

36-720: Latent Class Models

Brian Junker

October 17, 2007

- Latent Class Models and Simpson's Paradox
- Latent Class Model for a Table of Counts
- An Intuitive E-M Algorithm for Fitting Latent Class Models
- Deriving the E-M Algorithm
- Example
- Issues and Warnings

For more reading, see:

Bartholomew, D. J. & Knott, M. M. (1999). *Latent Variable Models and Factor Analysis*. Oxford Univ Press.

Hagenaars, J. A. & McCutcheon, A. L. (2003). *Applied Latent Class Analysis*. Cambridge Univ Press.

McLachlan, G. & Peel, D. (2000). *Finite Mixture Models*. NY: Wiley.

The Many Siblings of Latent Class Analysis (LCA)

- To the extent that we are “assigning” some observations to the group “ $Z = 0$ ” and some to the group “ $Z = 1$ ”, LCA is a kind of *model-based cluster analysis* and/or an *unsupervised classifier* for multivariate discrete data.
- We will see below that LCA basically models the distribution of $\{n_{00}, n_{01}, n_{10}, n_{11}\}$ as the mixture of two log-linear models (one for $Z = 0$ and one for $Z = 1$). Therefore LCA can also be viewed as a kind of *finite mixture modeling*.
 - Finite mixture models for continuous response data are perhaps more familiar: *mixture-of-normals density estimators*, *kernel density estimators*, etc.
 - Other finite mixtures for discrete data, such as *zero-inflated Poisson models*, are also closely related to LCA.
- *Bayes Network models* for discrete data with discrete hidden nodes may also be viewed as a kind of constrained LCA (each latent class is determined by a unique set of values for the hidden nodes).
 - These models are basic choices in AI/expert systems, computer-based tutoring, cognitive diagnosis, etc.

Latent Class Model for a Table of Counts

Latent class models are often used with item response data (like the Rasch model) so we will consider that case as well. Once again let

$$y_{ij} = \begin{cases} 1, & \text{if student } i \text{ got question } j \text{ correct} \\ 0, & \text{else} \end{cases}$$

say, for $i = 1, \dots, N$ students and $j = 1, \dots, J$ questions.

We begin by considering the 2^J table

$$\{n_{\underline{y}} : \text{as } \underline{y} \text{ ranges over all } 2^J \text{ possible patterns } (y_1, \dots, y_J)\}$$

directly.

The latent class model says that there are W layers, or latent classes, that we can split this table into:

$$\begin{aligned} n_{\underline{y}}^{(1)} &= t_{\underline{y}}^{(1)} n_{\underline{y}} \\ n_{\underline{y}}^{(2)} &= t_{\underline{y}}^{(2)} n_{\underline{y}} \\ &\vdots \\ n_{\underline{y}}^{(W)} &= t_{\underline{y}}^{(W)} n_{\underline{y}} \end{aligned}$$

where the $t_{\underline{y}}^{(w)}$ is simply the proportion of observations in cell \underline{y} that get assigned to layer, or latent class, w (so $\sum_w t_{\underline{y}}^{(w)} = 1$, for each \underline{y} ; and $t_{\underline{y}}^{(w)}$ may be different for different \underline{y}).

Within each latent class we assume the model of independence holds:

$$\begin{aligned} n_{\underline{y}}^{(w)} &\sim \text{Multinom}(p_{\underline{y}}^{(w)}, N_w) \\ \log m_{\underline{y}}^{(w)} &= \alpha^{(w)} + \beta_1^{(w)} y_1 + \cdots + \beta_J^{(w)} y_J \end{aligned}$$

where $N_w = \sum_{\underline{y}} n_{\underline{y}}^{(w)}$, and $m_{\underline{y}}^{(w)} = N_w p_{\underline{y}}^{(w)}$.

An Intuitive E-M Algorithm for Fitting Latent Class Models

One way to fit a latent class model is with a kind of E-M algorithm:

Initialization: Make an initial guess $\tilde{n}_{\underline{y}}^{(w)}$ for the counts in the w^{th} layer.

