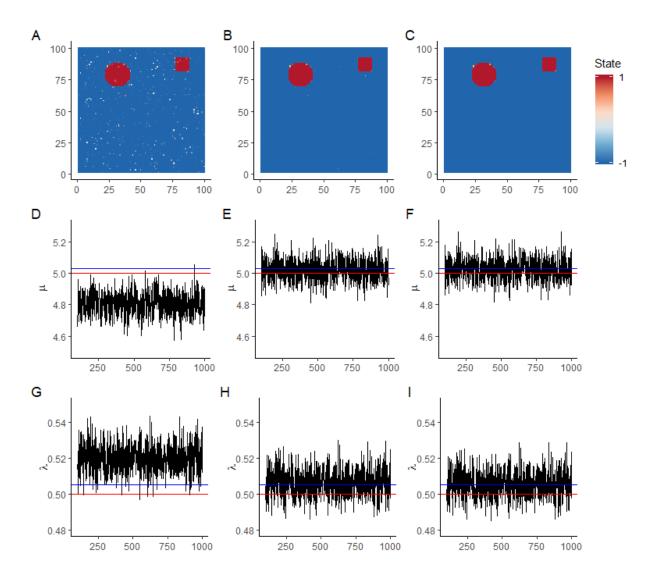


Figure 2: The left, middle, and right columns show the results for $\gamma=2, \gamma=4$, and $\gamma=6$ respectively. A-C show the average state of each cell over 900 iterations. D-F are the trace plots for μ , with the horizontal red line denoting $\mu=5$. G-I are the trace plots for λ , with the horizontal red line denoting $\lambda=0.5$. The horizontal blue lines denote the maximum likelihood estimates.

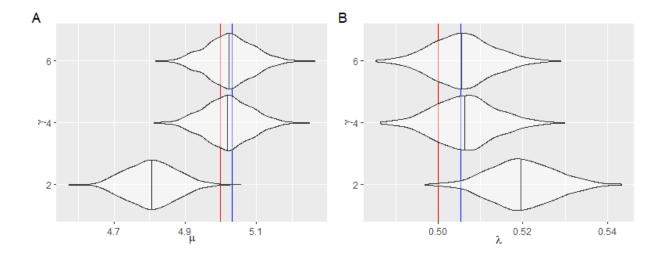


Figure 3: The posterior distributions of μ (A) and λ (B) are shown for each value of γ . The vertical red and blue lines indicate the true values and maximum likelihood estimates respectively of μ and λ .

2 Potts model

2.1 Description

The Potts model is a generalization of the Ising model that allows for more than two states. A cell's state can take values $z_i \in \{1, ..., q\}$. The expression is determined by

$$y_i = \sum_{k=1}^{q} \mu_k I(z_i = k) + \epsilon_i,$$

with priors

$$\mu_{k}|\mu_{0}, \lambda_{0} \sim \mathcal{N}\left(\mu_{0}, \lambda_{0}^{-1}\right) \ \forall \ k \in \{1, ..., q\}$$
$$\epsilon_{i}|\lambda \sim \mathcal{N}\left(0, \lambda^{-1}\right)$$
$$\lambda|\alpha, \beta \sim \operatorname{Gamma}\left(\alpha, \beta\right),$$

and known hyperparameters

$$\mu_0 = \bar{y}$$

$$\lambda_0 = \frac{1}{100}$$

$$\alpha = 1$$

$$\beta = 0.01.$$

Then

$$y_i|z_i, \boldsymbol{\mu}, \lambda \sim \mathcal{N}\left(\sum_{k=1}^q \mu_k I(z_i=k), \lambda^{-1}\right).$$

