

PRINCIPLES OF INFECTIOUS DISEASE EPIDEMIOLOGY

MODULE I – INTRODUCTION TO EPIDEMIOLOGY

This outline is provided as an aid to the student. It contains only the basic content of the module. To view the supporting material such as graphics, examples, etc. please see the module itself.

I. INTRODUCTION

A. What is the Purpose of Epidemiology:

- to better understand the burden and causes of health problems in human populations, and
- to make changes that decrease risk and improve health.

B. Epidemiology is comprised of a set of tools, including:

- Scientific methods for study/research
- Techniques for collecting and organizing information
- Information about the biological basis of health and illness
- Information about human behavior that affects health
- “People skills” needed to gain cooperation and gather solid information

C. Epidemiology can be applied to any aspect of human health, including:

- all types of diseases
- impairments
- disabilities
- injuries
- the utilization and outcomes of health services, and even
- good health.

D. Epidemiology has been used to study these conditions and many more:

- West Nile Virus infections
- Sexually transmitted diseases and HIV
- Surgical wound infections
- Hip fractures
- Childhood malnutrition
- Pertussis (whooping cough)
- Schizophrenia

E. Epidemiology can help us identify and understand the factors that influence the emergence, severity, and consequences of health problems. Examples include:

- tobacco use and lung cancer
- physical activity and heart disease
- diet and longevity
- lead exposure and developmental disabilities

- seat belt use and motor vehicle injuries/deaths
- childhood vaccine initiatives and child mortality
- statin drugs and outcomes of atherosclerotic heart disease

This course will concentrate on the use of epidemiology to understand, prevent and control infectious diseases.

F. Infectious diseases:

- Are caused by micro-organisms
- Are transmitted to humans from other humans, animals or the environment
- Usually follow recognizable patterns of symptoms, timing, etc.
- Evolve over time as new organisms emerge and human behavior and environments change

G. This module is designed to prepare public health workers to meet the following objectives:

1. Define key epidemiologic terms
2. Describe at least five factors that significantly influence the incidence and distribution of a disease in a human population
3. Describe the modern approach to epidemiology

II. KEY EPIDEMIOLOGICAL TERMS

A. EPIDEMIOLOGY is derived from three Greek root words:

- epi – on, upon
- demos – people
- logy – study

Epidemiology is, thus, the study of what is upon the people. **In modern terms, it is the science of the distribution of disease and its determinants (causes).**

Epidemiology is also a **process** that uses the facts at hand as clues to point to new knowledge and solutions. Epidemiologists have been called “disease detectives” and “medical sleuths” for this reason.

B. Three key terms are used to describe basic patterns of infectious disease occurrence. The terms are defined by the American Public Health Association (APHA) as:

Endemic: The habitual presence of a disease within a given geographic area; may also refer to the usual prevalence of a given disease within such an area.

Epidemic: The occurrence in a community or region of a group of illnesses of similar nature, clearly in excess of normal expectancy, and derived from a common or from a propagated source.

Pandemic: A worldwide epidemic.

Endemic disease levels are measured by ongoing surveillance systems. Some fluctuations usually occur.

The transition from endemic to epidemic can happen in days, weeks, months, or even years, depending on the disease. It may be hard to tell when the shift begins.

There are parallel terms referring to disease patterns in animals that may affect humans:

Zoonosis: An infection or an infectious disease transmissible under natural conditions between vertebrate animals and man (APHA).

Enzootic: “Endemic” among animal populations

Epizootic: “Epidemic” among animal populations

C. The term epidemiology is used to refer both to:

A method of study of diseases, and

A body of knowledge about the natural history of a disease

The natural history of a disease is a description of how that disease “behaves” and what factors affect its incidence and distribution. Here is a partial list of such factors:

1. Biological
2. Geographic
3. Nutritional
4. Meteorological
5. Social
6. Cultural
7. Religious
8. Behavioral
9. Economic
10. Historical
11. Political
12. Technological

Example:

“The Epidemiology of Measles” includes:

Factor	Example
Causative organism	Measles virus
Host	Humans
Mode of transmission	Airborne droplets and direct contact with secretions
Incubation period	About 10 days, range 7-18 days from exposure
Period of communicability	From a few days before onset of illness to 4 days after the appearance of the rash
Usual symptoms	Fever, conjunctivitis, coryza, cough and blotchy red rash
Complications	Otitis media, pneumonia, croup, diarrhea and encephalitis
Mortality rate	2-3 per 1,000 cases in USA; 3-5% in developing countries

Epidemiologic methods were used to compile most of this information.

III. EVOLUTION OF EPIDEMIOLOGY

Human beings have always sought to understand and explain the occurrence of disease and death.

A. Supernatural Causation

- Throughout most of human history, people believed in the supernatural as the cause of illness.
- Individuals and communities who became ill were thought to have angered the gods or spirits, or to be victims of their enemies' magic.
- Such beliefs are still held today, even in otherwise modern societies.

B. Environmental Explanations

- Hippocrates was the first to suggest that disease is caused by environmental elements, around 400 B.C.E.
- From 400 B.C.E. until the mid-19th century, many theories were developed to explain infection.
- Most of these theories were based on the concept of miasmas. Contagious matter was thought to create a gaseous form, a miasma, which spread infection through the atmosphere.

C. Host Factors

- Edward Jenner, in 1796, introduced the idea of host resistance to explain why some people were immune to smallpox.
- Peter Panum, in 1846, took this idea further when he studied measles in the Faroe Islands. The islanders had not been exposed to measles in 70 years, and many otherwise healthy adults died.
- By contrast, in mainland Denmark all were exposed to measles in childhood, and adult deaths from this disease were rare.

D. Toward a Modern Approach

I. John Snow, a British scientist, made the greatest strides toward modern epidemiology in the 19th Century.

- At that time, infectious diseases were the leading causes of death.
- Up to half of all children died before the age of five, mostly due to infectious diseases.
- a. Snow's first study was conducted in 1848 when an epidemic of cholera hit the Golden Square area of London.
 - Even before bacteria were discovered as a cause of disease, Snow was able to stop devastating outbreaks by observing and recording information about the distribution of the disease.
 - Snow was working from a theory that water had something to do with the spread of cholera.
 - In the Golden Square area, most people got their water from public pumps.

- Snow's first step was to gather information about the cholera patients in the area, and record their residence or place of work.
 - He developed a spot map that showed the distribution of cases in relation to the water pumps.
 - Snow investigated all the pumps in the area.
 - One pump was so grossly contaminated that people had avoided its use.
 - One pump was in an out-of-the way location, and therefore not used much.
 - Many of the cholera cases were clustered around the Broad Street Pump.
 - When Snow checked with the families of the cholera victims, he confirmed that they all used the Broad Street Pump.
 - Snow showed that by studying the distribution of cases, a source of infection could be found that explained the pattern. **He then took action to stop the epidemic, by having the pump handle removed.**
- b. Snow did more pioneering work during another cholera outbreak in 1854.
- This outbreak affected several areas of London that depended on water hauled in by wagon.
 - Two companies, the Southwark and Vauxhall (S&V) Company and the Lambeth Company, did the hauling.
 - Some districts were served mostly by S&V, other districts mostly by Lambeth, and some districts were served by both.
 - Snow compared the number of cholera deaths in the various districts.
 - Snow realized that the raw numbers of deaths could not tell him much.
 - If some districts had many more people than others, they couldn't really be compared.
 - So he collected population figures from the most recent census and calculated cholera death rates per 1,000 people.
 - The rates paint a very clear picture.
 - The death rate was 22 times higher in the districts served only by S&V, compared with those served only by Lambeth.
 - Further investigation showed that both companies were drawing their water from the Thames River.
 - However, S&V pulled water from a point just downstream from a major sewer outlet, while Lambeth drew theirs upstream from the outlet.
 - Snow had shown conclusively that water could serve as a vehicle for transmitting infection. He also showed that humans could intervene, in this case by changing the location of the S&V water intake.

2. William Farr, the father of modern vital records, was another important figure in the development of epidemiology. His contributions include:

- establishing the first registry of births and deaths in the 1830s.
- advancing population-based surveillance.
- distributing reports that led to public health interventions.

Example: Mortality in Liverpool, 1843. The median age of survival in Liverpool in 1843 was 6 years, compared with an average of 45 in the rest of England. This revelation led to policy and law changes to improve sanitation.

3. Establishment of the Germ Theory (1860 to 1890)

- Studies by Louis Pasteur and Robert Koch firmly established the germ theory.
- Attention and research efforts shifted
 - to biological agents, and
 - away from the environment or host resistance.
- This shift happened in spite of Pasteur's warning that the role of the environment was more important than that of the agent.
- Snow's work was not appreciated or expanded upon until the years after World War II.

4. Modern Epidemiology

a. The **modern approach** in describing an infectious disease focuses on:

- the interaction of the disease agent,
- the host, and
- the environment.

b. The disease agent, the host, and the environment interact dynamically to produce disease. All three of these are constantly changing:

- **Disease agent:** Microorganisms adapt to changing conditions, including human control efforts such as antibiotics.
- **Host:** Human populations are constantly growing and moving as people age, travel, and migrate into new environments.
- **Environment:** Changes occur locally and globally, both naturally and through human intervention.

(i) Disease Agents: There are many "agents" of disease and disabilities, including:

- nutritional components such as vitamin deficiency diseases and obesity.
- physical forces such as fire, radiation, and chemicals.
- biological agents such as bacteria and viruses.

In this course, however, we will be focusing on biological agents that will be covered more fully in other segments of the course.

(ii) Host Factors: Some host factors that influence susceptibility to disease are:

Sex
Race
Age
Occupation

Nutrition
Heredity
Marital status
Socioeconomic status
Religious and social customs
Immunization history
Previous history of disease

(iii) Environmental Factors: Many aspects of the environment influence both an agent's survival and growth, and a host's contact with a disease agent. Some of the environmental factors are shown below.

Temperature
Humidity
Altitude
Water
Radiation
Pollution
Housing conditions
Food/milk

Each disease is influenced by a particular set of factors.

Examples:

- 1) Some foodborne diseases, such as salmonella, are highly dependent on environmental factors such as cross-contamination and cooking/holding temperatures.
- 2) The spread of measles is influenced by immunization status, but also by housing conditions and nutritional status.
- 3) Disease history is a big factor with some diseases, such as hepatitis A, that confer lifetime immunity after infection.
- 4) Many other diseases can cause repeated infections in the same individual, for example gonorrhoea, shigellosis and malaria.
- 5) Some diseases are transmitted to humans only through arthropod vectors such as mosquitoes, ticks or lice.
 - These disease organisms may have complex lifecycles that pass through several different hosts.
 - For example, the spirochete that causes Lyme Disease is transmitted to humans from certain ticks, but its lifecycle includes rodents and large mammals such as deer.
 - Vectors are subject to agent, host and environmental factors too.

Epidemiology gives us tools to learn about how these factors interact to produce a particular disease in a particular population. Good epidemiology is most critical when investigating an outbreak or the emergence of a new disease - some of the most important tasks of the field epidemiologist.

A team approach is almost always used in epidemiologic investigations. The range of experts needed depends on the disease and the setting, but usually includes:

- Epidemiologists
- Medical professionals (physicians, nurses)
- Laboratory scientists
- Statisticians
- Environmental specialists

Summary

Epidemiology is a set of tools for understanding the burden and causes of health problems in human populations, so that we can make changes that decrease risk and improve health.

Epidemiology can help us identify and understand the factors that influence the emergence, severity, and consequences of health problems.

