

Ordered logistic regression R

```
plot(s, which=1:3, pch=1:3,  
      xlab='logit', main=' ',  
      xlim=range(s[,3:4]))
```

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- ▶ Once we are done assessing whether the assumptions of our model hold, we can obtain predicted probabilities, which are usually easier to understand than either the coefficients or the odds ratios.
- ▶ For example, we can vary gpa for each level of pared and public and calculate the probability of being in each category of apply.
- ▶ We do this by creating a new dataset of all the values to use for prediction.

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```
newdat <- data.frame(  
  pared = rep(0:1, 200),  
  public = rep(0:1, each = 200),  
  gpa = rep(seq(from = 1.9, to = 4,  
    length.out = 100), 4))  
  
newdat <- cbind(newdat, predict(m,  
  newdat, type = "probs"))
```

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```
# Show first few rows
```

```
head(newdat)
```

	pared	public	gpa	unlikely	somewhat	likely	very likely
1	0	0	1.900	0.7376		0.2205	0.04192
2	1	0	1.921	0.4932		0.3946	0.11221
3	0	0	1.942	0.7325		0.2245	0.04299
4	1	0	1.964	0.4867		0.3985	0.11484
5	0	0	1.985	0.7274		0.2285	0.04407
6	1	0	2.006	0.4802		0.4023	0.11753

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- ▶ Now we can reshape the data long with the reshape2 package and plot all of the predicted probabilities for the different conditions.
- ▶ We plot the predicted probabilities, connected with a line, coloured by level of the outcome, apply, and faceted by level of pared and public.
- ▶ We also use a custom label function, to add clearer labels showing what each column and row of the plot represent.

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```
library(reshape2)

lnewdat <- melt(newdat,
  id.vars = c("pared", "public", "gpa"),
  variable.name = "Level",
  value.name="Probability")
```

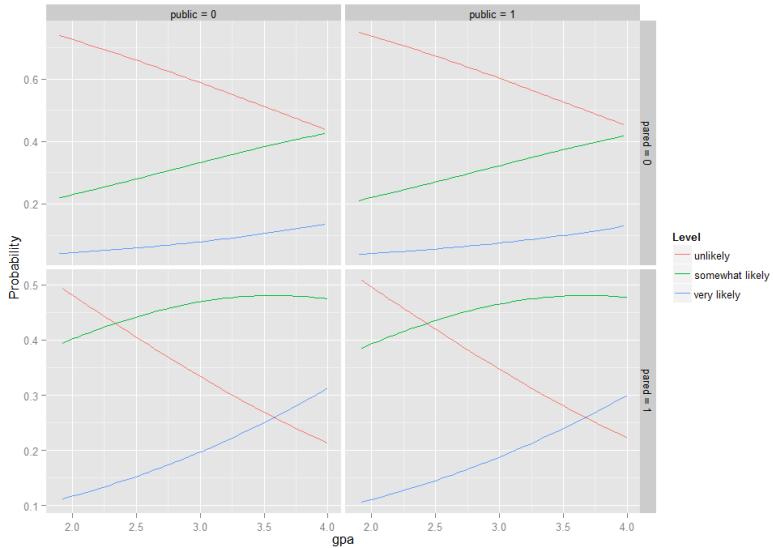
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```
### view first few rows
head(lnewdat)
```

	pared	public	gpa	Level	Probability
### 1	0	0	1.900	unlikely	0.7376
### 2	1	0	1.921	unlikely	0.4932
### 3	0	0	1.942	unlikely	0.7325
### 4	1	0	1.964	unlikely	0.4867
### 5	0	0	1.985	unlikely	0.7274
### 6	1	0	2.006	unlikely	0.4802

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```
ggplot(lnewdat, aes(x = gpa, y = Probability,
  colour = Level)) +
  geom_line() +
  facet_grid(pared ~ public, scales="free",
    labeller=function(x, y) sprintf("%s = %d", x, y))
```

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Things to consider **Perfect prediction:** Perfect prediction means that one value of a predictor variable is associated with only one value of the response variable.

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Sample size: Both ordered logistic and ordered probit, using maximum likelihood estimates, require sufficient sample size.

Empty cells or small cells: You should check for empty or small cells by doing a crosstab between categorical predictors and the outcome variable. If a cell has very few cases, the model may become unstable or it might not run at all.

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Pseudo-R-squared: There is no exact analog of the R-squared found in OLS. There are many versions of pseudo-R-squares. Please see Long and Freese 2005 for more details and explanations of various pseudo-R-squares.

Diagnostics: Doing diagnostics for non-linear models is difficult, and ordered logit/probit models are even more difficult than binary models.