

# Aster Models for Life History Analysis with Lessons for Teaching Statistics

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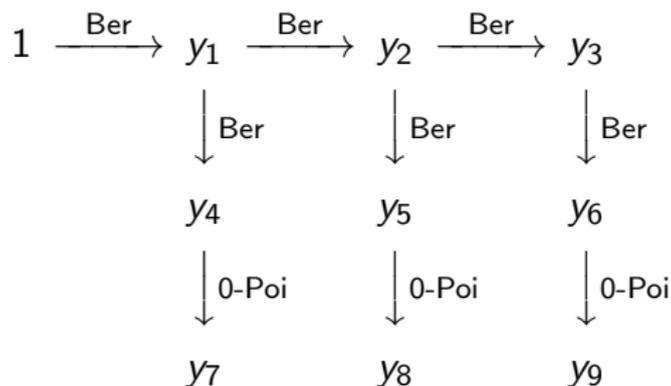
R contributed package `aster` on CRAN.

```
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.

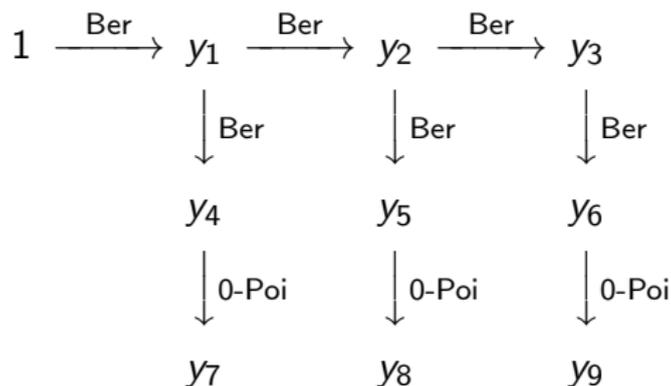
# An Aster Graph



$y_i$  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  $\xi_j$  to be the mean of one of those variables, so

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

$\xi_j$  are components of *conditional mean value parameter* vector  $\xi$ .

## Predecessor is Sample Size (cont.)

Define

$$\mu_j = E(y_j)$$

components of *unconditional mean value parameter* vector  $\mu$ .

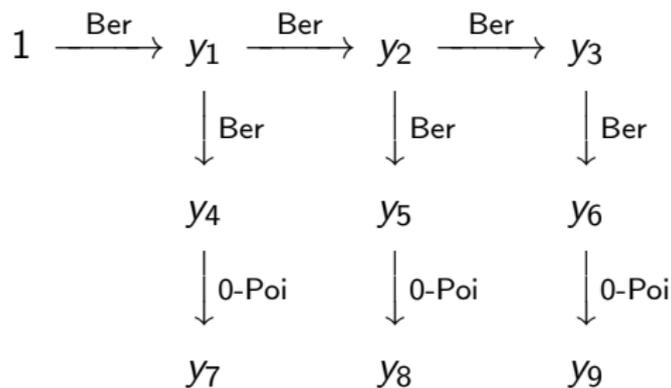
By iterated expectation theorem

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

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  $\theta$  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  $\theta_j$  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

$\theta$  is the *conditional canonical parameter vector*.

$\varphi$  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  $\theta_k$  on right-hand side have “already” been determined as function of  $\varphi$ . 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  $\nabla$  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  $\mu$ ,  $\xi$ ,  $\theta$ , and  $\varphi$ .

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  $\varphi$ , 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  $\beta$ .

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

|           |                 |               |            |
|-----------|-----------------|---------------|------------|
| $\mu$     | saturated model | unconditional | mean value |
| $\xi$     | saturated model | conditional   | mean value |
| $\varphi$ | saturated model | unconditional | canonical  |
| $\theta$  | saturated model | conditional   | canonical  |
| $\beta$   | submodel        | unconditional | canonical  |
| $\tau$    | submodel        | unconditional | mean value |

## Fisher Information

Fisher information for submodel canonical parameter vector  $\beta$  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  $\varphi_i$  increases, other components of  $\varphi$  being held fixed, then  $\mu_i$  increases (other components of  $\mu$  also change).

If  $\beta_i$  increases, other components of  $\beta$  being held fixed, then  $\tau_i$  increases (other components of  $\tau$  also change).

If  $\theta_i$  increases, then  $\xi_i$  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  $\mu$  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  $\beta$  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  $\beta$ .

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

Mean value parameter vector  $\mu$ .

Calling only  $\beta$  “the” parameter vector and referring to  $\varphi$  as the “linear predictor” and  $\mu$  as the “predicted values” is even more confusing in this context.

Confidence intervals for components of  $\varphi$  and  $\mu$  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.