Epidemiology is defined as the science of the distribution of disease and its determinants (causes).

Human beings have always sought to understand and explain the occurrence of disease and death. The modern approach to epidemiology has developed within the past 150 years.

The modern epidemiological approach to infectious diseases focuses on the interaction of the disease agent, the host and the environment.

PRINCIPLES OF INFECTIOUS DISEASE EPIDEMIOLOGY

MODULE II – THE INFECTIOUS DISEASE PROCESS

This outline is provided as an aid to the student. It contains only the basic content of the module. To view the supporting material such as graphics, examples, etc. please see the module itself.

I. INTRODUCTION

Module II is designed to prepare public health workers to meet the following objectives:

1. Describe the six major components of the infectious disease process
2. Demonstrate understanding of the concepts of the infectious disease spectrum

II. THE CHAIN OF INFECTION

In order for infection and disease to occur in an individual, a process involving six related components must occur. This process has been referred to as the “Chain of Infection.” The six steps or “links” in the chain are:

- Etiologic agent
- Reservoir
- Portal of Exit
- Mode of Transmission
- Portal of Entry
- Susceptible Host

In this module, we will examine each of these links and some other important concepts that help us understand infectious disease transmission. To stop the spread of disease, one or more of these links must be broken.

A. Etiologic Agents

There are seven categories of biological agents that can cause infectious diseases. Each has its own particular characteristics. The types of agents are:

1. Metazoa
2. Protozoa
3. Fungi
4. Bacteria
5. Rickettsia
6. Viruses
7. Prions

1. **Metazoa** are multicellular animals, many of which are parasites. Among the diseases they cause are:

- a. Trichinellosis, also called trichinosis, caused by an intestinal roundworm transmitted through undercooked meat.
 - b. Hookworm, transmitted through feces-contaminated water and soil. Infestation can cause chronic anemia that often results in retarded mental and physical development of children.
 - c. Schistosomiasis, caused by a blood fluke and transmitted through contaminated water. Symptoms are related to the number and location of eggs in the human body, and may involve the liver, intestines, spleen, urinary tract, and reproductive system.
2. **Protozoa** are single-cell organisms with a well-defined nucleus. Some of these are human parasites. Examples of diseases cause by protozoa include:
 - a. Malaria, a mosquito-borne disease that is one of the top three infectious diseases in the world (along with tuberculosis and HIV).
 - b. Giardiasis, an infection of the upper small intestine that causes a diarrheal illness. Outbreaks can be difficult to control, especially in child care settings
 - c. Toxoplasmosis, transmitted to humans from cats and undercooked meat. When this systemic disease infects a pregnant woman, it can cause the death of the fetus.
 - d. *Pneumocystis carinii* pneumonia or PCP, which is often fatal, especially in people with compromised immune systems such as those infected with HIV.
3. **Fungi** are nonmotile, filamentous organisms that cause diseases that can be very difficult to treat. Some examples important to public health are:
 - a. Histoplasmosis, transmitted by inhaling dust from soil that contains bird droppings. The severity varies widely, with the lungs the most common site of infection.
 - b. Candidiasis, transmitted by contact with human patients and carriers. This fungus causes lesions on the skin or mucous membranes, including “thrush” and vulvovaginitis. Symptoms can be severe in immunocompromised people.
4. **Bacteria** are single-celled organisms that lack a nucleus. They are responsible for a wide range of human diseases, including:
 - a. Tuberculosis, a chronic lung disease that is a major cause of disability and death in many parts of the world.
 - b. Staphylococcal disease, which can affect almost every organ system. Severity ranges from a single pustule of impetigo, through pneumonia, arthritis, endocarditis, etc., to sepsis and death.
 - c. Chlamydia and gonorrhea, the most widespread sexually transmitted diseases.
 - d. Tetanus and diphtheria, two diseases that were once major public health problems but are now well controlled through immunization.
 - d. Other vaccine-preventable diseases caused by bacteria are:
 - Pertussis
 - Haemophilus influenzae type b (Hib)
 - Pneumococcal disease.

5. **Rickettsia** are a genus of bacteria usually found in the cells of lice, ticks, fleas and mites. They are smaller than most bacteria and share some characteristics of viruses. Diseases cause by rickettsia include:

- a. Rocky Mountain Spotted Fever, a tick-borne systemic disease that can be hard to diagnose and that leads to death in 3-5% of US cases.
- b. Typhus, a louse-borne rash illness with a high case-fatality rate that has occurred historically in poor living conditions brought on by war and famine.

6. **Viruses** are very small, consisting of an RNA or DNA core and an outer coat of protein. They can reproduce and grow only inside of living cells. Many viral illnesses are significant to public health, including:

- a. Influenza, a respiratory illness that contributes to development of pneumonia and occurs in annual epidemics during the winter months
- b. HIV (human immunodeficiency virus), that causes Acquired Immunodeficiency Syndrome (AIDS). This severe, life-threatening pandemic disease has spread worldwide within the past 20-30 years.
- c. Rabies, that is spread to humans from animal bites or scratches. Rabies is almost always fatal in humans but is preventable by a vaccine.
- d. Measles, mumps, rubella, and poliomyelitis are all well controlled in the US through immunization.

7. **Prions** are infectious agents that do not have any genes. They seem to consist of a protein with an aberrant structure, which somehow replicates in animal or human tissue. Prions cause severe damage to the brain. Diseases associated with prions include:

- a. CWD, chronic wasting disease of mule, deer and elk;
- b. BSE, bovine spongiform encephalopathy in cows; and
- c. CJD, Creutzfeld-Jacob disease in humans.

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B. Reservoirs

The next essential link in the chain of infection is the reservoir, the usual habitat in which the agent lives and multiplies. Depending upon the agent, the reservoir may be:

- humans,
- animals, and/or
- environment

When working with any disease agent, it is important to learn about its usual reservoir(s).

1. Human Reservoirs

There are two types of human reservoirs, acute clinical cases and carriers.

- a. Acute clinical cases are people who are infected with the disease agent and become ill.
 - Because they are ill, their contacts and activities may be limited.

- They are also more likely to be diagnosed and treated than carriers are.
- b. Carriers, on the other hand, are people who harbor infectious agents but are not ill.
- Carriers may present more risk for disease transmission than acute clinical cases, because their contacts are unaware of their infection, and their activities are not restricted by illness.
 - Depending on the disease, any of the following types of carriers may be important:
 - Incubatory carriers
 - Inapparent infections (also called subclinical cases)
 - Convalescent carriers
 - Chronic carriers

Incubatory carriers are people who are going to become ill, but begin transmitting their infection before their symptoms start. Examples: measles: a person infected with measles begins to shed the virus in nasal and throat secretions a day or two before any cold symptoms or rash are noticeable. Many other diseases also have an incubatory carrier phase. Most notably, HIV infection may be present for years before the person develops any symptoms.

Inapparent infections: People with inapparent infections never develop an illness, but are able to transmit their infection to others. With some diseases, inapparent infections are more common than acute clinical cases. Example: Of every 100 individuals infected with the poliomyelitis virus, only one becomes paralyzed. Four others will have a mild illness with fever, malaise, headache, nausea and vomiting. But 95 out of the 100 will have no symptoms at all, although they pass the virus in their feces.

Sometimes the likelihood of an inapparent infection depends on another epidemiologic factor, such as age. Hepatitis A is a good example of this. Over 50% of adults infected with this virus develop symptoms. However, among children under 5, there may be 10 inapparent infections for every child who develops jaundice. So children are very effective spreaders of the hepatitis A virus, which is passed in the feces regardless of the presence of symptoms.

Subclinical infections: With some diseases, such as meningococcal meningitis, the number of subclinical cases may be quite high before a single clinical case appears. On some military bases where outbreaks have occurred, the carrier rate has been documented at 50% or more.

Convalescent carriers are people who continue to be infectious during and even after their recovery from illness. This happens with many diseases. Example: Salmonella patients may excrete the bacteria in feces for several weeks, and rarely even for a year or more. This is most common in infants and young children. Treatment with inappropriate antibiotics may prolong the convalescent carrier phase.

Chronic carriers are people who continue to harbor infections for a year or longer after their recovery. Example: the chronic carrier state is not uncommon following hepatitis B infection, whether or not the person became ill, and may be lifelong. The risk of developing chronic hepatitis B depends on the person's age at infection. About 90% of infants infected at birth become chronic carriers of the disease, compared with only 1-10% infected after age 5. That is why it is so important to give hepatitis B vaccine to newborns.

2. Animal Reservoirs

Animal reservoirs of infectious agents can be described in the same way as human reservoirs. They may be

- acute clinical cases, or
- carriers.

Depending upon the disease, different carrier phases may be important in transmission.

3. Environmental Reservoirs

Plants, soil and water may serve as the reservoir of infection for a variety of diseases.

- Most fungal agents (mycoses) live and multiply in the soil.
Examples:
 - The organism that causes histoplasmosis lives in soil with high organic content and undisturbed bird droppings.
 - The agents that cause tetanus, anthrax and botulism are widely distributed in soil.
 - The agent of Legionnaire's Disease lives in water, including hot water heaters.

C. Portal of Exit

The next link in the chain of disease transmission is the portal of exit, **the route by which the disease agent may escape from the human or animal reservoir**. While many disease agents have only one portal of exit, others may leave by various portals.

The portals most commonly associated with human and animal diseases are:

- Respiratory
- Genitourinary
- Alimentary
- Skin
Superficial lesions

Percutaneous

- Transplacental

1. **Respiratory**: This is the route of many disease agents that cause respiratory illnesses such as the common cold, influenza, and tuberculosis. It is also the route used by many childhood vaccine-preventable diseases, including measles, mumps, rubella, pertussis, *Haemophilus influenzae* type b (Hib), and pneumococcal disease. This is the ***most important portal and the most difficult to control***.

2. **Genitourinary**: This portal of exit is the route of sexually transmitted diseases, including syphilis, gonorrhea, chlamydia, and HIV. Schistosomiasis, a parasitic disease, and leptospirosis, a bacterial infection, are both spread through urine released into the environment.

3. **Alimentary**: The alimentary portal of exit may be the mouth, as in rabies and other diseases transmitted by bites. More commonly, disease agents are spread by the other end of the intestinal tract. These are referred to as enteric diseases. In general, enteric diseases may be controlled through good hygiene, proper food preparation and sanitary sewage disposal. Examples include:

- Hepatitis A
- Salmonella, including typhoid
- Shigella
- Cholera
- Giardia
- Campylobacter

4. **Skin**: Skin may serve as a portal of exit through superficial lesions or through percutaneous penetration.

- Superficial skin lesions that produce infectious discharges are found in smallpox, varicella (chickenpox), syphilis, chancroid, and impetigo.
- Percutaneous exit occurs through mosquito bites (malaria, West Nile virus) or through the use of needles (hepatitis B and C, HIV).

5. **Transplacental**: This portal of exit from mother to fetus is important in the transmission of rubella, HIV, syphilis, and cytomegalovirus (the most common infectious cause of developmental disabilities). It is, fortunately, not a factor for most diseases.

D. Mode of Transmission

A mode of transmission is necessary to bridge the gap between the portal of exit from the reservoir and the portal of entry into the host. The two basic modes are **direct** and **indirect**.

1. **Direct transmission** occurs more or less immediately. Many diseases are transmitted by direct contact with the human, animal or environmental reservoir. Prime

examples are sexually transmitted diseases and enteric diseases such as shigella, giardia and campylobacter. Contact with soil may lead to mycotic (fungal) diseases.

