# Data-analysis and Retrieval Ordinal Classification

#### Ad Feelders

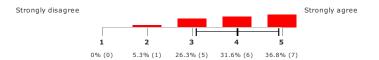
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# **Ordinal Classification**

- When a variable is ordinal, its categories can be ranked from low to high, but the distances between adjacent categories are unknown.
- In ordinal classification the class variable is ordinal.

#### Example: Likert scale





#### Logistic Regression Revisited

Consider the linear regression model

$$y^* = \beta^\top \mathbf{x} + \varepsilon, \qquad \qquad E[\varepsilon \mid \mathbf{x}] = 0$$

where  $y^*$  is an unobserved (latent) numeric variable. We only observe whether  $y^*$  is bigger than a given threshold:

$$y = \begin{cases} 1 & \text{if } y^* > 0 \\ 0 & \text{if } y^* \le 0 \end{cases}$$

Note the vector notation:  $\mathbf{x} = (1, x_1, \dots, x_p)^{\top}$  and  $\beta = (\beta_0, \beta_1, \dots, \beta_p)^{\top}$ , so

$$\beta^{\top}\mathbf{x} = \beta_0 + \sum_{j=1}^{p} \beta_j x_j$$

#### Logistic Regression Revisited

According to this model, the probability that y = 1 is

$$P(y = 1) = P(y^* > 0)$$
  
=  $P(\beta^{\top} \mathbf{x} + \varepsilon > 0)$   
=  $P(\varepsilon > -\beta^{\top} \mathbf{x})$ 

If the distribution of  $\varepsilon$  is symmetric around zero, then  $P(\varepsilon > a) = P(\varepsilon < -a)$ , so

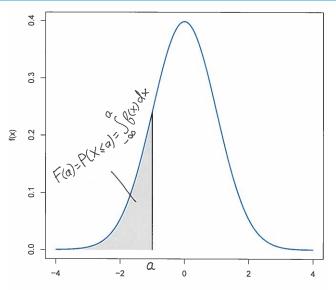
$$P(\varepsilon > -\beta^{\top} \mathbf{x}) = P(\varepsilon < \beta^{\top} \mathbf{x}) \equiv F(\beta^{\top} \mathbf{x})$$

Here *F* is the cumulative density function (cdf) of  $\varepsilon$ . The cdf is defined as

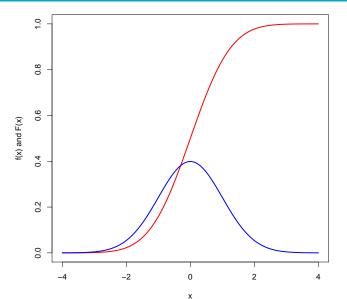
$$F(z) = P(Z \le z) = \int_{-\infty}^{z} f(Z) dZ$$

where f is the probability density function (pdf) of Z.

## Density and cumulative density



## Standard Normal density and cumulative density function



We have established that (under certain assumptions):

$$P(y=1) = F(\beta^{\top}\mathbf{x})$$

Depending on the choice of F (or f) we get different models.

• If we choose  $\varepsilon \sim N(0,1)$ , then we get the so-called probit model:

$$P(y=1) = \Phi(eta^ op \mathbf{x})$$

where  $\Phi(\cdot)$  denotes the standard normal cumulative density function.

• The assumption of unit variance is a harmless normalization.

Suppose we assume instead that  $\varepsilon \sim N(0, \sigma^2)$ , as is common in linear regression. First of all, note that

$$P(y = 1 \mid \mathbf{x}) = P(\varepsilon < \beta^{\top}\mathbf{x}) = P\left(rac{\varepsilon}{\sigma} < rac{eta^{\top}\mathbf{x}}{\sigma}
ight)$$

Define  $u = \frac{\varepsilon}{\sigma}$ . Then  $u \sim N(0, 1)$ . Furthermore, let  $\alpha_j = \frac{\beta_j}{\sigma}$ .

