Logistic Regression

$$Odds = \frac{\pi}{1 - \pi}$$

$$logit(\pi_i) = log(\frac{\pi_i}{1 - \pi_i}) = \alpha + \beta_i x_{ji}$$

Estimate α and β maximum likelihood

$$\eta_i = \alpha + \beta x_i$$

$$\frac{\hat{\pi}_i}{1 - \hat{\pi}_i} = \exp(\eta_i)$$

$$\hat{\pi}_i = \frac{\exp(\eta_i)}{1 - \exp(\eta_i)} = (1 + \exp(-\eta_i))^{-1}$$

Generalized Linear Models

$$g(\mu_i) = \eta_i = \boldsymbol{\beta}^\top \boldsymbol{x_i}$$

- Linear component: $\eta_i = oldsymbol{eta}^{ op} oldsymbol{x}_i$
- Link function: let univariate function g be monotonic (strictly decreases or increases) and differentiable
- Random component: y_i are independent and from an exponential family, which impose the variance of y_i depends on μ_i through a variance function $var(y_i) = \phi V(\mu_i)$ where ϕ is called the dispersion parameter.

Classic linear regression assumes normal dist.

Logistic regression assumes $g(\mu) = \log[\frac{\mu}{(1-\mu)}]$ and y_i is a Bernoulli trial ($\mathbb{V}(y) = \mu(1-\mu)$). Other possible links for binary responses include

- Probit:

 $g(\mu) = \Phi^{-1}(\mu)$, where Φ is the cumulative standard normal distribution

- Complementary log-log:

$$g(\mu) = \log(-\log(1-\mu))$$

Maximum Likelihood Estimation of Proportions

Suppose we draw a random sample of size n=5 from some population, send them an offer, and x=2 respond. What is our best guess of the response probability?

$$L(\pi) = {5 \choose 2} \pi^2 (1 - \pi)^3$$

- Let $l(\pi) = \log(L(\pi))$, log-likelihood

$$l(\pi) = \log(10) + 2\log(\pi) + 3\log(1 - \pi)$$

$$\frac{dl(\pi)}{d\pi} = \frac{2}{\pi} - \frac{3}{1 - \pi} = 0 \Rightarrow \hat{\pi} = \frac{2}{5}$$

Maximum Likelihood Estimation of Means

Suppose we draw a random sample of size n=3 from a normal population with unknown mean μ and known variance $\sigma^2=5$. The observed values are $x_1=4, x_2=5, x_3=6$. What is the best guess of μ ?

Pick μ so that the probability of observing the three values given μ is maximized:

$$f(x \mid \mu, \sigma^2) = \frac{1}{\sqrt{2\pi\sigma^2}} \exp[-\frac{1}{2\sigma^2}(x - \mu)^2]$$

$$L(\mu) = \prod_{i=1}^{3} \left(\frac{1}{\sqrt{2\pi\sigma^2}} \exp\left[-\frac{1}{2\sigma^2} (x_i - \mu)^2 \right] \right)$$

$$l(\mu) = \sum_{i=1}^{3} \left[-\frac{1}{2} \log(2\pi\sigma^2) - \frac{1}{2\sigma^2} (x_i - \mu)^2 \right]$$

$$= -k_1 - k_2 \sum_{i=1}^3 (x_i - \mu)^2$$

$$\frac{dl(\mu)}{d\mu} = 2k_2 \sum_{i=1}^{3} (x_i - \mu) = 0 \Rightarrow \mu = \frac{1}{3} \sum_{i=1}^{3} x_i$$

Note:
$$\frac{d^2l(\mu)}{du^2} = -6k_2 < 0$$
 thus maximum

Likelihood Function for Logistic Regression

$$\log\left(\frac{\pi_i}{1-\pi_i}\right) = \alpha + \beta x_i$$

Probability distribution: $f_i(y_i) = \pi_i^{y_i} (1 - \pi_i)^{1-y_i}$

$$\Rightarrow f(y_1, ..., y_n) = \prod_{i=1}^n \pi_i^{y_i} (1 - \pi_i)^{1 - y_i}$$

$$\log(f) = l = \log(L) = \text{Log-Likelihood}$$

$$\sum_{i=1}^{n} y_i(\alpha + \beta x_i) - \sum_{i=1}^{n} \log[1 + \exp(\alpha + \beta x_i)]$$

Maximize this with respect to α and β

Residual versus null deviance

Residual deviance: deviance for full model

Null deviance: deviance for the intercept-only model (think of as SST)

Difference between them measures how much variation is explained by the model and plays the role of extra sums of squares. Can test overall significance of the model b/c it has a **Chi-squared distribution**.