Droplet spread is also considered direct transmission. Infectious aerosols produced by coughing or sneezing can transmit infection directly to susceptible people up to three feet away. Many respiratory diseases are spread this way.

2. **Indirect transmission** may occur through animate or inanimate mechanisms.

- *Animate mechanisms* involve vectors. Flies may transmit infectious agents such as shigella in a purely mechanical way, by walking on feces and then on food. Mosquitoes, ticks or fleas may serve as reservoirs for the growth and multiplication of agents, for example in malaria or Lyme disease.
- *Inanimate mechanisms*: When disease agents are spread by environmental vehicles or by air, this is referred to as indirect transmission by *inanimate mechanisms*. Anything may be a vehicle, including objects, food, water, milk, or biological products.
 - Food is a common vehicle for salmonella infections
 - Water is the usual vehicle in cholera outbreaks
 - Surgical instruments and implanted medical devices may be the vehicles of staphylococcal infections

Indirect, airborne transmission is important in some respiratory diseases. This occurs when very tiny particles of respiratory material become suspended in the air (called aerosols). Such particles may remain suspended and stay infectious for varying periods of time. They are particularly dangerous because their size (1 to 5 microns) allows them to be drawn deep into the lungs and retained. Tuberculosis is spread this way, as is measles in certain settings such as doctors' offices. Air may also spread particles of various sizes from contaminated soil, or from objects such as clothing and floors.

E. Portals of Entry

The portal of entry into the host is usually the same as the portal of exit from the reservoir.

In some diseases, however, the exit and entry portals may differ. Example: staphylococcal bacteria may escape from one person's respiratory tract to infect another person's skin lesion. If that person is a foodhandler, the staphylococcal bacteria may escape from the infected skin lesion, contaminate food where it can incubate, and cause "food poisoning" in people eating the food.

F. Susceptible Host

The last essential component in the chain of infection is the susceptible host.

Susceptibility is affected by:

- Genetic factors
- General resistance factors
- Specific acquired immunity

1. **Genetic factors** The role of *genetic factors* in susceptibility to infectious diseases is not yet well understood. Genes do seem to play a role in the progression of HIV disease, and perhaps in individuals' susceptibility to meningococcal meningitis.

2. **General resistance factors** include many body functions that we take for granted. Intact skin and mucous membranes help us resist disease. So do the gastric acid in our stomachs, the cilia in our respiratory tracts, and the cough reflex.

3. **Specific acquired immunity** is the greatest influence on host susceptibility. This immunity is specific to a particular disease agent, and it may be acquired naturally or artificially.

- Natural immunity may be acquired by experiencing an infection, which is called "active natural immunity." Many diseases confer immunity after a single infection, but many others do not. A single bout of measles or chickenpox, for example, confers lifelong immunity to that disease. Influenza and salmonella are examples of infections that do not confer immunity and therefore may recur.

Another mechanism of natural immunity is the transfer of antibodies from the mother to the newborn child, via the placenta and/or breast milk. This is called "passive natural immunity," and it diminishes after varying lengths of time. It is very important in giving infants a good head start in life.

- Artificial immunity may be acquired through the use of vaccines, toxoids and immune globulins.
 - Active immunity: Receiving a vaccine or toxoid stimulates "active" immunity, since the recipient responds by producing his/her own antibodies.
 - Passive immunity: Receiving an antitoxin or immune globulin confers "passive" immunity, essentially by borrowing the antibodies of other people. Passive immunity lasts for only a short time, while active immunity usually lasts much longer, even for a lifetime.

III. The Infectious Disease Spectrum

By now, you probably appreciate the complexity of the factors that work together to cause the transmission of infectious agents. The *impact* of disease agents on human host populations is also a bit complex.

If a large number of individuals are equally exposed to an infectious agent, they do not all respond in the same manner. In fact, there may be a broad range of responses:

- Some do not become infected at all
- Some become infected but develop no symptoms
- Some become infected and develop mild or moderate symptoms
- Some become infected and develop severe symptoms
- Some die as a result of their infection

Part of this variation is due to the capacity of the agent to produce disease. Infection of a healthy adult population with salmonella is likely to result in mostly inapparent or mild cases, with only a few people with more severe symptoms and very few deaths. On the other end of the spectrum, infections with rabies almost always result in severe illness and death.

Part of the variation is due to differing levels of resistance of the hosts. If measles is introduced into a highly immunized population, then most individuals do not become infected. If measles is introduced into an unimmunized, nutritionally deprived population, the spectrum shifts toward severe symptoms and a high death rate.

The existence of the infectious disease spectrum can make it challenging to find out the extent of transmission in a particular population. Most cases with inapparent or mild symptoms will never be discovered or reported, since these people will not seek health care. So when moderate or severe cases are reported, they may represent the “tip of the iceberg.”

Another challenge is posed by the fact that many diseases look alike. A variety of agents may produce essentially similar clinical syndromes. For example, the signs and symptoms of tuberculosis, other mycobacteria, and histoplasmosis may be the same. However, effective treatment and control measures are very different for these three diseases. This is why laboratory identification of the specific disease agent is so important in any epidemiological investigation.

Summary

In order for infection and disease to occur in an individual, a process involving six related components must occur. This process has been referred to as the “Chain of Infection.”

To stop the spread of disease, one or more of these links must be broken.

The impact of disease agents on human host populations is complex. If a large number of individuals are equally exposed to an infectious agent, there may be a broad range of responses, from no infection at all to death.

Because of the infectious disease spectrum, it can be challenging to identify the extent of transmission in a particular population.

Many diseases share the same signs and symptoms, so laboratory studies are important to identify the specific disease agent.

PRINCIPLES OF INFECTIOUS DISEASE EPIDEMIOLOGY

MODULE III – Public Health Surveillance

This outline is provided as an aid to the student. It contains only the basic content of the module. To view the supporting material such as graphics, examples, etc. please see the module itself.

I. INTRODUCTION

Module III is designed to prepare public health workers to meet the following objectives:

- Define public health surveillance and describe its components
- Demonstrate understanding of the purposes of surveillance and how it relates to public health action
- Describe the two basic forms of surveillance and the most common sources of surveillance information
- Demonstrate knowledge of how to evaluate and improve surveillance systems

II. WHAT IS PUBLIC HEALTH SURVEILLANCE?

Surveillance is the continuous monitoring of the occurrence of a disease (or other important health event) in a population. It consists of:

- Ongoing, systematic collection of health data
- Data analysis
- Interpretation of data,
- Dissemination of the information, AND
- Linking the health data to public health practice

Public Health Surveillance

- is an important part of assessment, one of the three core functions of public health.
- takes many forms, depending on the disease(s) involved and the conditions under which it is being done.
- may focus on the entire population, or on subpopulations within it.
- may be longstanding or temporary, or even intermittent.
- may be “high-tech” or very rudimentary.

Whatever form it takes, surveillance is essential to the practice of public health, especially for infectious disease prevention and control.

The concept of public health surveillance grew out of the earlier practice of **medical surveillance**, which is still done in some situations.

Medical Surveillance	Public Health Surveillance
Close observation of people exposed to a communicable disease to detect symptoms early and provide treatment or require isolation	Always looks at populations , rather than individuals

III. WHO IS RESPONSIBLE FOR PUBLIC HEALTH SURVEILLANCE?

- To be effective, surveillance systems must involve
 - public health agencies,
 - healthcare providers, and
 - the public
- Active cooperation is essential so that accurate, timely information can be collected and acted upon.
- Communication and data must flow among healthcare providers and the public, but only public health agencies can pull information together from a variety of sources, analyze it and feed it back in a form that can guide action.

IV. WHY DO WE DO PUBLIC HEALTH SURVEILLANCE?

A well-designed surveillance system can provide critical information for public health action. Agencies use various kinds of surveillance data to:

- 1. Guide immediate action for cases of public health importance**, such as a rare disease or suspected bioterrorist agent
- 2. Describe and monitor health events and trends through surveillance systems** in their jurisdiction
 - Who is at risk?
 - When and where is disease occurring?
 - What exposures or risk factors may be related to the disease?
 - Is there an urgent problem developing, such as an outbreak?

- Is a new disease or health concern emerging?
- 3. Set priorities for the use of resources** (time, expertise, technology and money)
 - To what activities should staff time and expertise be directed?
 - How much money should be spent on a particular problem?
 - Do we need to respond immediately to investigate and control a problem in our community?
- 4. Assist in planning, implementing and evaluating public health interventions and programs**
 - To which populations should we target interventions or control measures?
 - Are new prevention programs needed?
 - How effective are our programs and control measures in reducing disease occurrence?
- 5. Evaluate public policy**
 - Are current rules or laws protecting the public?
 - Are new rules, laws, ordinances or other policies needed?
- 6. Generate questions and hypotheses that provide direction for further research**
 - Why is this disease occurring in this specific population?
 - Is the disease the result of an environmental exposure?
 - Is this group of people especially susceptible to the disease?
 - Has the disease agent changed somehow?

Let's look at each of these uses of surveillance data in more detail.

- 1. Guiding immediate action for cases of public health importance.**
 - A good surveillance system serves as an “early warning” system for the community's health.
 - When an unusual disease case occurs, the public health agency can investigate immediately and implement control measures if needed to prevent spread.
 - For example, even a single case of anthrax, measles or diphtheria should trigger a full-scale public health investigation.
- 2. Describing and monitoring health events and trends through surveillance systems** can allow us to:
 - Detect sudden changes in disease occurrence and distribution.
 - We may notice a sudden increase that could be an outbreak.

- Or a disease that previously occurred mainly in one group of people may move to another
- Follow long-term trends and patterns of disease.
 - Many diseases ebb and flow in cycles that vary by season or over several years.
 - Knowing this may help us interpret the current situation.
- Identify changes in agents and host factors.
 - Many infectious agents change over time. (E.g., shifts in predominant salmonella serotype by year, antibiotic resistant gonorrhoea).
 - Data from laboratory scientists who monitor these characteristics can be used to direct vaccine production, treatment, and other prevention and control measures.

3. Set priorities for the use of resources.

- Surveillance data are very important in determining the best use of a public health agency's time, staff and budget.
 - Which diseases or conditions have the greatest impact on the population?
- Infectious disease trends may change quickly, sometimes even overnight, and health agency resources must be flexible to respond effectively.

Example: a suspected bioterrorism incident or an outbreak of foodborne illness may supersede other routine activities. In such a situation, staff must be reassigned to:

 - intensify surveillance to identify other susceptible and potentially exposed people
 - search for the source of the problem
 - design and implement control measures (environmental controls, screening, immunization and/or treatment, restriction of activities)
- If the outbreak is large, or the disease very serious, the agency's emergency response plan may need to be activated. In that case many resources outside the agency can be brought into play.
- Surveillance data can also be used to identify when a public health problem has been solved, so that resources can be redirected to other, more pressing needs. However, if this is done too soon, the consequences can be costly. **Example:** it may be tempting to cut resources for tuberculosis

control when the incidence goes down. This was done in the 1980s, and the result was an upsurge in tuberculosis cases.

4. Assist in planning, implementing and evaluating public health interventions and programs.

Information from ongoing surveillance systems helps us identify public health problems and design programs to address them.

- In the planning stage, surveillance can pinpoint where a problem is occurring and who is affected by it so that efforts can be directed to the right area and the right people.
- In the implementation stage, good data can help persuade policymakers and community members that a program or policy change is needed and worth supporting.
- In the evaluation stage, surveillance data can help us determine the effect of our efforts. If the program or intervention is working, then disease incidence should go down.

