Logistic, multinomial, and ordinal regression

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Introduction

Testing a risk factor: we want to check whether a certain factor adds to the probability of outbreak of a disease.

This corresponds to the following contingency table:

risk factor	disease	sum	
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exposed	<i>n</i> ₁₁	n ₁₂	<i>n</i> ₁₀
unexposed	<i>n</i> ₂₁	n ₂₂	<i>n</i> ₂₀
sum	<i>n</i> ₀₁	n ₀₂	n

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sum	<i>n</i> ₀₁	n ₀₂	n

Odds of the outbreak for both groups are:

$$o_1 = \frac{n_{11}}{n_{10} - n_{11}} = \frac{\frac{n_{11}}{n_{10}}}{1 - \frac{n_{11}}{n_{10}}} = \frac{\hat{p}_1}{1 - \hat{p}_1}, \quad o_2 = \frac{\hat{p}_2}{1 - \hat{p}_2}$$

As a measure of the risk, we can form odds ratio:

$$\mathbf{OR} = \frac{o_1}{o_2} = \frac{\hat{p}_1}{1 - \hat{p}_1} \cdot \frac{1 - \hat{p}_2}{\hat{p}_2}$$
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Then, it holds

$$\log\left(o_{1}\right)=\log\left(o_{2}\right)+\log\left(\mathbf{OR}\right),$$

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Then, it holds

$$\log\left(o_{1}\right) = \log\left(o_{2}\right) + \log\left(\mathbf{OR}\right),$$

or

$$y = \log(o_2) + \log(\mathbf{OR}) \cdot x,$$

where $x \in \{0, 1\}$ and $y \in \{\log(o_1), \log(o_2)\}$.

This is the basic logistic model. Formally it is a regression model

$$y = \beta_0 + \beta_1 x$$

with baseline $\beta_0 = \log(o_2)$ and slope $\beta_1 = \log(\mathbf{OR})$ – effect of the exposure.

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If we denote the probability of the event (outbreak of the disease)
 by p, then

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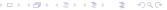
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It follows

$$\rho = \frac{\exp(\beta_0 + \beta_1 x)}{1 + \exp(\beta_0 + \beta_1 x)} = \frac{1}{1 + \exp(-\beta_0 - \beta_1 x)}$$



• Estimation is done via ML-method. Likelihood of observed frequencies for given β 's is

$$L(\beta) = \binom{n_{10}}{n_{11}} p_1^{n_{11}} (1 - p_1)^{n_{10} - n_{11}} \binom{n_{20}}{n_{21}} p_2^{n_{21}} (1 - p_2)^{n_{20} - n_{21}}$$

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Then

$$\ell(\beta) = \log L(\beta) = n_{11} \log (p_1) + n_{12} \log (1 - p_1) + + n_{21} \log (p_2) + n_{22} \log (1 - p_2)$$

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It is easy to derive that ML-equations are

$$\frac{\partial \ell}{\partial \beta_1} = n_{11} - n_{10}p_1 = 0, \quad \frac{\partial \ell}{\partial \beta_0} = n_{11} - n_{10}p_1 + n_{21} - n_{20}p_2 = 0$$



• In this case we already know the solution, $\hat{p}_1 = \frac{n_{11}}{n_{10}}$ and $\hat{p}_2 = \frac{n_{21}}{n_{20}}$ (and corresponding β 's).

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- Asymptotic variance matrix of $\hat{\beta}$'s can also be established. In this case it is

$$\operatorname{var}\begin{pmatrix} \hat{\beta}_0 \\ \hat{\beta}_1 \end{pmatrix} = J^{-1} = \frac{1}{ab}\begin{pmatrix} a & -a \\ -a & a+b \end{pmatrix},$$

where
$$a = n_{10}\hat{p}_1 (1 - \hat{p}_1), b = n_{20}\hat{p}_2 (1 - \hat{p}_2).$$

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, $b = n_{20}\hat{p}_2 (1 - \hat{p}_2)$.

• $J = \begin{pmatrix} a+b & a \\ a & a \end{pmatrix}$ is the information matrix.

