

Cross-sectional Studies

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Like cohort studies, cross-sectional studies conceptually begin with a population base. But unlike cohort studies, in cross-sectional studies we do not follow individuals over time. Instead, we only look at the prevalence of disease and/or exposure at one moment in time. These studies take a "snapshot" of the proportion of individuals in the population that are, for example, diseased and non-diseased at one point in time. Other health outcomes besides diseases may also be studied. Cross-sectional studies also differ from cohort studies in the populations that are studied. Cohort studies begin by selecting a population of persons who are at risk of for a specific disease or health outcome; cross-sectional studies begin by selecting a sample population and then obtaining data to classify all individuals in the sample as either having or not having the health outcome.

Ways to use cross-sectional studies

Cross-sectional studies are used both descriptively and analytically.

Descriptive cross-sectional studies simply characterize the prevalence of a health outcome in a specified population. Prevalence can be assessed at either one point in time (*point prevalence*) or over a defined period of time (*period prevalence*). Period prevalence is required when it takes time to accumulate sufficient information on a disease in a population, e.g. what proportion of persons served by a public health clinic over a year have hypertension. These prevalence measures are commonly used in public health; often the point or period aspect is not specified.

In *analytical cross-sectional studies*, data on the prevalence of both exposure and a health outcome are obtained for the purpose of comparing health outcome differences between exposed and unexposed.

Analytical studies attempt to describe the prevalence of, for example, disease or non-disease by first beginning with a population base. These studies differ from solely descriptive cross-sectional studies in that they compare the proportion of



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| | Cohort | Cross-sectional |
|-----------------|--------------------|---------------------------------|
| Study group | Population-at-risk | Entire population (or a sample) |
| Common Measures | Risks and Rates | Prevalence |

exposed persons who are diseased ($a/(a+b)$) with the proportion of non-exposed persons who are diseased ($c/(c+d)$).

Calculating prevalence

The prevalence of a health outcome is simply the proportion of individuals with the health outcome in a population.

$$\text{Prevalence} = \text{cases} / \text{total population}$$

For the following example, two different sub-measures of prevalence can be calculated: the prevalence of coronary heart disease (CHD) among the exposed (people who are not active) and the prevalence of CHD among the unexposed.

Example:

| | Present | Absent CHD | Total |
|------------|---------|------------|-------|
| Not active | 50 a | b 200 | 250 |
| Active | 50 c | d 700 | 750 |
| Total | 100 | 900 | 1000 |

$P_1 = a/a+b = 50/250 = 20.0\%$ prevalence of CHD among people who are not active.

$P_0 = c/c+d = 50/750 = 6.7\%$ prevalence of CHD among people who are active.

The prevalence odds ratio

The prevalence odds ratio (POR) is calculated in the same manner as the odds ratio.

$$\text{POR} = ad / bc$$

The prevalence ratio

The prevalence ratio (PR) is analogous to the risk ratio (RR) of cohort studies. The denominators for both ratios are fixed populations – fixed at the start of the study in the case of a cohort study, and fixed at the point or period of

time for the case-control study. The PR is similar to a RR when the outcome occurs over a short period of time. For example, one would calculate a prevalence ratio for an acute outbreak of tuberculosis in a prison population. This is in contrast to calculating the overall prevalence of positive tuberculin skin tests among the prisoners.

The prevalence ratio can also be calculated from the information on CHD and physical activity:

$$\text{PR} = (a/N_1) / (c/N_0)$$

$$\text{PR} = (50/250) / (50/750) = 3.0$$

In this case, a prevalence ratio of 3.0 can be interpreted to mean that the people who are not physically active are 3 times as likely as those who are physically active to have CHD.

Note that it is preferable to calculate the prevalence odds ratio when the period for being at risk of developing the outcome extends over a considerable time (months to years) as it does in this example. **POR vs. PR**

For chronic disease studies or studies of long-lasting risk factors, POR is the preferred measure of association in cross-sectional studies. For acute disease studies, PR is the preferred measure of association. If the prevalence of disease is low, i.e. 10% or less in exposed and unexposed populations, $\text{POR} = \text{PR}$. Since cross-sectional studies are particularly useful for investigating chronic diseases (e.g. prevalence of AIDS) where the onset of disease is difficult to determine, or for studying long lasting risk factors (such as smoking, hypertension, and high fat diets), the prevalence odds ratio will generally be the preferred measure of association.

Limitations of cross-sectional studies to evaluate risk

Recall that, under steady conditions, the prevalence of disease is influenced both by incidence and duration of disease (or survival with disease).

$$\text{Prevalence} = \text{Rate} \times \text{Average Duration of Disease}$$

Persons who survive longer with a disease will have a higher probability of being counted in the numerator of a prevalence proportion. Short-term survivors will be less likely to be counted as a case. Incidence is influenced only by exposure, whereas prevalence is influenced both by exposure and duration of disease.

If exposure influences survival time, then the POR or PR will not provide a valid estimate of the risk ratio or rate ratio. Thus, the interpretation of the POR or PR is subject to survival bias.

Even if incidence remains constant, either an improvement in disease treatment (that results in higher cure rates) or increased lethality (resulting in a higher case fatality rate) will result in decreased prevalence. The disease itself or the threat of developing the disease may cause outmigration of cases from an environment perceived as causing disease, e.g. workers affected by toxic exposures in a plant may quit, while more resistant workers will stay. This selective migration can bias measures of prevalence.