EXAMPLES

a. When surveillance data showed that a new arbovirus, the West Nile Virus, was spreading across the country, public health agencies developed plans to prevent and control it. They tried several new approaches including:

- public education campaigns (billboards, radio spots) to promote the use of mosquito protections, including DEET insect repellent
- public education and code enforcement activities to reduce mosquito breeding sites by emptying/draining standing water
- community cleanup activities to reduce mosquito habitat

Agencies then evaluated the results of these efforts through surveys as well as by continued, intensive surveillance activities.

b. Detailed surveillance data regarding HIV infection have guided community planning for prevention programs.

- In the 1980s when AIDS first emerged, health education messages were targeted toward the groups most heavily impacted by the epidemic, men who have sex with men and injecting drug users.
- In the late 1990s, the data showed that HIV was beginning to infect more women, especially minority women.
- Outreach and education programs were designed to reach out to women, for example, by distributing gender and culture-appropriate materials in places where women go, such as beauty shops.

5. Evaluate Public Policy

Public policies, such as laws, ordinances, and institutional policies, are important public health interventions. Surveillance information can help identify needed policy changes.

EXAMPLES

- a. Immunization requirements have evolved as disease transmission patterns change.
 - School immunization rules were implemented and enforced to reduce outbreaks of measles and rubella in schools.
 - As more children entered childcare centers and homes, the transmission of vaccine-preventable diseases among preschool children increased. In the 1980s, immunization requirements for childcare attendance were developed and incorporated into licensure requirements. Outbreaks in licensed childcare settings have since become rare.
- b. Standards for food protection are codified as state rules and local ordinances. Inadequacies in these food codes were identified in the 1990s through surveillance data, and the rules were strengthened.
 - Required cooking temperatures for beef were changed because of *E. coli* O157:H7 outbreaks.
 - The rules on the transport and handling of fresh eggs were changed because of outbreaks of *Salmonella enteritidis*.

6. Generate questions and hypotheses that provide direction for further research.

- Surveillance data by themselves cannot answer questions about the causes of illness.
- However, observation of the patterns of disease can help generate ideas about causes and potential control measures.
- These ideas can be developed into research hypotheses that lead to new knowledge in disease prevention and control.

EXAMPLES

- a. Toxic shock syndrome (TSS)
 - is a serious illness caused by certain toxin-producing strains of the *Staphylococcus aureus* bacterium.
 - It usually has a sudden onset of fever, chills, vomiting, diarrhea, muscle aches and rash, and can rapidly progress to multisystem dysfunction. About 5% of cases are fatal.

- An epidemic of TSS in women of childbearing age was detected through public health surveillance in the late 1970s.
 - Menstruating women and women using barrier contraceptives such as intra-uterine devices (IUDs) were at highest risk.
 - This observation led to further research on tampons, which found that the fibers used in certain highly absorbent brands encouraged bacterial growth.
 - These brands were removed from the market, the epidemic abated, and TSS incidence has declined ever since.
- b. Vaccine policies may be adjusted as a result of research prompted by surveillance findings.
 - A puzzling upsurge in measles cases occurred in the late 1980s, in Missouri and throughout the US, mainly in school-age children.
 - This upsurge prompted CDC to study the children's antibody levels, and they found that immunity waned among those who had only had one measles shot.
 - Two doses of measles vaccine have been required for school attendance since that time.

V. CONDUCTING PUBLIC HEALTH SURVEILLANCE: THE FIVE COMPONENTS

Our definition of surveillance consists of five components:

1. Ongoing, systematic collection of health data
2. Data analysis
3. Interpretation of data,
4. Dissemination of the information, AND
5. Linking the data to public health practice

Let's look at each of these in more detail.

1. ONGOING, SYSTEMATIC COLLECTION OF HEALTH DATA→How do we get the information?

A. Legal Requirements

- Most surveillance involves the collection of personal information from medical records.

- Healthcare providers can only share such information if the proper legal framework is in place to authorize it.
- Under the federal Health Insurance Portability and Accountability Act (known as HIPAA), public health agencies have the right to collect personal health information if state law authorizes them to do so.
- This right is known as the “public health exemption” from HIPAA. Without this provision, public health surveillance would be much more difficult.
- Missouri law requires the state Department of Health and Senior Services (DHSS) to designate which diseases and conditions must be reported to public health authorities and who must report (Revised Statutes of Missouri 192.020) <http://www.moga.mo.gov/statutes/c192.htm>.
- Through the Code of State Regulations, local public health agencies are designated to receive such information as well (19 CSR 20–20.010–20.080). In the large metropolitan areas, there may also be local ordinances and rules pertaining to disease reporting.
- Some key provisions of the Missouri reporting rules include:
 - a. **Mandated Reporters.** The following individuals and agencies are required to report diseases:
 - Physicians
 - Physician’s assistants
 - Nurses
 - Hospitals
 - Clinics
 - Laboratories
 - Other private or public institutions providing diagnostic testing, screening or care to any person with any reportable disease or condition
 - Persons in charge of a public or private school, summer camp or child or adult care facility
 - b. **Disease Reporting Categories.** There are several categories of reportable diseases listed in 19 CSR 20–20.020, based on the urgency of reporting and response. They are:
 1. Diseases, findings, or agents that shall be reported immediately upon knowledge or suspicion by telephone, facsimile or other rapid communication.

- 1A: Selected high priority diseases, findings, or agents that occur naturally, from accidental exposure, or as the result of a bioterrorism event.
 - 1B: Instances, clusters, or outbreaks of unusual disease or manifestations of illness and clusters or instances of unexplained deaths which appear to be the result of a terrorist act or the intentional or deliberate release of biological, chemical, radiological, or physical agents, including exposures through food, water, or air.
 - 1C: Instances, clusters, or outbreaks or unusual, novel, and/or emerging diseases or findings not otherwise named in this rule, appearing to be naturally occurring, but posing a substantial risk to public health and/or social and economic stability due to their ease of dissemination or transmittal, associated mortality rates, or the need for special public health actions to control.
2. Diseases, findings, or agents that shall be reported within one (1) day of first knowledge or suspicion.
 - 2A: Diseases, findings, or agents that occur naturally, or from accidental exposure, or as a result of an undetected bioterrorism event.
 - 2B: Diseases, findings, or adverse reactions that occur as a result of inoculation to prevent smallpox.
 3. Diseases, findings, or agents that shall be reported within three (3) days of first knowledge or suspicion.
 4. Diseases or findings that shall be reported weekly.
 5. Diseases or findings that shall be reported quarterly.

The list of reportable diseases can be found at:

<http://www.dhss.mo.gov/CommunicableDisease/reportablediseaselist2.pdf>

c. Flexibility.

Missouri's rules include the provision that any unusual outbreak of disease must be reported.

- Allow healthcare providers to share information even if the specific disease or condition isn't on the official list.
- Is especially important for emerging infections, or outbreaks of unknown cause including potential bioterrorist incidents.

d. Multiple entry points for information.

Reports may be made to the local public health agency where the healthcare provider is located, or where the patient lives, or to the state DHSS.

- Provides maximum flexibility for those reporting disease, in the hope of minimizing missed cases.
 - It means, however, that communication across health agency jurisdictions is critical.
- **The entire text of 19 CSR 20-20.020 may be viewed at:**
<http://www.sos.mo.gov/adrules/csr/current/19csr/19c20-20.pdf>
- **The mandated reporting rule** has evolved over time and will continue to do so.
 - New categories of reporters may be added as the healthcare system changes.
 - New diseases may be added as they emerge into public health importance, and old ones may be removed if they are no longer significant threats.
 - Sometimes old diseases take on new significance, as happened when smallpox was added back to the list because of its potential use as a bioterrorist agent.
- **There are no penalties** if a healthcare worker fails to comply with his/her obligations under the disease reporting rules. For this reason, achievement of voluntary compliance is very important. Constant efforts are needed to educate healthcare providers about surveillance in order to keep the information flowing.
- **National Notifiable Diseases Surveillance System.** The Centers for Disease Control and Prevention (CDC) designate nationally notifiable diseases, which can be viewed at:
www.cdc.gov/epo/dphsi/phs/infdis.htm#public. All of these diseases are included on Missouri's list. DHSS passes on reports of nationally notifiable diseases (with no personal identifying information) to the national system.

B. Sources Of Surveillance Data For Infectious Diseases

There are many sources of data for surveillance purposes. Some of the main ones are:

- Morbidity reports (mandated disease reports, laboratory test results, hospital and clinic data)

- Mortality reports (death certificates, medical examiner data)
- Surveillance systems for disease indicators (for example, animal rabies test results, bird and horse West Nile Virus test results)
- Environmental data (laboratory test results on water, milk, and food supplies)
- Student and employee data (for example, school or work absenteeism)
- Drug and biologic utilization (for example, prescriptions and sales figures on over-the-counter medications)
- Population-based surveys (for example, Behavioral Risk Factor Surveillance System, National Health and Nutrition Examination Survey)

C. Forms Of Surveillance

There are two forms of surveillance, passive and active.

1. Passive surveillance.

- Reports are initiated by the information source, such as a physician, laboratory, or hospital.
- Traditional mandated disease reporting is based on this approach.
- The public health agencies put a system in place and then wait for reports to come in by telephone, fax or mail.
 - The standard form for reporting in Missouri is the CD-1 Form. Click here to view the form: <http://www.dhss.mo.gov/CommunicableDisease/index.html>
- Passive reporting is the backbone of surveillance for many diseases, because it is easy and inexpensive for the public health agency.
- However, it can be cumbersome for the reporters, and opportunities to report are often missed.
- Systems that depend solely on passive reporting usually undercount disease incidence.

2. Active surveillance.

- The public health agency initiates contact with the reporting sources on a regular basis (daily, weekly, etc.).
- Contacts may be made by telephone, electronically, or through on-site record review by public health agency staff.
- With active surveillance, reports are solicited even if they indicate zero disease activity for that time period.

- Active surveillance often takes the form of sentinel surveillance. This can involve regular contact with a selected set of sentinel sites or providers, or using animals or insect vectors as “sentinels” for a particular disease.
- Electronic medical records systems have opened up many new possibilities for active surveillance.

Example: DHSS is developing a process that will allow hospital laboratories to submit electronic files of test results directly into the Missouri Health Surveillance Information System (MOHSIS) computer system. The data will then be processed and alert messages will be sent to local public health agency staff. This is expected to reduce the data entry workload and increase the completeness, accuracy and timeliness of laboratory reports.

D. Modern Surveillance Practice

Modern surveillance practice combines active and passive methods to get the most complete picture possible.

Example:

Influenza surveillance in Missouri includes the following information:

- Summary case counts collected by local public health agencies (LPHAs) from sentinel healthcare providers. This information is compared with reporting from previous time periods to track the seasonal epidemic.
- Weekly reports from healthcare providers (34 healthcare providers in 2005) who participate in CDC’s US Influenza Sentinel Provider Surveillance Network (US ISPSN). They report the number of patients they have seen that week, broken down by age group, and how many of them showed influenza-like illness. This information is compared to a national baseline.
- Results of laboratory tests performed by the State Public Health Laboratory, and of more specialized testing performed on some isolates by the CDC laboratory (passive surveillance). This information is used with other test results from around the world, to determine influenza vaccine strains for the coming year.

All of these sources of surveillance information are combined into a weekly report posted on the DHSS website.

Example:

For vector-borne diseases, some combination of active and passive surveillance of both humans and animals may be needed. Surveillance for arboviruses such as West Nile Virus (WNV) is a good example. In Missouri, arboviral surveillance includes humans, horses, birds and mosquitoes.

