

# Logistic regression

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Machine Learning

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# Generative vs discriminative models

- Goal: predict  $Y$

- Bayes rule:

$$p(Y|\mathbf{X}) = \frac{p(Y) \cdot p(\mathbf{X}|Y)}{p(\mathbf{X})}$$

- Generative classifiers

- Specify prior probability of  $p(Y)$
  - Assume conditional distribution  $p(\mathbf{X}|Y)$
  - Use Bayes rule to derive the posterior  $p(Y|\mathbf{X})$
  - **Example:** Linear discriminant analysis

- Discriminative classifiers

- Estimate the posterior the posterior  $p(Y|\mathbf{X})$
  - Do not assume the distribution on  $\mathbf{X}$
  - **Example:**  $Y$  binary: logistic regression

# Probability Distribution of a Binary Outcome $Y$

- In many situations, the response variable has only two possible outcomes
  - Disease ( $Y = 1$ ) vs Not diseased ( $Y = 0$ )
  - Employed ( $Y = 1$ ) vs Unemployed ( $Y = 0$ )
- Response is *binary or dichotomous*
- Can model response using Bernoulli dist

$Y_i$	Probability
1	$\Pr\{Y_1 = 1\} = \pi_i$
0	$\Pr\{Y_1 = 0\} = 1 - \pi_i$

- $E\{Y_i\} = \pi_i$
- $Var\{Y_i\} = \pi_i(1 - \pi_i)$

# Goal: express $E\{Y\}$ as function of a covariate $X$

- The simple regression is not appropriate

$$E\{Y_i\} = \beta_0 + \beta_1 X_i$$

It violates several assumptions:

(1) Does not enforce the constraint  $0 \leq E\{Y_i\} \leq 1$  is

(2) Non-normal (binary) distribution of  $\varepsilon \mid X$ :

$$\text{When } Y_i = 0 : \varepsilon_i = 0 - \beta_0 - \beta_1 X_i$$

$$\text{When } Y_i = 1 : \varepsilon_i = 1 - \beta_0 - \beta_1 X_i$$

(3) Non-constant variance

$$\begin{aligned} \text{Var}\{Y_i\} &= \pi_i(1 - \pi_i) \\ &= (\beta_0 + \beta_1 X_i)(1 - \beta_0 - \beta_1 X_i) \end{aligned}$$

# Solution: a Generalized Linear Model

- A generalized linear model is

$$E\{Y_i\} = g(\beta_0 + \beta_1 X_i), \text{ or}$$
$$g^{-1}(E\{Y_i\}) = \beta_0 + \beta_1 X_i$$

where  $g$  is a sigmoid function in  $(0,1)$ .

- $g$  is called the *mean response function*
  - $g^{-1}$  is called the *link function*
- 
- A choice of  $g$  produces different models
    - $g(t) = \text{Identity}$   
→ linear regression
    - $g(t) = \Phi(t) = \text{standard Normal CDF}$   
→ probit regression
    - $g(t) = \frac{\exp(t)}{1+\exp(t)} = \text{CDF of the logistic distrib.}$   
→ logistic regression

# Motivation for Probit

## Regression: Latent Variable

- Assume that the binary response is guided by a non-observed continuous variable
- Example: linear model for blood pressure:

$$bp = \beta_0 + \beta_1 \text{age} + \varepsilon$$

Only observe

$$Y = \begin{cases} 1 \text{ (disease),} & \text{if blood pressure} > c \\ 0 \text{ (healthy),} & \text{if blood pressure} \leq c \end{cases}$$

$$\Pr\{Y = 1\}$$

$$= \Pr\{bp > c\} = \Pr\{\beta_0 + \beta_1 \text{age} + \varepsilon > c\}$$

$$= \Pr\{\varepsilon < \beta_0 + \beta_1 \text{age} - c\}$$

$$= \Pr\left\{\frac{\varepsilon}{\sigma} < \frac{\beta_0 - c}{\sigma} + \frac{\beta_1}{\sigma} \text{age}\right\}$$