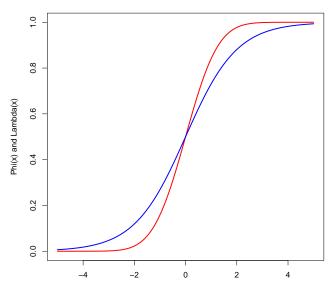
The model with coefficients  $\alpha_j$  and error term u is "observationally equivalent" to the model with coefficients  $\beta_j$  and error term  $\varepsilon$ . They are "observationally equivalent" because they produce the exact same probabilities for the different Y values. Since Y is all we observe (not  $Y^*$ ), the two models cannot be distinguished from each other on the basis of observations. For the logit (logistic regression) model

$$P(y=1) = \Lambda(eta^{ op} \mathbf{x}) = rac{e^{eta^{ op} \mathbf{x}}}{1+e^{eta^{ op} \mathbf{x}}}$$

where  $\Lambda(\cdot)$  denotes the logistic cumulative density function.

Note that this is the logistic response function we have already seen in one of the previous lectures.

# Normal (red) and logistic (blue) cumulative density



Instead of fixing the threshold at zero, we can also remove the intercept  $\beta_0$  from the model and make the threshold an unknown parameter. Then we get the model:

$$y^* = \sum_{j=1}^p \beta_j x_j + \varepsilon, \qquad \qquad E[\varepsilon \mid \mathbf{x}] = 0$$

where  $y^*$  is still an unobserved (latent) numeric variable. We only observe whether  $y^*$  is bigger than a threshold t:

$$y = \begin{cases} 1 & \text{if } y^* > t \\ 0 & \text{if } y^* \le t \end{cases}$$

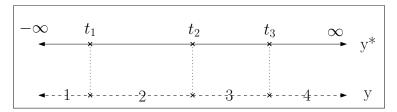
Let *m* denote the number of classes, where the classes are labeled  $\{1, 2, ..., m\}$ . Then *y* is defined as follows:

$$y = \begin{cases} 1 & \text{if } -\infty < y^* \le t_1 \\ 2 & \text{if } t_1 < y^* \le t_2 \\ \vdots & \vdots \\ m & \text{if } t_{m-1} < y^* < \infty \end{cases}$$

We only observe between which thresholds  $y^*$  falls.

Here  $t_1, \ldots, t_{m-1}$  are unknown thresholds that have to be estimated from the data (together with the coefficient vector  $\beta$ ).

We only observe y, which indicates the interval  $y^*$  falls into.



We observe y = 1 when  $y^*$  falls between  $t_0 = -\infty$  and  $t_1$ . Hence

$$P(y_i = 1 \mid \mathbf{x}_i) = P(t_0 \leq y_i^* < t_1 \mid \mathbf{x}_i)$$

Substituting  $y_i^* = \beta^\top \mathbf{x}_i + \varepsilon_i$ , (suppressing condition on  $\mathbf{x}_i$ ) we get

$$P(y_i = 1) = P(t_0 \leq \beta^\top \mathbf{x}_i + \varepsilon_i < t_1)$$

Now we subtract  $\beta^{\top} \mathbf{x}_i$  from all terms in the inequality to get

$$P(y_i = 1) = P(t_0 - \beta^\top \mathbf{x}_i \le \varepsilon_i < t_1 - \beta^\top \mathbf{x}_i)$$

Continuing from the previous slide:

$$P(y_i = 1) = P(t_0 - \beta^\top \mathbf{x}_i \le \varepsilon_i < t_1 - \beta^\top \mathbf{x}_i)$$
  
=  $P(\varepsilon_i < t_1 - \beta^\top \mathbf{x}_i) - P(\varepsilon_i < t_0 - \beta^\top \mathbf{x}_i)$   
=  $F(t_1 - \beta^\top \mathbf{x}_i) - F(t_0 - \beta^\top \mathbf{x}_i),$ 

because  $P(a \le Z \le b) = P(Z \le b) - P(Z \le a) = F(b) - F(a)$ .