- **Dead birds** are collected and submitted to DHSS by (LPHAs) for laboratory testing. This may be the only early warning that local spread of the WNV virus is occurring.
- **Mosquito trapping** is the best tool to quantify the intensity of virus transmission in an area. In 2004 over a dozen LPHAs (covering 59% of the Missouri population) trapped mosquitoes, collected them, and either shipped them for testing or tested them locally and provided the results to DHSS.
- **Equine (horse) surveillance** is a passive system that relies on veterinarians to report. Equine WNV is rare now that a vaccine is in wide use.
- **Active surveillance for human cases.** During the peak of WNV activity in 2001–2003, sentinel healthcare providers were contacted by LPHAs at least two times per week.
- **Passive surveillance for human cases.** This is mostly laboratory-based, with the majority of reports coming from the State Public Health Laboratory.

2. DATA ANALYSIS →How do we make sense of it?

- Before surveillance data can be interpreted, they must be organized and analyzed.
- Computer systems such as MOHSIS are frequently used to compile the information and analyze it.
- We must always take care to clearly define the questions we are trying to answer through data analysis, to make sure the results are useful for public health action.
- In some circumstances, for example the very early stages of investigating a suspected outbreak, hand tallies of information about suspected cases may be useful.

A. Determine the number of cases.

- Every reportable disease has an official case definition that is used to determine whether a case should be “counted” in the surveillance system.
- The case definition is usually a combination of symptoms and laboratory test results, and is defined by CDC for nationally notifiable diseases.
- Case definitions reflect different levels of certainty. There are definitions for suspect vs. probable vs. confirmed cases.
- Each case reported to the system should be evaluated in relation to the case definition, as a basic quality assurance mechanism.

B. Calculate incidence rates.

- Rates take into account the size of the population so comparisons can be made across geographic areas.
- By using rates, the incidence in one county can be compared to state or national incidence of the disease.
- For more information about rates, see Module IV, Statistical Measures.

C. Analyze the data by person, place and time.

- Person includes variables such as age, sex, and race, as well as risk factors such as childcare or food handler status.
- Place may be a nation, state, county, city or even zip code or census tract.
- Time is usually represented as date of onset of illness, grouped by day, week, month or year.
- More detailed information about these epidemiologic variables will be given in Module V, Displaying and Interpreting Epidemiologic Variables.

D. Current surveillance data are compared with some expected value to identify how they differ and to assess their importance.

One way to do this is by comparing the incidence rate in the current time period with past incidence in the same jurisdiction. This may be incidence as of the same time last year, or a measure of the “usual” incidence, such as a five-year median (see Module IV, Statistical Measures).

E. It is also useful to compare with other jurisdictions.

Is the current incidence rate in County A significantly higher than that in surrounding counties, or the state or national average? If so, that may indicate a problem that needs to be followed up.

F. Spatial analysis may be helpful.

- Modern technology such as Geographic Information Systems (GIS) can create very informative analyses.
- The locations of cases can be viewed in relation to environmental features and/or population characteristics.

3. INTERPRETATION OF DATA → What can we learn from all these numbers?

If the analysis shows that the incidence of a disease is different from what you would expect, then further investigation should be done.

- For some diseases, this is true even if the number of cases is small.
Example: even a single case of a potential bioterrorist agent (such as anthrax) or a vaccine-preventable disease (such as measles) should be investigated.
- There are several possible explanations for changes in surveillance data. Explanations that should be considered include (but are not limited to):
 - An outbreak
 - An intentional exposure (such as a bioterrorist attack)
 - A newly emerging infection
 - Improved diagnosis (new laboratory test, increased physician awareness)
 - Increased awareness of the disease and/or the need to report it
 - A gradual increase or decrease in incidence due to environmental or population changes, or changes in the disease agent
 - A disease following its natural seasonal or “secular” (years-long) cycle
 - Changes in the surveillance system (new data collection system, loss of a reporting source, addition of a new source, change in case definition, etc.)

4. DISSEMINATION OF THE INFORMATION → Sharing the Results

- The purpose of sharing surveillance information is to inform those who need to know, and to motivate those who need to do something.

Examples:

- keeping healthcare providers informed about current trends can help them diagnose and treat their patients better.
- Keeping policymakers informed can assist them in allocating resources and updating rules, ordinances and laws.

- And keeping public health professionals and administrators up to date will help them design and carry out better programs.
- Surveillance data generally flow “up” through the public health system, from local reporters to local public health agencies, to DHSS, to CDC, to the World Health Organization (WHO).
- As with any good communication, the information must be tailored to the audience.
 - Public health workers may need more detailed information than other groups.
 - In general, reports should be brief and clear, and should highlight the information most important to the reader.
 - **Examples** of reports issued by Missouri may be found at <http://www.dhss.mo.gov/Influenza/Reports.html>
 - **Examples** of reports issued by CDC may be found at www.cdc.gov/mmwr.

5. LINKS TO PUBLIC HEALTH PRACTICE → Information for Action

- The only reason to carry out public health surveillance is to use the information to improve the health of the public.
- Many of the ways this is done were discussed earlier in the section, “Why Do We Do Public Health Surveillance?”

VI. EVALUATION AND IMPROVEMENT OF PUBLIC HEALTH SURVEILLANCE

- An efficient, effective public health surveillance system is essential to protect the public’s health.
 - Because disease agents, host factors, the environment and the healthcare system are constantly changing, we must continuously assess and improve our surveillance systems.
 - A periodic evaluation can provide information for improvement.
- Probably the most important question to ask in evaluating a surveillance system is: “How useful is it?” Some ways to look at usefulness are:
 - Does the public health surveillance system contribute to the prevention and control of diseases?

- Does it contribute to an improved understanding of the public health implications of diseases (numbers of people affected, severity of illness, burden of hospitalization, etc.)?
- Can it help determine that a disease that was previously thought to be unimportant is actually important?
- Does it provide data for performance measurement, including health indicators used for assessing community needs and evaluating health programs?
- It may be helpful to list and describe the actions taken as a result of analysis and interpretation of the data from the system, and to identify who has used the information to make decisions and take actions.

CDC has identified the attributes of a good surveillance system. They are:

- ◆ Simplicity. Is the system simple in structure and easy to operate? Are there steps in the process that could be combined or eliminated? Surveillance systems should be as simple as possible while still meeting their objectives. The simpler the system, the more acceptable and timely it is likely to be. Simpler systems also take fewer resources to operate.
- ◆ Flexibility. Can the system adapt to changing information needs or operating conditions with little additional time, personnel, or allocated funds? Does the system use standard data formats (e.g., in electronic data interchange) that can be easily integrated with other systems?
- ◆ Data quality. Is the information in the system complete and valid? One easy way to check this is to examine the percentage of "unknown" or "blank" responses to items on surveillance forms. However, a full assessment might require a special study in some cases.
- ◆ Acceptability. Are people and organizations willing to participate in the surveillance system?
- ◆ Sensitivity. What proportion of actual cases are detected by the surveillance system? This may require a check against other data sources, such as hospital discharge data or a laboratory record review. Can the system monitor changes in the numbers over time, and detect outbreaks?
- ◆ Predictive value positive. What proportion of reported cases actually have the disease under surveillance? This can be assessed by looking at how many of the reported cases meet the case definition, or how many are eventually classified as confirmed.

- ◆ Representativeness. Does the system accurately describe the occurrence of the disease over time and its distribution in the population by place and person? Or is it “skewed” toward certain age groups, ethnic groups, geographic areas, or healthcare providers?
- ◆ Timeliness. How quickly does the system receive and process information? This can be assessed by looking at the speed between steps in the system.
- ◆ Stability. How reliable is the system (reliability is the ability to collect, manage, and provide data properly without failure). Is the system operational when it is needed?

By evaluating these attributes, we can identify ways to improve the system. For more complete information about these criteria and recommended methods for evaluation of surveillance systems, go to www.cdc.gov/mmwr/preview/mmwrhtml/rr5013a1.htm.

Limitations Of Disease Reporting Systems

- It is important to remember that no formal system can take the place of good relationships and communication networks in a community.
- There is no substitute for good collegial relationships among the public health agency, healthcare providers, and other key informants such as the school system.
- Time spent building those relationships can help assure that the public health agency is kept “in the loop” when problems occur.

PRINCIPLES OF INFECTIOUS DISEASE EPIDEMIOLOGY

MODULE IV – STATISTICAL MEASURES

Much of this module was adapted from the Centers for Disease Control and Prevention (CDC) “Principles of Epidemiology, Second Edition, An Introduction to Applied Epidemiology and Biostatistics.”

Please note: Because of formatting constraints, the formulas in the outline may not appear in correct mathematical format, however, they do appear correctly in the course module.

I. INTRODUCTION

Module IV is designed to prepare public health workers to meet the following objectives:

- Define the most common statistical frequency measures used in infectious disease epidemiology
- Construct a frequency distribution
- Calculate and interpret the following statistical measures:
 - Ratios
 - Proportions
 - Incidence rates, including attack and secondary attack rates
 - Prevalence
 - Mortality rates
 - Relative risk and odds ratio
- Choose and apply the appropriate statistical measures

II. HOW DO WE USE STATISTICAL MEASURES IN EPIDEMIOLOGY?

- Statistics are used to summarize the data collected through disease surveillance or an outbreak investigation, so we can learn from the data. Calculating statistics helps us to:
 - describe risk
 - make comparisons among communities and smaller definable groups
 - identify high-risk groups
 - develop hypotheses about the cause(s) of disease
- Why do we need to describe and compare risk?
 - Differences in risk among different populations can provide clues for investigation of what caused one group to have a higher risk.
 - If causes can be identified, then perhaps prevention and control measures can be identified too.

- The most common statistical measures used in field epidemiology are “frequency measures,” which are simply ways of counting cases and comparing their characteristics. In contrast with statistics used in epidemiological research, frequency measures are relatively easy to calculate and use.

III. FREQUENCY DISTRIBUTIONS

- When we collect data about disease cases, we must put them in some kind of order. The most basic way to do this is to organize a “line listing”. Example: <http://www.dhss.mo.gov/CDManual/CDsec30.pdf> (scroll down to page 77).
- A line listing is actually a simple database, in which each row represents a case of the disease we are investigating. Each column contains information about one characteristic, called a “variable.”
- Look at the data in **Table 1**

Table 1
Neonatal Listeriosis, General Hospital A, Costa Rica, 1989

ID	Sex	Culture Date	Symptom Date	DOB	Delivery Type	Delivery Site	Outcome	Admitting Symptoms
CS	F	6/2	6/2	6/2	Vaginal	Del Rm	Lived	Dyspnea
CT	M	6/8	6/8	6/2	C-section	Oper Rm	Lived	Fever
WG	F	6/15	6/15	6/8	Vaginal	Emer Rm	Died	Dyspnea
PA	F	6/15	6/12	6/8	Vaginal	Del Rm	Lived	Fever
SA	F	6/15	6/15	6/11	C-section	Oper Rm	Lived	Pneumonia
HP	F	6/22	6/20	6/14	C-section	Oper Rm	Lived	Fever
SS	M	6/22	6/21	6/14	Vaginal	Del Rm	Lived	Fever
JB	F	6/22	6/18	6/15	C-section	Oper Rm	Lived	Fever
BS	M	6/22	6/20	6/15	C-section	Oper Rm	Lived	Pneumonia
JG	M	6/23	6/19	6/16	Forceps	Del Rm	Lived	Fever
NC	M	7/21	7/21	7/21	Vaginal	Del Rm	Died	Dyspnea

Abbreviations:

Vaginal = vaginal delivery
 Del Rm = delivery room
 Oper Rm = operating room
 Emer Rm = emergency room

- How many of the cases were male? We can easily pick out that information because there are only a few cases.
- But with a larger database, we cannot get that information at a glance. We must summarize variables into tables called “**frequency distributions.**”
- **A frequency distribution**
 - shows the values a variable can take, and
 - the number of people or records with each value.

