

# Bayesian Clustered Ensemble Prediction for Multivariate Time Series<sup>1</sup>

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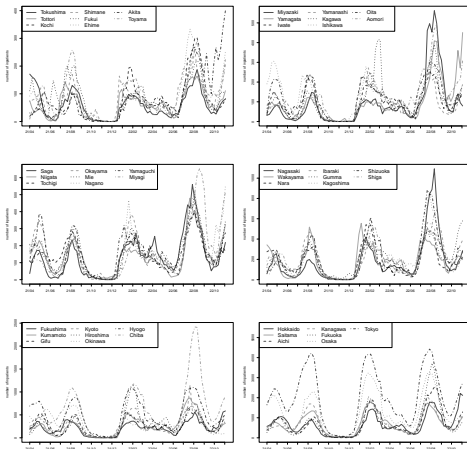
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<sup>1</sup>Joint work with Genya Kobayashi, Yuki Kawakubo, Dongu Han, Taeryon Choi

# Motivating Data

Weekly number of inpatients for COVID-19 by prefecture in Japan  
(from April 2021 to November 2022)



47 prefectures are grouped by the means of the series.

# Motivating Data

## Characteristics of the data

- Multiple time series count data for 47 prefectures
- Cross section and temporal correlation

## Purpose of data analysis

- Precise future prediction on the number of inpatients
- Uncertainty quantification of the prediction

## Strategy

- Ensemble prediction

In stead of relaying on a single model, combining multiple models is known to improve predictive performance.

# Bayesian Predictive Synthesis (BPS)

BPS for univariate time series,  $y_t$  ( $t = 1, \dots, T$ )

- $J$  models (agents) with the predictive density,  $h_{tj}(f_{tj})$  ( $j = 1, \dots, J$ )
- General form of BPS (e.g. West, 1992; McAlinn and West, 2019)

$$p(y_t | \boldsymbol{\theta}_t, 1 : t) = \int \alpha(y_t | \mathbf{f}_t, \boldsymbol{\theta}_t) \left[ \prod_{j=1}^J h_{tj}(f_{tj}) \right] d\mathbf{f}_t$$

- $\boldsymbol{\theta}_t = (\theta_{t0}, \theta_{t1}, \dots, \theta_{tJ})$ : weights for  $J$  predictive models
  - $\mathbf{f}_t = (f_{t1}, \dots, f_{tJ})^\top$ : draw from the predictive densities, regarded as latent factors
  - $\alpha(y_t | \mathbf{f}_t, \boldsymbol{\theta}_t)$ : synthesis function that controls how to combine  $J$  predictions
- BPS includes existing ensemble methods (e.g. Bayesian model averaging) by appropriately specifying the synthesis function.

# Bayesian Predictive Synthesis

## Specification of the synthesis function

- Dynamic linear model with latent factor (McAlinn and West, 2019)

$$\alpha(y_t | \mathbf{f}_t, \boldsymbol{\theta}_t) = \phi\left(y_t; \theta_{t0} + \sum_{j=1}^J \theta_{tj} f_{tj}, \sigma_t^2\right), \quad \boldsymbol{\theta}_t \sim \text{random walk}$$

$\phi(x; a, b)$ : normal density with mean  $a$  and variance  $b$

- BPS for multivariate time series,  $y_{it}$  ( $i = 1, \dots, n; t = 1, \dots, T$ ) (McAlinn et al., 2020)
  - Use multivariate dynamic linear models as the synthesis function.
  - Computationally intensive when  $n$  is large ( $n = 47$  in our example).
- Methodological motivation: Any scalable approach for multivariate time series (of count)?

# Proposal: Mixture of BPS

Idea: (soft) clustering multiple time series in terms of model importance

- $J$  predictive models with the predictive densities  $h_{itj}(f_{itj})$  ( $i = 1, \dots, n; j = 1, \dots, J$ ).
- **Mixture of BPS (MBPS)**

$$\alpha(y_{it}|\mathbf{f}_{it}, \boldsymbol{\theta}_{t,1:K}, \pi_{1:K}) = \sum_{k=1}^K \pi_k \alpha_k(y_{it}|\mathbf{f}_{it}, \boldsymbol{\theta}_{tk}), \quad \sum_{k=1}^K \pi_k = 1$$

- $\alpha_k(y_{it}|\mathbf{f}_{it}, \boldsymbol{\theta}_{tk})$ :  $k$ th component of the synthesis function
- $\boldsymbol{\theta}_{tk}$ : Parameters in the  $k$ th synthesis function
- MBPS reduces the number of parameters in multivariate BPS while sharing cross sectional information

# Proposal: Mixture of BPS

## Structure of MBPS

- For two time series  $i$  and  $i'$  in the same cluster  $k$ , MBPS places the same weight  $\theta_{tjk}$  on  $j$ th model whose forecasts are generally different between  $i$  and  $i'$ .  
 $\Rightarrow$  Clustering  $n$  series in terms of the importance of the  $k$ th model
- Within the same cluster, the contribution of the models to the BPS forecast is the same.
- The weights  $\theta_{tk}$  are estimated from the set of time series belonging to the same component.