$$= \Pr\{z < \beta'_0 + \beta'_1 \text{age}\}$$

$$= \Phi(\beta'_0 + \beta'_1 \text{age})$$

# Logistic Response Function and Logistic Regression

- A sigmoidal response function

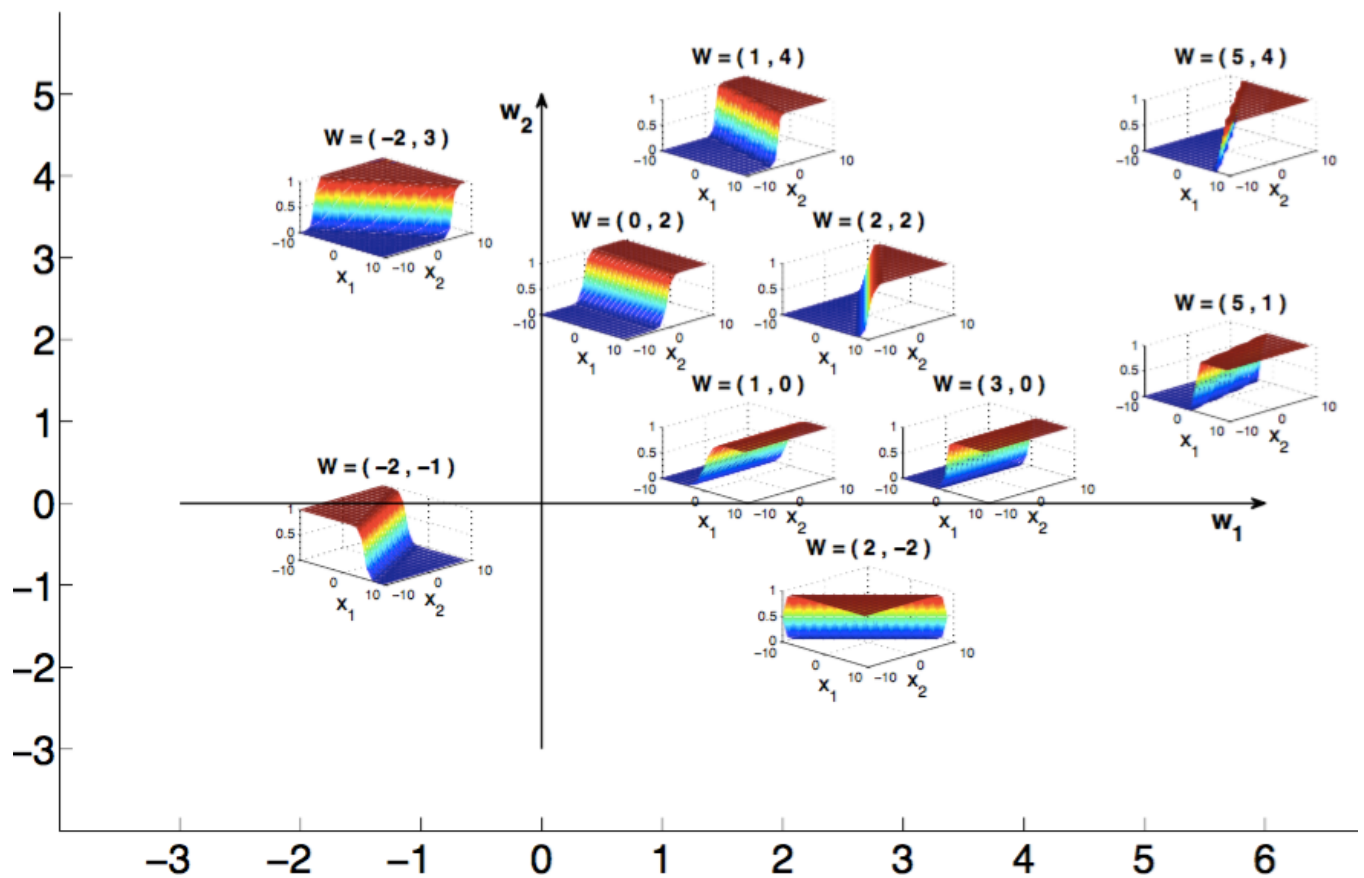
$$\begin{aligned} E\{Y_i\} &= \frac{\exp(\beta_0 + \beta_1 X_i)}{1 + \exp(\beta_0 + \beta_1 X_i)} \\ &= \frac{1}{1 + \exp(-\beta_0 - \beta_1 X_i))} \end{aligned}$$

- A monotonic increasing/decreasing function
- Explicit functional form
- Restricts  $0 \leq E(Y_i) \leq 1$
- Example of a **nonlinear** model

