Transition Models

 $t'_{ij}s$ are assumed to be equally spaced.

Let
$$H_i = \{y_k, k = 1, \dots, j - 1\}.$$

Consider

$$f(y_{ij} \mid H_{ij}, \alpha, \beta) = \exp \left\{ \frac{y_{ij} - \psi(\theta_{ij})}{\phi} + c(y_{ij}, \phi) \right\},$$

where $\psi(\theta_{ij})$ and $c(y_{ij}, \phi)$ are known functions.

One has

$$\mu_{ij}^{c} = E\left[y_{ij} \mid H_{ij}\right] = \psi'\left(\theta_{ij}\right)$$

and

$$v_{ij}^{c} = V\left[y_{ij} \mid H_{ij}\right] = \psi''\left(\theta_{ij}\right)\phi$$

with

$$h(\mu_{ij}^c) = x_{ij}^T \beta + \sum_{r=1}^s f_r(H_{ij}; \alpha)$$
 for suitable functions $f_r(\cdot)'s$,

and

$$v_{ij}^c = v(\mu_{ij}^c)\phi.$$

Problem 1. Fitting transition models: (A markov model of order q)

Ву

$$L_i(y_{i1}, \dots, y_{im_i}) = f(y_{i1}, \dots, y_{iq}) \prod_{j=q+1}^{m_i} f(y_{ij} \mid y_{ij-1}, \dots, y_{ij-q}), i = 1, \dots, n,$$

one can get the likelihood function

$$L(\alpha, \beta) = \prod_{i=1}^{n} f(y_{i1}, \dots, y_{iq}) \prod_{j=q+1}^{m_i} f(y_{ij} \mid H_{ij}, \alpha, \beta),$$

where

$$H_{ij} = \{y_{i\,j-1}, \cdots, y_{i\,j-q}\}.$$

Since the term $f(y_{i1}, \dots, y_{iq})$ is always unavailable, the estimators of (α, β) are obtained via maximizing the conditional likelihood

$$\prod_{i=1}^{n} \prod_{j=q+1}^{m_{i}} f\left(y_{ij} \mid H_{ij}, \alpha, \beta\right).$$

Let $\theta = (\alpha, \beta)$.

Show that the log-conditional likelihood or conditional score function has the form

$$S^{c}(\theta) = \sum_{i=1}^{n} \sum_{j=(q+1)}^{m_{i}} \frac{\partial \mu_{ij}^{c}}{\partial \theta} v_{ij}^{c-1} (y_{ij} - \mu_{ij}^{c}).$$

$$L^{c}(\theta) = \prod_{i=1}^{n} \prod_{j=q+1}^{m_{i}} f\left(y_{ij} \mid H_{ij}, \alpha, \beta\right).$$

$$l^{c}(\theta) = \ln L^{c}(\theta) = \frac{\sum_{i=1}^{n} \sum_{j=(q+1)}^{m_{i}} (y_{ij}\theta_{ij} - \psi(\theta_{ij}))}{\phi} + \sum_{i=1}^{n} \sum_{j=(q+1)}^{m_{i}} c(y_{ij}, \phi).$$

We have

$$S^{c}(\theta) = \frac{\partial l^{c}(\theta)}{\partial \theta} = \frac{\sum_{i=1}^{n} \sum_{j=(q+1)}^{m_{i}} (y_{ij} - \psi'(\theta_{ij}))}{\phi}$$
$$= \sum_{i=1}^{n} \sum_{j=(q+1)}^{m_{i}} \frac{1}{\phi} (y_{ij} - \mu_{ij}^{c})$$
$$= \sum_{i=1}^{n} \sum_{j=(q+1)}^{m_{i}} \frac{\partial \mu_{ij}^{c}}{\partial \theta} v_{ij}^{c}^{-1} (y_{ij} - \mu_{ij}^{c}),$$

where

$$\mathbb{E}[y_{ij} \mid H_{ij}] = \psi'(\theta_{ij}) \triangleq \mu_{ij}^c,$$

$$\mathbf{V}[y_{ij} \mid H_{ij}] = \psi''(\theta_{ij})\phi = \frac{\partial \mu_{ij}^c}{\partial \theta}\phi \triangleq v_{ij}^c \implies \frac{1}{\phi} = \frac{\partial \mu_{ij}^c}{\partial \theta}v_{ij}^{c^{-1}}.$$

Problem 2. Ordered Categorical data

Y: ordinal response with categories labeled $1, 2, \dots, k$.

Let

$$F(a \mid x) = P(Y \le a \mid x),$$

where $a = 1, \dots, (k - 1), x = (x_1, \dots, x_p)^T$.

