GLMs and Causal Inference

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GLM examples

Causal Inference

GLM Interlude



- Some more GLM examples...
- Before looking at details for non-parametrics, let us re-visit the details of the GLM specification.
- Recall for a Bernoulli random variable has pmf

$$f_Y(y) = \theta^y \{1 - \theta\}^{1 - y}$$

$$= \exp\{y \log \frac{\theta}{1 - \theta} + \log(1 - \theta)\}. \tag{1}$$

Clearly here we set $\phi = \log \frac{\theta}{1-\theta}$. We can solve for $\exp(\phi) = \frac{\theta}{1-\theta}$, with

$$heta = rac{1}{1 + \mathsf{exp}(\phi)} \Rightarrow 1 - heta = rac{\mathsf{exp}(\phi)}{1 + \mathsf{exp}(\phi)}.$$

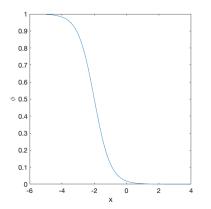
For example we could look at a single covariate x_i and set

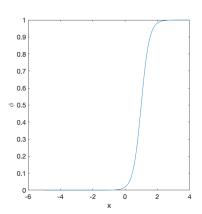
$$\phi_i = \beta_0 + \beta_1 x_i.$$

We see directly that as ϕ ranges across any value, θ is constrained to lie between zero and unity.

GLM Interlude







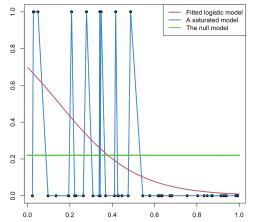
GLM Interlude



• We can examine the improvement using the deviance. Recall that

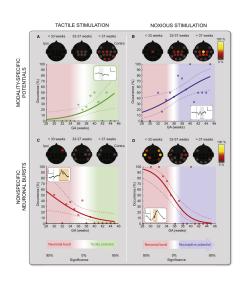
$$D = 2\{\ell_n(\widehat{\phi}) - \ell_n(\widehat{\beta})\}.$$

So we can fit both models and compare



Bernoulli and Binomial





Poisson observations



Looking at the Poisson pmf we have

$$f_Y(y) = \frac{e^{-\mu} \mu^y}{y!} = \exp\{-\mu + y \log(\mu) - \ln y!\}.$$
 (2)

Here we clearly set $\phi = \log(\mu)$. Solving for μ just gives us $\mu = \exp(\phi)$. We only need the mean to remain positive so this will fix our problem.

• Again we use the deviance to assess the fit; and would compare to the model μ_i is different for each value of i.

What about the sparse GLM?



• Hastie and Park (2007) estimate the parameters of the GLM using

$$\hat{\beta}_L(\lambda) = \arg\min_{\beta} \{ -\log L(\beta) + \lambda \|\beta\|_1. \}$$

- This mimics using the Lasso for the Gaussian linear model.
- We can study the geometry of this space in β . Unfortunately unlike the LASSO it is not a convex optimisation problem. This means we are not seeing the possibility of a polynomial-time algorithm solving our problem. We could also end up with multiple optima.
- Hastie and Park also extended the <u>elastic net</u> to this setting

$$\hat{\beta}_{EN}(\lambda_1, \lambda_2) = \arg\min_{\beta} \left\{ -\log L(\beta) + \lambda_1 \|\beta\|_1 + \lambda_2 \|\beta\|_2^2 \right\}. \tag{3}$$

• This has two penalties. Problems that arise when X has linearly dependent columns; the coefficient estimates are highly unstable.

What about the sparse GLM?



- When λ_2 is a constant, and λ_1 varies in an open set, such that the current active set remains the same, a unique, continuous and differentiable function.
- The additional penalization of the elastic net, is not either yielding a convex problem.
- Just optimizing the GLM likelihood can be problematic on its own.
- This brings us back to the penalized GLM. Augugliaro et al (2013) looked at the differential geometry of this problem.