- **Logit** link function

$$\log \left( \frac{E\{Y_i\}}{1 - E\{Y_i\}} \right) = \log \left( \frac{\pi_i}{1 - \pi_i} \right) = \beta_0 + \beta_1 X_i$$

$$\text{Sigm}(\beta_1 x_1 + \beta_2 x_2)$$



K. Murphy, Fig 8.1



# Probability Distribution of $Y$ in Logistic Regression

- $Y_i$  are independent but not identically distributed Bernoulli random variables

$$Y_i \stackrel{ind}{\sim} \text{Bernoulli}(\pi_i) \text{ where}$$
$$\pi_i = \frac{\exp(\beta_0 + \beta_1 X_i)}{1 + \exp(\beta_0 + \beta_1 X_i)}$$

– note no more error term!

- Probability density of  $Y_i$

$$f(Y_i) = \pi_i^{Y_i} (1 - \pi_i)^{1-Y_i}$$

- Least Squares Estimates are inappropriate
  - use maximum likelihood for parameter estimation

# Estimation by Maximum Likelihood

- $Y_i \overset{ind}{\sim} \text{Bernoulli}(\pi_i)$  where

$$\pi_i = \frac{\exp(\beta_0 + \beta_1 X_i)}{1 + \exp(\beta_0 + \beta_1 X_i)}$$

- Log likelihood:  $\log_e(L) =$

$$\begin{aligned} &= \log \left\{ \prod_{i=1}^n \pi_i^{Y_i} (1 - \pi_i)^{1-Y_i} \right\} \\ &= \sum_{i=1}^n Y_i \log(\pi_i) + \sum_{i=1}^n (1 - Y_i) \log(1 - \pi_i) \\ &= \sum_{i=1}^n Y_i \log\left(\frac{\pi_i}{1 - \pi_i}\right) + \sum_{i=1}^n \log(1 - \pi_i) \\ &= \sum_{i=1}^n Y_i(\beta_0 + \beta_1 X_i) - \sum_{i=1}^n \log(1 + \exp(\beta_0 + \beta_1 X_i)) \end{aligned}$$

- MLEs do not have closed forms

# Equivalent specification:

## Binomial distribution

- Change in notation
  - Data:  $(Y_{ij}, n_i, X_i)$ ,  $i = 1, 2, \dots, c$
  - $X_i$  : predictor for observation  $i$
  - $n_i$  : # of Bernoulli trials in observation  $i$
  - $Y'_i := \sum_{j=1}^{n_i} Y_{ij}$
  - Model:

$$Y'_i \stackrel{\text{ind}}{\sim} \text{Binomial}(n_i, \pi_i), \text{ where}$$

$$\pi_i = \frac{\exp(\beta_0 + \beta_1 X_i)}{1 + \exp(\beta_0 + \beta_1 X_i)}$$

- Log-Likelihood:  $\log_e(L) =$

$$= \log \prod_{i=1}^c \left\{ \binom{n_i}{Y'_i} \pi_i^{Y'_i} (1 - \pi_i)^{n_i - Y'_i} \right\}$$

$$= \sum_{i=1}^c \left\{ Y'_i \log(\pi_i) + (n_i - Y'_i) \log(1 - \pi_i) + \log \left( \binom{n_i}{Y'_i} \right) \right\}$$

$$= \sum_{i=1}^c \left\{ Y'_i \log \frac{\pi_i}{1 - \pi_i} + n_i \log(1 - \pi_i) + \log \left( \binom{n_i}{Y'_i} \right) \right\}$$

# Equivalence of Bernoulli and Binomial Models

- Binomial Log-Likelihood equals Bernoulli Log-Likelihood, up to a constant:

$$\begin{aligned}\log_e(L)^{Binomial} &= \\&= \sum_{i=1}^c \{Y'_i \log(\pi_i) + (n_i - Y'_i) \log(1 - \pi_i)\} + constant \\&= \sum_{i=1}^c \left\{ \sum_{j=1}^{n_i} Y_{ij} \log(\pi_i) + (n_i - \sum_{j=1}^{n_i} Y_{ij}) \log(1 - \pi_i) \right\} + constant \\&= \sum_{i=1}^c \sum_{j=1}^{n_i} \left\{ Y_{ij} \log\left(\frac{\pi_i}{1 - \pi_i}\right) + \log(1 - \pi_i) \right\} + constant \\&= \log_e(L)^{Bernoulli} + constant\end{aligned}$$

- Both models lead to same parameter estimates and inferences, but have different deviances.