This derivation can be generalized to compute the probability of any observed outcome  $y_i = j$  given  $\mathbf{x}_i$ :

$$P(y_i = j \mid \mathbf{x}_i) = F(t_j - \beta^\top \mathbf{x}_i) - F(t_{j-1} - \beta^\top \mathbf{x}_i), \qquad j = 1, \dots, m.$$

So for a model with four possible classes, the formula's for the different outcomes are:

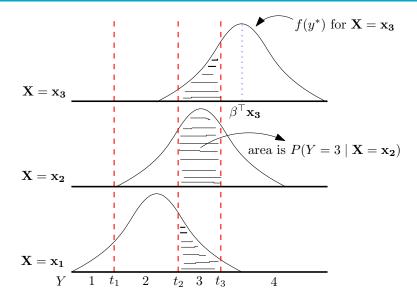
$$P(y_i = 1 | \mathbf{x}_i) = F(t_1 - \beta^\top \mathbf{x}_i)$$

$$P(y_i = 2 | \mathbf{x}_i) = F(t_2 - \beta^\top \mathbf{x}_i) - F(t_1 - \beta^\top \mathbf{x}_i)$$

$$P(y_i = 3 | \mathbf{x}_i) = F(t_3 - \beta^\top \mathbf{x}_i) - F(t_2 - \beta^\top \mathbf{x}_i)$$

$$P(y_i = 4 | \mathbf{x}_i) = 1 - F(t_3 - \beta^\top \mathbf{x}_i)$$

#### **Class Probabilities**



Verbal description:

- Depending on the value of  $\mathbf{x}$ , the distribution of  $y^*$  is shifted.
- The expected value of  $y^*$  is  $\beta^{\top} \mathbf{x}$ .
- The class probabilities are defined by the area under  $f(y^*)$  between the different thresholds.
- In this way, the class probabilities depend on the value of x.

The likelihood function is

$$L(\beta, t ; \mathbf{X}, \mathbf{y}) = \prod_{j=1}^{m} \prod_{i:y_i=j} P(y_i = j | \mathbf{x}_i, \beta, t)$$
$$= \prod_{j=1}^{m} \prod_{i:y_i=j} \left[ F(t_j - \beta^{\top} \mathbf{x}_i) - F(t_{j-1} - \beta^{\top} \mathbf{x}_i) \right],$$

where  $\prod_{i:y_i=j}$  indicates we multiply over all cases where y is observed to have value j.

Taking logs, the log likelihood is equal to

$$\log L(\beta, t ; \mathbf{X}, \mathbf{y}) = \sum_{j=1}^{m} \sum_{i: y_i = j} \log \left[ F(t_j - \beta^\top \mathbf{x}_i) - F(t_{j-1} - \beta^\top \mathbf{x}_i) \right].$$

This expression can be maximized with numerical methods to estimate the thresholds  $t_i$  and vector of coefficients  $\beta$ .

The likelihood function is

$$L(\beta, t ; \mathbf{X}, \mathbf{y}) = \prod_{j=1}^{m} \prod_{i: y_i = j} P(y_i = j \mid \mathbf{x}_i, \beta, t)$$

Note that the likelihood score of a model (choice of  $t, \beta$ ) only depends on the probability that it assigns to the correct class.

Homework: Can you think of an argument against using MLE in *ordinal* classification?

Also, note that:

$$P(y_i \leq 1 \mid \mathbf{x}_i) = F(t_1 - \beta^\top \mathbf{x}_i)$$
$$P(y_i \leq 2 \mid \mathbf{x}_i) = F(t_2 - \beta^\top \mathbf{x}_i)$$
$$P(y_i \leq 3 \mid \mathbf{x}_i) = F(t_3 - \beta^\top \mathbf{x}_i)$$
$$P(y_i \leq 4 \mid \mathbf{x}_i) = 1$$

In general we have  $P(y_i \leq j | \mathbf{x}_i) = F(t_j - \beta^\top \mathbf{x}_i)$ .

