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Introduction to Biostatistics
Lecture 6

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What shall we learn today?

Many analyses and concepts that relates to count data (tables), in particular odds.

An overly optimistic description: Lectures 3-5 but for categorical, rather than numerical, data.

For tables relating counts of a categorical variable, how can we test the distribution of values against

- a model (probability function), or
- between subgroups.

We will start with the special case of dichotomous/binary data (event or non-event).

A note on categorical data

We generally talk differently about categorical data depending on the number of unique values.

Dichotomous/binary data is typically yes/no- or event/non-event data. We talk about this data in terms of the probability of one of the events (often the one with the smallest probability). If Prob(event) is modelled, then Prob(non-event) is implicit (since Prob(non-event)=1-Prob(event)).

"Non-binary" categorical data has more than 2 values (A, B, C, ...). We talk about this data in terms of the entire *distribution*, i.e. the probability function $(\text{Prob}(A), \text{Prob}(B), \text{Prob}(C), \ldots)$. Of course we could omit one of these, since it would be implicit, but it is unconvenient.

Numeric data verses categorical

1 sample:

Suppose $x_1, x_2, \ldots, x_n$ is a sample from some population.

Numeric: population mean $= \mu$? Answered by 1-sample t-test (parametric) or Wilcoxon signed rank test (non-parametric).

Categorical: is the probability function $= p(k)$? Or, if binary, is Prob(event) $= p$? Answered by e.g. confidence interval for p or $\chi^2 - test$.

2-sample paired:

Numeric: taking differences reduces this to the 1-sample situation.

Categorical: McNemars test.

Numeric data verses categorical

2 samples:

Suppose $x_1, x_2, \ldots, x_n$ and $y_1, y_2, \ldots, y_n$ are samples from possibly different populations.

Numeric: Are the population means the same? Answered by 2-sample t-test (parametric) or Mann-Whitney (non-parametric).

Categorical: are the probability function the same? Or, if binary, is Prob(event) the same? Answered by Fishers exact test or χ^2-test.

Many samples:

Numeric: Are all means the same? Answered by ANOVA (parametric) or Kruskal-Wallis (non-parametric)

Categorical: Are all probability functions the same? Answered by χ^2-test.

Dabigatran data

Dagbigatran is an anticoagulant used for e.g. stroke prevention in patients with atrial fibrillation. *The following example only looks at side effects.*

718 people were randomized to Dabigatran or placebo and observed for some set time for bleeding.

id	intervention	bleeding
1	dabigatran	Yes
2	placebo	No
3	placebo	No
4	dabigatran	No
⋮	⋮	⋮
718	placebo	No

Tabulated data

	Bleeding		
	Yes	No	Sum
dabigatran	27	320	347
placebo	8	363	371
Sum	35	683	718

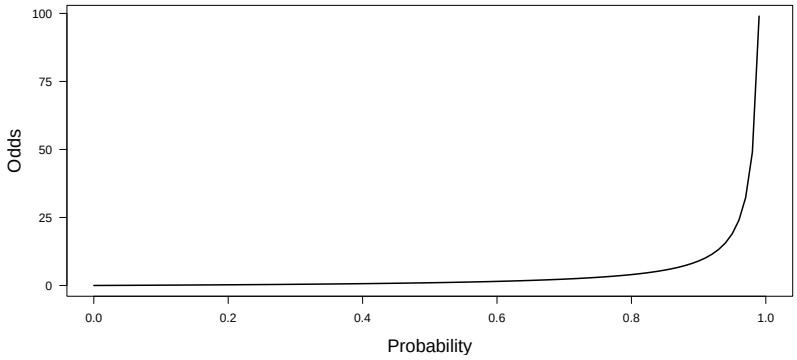
Measures for dabigatran:

- **Risk** (probability of an unwanted event)
  Risk of bleeding $= 27/347 = 0.078$
- **Odds** (how much more likely it is, versus not, to experience an event)
  Odds of bleeding
  $$= \frac{27/347}{320/347} = \frac{27}{320} = 0.084$$

(Odds? Sometimes this is easier to model.)

