Space-Time Areal Mixture Model: Relabeling Algorithm and Model Selection Issues

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Areal model (Stat in Med, 2006; EES, 2005): Local-likelihood cluster (LLC) model Compared with the BYM model Results: For detecting clusters of low and medium risk areas, LLC models signal better than BYM model Cluster detection diagnostics



Spatio-temporal Areal model ((EES, 2012; EES, 2010) : Space-time local-likelihood cluster (LLCST) model Space-time mixture of Poisson (MPST) model Space-time cluster detection diagnostics

Space-time stick-breaking process (SBPST) Compared with the SREST model



Motivation

With the growing popularity of using spatial mixture model in cluster analysis, using model selection criteria to find the most parsimonious model is an established technique.

- Label-switching is an inherent problem with the mixture models and a variety of relabeling algorithms have been proposed over the decade.
- We used a space-time mixture of Poisson regression model with homogeneous covariate effects
- The results are illustrated for real and simulated datasets.
- The objective is to aware the researcher that if the purpose of statistical modeling is to identify the clusters, applying the relabeling algorithm to the best fitted model may not generate the optimum labeling.



Space-Time Mixture of Poisson Model

$$o_{it} \sim \sum_{l=1}^{L} \omega_{itl} Poisson(\theta_l e_{it} \exp(\eta_{it}))$$
$$\eta_{it} = \beta_{i0} x_{it0} + \beta_{i1} x_{it1} + \dots + \beta_{ip} x_{itp}$$
$$0 < \omega_{itl} < 1, \text{ and } \sum_{l=1}^{L} \omega_{itl} = 1$$

The weights are determined by the unobserved/hidden allocation variable:

$$\begin{split} \boldsymbol{\omega}_{itl} &= p\left(Z_{it} = l\right) \\ \boldsymbol{\omega}_{itl} &= \exp(h_{itl} / \phi) / \sum_{l=1}^{L} \exp(h_{itl} / \phi) \\ h_{itl} &= \kappa_{il} + \gamma_{tl} \end{split}$$



Model Selection Criteria

Deviance information criteria (Spiegelhalter et al., 2002):

$$DIC_M = \overline{D(\Theta_M)} + p_M$$

Deviance information criteria (Celeux et al., 2006):

$$\mathrm{DIC}_{3M} = \overline{D(\Theta_M)} + \left[\overline{D(\Theta_M)} + 2\log \hat{p}(\mathbf{o} | \Theta_M)\right]$$

Mean square predictive error (Gelman and Ghosh, 1998):

$$MSPE_{M} = \sum_{g=1}^{G} \sum_{i=1}^{n} \sum_{t=1}^{T} \left(O_{it} - O_{it}^{(g,M)} \right)^{2} / nTG$$



Relabeling Algorithm

Posterior similarity matrix:

$$\pi_{ijt} = \Pr\left(Z_{it} = Z_{jt} \mid \mathbf{o}\right) \approx \frac{1}{G} \sum_{g=1}^{G} I\left(Z_{it}^{(g)} = Z_{jt}^{(g)}\right)$$
$$E\left[\left(Z_{it} = Z_{jt}\right) \mid \mathbf{o}\right] = \pi_{ijt} = E\left[I\left(Z_{it} = Z_{jt}\right) \mid \mathbf{o}\right]$$

Viewed as a similarity matrix of posterior expected clustering. Regarded as a similarity matrix for unknown true clustering. Binder's loss:

$$L(\mathbf{Z}'_{t}, \mathbf{Z}_{t}) = \sum_{i < j} l_{1} \cdot I(\mathbf{Z}'_{it} \neq \mathbf{Z}'_{jt}) I(\mathbf{Z}_{it} = \mathbf{Z}_{jt}) + l_{2} \cdot I(\mathbf{Z}'_{it} = \mathbf{Z}'_{jt}) I(\mathbf{Z}_{it} \neq \mathbf{Z}_{jt})$$
$$E[L(\mathbf{Z}'_{t}, \mathbf{Z}_{t}) | \mathbf{o}] = \sum_{i < j} |I(\mathbf{Z}'_{it} = \mathbf{Z}'_{jt}) - \pi_{ijt}|$$
$$\hat{\mathbf{Z}}_{t} = \arg\min_{g} E[L(\mathbf{Z}'_{t}, \mathbf{Z}_{t}) | \mathbf{o}]$$