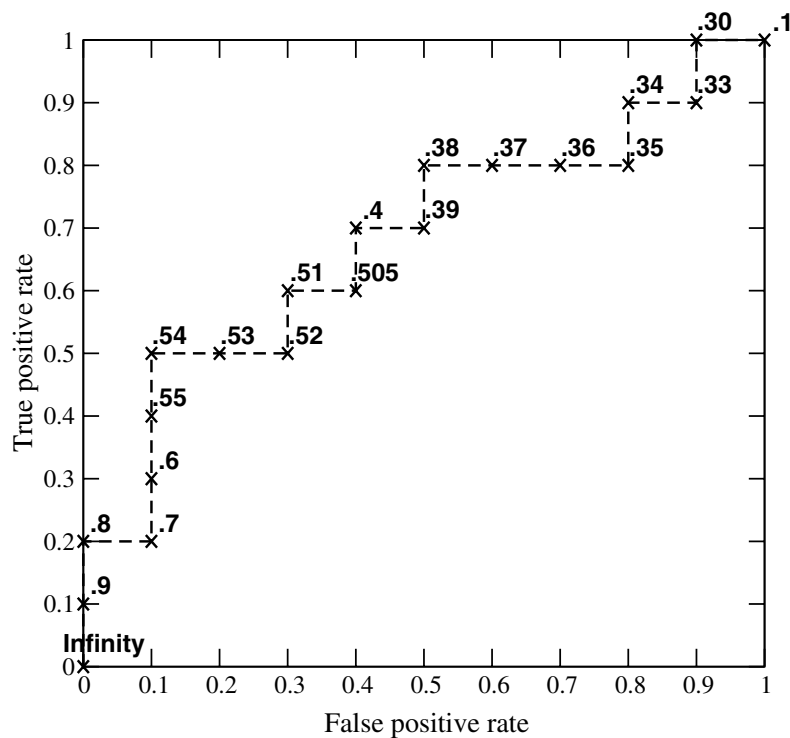
# Summaries of prediction/classification

		<u>True class</u>			
		<b>p</b>	<b>n</b>		
<u>Hypothesized class</u>	<b>Y</b>	True Positives	False Positives	$fp\ rate = \frac{FP}{N}$	$tp\ rate = \frac{TP}{P}$
	<b>N</b>	False Negatives	True Negatives	$precision = \frac{TP}{TP+FP}$	$recall = \frac{TP}{P}$
<b>Column totals:</b>		<b>P</b>	<b>N</b>	$accuracy = \frac{TP+TN}{P+N}$	
				$F\text{-measure} = \frac{2}{1/precision+1/recall}$	

- Results over multiple score cutoffs are summarized in a Receiver Operating Characteristic (ROC) curve
- Vary  $c$ , and for all  $c$  plot sensitivity vs 1-specificity. Evaluate models by area under the curve.
- Area = 1 → perfect classification  
Area = .5 → random classification.

Fawcett, "An introduction to ROC analysis". *Pattern Recognition Letters*, 2005

Inst#	Class	Score	Inst#	Class	Score
1	p	.9	11	p	.4
2	p	.8	12	n	.39
3	n	.7	13	p	.38
4	p	.6	14	n	.37
5	p	.55	15	n	.36
6	p	.54	16	n	.35
7	n	.53	17	p	.34
8	n	.52	18	n	.33
9	p	.51	19	p	.30
10	n	.505	20	n	.1