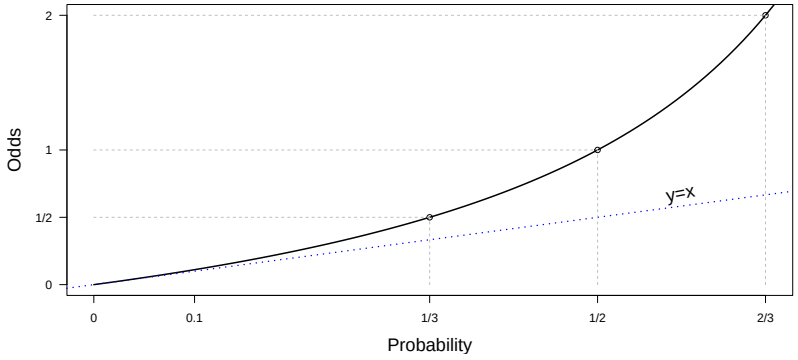
Odds

For an event with probability p, the odds is $p/(1-p)$.



Odds

For small probabilities: odds $\approx$ probability.



A few remarks on odds

- if an event has odds θ, then its probability p is $p = \theta/(1+\theta)$
  E.g.
  - $\theta = 2$ corresponds to $p = 2/3$.
  - $\theta = 1$ corresponds to $p = 1/2$.
  - $\theta = 1/100$ corresponds to $p = 1/101$.
- there are multiple systems of betting (sports) 'odds', that are not odds in the sense of this course!
- in a betting game where you stand to win 1 unit of money, your stake S (*if this is kept when winning*) should not exceed the odds
  - 'expected' profit $= 1p - S(1-p) \geq 0$ is equivalent to $S \leq \frac{p}{1-p} = \theta$
  - E.g. if you are offered x units of money for a game you think has odds 2 (in your favor) then do not bet more than $2x$.
    (Betting $2x$ makes the game *fair*.)

Relational measures

	Bleeding		
	Yes	No	Sum
dabigatran	27	320	347
placebo	8	363	371
Sum	35	683	718

Measures for risk of dabigatran versus placebo

- (**Risk ratio (RR)** $= \frac{27/347}{8/371} \approx 3.6$)
- **Odds ratio (OR)** $= \frac{27/320}{8/363} \approx 3.8$
- **Risk difference** $= 27/347 - 8/371 \approx 0.056$

Odds ratio (OR)

Probabilities cannot be retrieved from the OR alone.

N.B.

- Odds = 1 means p = 0.5 (as likely to experience the event as to not experience the event)
- OR = 1 means events are equally likely in both groups

If you know the 'denominator' probability (p_2) then the 'numerator' probability (p_1) can be calculated

$$p_1 = \frac{\text{OR} \cdot p_2}{1 + (\text{OR} - 1) \cdot p_2}.$$

For small values of p_2 and 'moderate' values of OR

$$p_1 \approx \text{OR} \cdot p_2,$$

i.e.

$$\text{OR} \approx \frac{p1}{p2} = \text{RR}.$$

Fishers exact test

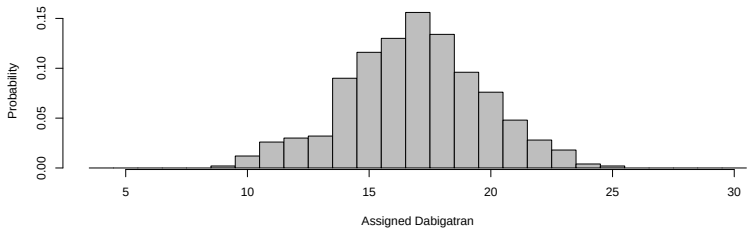
	Bleeding		
	Yes	No	Sum
dabigatran	X	(347-X)	347
placebo	(35-X)	(683-347+X)	371
Sum	35	(683)	718

Suppose that whether a person bleeds or not is independent of intervention. (H_0: "odds ratio = 1".)