> fit\$null.deviance - fit\$deviance
[1] 1095.99

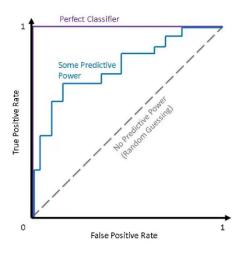
Likelihood-Ratio Test

We can test H_0 : $\beta_{p+1}=\cdots=\beta_{p+q}=0$ with the test statistic D_1-D_2 , which has a chi square distribution with q degrees of freedom (the D's represent deviance)

Predictive Accuracy of Classifiers

- GLMs minimize deviance
- AIC = deviance + 2p common penalized measure
- Classification rate: $\frac{TN+TP}{TN+FP+FN+TP}$
- Recall: $\frac{TP}{FN+TP}$
- TPR (True Positive Rate) (Sensitivity): $\frac{TP}{TP+FN}$
- FPR (False Positive Rate) (1-Specificity): $\frac{FP}{TN+FP}$
- Precision: $\frac{TP}{FP+TP}$
- **F1**: $\frac{2}{\frac{1}{Recall} + \frac{1}{Precision}} = 2 \frac{P*R}{P+R}$

- AUC (Area under curve): ROC curve is produced by calculating and plotting the true positive rate against the false positive rate for a single classifier at a variety of thresholds



Scoring Model:

- **Proxy behavior**: behavior that has been observed in the past that is similar to future behavior you would like to predict, e.g., response to similar offer sent yesterday

Warning: your model may not work because this is only a proxy behavior. Seasonality, the state of the economy, what competition is doing, etc. usually all affect response

- *Target period*: time period when proxy offer was active
- **Base period**: a period of time prior to the target period. Information from the base period will be used to predict proxy behavior

Performance usually assessed with a gains table

- 1. Find quantiles of predicted values \hat{y}
- 2. Compute number of responders and revenue by quantile, also averages
- 3. Compute cumulative counts and revenues by quantile, also averages and lifts

A	В	C	D	E	F	G	H	1	J	K	L	M
Quantile		Amount by Quantile			Cumulative				Lift			
%	of ŷ	Num Resp				Num Cont		Rev Amt		Avg Amt		
1	10569	1681	159501	0.159		10569		159501				3.18
2	10569	477	40241	0.0451	3.81	21138	2158	199742	0.102	9.45	1.91	1.99
3	10568	307	23716	0.0290	2.24	31706	2465	223458	0.0777	7.05	1.45	1.49
4	10569	201	15484	0.0190	1.46	42275	2666	238942	0.0631	5.65	1.18	1.19
5	10569	159	11608	0.0150	1.10	52844	2825	250549	0.0535	4.74	1	1

- Columns A and B: Quantiles of \hat{y}
- **Columns C and D**: number of responders and total revenue by quantile
- **Columns E and F**: response rate and average revenue in quantile, e.g., $\frac{1681}{10569}=15.9\%$ and $\frac{96728}{10569}=\$15.1$ per contact
- **Column G**: *Depth* of contacts or cumulative counts, e.g., 10,569 + 10,569 = 21,138
- **Columns H and I**: cumulative responders and revenue by quantile:

Row 2: 1,681 + 444 = 2,158 responders and 159,501 + 40,241 = 199,742 revenue at 40%

Last row: 2,825 total responders and 250,549 total

- **Columns J and K**: cumulative response rate and average revenue in quantile:

Row 2:
$$\frac{2,158}{21,138} = 9.45\%$$
 and $\frac{199,742}{21,138} = \9.45 per contact at 40%

Last row: **contacting at random** gives 5.35% respond rate, \$4.74 per contact

- **Columns L and M**: lift of model over random guessing, e.g. $\frac{15.1\%}{5.35\%}=2.98$ indicates the response rate from using model to pick best 20% of the names is improved by 57% over picking names at random. Revenue more than tripled (3.93)!

