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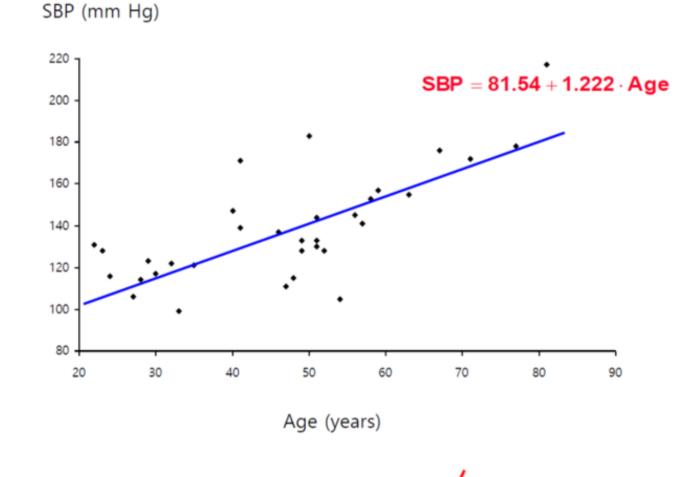
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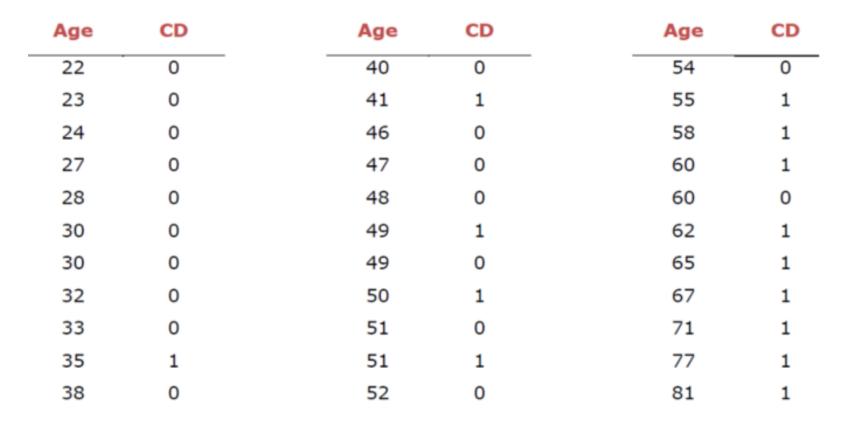
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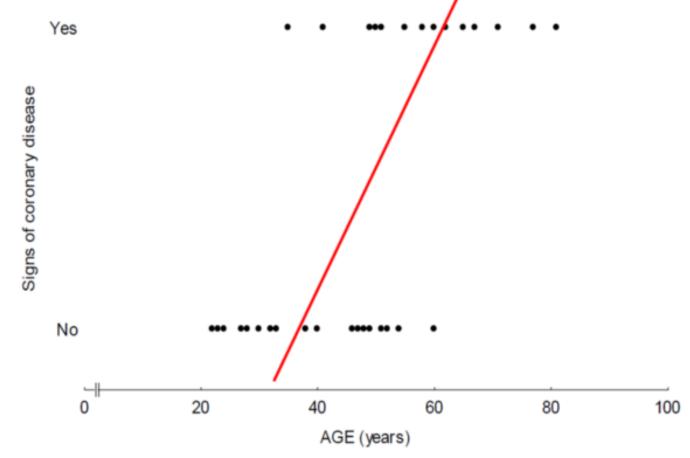
Introduction

Age	SBP
22	131
23	128
24	116
27	106
28	114
29	123
30	117
32	122
33	99
35	121
40	147

Age	SBP	Ag	e SBP
41	139	52	128
41	171	54	105
46	137	56	145
47	111	57	7 141
48	115	58	3 153
49	133	59	157
49	128	63	155
50	183	67	7 176
51	130	71	172
51	133	77	178
51	144	81	217







- Linear Regression
- $y = X\beta + \epsilon$
- Estimates continuous variable
 (blood pressure in the example)
- What about estimating categorical (count) variable or probability (proportion) ??
- Could be **{0, 1}** or **[0, 1]**
- If applying linear regression,?!