Fawcett, "An introduction to ROC analysis". *Pattern Recognition Letters*, 2005

# Automatic Variable Selection

- Exhaustive search. Minimize:

$$-2 \log_e L(\mathbf{b})$$

$$AIC_p = -2 \log_e L(\mathbf{b}) + 2p$$

$$BIC_p = -2 \log_e L(\mathbf{b}) + p \log_e(n)$$

- Heuristic search

- forward selection; backward elimination; step-wise selection

- Statistical regularization

- Negative log-likelihood penalized with ridge, lasso, elastic net etc

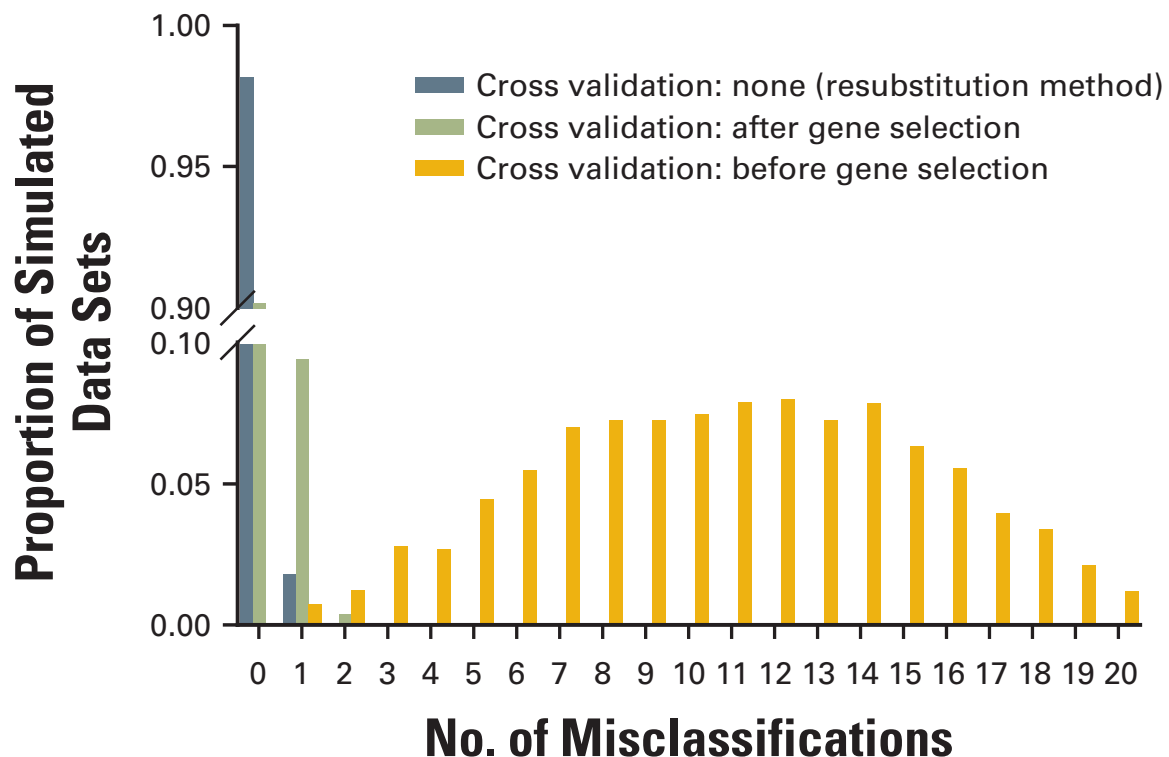
# Variable Selection Should be Done as Part of Cross-Validation

- Example from Simon *et al.*, JNCI, 2003.
- Simulated data with no structure
  - 20 observations with random labels
  - 6,000 possible but unrelated predictors
  - Repeated 200 times
- Estimated predictive accuracy using
  - no cross-validation
  - selecting features on full dataset, then using cross-validation
  - selecting features at each step of cross-validation



# Variable Selection Should be Done as Part of Cross-Validation

Example from Simon *et al.*, JNCI, 2003.



- Conclusion

- Incorporating selection of predictors within the cross-validation procedure is key

# More than 2 groups: conditional multinomial distributions

- $Y|X = x \sim \text{Multinomial}(n_{i+}, \pi_1(x), \dots, \pi_J(x))$
- $X$  - predictor;  
 $Y$  - multinomial response with  $J$  categories

Row	Column			Total
	1	$\dots$	$J$	
1	$\pi_{11}$ $(\pi_{1 1})$	$\dots$	$\pi_{1J}$ $(\pi_{J 1})$	$\pi_{1+}$
$\vdots$	$\vdots$	$\vdots$	$\vdots$	$\vdots$
$I$	$\pi_{I1}$ $(\pi_{1 I})$	$\dots$	$\pi_{IJ}$ $(\pi_{J I})$	$\pi_{I+}$
Total	$\pi_{+1}$	$\dots$	$\pi_{+J}$	$\pi_{++}$