# Model Specification

- Consider a synthesis function based on the Poisson distribution as  $y_{it}$  is count in our example.
- Cluster assignment indicator,  $z_i \in \{1, \dots, K\}$  for each time series
- Hierarchical model of MBPS for count response

$$\begin{aligned}
 y_{it} | (z_i = k, \boldsymbol{\theta}_{tk}, \mathbf{f}_{it}) &\sim \text{Po}(\exp(\boldsymbol{\theta}_{tk}^\top \mathbf{F}_{it})), \quad \mathbf{F}_{it} = (1, \mathbf{f}_{it}^\top)^\top \\
 \Pr(z_i = k) &= \pi_k, \quad k = 1, \dots, K, \quad (\pi_1, \dots, \pi_K) \sim \text{Dir}(\mathbf{a}_0) \\
 f_{itj} &\sim N(m_{itj}, s_{itj}^2), \quad \boldsymbol{\theta}_{tk} = \boldsymbol{\theta}_{t-1,k} + \mathbf{e}_{tk}, \quad \mathbf{e}_{tk} \sim N(\mathbf{0}, \boldsymbol{\Sigma}_{tk})
 \end{aligned}$$

- $(m_{itj}, s_{itj}^2)$ : fixed values (prediction and its variance of log-intensity)
- $\mathbf{a}_0 = (1/K, \dots, 1/K)$ : it tends to produce empty clusters.



## Posterior Computation (MCMC)

- Use the negative binomial approximation of the Poisson distribution with a large dispersion parameter (Hamura et al., 2025) and apply the Pólya-gamma (PG) augmentation (Polson et al., 2013).

$$\begin{aligned}\alpha_k(y_{it}|\mathbf{f}_{it}, \boldsymbol{\theta}_{tk}) &\approx \tilde{\alpha}_k(y_{it}|\mathbf{f}_{it}, \boldsymbol{\theta}_{tk}, r) = \frac{\Gamma(y_{it} + r)}{\Gamma(r)y_{it}!} \frac{(e^{\psi_{itk}})^{y_{it}}}{(1 + e^{\psi_{itk}})^{y_{it}+r}} \\ &= 2^{-b_{it}} \exp\{\kappa_{it}\psi_{itk}\} \int_0^\infty \exp\left\{-\frac{\omega_{itk}\psi_{itk}^2}{2}\right\} p(\omega_{itk}|b_{it}, 0) d\omega_{itk}\end{aligned}$$

- $b_{it} = y_{it} + r$ ,  $\kappa_{it} = (y_{it} - r)/2$ ,  $\psi_{itk} = \boldsymbol{\theta}_{tk}^\top \mathbf{f}_{it} - \log r$
  - $\omega_{itk}$  follows the PG distribution.
- The resulting joint distribution can be seen as a Gaussian dynamic linear model.
- ⇒ Forward filtering and backward sampling can be applied.

# Extension: MBPS with Heterogeneous Intercept (MBPSH)

- Intercept in BPS: adjusting the level of inadequacy of ensemble prediction
- MBPS assumes the same intercept within a cluster, which may be restrictive in practice.
- MBPS with heterogeneous intercept (MBPSH)

$$y_{it}|(z_i = k, \boldsymbol{\theta}_{tk}, \mathbf{f}_{it}, u_{it}) \sim \text{Po}(\exp(\boldsymbol{\theta}_{tk}^\top \mathbf{f}_{it} + u_{it})), \quad u_{it}|(z_i = k) \sim N(0, \tau_{tk}^2),$$

$$\tau_{tk}^2 = \frac{\beta_\tau}{\gamma_t} \tau_{t-1,k}^2, \quad \gamma_t \sim \text{Beta}\left(\frac{\beta_\tau n_{t-1}}{2}, \frac{(1 - \beta_\tau) n_{t-1}}{2}\right).$$

It can be regarded as an intermediate model between the univariate BPS and MBPS.