- Example: suppose we are investigating an outbreak in an elementary school. We could construct a frequency table that shows how many of the ill children were in each classroom.

Table 2
Distribution of cases by classroom
Brown School, Missouri, June 2005

Classroom	Number of Cases
101	25
102	43
103	32
104	0
105	8
106	2
Total	110

Please notice some key points about the table format:

- All possible values of the variable, in this case all classrooms, are listed, even if there were no cases for some values.
- Each column is properly labeled.
- The total is given in the bottom row.

Now we can tell at a glance which rooms were most affected, and which were not affected at all.

- **Variables**

- The **values of a variable** may be
 - numbers (for example, number of tacos eaten);
 - an ordered numerical scale (for example, age); or,
 - categories (for example, ill or well), called a “nominal scale” because the categories are named.
- In epidemiology we often deal with variables that have only two categories, like alive or dead, ill or well, did or did not eat the potato salad.
- Any of these types of data may be summarized in a frequency distribution (See **Table 3**, which shows a variable with only two possible values).

Table 3
Influenza vaccination status among residents
Nursing Home A, Missouri, December 2005

Vaccinated?	Number
Yes	76
No	125
Total	201

IV. RATIOS, PROPORTIONS AND RATES

- Three kinds of frequency measures are used with two-category variables (also called dichotomous variables). These frequency measures are
 - Ratios
 - Proportions, and
 - Rates.
- Before you learn about specific measures, it is important to understand the relationship between the three types of measures and how they differ. All three measures are based on the same formula:

$$\text{Ratio, proportion, rate} = x/y \times 10^n$$

- In this formula, x and y are the two quantities that are being compared.
- The formula shows that x is divided by y .
- 10^n is a constant that we use to transform the result of the division into a uniform quantity.
- The size of 10^n may equal 1, 10, 100, 1000 and so on depending on the value of n .

Example:

$$10^2 = 10 \times 10 = 100$$

$$10^3 = 10 \times 10 \times 10 = 1000$$

$$10^5 = 10 \times 10 \times 10 \times 10 \times 10 = 100,000$$

A. Ratios:

- A ratio is used to compare the occurrence of a variable in two different groups.
- These may be two completely independent groups, or one may be included in the other.
- For example, we could compare the sex of children attending an immunization clinic in either of the following ways:

$$1) \quad \frac{\text{female}}{\text{male}} \qquad \text{or} \qquad 2) \quad \frac{\text{female}}{\text{all}}$$

In the first example, x (female) is completely independent of y (male). In the second example, x (female) is included in y (all). This second type of ratio, called a proportion, is examined in more detail in the next section of this module.

B. Proportions:

- The second type of ratio, in which x is part of y , is also called a proportion (as in the previous (female/all) example).
- Proportions are usually expressed as percentages.

Examples

➤ Independent x and y :

During the first 9 months of national surveillance for eosinophilia-myalgia syndrome (EMS), CDC received 1,068 case reports that specified sex; 893 cases were in females, 175 in males. Here is the method for calculating the **female-to-male ratio** for EMS.

1. Define x and y :
 x = cases in females
 y = cases in males
2. Identify x and y :
 x = 893
 y = 175
3. Set up the ratio x/y : 893/175
4. Reduce the fraction so that one value equals 1. Female to male =
 $893/175 = 5.1/1$
5. Express the ratio in one of the following ways: 5.1 to 1, or 5.1:1, or 5.1/1

Thus, there were just over 5 female EMS patients for each male EMS patient reported to CDC.

➤ x included in y :

Based on the same data, here is the method for calculating the **proportion** of EMS cases that were male.

1. Define x and y :
 x = cases in males
 y = all cases
2. Identify x and y :
 x = 175
 y = 1,068
3. Set up the ratio x/y : 175/1,068
4. Reduce the fraction so that one value equals 1. Divide the smaller number by the larger number: $175/1,068 = 0.16/1$

5. Proportions are usually expressed as percentages, so the value of the constant (10^n) = 10^2 = 100:
 $0.16 \times 100 = 16$ (16%)

Thus, 16% of the reported EMS cases were in males.

C. Rates:

- The third type of frequency measure used with two-category (dichotomous) variables is a rate.
- Rates have the added dimension of time. Rates measure the occurrence of an event in a population over time.
- The basic formula for a rate is:

$$\text{Rate} = \frac{\text{number of cases occurring during a given time period}}{\text{population at risk during the same time period}} \times 10^n$$

- Rates are always specific to a particular population. They reflect groupings of people based on time, place and person.
 - Time: a specific year, month, week, day or hour
 - Place: country, state, county, city, township, school, institution, area
 - Person: age, sex, membership in some group or class
- Rates take into account the size of the population, so comparisons can be made across different population groups.
 - By using rates instead of raw numbers, the occurrence of disease in one group can be fairly compared with another.
 - Example: males with females; one county with another; Missouri with Arkansas or the US.
- To calculate a rate, we must have an estimate of the population at risk during a specific time period for the denominator.
 - Ratios and proportions do not require this.
 - Earlier, we calculated ratios and proportions of EMS cases without knowing the number of people at risk of EMS.
 - Rates may be harder to get, because accurate denominator data may not be available for small, localized population groups.

To summarize:

- All three of these frequency measures are calculated in basically the same way. In practice, we use:
 - a **ratio** to compare two independent groups,
 - a **proportion** to compare one group with a larger one to which it belongs, and
 - a **rate** to measure an event in a population over time.
- Ratios, proportions, and rates are used in infectious disease epidemiology to describe **morbidity** (disease) and **mortality** (death).

V. MORBIDITY FREQUENCY MEASURES

- Several standard measures are used to measure and describe the frequency of disease.
- Each measure has its appropriate uses, depending on the situation and the information available to the epidemiologist.
- The two main types of rates are:
 - **incidence**
 - **prevalence**.

A. Incidence Rates

- Incidence rates are the most common way of measuring and comparing the frequency of disease in populations.
- Incidence is a measure of risk.
- When Population A has a higher incidence of a disease than Population B, we can say that Population A has a higher risk of developing the disease than Population B. If it is a lot higher, we could say that Population A is a **high-risk** group for that disease.
- **Table 4** shows the three types of incidence rates we will study, along with their formulas. We will discuss each of these in more detail.

Table 4
Frequently Used Measures of Morbidity

Measure	Numerator (x)	Denominator (y)	Expressed per Number at Risk (10^n)
Incidence Rate	# new cases of a specified disease reported during a given time interval	Average population during time interval	Varies: 10^n where $n = 2, 3, 4, 5, 6$
Attack Rate	# new cases of a specified disease reported during an epidemic period	Population at start of the epidemic period	Usually a percentage: 10^n where $n = 2$
Secondary Attack Rate	# new cases of a specified disease among contacts of known cases	Size of contact population at risk	Usually a percentage: 10^n where $n = 2$

We will discuss each of these in more detail.

1) Incidence

The basic **incidence rate** (sometimes called just **incidence**) is a measure of the frequency with which a disease occurs in a population over a period of time. The formula for calculating an incidence rate is:

$$\text{Incidence Rate} = \frac{\text{new cases occurring during a given time period}}{\text{population at risk during the same time period}} \times 10^n$$

- **The numerator (x)**
 - should include only **new** cases of the disease that occurred during the specified period.
 - should not include cases that occurred or were diagnosed earlier.
 - This is very important when working with chronic infectious diseases such as tuberculosis, malaria and HIV.

- **The denominator (y) is the population at risk.**
 - This means that the people included in the denominator should be able to develop the disease in question during the time period covered. In practice, we usually use census data for the denominator.
 - The denominator should also represent the population from which the cases in the numerator arose. The population may be defined by geographic area (e.g., St. Francois County) or by membership in a specific group (e.g., employee of Company X, student at School Y). If we are studying a specific group such as students in a school or residents in a long term care facility, we should use a census of that population for an exact denominator.

- **Any value of *n*** may be used in calculating incidence.
 - The epidemiologist should always make it clear what *n* value was used. National surveillance systems use a value of 10^5 or 100,000.
 - A good rule of thumb is to choose a value for 10^n so that the smallest rate calculated is a small whole number (for example, $4.2/100$, not $0.42/1,000$ or $0.042/10,000$). This is just easier for the reader to understand.
 - When comparing two incidence rates, always be sure that the same value of *n* was used in calculating both rates.

Disease incidence rates imply a change over time, from health to disease. So **the period of time must be specified**. For surveillance purposes this is usually one calendar year, but any time period may be used as long as it is stated.

Example

In 2003, 335,104 new cases of gonorrhea were reported among the US civilian population. The 2003 mid-year US civilian population was estimated to be 290,788,976. For these data we will use a value of 10^5 for 10^n . We will

calculate the 2003 gonorrhea incidence rate for the US civilian population using these data.

1. Define x and y : x = new cases of gonorrhea in US civilians during 2003
 y = US civilian population in 2003
2. Identify x , y , and 10^n : $x = 335,104$
 $y = 290,788,976$
 $10^n = 10^5 = 100,000$
3. Calculate $(x/y) \times 10^n$:

$$\frac{335,104}{290,788,976} \times 10^5 = .001152 \times 100,000 = 115.2 \text{ per } 100,000 \text{ or}$$

approximately 1 reported case per 1,000 population

The **numerator** of this incidence rate

- reflects new cases of gonorrhea that occurred or were diagnosed during the specified period.
- does **not** include cases that occurred or were diagnosed earlier.

The **denominator** is the population at risk.

- it represents the population from which the gonorrhea cases arose, in this case the US civilian population.

Notice that the numerator was limited to civilian cases. Therefore, we had to restrict the denominator to civilians as well.

2) Attack Rate

An **attack rate** is a specific type of incidence rate. It is calculated for a narrowly defined population observed for a limited time, such as during an outbreak. It is usually expressed as a percentage, so 10^n equals 100.

For a defined population (the population at risk), during a limited time period:

$$\text{Attack Rate} = \frac{\text{\# of new cases among the population during the period}}{\text{population at risk at the beginning of the period}} \times 100$$

- The attack rate is a measure of the **probability** or **risk** of becoming a case.
- Remember, attack rates have some special characteristics:
 - Highly specific by “person” variables
 - Limited by “place” variables
 - Time period is usually brief
 - Usually expressed as a percentage

Example

Of 75 persons who attended a church picnic, 46 subsequently developed a gastrointestinal illness. To calculate the attack rate of GI illness we first define the numerator and denominator:

x = Cases of GI illness occurring within the incubation period
 for GI illness among persons who attended the picnic = 46
 y = Number of persons at the picnic = 75
 Then, the attack rate for GI illness is $\frac{46}{75} \times 100 = 61\%$

In this example, we could say that among persons who attended the picnic, the probability of developing GI illness was 61%, or the risk of developing GI illness was 61%.