 Example: Baystate Medical Center in Springfield, MA, studied factors influencing low birth weights of babies. Let us take as a risk factor smoking during pregnancy. We get the following contingency table:

smoking	low birt	sum	
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exposed	30	44	74
unexposed	29	86	115
sum	59	130	159

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• Odds for the unexposed is $o_1 = 30/44 = 0.681818$, for the exposed $o_2 = 29/86 = 0.337209$, **OR** = 2.021644. Sakoda coefficient S = 0.225344 indicates moderate association, but statistically significant ($\chi^2 = 4.92 > 3.84 = \chi_1^2(0.05)$).

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Software gets

	coefficient	std. error	Z	P > z	95% conf. interval	
smoking	0.7040592	0.319639	2.20	0.028	0.077579	1.330539
constant	-1.087051	0.21473	-5.06	0.000	-1.50791	-0.66619

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• We can also get 95% CI for the **OR**:

$$\left[e^{0.077579};e^{1.330539}\right] = \left[1.08;\ 3.78\right]$$



General categorical predictor:

There are more probabilities to estimate. Let the predictor have m categories:

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Then,

$$y = \beta_0 + \beta_1 x_1 + \cdots + \beta_{m-1} x_{m-1}$$
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where all $x_i \in \{0, 1\}$, but only one of them can be 1 at a time. One category out of m has to be a reference category. One explanatory variable is replaced by m1 indicator variables of other categories, which are mutually exclusive. Significance of a regression coefficient indicates significant difference between corresponding category and the reference category (differential effect of the category).

Example: Let us take mother's weight (grouped in 3 categories) as a risk factor of child's low birth weight. We get

weight group	low birth weight		sum	row percentages	
weight group	yes	no	Sum	10W per	Jernages
≤ 110	25	28	53	47.2%	52.8%
(110; 150]	27	73	100	27.0%	73.0%
> 150	7	29	36	19.4%	80.6%
sum	59	130	189	31.2%	68.8%

Changing row percentages show that mother's weight can be a risk factor.

Software output for the 3rd category as the reference:

variable	coefficient	std. error	Wald	df	p-value
wt_groups			9.073891	2	0.010706
wt_groups(1)	1.308056995	0.503044916	6.761449	1	0.009315
wt_groups(2)	0.426763105	0.477572579	0.798537	1	0.371531
constant	-1.42138568	0.421117444	11.39246	1	0.000737

Software output for the 1st category as the reference:

variable	coefficient	std. error	Wald	df	p-value
wt_groups			9.073891	2	0.010706
wt_groups(2)	-0.88129389	0.355598021	6.142184	1	0.013199
wt_groups(3)	-1.308057	0.503044916	6.761449	1	0.009315
constant	-0.11332869	0.27516229	0.16963	1	0.680441

Symbolically, for the impact of weight groups holds 1 \neq {2,3} at 5% level. Weight as a factor also is significant.



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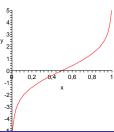
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Logistic transformation $p \to \log\left(\frac{p}{1-p}\right)$ is from (0; 1) to $(-\infty; +\infty)$, so that it removes restrictions harming regression.



Example: Let us take mother's weight as a continuous risk factor of child's low birth weight.

Software gets:

variable	coefficient	std. error	Wald	df	p-value
lwt	-0.01405826	0.006169588	5.192193	1	0.022689
constant	0.998314313	0.78529092	1.616119	1	0.203634

Mother's weight is again significant.

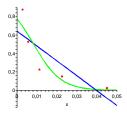
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Example: Effect of anti-pneumococcus serum on survival of ill mice was studied. Five different doses were administered to five groups of 40 mice. Plot shows the death rates, simple linear regression line, and logistic regression curve.



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or (taking $X_0 = 1$)

$$p_i = \frac{\exp\left(\sum_{j=0}^k \beta_j x_{ij}\right)}{1 + \exp\left(\sum_{j=0}^k \beta_j x_{ij}\right)} = \frac{1}{1 + \exp\left(-\sum_{j=0}^k \beta_j x_{ij}\right)}$$

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The ML-equations are

$$\frac{\partial \ell}{\partial \beta_j} = \sum_{i=1}^n (y_i - p_i(\beta)) x_{ij} = 0 \ \forall j$$

 In general, these equations are solved numerically (Newton-Raphson type algorithm).

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- Pros & cons:
 - no closed form formulas, iterative estimation
 - + approximate variances, p-values and CI available

• Wald test of a coefficient: If H_0 : $\beta_i = 0$ holds, then

$$Z = \frac{\hat{\beta}_i}{s_{\hat{\beta}_i}}$$

has asymptotically N(0; 1). There is alternative chi-square form (Z^2) .