We have seen that:

$$P(y \leq j \mid \mathbf{x}) = F(t_j - \beta^{\top} \mathbf{x}).$$

In logistic regression we choose for F the logistic cdf

$$\Lambda(z) = \frac{\exp(z)}{1 + \exp(z)},$$

so we get

$$P(y \leq j \mid \mathbf{x}) = rac{\exp(t_j - eta^{ op} \mathbf{x})}{1 + \exp(t_j - eta^{ op} \mathbf{x})},$$

Set of *parallel* logistic regression models for  $y \le j$  against y > j:

$$\log\left[\frac{P(y \le j \mid \mathbf{x})}{P(y > j \mid \mathbf{x})}\right] = t_j - \beta^{\top} \mathbf{x}$$

#### Interpretation: effect of increase in $x_k$

We have

$$P(y \leq j \mid \mathbf{x}) = F(t_j - \beta^\top \mathbf{x}).$$

Hence

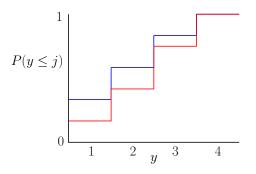
$$\frac{\partial P(\mathbf{y} \leq j \mid \mathbf{x})}{\partial x_k} = \frac{\partial F(t_j - \beta^\top \mathbf{x})}{\partial x_k} = \frac{\partial F(z)}{\partial z} \frac{\partial z}{\partial x_k}$$
$$= f(z) \times -\beta_k = -\beta_k f(t_j - \beta^\top \mathbf{x}).$$

 $f(t_j - \beta^\top \mathbf{x})$  is always positive, since f is a probability density function. So if  $\beta_k$  is positive, an increase in  $x_k$  will lead to a decrease in  $P(y \le j)$  for all j = 1, ..., m - 1.

Or (same thing), an increase in  $x_k$  will lead to an increase in  $P(y \ge j)$  for all j = 2, ..., m.

In this specific sense, one can say that if  $x_k$  increases, higher values of y become more likely.

## Interpretation for $\beta_k$ positive



Blue:  $P(y \le j)$  for  $x_k$ . Red:  $P(y \le j)$  for  $x_k + 1$ .

The cumulative distribution of y for  $x_k + 1$  is *entirely below* the cumulative distribution of y for  $x_k$ .

How exactly are the ordered and unordered (= multinomial) logistic regression model different?

- The ordinal model has a single coefficient vector β for all classes, whereas the multinomial model has a coefficient vector β<sub>k</sub> for each class k (except one).
- As a consequence the decision boundaries are restricted to be parallel to each other in the ordinal model. This is quite a strong constraint!
- In the ordinal model the relation between predictor and class label is monotone, either increasing or decreasing.
- For example: if β<sub>k</sub> is positive, then (all else equal) an increase in x<sub>k</sub> makes the higher classes more likely and a decrease in x<sub>k</sub> makes the lower classes more likely.

#### Fitting the Ordinal Logistic Regression Model in R

```
> library(MASS)
```

- # fit proportional odds logistic regression model
- > wine.polr2 <- polr(quality~density+alcohol,data=wine.dat, Hess=T)</pre>
- > summary(wine.polr2)

#### Coefficients:

	Value	Std.	Error	t value
density	106.27	0	.30531	348.08
alcohol	1.11	0	.05267	21.07

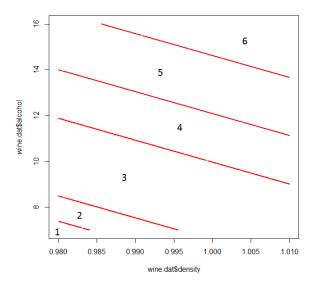
#### Intercepts:

	Value	Std. Error	t value	
1 2	111.9811	0.3032	369.3438	
2 3	113.8616	0.3057	372.4595	
3 4	117.2274	0.3253	360.3388	
4 5	119.7431	0.3667	326.5169	
5 6	122.6179	0.4476	273.9202	