Other problems with interpretation of cross-sectional studies

Cross-sectional studies as well as case-control studies are affected by the *antecedent-consequent bias*, similar to the chicken and egg question (i.e. "which came first?"). This bias occurs when it cannot be determined that exposure preceded disease, since both are ascertained at the same time (unlike cohort studies or clinical trials). Antecedent-consequent bias does not affect cohort studies because subjects in cohort studies are selected for study because they are disease-free. Exposure is actually observed to precede disease only in a cohort design, including randomized trials.

Uses of cross-sectional studies

Descriptive studies are an important method to evaluate the proportion of a population with disease or with risk factors for disease, such as the prevalence of asthma in children or the prevalence of elevated blood lead in toddlers.

Descriptive cross-sectional studies are widely used to estimate the occurrence of risk factors in segments of the population characterized by age, sex, race or socioeconomic status (SES). National examples of cross-sectional studies of great importance are the *decennial census* and the National Health and Nutrition Surveys (NHANES). Opinion polls and political polls are basically cross-sectional studies. Surveillance of changes in smoking habits or of other behavioral risk factors are sequential cross-sectional studies. The US National Health and Nutrition Examination Survey (NHANES) is one such example. Similarly, surveillance of long lasting diseases such as AIDS is cross-sectional. Descriptive cross-sectional studies are useful for planning or administering preventive or health care services, surveillance programs, and surveys and polls.

Descriptive/analytical cross-sectional studies are useful for establishing preliminary evidence for a causal relationship. These studies are also useful for examining the association between exposure and disease onset for chronic diseases where researchers lack information on time of onset. Examples might include diet and arthritis, smoking and chronic bronchitis, and asthma and exposure to air pollution. Interpretation requires caution regarding potential association of duration of disease with exposure status (*survival bias*).

Survival bias may be minimized if information can be obtained on exposures that clearly preceded the first symptoms of a chronic disease such as arthritis, diabetes, or chronic bronchitis. This depends on access to medical records before the onset of a chronic disease. In addition, it may be necessary to have historical records on an individual's exposure status prior to these first medical visits, e.g. where the person lived or where the person was employed.

Terminology

Antecedent-consequent bias: occurs in cross-sectional studies when it cannot be determined if exposure preceded disease.

Prevalence: the proportion of diseased individuals in a population.

Survival bias: occurs in cross-sectional studies when the exposure influences survival time, and the distribution of that exposure will be distorted among a sample of survivors. (a.k.a. Neyman bias, incidence-prevalence bias, or selective survival bias)

Practice Questions

Answers are located at the end of this notebook

1) Which of the following health outcomes could be studied using a cross-sectional study design? Choose all that apply.

- a) The prevalence of diabetes among adults in the United States in 2014
- b) The prevalence of diabetes among all patients seen at a particular health clinic on one day in 2014
- c) The number of new cases of diabetes diagnosed among at risk adults in the United States in 2014
- d) The number of people in a population with diabetes who are obese and the number of people in a population with diabetes who are not obese, in the United States in 2014

2) Researchers are studying HIV prevalence using a cross-sectional study design. Which of the following factors may affect the researchers' assessment of HIV prevalence in their study population? Choose all that apply.

- a) Changes in HIV treatment
- b) Changes in HIV virulence (HIV virulence refers to the ability of the virus to cause disease)
- c) Population changes (e.g. migration) due to HIV infection, for example people who leave their community to live closer to physicians specialized in treating HIV
- d) Other factors which may affect survival of HIV-infected persons (e.g. changes in healthcare practices)
- e) Changes in HIV diagnostics
- f) Changes in HIV awareness and education

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Answers to Practice Questions

1. Answer choices a, b and d are correct. A cross-sectional study design could be used to assess the prevalence of diabetes among adults in the United States in 2014. A cross-sectional study design could also be used to assess the prevalence of diabetes among all patients seen at a particular health clinic on one day in 2014 by simply recording which patients are known to have been diagnosed with diabetes divided by the number of total patients seen that day. However, the number of new cases of diabetes diagnosed among at risk adults in the United

States in 2014 could not be assessed using a cross-sectional study. The incidence of new cases cannot be measured using a cross-sectional “snapshot” in time design. Instead, a cohort study could be used to follow at risk participants over time to see which people were newly diagnosed with diabetes in 2014. The number of people in a population with diabetes who are obese and the number of people in a population with diabetes who are not obese could be assessed using a cross-sectional design, this would be an example of an analytical cross-sectional study.

2. Answer choices b, c, d, e and f are correct. Changes in HIV virulence could affect HIV-related mortality (survival time) and, thus, would affect prevalence. Any other factors which affect survival of HIV-infected persons would also affect prevalence. Migration could also affect assessment of HIV prevalence if people migrate in or out of the study area based on their HIV status. Changes in HIV diagnostics could affect the timing of HIV diagnosis and/or how likely people are to get tested and, thus, could affect prevalence. Answer choice a is not correct because HIV is not a curable disease. HIV treatment may help an HIV-infected person live longer but will not affect whether or not they have HIV. Therefore, HIV prevalence is not affected by HIV treatment. Note that if a disease was curable with treatment then that disease’s prevalence could be affected by treatment.