Generalized Logistic Regression Model

Let Y in $\{1,2,\ldots,K\}$ be a multinomial r.v. for the outcome with K values. The probability that observation i comes from class k is $\pi_{ik}=P(Y_i=k)$, where $\pi_{i1}+\cdots+\pi_{iK}=1$

Approaches:

- $\mathit{One}\ \mathit{vs}\ \mathit{all}$: fit $\mathit{K}\ \mathit{separate}\ \mathsf{logistic}\ \mathit{regression}\ \mathsf{models}$
- Generalized logit: fit K-1 models. Pick class 1 as the base category, but, as with the binary logit, this choice is arbitrary (models equivalent with a different base category)

Generalized logit model math:

- Model (k = 2, ..., K):

$$\log\left(\frac{\pi_k}{\pi_1}\right) = \boldsymbol{\beta}_k^{\mathsf{T}} \mathbf{x} \qquad (k = 2, ..., K)$$

- Solving for π_k :

$$\pi_k = \pi_1 \exp(\boldsymbol{\beta}_k^{\mathsf{T}} \mathbf{x}) \qquad (k = 2, ..., K)$$

- The probabilities must sum to 1:

$$\begin{split} 1 &= \Sigma_{j=1}^K \pi_j = \pi_1 + \Sigma_{(j=2)}^K \pi_1 \exp \left(\boldsymbol{\beta}_j^\top \mathbf{x} \right) \\ &= \pi_1 [1 + \Sigma_{j=2}^K \exp \left(\boldsymbol{\beta}_j^\top \mathbf{x} \right)] \end{split}$$

- Solving for π_1 :

$$\pi_1 = \frac{1}{1 + \Sigma_{j=2}^K \exp(\boldsymbol{\beta}_j^\mathsf{T} \mathbf{x})}$$

- Substituting back into formula for π_k :

$$\pi_k = \frac{\exp(\boldsymbol{\beta}_k^\mathsf{T} \mathbf{x})}{1 + \Sigma_{i=2}^K \exp(\boldsymbol{\beta}_i^\mathsf{T} \mathbf{x})} \qquad (k = 2, \dots, K)$$

Generalized logit estimation:

- Likelihood function:

$$L(\mathbf{B}) = \prod_{i=1}^{n} \prod_{k=1}^{K} \pi_{ik}^{z_{ik}}$$

Where $z_{ik}=1$ when $y_i=k$ and $z_{ik}=0$ otherwise

- The log-likelihood is the cost function

$$\log(L(\mathbf{B})) = l(\mathbf{B}) = \sum_{i=1}^{n} \sum_{k=1}^{K} z_{ik} \log(\pi_{ik})$$

- Denote optimal values of β_k by b_k . Compute estimated probabilities that each observation is in class k as:

$$p_{i1} = \frac{1}{1 + \Sigma_{i=2}^K \exp(\mathbf{b}_i^\mathsf{T} \mathbf{x}_i)}$$
 and $p_{ik} = \exp(\mathbf{b}_k \mathbf{x}^\mathsf{T}) \, p_{i1}$

- The maximum likelihood classifier assigns i to the class with the largest p_{ik}

Base category is y = 1

$$\log\left(\frac{\pi_2}{\pi_1}\right) = -8.69 + 1.42x_1 + 0.99x_2$$

$$\log\left(\frac{\pi_3}{\pi_1}\right) = -7.64 + 0.66x_1 + 1.52x_2$$

All four models give similar results

- There are three possible base categories and all give equivalent models

Call: multinom(formula = y ~ x1 + x2, data = train)