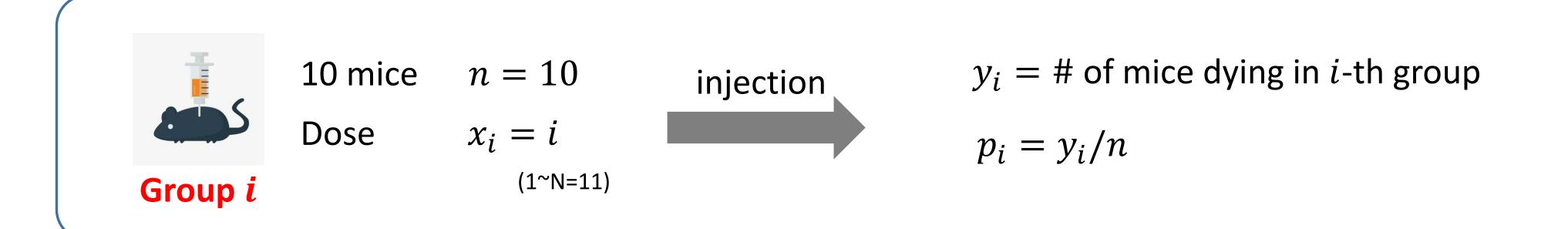
Odds and Logistic Transformation

$$Odds = \frac{P(A)}{P(A^c)} = \frac{\pi}{1-\pi} \qquad \Rightarrow \text{ in the range } [0, \infty] \qquad \lambda = \log\{odds\} = \log\frac{\pi}{1-\pi} \qquad \Rightarrow \text{ in the range } [-\infty, \infty]$$

$$\pi = \alpha_0 + \alpha_1 X \qquad \qquad \log \left\{ \frac{\pi}{1-\pi} \right\} = \alpha_0 + \alpha_1 X$$

Now, we can consider count or proportion data as holding linear regression frames

Example: dose response



We are going to model y_i as independent binomials

$$y_i \stackrel{\text{ind}}{\sim} \text{Bi}(n_i, \pi_i)$$
 for $i = 1, 2, ..., N$

Assume that the logit follows linear function of dose

$$\lambda_i = \log\left\{\frac{\pi_i}{1 - \pi_i}\right\} = \alpha_0 + \alpha_1 x_i$$

Example: dose response



10 mice

$$n = 10$$

injection

 $y_i = \#$ of mice dying in *i*-th group

(1~N=11)

 $p_i = y_i/n$

(blue dots)

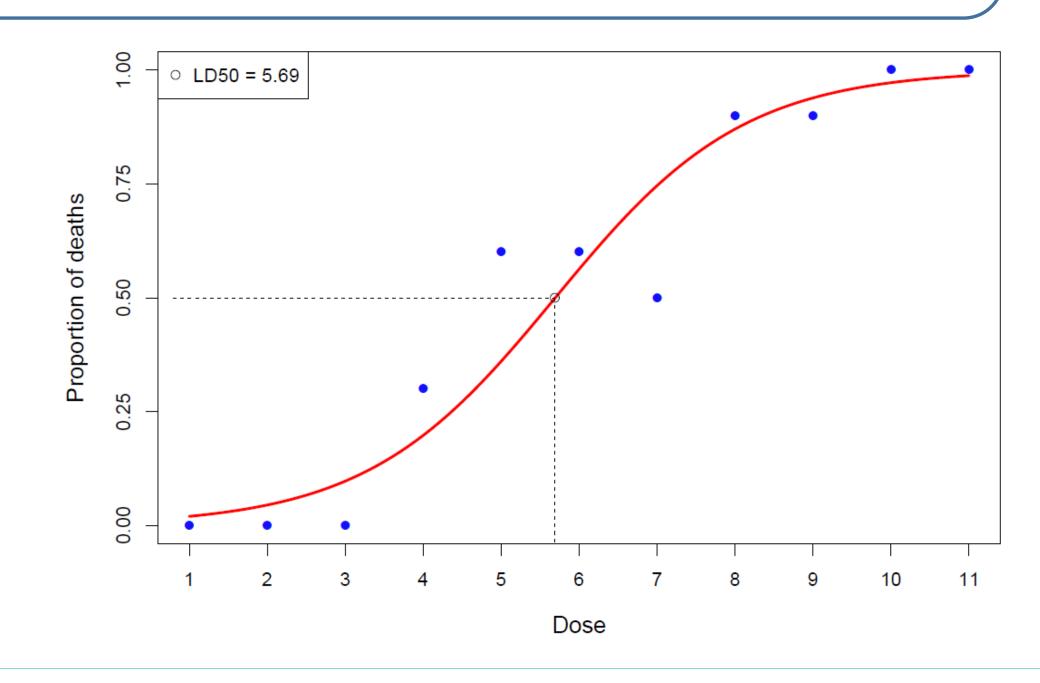
Group i

MLE provides $(\hat{\alpha}_0, \hat{\alpha}_1)$ which yields the following equation

$$\hat{\lambda} = \log \left\{ \frac{\hat{\pi}}{1 - \hat{\pi}} \right\} = \hat{\alpha}_0 + \hat{\alpha}_1 \mathbf{x}$$