So far: how to estimate $g:\mathbb{R} o \mathbb{R}$ (assumed smooth) in

$$Y_i = g(x_i) + \varepsilon_i, \quad \varepsilon_i \stackrel{iid}{\sim} \mathcal{N}(0, \sigma^2), \text{ given data } \quad \{(Y_i, x_i)\}_{i=1}^n.$$

Can extend to GLM setting as:

$$Y_i|x_i \overset{indep}{\sim} \exp\left\{g(x_i)y - \gamma(g(x_i)) + S(y)\right\}$$

ullet Parametrise candidate g via spline

$$s(x) = \sum_{j=1}^n \gamma_j B_j(x).$$

 \bullet Define matrices B and Ω as before,

$$B_{ij} = B_j(x_i), \quad \Omega_{ij} = \int B_i''(x)B_j''(x)dx$$

And consider penalised likelihood, similarly as with penalised GLM

$$\ell_n(\gamma) + \lambda \gamma^ op \Omega \gamma = \gamma^ op B^ op Y - \sum_{i=1}^n \gamma(b_i^ op \gamma) + \lambda \gamma^ op \Omega \gamma.$$



How can we generalise to multivariate covariates?

▶ "Immediate" Generalisation: $g: \mathbb{R}^p \to \mathbb{R}$ (smooth)

$$Y_j = g(x_{j1}, \dots, x_{jp}) + \varepsilon_j, \quad \varepsilon_j \stackrel{iid}{\sim} \mathcal{N}(0, \sigma^2)$$

- ▶ Estimation by (e.g.) multivariate kernel method.
- ▶ Two basic drawbacks of this approach . . .
- ← Curse of dimensionality





What is "local" in \mathbb{R}^p , though?

- → Need some definition of "local" in the space of covariates
- \hookrightarrow Use some metric on $\mathbb{R}^p \ni (x_1, \dots, x_p)^\top$!

But which one?

- Choice of metric choice of geometry
 - ← e.g., curvature reflects intertwining of dimensions
- Geometry \implies reflects structure in the covariates
 - potentially different units of measurement (variable stretching of space)
 - a may be of higher variation in some dimensions (need finer neighbourhoods there)
 - statistical dependencies present in the covariates ("local" should reflect these)



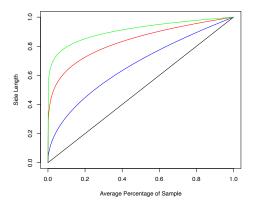


Figure: Curse of Dimensionality (Unif[0, 1]^p): p = 1, p = 2, p = 5, p = 10





Curse of Dimensionality

"neighbourhoods with a fixed number of points become less local as the dimensions increase"

Bellman (1961)

- Hence to allow for reasonably small bandwidths
 → Density of sampling must increase.
- → Density of sampling must increase
- Need to have ever larger samples as dimension grows.



Attempt to find a link/compromise between:

- our mastery of 1D case (at least we can do that well ...),
- and higher dimensional covariates (and associated difficulties).

Perhaps something that can be fitted/interpreted variable-by-variable?

▶ Compromise: Additive Model

$$Y_i = lpha_i + \sum_{k=1}^p f_k(x_{ik}) + arepsilon_i, \quad arepsilon_i \stackrel{iid}{\sim} \mathcal{N}(0, \sigma^2),$$

with f_k 's univariate smooth functions, $\sum_i f_k(x_{ik}) = 0$.

► Can extend to Generalised Additive Model:

$$|Y_i|x_i^{ op} \stackrel{indep}{\sim} \exp\left\{lpha_i y + y \sum_{k=1}^p f_k(x_{ik}) - \gamma\left(lpha_i + \sum_{k=1}^p f_k(x_{ik})
ight) + S(y)
ight\}$$



- ▶ How to fit additive model? Consider Gaussian case only for simplicity.
- \hookrightarrow Know how to fit each f_k separately quite well
- ► Consider *i*th response:

$$\mathbb{E}\left[Y_i - \alpha - \sum_{m \neq k} f_m(x_{im})\right] = f_k(x_{ik})$$

- ▶ Suggests the Backfitting Algorithm:
- (1) Initialise: $\alpha = \overline{Y}$, $f_k = f_k^0$, $k = 1, \ldots, p$.
- (2) Cycle: Get f_k by 1D smoothing of partial residual scatterplot

$$\left\{\left(Y_i-lpha-\sum_{m
eq k}f_m(x_{im}),x_{ik}
ight)
ight\}_{i=1}^n=\{e_{ik},x_{ik}\}_{i=1}^n.$$

- (3) Stop: when individual functions don't change
- ▶ Any smoother can be used, usually splines.



