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# Outline

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# The exponential family

A probability distribution belongs to the exponential family if its density can be modeled as:

$$p(\mathbf{y}|\boldsymbol{\eta}) = \frac{1}{Z(\boldsymbol{\eta})} h(\mathbf{y}) \exp\left[\boldsymbol{\eta}^{\top} \mathcal{T}(\mathbf{y})\right] = h(\mathbf{y}) \exp\left[\boldsymbol{\eta}^{\top} \mathcal{T}(\mathbf{y}) - A(\boldsymbol{\eta})\right]$$
(1)

where:

- $Z(\eta)$  is the partition function (normalization constant)
- h(y) is the base measure (scaling constant; typically 1)
- ullet  $\eta$  are the natural/canonical parameters
- $A(\eta) = \ln Z(\eta)$  is the log-partition function

The log-likelihood is then given by:

$$\log p(\mathbf{y}|\boldsymbol{\eta}) = \log h(\mathbf{y}) + \boldsymbol{\eta}^{\top} \mathcal{T}(\mathbf{y}) - A(\boldsymbol{\eta}) + \text{const}$$
 (2)

# Properties of exponential family

• Generalization: we define  $\eta = f(\phi)$ , thus:

$$p(\mathbf{y}|\phi) = h(\mathbf{y}) \exp\left[f(\phi)^{\top} \mathcal{T}(\mathbf{y}) - A(f(\phi))\right]$$
(3)

- If  $f(\phi)$  is nonlinear, then the model is in the curved exponential family
- If  $\eta = f(\phi) = \phi$ , the model is in **canonical form**
- If  $\mathcal{T}(y) = y$ , the model is in the natural exponential family

$$p(\mathbf{y}|\boldsymbol{\eta}) = h(\mathbf{y}) \exp\left[\boldsymbol{\eta}^{\top} \mathbf{y} - A(\boldsymbol{\eta})\right]$$
(4)

# Bernoulli distribution in exponential family form (1/2)

The Bernoulli distribution is given by:

$$p(y|\mu) = \mu^{y}(1-\mu)^{1-y}, \quad y \in \{0,1\}, \quad 0 < \mu < 1$$
 (5)

where  $\mu = \mathbb{E}(y)$  is the probability of success. Rewriting:

$$p(y|\mu) = (1-\mu) \left(\frac{\mu}{1-\mu}\right)^y = (1-\mu) \exp\left[y \log\left(\frac{\mu}{1-\mu}\right)\right]$$

$$= (1-\mu) \exp\left[y \log\left(\frac{\mu}{1-\mu}\right) - 0\right]$$

Comparing to the exponential family form:

$$h(y) = 1 - \mu$$
 (base measure)  
 $\mathcal{T}(y) = y$  (sufficient statistic)  
 $\eta = \log\left(\frac{\mu}{1-\mu}\right)$  (natural parameter)  
 $A(\eta) = 0$  (log-partition function)

## Cumulant generating function

- Cumulants  $\kappa_n(\mathbf{y})$  are functions of the central moments of a distribution
- ullet For example,  $\kappa_1(oldsymbol{y}) = \mathbb{E}(oldsymbol{y})$  and  $\kappa_2(oldsymbol{y}) = \mathbb{V}(oldsymbol{y})$
- Higher order cumulants are polynomial functions of the central moments
- The cumulants of a distribution are defined by the cumulant generating function (CGF):

$$K_{\mathbf{y}}(t) = \log \mathbb{E}(\exp(t\mathbf{y}))$$
 (6)

where  $\mathbb{E}(\exp(t\mathbf{y}))$  is the moment generating function (MGF) of  $\mathbf{x}$ 

- In the exponential family, the log-partition function  $A(\eta)$  is the CGF of the sufficient statistics  $\mathcal{T}(\mathbf{y})$
- Thus, the cumulants can be obtained by differentiating  $A(\eta)$ :

$$\kappa_1(\mathcal{T}(\mathbf{y})) = \mathbb{E}(\mathcal{T}(\mathbf{y})) = \nabla_{\boldsymbol{\eta}} A(\boldsymbol{\eta})$$

$$\kappa_2(\mathcal{T}(\mathbf{y})) = \operatorname{Cov}(\mathcal{T}(\mathbf{y})) = \nabla_{\boldsymbol{\eta}}^2 A(\boldsymbol{\eta})$$

# Unique global maximum of the likelihood

From the CGF properties, we have:

$$\nabla_{\boldsymbol{\eta}}^2 A(\boldsymbol{\eta}) = \mathsf{Cov}(\mathcal{T}(\boldsymbol{y})) > 0 \tag{7}$$

This implies that the log-partition function  $A(\eta)$  is strictly convex. Thus, the log-likehood

$$\log p(\mathbf{y}|\boldsymbol{\eta}) = \log h(\mathbf{y}) + \boldsymbol{\eta}^{\top} \mathcal{T}(\mathbf{y}) - A(\boldsymbol{\eta}) + \text{const}$$
 (8)

is guaranteed to have a unique global maximum.