# Analysis of COVID-19 Hospitalization in Japan

- Total number of inpatients including severe conditions, for  $n = 47$  prefectureds in Japan
- Total data period: 2020/05/07 – 2022/11/23 (134 weeks)
- $J = 4$  (number of prediction models),  $K = 47$  (maximum number of clusters)
- **Prediction steps**
  - The four agent models are estimated using the first 50 weeks up to 2021/04/14.
  - MBPS is first run using the data from 2021/04/21 to 2022/03/30 (50 weeks) to produce one step (week) ahead forecast.
  - By expanding the window of past data, the one step forecasts from 2022/04/06 to 2022/11/23 (34 weeks).

# Agent Models

- **Model 1:** Poisson dynamic generalized linear model (DGLM):

$$y_{it} \sim \text{Po}(\lambda_{it}), \quad \log \lambda_{it} = \mathbf{x}_{it}^{\top} \boldsymbol{\beta}_{it}, \quad \boldsymbol{\beta}_{it} \sim N(\boldsymbol{\beta}_{i,t-1}, \mathbf{V}_{it})$$

where  $\mathbf{x}_{it} = (1, \tilde{l}_{it}, \tilde{l}_{it}^2)$  and  $\tilde{l}_{it}$  is the log of the 7 days lag of the 14 days moving average of the number of infected.

- **Model 2:** Poisson generalized additive model (GAM):

$$y_{it} \sim \text{Po}(\lambda_{it}), \quad \log \lambda_{it} = \mu + s(\tilde{l}_{it}) + s(t),$$

where  $s$  denotes the smoothing splines. The model is fitted by `gam`.

# Agent Models

- **Model 3:** Poisson integer autoregressive model (INAR):

$$y_{it} \sim \text{Po}(\lambda_{it}), \quad \log \lambda_{it} = \gamma_i y_{i,t-1} + \mathbf{x}_{it}^\top \boldsymbol{\beta}_i$$

where  $\mathbf{x}$  is the same as DGLM. The model is fitted by `tscount`.

- **Model 4:** Power-weighted Poisson-SIHR model:

$$p(y_{i,1:t} | \lambda_T) = \prod_{s=0}^T \left[ \frac{\lambda_{i,T-s}^{y_{i,T-s}} e^{-\lambda_{i,T-s}}}{y_{i,T-s}!} \right]^{\rho_s}$$

$\lambda_{i,T-s}$  is the solution of the SIHR  
(susceptible-infectious-hospitalized-recovered) model:

$$\begin{aligned} dS/dt &= -\alpha I \cdot S/n, & dI/dt &= \alpha I \cdot S/n - (\beta + \delta_I)I \\ dH/dt &= \beta I - \delta_H H, & dR/dt &= \delta_I I + \delta_H H \end{aligned}$$

## Comparative Methods

- **MBPS, MBPSH** (proposed methods)
- **BPS**: Univariate BPS (with count response) separately in each prefecture, with the same four models as MBPS and MBPSH.
- **FMPR**: Finite mixture of Poisson regression

$$y_{it}|(z_i = k) \sim \text{Po}(\exp(\mathbf{x}_{it}^\top \boldsymbol{\beta}_k)), \quad \mathbf{x}_{it} = (1, \tilde{l}_{it}, \tilde{l}_{it}^2, y_{i,t-1})^\top$$
$$\Pr(z_i = k) = \pi_k, \quad k = 1, \dots, K$$

## Performance Measures

- Cumulative absolute prediction errors (CAPE):

$$\text{CAPE}_t = \sum_{i=1}^n \sum_{t^*=T}^t |y_{i,t^*+k} - \hat{y}_{i,t^*+k}|$$

- Log predictive density ratios (LPDR):

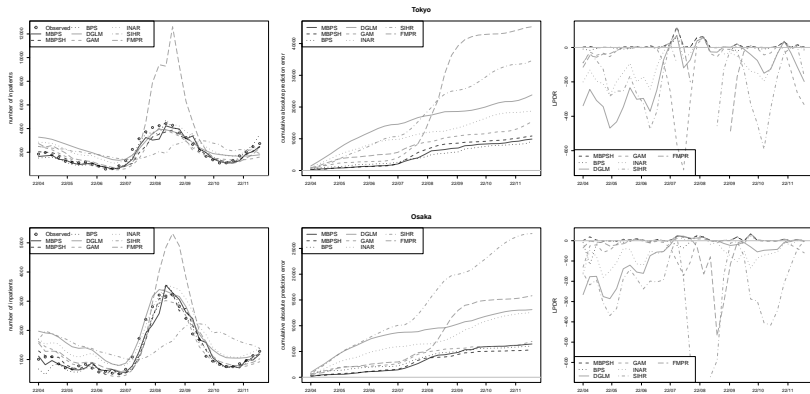
$$\text{LPDR}_t = \log \frac{p_j(y_{t+k}|y_{1:t})}{p_{\text{MBPS}}(y_{t+k}|y_{1:t})}$$

- Coverage:

$$\frac{1}{T^*} \sum_t I\{y_{t+k} \in 95\% \text{ prediction interval}\}$$

where  $T^*$  is the length of prediction periods.

