# Measures of Association

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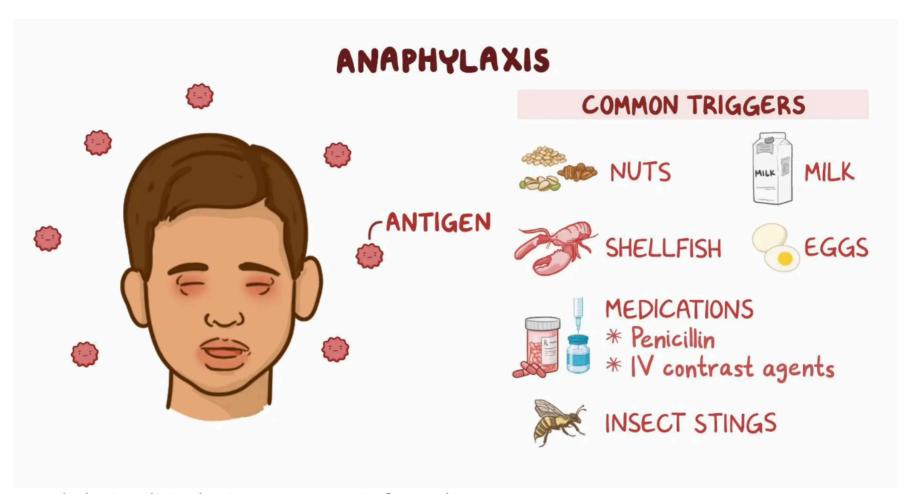
Updated: 2025-09-08

# **Expected Competencies**

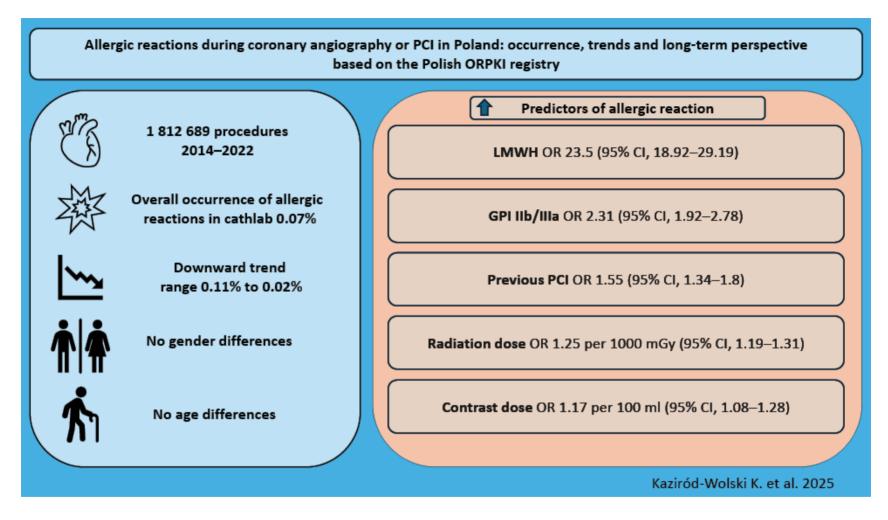
- Knows what are measures of frequency, association and effect.
- Know the difference between relative and absolute measures.
- Recognize and correctly interpret measures of association.

# **Objetives**

- To clarify potential misconceptions about measures of association.
- To introduce the counterfactual and potential outcomes framework.
- To identify advantages and adequate use of different measures.



Anaphylaxis: Clinical sciences. Osmosis from Elsevier



Allergic reactions during coronary angiography or PCI in Poland: Occurrence, trends, and long-term perspective based on the Polish ORPKI registry. Polish Heart Journal (Kardiologia Polska) Vol 83, No 3 (2025)

#### What's an Association?

Our common objective of epidemiologic research:

- ullet The effect of exposure X on the occurrence of outcome Y
  - But we can rarely observe or even estimate this effect directly.
- It involves the same people at the same time in contrasting exposures, which is impossible.
- We **observe an association** between the *Exposure* and *Outcome* among study subjects, which estimates a population association.

"The observed association will be a poor substitute for the desired effect, if it is a poor estimate of the population association, or if the population association is not itself close to the effect of interest." ME4 (2020)

#### What's an Effect?

Effect here means the end point of a causal mechanism, i.e., identifying the type of outcome that a cause produces.

EXAMPLE: "Liver cirrhosis is an effect of chronic excessive alcohol consumption".

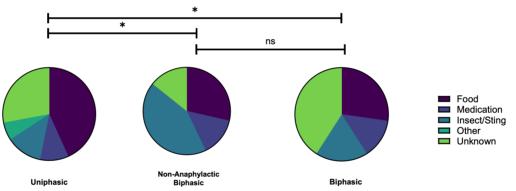
- This use of the term effect **merely** identifies liver cirrhosis as **one consequence of chronic excessive alcohol consumption**.
  - **Compared** to something else (e.g., Abstinence or another level of consumption).
  - Cirrhosis may Not be the only effect of of chronic excessive alcohol consumption.
  - May change across populations and or over time.

"An effect of some factor is thus relative to the outcomes, to the population, and to the time frame." [ME4 (2020)]

#### **Exposure vs Cause**

An exposure (usually denoted as X) is a *potential* causal characteristic, "a factor that produces an outcome"

• Could be the *sole* or *compounded* cause <sup>1</sup> of an Outcome. Can be a behavior, a treatment/intervention, a social condition, a health condition, a genetic trait...

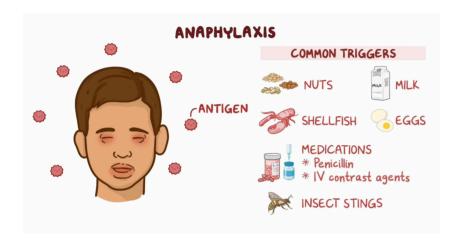


**Fig. 1** Reaction profiles. The proportion of anaphylactic triggers across the responder groups is represented by pie charts. The reaction profiles significantly differed between uniphasic responders and non-anaphylactic biphasic and biphasic responders (p=0.0179 and p=0.0375, 2-way ANOVA with Tukey's Multiple Comparisons test). Reaction profiles were comparable between non-anaphylactic biphasic and biphasic responders (p=0.8665, 2-way ANOVA with Tukey's Multiple Comparisons test)

<sup>1</sup> For more on Sufficient Component Causal Framework and Bradford Hill criteria, in this course we focus on potential outcomes and causal DAGs for approaches to causal inference.

• Ellis A.K, et al. (2025) Biphasic anaphylaxis in a Canadian tertiary care centre: an evaluation of incidence and risk factors from electronic health records and telephone interviews

# **Exposures vs Causes**



Anaphylaxis: Clinical sciences. Osmosis from Elsevier

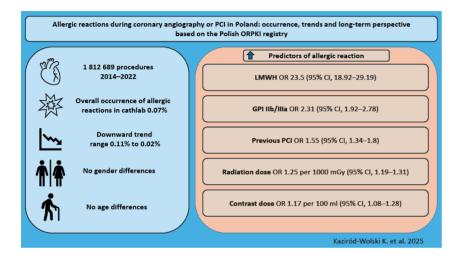
#### **Anaphylaxis Etiology**

"Common inciting sources may include **exposure** to certain medications, foods, or insect stings."

"Occasionally, the offending agent is not identified; these reactions are idiopathic anaphylaxis."

"The most common causes include bee stings, fire ant bites, latex, and foods (peanuts, tree nuts, fish, shellfish, milk, eggs, wheat, spelt, rye, barley, soy, red meat, and sesame)."