#### **Prediction Accuracy**

```
# predict class labels
> wine.pred <- predict(wine.polr,wine.dat,type="class")</pre>
# construct confusion matrix
> wine.confmat <- table(wine.dat[,12],wine.pred)</pre>
> wine confmat
  wine.pred
     1
        2 3 4 5 6
     0 0 9 1 0 0
 1
 2 0 0 36 17 0 0
 3 0 2 503 173 2 1
 4 0 0 213 392 33 0
 5 0 0 7 138 54 0
 6 0 0 0 10 8 0
# compute accuracy
> sum(diag(wine.confmat))/1599
[1] 0.5934959
> summary(wine.dat[,12])
 1 2 3 4 5 6
 10 53 681 638 199 18
> 681/1599
[1] 0.4258912
```

# Decision Boundary Ordinal LR on Wine Data



#### Fitting the Multinomial Logistic Regression Model

```
> library(nnet)
```

```
# fit multinomial logistic regression model
```

```
> wine.multi2 <- multinom(quality~density+alcohol,data=wine.dat)</pre>
```

```
> summary(wine.multi2)
```

```
Coefficients:
```

	(Intercept)	density	alcohol	
2	43.003814	-45.879145	0.4365809	
3	68.378492	-62.949698	-0.1396656	
4	2.385932	-7.137190	0.8667334	
5	-108.347472	93.833526	1.6793355	
6	-17.906887	-4.108423	2.0746746	

Std. Errors:

	(Intercept)	density	alcohol	
2	2.260309	2.274173	0.4522562	
3	2.137517	2.146860	0.4289871	
4	2.139384	2.141155	0.4282659	
5	2.166738	2.182500	0.4334476	
6	2.507561	2.531655	0.4810792	

### Prediction Accuracy with Multinomial Logit

```
# predict class labels
```

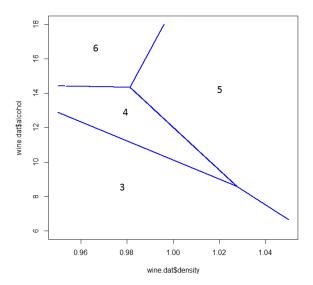
- > wine.pred.m <- predict(wine.multi2,wine.dat,type="class")</pre>
- # construct confusion matrix
- > wine.confmat.m <- table(wine.dat[,12],wine.pred.m)</pre>
- > wine.confmat.m

wine.pred.m

	1	2	3	4	5	6
1	0	0	7	3	0	0
2	0	0	30	22	1	0
3	0	0	514	161	6	0
4	0	0	267	347	24	0
5	0	0	23	159	17	0
6	0	0	2	10	6	0

# compute accuracy of predictions
> sum(diag(wine.confmat.m))/1599
[1] 0.5490932

# Decision Boundary Multinomial LR on Wine Data



#### Comparison of Ordinal and Multinomial Model

- > wine.polr.corr <- as.numeric(wine.pred==wine.dat[,12])</pre>
- > wine.multi.corr <- as.numeric(wine.pred.m==wine.dat[,12])</pre>

# is the difference in error rate (= 1-accuracy) significant? Null hypothesis:

$$H_0: e_{\text{polr}} = e_{\text{multi}}, \qquad H_a: e_{\text{polr}} \neq e_{\text{multi}}$$

If the null hypothesis is correct then  $P(\text{cell } (1,0)) = P(\text{cell } (0,1)) = \frac{1}{2}$  (the other two cells are ignored).

#### Comparison of Ordinal and Multinomial Model

Hence the p-value is

 $P(X \le 104) + P(X \ge 175)$ , where  $X \sim \text{Binom}(\pi = \frac{1}{2}, n = 279)$ 

In R we can compute this as

```
> 2*pbinom(104,175+104,prob=0.5)
[1] 2.531092e-05
```

```
# yes, the p-value is smaller than 0.01, which is
# already a very strict significance level
```

The p-value is very small, so we conclude that the ordinal model has "significantly higher" accuracy than the multinomial model.

# Exploiting dependence of predictions

If predictions were independent, we would have gotten the following table:

> 2\*pbinom(357,357+428,prob=0.5)
[1] 0.0124263

Now the p-value is much higher for the same difference in accuracy!