- Consider the new notation:  
 $\pi_j(X) = P(Y = j|X = x), \sum_{j=1}^J \pi_j = 1$
- If  $Y$  is ordered, we can also be interested in cumulative probabilities:  
 $P_j(X) = P(Y \leq j|X = x)$

# Baseline-Category Logistic Regression

- *Data:*

$$(y_{ij}, x_i); \quad y_{ij} = I_{\{y_i=j\}}; \quad \sum_{j=1}^J y_{ij} = 1; \\ j = 1, \dots, J, \quad i = 1, \dots, n$$

- The model describes the effects of covariates  $X$  on the  $J - 1$  logits.

$$\log \frac{\pi_j(X)}{\pi_1(X)} = \alpha_j + \beta_j X, \quad j = 2, \dots, J$$

- An arbitrary category ( $j = 1$  or  $j = J$ ) is chosen as the baseline category
- Each other category  $j$  is paired with the baseline to build a logistic model
- Separate set of parameters  $\beta_j$  for each  $\pi_j$
- Separate linear relationship between  $X$  and  $\log \frac{\pi_j(X)}{\pi_1(X)}$
- Values of  $\beta_j$  depend on the baseline

# Predicted Probability

- Since  $\pi_j(X) = \pi_1(X)e^{\alpha_j + \beta_j X}$  and  $\sum_{j=1}^J \pi_j(X) = 1$

$$\left\{ \begin{array}{l} \pi_1(X) = \frac{1}{1 + \sum_{k=2}^J \exp(\alpha_k + \beta_k X)}, \quad j = 1 \\ \pi_j(X) = \frac{\exp(\alpha_j + \beta_j X)}{1 + \sum_{k=2}^J \exp(\alpha_k + \beta_k X)}, \quad j = 2, 3, \dots, J \end{array} \right.$$

- Same denominator for all  $j$
- For  $J = 2$  - an ordinary logistic regression
- Can be viewed as a classification model
  - the observation is assigned to the category with the highest predicted probability
  - can plot ROC curves for a particular category versus all other categories combined

# Max. Likelihood Estimation

Since

$$\pi_1(x_i) = [1 + \sum_{j=2}^J \exp(\alpha_j + \beta_j x_i)]^{-1} \text{ and } y_{i1} = (1 - \sum_{j=2}^J y_{ij}),$$

contribution of  $(y_{ij}, x_i)$  to the log-likelihood is:

$$\begin{aligned} l_i &= \log \left[ \prod_{j=1}^J \pi_j(x_i)^{y_{ij}} \right] \\ &= \sum_{j=2}^J y_{ij} \log \pi_j(x_i) + \left( 1 - \sum_{j=2}^J y_{ij} \right) \log \pi_1(x_i) \\ &= \sum_{j=2}^J y_{ij} \log \frac{\pi_j(x_i)}{1 - \sum_{k=2}^J \pi_k(x_i)} + \log \pi_1(x_i) \\ &= \sum_{j=2}^J y_{ij} (\alpha_j + \beta_j x_i) - \log \left[ 1 + \sum_{j=2}^J \exp(\alpha_j + \beta_j x_i) \right] \end{aligned}$$

- Maximize  $\sum_{i=1}^I l_i$  with respect to  $\alpha_j$  and  $\beta_j$

# Multinomial Vs Binary Logistic Regression

- $\log \frac{\pi_j(X)}{\pi_1(X)} = \alpha_j + \beta_j X, \quad j = 2, \dots, J$
- Can we fit separate  $J-1$  logistic regressions for  $J-1$  response categories vs baseline?
  - Same model in principle
- Separate-fitting ML parameter estimates
  - Differ from the joint-fitting ML estimates
  - Tend to have larger standard errors
  - Loss of efficiency is minor when the baseline is the most common category

# Example: Dose Response

```
#-----read the data-----
x <- data.frame(
  count = c(59, 25, 46, 48, 32, 48, 21, 44, 47, 30, 44,
            14, 54, 64, 31, 43, 4, 49, 58, 41),
  dose = c(rep(0, 5), rep(1, 5), rep(2, 5), rep(3, 5)),
  response = as.factor(rep(c(0:4), 4))
)
m <- matrix(x$count, byrow=TRUE, ncol=5,
  dimnames=(list(0:3, 0:4)))
> m
  0  1  2  3  4
0 59 25 46 48 32
1 48 21 44 47 30
2 44 14 54 64 31
3 43  4 49 58 41
```