M step: Fit the log-linear model of independence to each table $\tilde{n}_{\underline{y}}^{(w)}$, using `glm()`, obtaining fitted values $\tilde{m}_{\underline{y}}^{(w)}$;

E step: Recalculate

$$\tilde{t}_{\underline{y}}^{(w)} = \frac{\tilde{m}_{\underline{y}}^{(w)}}{\tilde{m}_{\underline{y}}^{(1)} + \tilde{m}_{\underline{y}}^{(2)} + \cdots + \tilde{m}_{\underline{y}}^{(W)}} \quad , \quad \tilde{n}_{\underline{y}}^{(w)} = \tilde{t}_{\underline{y}}^{(w)} n_{\underline{y}}$$

Convergence check: If this iteration does not produce much change from the previous iteration, then stop. Otherwise start over, with the “M” step.

Note that all of the data $n_{\underline{y}}^{(w)}$ is missing, in every layer of the table along W .

- Clearly the “E” step above is imputing this missing data by first calculating the fitted probabilities $\tilde{t}_{\underline{y}}^{(w)}$ that \underline{y} will be classified into class w , and then computing the expected number $n_{\underline{y}}^{(w)} = \tilde{t}_{\underline{y}}^{(w)} n_{\underline{y}}$.
- The “M” step is just fitting the complete-data model (log-linear model of independence) to the imputed tables $n_{\underline{y}}^{(w)}$.
- Finally, any reasonable measure of fit can be used in the “Convergence check”. One interesting possibility is to calculate expected cell counts $m_{\underline{y}} = \sum_w m_{\underline{y}}^{(w)}$ and then keep track of $G^2 = 2 \sum_{\underline{y}} n_{\underline{y}} \log(n_{\underline{y}}/m_{\underline{y}})$ for the original table. When G^2 stops changing, we can quit the E-M algorithm.

Deriving the E-M Algorithm

Imitating our work for the Rasch model, we can see that the likelihood for examinee i in latent class w is

$$P[y_{i1}, \dots, y_{iJ} \mid w, p_{i1}, \dots, p_{iJ}] = \prod_{j=1}^J p_{wj}^{y_{ij}} (1 - p_{wj})^{y_{ij}}$$

If we treat w as a random effect (like we treated θ in the Rasch model) we get the mixture model

$$P[y_{i1}, \dots, y_{iJ} \mid p_{i1}, \dots, p_{iJ}, \lambda_1, \dots, \lambda_W] = \sum_w \lambda_w \prod_{j=1}^J p_{wj}^{y_{ij}} (1 - p_{wj})^{y_{ij}}$$

where λ_w is the (prior) probability that an examinee belongs to class w . This equation is analogous to equation

$$P[y_{i1}, \dots, y_{iJ} \mid \beta_1, \dots, \beta_J; \sigma] = \int_{-\infty}^{\infty} \prod_{j=1}^J \frac{\exp\{y_{ij}(\theta_i - \beta_j)\}}{1 + \exp\{\theta_i - \beta_j\}} f(\theta_i \mid \sigma) d\theta_i$$

for the Rasch model.

The joint model for all examinees

$$P[\mathcal{Y} \mid p_{ij}, \text{ all } i, j; \lambda_1, \dots, \lambda_W] = \prod_{i=1}^N \sum_w \lambda_w \prod_{j=1}^J p_{wj}^{y_{ij}} (1 - p_{wj})^{1-y_{ij}}$$

is kind of unwieldy but it simplifies if we introduce a *data augmentation variable* $z_{iw} = 1$ if examinee i is in latent class w and $z_{iw} = 0$ otherwise. With this the complete data likelihood becomes

$$P[\mathcal{Y}, \mathcal{Z} \mid \text{all } p_{ij}, \text{ all } \lambda_w] = \prod_i \prod_w \left\{ \lambda_w \prod_j p_{wj}^{y_{ij}} (1 - p_{wj})^{1-y_{ij}} \right\}^{z_{iw}}$$

Now if we apply the “standard” E-M algorithm with

$$Q(\varphi, \varphi^{(k)}) = E[\log P(\mathcal{Y}, \mathcal{Z} \mid \varphi) \mid \varphi^{(k)}, \mathcal{Y}]$$

[where now $\varphi = (\text{all } p_{ij}, \text{ all } \lambda_w)$] we will eventually obtain the steps above, with $\tilde{t}_{\underline{y}_i}^{(w)} = P[z_{iw} = 1 \mid \underline{y} = \underline{y}_i]$.