And we finally obtain the linear logistic regression curve

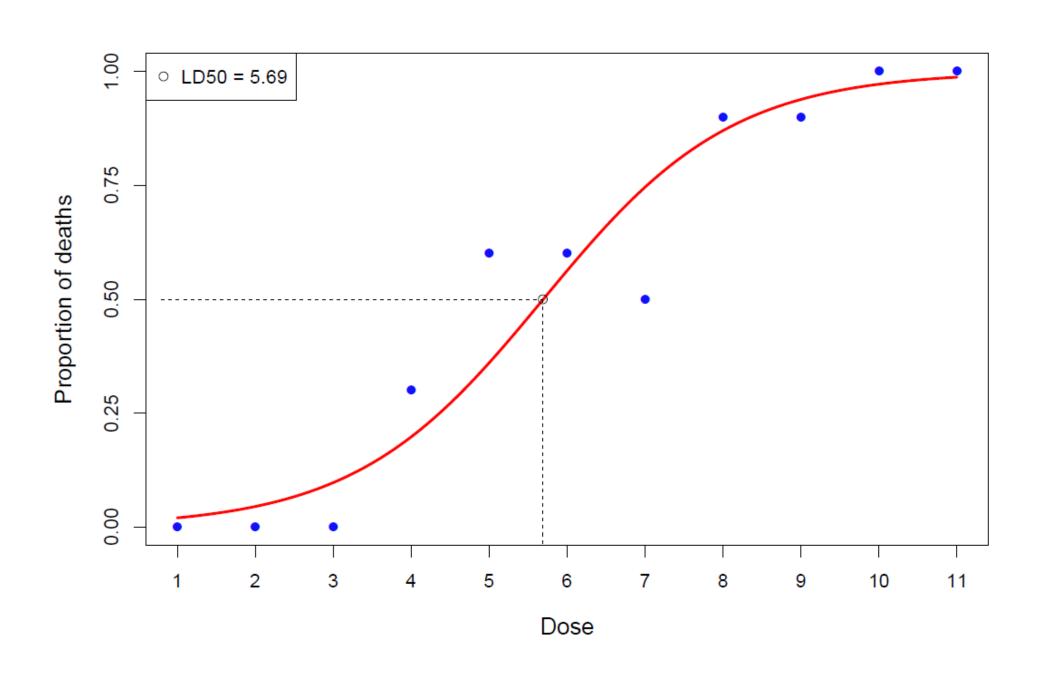
$$\widehat{\pi}(x) = \left(1 + e^{-(\widehat{\alpha}_0 + \widehat{\alpha}_1 x)}\right)^{-1}$$



Note: dose response

- The regression has reduced error (see the table on the right side)
- It is ture unless logit linear model seriously goes wrong
- λ is NOT restricted to the range [0, 1]
- Able to utilize exponential family properties!

x	1	2	3	4	5	6	7	8	9	10	11
$\operatorname{sd} \hat{\pi}(x)$ $\operatorname{sd} p_i$											



Merging Into Exponential Family

The probability density function of Bi(n, y) is given

$$\binom{n}{y}\pi^{y}(1-\pi)^{n-y} = e^{\lambda y - n\psi(\lambda)}\binom{n}{y}$$
 \rightarrow one parameter exponential family (see chapter 5.5; eq 5.54 or 5.46)

, where $\psi(\lambda) = \log\{1 + e^{\lambda}\}$

The independence of the data gives the probability density of full data set y as a function of (α_0, α_1) ,