Then the 35 individuals who bled should be a random sample of the study population (of size 718) and we would expect that $X/35 = 347/718 \approx 48\%$.

We would expect X to be around 17, but is $X = 27$ within some acceptable range of possiblities?

Simulate (500 times) the experiment of randomly selecting 35 people from the study population and record the number who got dabigatran (15, 15, 16, 18, 16...)

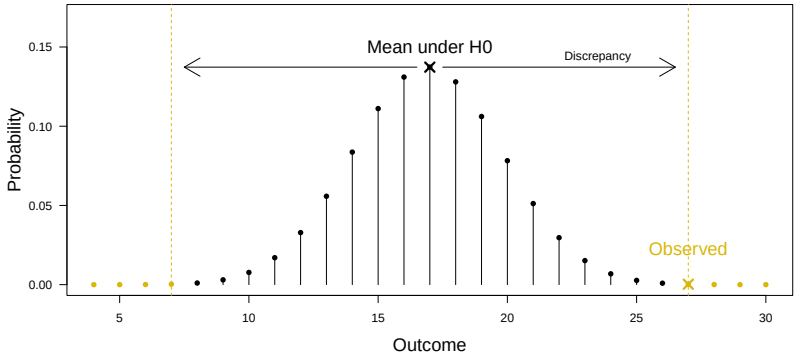


However, we can calculate *exactly* what the distribution of X is *given* H_0 (in this case).

The p-value is the probability of a discrepancy the size of that between the observed and the expected.

p-value in Fishers exact test

Sum the yellow values to get $p = 0.00045$.



More on Fishers exact test

My software produced the following output:

```
   Fisher's Exact Test for Count Data

data:  Dabigatran_example
p-value = 0.0004458
alternative hypothesis: true odds ratio is not equal to 1
95 percent confidence interval:
 1.659358 9.877595
sample estimates:
odds ratio
  3.821942
```

So odds ratio is between 1.7 and 9.9. (Allows for hypothesis testing.)
Probabilities are small, so risk of dabigatran is (approx.) between 1.7 and 9.9 times larger than placebo risk.

Risk difference

What is the *difference* in risk between dabigatran and placebo?
This has (almost) been covered by Lars. One needs to know that for two **independent** estimators (having SE_1 and SE_2) the SE for their difference is given by

$$\sqrt{SE_1^2 + SE_2^2}.$$

Risk	Estimate	Standard error
dabigatran	$p_1 = 27/347 = 0.078$	$\sqrt{p_1(1-p_1)/347} = 0.0144$
placebo	$p_2 = 8/371 = 0.022$	$\sqrt{p_2(1-p_2)/371} = 0.0075$
difference	$p_1 - p_2 = 0.056$	$\sqrt{0.0144^2 + 0.0075^2} = 0.0162$

We get a 95% confidence interval for the difference with

$$(0.056 \pm 1.96 \cdot 0.0162) = (0.024, 0.088).$$

(This allows for hypothesis testing.)

Absolute risk

What can we say about the *absolute* risk of bleeding with dabigatran?

This was covered by Lars in Lecture 3! (Genotype example.)

The risk estimate $27/347 = 0.078$ has a standard error (SE) given by

$$\sqrt{\frac{0.078(1-0.078)}{347}} = 0.0144.$$

This yields a 95% confidence interval given by

$$(0.078 \pm 1.96 \cdot 0.0144) = (0.050, 0.11).$$

(This allows for hypothesis testing.)

Rule of thumb: $n * \min(p, 1-p) \geq 5$.

Have we exhausted the Dabigatran example yet?

It certainly seems so (but it will actually return again later in the lecture!)

Summary of the dabigatran example:

Quantity	Estimate	Confidence interval
p_1	0.078	$(0.050, 0.11)$
$p_1 - p_2$	0.056	$(0.024, 0.088)$
OR (p1 vs. p2)	3.82	$(1.7, 9.9)$

Cosmetic skin testing

To prove a new product is hypoallergenic it should provoke no more skin reactions than current market leader.