Proportional odds model:

logit
$$F(a \mid x) = \theta_a + x^T \beta$$
, $a = 1, \dots, (k-1)$.

Define $Y^* = (Y_1^*, \cdots, Y_{k-1}^*)$ with $Y_a^* = 1_{(Y \le a)}$.

Then,

$$\operatorname{logit} F(a \mid x) = \operatorname{logit} P\left(Y_a^* = 1 \mid x\right).$$

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Example:

Assume that

logit
$$P(Y_j \le b \mid Y_{ij-1} = a) = \theta_{ab} + x_i^T \beta_a, \quad a, b = 1, \dots, (k-1).$$

It can be derived that

$$\operatorname{logit} P(Y_{ij} \leq b \mid Y_{ij-1}^* = y_{ij-1}^*) = \theta_b + \sum_{l=1}^{k-1} \alpha_{lb} y_{i(j-1)l}^* + x_{ij}^T (\beta + \sum_{l=1}^{k-1} r_l y_{i(j-1)l}^*),$$

where $\begin{cases} \theta_{kb} = \theta_b, \\ \alpha_{lb} = \theta_{lb} - \theta_{l+1b}, \\ \beta_k = \beta, \\ r_l = \beta_l - \beta_{l+1} \end{cases}.$

$$Y_{ij} = a$$

$$\Rightarrow Y_{ij}^* = (\underbrace{0 \cdots 0}_{a-1} \underbrace{1 \cdots 1}_{k-a}).$$

$$\begin{aligned} \log it \, P(Y_{ij} \leq b \mid Y_{ij-1}^* = y_{ij-1}^*) &= \theta_b + (\theta_{1b} - \theta_{2b}) y_{i(j-1)1}^* \\ &\quad + (\theta_{2b} - \theta_{3b}) y_{i(j-1)2}^* \\ &\quad + (\beta_2 - \beta_3) y_{i(j-1)2}^* \\ &\quad + \cdots \\ &\quad + (\theta_{a-1b} - \theta_{ab}) y_{i(j-1)(a-1)}^* \\ &\quad + (\beta_{a-1} - \beta_a) y_{i(j-1)(a-1)}^* \\ &\quad + (\beta_{a-1} - \beta_a) y_{i(j-1)(a-1)}^* \\ &\quad + (\beta_a - \beta_{a+1}) y_{i(j-1)a}^* \\ &\quad + (\theta_{a+1b} - \theta_{a+2b}) y_{i(j-1)(a+1)}^* \\ &\quad + (\beta_{a+1} - \beta_{a+2}) y_{i(j-1)(a+1)}^* \\ &\quad + \cdots \\ &\quad + (\theta_{(k-1)b} - \theta_{kb}) y_{i(j-1)(k-1)}^* \end{aligned}$$

$$\log \operatorname{id} P(Y_{ij} \leq b \mid Y_{ij-1} = a) = \theta_b + (\theta_{ab} - \theta_{a+1b}) + x_{ij}^T \{ \beta + (\beta_a - \beta_{a+1}) + (\theta_{a+1b} - \theta_{a+2b}) + (\beta_{a+1} - \beta_{a+2}) + \cdots + (\theta_{(k-1)b} - \theta_{kb}) + (\beta_{k-1} - \beta_{k}) \}.$$

$$\begin{split} \log & \operatorname{logit} P(Y_{ij} \leq b \mid Y_{i\,j-1} = a) = \theta_b + \theta_{ab} - \theta_{kb} + x_{ij}^T \{\beta + \beta_a - \beta_k\} \\ &= \theta_{ab} + x_{ij}^T \beta_a. \end{split}$$

Problem 3. Log-linear transition models for count data

 $Y_{ij} \mid (H_{ij}, x_{ij}) \sim \text{Poissom } (\mu_{ij}^c).$

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Model 1. Wong (1986) proposed that

$$\mu_{ij}^c = \exp\left(x_{ij}^T \beta\right) \left\{1 + \exp\left(-\alpha_0 - \alpha_1 y_{ij-1}\right)\right\},\,$$

 $\alpha_0, \alpha_1 > 0$, where β is the influence of x_{ij} as $y_{ij-1} = 0$.

Remark. When $y_{ij-1} > 0$, μ_{ij}^c decreases as y_{ij-1} increases. A negative association is implied between the prior and current responses.

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Model 2.
$$\mu_{ij}^c = \exp\left(x_{ij}^T \beta + \alpha y_{ij-1}\right)$$
.