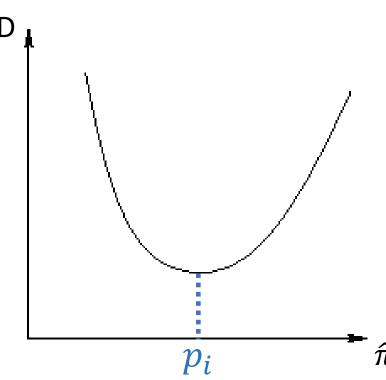
$$f_{\alpha_0,\alpha_1}(\mathbf{y}) = \prod_{i=1}^N e^{\lambda_i y_i - n_i \psi(\lambda_i)} \binom{n_i}{y_i}$$
$$= e^{\alpha_0 S_0 + \alpha_1 S_1} \cdot e^{-\sum_{i=1}^N n_i \psi(\alpha_0 + \alpha_1 x_i)} \cdot \prod_{i=1}^N \binom{n_i}{y_i}$$

, where $S_0 = \sum_{i=1}^N y_i$ and $S_1 = \sum_{i=1}^N x_i y_i$

Merging Into Exponential Family

Suppose that the deviance is given as follows

$$D(p_i, \hat{\pi}_i) = 2n_i \left[p_i \log \left(\frac{p_i}{\hat{\pi}_i} \right) + (1 - p_i) \log \left(\frac{1 - p_i}{1 - \hat{\pi}_i} \right) \right]$$



The deviance gives us the intuition: it is zero at $\hat{\pi}_i = p_i$, otherwise it increases as $\hat{\pi}_i$ departs further from p_i

The logistic regression MLE value $(\hat{\alpha}_0, \hat{\alpha}_1)$ has to do with minimizing the total deviance between p_i and $\hat{\pi}_i = \pi_{\alpha_0, \alpha_1}(x_i)$

$$(\hat{\alpha}_0, \hat{\alpha}_1) = \underset{(\alpha_0, \alpha_1)}{\operatorname{arg \, min}} \sum_{i=1}^N D\left(p_i, \pi_{\alpha_0, \alpha_1}(x_i)\right)$$

Example: cell infusion

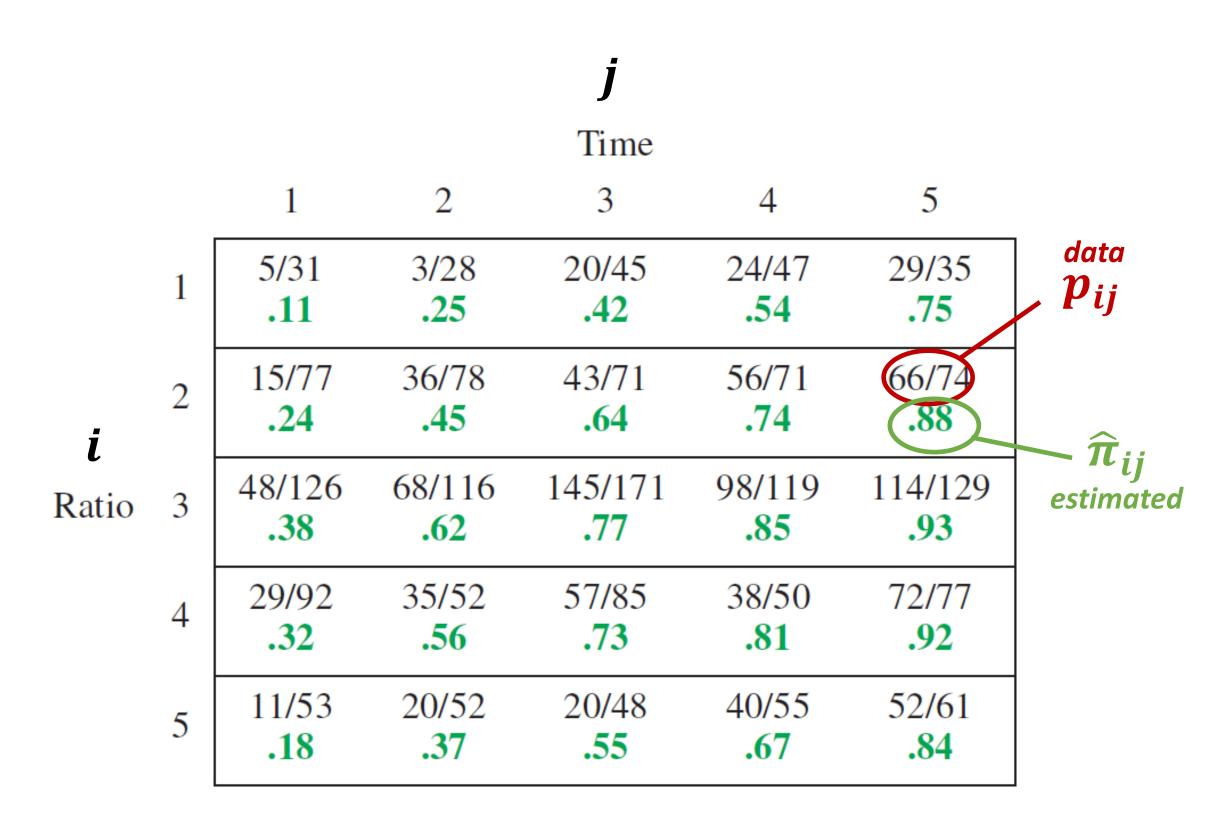
Let π_{ij} denote the true probability of thriving of ratio i during time period j