To test a new product 40 individuals got both products applied to patches of skin and observed for reaction (yes/no)

id	new	market
1	no	no
2	yes	no
3	no	no
⋮	⋮	⋮
40	yes	yes

Is the new product as good as the market leader?

The following table is *not* appropriate to answer that question.

	no	yes
old	22	18
new	32	8

The rows are dependent.

Note:

One *can* estimate (and get confidence intervals) for p_1 and p_2 (risk of skin reaction with new and old, respectively).

But it is harder to quantify the SE for risk difference and OR, due to the dependence.

McNemars test for paired data

		new		
		no	yes	$\sum$
market	no	17	5	22
	yes	15	3	18
	$\sum$	32	8	40

McNemars test only considers the pairs were the results are different.

With new product there are 8 reactions but 3 of them would have happened anyway. The new product 'creates' 5 reactions, whereas the market leader 'creates' 15 reactions.

Switching from the market leader to the new product would benefit 15, make it worse for 5, and have no effect on 20. (In terms of skin reactions.)

A test statistic can be calculated. p-value for H_0:'no difference' is approx 4%.

Other situations

Twins being randomized to intervention or placebo:

		Placebo	
		improvement	non
Intervention	improvement	a	b
	non	c	d

Before/after data:

		Before	
		event	non
After	event	a	b
	non	c	d

Mendel's pea experiment

One of Mendel's pea-experiments was a (dihybrid) cross between the genes for round/wrinkled seeds and yellow/green seeds.

Type	RY	RG	WY	WG	Sum
Count (O)	315	108	101	32	558

According to his theory, these should appear in ratios of 9:3:3:1.
So, we have a model for X ="the type":

Value v	RY	RG	WY	WG
Prob($X = v$)	9/16	3/16	3/16	1/16

χ^2-tests

χ^2 tests are applied to tabulated data (i.e. the 'counts'), typically categorical data. (E.g. the Dabigatran data.)

Like the t-test, we can use χ^2 to compare a sample against a model or, compare 2 or more samples against each other.

χ^2 tests typically calculate a test statistic Q according to the formula

$$Q = \sum \frac{(\text{observed - expected})^2}{\text{expected}}.$$

Q is compared to a χ^2 distribution with a parameter (degrees of freedom) that depends on the situation.

This is not an exact test. Rule of thumb: expected cell count ≥ 5.

Comparing data to a model

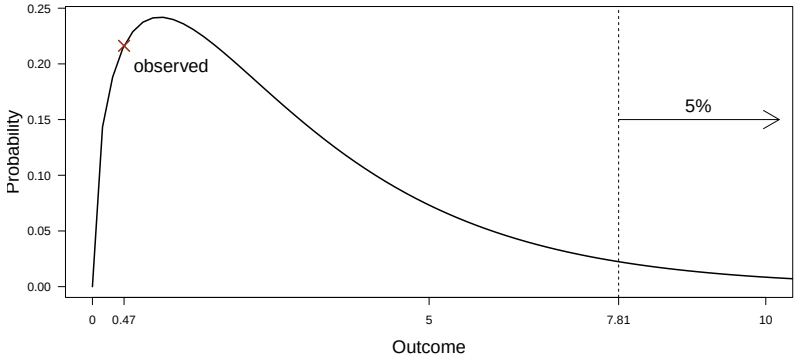
χ^2-analysis:

Type	RY	RG	WY	WG	Sum
Data (O)	315	108	101	32	558
H_0 model (p)	9/16	3/16	3/16	1/16	1
Expected ($E = 558 \times p$)	313.9	104.6	104.6	34.9	558
Q, i.e. $(O-E)^2/E$	0.004	0.111	0.124	0.241	0.479

If H_0 is correct then Q should be (approximately) $\chi^2(3)$.
(3 = the number of categories -1.)

Mendels hypothesis seems ok

The observed test statistic 0.47 is compatible with H_0.