#### **Exposures vs Causes**

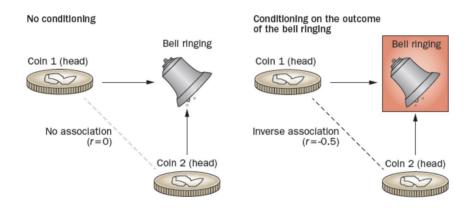


**Table 3.** Clinical and periprocedural factors affecting incidence allergic reaction

Variables	Univariable		
	OR (95% CI)	<i>P</i> -value	
Male sex	0.93 (0.83–1.04)	0.21	
Age, years	1 (0.99–1)	0.24	
Weight, kg	1 (1–1)	0.49	
Diabetes	0.87 (0.75–1)	0.06	
Previous stroke	1.14 (0.83–1.57)	0.42	
Previous myocardial infarction	1.27 (1.12–1.44)	< 0.001	
Previous PCI	1.37 (1.22–1.55)	< 0.001	
Previous CABG	0.89 (0.68-1.16)	0.37	
Smoking status (active)	0.8 (0.68-0.94)	0.01	
Arterial hypertension	0.96 (0.85-1.09)	0.56	
Kidney disease	0.68 (0.5-0.92)	0.01	
COPD	0.46 (0.27-0.78)	0.003	
ACS (yes/no)	1.29 (1.15–1.45)	< 0.001	
UFH during angiogram	0.67 (0.52-0.87)	0.003	
LMWH during angiogram	0.53 (0.2-1.41)	0.21	
Cardiac arrest at baseline	1.24 (0.62-2.48)	0.55	
Direct transport to cathlab	0.64 (0.41-0.99)	0.045	
UFH	0.34 (0.3-0.39)	< 0.001	
LMWH	49.18 (43.85-55.15)	< 0.001	

#### Not every Association between Exposure an Outcome is "Causal"

#### **Recall:**



At the population level, we assess the effects with measures of occurrence and we **estimate** the associations by contrasting such measures of occurrence in the population.

### Absolute vs. Relative measures

- Absolute effect measures are **differences** in occurrence measures.
- Relative effect measures are **ratios** of occurrence measures.



#### **Null Value**

Absence of contrast on either the Absolute (Difference) and Relative (Ratio) Scales

$$R_{exp} - R_{Non-exp} = 0$$

and

$$\left(rac{R_{exp}}{R_{Non-exp}}
ight)=1$$

### Absolute vs. Relative measures

- Absolute effect measures are **differences** in occurrence measures.
- Relative effect measures are **ratios** of occurrence measures.



Sample	Outcome	Outcome	Risk Among	Risk Among Non-	Risk	Risk
	+	-	Exposed	Exposed	Difference	Ratio
63	9/36	6/27	0.25	0.22	0.03	1.12

Key elements: Exposure, Outcome & Measures of Occurrence!

#### The 2x2 Table

• A summary table of observations.

	Outcome	No.Outcome	Total
Exposed	A	В	A+B
No Exposed	С	D	C+D
Total	A+C	B+D	A+B+C+D

#### The 2x2 Table

#### From the previous example:

	Outc		
	Dead	Alive	Total
Exposed	9	27	36
No Exposed	9	18	27
Total	18	45	63

#### The measures are:

Sample	Outcome	Outcome	Risk Among	Risk Among Non-	Risk	Risk
	+	-	Exposed	Exposed	Difference	Ratio
63	9/36	6/27	0.25	0.22	0.03	1.12

Key elements: Exposure, Outcome & Measures of Occurrence!

# Absolute measures: Risk Differences

The RD provides the absolute change in risk

- Indicates how much of the effect is attributable to exposure
- It does not provide information about the magnitude of the shift on the estimates 1 to 16% or 81 to 96%?
- Clinical vs statistical importance
- Public Health Relevance

# Relative Measures: Risk Ratios

- Relative measures are popular and practical
- Easier to obtain
  - Dichotomous outcomes!
- Useful in both causal inference and prediction

#### **Interpretability?**

• RR's magnitude change according to the coding scheme

Example: (0 to 1)  $\neq$  (2 to 1)

# Why not both?

• It's possible to see a reduction in absolute estimates, but an increase in relative measures (and vice versa)

These are complimentary estimators! Both tell you something different about the data

 In fact, STROBE and CONSORT guidelines now advise researchers to publish both measures

# 695 reactions vs 482 = 29% increase vs 20 more cases per 1000 people?

In an epidemiological utopia, researchers would run the model of their choice, obtain relative and absolute estimates, and publish these along with the baseline/background risk

But we live in the real world, so we're more likely to encounter...

### ...Odds ratios

$$Odds = \left(\frac{P}{1-P}\right)$$

The Odds ratio is the relative contrasts of the Odds among Exposed and the Odds among Non-Exposed

$$OddsRatio = \left(rac{\left(rac{P_{exp}}{1-P_{exp}}
ight)}{\left(rac{P_{No-exp}}{1-P_{No-exp}}
ight)}
ight)$$

### ...Odds ratios

We are already aware of some key problems with odds and therefore odds ratios

- They overestimate risks
  - $\circ$  While probabilities are bounded [0, 1], **odds** can range from 0 to  $\infty$
- They're not intuitive (except as an approximation of the risk ratio <sup>1</sup>)
- And most of the time we care about probabilities, not odds

When probability is small (<0.10) or given the study design (e.g., case-cohorts) with rare outcomes

# Risk Differences and Risk Ratios

Sample	Risk Among Exposed	Risk Among Non- Exposed	Risk Difference	Risk Ratio	Odds Ratio
63	0.25	0.22	0.03	1.12	1.17
63	0.17	0.15	0.02	1.12	1.15
630	0.017	0.015	0.002	1.12	1.15
630	0.25	0.22	0.03	1.12	1.17

#### **Null Value**

Absence of contrast on either the Absolute (Difference) and Relative (Ratio) Scales

$$R_{exp}-R_{Non-exp}=0$$
 ; or  $\left(rac{R_{exp}}{R_{Non-exp}}
ight)=1$ 

#### Example:

Sample	Outcome	Outcome	Risk	Risk Non-	Risk	Risk	Odds
	+	-	Exposed	Exposed	Difference	Ratio	Ratio*
63	4/36	3/27	0.11	0.11	0	1	1

- Null Value (Absolute) = (4/36) (3/27) = 0
- Null Value (Relative) = (4/36) / (3/27) = 1

#### **Note**

- 1. When the RR is above or below the null (>1 or <1) the ORs is FURTHER away form the Null.
- 2. When the absolute risk in each exposure groups are high, the OR will considerable overestimate the RR.

• Generating 10 data-sets with the *same* structure.

Verification of "mean" values across datasets.

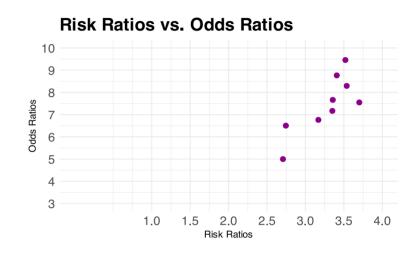
```
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```

#### Using "hand calculations" formulas.

