

Aster Models for Life History Analysis with Lessons for Teaching Statistics

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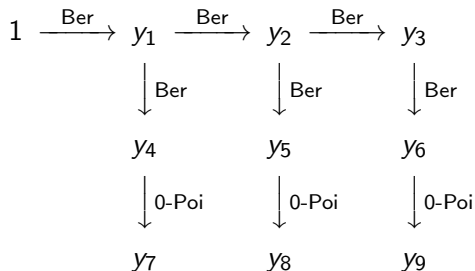
R contributed package `aster` on CRAN.

```
install.packages("aster")  
library(aster)
```

Function `aster` fits models. Generic functions `summary`, `predict`, and `anova` work like those for linear and generalized linear models.

<http://www.stat.umn.edu/geyer/aster/> has links to papers and tech reports. All tech reports done with Sweave so everything is exactly reproducible.

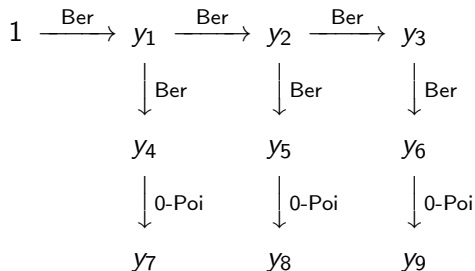
An Aster Graph



y_i are components of response vector for one individual (all individuals have isomorphic graphs). 1 is the constant 1.

Arrows indicate conditional distributions of variable at head of arrow (successor) given variable at tail of arrow (predecessor).
Ber = Bernoulli, 0-Poi = zero-truncated Poisson.

An Aster Graph (cont.)



Graph for *Echinacea angustifolia* example in Geyer, Wagenius and Shaw (*Biometrika*, 2007).

y_1, y_2, y_3 indicate survival in each of three years (2002–2004).

y_4, y_5, y_6 indicate flowering status (1 = some flowers, 0 = no flowers) in corresponding years.

y_7, y_8, y_9 are flower counts in corresponding years.

Abstract Aster Graph

Nodes (variables) have at most one predecessor, hence graph is specified by function p that maps from set J of non-initial nodes to set N of all nodes. $y_{p(j)}$ is predecessor of y_j .

y_j at initial nodes treated as constants. Then joint distribution factors as product of conditionals

$$f_{\theta}(y) = \prod_{j \in J} f_{\theta}(y_j \mid y_{p(j)})$$

because graph is not allowed to have loops. Log likelihood is

$$l(\theta) = \sum_{j \in J} \log f_{\theta}(y_j \mid y_{p(j)})$$

Predecessor is Sample Size

In subgraph

$$y_{p(j)} \longrightarrow y_j$$

y_j is sum of $y_{p(j)}$ independent and identically distributed random variables.

Define ξ_j to be the mean of one of those variables, so

$$E(y_j \mid y_{p(j)}) = y_{p(j)} \xi_j$$

ξ_j are components of *conditional mean value parameter* vector ξ .

Predecessor is Sample Size (cont.)

Define

$$\mu_j = E(y_j)$$

components of *unconditional mean value parameter* vector μ .

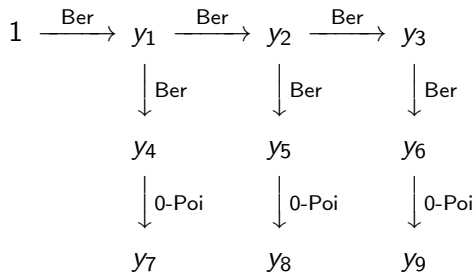
By iterated expectation theorem

$$\begin{aligned} E(y_j) &= E\{E(y_j \mid y_{p(j)})\} \\ &= E(y_{p(j)}\xi_j) \\ &= \xi_j E(y_{p(j)}) \end{aligned}$$

that is

$$\mu_j = \xi_j \mu_{p(j)}$$

Predecessor is Sample Size (cont.)