Coefficients:
(Intercept) x1 x2
2 -8.693704 1.4200155 0.989357
3 -7.644341 0.6586461 1.5195375
Residual Deviance: 68.61044
multinom(formula = factor(y, levels = c(2, 1, 3)) ~ x1 + x2, data = train)
Coefficients:
(Intercept) x1 x2
1 8.691776 -1.4197191 -0.9891887
3 1.047763 -0.7623966 0.5302961
Residual Deviance: 68.61043
multinom(formula = factor(y, levels = 3:1) ~ x1 + x2, data = train)
Coefficients:
(Intercept) x2
2 -1.047771 0.7622246 -0.5302758
1 7.643941 -0.6567638 -1.5194762

- All three models have the same deviance (cost function value)
- Let $\pmb{\beta}_{ij}$ be slopes for $\log\left(\frac{\pi_i}{\pi_j}\right) = \pmb{\beta}_{ij}^{\mathsf{T}} \mathbf{x}$. Note that $\frac{\pi_i}{\pi_i} = e^{\pmb{\beta}_{ij}^{\mathsf{T}} \mathbf{x}}$
- Clearly, $\boldsymbol{\beta}_{ij} = -\boldsymbol{\beta}_{ij}$, e.g.,

$$\log\left(\frac{\pi_3}{\pi_1}\right) = \beta_{31}^{\mathsf{T}} \mathbf{x} \Rightarrow \frac{\pi_3}{\pi_1} = e^{(\beta_{31}^{\mathsf{T}} \mathbf{x})} \Rightarrow \frac{\pi_1}{\pi_3} = e^{-\beta_{31}^{\mathsf{T}} \mathbf{x}}$$

 $- \beta_{23} = \beta_{21} - \beta_{31}$, e.g.,

$$0.7629246 = 1.4200155 - 0.6568461$$

$$\begin{split} \frac{\pi_2}{\pi_3} &= \frac{\pi_2}{\pi_3} * \frac{\pi_1}{\pi_1} = \frac{\pi_2}{\pi_1} * \frac{\pi_1}{\pi_3} = e^{\beta_{21}^\mathsf{T} \mathbf{x}} * e^{-\beta_{31}^\mathsf{T} \mathbf{x}} \\ &= e^{(\beta_{21} - \beta_{31})^\mathsf{T} \mathbf{x}} \end{split}$$

Making predictions

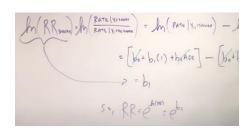
- Binary logit: individual chooses between two options and selects the one that provides greater utility
- Multinomial logit: individual chooses among more than two alternatives and selects the one that provides the greatest utility
- Ordered logit: individual reveals the strength of his or her preferences with respect to a single outcome
- **Conditional logit**: allows variables that <u>vary across</u> <u>alternatives</u> and possibly across the individuals as well, e.g., choice of mode of transportation (e.g., train, bus, car). Characteristics or attributes of these include time waiting, how long it takes to get to work, and cost.
- Log-linear analysis: class of models that subsumes (includes or absorbs) the logit models and more.

Poisson

- The PMF of a Poisson distribution:

$$P(Y = y) = \frac{\mu^{y}e^{-\mu}}{y!}, \mu > 0, y = 0,1,2,...$$

- Suppose we observe a random sample of ordered pairs (x_i,y_i) , i=1,...,n from where y_i has a Poisson distribution with mean μ_i . More generally, x_i could be a vector of predictors. Assume (log link function) (log(mu) -> rate ratio)



$$\log(\mu_i) = \eta_i = \alpha + \beta x_i \Longleftrightarrow \mu_i = e^{\eta_i}$$

- The likelihood is:

$$L = \prod_{i=1}^{n} \frac{e^{-\mu_i} \mu_i^{y_i}}{y_i!}$$

- The log-likelihood of a Poisson model is:

$$l = \log(L) = \sum_{i=1}^{n} [y_i \log(\mu_i) - \mu_i - \log(y_i!)]$$

- The log-likelihood of a saturated Poisson model is:

$$\hat{\mu}_i \equiv y_i \Rightarrow l_s = \sum_{i=1}^n [y_i \log(y_i) - y_i - \log(y_i!)]$$