Example

We previously analyzed the Stouffer-Toby data using graphical log-linear models and found the decomposable model

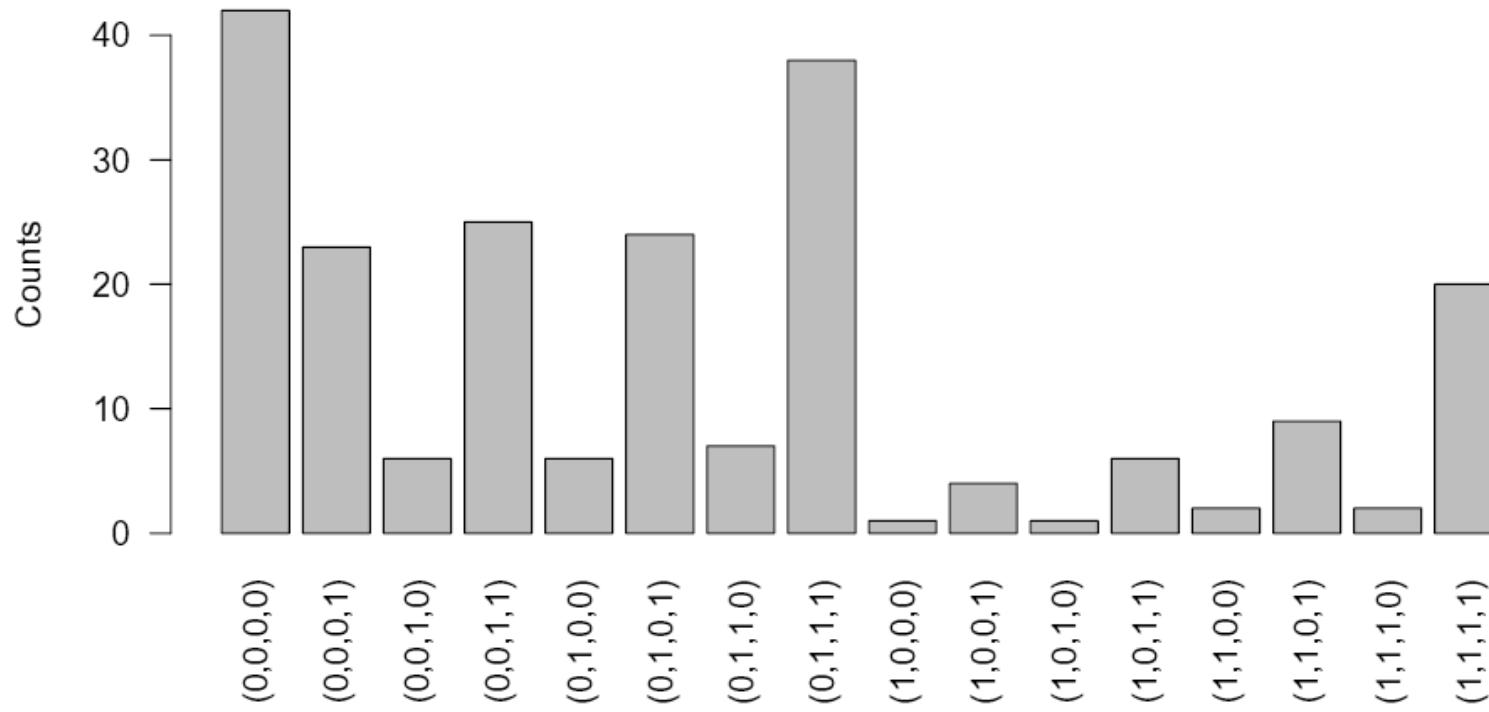
$$\begin{array}{ccccc} 1 & -- & 2 & -- & 3 \\ & & \backslash & & | \\ & & \backslash & & | \\ & & 4 & & \end{array}$$

Recall that the data consist of role-conflict responses of 216 respondents to four situations, with response “0” being “universalistic” and “1” being “particularistic”. Goodman (2002, Chapter 1 of *Applied Latent Class Analysis*, Hagenaars & McCutcheon, eds, NY: Cambridge Univ Press) considers a latent class analysis of the same data.