Comparing distributions

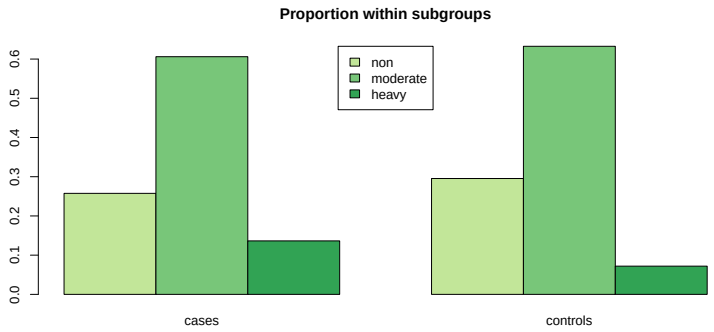
A case control study of coronary heart disease and drinking (none, moderate, heavy). Cases were matched on age, gender and smoking habits.

	non	moderate	heavy	sum
cases	34	80	18	132
controls	156	334	38	528
sum	190	414	56	660

Does drinking habits differ between cases and controls?
If they do not (H_0), their distributions should be close to

non	moderate	heavy
28.8% (190/660)	62.7% (414/660)	8.48% (56/660)

Visualizing the distributions



Are drinking categories equidistributed for cases and controls?

	non	moderate	heavy	Sum
observed cases	34	80	18	132
observed controls	156	334	38	528
sum	190	414	56	660
prop. (p=sum/660)	0.29	0.63	0.08	(1)
expected cases (132·p)	38.1	82.9	11.0	(132)
expected controls (528·p)	152.0	331.0	44.0	(527)
Q cases ((Obs.-Exp.)^2/Exp.)	0.43	0.10	4.4	Tot:
Q controls	0.11	0.026	1.1	6.2

The test statistic $Q = 6.2$ should be compared to a χ^2 with (rows-1)×(columns-1)=1*2=2 degrees of freedom.
$p = \text{Prob}(Q > 6.2) = 0.045$.

So the difference between cases and controls is statistically significant.

The large sample size gives this test a lot of power (ability to find differences).

Do not forget to look at the estimates!

	non	moderate	heavy
proportion cases	0.26	0.61	0.13
proportion controls	0.30	0.63	0.07
proportion total	0.29	0.63	0.08

Whether these differences are significant in any other sense is for the researcher to discuss.

χ^2 on the dabigatran data

The χ^2 test can also be applied to our dabigatran data.

It tests if the distribution of complications (bleeding/not) is the same for the two groups.

Output from my software:

```
#  Pearson's Chi-squared test with Yates'
#  continuity correction
#
#  data:  bord
#  X-squared = 11.05, df = 1, p-value = 0.0008869
```

(Recall that Fisher's exact test gave $p = 0.0004458$.)

Adjusting for multiple confounders

In observational studies we typically gather more information. E.g.

Ind.	Bleeding	DE Dose	Age	Gender	Weight	...
1	Yes	50	75	M	83	...
2	No	75	64	F	77	...
⋮	⋮	⋮	⋮	⋮	⋮	

When medicine is **not** randomized a simple cross tabulation analysis of 'Bleeding' versus 'DE Dose' is at risk of confounding.

One way to deal with this is logistic regression (more on that in a later lecture).

Adjusting for a confounder

Comparison of open surgery (OS) and percutaneous nephrolithotomy (PN) for removal of kidney stones.
(Data illustrates Simpson's paradox.)

			Adjusted for size			
	Total		Small stones		Large stones	
	OS	PN	OS	PN	OS	PN
Success	273	289	81	234	192	55
Failure	77	61	6	36	71	25
Odds (for success)	3.5	4.7	13.5	6.5	2.7	2.2
Odds ratio (OS / PN)	0.75		2.1		1.2	
(Percent success	78%	83%	93%	87%	73%	69%)

Here it seems like we should adjust for stone size.
(**The Mantel-Haenszel** test is a way to analyse several contingency tables.)