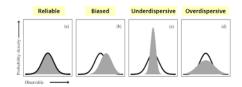
- Let $\mu_i \equiv \exp\!\left(\hat{\alpha} + \hat{\beta} x_i\right)$ be the MLEs. The deviance is:

$$D^{2} = -2(l - l_{s}) = -2\left[\sum_{i=1}^{n} y_{i} \log\left(\frac{\hat{\mu}_{i}}{y_{i}}\right) + \sum_{i=1}^{n} (y_{i} - \hat{\mu}_{i})\right]$$

Where
$$y_i \log \left(\frac{\hat{\mu}_i}{y_i}\right) = 0$$
 for $y_i = 0$

Beyond Poisson

- Recall that for Poisson Y, $\mathbb{E}(Y) = \mathbb{V}(Y) = \mu$, but in practice we may find $\mathbb{E}(Y) < \mathbb{V}(Y)$, called the problem of **overdispersion** (or **underdispersion** $\mathbb{E}(Y) > \mathbb{V}(Y)$)



- Negative binomial distribution (NBD) regression models allow for overdispersion
- Beyond overdispersion, we often observe too many zero values for either Poisson or NBD. What to do?

Zero-inflated Poisson (ZIP) assumes a mixture distribution

$$P(Y = 0) = \pi + (1 - \pi)e^{-\mu},$$

$$P(Y = y) = (1 - \pi) \frac{e^{-\mu} \mu^{y}}{y!}, \mu > 0, y = 1, 2, ...$$

Where $\pi \in [0,1]$ is the probability of extra (structural) zeros

Zero-inflated negative binomial (ZINP)

Hurdle models

- GLMs can accommodate other distributions including exponential and gamma

Gamma regression

- The PMF of a gamma distribution is:

$$f(y) = \frac{1}{\beta^{\alpha}\Gamma(\alpha)} y^{\alpha-1} e^{-\frac{y}{\beta}}, \alpha > 0, \beta > 0, y > 0$$

- Where α is the *shape* parameter and β is the *scale* parameter. The mean of a gamma distribution is $\mu=\alpha\beta$, so $\beta=\frac{\mu}{\alpha}$. The pdf can be written in terms of μ :

$$f(y) = \left(\frac{\alpha}{\mu}\right)^{\alpha} \Gamma(\alpha)^{-1} y^{\alpha - 1} e^{\frac{y\alpha}{\mu}}$$

- The log pdf is given by:

$$\log(f(y)) = \alpha \log(\alpha) - \alpha \log(\mu) - \log[\Gamma(\alpha)]$$
$$+(\alpha - 1)\log(y) - \frac{y\alpha}{\mu} = \alpha \left(-\frac{y}{\mu} - \log(\mu)\right)$$
$$-\log(\Gamma(\alpha)) + \alpha \log(\alpha y) - \log(y)$$

- The log-likelihood is:

$$\begin{split} l &= \Sigma_{i=1}^n \left[\alpha \left(-\frac{y_i}{\mu_i} - \log(\mu_i) \right) - \log[\Gamma(\alpha)] \right. \\ &+ \alpha \log(\alpha y_i) - \log(y_i) \right] \end{split}$$

- The log-likelihood of the saturate model is:

$$l_s = \sum_{i=1}^{n} [\alpha(-1 - \log(y_i)) - \log[\Gamma(\alpha)] + \alpha \log(\alpha y_i) - \log(y_i)]$$

- The deviance is $-2(l-l_s)$, but notice how the last three terms of l and l_s are identical and thus cancel out. Thus the deviance is:

$$-2(l - l_s) = -2\alpha \sum_{i=1}^{n} \left(-\frac{y_i}{\mu} - \log(\mu_i) + 1 + \log(y_i) \right)$$

$$= -2\alpha \Sigma_{i-1}^n \left(\log \left(-\frac{y_i}{\mu_i} \right) - \frac{y_i - \mu_i}{\mu_i} \right)$$

Regression Terms and Symbols:

Term	ACT	JWHT	Other
Sum of	SSE	RSS	
squared			
errors			
Total	SST	TSS	
sum of			
squares			
Mean	MSE		S_e^2
squared			
error			
Residual		RSE	
Standard			
Error			

