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Motivation

Start writing what might be a good introduction for any paper or chapter.

Include citations!

Definitions

- ASV
- OTU
- Zero-Inflation
- Phylogenetic tree
- Compositional
- Covariates/variables/ coefficients

Notation

 \bullet Number of samples: n

• Sample index: $i:i=1,\ldots,n$

 \bullet Number of ASVs: k

• ASV index: $j:j=1,\ldots,k$

 \bullet Number of covariates: q. Note that q does not include the intercept.

• Covariate index: $k: k = 1, \dots, q$

• Responses: Transformed counts into relative abundances.

Note that an alternative transformation to relative abundance might be good.

$$y_{ij}$$

$$\mathbf{y}_{np imes 1} = \begin{pmatrix} y_{i=1,j=1} \\ \vdots \\ y_{i=1,j=p} \\ \vdots \\ y_{i=n,j=1} \\ \vdots \\ y_{i=n,j=p} \end{pmatrix}$$

• Design matrix \mathbf{x} ('little \mathbf{x} ').

$$\mathbf{x}_{n \times q} = \begin{pmatrix} x_{i=1,k=1} & \cdots & x_{i=1,k=q} \\ x_{i=2,k=1} & \cdots & x_{i=2,k=q} \\ \vdots & \ddots & \vdots \\ x_{i=n,k=1} & \cdots & x_{i=n,k=q} \end{pmatrix}$$

We don't have coefficient values measured on each ASV (i.e. no j index), they are only measured on each sample. Thus $x_{ij} = x_i$ for all j = 1, ..., p. Then we convert to design matrix big X:

• Design matrix X:

• Parameter vector β with entries β_{ik}

$$\boldsymbol{\beta}_{pq\times 1} = \begin{pmatrix} \beta_{j=1,k=1} \\ \beta_{j=2,k=1} \\ \vdots \\ \beta_{j=p,k=1} \\ \beta_{j=1,k=2} \\ \vdots \\ \beta_{j=1,k=q} \\ \vdots \\ \beta_{j=p,k=q} \end{pmatrix}$$

• Link:

Link covariates to response.

First assume that **y** has the same mean and covariance structure as if $\mathbf{y}_i \sim Dir(\alpha_{1p}, \dots, \alpha_{ip})$ for all $i = 1, \dots, n$.

Then,

$$g(\alpha) = \log(\alpha) = X\beta$$
$$\log(\alpha_{ij}) = x_i\beta_j$$

Where $\beta_j = (\beta_{j,k=1}, \dots, \beta_{j,k=q})^t$, and $x_i = (x_{i1}, \dots, x_{iq})$

This link function makes sense since $\alpha > 0$.

Dirichlet ideas

Assume
$$\eta_i = (\eta_{i1}, \dots, \eta_{ip}) \sim Dir(\alpha_{i1}, \dots, \alpha_{ip})$$
, and $\sum_{j=1}^p \alpha_{ij} = \alpha_{i0}$

Dirichlet Expected value

$$E(\eta_{ij}) = \frac{\alpha_{ij}}{\alpha_{i0}}$$

Dirichlet variance

$$Var(\eta_{ij}) = \frac{\alpha_{ij}(\alpha_{i0} - \alpha_{ij})}{\alpha_{i0}^2(\alpha_{i0} + 1)}$$

Note: $0 \le Var(\eta_{ij}) \le 1$

Dirichlet covariance

For $i \neq j$

$$Cov(\eta_{is}, \eta_{it}) = -\frac{\alpha_{is}\alpha_{it}}{\alpha_{i0}^2(\alpha_{i0} + 1)}$$

Note this matrix is singular. Note this quantity is always negative

Dirichlet correlation

For $i \neq j$

$$Cor(\eta_{is}, \eta_{it}) = \frac{Cov(\eta_{is}, \eta_{it})}{\sqrt{Var(\eta_{is})Var(\eta_{it})}}$$

$$= -\frac{\alpha_{is}\alpha_{it}}{\alpha_{i0}^2(\alpha_{i0} + 1)} \cdot \sqrt{\frac{\alpha_{i0}^2(\alpha_{i0} + 1)}{\alpha_{is}(\alpha_{i0} - \alpha_{is})}} \sqrt{\frac{\alpha_{i0}^2(\alpha_{i0} + 1)}{\alpha_{it}(\alpha_{i0} - \alpha_{it})}}$$

$$= -\sqrt{\frac{\alpha_{is}\alpha_{it}}{(\alpha_{i0} - \alpha_{is})(\alpha_{i0} - \alpha_{it})}}$$

Note this is always negative

GEEs

The collection of measurements taken on a single sample are assumed to be correlated. The entries in \mathbf{y}_i for a given i are correlated.

One such analysis method useful for correlated data are Generalized Estimating Equations. Originally proposed by

The GEE equations are

$$\sum_{i=1}^n \left(\frac{\partial \boldsymbol{\mu}_i}{\partial \boldsymbol{\beta}}\right)_{pq \times p}^t \mathbf{V}_{i_{p \times p}}^{-1} (\mathbf{Y_i} - \boldsymbol{\mu}_i)_{p \times 1} = 0$$

- $diag(A_i) = \sqrt{Var(\eta_i)}$: Square root of dirichlet variance.
- R_i : working correlation matrix, see below, mixture of dirichlet and phylogenetic correlation.
- $\left(\frac{\partial \mu_i}{\partial \beta}\right)$: Matrix of partial derivatives, defined below
- Y_i : response ASV proportions $0 < Y_i < 1$
- μ_i : Expected mean: $E(Y_i) = \frac{\alpha_i}{\alpha_{i0}}$

Derivation of $\left(\frac{\partial \boldsymbol{\mu}_i}{\partial \boldsymbol{\beta}}\right)$

GEE algorithm

Let G be the GEE equation, and H be the hessian.

Identifiability issues

Note that we can write the mean:

$$\mu_{ij} = \frac{\alpha_{ij}}{\alpha_{i0}}$$

$$= \frac{e^{x_i\beta_j}}{\sum_{s=1}^p e^{x_i\beta_s}}$$

$$= \frac{e^{x_i\beta_j}}{e^{x_i\beta_1} + \dots + e^{x_i\beta_p}}$$

$$= \frac{e^{\gamma}}{e^{\gamma}} \frac{e^{x_i\beta_1} + \dots + e^{x_i\beta_p}}{e^{x_i\beta_1} + \dots + e^{x_i\beta_p}}$$

$$= \frac{e^{\gamma}}{e^{\gamma}} \frac{e^{x_i\beta_1} + \dots + e^{x_i\beta_p}}{e^{x_i\beta_1} + \dots + e^{x_i\beta_p}}$$

Thus μ_{ij} is the same regardless of if $\beta = (\beta_1, \dots, \beta_p)$ or $\beta = (\beta_1 + \gamma, \dots, \beta_p + \gamma)$.

Thus the mean is non-identifiable.

Check if this is the same for the variance.

Penalty

To solve this problem, we first attempt to add a penalty.

Penalty:

$$\lambda \left(\sum_{j=1}^{p} \beta_{p} \right)^{2} = \lambda \beta^{t} 11^{t} \beta$$

New GEE eqn

$$G^* = G + \frac{\partial \text{Penalty}}{\partial \beta}$$
$$= G + 2\lambda 11^t \beta$$

New Hessian:

$$H^* = H + \frac{\partial^2 \text{Penalty}}{\partial^2}$$
$$= H + 2\lambda 11^t$$