- Row='dose'; column='response'
- View 'dose' as a categorical predictor
  - introduce 3 indicators for predictors
- View 'dose' as a continuous predictor
  - assign scores to categories on an arbitrary scale
- More R examples: Venable & Ripley p. 203

# Example: Dose Response

## 'Dose' Viewed as Categorical

```
library(nnet)
fit1 <- multinom(response ~ as.factor(dose), weights=count,
  data=x)
```

```
> summary(fit1)
```

Coefficients:

	(Intercept)	...(dose)1	...(dose)2	...(dose)3
1	-0.8586335	0.03194971	-0.2864809	-1.5161958
2	-0.2488754	0.16185828	0.4536705	0.3794879
3	-0.2063195	0.18526707	0.5810140	0.5055581
4	-0.6117850	0.14178037	0.2615807	0.5641507

Std. Errors:

	(Intercept)	...(dose)1	...(dose)2	...(dose)3
1	0.2386396	0.3541205	0.3887204	0.5746170
2	0.1966936	0.2867909	0.2827264	0.2869711
3	0.1943777	0.2826526	0.2759257	0.2797853
4	0.2195434	0.3199468	0.3212239	0.3095891

Residual Deviance: 2443.166

AIC: 2475.166



# Plot Predicted Probabilities

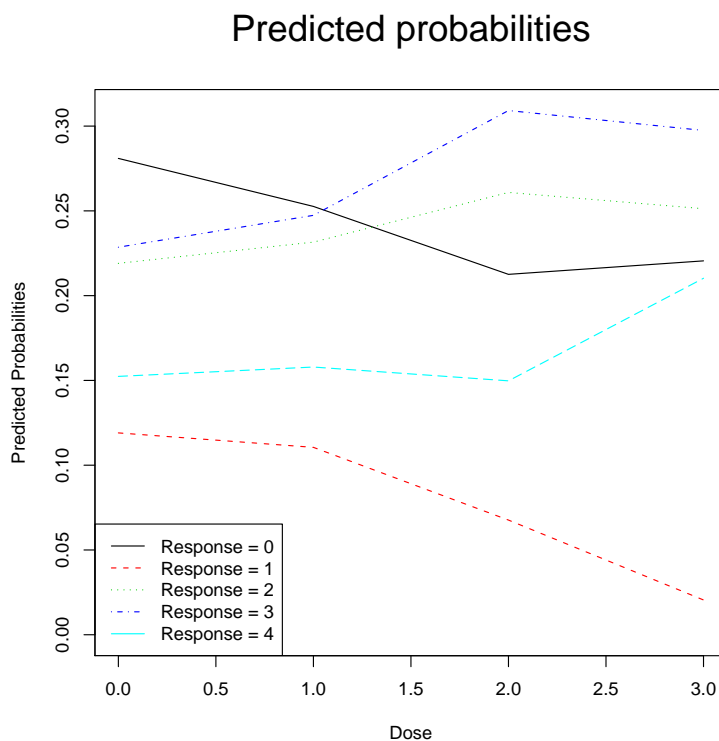
```
predProb <- unique(fit1$fitted.values)
```

```
> predProb
```

	0	1	2	3	4
0.2809484	0.11904927	0.2190490	0.2285720	0.1523813	
0.2526320	0.11052594	0.2315780	0.2473691	0.1578949	
0.2125601	0.06763396	0.2608694	0.3091786	0.1497580	
0.2205135	0.02051445	0.2512809	0.2974355	0.2102557	

```
matplot(predProb)
```

```
legend("bottomleft", lty=c(1:4), col=c(1:5),  
      paste("Response =", c(0:4)))
```



# Example: Dose Response 'Dose' Viewed as Continuous

```
library(nnet)
fit2 <- multinom(response ~ dose, weights=count, data=x)
> summary(fit2)
Coefficients:
      (Intercept)      dose
1  -0.6999134 -0.3544346
2  -0.2194566  0.1470232
3  -0.1772963  0.1945578
4  -0.6544057  0.1914772
...
Residual Deviance: 2449.145
AIC: 2465.145
```