We will reproduce, with some modifications, Goodman’s analysis.

A pdf of Goodman’s chapter is included with these lecture notes.

We begin with a barplot of the actual counts in each cell:



- There is clearly dependence in the data: for example $P[y_2, y_3, y_4 | y_1 = 0] \neq P[y_2, y_3, y_4 | y_1 = 1]$.
- This also suggests perhaps some latent class structure, driven somehow by response to item 1.

An E-M algorithm to fit the LC model can be coded easily in R:

```
lcm <- function(data,tstar=runif(dim(data)[1]),
                 fla1=n~, fla2=n~, tol=1, reps=100) {
  cells <- dim(data)[1]
  if (length(tstar)!=cells) {
    tstar <- rep(abs(tstar[1]),cells)
  }
  tstar <- tstar/sum(tstar)
  d1 <- data
  d2 <- data
  G1 <- 1e6
  G2 <- 1e6
  K <- 0
  err <- 1e6
  while ((err > tol)&&(K<reps)) {
    K <- K+1
    G1.old <- G1
    G2.old <- G2
    d1$n <- tstar*data$n
    d2$n <- (1-tstar)*data$n
    fit1 <- glm(fla1,data=d1,family=poisson)
    fit2 <- glm(fla2,data=d2,family=poisson)
    m1 <- fitted(fit1)
    m2 <- fitted(fit2)
    tstar <- m1/(m1+m2)
    G1 <- summary(fit1)$deviance
    G2 <- summary(fit2)$deviance
    err <- abs((G1+G2)-(G1.old+G2.old))
  }
  return(list(K=K,err=err,tstar=tstar,
             fit1=fit1,fit2=fit2,d1=d1,d2=d2))
}
```

Fitting in R we get

```
> result <- lcm(stouffer, tol=0.00001)
```

There were 50 or more warnings

(use warnings() to see the first 50)

```
> warnings()
```

Warning messages:

1: non-integer x = 0.450371

2: non-integer x = 1.485916

[...]

50: non-integer x = 21.600592

- The non-integer cell counts are a consequence of the E-M algorithm; you can verify that maximizing the Poisson or multinomial likelihood still makes sense mathematically.
- We can look at barplots in each of the two latent classes to see how the LC model split up the data [see next slide]...