A different approach is inspired by tomography. Model Gaussian response as:

$$Y_i = \underbrace{\sum_{k=1}^K h_k(x_i^ op eta_k)}_{=g(x_i^ op)} + arepsilon_i, \quad ||eta_k|| = 1, \,\, arepsilon_i \,\, \stackrel{ ext{i.i.d.}}{\sim} \,\, \mathcal{N}(0,\sigma^2).$$

- Also additively decomposes g into smooth functions $h_k : \mathbb{R} \to \mathbb{R}$.
- ullet But each function now depends on a global linear feature $oldsymbol{x}_i^ opoldsymbol{eta}_k$
 - → a linear combination of the covariates
 - $\hookrightarrow ||\beta_k|| = 1$ for identifiability.
- Projections directions to be chosen for best fit (nonlinear problem)
- Each h_k is a ridge function of x_i^{\top} : varies only in the direction defined by β_k

Pros and Cons:

- (+) By classical Fourier series, can show that any $C^1([0,1]^p) \to \mathbb{R}$ function is uniformly approximated arbitrarily well as $K \to \infty$. Useful for prediction.
- (-) Interpretability? What do terms mean within problem?



How is the model fitted to data?

Assume only one term, K = 1 and consider penalized likelihood:

$$\min_{h_1 \in C^2[[0,1], ||\beta|| = 1} \qquad \left\{ \sum_{i=1}^n \{ Y_i - h_1(\boldsymbol{x}_i^\top \boldsymbol{\beta}) \}^2 + \int_0^1 \{ h_1''(t) \}^2 dt \right\}.$$

Two steps:

- Smooth: Given a direction eta, fitting $h_1(x_i^{ op}eta)$ is done via 1D smoothing.
- Pursue: Given h_1 , have a non-linear regression problem w.r.t. β .

Hence, iterate between the two steps

- \hookrightarrow Complication is that h_1 not explicitly known, so need numerical derivatives.
- → Computationally intensive (impractical in the '80's but doable today).

Further terms added in forward stepwise manner.



If β_k needs to be estimated non-linearly anyway...

$$g(x_i^ op)pprox \sum_{k=1}^K h_k(x_i^ op oldsymbol{eta}_k)$$

 \dots do we really need to estimate the h_k or can we fix them?

Theorem (Nonlinear Sigmoidal Approximation)

Let $\Psi: \mathbb{R} \to [0,1]$ be a strictly increasing distribution function and $g: [0,1]^p \to \mathbb{R}$ be an arbitrary continuous function. Then, for any $\epsilon > 0$, there exists $K < \infty$ and vectors $\alpha, t \in \mathbb{R}^K$ and $\{\beta_1,...,\beta_K\} \subset \mathbb{R}^p$ such that

$$\sup_{x \in [0,1]^d} \left| g(x) - \sum_{k=1}^K lpha_k \Psi(t_k + oldsymbol{x}^ op eta_k)
ight| < \epsilon.$$

- Can take h_k to be translations of the same known function $\Psi!$
- The tradeoff is that K may need to be quite large (interpretability?)
- Called a (single layer) neural network by analogy to synaptic function.
- A parametric model with many parameters fit by nonlinear least squares (gradient descent)



What about including transformations of the original covariates?

① Can of course include J transformations $w_i: \mathbb{R}^p \to \mathbb{R}$

$$(u_1,...,u_p)\mapsto w_j(u_1,...,u_n), \qquad j=1,...,J,$$

of the original variables as additional covariates by suitably enlarging the design matrix \boldsymbol{X} .

② We simply adjoin to X another J columns of dimension $n \times 1$ each:

$$\left(egin{array}{c} w_j(oldsymbol{x}_1^ op) \ dots \ w_j(oldsymbol{x}_n^ op) \end{array}
ight) \qquad j=1,...,J.$$

3 Which functions w_i should we pick though?

Since we've gone nonlinear anyway,

why not attempt to learn which transformations to include from the data?



How?

- Instead of including our original covariates (p columns of X)...
- ... use q derived covariates (q can be larger than p)

$$\left(egin{array}{c} w_1(x_1^ op) \ dots \ w_1(x_n^ op) \end{array}
ight), \left(egin{array}{c} w_2(x_1^ op) \ dots \ w_2(x_n^ op) \end{array}
ight), \quad \cdots \quad , \left(egin{array}{c} w_q(x_1^ op) \ dots \ w_q(x_n^ op) \end{array}
ight)$$

ullet ... where the q transformations $\{w_j\}_{j=1}^q$ are to be estimated from the data.

Recycling our nonlinear approximation theorem, write

$$w_j(x^ op)pprox \sum_{m=1}^{M_j} \delta_{m,j} \Psi(s_{m,j}+x^ op \gamma_{m,j})$$

using the same Ψ , and needing to estimate $(\delta_j, s_j, \gamma_{1,j}, ..., \gamma_{M_j,j})$, for j=1,...,q.