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- Likelihood ratio test: Notions:
 - estimated model model with predictors
 - empty model model without predictors, just with intercept
 - full model model predicting $n_{i0}\hat{p}_i = n_{i1} \ \forall i, \text{t.j.} \ \hat{y}_i = y_i \ \forall i$

Deviance:

$$\hat{\ell}_m = \log \hat{L}_m = \log \hat{L} ext{(estimated model)},$$
 $\hat{\ell}_f = \log \hat{L}_f = \log \hat{L} ext{(full model)}$
 $D_m = 2 \left(\hat{\ell}_f - \hat{\ell}_m \right) = 2 \log \left(\hat{L}_f / \hat{L}_m \right)$

For binomial data, D_m is a measure of goodness-of-fit of the model. Asymptotically,

$$D_m \approx \chi^2_{n-k-1}$$

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For all kinds of models, deviance difference is used for comparison of nested models (LRT of significance of added predictors):

$$D_{m_1} - D_{m_2} = 2\left(\hat{\ell}_{m_2} - \hat{\ell}_{m_1}\right) \approx \chi^2_{k_2 - k_1}$$

Example: Let us consider low birth weight data with risk factor smoking. Software tell us that

$$\begin{split} \hat{\ell}_1 &= \log \hat{L}_1 = \log \hat{L} (\text{estimated model}) = -114.9023, \\ \hat{\ell}_0 &= \log \hat{L}_0 = \log \hat{L} (\text{empty model}) = -117.336. \end{split}$$

If H_0 : $\beta_1 = 0$ holds, then

$$2\left(\hat{\ell}_1 - \hat{\ell}_0\right) = 2\log\left(\frac{\hat{L}_1}{\hat{L}_0}\right) \approx \chi_1^2$$

We have

$$\Delta D = 2(-114.9023 + 117.336) = 4.8674 > 3.84 = \chi_1^2(0.05),$$

so that the association between smoking and low birth weight is significant.

Interactions

Logistic regression model allows to include (and test) interactions of categorical variables. If variable X has c categories and variable Z d categories, then the interaction X * Z has (c-1)(d-1) categories. They are all possible combinations of non-reference categories of X and Z.

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If an interaction is significant, the significance of the original constituent variables has no interpretation, nor meaning. The effects are crossed, and we have only two ways to handle the situation:

- Perform stratification and do separate analyses in different strata.
- Introduce a new variable, which operates on the cross-product of the crossed variables (the interaction variable), and omit the interacting variables.

Measures of goodness-of-fit

Let $\hat{\ell}_0 = \log \hat{L}_0 = \log \hat{L} \text{(empty model)}.$

• McFadden
$$R_{MF}^2 = 1 - rac{\hat{\ell}_m}{\hat{\ell}_0}$$

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- Nagelkerke $R_N^2 = \frac{R_{\text{CS}}^2}{1 \hat{L}_0^{\frac{2}{0}}}$

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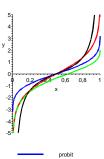
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- Cox & Snell $R_{CS}^2 = 1 \left(\frac{\hat{L}_0}{\hat{L}_m}\right)^{\frac{2}{n}}$
- Nagelkerke $R_N^2 = \frac{R_{CS}^2}{1 \hat{L}_0^{\frac{2}{n}}}$
- Hosmer-Lemeshow test (chi-square test of goodness-of-fit in contingency table between the outcome variable and groups of predicted values)

Alternatives

Logistic function is not the only one used for transformation of probabilities of binary outcomes. Most used are:

- logistic function $\log \left(\frac{p}{1-p} \right)$
- probit function $\Phi^{-1}(p)$
- complementary log-log function log(-log(1 - p))
- negative log-log functionlog(-log(p))
- ullet cauchit function $an\left(\left(p-rac{1}{2}
 ight)\pi
 ight)$

They are called link functions.







 Example: From 1991 U.S. General Social Survey data, we want to check whether sex of a respondent influences the probabilities of life satisfaction feelings.

We get the following contingency table:

sex	life	sum		
367	exiting	routine	dull	Suili
male	213	200	12	425
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 - onot exiting life: routine dull



Or, we have to consider more probabilities and more odds. In our example, we have to consider two multinomial distributions (p₁₁, p₁₂, p₁₃) and (p₂₁, p₂₂, p₂₃), describing the probabilities of life satisfaction feelings for males and females, respectively. The simplest way is to choose one response category as a reference – say exciting life – because one of the probabilities in each row is redundant.