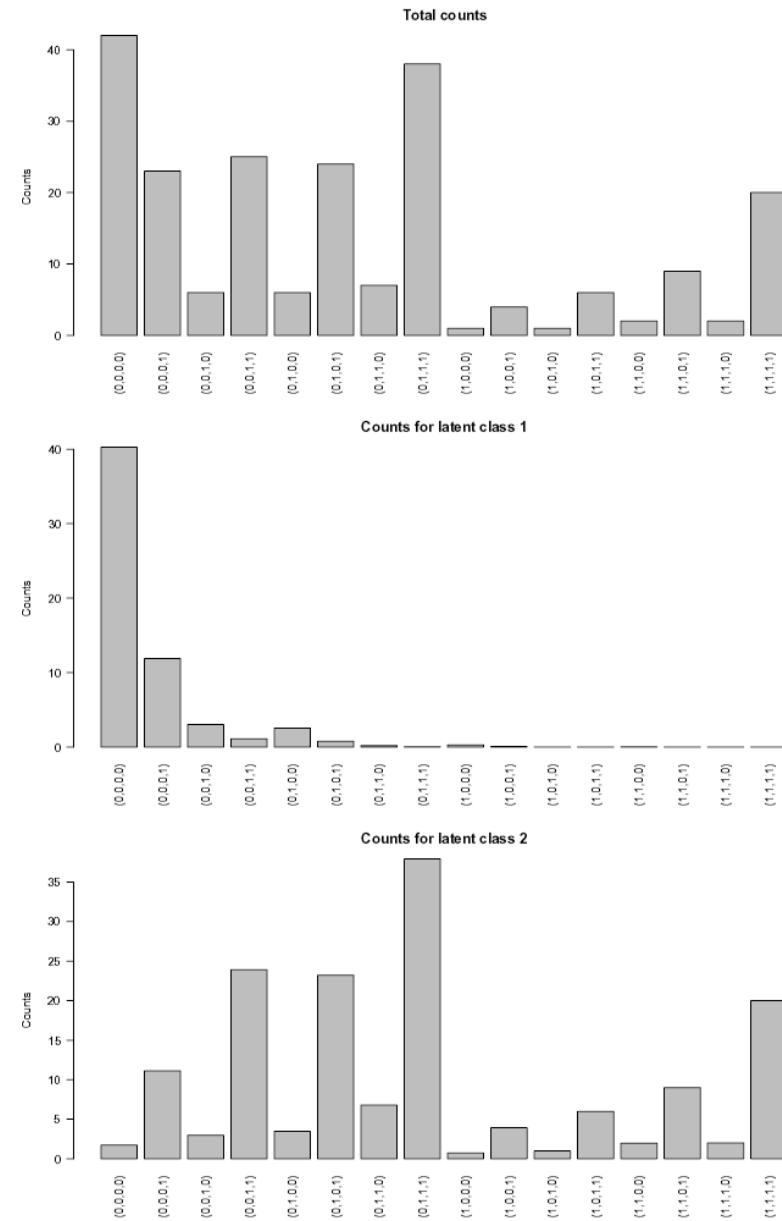
- LC #1: individuals excessively likely to respond 0 to most questions...
- LC #2: $y_1 = 0$ frequencies are roughly proportional to $y_1 = 1$ frequencies, allowing cond'l indep.
- Note that the deviances in each class add up to the overall deviance:

```

> m <- fitted(result$fit1) + fitted(result$fit2)
> 2*sum(n*log(n/m))
[1] 2.719957
> result$fit1$deviance + result$fit2$deviance
[1] 2.719960

```

You should be able to prove that this must happen!



Issues and Warnings

Many other issues remain for latent class analysis...

- How many latent classes? We chose two...
 - The best approaches to choosing the number of latent classes tend to involve Bayes Factor (or BIC) comparisons of models with different numbers of latent classes.
- Reliable MLE optimization...
 - LC models tend to have one or more secondary maxima, so that it is difficult to see that you are at the global maximum without a lot of trial and error with starting values.
- Label-switching issues
 - “The” global maximum will actually occur at $W!$ different locations on the likelihood surface, due to equivalent re-labellings of the latent classes.
 - This is an annoyance for careful inference using MLE methods...

- For MCMC methods it can be a disaster, since the Markov Chain will visit different ones of these modes and average over them, producing mashed potatoes for parameter estimates . . .
- Interpretation . . .
 - LC models for prediction have all of the above problems.
 - In Social Science (and other) applications, we also often want to interpret the latent classes in terms of the underlying science.
 - This can be very useful, but it can also be dangerous—reifying apparent structure in the latent classes that isn't scientifically justifiable.
 - Similar problems occur with mixture-of-normal models, factor analysis models, etc.
- Etc., etc., . . .

Details and Difficulties

Marin, J.M., Mengersen, K. & Robert, C.P. (2004). Bayesian modelling and inference on mixtures of distributions. *Handbook of Statistics 25*, D. Dey and C.R. Rao (eds). Elsevier-Sciences (to appear).

only deals with continuous-data mixtures, but it addresses most of the computational and inferential issues found in current discussions of mixture models by statisticians; the same issues apply to LCA. *A pdf is included with these lecture notes.*

Fienberg, S. E., Hersh, P., Rinaldo, A., & Zhou, Y. (2007). Maximum Likelihood Estimation in Latent Class Models For Contingency Table Data. Document arXiv:0709.3535v1, available at <http://arxiv.org/abs/0709.3535>

provides an account of LC models from the point of view of algebraic statistics and gives new examples suggesting that ML estimation of LC models can be quite delicate. *A pdf is also included with these lecture notes.*