- Attack rates are usually calculated several times during the course of an outbreak investigation. The first time, early in the outbreak, the attack rate might be calculated as follows:

$$\text{Attack Rate} = \frac{\# \text{ of new cases in the community during the period}}{\text{the population of the community}} \times 100$$

- This would help us see whether the current situation is “unusual” compared with other time periods or communities.
- In the course of an investigation, attack rates will be recalculated as new cases are identified, diagnoses are confirmed, and other information comes to light.
- Attack rates may be needed for various subgroups within the community.
- Selecting the appropriate numerators and denominators is very important and can be a challenge.
 - Ideally, the denominator should include the smallest definable area or group that contains all the known cases.
 - In practice, however, accurate population counts may not be available.
 - For example, we may not be able to find out exactly how many people ate in a particular restaurant during a specified time.

3. Secondary Attack Rate

A **secondary attack rate** measures the frequency of new cases of a disease among the contacts of known cases. This can be very important for diseases that are spread from person to person, such as tuberculosis, measles, shigellosis, and varicella. The formula is:

$$\text{Secondary Attack Rate} = \frac{\# \text{ cases among contacts of primary cases during the period}}{\text{total number of contacts}} \times 100$$

- Secondary attack rates are often calculated for households.
 - To calculate the number of household contacts (denominator), subtract the number of primary cases from the total number living in the households.

- In some situations, contacts in other settings may be investigated (for example, residents of a homeless shelter, or people who work in a specific building). The calculation is done in the same way as for household contacts.

Example

Seven cases of hepatitis A occurred among 70 children attending a childcare center. Each infected child came from a different family. The total number of persons in the 7 affected families was 32. One incubation period later, 5 family members of the 7 infected children also developed hepatitis A. We will calculate the attack rate in the childcare center and the secondary attack rate among family contacts of those cases.

1. Attack rate in childcare center:

x = cases of hepatitis A among children in childcare center = 7
 y = number of children enrolled in the childcare center = 70

$$\text{Attack rate} = \frac{x}{y} \times 100 = \frac{7}{70} \times 100 = 10\%$$

2. Secondary attack rate (refer to Figure 1):

x = cases of hepatitis A among family contacts of children with hepatitis A = 5

y = number of persons at risk in the families (total number of family members – children already infected) = 32 – 7 = 25

$$\text{Secondary attack rate} = \frac{x}{y} \times 100 = \frac{5}{25} \times 100 = 20\%$$

B. Rate Ratio is another tool that is helpful for comparing rates between groups.

A **rate ratio** compares the rates of disease in two groups that differ by demographic characteristics or exposure history. The rate for the group of primary interest is divided by the rate for a comparison group.

$$\text{Rate Ratio} = \frac{\text{rate for group of primary interest}}{\text{rate for comparison group}} \times 1$$

Rate ratios may be calculated for incidence rates (including attack rates) or for mortality rates, discussed later.

Example

The Association of Interested Persons held their annual conference during the first week in June. There were two events: a dinner meeting on Wednesday evening (75 attendees), and a luncheon awards ceremony on Thursday at noon (60 attendees). Twenty (20) of the 75 Wednesday dinner participants subsequently developed signs and symptoms of gastrointestinal illness; 5 of the 60 luncheon participants became ill. Calculate the rate ratio to help determine which event may have been the source of the illness. The rate ratio is calculated as follows:

1. Calculate the attack rate for the dinner meeting:
 x = number of ill persons attending the dinner meeting
 y = number of persons attending the dinner meeting
 $\text{attack rate} = (x/y) \times 100 = (20/75) \times 100 = 27\%$
2. Calculate the attack rate for the luncheon:
 x = number of ill persons attending the luncheon
 y = number of persons attending the luncheon
 $\text{attack rate} = (x/y) \times 100 = (5/60) \times 100 = 8\%$
3. Calculate the rate ratio:

$$\text{Rate ratio} = \frac{\text{rate for group of primary interest}}{\text{rate for comparison group}} \times 1 = (27/8) \times 1 = 3.4$$

The dinner meeting attendees were 3.4 times more likely to become ill than those who attended the luncheon.

Now that you know how to calculate and use each type of morbidity rate, you have mastered some important tools for investigating infectious diseases.

C. Prevalence (Prevalence Rates)

Prevalence is the proportion of people in a population who have a particular disease at a specified point in time, or over a specified period of time.

- The numerator includes not only new cases, but also old cases (people who remained ill during the specified point or period in time). A case is counted in prevalence until death or recovery occurs.
- This makes prevalence different from incidence, which includes only new cases in the numerator.
- Prevalence is most useful for measuring the burden of chronic diseases such as tuberculosis, malaria and HIV in a population. The formula for calculating prevalence is:

$$\text{Prevalence} = \frac{\text{all new and pre-existing cases during a time period}}{\text{population during the same time period}} \times 10^n$$

Point vs. Period Prevalence

The amount of disease present in a population obviously changes over time. Sometimes, we want to know how much of a particular disease is present in a population at a single point in time, a sort of snapshot view.

- **Point Prevalence:** For example, we may want to find out the prevalence of TB in Community A today. To do that, we need to calculate the **point prevalence** on a given date.
 - The numerator would include all known TB patients who live in Community A that day. That information could be determined from a TB case registry.
 - The denominator would be the population of Community A that day.

- Example: A review of patients reported to the tuberculosis registry in Midville revealed that as of July 1, 2005 there were 35 cases that had not yet completed therapy. The most recent population estimate for Midville was 57,763. The prevalence of TB in Midville on July 1, 2005 was:

$$\frac{35}{57,763} \times 10,000 = 6.1 \text{ per } 10,000 \text{ people}$$

- Point prevalence is useful in comparing different points in time to help determine whether an outbreak is occurring. In this case, we could also calculate point prevalence of TB for July 1, 2004, July 1, 1995 or other relevant points of comparison.

- **Period Prevalence:** At other times, we want to know how much of a particular disease is present in a population over a longer period. We would use **period prevalence** to do that.

- Period prevalence is calculated in exactly the same way as point prevalence, except the numerator is the number of people who had the disease at any time during a specified time period.
- Period prevalence can be calculated for a week, month, year, decade, or any other specified length of time.
- Example: Midville's TB registry indicates that during 2004, there were 89 new and pre-existing TB cases. The prevalence of TB in Midville in 2004 was:

$$\frac{89}{57,763} \times 10,000 = 15.4 \text{ per } 10,000 \text{ people}$$

- When comparing prevalence figures:
 - be sure that the length of the time period is the same. The prevalence of TB in Midville in 2004 should only be compared with the prevalence during other one-year time periods (2003, 1994, etc.).
 - It cannot be compared with the point prevalence on July 1, 2005, or with the period prevalence during a week or month.
 - Prevalence may be compared among different diseases or different populations, as long as the same length of time is used.

Example: Comparing Prevalence and Incidence

Two surveys were done of the same community 12 months apart. Of 5,000 people surveyed the first time, 25 had antibodies to histoplasmosis. Twelve months later, 35 had antibodies, including the original 25. We will calculate the prevalence at the second survey, and compare the prevalence with the 1-year incidence.

1. Prevalence at the second survey:
 - x = antibody positive at second survey = 35
 - y = population = 5,000
 - $(x/y) \times 10^n = 35/5,000 \times 1,000 = 7 \text{ per } 1,000$

2. Incidence during the 12-month period:

x = number of new positives during the 12-month period = $35 - 25 = 10$

y = population at risk = $5,000 - 25$ (already infected) = $4,975$

$(x/y) \times 10^n = 10/4,975 \times 1,000 = 2$ per 1,000

Prevalence is based on both incidence (risk) and duration of disease.

High prevalence of a disease within a population may reflect high risk, or it may reflect prolonged survival without cure. Conversely, low prevalence may indicate low incidence, a rapidly fatal process, or rapid recovery.

VI. MORTALITY FREQUENCY MEASURES

- Mortality rates measure the frequency of occurrence of death in a defined population during a specified interval.
- There are several specific kinds of mortality rates, but we will focus only on the ones that are used most often in infectious disease epidemiology.
- To calculate a simple mortality rate, we need to know the number of deaths in a given population during a specified time period, and the size of the population in which the deaths occurred. The basic formula is:

$$\text{Mortality rate} = \frac{\text{deaths occurring during a given time period}}{\text{size of the population in which the deaths occurred}} \times 10^n$$

The most commonly used values for 10^n are 1,000 and 100,000.

A. Crude Mortality Rate

- The crude mortality rate is the mortality rate from all causes of death for a population during a specified time period.
- The denominator is the population at the mid-point of the time period.
- For example, the crude mortality rate for Missouri in 2003 was 896 deaths per 100,000 people.

B. Cause-specific Mortality Rate

- This is the mortality rate from a specified cause for a population during a specified time period.
- The numerator is the number of deaths from that cause, and the denominator remains the size of the population at the mid-point of the time period.
- For example, the tuberculosis death rate for the US in 2002 was 0.3 per 100,000 (or 3 per 1,000,000).

C. Other Specific Mortality Rates

- Specific mortality rates may be calculated for population subgroups defined by age, sex, race, or other demographic factors.
- Combinations of factors are often used.

- For example, the mortality rate attributed to HIV among 25-to-44-year-olds in the US in 1987 was:

$$\text{HIV Mortality Rate} = \frac{9,280 \text{ deaths}}{77,600,000 \text{ aged 25-44 yrs}} \times 100,000 = 12 \text{ per } 100,000$$

This is an example of a cause- and age-specific mortality rate.

VII. RELATIVE RISK AND ODDS RATIO

The last two types of frequency measures we will study are **relative risk** (also called **risk ratio**) and **odds ratio**. These statistics are used in outbreak investigations and will be discussed again in the workshop portion of this course.

A. Relative Risk or Risk Ratio (RR)

- Compares the risk of disease or death in two groups.
 - The two groups may be defined by a demographic factor such as sex (for example, male vs. female).
 - More commonly, they are defined by a difference in their exposure to a suspected risk factor for disease (for example, ate the potato salad or didn't).
 - Often, the group of primary interest is labeled "exposed," and the comparison group is called "unexposed."
 - The group of primary interest goes into the numerator, and the comparison group is the denominator.

$$\text{Risk Ratio (Relative Risk)} = \frac{\text{Risk for group of primary interest}}{\text{Risk for comparison group}} \times 1$$

- "Risk" is defined as an incidence rate or attack rate of the disease in each group. To calculate RR, a two-by-two table is set up as shown:

Table 5
Number of Cases of Disease X by Sex, Smallville, 2004

	Disease X		Total
	Yes	No	
Female	a 46	b 1,438	1,484
Male	c 18	d 1,401	1,419

The term "two-by-two" refers to the two variables (sex and disease status), each with two categories. The outcome (illness or not) is shown at the top of the table and exposure or risk factor is shown along the left side. Note the letters assigned to each cell of the table (a-d). They are important in calculating the risk in each group.

Example

Using the data in the table above, we can calculate the relative risk of Disease X for females vs. males. First, we must calculate the risk of illness among females and among males:

$$\text{Risk among females (incidence)} = \frac{a}{a+b} = \frac{46}{1,484} = .031 \times 100 = 3.1\%$$

$$\text{Risk among males (incidence)} = \frac{c}{c+d} = \frac{18}{1,419} = .013 \times 100 = 1.3\%$$

To calculate the RR for females vs. males, females are considered the group of primary interest and males are the comparison group. The formula is:

$$\text{Risk ratio (Relative Risk)} = \frac{3.1\%}{1.3\%} = 2.4$$

So we can say that the risk of Disease X in females appears to be 2.4 times higher than the risk in males. If the RR is 1.0, that means the risk of disease is equal in the two groups. If the RR is greater than 1.0, then the group of interest has a higher risk of disease. If the RR is less than 1.0, then the group of interest has a lower risk of disease.