SSE =
$$\sum_{i=1}^{n} [y_i - f(X_i)]^2$$

$$MSE = \frac{SSE}{n}$$

$$R^2 = 1 - \frac{SSE}{SST} = 1 - \frac{RSS}{(n-1)var(y)}$$

Penalized Estimates

$$R^2 = 1 - \frac{\frac{\overline{SSL}}{n-p-1}}{\frac{\overline{SST}}{n-1}}$$
 or AIC = deviance + 2p

Multicollinearity

Pipe: e.g., $x_1 \rightarrow x_2 \rightarrow y$. If you are studying $x_1 \rightarrow y$ then do not control for x_2

Latent construct: predictors manifestations of common, underlying **latent construct**, e.g., $w \rightarrow x_1$ and $w \rightarrow x_2$. Often, estimate w and use it instead of x_1 and x_2

Back door confound (fork): include control to block back-door path, e.g., if $w \to y$ and $w \to x \to y$ then control for w to study $x \to y$

Collider: usually do not control for colliders, e.g., if $x \to w$ and $y \to w$, then do not control for collider w when studying $x \to y$

Extras

$$\mathsf{F} = \frac{\frac{SST - SSE}{p}}{\frac{SSE}{n - p - 1}} = \frac{\left(\frac{\Delta SSE}{\Delta df}\right)}{S_e^2}, \text{ bottom from full model}$$

1. (9 points) The Poisson distribution has the following PMF

$$P(X = x) = \frac{\lambda^x e^{-\lambda}}{x!}, \quad (\lambda > 0; \quad x = 0, 1, 2, ...).$$

Suppose that we have a random sample of n=2 observations, x_1 and x_2 , from a Poisson distribution with parameter λ , where the observations are independent. This problem will derive the maximum likelihood estimate of λ .

(a) (2 points) Find $L(\lambda) = P(X_1 = x_1 \cap X_2 = x_2)$ Answer.

$$L(\lambda) = \frac{\lambda^{x_1}e^{-\lambda}}{x_1!} \cdot \frac{\lambda^{x_2}e^{-\lambda}}{x_2!} = \frac{\lambda^{x_1+x_2}e^{-2\lambda}}{x_1!x_2!}$$

(b) (2 points) Find $l(\lambda) = \log[L(\lambda)]$. Answer: $l(\lambda) = (x_1 + x_2) \log \lambda - 2\lambda - \log(x_1!x_2!)$

(c) (3 points) Find the value of λ that maximizes l(λ). Answer:

$$\frac{dl(\lambda)}{d\lambda} = \frac{x_1 + x_2}{\lambda} - 2 = 0$$

so we find $\lambda = (x_1 + x_2)/2$, the sample mean.

(d) (2 points) How do you know that your value of λ is a maximum? Answer: The second derivative is

$$\frac{d^2l(\lambda)}{d\lambda^2} = -\frac{x_1 + x_2}{\lambda^2} < 0,$$

since both x_j and λ are not negative.

- 2. (30 points) Data from 37 patients receiving a non-depleted allogencic bone marrow transplant were examined to see which variables were associated with the development of acute graft-versus-host disease (GvHD), which is a binary response variable. The predictor variables are the age of the recipient (Rage), the age of the donor (Dage), whether or not the donor had been pregnant (preg), an index of mixed epidermal cell-lymphocyte reactions (index) and the type of leukemia (here are 3 types, 1= acute myeloid leukemia, 2=acute lymphocytic leukemia and 3=chronic myeloid leukemia).
 - (a) (3 points) I first estimated a full logistic regression model, with the output below

	Estimate	Std. Error	z value	Pr(> z)
(Intercept)	-5.343720	2.624293	-2.036	0.0417
Rage	0.003882	0.084106	0.046	0.9632
Dage	0.112187	0.081436	1.378	0.1683
preg	1.705127	1.199851	1.421	0.1553
log(index)	1.835935	0.892520	2.057	0.0397
as.factor(type)2	-0.405541	1.256857	-0.323	0.7470
as.factor(type)3	1.676694	1.297599	1.292	0.1963