Assuming that we've constructed our new variables, we have a new design matrix

$$\left(egin{array}{ccc} w_1(x_1^ op) & \dots & w_q(x_1^ op) \ dots & & dots \ w_1(x_n^ op) & \dots & w_q(x_n^ op) \end{array}
ight)$$
 .

Summarising, we have defined a hierarchical nonlinear regression model:

$$Y_i = \sum_{k=1}^K lpha_k \Psi \Big(t_k + (w_i(oldsymbol{x}_1^ op),...,w_i(oldsymbol{x}_n^ op))eta_k \Big) + arepsilon_i =$$

$$= \sum_{k=1}^K \alpha_k \Psi\left(t_k + \left(\sum_{m=1}^{M_1} \delta_{m,1} \Psi(s_{m,1} + x^\top \gamma_{m,1}),...,\sum_{l=1}^{M_q} \delta_{l,q} \Psi(s_{l,q} + x^\top \gamma_{l,q})\right) \beta_k\right) + \varepsilon_i$$

... known these days as a two-layer neural network.

- Can add more layers ("deep neural network").
- Highly non-linear and non-convex cascade of simple nonlinearities applied to linear transformations.
- More easily perceived visually through a graphical representation

Causal Inference



- If we say "X causes Y"; mathematically this means *changing* the value of x *changes* the distribution of Y.
- When X causes Y then X and Y will be associated (one type of association is correlation), but the converse is generally <u>not</u> true.
- We shall discuss this in terms of <u>counterfactual</u> random variables.
- Let us start by a simple binary setup. Let X=1 denote the event that a unit was "treated" and X=0 denote the event that a unit was not "treated".
- We use the term "treated" in a very broad sense. Instead we might have used "exposed" and "not-exposed".
- Let *Y* be some <u>outcome variable</u>. To distinguish between association and causation we need to enhance our vocabulary.

Causal Inference II



- Two new symbols C_0 and C_1 are introduced to denote potential outcomes.
- C_0 is the outcome if the unit was not treated, and similarly, C_1 is the outcome if the unit was treated. These are both random variables. Thus

$$Y=C_X. (4)$$

This is the consistency relationship.

- Note that many things are unobserved in this model. When X=1 then we do not observe C_0 for those cases; also when X=0 we do not observe C_1 . We call those outcomes <u>counterfactual</u>.
- Thus (C_0, C_1) are hidden or latent variables.

Causal Inference III



• Define the average causal effect to be

$$\theta = \mathbb{E}\{C_1\} - \mathbb{E}\{C_0\}. \tag{5}$$

 θ is the difference in effect if everyone was treated versus if everyone was not. If C_0 and C_1 were binary then we can define the <u>causal odds ratio</u>

$$\frac{\frac{\Pr\{C_1=1\}}{\Pr\{C_1=0\}}}{\frac{\Pr\{C_0=1\}}{\Pr\{C_0=0\}}}.$$

• We also define the causal relative risk:

$$\frac{\Pr\{C_1 = 1\}}{\Pr\{C_0 = 1\}}.$$

Define the association of Y with X to be

$$\alpha = \mathbb{E}\{Y \mid X = 1\} - \mathbb{E}\{Y \mid X = 0\}. \tag{6}$$

Causal Inference III



- Theorem (Association is not causation): In general $\theta \neq \alpha$.
- Example: Suppose that we have observed the following units for a treatment:

Table: Causation vs association.

X	Y	C_0	C_1
0	0	0	0*
0	0	0	0*
0	0	0	0*
0	0	0	0*
1	1	1*	1
1	1	1*	1
1	1	1*	1
1	1	1*	1

Asterisks are indicating unobserved values.

Causal Inference IV



• For every experimental unit $C_0 = C_1$ and so the "treatment" has no effect.

$$\theta = \mathbb{E}\{C_1\} - \mathbb{E}\{C_0\} \tag{7}$$

$$= \frac{1}{8} \sum_{i=1}^{8} C_{1i} - \frac{1}{8} \sum_{i=1}^{8} C_{0i} = \frac{1}{8} \sum \{C_{1i} - C_{0i}\} = 0$$
 (8)

Thus the average causal effect is zero.

• We can also estimate the association:

$$\alpha = \mathbb{E}\{Y \mid X = 1\} - \mathbb{E}\{Y \mid X = 0\} = \frac{1 + 1 + 1 + 1}{4} - \frac{0 + 0 + 0 + 0}{4}$$
$$= 1.$$

Thus in this example $\theta \neq \alpha$.