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- Then, the model is

$$\log\left(\frac{p_{ij}}{p_{i1}}\right) = \beta_{0j} + \beta_{1j}x_i, \quad j = 2, 3,$$

where $x_i \in \{0, 1\}$ is the indicator of sex.

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 The previous formula coincides with the simple linear logistic model in the case of dichotomic outcome.



Based on our frequencies, we get the following odds and log-odds:

ode	log-odds		
200/213 = 0,938967	-0,06297	-2,87639	
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The differences of the log-odds in the last two columns are -0,385123874 and -0,845518644. Thus, we can write two models

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, $y = -2.031 - 0.846x$

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L	ife	В	Std. Error	Wald	df	Sig.	Exp(B)	95% CI	
	Intercept	0,322149	0,088338	13,29904	1	0,000266			
Routine	[sex=1]	-0,38512	0,132282	8,476221	1	0,003598	0,680366	0,524982	0,881741
	[sex=2]	0			0				
	Intercept	-2,03087	0,197504	105,7336	1	0			
Dull	[sex=1]	-0,84552	0,356421	5,627561	1	0,01768	0,429335	0,213506	0,86334
	[sex=2]	0			0				

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Then, general multinomial regression model is

$$\log\left(\frac{\rho_{ij}}{\rho_{ij^*}}\right) = \mathbf{x}_i'\beta_j\,,\quad j \neq j^*$$

Inverse formulas are

$$p_{ij} = \frac{\exp(x_i'\beta_j)}{1 + \sum_{\substack{k=1 \ k \neq j^*}}^r \exp(x_i'\beta_k)}, j \neq j^*$$

and

$$p_{ij^*} = \frac{1}{1 + \sum\limits_{\substack{k=1 \\ k \neq j^*}}^{r} \exp\left(x_i' \beta_k\right)}.$$

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The log-likelihood function is

$$\ell(\beta) = \log\left(\frac{n_i!}{\prod_{j=1}^r y_{ij!}}\right) + \sum_{i=1}^n \sum_{j=1}^r y_{ij} \log\left(\rho_{ij}\right)$$

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• Hessian matrix of the estimates of $\beta = \left(\beta_j{}', j = 1, \dots, r, j \neq j^*\right)'$ is

$$H = -\sum_{i=1}^{n} (I_{r-1} \otimes \mathbf{x}_i) \, \hat{V}_i^* \, (I_{r-1} \otimes \mathbf{x}_i)' ,$$

where $\hat{V}_i^* = n_i (\text{diag}(\hat{p}_i^*) - \hat{p}_i^* \hat{p}_i^{*'})$ and \hat{p}_i^* is the vector of all estimates of probabilities p_{ij} except p_{ij^*}

Chi=square of the estimated model is

$$\chi^{2} = \sum_{i=1}^{n} \sum_{j=1}^{r} \frac{(y_{ij} - n_{i}\hat{p}_{ij})^{2}}{n_{i}\hat{p}_{ij}}$$

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 The same pseudo-R² statistics as in logistic regression model can be used.

If actual variance matrix of y_i are substantially larger than $V_i = n_i \, (\text{diag} \, (p_i) - p_i p_i') \, (\text{given by the multinomial model})$, we speak about **overdispersion**. Then, we introduce scale parameter σ^2 , such that var $y_i = \sigma^2 \, V_i$.

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- Use of σ^2 does not change estimators of β .
- Variance of the estimators is

$$\operatorname{var} \hat{\beta} = \hat{\sigma}^2 \left[\sum_{i=1}^n \left(I_{r-1} \otimes \mathbf{x}_i \right) \hat{\mathbf{V}}_i^* \left(I_{r-1} \otimes \mathbf{x}_i \right)' \right]^{-1}$$

Tests

• For any $L_{q \times k+1}$ of full rank, it holds

$$\hat{\beta}_j' L' \left[L \operatorname{var} \hat{\beta}_j L' \right]^{-1} L \hat{\beta}_j \approx \chi_q^2$$