Properties:

- 1. μ_{ij}^c increases as an exponential function of time as $\alpha > 0$.
- 2. When $\exp(x_{ij}^T\beta) = \mu$ and $\alpha < 0$, it leads to a stationary process.

$$\mathbb{E}[Y_{ij} \mid (H_{ij}, x_{ij})] = Var[Y_{ij} \mid (H_{ij}, x_{ij})] = \mu_{ij}^c = \exp\left(x_{ij}^T \beta + \alpha y_{ij-1}\right).$$

$$= \mu \exp\left(\alpha y_{ij-1}\right)$$

$$\in [0, \mu].$$

$$\begin{split} \mathbb{E}[Y_{ij}] &= \mathbb{E}[\mu_{ij}^c] = \mathbb{E}[\exp\left(x_{ij}^T \beta + \alpha y_{ij-1}\right)] \\ &= \mu \, \mathbb{E}[\exp\left(\alpha y_{ij-1}\right)] \end{split}$$

Problem 4.

Moded 3.

$$\mu_{ij} = \exp(x_{ij}^T \beta + \alpha \{\ln(y_{ij-1}^*) - x_{ij-1}^T \beta\}),$$

where $y_{i\,j-1}^* = \max\left\{y_{i\,j-1}, d\right\}, 0 < d < 1.$

 $\begin{array}{l} \text{Property:} \left\{ \begin{array}{l} \alpha = 0 : \text{ it reduces to an oedinary log-tinear model.} \\ \alpha < 0 : \text{ negative correlation between } y_{i\,j-1} \text{ and } y_{ij} \\ \alpha > 0 : \text{ positive correlation between } y_{i\,j-1} \text{ and } y_{ij} \end{array} \right. \\ \end{array}$

Application to a size-independent branching process:

$$\exp(x_{ij}^T \beta) = \mu$$

 y_{ij} : the number of individuals in the *i*-th population at generation j $Z_k(y_{i\;j-1})$: the number of offspring for person k in generation (j-1)

For $y_{ij-1} > 0$,

$$y_{ij} = \sum_{k=1}^{y_{i(j-1)}} Z_k(y_{i(j-1)}),$$

where

$$Z_k \overset{iid}{\sim} Poisson\left(\left(\frac{\mu}{y^*_{ij-1}}\right)^{1-\alpha}\right).$$

One can get

$$\mu_{ij}^c = \mu \cdot \left(\frac{y_{i(j-1)}}{\mu}\right)^{\alpha}.$$

Property:

- $\alpha < 0$: the sample paths oscillate back and forth about their long-term average level.
- $\alpha > 0$: the sample paths have sharper peaks and broader valleys.

R10A21126 HW05 Q4

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```
[]: import numpy as np
     import matplotlib.pyplot as plt
[]: | # Function to calculate lamda for Poisson distribution
     def calculate_lamda(yij_prev, mu, alpha):
         return (mu /yij_prev ) ** (1-alpha)
     # Function to generate sample paths
     def generate_sample_paths(num_generations, initial_values, mu, alpha, d):
         num_populations = len(initial_values)
         sample_paths = np.zeros((num_populations, num_generations))
         sample_paths[:, 0] = initial_values
         for j in range(1, num_generations):
             for i in range(num_populations):
                 yij_prev = sample_paths[i, j-1]
                 # print(yij_prev)
                 Zk = np.random.poisson(calculate_lamda(max(yij_prev, d), mu[i],
      →alpha), size = round(yij_prev))
                 yij = np.sum(Zk)
                 sample_paths[i, j] = yij
         return sample_paths
     def simulate_and_plot(num_generations, initial_values, mu, alpha, d):
         # Generate sample paths
         sample_paths = generate_sample_paths(num_generations,initial_values, mu,__
      ⇔alpha, d)
         # Plotting
         plt.figure(figsize=(10, 6))
         for i in range(len(initial_values)):
```

```
plt.plot(sample_paths[i, :], label=f'Population {i + 1}')
    plt.axhline(y=np.average(sample_paths[i, :]), color='grey', u

slinestyle='--')

plt.title(f'Sample Paths for a = {alpha}')
    plt.xlabel('Generation')
    plt.ylabel('Population Size')
    plt.legend(bbox_to_anchor=(1, 1))
```

```
[]: # Parameters
   num_generations = 200  # Number of generations

# Set initial values of population
   initial_values = np.array([800, 500, 300])
   mu = initial_values

d = 0.1

# Parameter alpha
   alpha_values = [-0.9,-0.5,-0.1]

for alpha in alpha_values:
        simulate_and_plot(num_generations, initial_values, mu, alpha, d)

plt.tight_layout()
   plt.show()
```