```
# Using lapply
tabs <- lapply(1:10,FUN=function(x) table(e[,x],y[,x]))
# With a loop create an empty data frame with the right dimensions
ests <- as.data.frame(matrix(NA,nrow=10,ncol=3))
# Name columns
names(ests) <- c("RD","RR","OR")

# Loop it up
for (i in 1:10) {
    x <- tabs[[i]]
    ests[i,1] <- x[2,2]/sum(x[2,]) - x[1,2]/sum(x[1,]) #RD
    ests[i,2] <- (x[2,2]/sum(x[2,]))/ (x[1,2]/sum(x[1,])) #RR
ests[i,3] <- (x[2,2]/sum(x[2,1]))/ (x[1,2]/sum(x[1,1])) #OR
}</pre>
```

RD	RR	OR
0.43	3.70	7.55
0.36	2.71	5.00
0.43	3.35	7.17
0.50	3.52	9.46
0.47	3.54	8.29
0.45	3.36	7.66
0.49	3.41	8.77
0.43	2.74	6.50
0.52	3.73	10.50
0.43	3.17	6.76



#### Using regressions <sup>1</sup> to obtain the estimates (*Rare-ish* outcomes)

```
## E Y

## Min. :0.0000 Min. :0.00

## 1st Qu.:0.0000 1st Qu.:0.00

## Median :1.0000 Median :0.00

## Mean :0.5025 Mean :0.11

## 3rd Qu.:1.0000 3rd Qu.:0.00

## Max. :1.0000 Max. :1.00
```

<sup>1</sup> Some of you may have advanced knowledge on regression analysis but since we have not explained it during the course this resource is only for illustration purposes.

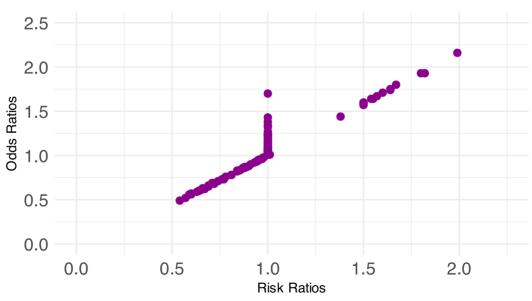
```
RRs <- sapply(sim100,FUN=function(x) {
    results <- logbin(Y ~ E , data=x)$coef
    return(round(exp(results[names(results)=="E"]),2))
})

ORs <- sapply(sim100,FUN=function(x) {
    results <- glm(Y ~ E , family="binomial", data=x)$coef
    return(round(exp(results[names(results)=="E"]),2))
})

sim_RRs<-round(quantile(RRs,probs = c(0.05,0.5,0.95)),2)
sim_ORs<-round(quantile(ORs,probs = c(0.05,0.5,0.95)),2)</pre>
```