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• LR test of nested models with k_1 and k_2 ($k_1 < k_2$) regression parameters is based on

$$\frac{1}{\hat{\sigma}^2}(D_{m_1} - D_{m_2}) = \frac{2}{\hat{\sigma}^2} \left(\hat{\ell}_{m_2} - \hat{\ell}_{m_1}\right) \approx \chi^2_{k_2 - k_1}$$

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Overdispersion is not observed. Software gets:

$$\hat{\ell}_{m_1} = -25,8165, \quad \hat{\ell}_{m_2} = -24.332$$

Therefore,

$$\Delta D_m = 2(-24.332 + 25,8165) = 2.969 < 9.488 = \chi^2_{8-4}(0.05)$$

Corresponding p-value is 0.563. The race factor proves to be non-significant.

Example: Random sample of Vermont citizens was asked to rate the work of criminal judges in the state. The scale was Poor (1), Only fair (2), Good (3), and Excellent (4). At the same time, they had to report whether somebody of their household had been a crime victim within the last 3 years.

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The data:

Household victim	Judges' performance					
riouseriola victim	Poor	Only fair	Good	Excellent	sum	
Yes	14	28	31	3	76	
No	38	170	248	34	490	
sum	52	198	279	37	566	

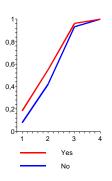


With ordinal data, it is natural to consider probabilities of cumulative events, like specific score or worse. Table of cumulative frequencies is as follows:

	Judges' performance							
Household victim	Poor	Only fair	Good	Excellent				
	1 001	or worse	or worse	or worse				
Yes	14	42	73	76				
row percentage	18,42%	55,26%	96,05%	100,00%				
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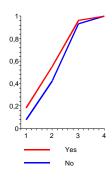


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The graph suggests that having a crime victim in the household implies more negative opinion on judges' performance.

The lines must meet at 100%. Otherwise they look almost parallel. That suggest model with common slope for both categories.



Let us denote $p_{ij}^c = P(\text{score} \le j)$, i = 1(No), 2(Yes), j = 1, 2, 3 the non-trivial cumulative probabilities. Then, our model is

$$\log\left(\frac{p_{1j}^c}{1 - p_{1j}^c}\right) = \alpha_j \quad \text{and} \quad \log\left(\frac{p_{2j}^c}{1 - p_{2j}^c}\right) = \alpha_j + \beta,$$

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Software gets $\alpha_1 = -2.39$, $\alpha_2 = -0.32$, $\alpha_2 = 2.59$, $\beta = 0.63$. Using standard inverse formula for logits, we obtain the following estimates:

	≤ 1	≤ 2	≤ 3
Yes	14,69%	57,85%	96,18%
No	8,38%	42,15%	93,04%

Proportional odds model

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Y be ordinal response variable with possible values 1,...,r

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Model: logits of cumulative probabilities $p_j^c(x) = P(Y \le j | X = x)$ satisfy

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Because log of cumulative odds ratio of making the same responses at different x-points is proportional to the distance of the points, the model is called proportional odds model:

$$\log \left(\frac{p_j^c(x_1)}{1 - p_j^c(x_1)} \cdot \frac{1 - p_j^c(x_2)}{p_j^c(x_2)} \right) = \beta'(x_1 - x_2)$$

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- All standard goodness-of-fit measures apply.
- The fit is different than separate logit models for all j's.

Example: Software output for Vermont crime data:

		Estimate	Std. Error	Wald	df	Sig.	95% con	f. interval
Threshold	[rating = 1]	-2,39221	0,15177	248,44332	1	0,00000	-2,68968	-2,09475
	[rating = 2] [rating = 3]	-0,31651 2,59316	0,09082 0,17163	12,14637 228,28667	1	0,00049 0,00000	-0,49451 2,25678	-0,13852 2,92955
Location	[hhcrime=1]	-0,63298	0,23198	7,44539	1	0,00636	-1,08765	-0,17831
	[hhcrime=2]	0			0			

Notice opposite sign of the coefficient β (hhcrime=1). Many work with the model $\alpha_j - \beta x$ because of interpretation reasons: in such a case, higher coefficients indicate association with higher scores.