# The generalized linear model (GLM)

• Conventional linear regression models have the form:

$$p(y|\mathbf{x}, \mathbf{w}) \sim \mathcal{N}(y|\mathbf{x}^{\top}\mathbf{w}, \sigma^2)$$
 (9)

#### where

- y<sub>i</sub> is a continuous response
- $x_i$  is a vector of quantitative and/or qualitative explanatory variables
- Generalized linear models (GLMs) were introduced to extend this framework to allow y<sub>i</sub> to be modeled by other exponential family distributions besides the normal/Gaussian, e.g.
  - exponential
  - binomial/multinomial (with fixed number of trials)
  - Poisson
- In the GLM framework:
  - The mean of  $y_i$  is given by  $\mu_i$
  - $\mu_i$  can be specified by a nonlinear function of  $\mathbf{x}_i^{\mathsf{T}} \mathbf{w}$
  - Note that the simple linear regression is a special case of GLM in which  $\mu_i = \mathbf{x}_i^{\top} \mathbf{w}$  and  $y_i$  follows a Gaussian distribution

### **GLM** formulation

The GLM is a version of the exponential family distribution in which the natural parameters  $\eta_n$  are a **linear function** of the output. It is given by:

$$p(y_n|\mathbf{x}_n, \mathbf{w}, \sigma^2) = \exp\left[\frac{y_n\eta_n - A(\eta_n)}{\sigma^2} + \log h(y_n, \sigma^2)\right]$$
(10)

where:

- $\eta_n = \mathbf{w}^{\top} \mathbf{x}_n$  is the natural parameter (input)
- $y_n = \mathcal{T}(y_n)$  is the sufficient statistic
- $A(\eta_n)$  is the log-partition function (or log normalizer)
- $h(y_n, \sigma^2)$  is the base measure
- $\sigma^2$  is the dispersion parameter (typically known or set to 1)

## Link and mean functions

Exponential family

Recalling that the mean and variance of the sufficient statistics  $\mathcal{T}(y_n) = y_n$  are given by the first and second derivatives of the log-partition function  $A(\eta_n)$ , we have:

$$\mathbb{E}(y_n|\mathbf{x}_n,\mathbf{w},\sigma^2) = A'(\eta_n) = \ell^{-1}(\eta_n)$$
 (11)

$$Var(y_n|\mathbf{x}_n,\mathbf{w},\sigma^2) = A''(\eta_n)\sigma^2$$
 (12)

We define the **mean function** as

$$\mu_n = \ell^{-1}(\eta_n) \tag{13}$$

and the link function as its inverse:

$$g(\mu_n) = \ell(\mu_n) = \eta_n \tag{14}$$

The link function is thus the inverse of the mean function.

Linear regression (1/2)

Linear regression has the form:

$$p(y_n|\mathbf{x}_n, \mathbf{w}, \sigma^2) = \frac{1}{\sqrt{2\pi\sigma^2}} \exp\left(-\frac{1}{2\sigma^2}(y_n - \mathbf{w}^\top \mathbf{x}_n)^2\right)$$
(15)

Taking logs:

$$\log p(y_n|\mathbf{x}_n, \mathbf{w}, \sigma^2) = -\frac{1}{2}\log(2\pi\sigma^2) - \frac{1}{2\sigma^2}(y_n - \mathbf{w}^\top \mathbf{x}_n)^2$$
 (16)

Setting  $\eta_n = \mathbf{w}^{\top} \mathbf{x}_n$ , we can write in GLM form as:

$$\log p(y_n|\mathbf{x}_n, \mathbf{w}, \sigma^2) = \frac{y_n \eta_n - \eta_n^2/2}{\sigma^2} - \frac{1}{2} \left( \frac{y_n^2}{\sigma^2} + \log(2\pi\sigma^2) \right)$$
(17)