And take logistic regression

$$\lambda_{ij} = \log\left\{\frac{\pi_{ij}}{1 - \pi_{ij}}\right\} = \mu + \alpha_i + \beta_i$$

MLE and the data set $\{p_{ij}\}$ give estimation $\hat{\pi}_{ij}$ as follows

$$\hat{\pi}_{ij} = \frac{1}{1 + e^{-(\hat{\mu} + \hat{\alpha}_i + \hat{\beta}_j)}}$$



Cell infusion data; human cell colonies infused with mouse nuclei in five ratios over 1 to 5 days and observed to see whether they did or not thrive.

Example: spam filter

George labeled N=4601 emails whether spam or ham(i.e. non-spam) He used 57 words as predictors in the table.

 x_{ij} : relative frequency of keyword j in email i

 $\pi_{i\,i}$: true probability that email i is spam

 λ_i : the logit transformation of π_{ij}

$$\lambda_{ij} = \log\left\{\frac{\pi_{ij}}{1 - \pi_{ij}}\right\} = \alpha_0 + \sum_{j=1}^{57} \alpha_j x_{ij}$$

- Then you are able to predict whether future emails are spam or ham by using these keywords
- The table provides the estimated $\hat{\alpha}_i$ and its se value (by MLE)
- It seems that 'free' and 'your' are good spam predictors large $\hat{\alpha}_i$ and small se; large z-value
- The occasional very large $\hat{\alpha}_i$ may bother MLE

se								
	Estimate	se	z-value		Estimate	se	z-value	
intercept	-12.27	1.99	-6.16	lab	-1.48	.89	-1.66	
make	12	.07	-1.68	labs	15	.14	-1.05	
address	19	.09	-2.10	telnet	07	.19	35	
all	.06	.06	1.03	857	.84	1.08	.78	
3d	3.14	2.10	1.49	data	41	.17	-2.37	
our	.38	.07	5.52	415	.22	.53	.42	
over	.24	.07	3.53	85	-1.09	.42	-2.61	
remove	.89	.13	6.85	technology	.37	.12	2.99	
internet	.23	.07	3.39	1999	.02	.07	.26	
order	.20	.08	2.58	parts	13	.09	-1.41	
mail	.08	.05	1.75	pm	38	.17	-2.26	
receive	05	.06	– .86	direct	11	.13	84	
will	12	.06	-1.87	cs	-16.27	9.61	-1.69	
people	02	.07	35	meeting	-2.06	.64	-3.21	
report	.05	.05	1.06	original	28	.18	-1.55	
addresses	.32	.19	1.70	project	98	.33	-2.97	
free	.86	.12	7.13	re	80	.16	-5.09	
business	.43	.10	4.26	edu	-1.33	.24	-5.43	
email	.06	.06	1.03	table	18	.13	-1.40	
you	.14	.06	2.32	conference	-1.15	.46	-2.49	
credit	.53	.27	1.95	char;	31	.11	-2.92	
your	.29	.06	4.62	char(05	.07	75	
font	.21	.17	1.24	char_	07	.09	78	
000	.79	.16	4.76	char!	.28	.07	3.89	
money	.19	.07	2.63	char\$	1.31	.17	7.55	
hp	-3.21	.52	-6.14	char#	1.03	.48	2.16	
hpl	92	.39	-2.37	cap.ave	.38	.60	.64	
george	-39.62	7.12	-5.57	cap.long	1.78	.49	3.62	
650	.24	.11	2.24	cap.tot	.51	.14	3.75	

THANK YOU

Q&A