```
## 5% 50% 95%
## sim_RRs 0.63 1.00 1.80
## sim_ORs 0.59 1.02 1.93
```

#### Risk Ratios vs. Odds Ratios



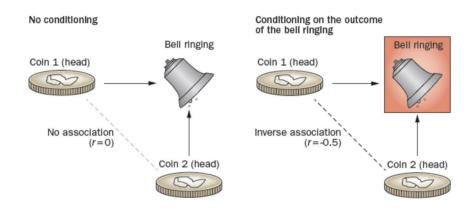
#### **Associations Vs Causes**

Pearl (2000) uses Pr(Y=y|SET[X=x]) to define the probability of an event if the condition X=x were **enforced uniformly** over a population.

- The key to this definition: it involves intervention, not observation.
- Measures of effect can be built based on  ${\bf SET}$  notation by creating contrasts of probabilities (or risks) across different X values.
- The What If? Book, express the same notion with  $Y^{X=x}$ .

### **Associations Vs Causes**

#### **Recall:**



Be aware of the difference!

#### Measures of causal effect

Measures of causal effect require a contrast of two counterfactual quantities:

• 
$$Pr(Y_i = y_i | SET[X_i = 1]) - Pr(Y_i = y_i | SET[X_i = 0])$$

Measures of association involve a contrast of two observed quantities:

• 
$$Pr(Y_i = y_i | X_i = 1) - Pr(Y_i = y_i | X_i = 0)$$

### Who are we interested in?

**Target Population:** The group of people about which the scientific or public health question is asked, in the **relevant etiologic time period.** 

#### "Target Population" in Encyclopedia of Biostatistics, 2005 [Sander Greenland]

"The concept of a target population is an informal one, sometimes defined as "the population about which information is wanted" [1] or the "totality of elements which are under discussion and about which information is desired" [4] ...The word "target" emphasizes, however, that this population is not necessarily the same as the one that we end up sampling. The latter population is sometimes called the sampled population [1, 4] or (in epidemiology) the source population [6]". [1] Cochran, W.G. (1977). Sampling Techniques, 3rd Ed. Wiley, New York. [4] Mood, A.M., et al. (1974). Introduction to the Theory of Statistics. McGraw-Hill, New York. [6] Rothman, K.J. & Greenland, S. (1997). Modern Epidemiology, 2nd Ed. Lippincott, Philadelphia.

### **Potential Outcomes Framework**

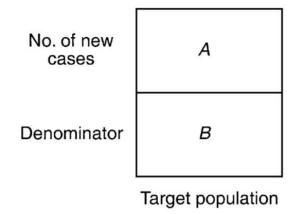
We are interested in the effect of exposure  $\left(A=1\right)$  on the occurrence of disease  $\left(Y=1\right)$ 

- Suppose everyone in the **target population** of inference is unexposed (A=0) and we can observe the distribution of Y in the population.
- ullet We would like to also observe the distribution of Y had these same people been all exposed (A=1)
  - This is "counter-to-fact", and we call this condition the counterfactual
  - Each individual has their own counterfactual exposure
  - (what would have happened to me if ...)

### **Potential Outcomes Framework**

- We can never observe both conditions in the same population (or individual).
  - $\circ$  That is, we cannot observe the distribution of disease under A=1 and A=0 within the same time period in the same cohort.
  - Thus we need to make an estimate under the condition we do not observe.
- To do so, we use a **substitute population**.
- Our goal is to choose a substitute population that will best mimic what would have happened to the target population had they experienced the other exposure condition.

## Estimating Causal Effects <sup>1</sup>



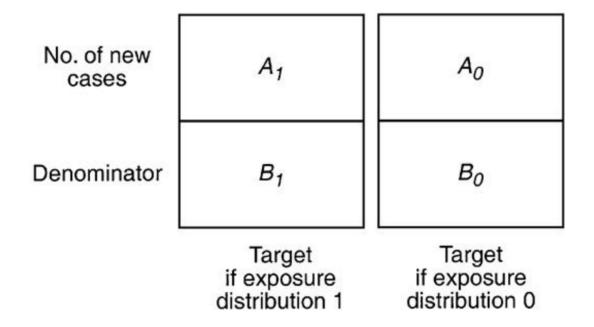
- If B = people at risk at the *beginning of the period*  $^2$  = **incidence proportion**, **average risk**.
- If B = person-time at risk during the period, R = person-time incidence rate.
- If B = people who do NOT get disease by the end of the period, R = incidence odds.

<sup>&</sup>lt;sup>1</sup> International Journal of Epidemiology, Volume 31, Issue 2, April 2002, Pages 422–429, https://doi.org/10.1093/intjepid/31.2.422

<sup>&</sup>lt;sup>2</sup> And all individuals are followed throughout the etiologic time period.

### Define the causal effect

Define the counter-to-fact condition and outcome in the target:



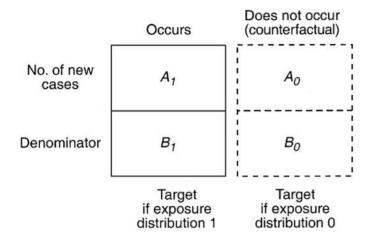
Only possible to, at best, observe one of these conditions.

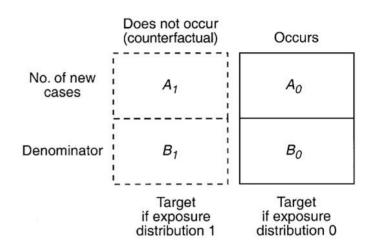
## Define the causal effect

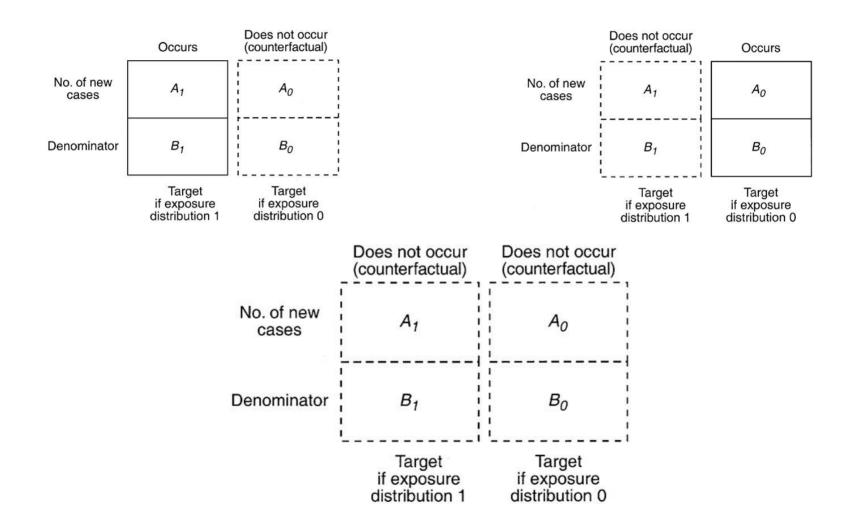
Let 
$$R_1=A_1/B_1$$
 and  $R_0=A_0/B_0$ 

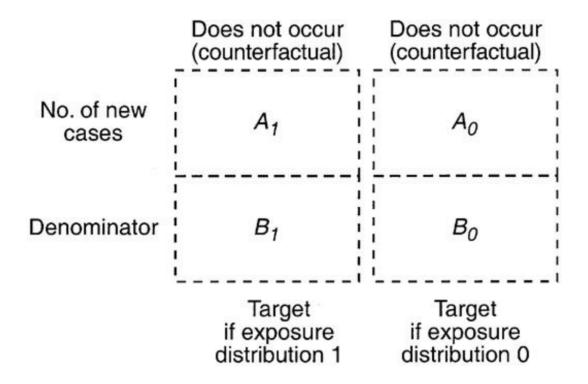
- $R_1-R_0$  is the causal **difference measure** and
- $R_1/R_0$  is the causal ratio measure
- Both of these are causal contrasts (measures of effect).
- Here, the only possible reason for a difference between  $R_1$  and  $R_0$  is due to **exposure** because we are contrasting the **exact same people over the exact same time period**.
- But, we cannot observe the causal contrast because we cannot observe both conditions.

We require a substitute with observable information!

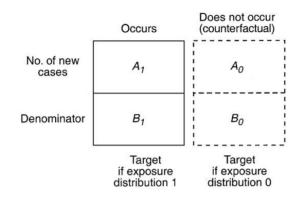








"Both  $R_1$  and  $R_0$  are counterfactual disease frequencies, both are hypothetical alternatives to the actual disease frequency that occurs under the actual exposure distribution (which is neither exposure distribution 1 nor 0), and therefore neither R1 nor R0 can occur and be observed." Maldonado & Greenland 2002



The impossibility of observing both halves of the causal contrast leads to the idea of substitute populations.

#### These are often:

- Different people observed during the same etiologic time period.
- The same people observed over two different time periods (case-crossover design).

In a substitute population under **exposure distribution** 1, let

- ullet  $C_1$  be the name for the numerator of the disease-frequency measure,
- $D_1$  be the denominator (number of people or amount of person-time at risk).

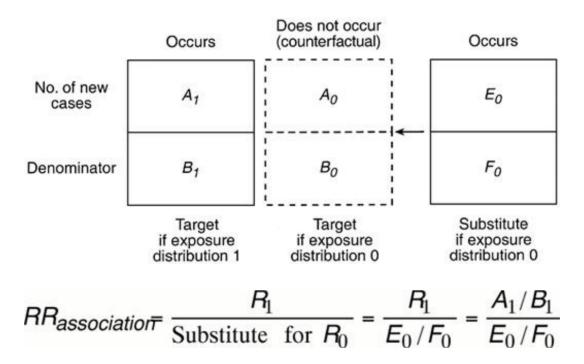
In a substitute under **exposure distribution** 0, let

- $E_0$  be the numerator,
- $F_0$  be the denominator.

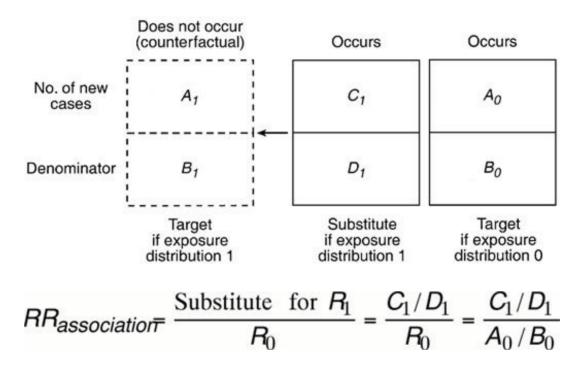
Exposure = 1, 
$$C_1/D_1$$

Exposure = 0, 
$$E_0/F_0$$

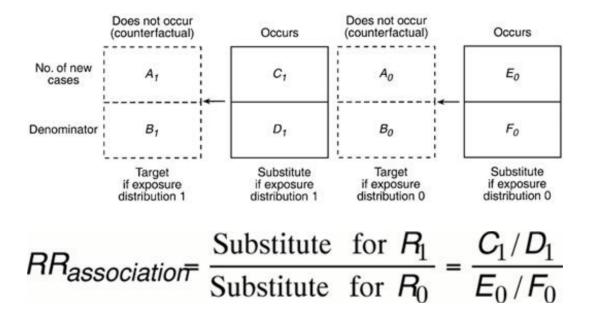
Target experiences exposure distribution 1



Target experiences exposure distribution 0



Target experiences neither exposure distribution 1 or 0



### **Other Notation Used**

- ullet Greenland employs a probabilistic model of disease such that each individual i has a risk  $r_{1_i}$  of disease when E=1 and a risk of  $r_{0_i}$  when E=0.
- ullet Survival probabilities:  $S_{1_i}=1-r_{1_i}$  and  $S_{0_i}=1-r_{0_i}$
- ullet Odds:  $w_{1_i}=r1_i/s_{1_i}$  and  $w_{0_i}=r_{0_i}/s_{0_i}.$ 