Test whether the overall model is significant at the 5% level. State the null and alternative and your decision. Hint: the 95th percentile of a chi-square distribution with 6 degrees of freedom is 12.59. Answer: $H_0: \beta_1 = r - \beta_2 = 0$ reversed. $H_1: z_1$ least one $\beta_1 \neq 0$. The test statistic is 51.049 - 26.288 = 24.761 > 12.59 so we reject $H_1: z_1 = z_2 = z_3 = z_3$

- so we reject H₀.
 (b) (3 points) Test whether the type of leukemia has an effect, i.e., do we need the two leukemia dummies? A model without the two leukemia dummies has a residual deviance of 29.27. Hint: the 95% percentile of a chi-square distribution with 2 df is 5.992. Answer: H₀: β₀ = β₀ = 0 versus H₁: at least one of β₀ and β₀ is nonzero. The test statistics is 29.27 − 26.288 = 2.984 < 5.992 so we cannot reject H₀.
- (c) (2 points) I created three dummies for the three types of leukemia (they equal I for that type and 0 otherwise): aml, all and cml). I entered all variables into a backward selection logistic regression model giving the following model:

Null deviance: 51.049 on 36 degrees of freedom Residual deviance: 28.848 on 33 degrees of freedom

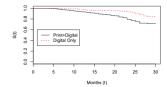
State the estimated regression equation. Answer

$$\log\left(\frac{\pi}{1-\pi}\right) = -2.5465 + 1.4877\log(\text{index}) + 2.2506\text{cml} + 2.4955\text{preg}$$

- (d) (3 points) Use this model to estimate the probability if GvHD for someone with an index of 1.10, acute lymphocytic leukemia, and whose donor had been pregnant. $Answer: ~\ddot{\eta} = -2.5464 + 1.4877* \log(1.1) + 2.4955 = 0.0909~and~\ddot{\pi} = e^{0.0909} = 0.52.$
- (c) (2 points) Using the model from the previous part, how many times greater are the odds (not probability or log odds) of GvHD for someone with chronic myeloid leukemia compared with one of the other two types of leukemia? Answer: e^{2,2006} = 9.49 times more likely.
- 3. (8 points) Complete the following table giving the Kaplan-Meier estimates of the survival function and retention rate

П	Number	Number	Number	Retention	Survival			
t	Cancels	Censored	at Risk	Rate	Function			
1	0	5	1000	1 - 0/1000 = 1	1			
2	5	10	995	1 - 5/995 = .995	.995			
3	7	8	980	1 - 7/980 = .993	.995(.993) = .988			

- 4. (6 points) Multiple logistic regression was used to construct a prognostic index to predict significant coronary artery disease from data on 348 patients with valvular heart disease who had undergone routine coronary arteriography before value replacement. Forward stepwise selection was used, with a significance level for entry into the model of 0.05. The prognostic index obtained was based on a model with seven variables.
 - (a) (2 points) The regression coefficient for a family history of ischaemic heart disease (coded 0=no and 1=yes) was 1.167. What is the estimated "odds ratio" for having significant coronary artery disease associated with a positive family history? Hint: c². Answer: cpt/1.167] = 3.21.
 - c. Assect: csp(1.16) = 3.21.
 (b) (4 points) One of the variables in the model was the estimated total number of cigarettes ever smoked, calculated as the average number smoked annually × the number of years smoking. The regression coefficient was 0.0106 per 1000 cigarettes. What total number of cigarettes ever smoked carries the same risk as a family history of ischaemic heart disease? Convert this figure into years of smoking 20 cigarettes per day. (Assume there are 365 days in a year.) Answer: 1.167/(.0106*20*365/1000) = 15.08 years.
- The plot below is from a KM model of the newspaper data, stratifying by the publication type (print+digital versus digital only). What does the plot tell you?



 $Answer:\ Digital\ only\ survive\ longer\ than\ print+digital.$

- Expect a question interpreting a Poisson regression similar to the news desert homework 7, but not as complicated
- $7. \ \ {\rm Expect\ something\ interpreting\ a\ multinomial\ logic}$