Example: Software output for Vermont crime data:

		Estimate	Std. Error	Wald	df	Sig.	95% con	f. interval
Threshold	[rating = 1] [rating = 2]	-2,39221 -0,31651 2,59316	0,15177 0,09082 0,17163	248,44332 12,14637 228,28667	1	0,00000 0,00049 0.00000	-2,68968 -0,49451 2,25678	-2,09475 -0,13852 2,92955
Location	[rating = 3] [hhcrime=1] [hhcrime=2]	-0,63298 0	0,17163	7,44539	1 0	0,00636	-1,08765	-0,17831

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Let us now add another predictor variable, sex:

		Estimate	Std. Error	Wald	df	Sig.	95% con	f. interval
Threshold	[rating = 1] [rating = 2] [rating = 3]	-2,57419 -0,48730 2,43740	0,17641 0,12326 0,18672	212,93519 15,62868 170,40298	1 1 1	0,00000 0,00008 0,00000	-2,91995 -0,72890 2,07143	-2,22844 -0,24571 2,80336
Location	[hhcrime=1] [hhcrime=2] [sex=1] [sex=2]	-0,62074 0 -0,34145 0	0,23228 0,16030	7,14177 4,53709	1 0 1 0	0,00753 0,03317	-1,07599 -0,65563	-0,16548 -0,02726

We suspect that sex may influence sensitivity to crime victims, so that we add the interaction:

		Estimate	Std. Error	Wald	df	Sig.	95% con	f. interval
Threshold	[rating = 1]	-2.64904	0.18097	214.26179	1	0.00000	-3.00374	-2.29434
11110011010	[rating = 1]	-0.55150	0.12873	18,35418	1	0.00002	-0.80381	-0.29920
	[rating = 3]	2,38107	0,18819	160,07877	1	0,00000	2,01222	2,74993
Location	[hhcrime=1]	-1,13654	0,33008	11,85565	1	0,00057	-1,78350	-0,48959
	[hhcrime=2]	0			0			
	[sex=1]	-0,46925	0,17330	7,33183	1	0,00677	-0,80891	-0,12959
	[sex=2]	0			0			
	[hhcrime=1] * [sex=1]	0,95889	0,46413	4,26832	1	0,03883	0,04921	1,86857
	[hhcrime=1] * [sex=2]	0			0			
	[hhcrime=2] * [sex=1]	0			0			
	[hhcrime=2] * [sex=2]	0			0			

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	[hhcrime=1] * [sex=2]	0			0			
	[hhcrime=2] * [sex=1]	0			0			
	[hhcrime=2] * [sex=2]	0			0			

But, since the interaction is significant, the two individual variables don't have good meaning any more:

			Std. Error	Wald	df	Sig.	95% con	f. interval
Threshold	[rating = 1] [rating = 2]	-2,64904 -0.55150	0,18097 0.12873	214,26179 18.35418	1	0,00000 0.00002	-3,00374 -0.80381	-2,29434 -0,29920
	[rating = 3]	2,38107	0,18819	160,07877	1	0,00000	2,01222	2,74993
Location	[hhcrime=1] * [sex=1]	-0,64690	0,32950	3,85460	1	0,04961	-1,29270	-0,00110
	[hhcrime=1] * [sex=2]	-1,13654	0,33008	11,85565	1	0,00057	-1,78350	-0,48959
	[hhcrime=2] * [sex=1]	-0,46925	0,17330	7,33183	1	0,00677	-0,80891	-0,12959
	[hhcrime=2] * [sex=2]	0			0			

Redundant parameters are not estimated, so that interaction itself is enough. This model has the same χ^2 , deviance, and pseudo- R^2 as the previous one.

Other ordinal regression models

General cumulative logit model is

$$\log\left(\frac{p_j^c(x)}{1-p_j^c(x)}\right) = \alpha_j + \beta_j'x \quad \forall j = 1, \dots, r-1$$

Thus, every group has its own slope. Proportional odds model is a special case, and can be tested by LR test.

Other ordinal regression models

General cumulative logit model is

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Thus, every group has its own slope. Proportional odds model is a special case, and can be tested by LR test.

Adjacent categories model is

$$\log\left(\frac{p_j(\mathbf{x})}{p_{j+1}(\mathbf{x})}\right) = \alpha_j + \beta' \mathbf{x} \quad \forall j = 1, \dots, r-1$$

This model recognizes the ordering, since

$$\log\left(\frac{p_j(x)}{p_r(x)}\right) = \sum_{m=j}^r \alpha_m + \beta'(r-j)x \quad \forall j$$

The suffering is over...

Thank you for your attention!

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