  - Only defined when survival probabilities are not equal to zero.

#### **Notation Used**

- The effect of exposure on the risk of an individual can be measured in terms of the risk difference  $r_{1_i}-r_{0_i}$ , risk ratio  $r_{1_i}/r_{0_i}$ , or the risk-odds ratio  $w_{1_i}/w_{0_i}$
- The ratios will be undefined if the risk in the exposed group is 0 and
- The risk-odds ratio will be undefined if either survival probability is 0

## **Other Notation Used**

In a cohort with  $N_1$  E+ individuals and  $N_0$  E- individuals:

	E+	E-
D+	$A=\Sigma_1 r_{1i}$	$B=\Sigma_0 r_{0i}$
D-	$C=\Sigma_1 s_{1i}$	$D=\Sigma_0 s_{0i}$
Total	$N_1$	$N_0$

- ullet Incidence proportions:  $A/N_1$  and  $B/N_0$ , interpretable as average risks in their respected groups
- ullet Incidence odds: A/C and B/D, interpretable as ratios of the average risk to the average survival probabilities.

# Defining the counterfactual

- Assuming no confounding <sup>1</sup>
- Had the **exposure** been absent from the E+ group,
- The average risk would have been **the same** among the sub-cohorts that were in fact exposed and unexposed.

$$\left(rac{\sum_1 r_{0i}}{N_1}
ight) = \left(rac{\sum_0 r_{0i}}{N_0}
ight)$$

<sup>&</sup>lt;sup>1</sup> More on confounding next lecture.

#### **Risk Difference**

Thus: The risk difference is interpretable as both:

1) the absolute change in the average risk of the exposed sub-cohort produced by exposure,

$$\left(rac{\sum_1 r_{1i}}{N_1}
ight) - \left(rac{\sum_1 r_{0i}}{N_1}
ight)$$

2) the average absolute change in risk produced by exposure among exposure individuals

$$\left(rac{\sum_1(r_{1i}-r_{0i})}{N_1}
ight)$$

Expressions 1 and 2, on: Interpretation and choice of effect measures in epidemiologic analyses. S. Greenland (1987)

#### **Risk Ratio**

The incidence proportion ratio is given by

$$\left(\frac{A}{N_1}\right) / \left(\frac{B}{N_0}\right)$$

The risk ratio is interpretable as:

1) the **proportionate change in the average risk** of the exposed subcohort produced by exposure,

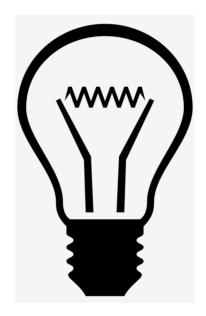
$$\left(rac{\sum_1 r_{1i}}{N_1}
ight)/\left(rac{\sum_0 r_{0i}}{N_0}
ight)=\left(rac{\sum_1 r_{1i}}{N_1}
ight)/\left(rac{\sum_1 r_{0i}}{N_1}
ight)$$

It **is not interpretable** as the average proportionate change in risk produced by exposure among exposed individuals:

$$\left(rac{\sum_1 (r_{1i}/r_{0i})}{N_1}
ight)$$

Expressions 3 and 4, on: Interpretation and choice of effect measures in epidemiologic analyses. S. Greenland (1987)

## **Incidence Proportion Ratio**



However, **if the individual risk ratios are all equal** then the ratio of the average risks across exposure will be equal to the average of the individual risk ratios

#### A note on Risk Ratios

Risk Differences have a symmetric range [-1, 1]

**But** the risk Ratios have an *asymmetric* range:

- From 0 to 1, below the null
- From 1 to infinity above the null

This presents a challenge in the interpretation...

What's more "impressive" (RR = 2) or (RR= 0.2)?...

#### What's more "impressive" a (RR = 2) or a (RR = 0.2)?

Two ways to find out:

**Simple:** the reciprocal of the value below the null = 1/0.2 = 5, since 5 > 2, then

a RR of 0.2 is of larger magnitude (further away form the null) than a RR = 2

**Elaborated:** Take the absolute values of the natural logarithm (log or ln) of each value:

- log(2) = 0.693
- log(0.2) = -1.609

In absolute terms,  $|\log(0.2)| > |\log(2)| = |-1.609| > |0.693|$ 

- Regression models for ratio measures generally operate on the log-scale (that's why we exponentiate to provide estimates and graph on the log scale).
- Note: Recall that on the log10 or ln scale, the null for a ratio measure is 0, not 1 because log(1) = ln(1) = 0.
- Try it with 3 and 0.3, and with 5 and 0.5 and see what happens! :)

The incidence odds ratio is given by:

$$\left(\frac{A}{C}\right)/\left(\frac{B}{D}\right)$$

Thus, the odds ratio is interpretable as:

1) the proportionate change in the incidence odds in the exposed subcohort produced by exposure,

$$\left(rac{\sum_1 r_{1i}}{\sum_1 s_{1i}}
ight) ig/ \left(rac{\sum_0 r_{0i}}{\sum_0 s_{0i}}
ight)$$

$$\left(rac{\sum_1 r_{1i}}{\sum_1 s_{1i}}
ight) / \left(rac{\sum_1 r_{0i}}{\sum_1 s_{0i}}
ight)$$

It **is not interpretable** as the proportionate change in the average odds in the exposed produced by exposure:

• Furthermore, neither of the last two expressions is equivalent to the average of the individual odds ratios among the exposed

$$\left(rac{\sum_1 (w_{1_i}/w_{0_i})}{N_1}
ight)$$

• The incidence odds ratio (**that we calculate**) lacks any simple interpretation in terms of exposure effect on the average risk or odds, or average exposure effect on individual risk or odds.

The incidence odds do not equal the simple averages of the risk odds:

This severely handicaps the interpretability of measures based on the incidence odds.

- It is not a measure of average causal effect (the RR and RD are) (Greenland 1987)
- Cannot be relied upon to reveal confounding (Greenland et al., 1999)

• If the individual ORs are all equal (which is the assumption made by a logistic model), then the ratio of the average odds will equal the average of the individual odds ratios.

But, the incidence odds ratio will need not equal that value!

- ullet For example: Define a population where 10% of people have  $r_{1i}=$  0.60 and
- ullet  $r_{0i}=$  0.20 and 90% of the people have  $r_{1i}=$  0.035 and  $r_{0i}=$  0.006
- Here, the individual ORs = 6.0 for every individual
  - the average of the individual ORs = 6.0
- Also, the ratio of the average odds equals 6.0 as well

But, the incidence odds ratio is equal to 3.9

Want to give it a try and calculate it?

- Because of this fact, the crude odds ratio can be smaller than any of the stratum-specific odds ratios, even if confounding is entirely absent
- This paradoxical behaviour will not occur with the risk difference or the risk ratio
- Unless, equal to the null (when OR = RR = 1) the OR will almost always be further away from the null than RRs.
  - 2nd exception is when OR = RR = 0, as would occur when risk in the exposed is 0 (zero), and risk in the unexposed is for example 0.6. In this case, the risk ratio is 0 and the OR is odds(0)/odds(0.6), which is also 0.

More on Epidemiology by design by Daniel Westreich

## Odds difference?