Relabeling Algorithm

Lau and Green (2007) criteria:

$$E\left[L\left(\mathbf{Z}_{t}^{\prime},\mathbf{Z}_{t}\right)\mid\mathbf{o}\right]=\sum_{i< j}I\left(Z_{it}^{\prime}=Z_{jt}^{\prime}\right)\left(1-\pi_{ijt}\right)$$

Posterior expected adjusted Rand (PEAR):

$$AR[\mathbf{Z}'_{t}, E(\mathbf{Z}_{t} | \mathbf{0})] = \frac{\sum_{i < j} I(Z'_{it} = Z'_{jt}) \pi_{ijt} - \sum_{i < j} I(Z'_{it} = Z'_{jt}) \sum_{i < j} \pi_{ijt} / \binom{n}{2}}{\frac{1}{2} \left[\sum_{i < j} I(Z'_{it} = Z'_{jt}) + \sum_{i < j} \pi_{ijt} \right] - \sum_{i < j} I(Z'_{it} = Z'_{jt}) \sum_{i < j} \pi_{ijt} / \binom{n}{2}}$$

$$\hat{\mathbf{Z}}_{t} = \arg\max_{g} AR\left[\mathbf{Z}_{t}', E\left(\mathbf{Z}_{t} \mid \mathbf{0}\right)\right]$$



Variable Selection

Kuo and Mallick (1998):

$$\eta_{ij} = \beta_{i0} x_{ij0} + I_1 \beta_{i1} x_{ij1} + \dots + I_p \beta_{ip} x_{ijp}$$

Covariate being selected or not is independent of covariate effect a priori:

$$p(I_{q}\beta_{iq}) = p(I_{q})p(\beta_{iq})$$



Simulation-1: 2 clusters Simulation-2: 5 clusters



 $x_{itq} \sim \text{Uniform}(0,1); i = 1, ..., n, t = 1, ..., T, and q = 1, ..., 3$ Ohio geography: Expected lung cancer (year: 1968-88) Realization: 100



Simulation-1

Indicator Posterior Standard variable for mean error covariate effect 0.09778 0.22036 1 0.72674 0.20037 2 0.01051 0.98873 3

Simulation-2

Indicator	Posterior	Standard
variable for	mean	error
covariate		
effect		
1	0.67662	0.00937
2	0.83869	0.01011
3	0.95003	0.01030



Simulation-1

Covariate	DIC	DIC3	MSPE	Binder's	Lau &	Maximum
				loss	Green loss	PEAR
Intercept only	11618.51	11944.69	189.7469	1174.993	1173.514	0.37789
Intercept and 1	11619.55	11953.45	185.6524	1093.262	1091.638	0.39438
Intercept and 2	11986.08	12277.73	214.2977	337.1096	336.5138	0.57292
Intercept and 3	11541.8	11873.84	173.6958	221.3878	220.0612	0.545087
Intercept, 1 and 2	11573.86	11915.37	180.5074	291.3908	291.0096	0.4914627
Intercept, 1 and 3	11615.46	11958.0	174.4494	199.1183	198.8899	0.5548457
Intercept, 2 and 3	11580.64	11920.02	175.5993	137.805	137.7711	0.6375331
Full model	11564.58	11906.14	175.4762	175.4762	167.6591	0.4951442



Simulation-2

Covariate	DIC	DIC3	MSPE	Binder's	Lau &	Maximum
				loss	Green loss	PEAR
Intercept only	13089.64	13635.52	422.1604	1237.826	1084.311	0.2069
Intercept and 1	14840.08	15964.27	1221.702	1030.294	991.5413	0.3707
Intercept and 2	15512.65	16973.48	1487.344	873.979	851.8798	0.4744
Intercept and 3	16293.81	17920.75	1981.996	873.1445	850.3804	0.4700
Intercept, 1 and 2	15574.62	16992.66	1523.662	901.3975	877.3205	0.4447
Intercept, 1 and 3	15287.31	16695.92	1358.119	877.1459	853.2936	0.4658
Intercept, 2 and 3	15235.93	16609.79	1388.221	878.995	854.6614	0.4707
Full model	14131.80	15034.43	970.9007	1091.618	1037.593	0.3477



Real Data Example

South Carolina low birth weight data (Year: 1997-2007)

Component	DIC	DIC ₃	MSPE
2	3660.9	3783.7	270.8
3	3620.1	3734.1	252.8
4	3646.5	3771.2	251.9
5	3609.0	3720.7	241.4
6	3633.9	3754.6	253.8
7	3650.0	3780.6	250.9