$$\mu_1 = \xi_1$$

$$\mu_2 = \xi_2 \xi_1$$

$$\mu_3 = \xi_3 \xi_2 \xi_1$$

$$\mu_4 = \xi_4 \xi_1$$

$$\mu_5 = \xi_5 \xi_2 \xi_1$$

$$\mu_6 = \xi_6 \xi_3 \xi_2 \xi_1$$

$$\mu_7 = \xi_7 \xi_4 \xi_1$$

$$\mu_8 = \xi_8 \xi_5 \xi_2 \xi_1$$

$$\mu_9 = \xi_9 \xi_6 \xi_3 \xi_2 \xi_1$$

Exponential Families of Distributions

An *exponential family* of distributions is a statistical model with log likelihood

$$\langle z, \theta \rangle - c(\theta)$$

when terms not containing the parameter have been dropped, and

$$\langle z, \theta \rangle = z^T \theta = \theta^T z$$

Statistic vector z and parameter vector θ that give log likelihood of this form are called *canonical*. c is called *cumulant function*.

Log likelihood for z_1, \dots, z_n independent and identically distributed is

$$\langle z_1 + \dots + z_n, \theta \rangle - nc(\theta)$$

Exponential Families and Predecessor is Sample Size

Each conditional distribution of y_j given $y_{p(j)}$ is one-parameter exponential family having y_j as the canonical statistic. In

$$l(\theta) = \sum_{j \in J} \log f_{\theta}(y_j \mid y_{p(j)})$$

j -th term of is

$$y_j \theta_j - y_{p(j)} c_j(\theta_j)$$

(compare with)

$$\langle z_1 + \cdots + z_n, \theta \rangle - n c(\theta)$$

Have subscripts on c_j and θ_j because each arrow can have different exponential family and different parameter.

Aster Model Log Likelihood

$$\begin{aligned}l(\theta) &= \sum_{j \in J} [y_j \theta_j - y_{p(j)} c_j(\theta_j)] \\ &= \sum_{j \in J} y_j \left[\theta_j - \sum_{\substack{k \in J \\ j=p(k)}} c_k(\theta_k) \right] - \sum_{\substack{k \in J \\ p(k) \notin J}} y_{p(k)} c_k(\theta_k)\end{aligned}$$

This is recognizable as log likelihood for joint exponential family.
Blue term is j -th component of joint canonical parameter vector.
Red term is cumulant function of joint family.

Aster Model Log Likelihood (cont.)

$$l(\varphi) = \langle y, \varphi \rangle - c(\varphi)$$

where

$$\varphi_j = \theta_j - \sum_{\substack{k \in J \\ j = p(k)}} c_k(\theta_k), \quad j \in J$$

and

$$c(\varphi) = \sum_{\substack{k \in J \\ p(k) \notin J}} y_{p(k)} c_k(\theta_k)$$

Aster Transform

θ is the *conditional canonical parameter vector*.

φ is the *unconditional canonical parameter vector*.

Map between them is invertible.

$$\theta_j = \varphi_j + \sum_{\substack{k \in J \\ j = p(k)}} c_k(\theta_k)$$

where θ_k on right-hand side have “already” been determined as function of φ . Use in any order that does successors before predecessors (always is one because graph has no loops).

Exponential Family Canonical and Mean Value Parameters

By properties of exponential families

$$\xi_j = c_j'(\theta_j)$$
$$\mu = \nabla c(\varphi)$$

where prime denotes univariate derivative and ∇ denotes multivariate derivative (vector of partial derivatives).

By properties of exponential families these changes of parameters are also invertible (although no closed-form expression in general, inversion equivalent to doing maximum likelihood).

A Plethora of Parameters

Four different parameterizations μ , ξ , θ , and φ .

All are important. All play a role in some scientific arguments.
Users have to understand all four.

But wait, there's more!

Canonical Linear Submodels

In an exponential family, with canonical parameter φ , the change of parameter

$$\varphi = M\beta$$

where M is a known matrix (model matrix or design matrix) gives a new exponential family because

$$\langle y, M\beta \rangle = y^T (M\beta) = y^T M\beta = (M^T y)^T \beta = \langle M^T y, \beta \rangle$$

and

$$l(\beta) = \langle M^T y, \beta \rangle - c(M\beta)$$

Submodel canonical parameter vector is β .