The conditional posterior distributions are given by

$$\mu_k | \boldsymbol{y}, \boldsymbol{z}, \lambda \sim \mathcal{N} \left(\frac{\lambda_0 \mu_0 + \lambda \sum_{i=1}^n y_i I(z_i = k)}{\lambda_0 + \lambda \sum_{i=1}^n I(z_i = k)}, \left(\lambda_0 + \lambda \sum_{i=1}^n I(z_i = k) \right)^{-1} \right)$$
(4)

$$\lambda | \boldsymbol{y}, \boldsymbol{z}, \boldsymbol{\mu} \sim \operatorname{Gamma}\left(\alpha + \frac{n}{2}, \beta + \frac{\sum_{k=1}^{q} \sum_{i=1}^{n} I(z_i = k) \left[y_i - \mu_k\right]^2}{2}\right).$$
 (5)

Using the Potts model, the energy of the lattice system is given by

$$H(z) = \frac{\gamma}{|\langle i \ j \rangle|} \times 2 \sum_{\langle i \ j \rangle} I(z_i = z_j) - 0.5,$$

and the probability of the system is given by

$$p(\boldsymbol{z}|\boldsymbol{y}) \propto \exp(H(\boldsymbol{z})) p(\boldsymbol{y}|\boldsymbol{z}).$$

For each proposal z', the Metropolis-Hastings acceptance probability is given by

$$\alpha(z', z) = \min \left\{ \frac{\exp(H(z')) p(y|z')}{\exp(H(z)) p(y|z)}, 1 \right\}.$$
(6)

The hyperparameters can be estimated using the following procedure:

- 1. Initialize $\mu_k = \bar{y} \ \forall \ k, \ \lambda = \left[\frac{\sum_{i=1}^n (y_i \bar{y})^2}{n}\right]^{-1}, \ z_i = 1 \ \forall \ i \in \{1,...,n\}.$
- 2. For each k, given y, z, λ , sample μ_k from the conditional distribution given in (4).
- 3. Given y, z, μ , sample λ from the conditional distribution given in (5).
- 4. Given $\mathbf{y}, \{z_2, ..., z_n\}, \boldsymbol{\mu}, \lambda$, sample z_1 (transition from state k to state $k' \in \{1, ..., q\} \setminus \{k\}$) with acceptance probability given by (6).
- 5. Repeat step 4 for all other indices of z.
- 6. Repeat steps 2-5 for N iterations.

2.2 Simulation

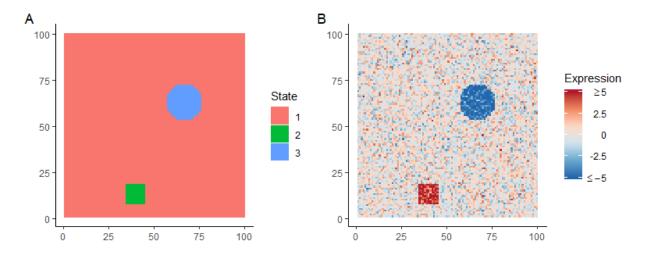


Figure 4: The ground truth (A) and simulated data (B).

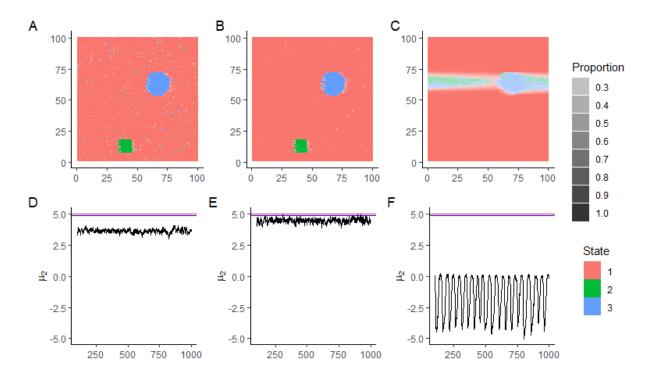


Figure 5: The left, middle, and right columns show the results for $\gamma=2,\,\gamma=3,$ and $\gamma=4$ respectively. A-C show the average state of each cell over 900 iterations. D-F are the trace plots for μ_2