Indicator variable for	Posterior	Standard
covariate effect	mean	error
PD	0.2347	0.4238
PAA	0.1835	0.3870
МНІ	0.4565	0.4981
PP	0.1862	0.3892
UR	0.1797	0.3839
Time	0.5883	0.4921



Real Data Example





Real Data Example

South Carolina low birth weight data (Year: 1997-2007)

Covariate	DIC	DIC ₃	MSPE	Binder's	Lau &	Maximum
				loss	Green loss	PEAR
Intercept only	3568.6	3662.4	223.7	170.78	168.59	0.607
Intercept, MHI	3597.0	3700.4	232.2	105.62	102.43	0.780
Intercept, time	3572.2	3663.0	227.7	497.71	497.10	0.037
Intercept, MHI, time	3582.2	3677.7	232.9	177.38	177.30	0.645
Full model	3609.0	3720.7	241.4	306.64	265.10	0.270



Conclusions

Previously, Best et al. (2005) showed that the spatial model with convolution prior (e.g. Besag et al., 1991) overestimate the risk surface for the high risk areas and the best model selected by the DIC is not always able to select the right clusters.

- We used a space-time mixture of Poisson regression model with homogeneous covariate effects.
- We designed two simulation studies with smaller and larger numbers of clusters, and with common covariate effects. The covariates are generated with stronger to weak levels of spatial correlation.
- In our simulated and real datasets, we observed that model selection criteria do not indicate to the right cluster model.



Conclusions

Limitations:

Simultaneous estimation of the number of clusters and variable selection for spatial data

Improvement the DIC performance:

Variational Bayes approach (McGrory and Titterington, 2007) Viewing DIC as an approximate penalized loss function (Plummer, 2008)

Model based relabeling algorithm where the label for each observation is chosen by maximizing the classification probability (Yao, 2012)



Lancashire larynx cancer (58 cases) and lung cancer (978 controls)

Recorded in Chorley and South Ribble Health Authority during 1974-83







Point process modeling (CSDA, 2010): Approximate likelihood Berman-Turner (BT) model Conditional logistic (CL) model Binomial mesh (BM) model Poisson mesh (PM) model



Point process modeling:

	BT me	ethod		PM model		BM model	
Parameter	$\phi = 0.0001$	$\phi = 0.001$	CL model	20X20	30X30	20X20	30X30
$oldsymbol{eta}_{0}$	-0.641	-0.648	-0.739	-0.868	-0.843	-0.915	-0.773
	(-4.112, 1.474)	(-4.018, 1.489)	(-4.249, 1.582)	(-5.027, 1.672)	(-4.874, 1.755)	(-5.255, 1.872)	(-4.965, 2.087)
eta_1	-0.00078	-0.00078	-0.00077	-0.00050	-0.00048	-0.00048	-0.00059
	(-0.00383,	(-0.00384,	(-0.00432,	(-0.00388,	(-0.00388,	(-0.00398,	(-0.00424,
	0.002161)	0.00221)	0.00252)	0.00307)	0.00311)	0.00317)	0.00313)
$\log(ho)$	-3.155 (-3.752, -2.758)	-3.161 (-3.684, - 2.767)	-3.374 (-4.076, -2.885)	-3.200 (-3.737, - 2.808)	-3.215 (-3.755, - 2.804)	-3.145 (-3.705, -2.717)	-3.173 (-3.726, -2.756)
$\sigma_{_{u}}$	0.390	0.411	0.771	0.165	0.141	0.216	0.242
	(0.127, 0.891)	(0.095, 0.932)	(0.241, 1.475)	(0.004, 0.473)	(0.001, 0.522)	(0.009, 0.633)	(0.002, 0.728)
$\sigma_{_{V}}$	0.050	0.0331	0.224	0.529	0.637	0.537	0.647
	(0.025, 0.087)	(0.005, 0.071)	(0.078, 0.510)	(0.111, 1.305)	(0.095, 1.440)	(0.014, 1.342)	(0.090, 1.632)



Current Works (Childhood Cancer)

 $\lambda \hat{K}$ (Avg Number of Nearby Cancer Incidences) $\overset{0}{5}$ Observed Random 500 -0 10 30 n 20 Distance (r) in mile

Childhood Cancer (age: 0-24, year: 1996-2009)



Current Works (Childhood Cancer)



envelope(fit, Linhom, 99)



Thanks!