If we set:

$$A(\eta_n) = \eta_n^2/2 \tag{18}$$

$$h(y_n, \sigma^2) = \frac{1}{\sqrt{2\pi\sigma^2}} \exp\left(-\frac{1}{2\sigma^2}y_n^2\right)$$
 (19)

then we can write:

$$\log p(y_n|\mathbf{x}_n,\mathbf{w},\sigma^2) = \frac{y_n\eta_n - A(\eta_n)}{\sigma^2} + \log h(y_n,\sigma^2)$$
 (20)

And thus, the cumulants are given by:

$$\mathbb{E}(y_n|\mathbf{x}_n,\mathbf{w},\sigma^2) = A'(\eta_n) = \eta_n = \mathbf{w}^\top \mathbf{x}_n$$
 (21)

$$Var(y_n|\mathbf{x}_n,\mathbf{w},\sigma^2) = A''(\eta_n)\sigma^2 = \sigma^2$$
 (22)

## GLM components

A GLM can be considered as consisting of three parts:

- Random component: this is the probability distribution of the response variable
- Systematic component: specifies the explanatory variables within the linear combination of their coefficients (Xw)
- Link function  $g(\mu)$ : defines the relationship between the random and systematic components:
  - Simple linear regression (identity link function):

$$g(\mu_n) = g(\mathbb{E}(y_n)) = \mathbf{x}_n^{\top} \mathbf{w}$$
 (23)

Binary logistic regression (logit link function):

$$g(\mu_n) = g(p(\mathbf{x}_n)) = \operatorname{logit}(p(\mathbf{x}_n)) = \operatorname{ln}\left(\frac{p(\mathbf{x}_n)}{1 - p(\mathbf{x}_n)}\right) = \mathbf{x}_n^{\top} \mathbf{w}$$
 (24)

# Assumptions of GLM

- The observations of the response variable y are i.i.d.
- Response variable  $y_n$  is typically exponentially distributed (not restricted to being normally distributed)
  - Implies that errors need not be normally distributed (but should be independent)
- Link function is linear with respect to the coefficients  $(w_d)$ 
  - Relationship between response and explanatory variables does not have to be linear
  - Explanatory variables can be nonlinear transformations of original values (as in simple linear regression)
- Variance may not homogeneous (i.e. homoscedasticity is not a requirement)
- Parameters are estimated via MIF

# Commonly used GLM models and their components

Model	Random component	Link function
Linear regression	Gaussian	Identity: $g(\mu_n) = \mu_n = \mathbf{w}^{ op} \mathbf{x}_n$
Binary logistic regression	Bernoulli	Logit: $g(\mu_n) = \log\left(\frac{\mu_n}{1-\mu_n}\right)$
Probit regression	Bernoulli	Probit: $g(\mu_n) = \Phi^{-1}(\mu_n)$
Multinomial logit/logistic	Categorical	Multinomial logit: $g(\mu_{nc}) = \log\left(rac{\mu_{nc}}{\mu_{nc}} ight)$
Poisson regression	Poisson	$Log: \ g(\mu_{n}) = log(\mu_{n})$

Note that in all cases, the link function always results in:

$$g(\mu_i) = w^{\top} x_i \tag{25}$$

Its job is to "link" the response to the systematic component via a suitable transformation that results in a linear function of the w's.

## MLE of GLM parameters

The negative log-likelihood (ignoring constant terms) is given by

$$NLL(\boldsymbol{w}) = -\log p(\mathcal{D}|\boldsymbol{w}) = -\sum_{n=1}^{N} \log p(y_n|\boldsymbol{x}_n, \boldsymbol{w}) = \sum_{n=1}^{N} \frac{A(\eta_n)}{\sigma^2} - \frac{y_n \eta_n}{\sigma^2}$$
 (26)

If we set  $\ell_n = \eta_n y_n - A(\eta_n)$ , then the NLL can be written as:

$$NLL(\boldsymbol{w}) = -\sum_{n=1}^{N} \frac{\ell_n}{\sigma^2}$$
 (27)

where  $\eta_n = \mathbf{w}^{\top} \mathbf{x}_n$ .

The gradient of the NLL is then given by:

$$\mathbf{g}_n = \sum_{n=1}^N \frac{y_n - \mu_n}{\sigma^2} \mathbf{x}_n \tag{28}$$

where  $\mu_n = A'(\eta_n) = \ell^{-1}(\eta_n)$  is the mean function.