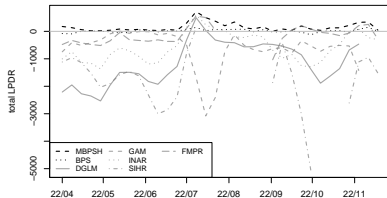
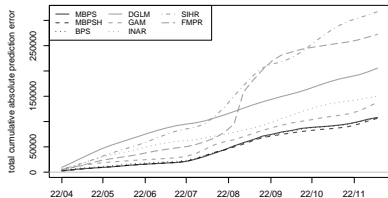
# One-step-ahead forecasting (Tokyo, Osaka)



Predictions (left), CAFE (middle), LPDR (right)



# One-step-ahead forecasting (Total)

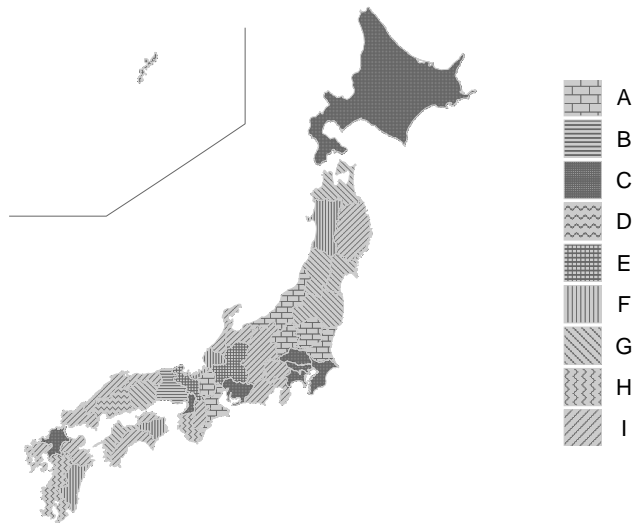


## Coverage

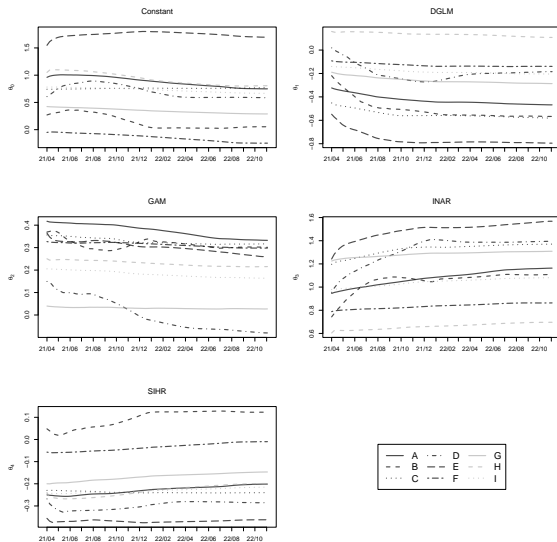
MBPS	MBPSH	BPS	DGLM	GAM	INAR	SIHR	FMPR
0.556	<b>0.928</b>	0.583	0.158	0.122	0.236	0.036	0.287

- The proposed methods provide better prediction accuracy.
- MBPSH provides much better coverage performance than MBPS.

## Clustering Result on 2022/11/23



# Synthesis weights by cluster



# Summary

- Mixture of Bayesian predictive synthesis (MBPS)
  - BPS for multivariate time series, sharing the common synthesis weights in the same cluster
  - Introduction of heterogeneity in the intercept improves the coverage.
  - MBPS is useful for large-dimensional multivariate time series prediction.
- For more information:

Kobayashi, G., Sugawara, S., Kawakubo, Y., Han, D. and Choi, T. (2024). Predicting COVID-19 hospitalisation using a mixture of Bayesian predictive syntheses. *The Annals of Applied Statistics* 18, 3383-3404.

## Supplement: Analysis of COVID-19 Isolation in Korea

- Daily numbers of isolated cases by first division obtained from `data.go.kr`
- Total data period: 2020/08/01 – 2021/11/30 (487 days)
- $K = n = 17$
- Multistep-ahead forecasts

Coverage								
$s$ -step	MBPS	MBPSH	BPS	DGLM	GAM	INAR	SIHR	FMPR
$s = 1$	0.918	0.964	0.932	0.605	0.169	0.848	0.028	0.351
$s = 3$	0.724	0.944	0.767	0.590	0.164	0.654	0.022	0.354
$s = 7$	0.637	0.953	0.697	0.575	0.169	0.456	0.028	0.360