```
▲ !! Nope, nope, nope!!!!!!! ▲ !!
```

X!! We never do this.!! X

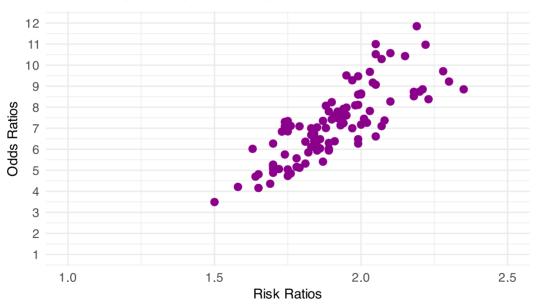
#### Some simulated Examples - Common Outcome

```
set.seed(7042025)
vea.dat1 <- function(n) {</pre>
          E <- rbinom(n,1,0.55) #parameters for E
          Y \leftarrow ifelse(E==1, rbinom(n, 1, 0.85), rbinom(n, 1, 0.45)) #parameters for Y
    return(data.frame(E=E,Y=Y)) #ask to return a data set with those parameters
sim100 <- lapply(1:100,FUN=function(x) yea.dat1(400))</pre>
RRs <- sapply(sim100, FUN=function(x) {
   results <- logbin(Y ~ E , data=x)$coef
   return(round(exp(results[names(results)=="E"]),2))
 })
ORs <- sapply(sim100, FUN=function(x) {
   results <- glm(Y ~ E , family="binomial", data=x)$coef
   return(round(exp(results[names(results)=="E"]),2))
 })
sim_RRs1 < -round(quantile(RRs, probs = c(0.05, 0.5, 0.95)), 2)
sim_ORs1 < -round(quantile(ORs, probs = c(0.05, 0.5, 0.95)), 2)
```

### Some simulated Examples - Common Outcome

## 5% 50% 95% ## sim\_RRs1 1.65 1.90 2.22 ## sim\_ORs1 4.73 7.12 10.52

#### Risk Ratios vs. Odds Ratios



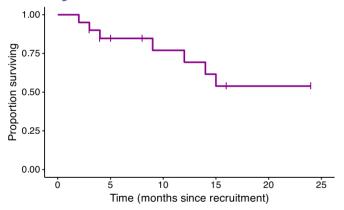
# **Key Takeaways**

Only incidence difference and incidence ratios possess direct interpretations as measures of impact on average risk/hazard

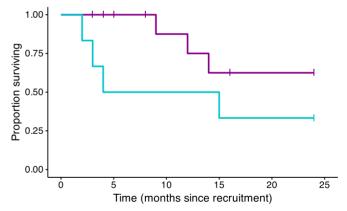
Consequently, odds ratios are useful only when they serve as incidence ratio estimates

#### **Incidence Rate Differences and Ratio**

#### From everyone's time...



#### ... To contrast outcomes and time



#### **Incidence Rate Differences and Ratio**

- ullet Incidence rate of outcome Y when X=1 is IR(Y=1|X=1)=IR(Y|X=1)
- ullet Incidence rate of outcome Y when X=0 is IR(Y=1|X=0)=IR(Y|X=0)
- Incidence rate difference:
  - ∘ Range: -∞ to ∞
  - Null value: 0

- Incidence rate ratio:
  - ∘ Range: 0 to ∞
  - Null value: 1

#### Relationship among Risk, Odds, and Incidence Rates

Question: If we calculated the risk ratio, odds ratio and rate ratio, which will be closest to the null? Furthest from the null?

#### Notation:

- R = Incidence Proportion ("Risk"),
- S=1-R ("Survival Proportion"),
- I = Incidence Rate
- T = interval length
- i = 1 if exposed; i = 0 if unexposed

#### Relationship among Risk, Odds, and Incidence Rates

Relations among relative risks

• In a closed population where the population at risk declines only slightly over the interval (implying that R must be small and S is close to 1):  $R\cong I\Delta T\cong R/S$ 

This implies:

$$\left(rac{R_1}{R_0}
ight)\cong \left(rac{I_1\Delta T_1}{I_0\Delta T_0}
ight)\cong \left(rac{I_1}{I_0}
ight)\cong \left(rac{R_/S_1}{R_0/S_0}
ight)$$

[Numerators] Holds if  $R_1$  and  $R_0$  are small enough so that  $S_1$  and  $S_0$  are close to 1

[Two Denominators on the right] Holds if exposure only has negligible effects on the persontime at risk

# Relationship among relative risks

If exposure causes the outcome, then R1>R0 and S1<S0.</li>

$$1<\left(rac{R_1}{R_0}
ight)<\left(\left(rac{R_1}{R_0}
ight) imes\left(rac{S_0}{S_1}
ight)
ight)=\left(rac{R_1/S_1}{R_0/S_0}
ight)$$

• If exposure prevents the outcome, R1 < R0 and S1 > S0, such that:

$$1 > \left(rac{R_1}{R_0}
ight) > \left(\left(rac{R_1}{R_0}
ight) imes \left(rac{S_0}{S_1}
ight)
ight) = \left(rac{R_1/S_1}{R_0/S_0}
ight)$$

In words: The odds ratio is further from the null than the risk ratio

## Relationship among relative risks

- ullet Now, if exposure is harmful (R1>R0) then we would ordinarily expect exposure to reduce the person-time at risk (T1< T0),
- and if exposure is preventive (R1 < R0) then we expect exposure to increase the persontime at risk (T1 > T0).
- Thus, when exposure is harmful:

$$1 < \left(rac{R_1}{R_0}
ight) \cong \left(rac{I_1 \Delta T_1}{I_0 \Delta T_0}
ight) < \left(rac{I_1}{I_0}
ight)$$

# Relationship among relative risks

And when exposure is preventive:

$$1 > \left(rac{R_1}{R_0}
ight) \cong \left(rac{I_1 \Delta T_1}{I_0 \Delta T_0}
ight) > \left(rac{I_1}{I_0}
ight)$$

• In words: We would **ordinarily** expect the risk ratio to be closer to the null than the rate ratio. Under further conditions, the rate ratio will be closer to the null than the odds ratio (Greenland and Thomas, 1982)

# Relationship among relative risks

#### Thus, we usually expect:

- Risk ratio nearest to the null
  - o *implicitly* suggesting all events occur at the end of follow up
- Odds ratio furthest from the null
  - implicitly suggesting all events occur at the beginning of follow up
- Rate ratio somewhere in between
  - allows event to occur at any point in time
  - 1 < Risk Ratio < Rate Ratio < Odds Ratio
- <sup>1</sup> More on Epidemiology by design by Daniel Westreich

### **Prevalence Ratios**

Recall that the prevalence odds is equal to the incidence rate multiplied by the average duration in a stationary, closed population. This implies:

$$POR = \left(\frac{PO_1}{PO_0}\right) = \left(\frac{I_1\overline{D_1}}{I_0\overline{D_0}}\right) = \left(\frac{I_1}{I_0}\right)$$

if the average duration of disease is unaffected by exposure.

## **Prevalence Ratios**

• If prevalence is low, then the prevalence ratio is approximately equal to the incidence rate multiplied by average duration. This implies:

$$PR = \left(rac{P_1}{P_0}
ight) \cong \left(rac{I_1\overline{D_1}}{I_0\overline{D_0}}
ight) = \left(rac{I_1}{I_0}
ight)$$

if the average duration of disease is unaffected by exposure

#### ...Still breathing?:)



#### **Key Points**

- "There was a decreasing incidence of allergic reactions overall, in both sexes and all age subgroups. The decrease in frequency did not depend on age or sex."
- "Our multivariable logistic regression analysis showed that GPI IIb/IIIa, LMWH, previous PCI, contrast dose, and x-ray dose were independent predictors of allergic reaction occurrence".

<sup>&</sup>lt;sup>1</sup> Allergic reactions during coronary angiography or PCI in Poland: Occurrence, trends, and long-term perspective based on the Polish ORPKI registry. Polish Heart Journal (Kardiologia Polska)Vol 83, No 3 (2025)

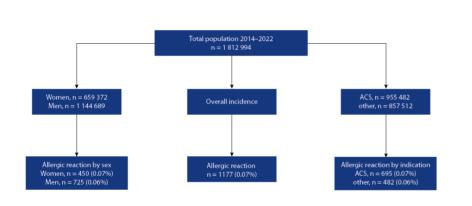
#### **Illustrated Example:**

**Table 3.** Clinical and periprocedural factors affecting incidence allergic reaction

Variables	Univariable	
	OR (95% CI)	<i>P</i> -value
Male sex	0.93 (0.83–1.04)	0.21
Age, years	1 (0.99–1)	0.24
Weight, kg	1 (1–1)	0.49
Diabetes	0.87 (0.75-1)	0.06
Previous stroke	1.14 (0.83-1.57)	0.42
Previous myocardial infarction	1.27 (1.12–1.44)	< 0.001
Previous PCI	1.37 (1.22–1.55)	< 0.001
Previous CABG	0.89 (0.68-1.16)	0.37
Smoking status (active)	0.8 (0.68-0.94)	0.01
Arterial hypertension	0.96 (0.85-1.09)	0.56
Kidney disease	0.68 (0.5–0.92) 0.01	
COPD	0.46 (0.27-0.78)	0.003
ACS (yes/no)	1.29 (1.15–1.45)	< 0.001
UFH during angiogram	0.67 (0.52-0.87)	0.003
LMWH during angiogram	0.53 (0.2-1.41)	0.21
Cardiac arrest at baseline	1.24 (0.62-2.48)	0.55
Direct transport to cathlab	0.64 (0.41-0.99)	0.045
UFH	0.34 (0.3-0.39)	< 0.001
LMWH	49.18 (43.85–55.15)	< 0.001

<sup>&</sup>lt;sup>1</sup> Allergic reactions during coronary angiography or PCI in Poland (2025)

#### Illustrated Example... Let's reproduce the analysis



	Outcome		
	Anaphylaxis	Non.Anaphylaxis	TOTAL
Male	687	1144002	1144689
Female	462	688910	659372
Total	1149	1832912	1804061

Allergic reactions during coronary angiography or PCI in Poland. Polish Heart Journal (Kardiologia Polska)Vol 83, No 3 (2025)

#### Illustrated Example... Let's reproduce the analysis from Figure 1