Submodel canonical statistic vector is $M^T y$.

Submodel mean value parameter vector is $\tau = E(M^T y) = M^T \mu$.

A Plethora of Parameters (cont.)

Six different parameterizations

μ	saturated model	unconditional	mean value
ξ	saturated model	conditional	mean value
φ	saturated model	unconditional	canonical
θ	saturated model	conditional	canonical
β	submodel	unconditional	canonical
τ	submodel	unconditional	mean value

Fisher Information

Fisher information for submodel canonical parameter vector β is

$$I(\beta) = -\nabla^2 \log l(\beta) = M^T \nabla^2 c(M\beta) M$$

Computer can convert to any other parameterization. And compute derivatives for applying the delta method to transfer standard errors.

Interpretation

In any exponential family, map between canonical and mean value parameter vectors is strictly multivariate monotone. Hence

$$\langle \mu_1 - \mu_2, \varphi_1 - \varphi_2 \rangle > 0$$

$$\langle \tau_1 - \tau_2, \beta_1 - \beta_2 \rangle > 0$$

$$\langle \xi_1 - \xi_2, \theta_1 - \theta_2 \rangle > 0$$

where $\mu_1, \varphi_1, \tau_1, \beta_1, \xi_1, \theta_1$ are different parameter vectors corresponding to the same aster model, ditto for $\mu_2, \varphi_2, \tau_2, \beta_2, \xi_2, \theta_2$, and the two models are distinct.

Interpretation (cont.)

Multivariate monotonicity dumbed down.

If φ_i increases, other components of φ being held fixed, then μ_i increases (other components of μ also change).

If β_i increases, other components of β being held fixed, then τ_i increases (other components of τ also change).

If θ_i increases, then ξ_i increases.

Interpretation (cont.)

Maximum likelihood estimators (MLE) in exponential families have the “observed equals expected” property. If $\hat{\mu}$ and $\hat{\tau}$ are MLE, then

$$\hat{\tau} = M^T \hat{\mu} = M^T y$$

Submodel always fits perfectly those aspects of the data that are submodel canonical statistics (components of $M^T y$).

Interpretation (cont.)

Exponential families have the maximum entropy property (Jaynes).

Each distribution in the family is as random as possible (maximizes entropy) subject to having a particular value of the mean value parameter vector $\tau = M^T \mu$.

If the components of the submodel canonical statistic vector $M^T y$ are scientifically interpretable, then the submodel is scientifically interpretable.

In teaching linear regression we usually introduce models with

$$y = M\beta + \text{error}$$

because students don't understand parametric statistic models.

Must be unlearned to understand generalized linear models.

Lessons (cont.)

In teaching linear regression we usually introduce confidence intervals for the mean value parameter

$$\mu = M\beta$$

by not calling μ a “parameter” but “predicted values.” Similarly for values of the regression function $x^T\beta$ for hypothetical data x .

This allows students to think of β as “the” vector of parameters and never think about other parameters.

But it gives students wrong intuitions about confidence intervals. Confidence intervals are always for *parameters*. That there are confidence intervals for things you don't want to call parameters (even though they are) must also eventually be unlearned.

Lessons (cont.)

In generalized linear models have three essential parameters.

Submodel parameter vector β .

Saturated model canonical parameter vector $\varphi = M\beta$.

Mean value parameter vector μ .

Calling only β “the” parameter vector and referring to φ as the “linear predictor” and μ as the “predicted values” is even more confusing in this context.

Confidence intervals for components of φ and μ are often scientifically important.

Lessons (cont.)

The map

$$\varphi = M\beta$$

between submodel and saturated model canonical parameter vectors is much woofed about.

(In linear regression, this is $\mu = M\beta$ because $\varphi = \mu$.)

In intro courses, where students don't know about matrices, this is presented as

$$\mu_i = x_{i1}\beta_1 + \cdots + x_{ip}\beta_p$$

or something of the sort.

Students are told this is how to interpret linear, generalized linear, and all regression-like models (including aster models).

The dual map

$$\tau = M^T \mu$$

between saturated model and submodel mean value parameter vectors just as important, if not more important, to scientific interpretation, but ignored in teaching.