```
pcidat1<-c(687, 1144002, 462, 658943)
pciOR1<- epi.2bv2(pcidat1, method = "cross.sectional")</pre>
pciOR1$tab
                                                                Prev risk *
##
               Outcome +
                            Outcome -
                                           Total
## Exposed +
                     687
                              1144002
                                          1144689
                                                        0.06 (0.06 to 0.06)
## Exposed -
                     462
                               658943
                                           659405
                                                        0.07 (0.06 to 0.08)
## Total
                    1149
                              1802945
                                          1804094
                                                        0.06 (0.06 to 0.07)
```

#### "No gender differences" (???)

```
round(pciOR1$massoc.detail$OR.strata.wald, 2)

## est lower upper
## 1 0.86 0.76 0.96
```

#### Illustrated Example... Let's reproduce the analysis

```
pciOR1<- epi.2bv2(pcidat1, method = "cross.sectional")</pre>
pciOR1
                                    Total
                                                        Prev risk *
##
             Outcome +
                        Outcome -
## Exposed +
                                    1144689
                                                 0.06 (0.06 to 0.06)
                   687
                        1144002
           462 658943 659405
## Exposed -
                                                 0.07 (0.06 to 0.08)
## Total
                                                 0.06 (0.06 to 0.07)
         1149 1802945
                                    1804094
##
## Point estimates and 95% CIs:
## Prev risk ratio
                                            0.86 (0.76, 0.96)
## Prev odds ratio
                                      0.86 (0.76, 0.96)
                             -0.01 (-0.02, -0.00)
## Attrib prev in the exposed *
## Attrib fraction in the exposed (%) -16.74 (-31.35, -3.76)
## Attrib prev in the population *
                                          -0.01 (-0.01, 0.00)
## Attrib fraction in the population (%) -10.01 (-18.04, -2.52)
## Uncorrected chi2 test that OR = 1: chi2(1) = 6.635 Pr>chi2 = 0.010
## Fisher exact test that OR = 1: Pr>chi2 = 0.011
## Wald confidence limits
## CI: confidence interval
## * Outcomes per 100 population units
```

How do I obtain the  $\chi^2$  ?

#### Illustrated Example... What's in the Table 3?

```
pcidat1a<-c(725, 1144689, 450, 659372)
pciOR1a<- epi.2bv2(pcidat1a, method = "cross.sectional")</pre>
pciOR1a
##
                          Outcome - Total
                                                             Prev risk *
               Outcome +
## Exposed +
                    725 1144689 1145414
                                                     0.06 (0.06 to 0.07)
## Exposed -
            450 659372 659822
                                                     0.07 (0.06 to 0.07)
## Total
                                                     0.07 (0.06 to 0.07)
          1175 1804061
                                       1805236
##
## Point estimates and 95% CIs:
## Prev risk ratio
                                               0.93 (0.83, 1.04)
                                         0.93 (0.83, 1.04)
## Prev odds ratio
## Attrib prev in the exposed *
                               -0.00 (-0.01, 0.00)
## Attrib fraction in the exposed (%) -7.75 (-21.19, 4.20)
## Attrib prev in the population * -0.00 (-0.01, 0.00) ## Attrib fraction in the population (%) -4.78 (-12.67, 2.55)
## Uncorrected chi2 test that OR = 1: chi2(1) = 1.548 Pr>chi2 = 0.213
## Fisher exact test that OR = 1: Pr>chi2 = 0.214
## Wald confidence limits
## CI: confidence interval
## * Outcomes per 100 population units
```

#### What's clear and what's not here?

#### **Results**

of significance of the difference in years (Table 2). Among clinical and pharmacological factors, the risk of allergic reactions was increased by a history of myocardial infarction, a history of coronary artery by-pass grafting, a diagnosis of ST-segment elevation myocardial infarction (STEMI), a diagnosis of ACS, the use of IIb/IIIa glycoprotein inhibitor (GPI IIb/IIIa), low molecular weight heparin (LMWH), and earlier PCI. The risk of allergic reaction was reduced by active smoking, renal disease, chronic obstructive pulmonary disease, UFH, and performing coronary angiography only (Table 3).

Allergic reactions during coronary angiography or PCI in Poland. Polish Heart Journal (Kardiologia Polska)Vol 83, No 3 (2025)

#### **Methods**

#### Statistical analysis

In this study, nominal variables were represented as percentages along with their respective counts. The temporal variation in the proportion of allergic reactions was analyzed using the Cochran–Armitage test for trend. The logistic regression model was employed to identify the predictors of periprocedural allergic reactions. All potential demographic, baseline, and procedural characteristics were incorporated into a multivariable model if the *P*-value in a univariable model was less than 0.2 or if they were considered to have clinical significance. The final model was derived by minimizing the Akaike information criterion. The robustness of the model was validated using bootstrap resampling. To evaluate the presence of multicollinearity, variance inflation factors were analyzed. The results were expressed as odds ratios with 2-sided 95% confidence intervals. The entire statistical analysis was conducted using R software (version 4.3.1)

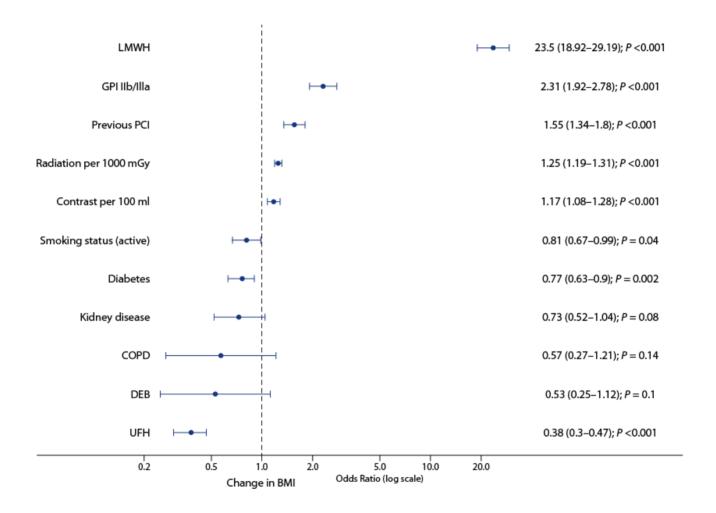
#### **Illustrated Example**

**Table 3.** Clinical and periprocedural factors affecting incidence allergic reaction

Variables	Univariable	
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LMWH	49.18 (43.85–55.15)	<0.001

<sup>&</sup>lt;sup>1</sup> Allergic reactions during coronary angiography or PCI in Poland. (2025)

#### What's clear and what's not here?



### What would be the actual interpretation of these ORs?

#### **Illustrated Example**

pcidat2<-c(695, 955482, 482, 857512)

"When assessing the incidence of allergic reactions ..., there was a decrease in the frequency of allergic reactions regardless of acute coronary syndromes (ACS) (Table 1)." "...the risk of allergic reactions was increased by... a diagnosis of ACS"

```
pciOR2<- epi.2by2(pcidat2, method = "cross.sectional")</pre>
pciOR2
##
                          Outcome - Total
                                                             Prev risk *
               Outcome +
## Exposed +
                                                     0.07 (0.07 to 0.08)
                    695
                              955482 956177
## Exposed -
                                                     0.06 (0.05 to 0.06)
            482 857512 857994
## Total
          1177 1812994
                                                     0.06 (0.06 to 0.07)
                                       1814171
##
## Point estimates and 95% CIs:
## Prev risk ratio
                                               1.29 (1.15, 1.45)
## Prev odds ratio
                                               1.29 (1.15, 1.45)
## Attrib prev in the exposed * 0.02 (0.01, 0.02) ## Attrib fraction in the exposed (%) 22.71 (13.19, 31.19)
## Attrib prev in the population * 0.01 (0.00, 0.01)
## Attrib fraction in the population (%) 13.41 (7.26, 19.15)
## Uncorrected chi2 test that OR = 1: chi2(1) = 19.007 Pr>chi2 = <0.001
## Fisher exact test that 0R = 1: Pr > chi2 = <0.001
  Wald confidence limits
## CI: confidence interval
  * Outcomes per 100 population units
```

### What did the authors do with the covariates?



### When to use which measure of Association

- Research Question
- Public Health Relevance
- Intervention (Design of/ Intervenable exposure?)
- Study design
- Contrasts requires assignment to group, which requires measurement of group membership For example, measurement of exposure
  - What will happen if the exposure is measured poorly?
- Since risks are only well-defined within a specific time-period, state that time-period.

## Things to consider when measuring associations

- Be aware that in real life, we encounter:
  - Random Error
  - Systematic error
  - Competing Risks, Confounding, Selection Bias, Measurement Error
  - Methods related limitations
  - Clinical vs Statistical hurdles

## **Contrast (Association vs Impact)**

Most contrasts can be used to assess either:

- Association (which is agnostic on the question of causality)
- Impact (which is causal). The risk difference, for example.

Certain measures, however, are implicitly causal and (probably) shouldn't be used to merely describe an association.

Among these latter measures are number needed to treat, and attributable contrasts.

## Population Attributable Fraction (PAF)

A population attributable fraction (PAF) can be thought of as

- "The proportion of disease burden among the total population which is caused by the exposure".
- That definition is explicitly causal.
- PAF is implicitly causal.
- "Attribution" implies cause, though we can argue over usage.

Calculated as  $(P(Y) - P(Y^{x=0}))/P(Y)$  where,

- P(Y) is the risk of the outcome in the whole population,
- P(Y|X=0) is the risk of the outcome in the unexposed.

Note that because most outcomes are caused by more than one thing, the sum of PAFs can be (and often are) greater than 100%. Epidemiology by design by Daniel Westreich

#### **Example:**

- Exposure to TB is a necessary cause of active TB: by definition.
  - $\circ$  So from the above,  $P(Y_{TB}exposure = 0) = 0$ , and so PAF = 100%.
- But not everyone exposed to TB develops active TB. There are other causes. E.g., being immunocompromised.
  - RD for immunocompromised status > 0; PAF>0.
- $PAF_{TB-exposure} + PAF_{immunocompromised} > 100$

This concept can be tied to Rothman's causal pies model of causality Epidemiology by design by Daniel Westreich

### A measure related to PAFs

The **population attributable risk difference** is the difference between the risk of the outcome in the observed population and the risk of the outcome **if all exposure were removed**, that is:

$$P(Y) - P(Y^{x=0})$$

-The potential outcomes notation is *deliberate* here.

## Remember our anaphylaxis by sex example?

#### **Assuming a cohort design:**

```
pcidat3<-c(687, 1144002, 462, 658943)

pciOR3<- epi.2by2(pcidat3, method = "cohort.count")
kbl(pciOR3$massoc.summary, digits = 2)</pre>
```

var	est	lower	upper
Inc risk ratio	0.86	0.76	0.96
Inc odds ratio	0.86	0.76	0.96
Attrib inc risk *	-0.01	-0.02	0.00
Attrib fraction in exposed (%)	-16.74	-31.35	-3.76
Attrib inc risk in population *	-0.01	-0.01	0.00
Attrib fraction in population (%)	-10.01	-18.04	-2.52

#### Attributable Risk $\cong$ Risk Difference

### **Number Needed to Treat**

The number needed to treat (NNT) is the number of individuals who we would need to treat in order to **prevent** one bad outcome.

The NNT is calculated as  $\left|RD\right|^{-1}$ 

- The inverse of the absolute value of the risk difference.
  - For a harmful exposure, we keep the absolute value, but describe the measure as a number needed to harm.

### **NNT**

If the 5-year risk of death among treated is 10%, and among untreated is 20%,

- then how many people do you need to treat to prevent one death over five years?
- $ullet \ |10\% 20\%|^{-1}$  =  $|-10\%|^{-1} = 0.10^{-1} = 10.$ 
  - We must treat 10 people to prevent one death over five years.

Notation: 
$$NNT = 1/|(P(Y|X=1) - P(Y|X=0))|$$

What's the null value?

### **NNT**

Often by convention, round NNT up to nearest integer - Can't treat half a person

- Conservative approach; doesn't always make sense.
- NNT implies causality more strongly than risk difference.
- Risk differences can be viewed as descriptive (although you should be specific and cautious about that)
- Just the difference in observed risks between two groups: not necessarily due to group identification.
- In contrast, NNTs explicitly discuss a treatment having a result: thus are expressing a causal effect.
- Explaining an NNT can become tricky outside of a trial setting because exposure (smoking) isn't always the same as treatment (cognitive behavioral therapy for smoking cessation).

**QUESTIONS?** 

**COMMENTS?** 

**RECOMMENDATIONS?** 

# 5 emergency steps







## **Incidence Odds Ratio**

- ullet For example: Define a population where 10% of people have  $r_{1i}=$  0.60 and
- ullet  $r_{0i}=$  0.20 and 90% of the people have  $r_{1i}=$  0.035 and  $r_{0i}=$  0.006
- Here, the individual ORs = 6.0 for every individual
- the average of the individual ORs = 6.0
- Also, the ratio of the average odds equals 6.0 as well

But, the incidence odds ratio is equal to 3.9

## Incidence Odds Ratio - Calculations

#### **Rudimentary calculations**

```
#10% pop
 0.6/(1-.6) #= 1.5 #r1
 0.2/(1-.2) #= 0.25 #r0
#OR in 10%
1.5/0.25 #= 6
#OR 90%
0.035 / (1-0.035) \# = 0.03626943 \# r1
 0.006/(1-0.006) # = 0.006036217 #r0
#OR in 90%
 0.03626943/0.006036217 #= 6.008636
# OR average
 1.5+0.03626943 #= 1.536269
1.536269/2 #= 0.7681345
 0.25+0.006036217 #= 0.2560362
 0.2560362/2 #= 0.1280181
 0.7681345/0.1280181 #= 6.000202
```

```
## IOR
(0.6*0.1) + (0.035 *0.9) #= 0.0915
(0.2*0.1) + (0.006*0.9) #= 0.0254
#OR1
0.0915/(1- 0.0915) #=0.1007155
#OR0
0.0254/(1-0.0254) #=0.02606197
#IOR
0.1007155/0.02606197 #3.864462
```