- However, before we can interpret RR figures, they must be subjected to a test of statistical significance such as the Chi square or some variation of it. This helps us judge the probability that the result could have occurred by chance alone. A probability of less than 5%, expressed as $p < .05$, is commonly used as a cutoff for statistical significance in field epidemiology.
- We will not teach more about statistical significance in this course, but the student should be aware that RR is affected by factors such as population size, and cannot stand alone. Statistical consultation is readily available from DHSS for the field epidemiologist.

B. Odds Ratio

The RR can only be calculated if incidence data are available. The **Odds Ratio (OR)** may be used in situations where we do not have denominator data to calculate incidence rates.

- The odds ratio is used frequently in case/control studies, which we will cover in more detail in the workshop portion of this course.
- In a case/control study, ill persons' characteristics and exposures are compared with those of well persons ("controls") selected from the same population in which the outbreak occurred.
 - Example: in an outbreak suspected to stem from exposure to contaminated food at a restaurant, the ill persons' food selections

could be compared with those of some well people who also ate at the restaurant the same day. This could be done even if we didn't know exactly how many people ate at the restaurant that day.

- A two-by-two table is constructed, just like the one used to calculate RR, with the same letters (a-d) used to label the cells. The OR is calculated by multiplying across the cells.

Example:

Table 7
Number of Cases of Disease X by Exposure History, Smallville, 2004

	Disease X		Total
	Yes	No	
Ate Tuna Casserole	a 46	b 25	71
Didn't Eat Tuna Casserole	c 18	d 40	58

The formula for OR is:

$$\text{Odds Ratio} = \frac{ad}{bc}$$

Where

- a* = number of persons with disease and with exposure of interest
- b* = number of persons without disease, but with exposure of interest
- c* = number of persons with disease, but without exposure of interest
- d* = number of persons without disease and without exposure of interest

- a+c* = total number of persons with disease ("cases")
- b+d* = total number of persons without disease ("controls")

The OR in this example is:

$$\text{Odds Ratio} = \frac{46 \times 40}{25 \times 18} = \frac{1840}{450} = 4.1$$

So those who became ill were 4.1 times as likely to have eaten the tuna casserole. We should probably look a little more deeply into the tuna casserole! We would still need to subject this result to a test of statistical significance (just like we do with the RR) to judge the probability that the result could have occurred by chance alone.

PRINCIPLES OF INFECTIOUS DISEASE EPIDEMIOLOGY

MODULE V – DISPLAYING AND INTERPRETING EPIDEMIOLOGIC VARIABLES

Portions of this module were adapted from the Centers for Disease Control and Prevention (CDC) “Principles of Epidemiology, Second Edition, An Introduction to Applied Epidemiology and Biostatistics, 1998.”

Note: You will need to access the course to view the many examples associated with this module, as they are not included in this outline.

I. INTRODUCTION

Module V is designed to prepare public health workers to meet the following objectives:

- Define the three categories of epidemiologic variables
- Identify the three main methods used to organize epidemiologic data
- Correctly interpret graphic presentations of epidemiologic data
- Choose appropriate display methods and formats for specific kinds of epidemiologic data

II. WHAT ARE EPIDEMIOLOGIC VARIABLES?

A. Epidemiologic variables are characteristics that can be observed and/or measured. They may be characteristics of:

- **Time** - the time of illness or of a relevant event.
Examples: date of exposure or onset of illness.
- **Place** - the environment in which illness occurs.
Examples: place of residence, of work, or of suspected exposure (such as a retail food establishment).
- **Person** - individuals who are infected, ill, or at risk. Examples: age, gender and occupation.

B. We look at epidemiologic variables to:

- identify characteristics that might be important, and
- to form hypotheses about the
 - source,
 - causative agent, and
 - mode of transmission of illness.

C. When we review surveillance data or investigate an outbreak, we are trying to answer these questions:

WHO? Identify individuals and sub-populations at risk of exposure or transmission

HOW? Identify the modes of disease transmission, especially any changes from known transmission patterns

HOW CAN WE INTERVENE? Identify factors or conditions that can be manipulated to modify or prevent disease occurrence and spread

WHAT ARE OUR PRIORITIES? Identify the usual patterns of disease in a population, so we can set priorities and respond more quickly to outbreaks

Remember, ***the ultimate goal of epidemiology is to prevent disease.*** This is done by finding associations between a disease and the characteristics of

- time,
- place, and
- person.

III. METHODS FOR ORGANIZING EPIDEMIOLOGIC DATA

- The field epidemiologist needs to organize data for several reasons:
 - It is a necessary step in data analysis.
 - It helps the epidemiologist visualize patterns and trends, and identify variations from those trends.
 - It provides a useful way to communicate information to others.
- There are three basic methods of organizing epidemiologic data
 - Tables
 - Charts and
 - Graphs

A. Tables

1. In General:

- A table is a set of data, organized into rows and columns.
- Tables are useful for identifying patterns, exceptions, differences and other relationships.
- Tables also serve as the basis for charts and graphs.

A table should be self-explanatory. It should convey all the information the reader needs to understand the data, including:

- A clear and concise title
- Row and column labels (clear and concise)
- Totals for rows and columns
- Footnotes to explain any codes, abbreviations, or symbols

- Footnotes that note any data excluded from the analysis
- Clear identification of the source of the data in a footnote

2. Types of Tables

a) *One Variable Tables.* The simplest form of table has only one variable. That is the frequency distribution, which was discussed in Module IV, Statistical Measures.

- To review briefly, in a frequency distribution table, the first column shows the values (or categories) of a variable, and the second column shows the number of people or records that fall into each value or category.
- Often, there is a third column that lists the percentage of persons or events in each category.
- Sometimes a one-variable table shows the cumulative frequency or cumulative percent.

b) *Two- and Three-Variable Tables.*

- Additional columns are added to a table to show counts by a second variable, for example age and sex.
- As you learned in Module IV, a two-by-two table is used to show cross-tabulated data. Another name for a two-by-two table is a contingency table. Such tables display two variables, each with two categories.
- It is usually better to use only one- or two-variable tables. Sometimes, though, a third variable is needed to show a set of data more completely, for example, age, sex and race. A three-variable table is hard for the reader to interpret. No table should attempt to show more than three variables.

c) *Tables of Other Statistical Measures.*

- Although all the examples used so far show counts (frequency) of the variables, tables can also be used to show other statistical measures.
- The cells can contain rates, means, relative risks or other measures.
- Just be sure the titles and row/column headings clearly identify what data is being presented.

3. *Creating Class Intervals.*

- Some variables such as sex or “ate potato salad?” have a limited number of possible values.
- Others have a broader range of possible responses, and categories or “class intervals” are needed to group them.
- The following rules are important when creating class intervals:

- Create categories that are mutually exclusive and include all of the data. For example, if your first category is 0-5, the next one must start with 6, not 5.
- Use a relatively large number of narrow categories for the initial analysis, since you can always combine them later.
- Try to use standard groupings if they are available – for example, age categories used by CDC for a particular disease.
- Create a category for unknowns, since there will usually be missing information for some of the cases.

If there are no standard or natural class intervals for a particular variable, there are several ways to create class intervals. These are discussed in more detail in the CDC Principles of Epidemiology home study course.

B. Charts

1. In General

Charts:

- Are a method of organizing and illustrating data using only one coordinate.
- Are best used for comparing data with discrete categories.

Several types of charts may be produced using common spreadsheet software such as Excel.

2. Types of Charts

a) Bar Charts. Bar charts are used to create a visual display of the data from a table. The bars may be either horizontal or vertical.

- ***Simple bar charts***
 - Used to display the data from a frequency distribution (one-variable table).
 - Each bar represents one value of the variable.
 - This makes it very easy to compare the relative magnitude of the different values.
- ***Grouped bar charts***
 - Used to illustrate data from two- or three-variable tables.
 - The bars must be shaded or colored differently and described in a legend.
 - It is best not to use more than three bars per group.
 - Leave a space between adjacent groups of bars, but not between bars in a group.
- ***Stacked bar charts***
 - Another way of showing two variables.

- o The values of the second variable make up segments of the bars that represent the first variable.
- o These charts can be hard to interpret, since only the first segment rests on a flat baseline.
- **Deviation bar charts**
 - o Used to show how a variable deviates from a baseline, in both positive and negative directions.
 - o The bars are usually positioned horizontally.
 - o Each week, CDC's *Morbidity and Mortality Report* (MMWR) uses a deviation bar chart to show the number of cases of several diseases reported during the last four weeks, compared to the number reported during the same four weeks for the past five years.
- **A few simple rules for constructing bar charts:**
 - o Arrange the categories that define the bars in a natural order (for example, alphabetically or by increasing age), or in an order that produces increasing or decreasing bar lengths.
 - o Make all of the bars the same width, whatever looks good.
 - o Make the length of the bars in proportion to the frequency of the event.
 - o Code different variables by differences in bar color or shading, and include a legend that interprets your code.

b) Pie Charts.

- A pie chart is simple and easily understood.
 - o Very useful for showing the component parts of a single group or variable.
 - o The size of the pie "slices" shows the percentage for each component part of the whole.
 - o Pie charts are easily generated using spreadsheet software such as Excel.
- A few simple rules for constructing pie charts:
 - o Start at "12:00 o'clock" (straight up) and arrange the component slices from the largest to the smallest, going clockwise.
 - o Put the categories "other" and "unknown" last.
 - o Use different colors or shading for each "slice."
 - o Show somewhere on the graph what 100% of the pie represents (for example, the total number of cases).
 - o Indicate the percentage that each "slice" represents.
- Multiple pie charts are sometimes used to compare the same components in two different groups or variables. However, it is hard to accurately compare two or more pie charts visually.

c) Maps.

- Maps are a very widely used type of chart.
- They are also called geographic coordinate charts.
- Spot maps and area maps are commonly used in field epidemiology.

- **Spot maps** use dots or other symbols to show where an event took place, or where a disease condition exists.
 - Spot maps are good for detecting clusters of disease cases.
 - However, we must remember that a spot map does not take into account the size of the population at risk.
 - So it does not show the risk of the event occurring in that particular place.
 - A heavy clustering of dots may simply mean that more people live in that area and therefore more cases appear there.

- **Area maps**, however, can be used to illustrate differences in risk in different areas.
 - An area map uses shaded areas to show either the incidence of an event, or the distribution of some condition over a geographic area.
 - We can show either rates or numbers with an area map.
 - When we calculate and show a specific rate for each sub-area, we can make direct comparisons of risk between them.

- Mapping technology has grown very sophisticated in recent years. The widespread availability and use of Global Positioning Systems (GPS) devices and Geographic Information Systems (GIS) software has made it easier to produce accurate, up-to-date maps to aid in epidemiologic investigations.

C. GRAPHS

1. In General:

- A graph is a way to show numerical data visually, using a system of coordinates.
- A graph can help us see patterns, trends, aberrations, similarities, and differences in the data.
- People usually understand and remember the important aspects of data much more easily from looking at a graph than a table.
- Graph format:
 - Most graphs used in epidemiology have two lines, one horizontal and one vertical.
 - The horizontal line is called the x-axis
 - The vertical line is the y-axis.

- o The x-axis (horizontal) is used to show the values of the method of classification, for example, time in years, which is called the independent variable.
- o The y-axis (vertical) is use to show the dependent variable, usually a frequency measure such as number of cases or rate of disease.
- o Each axis must be clearly labeled and the scale of measurement marked.

2. Types of Graphs

Arithmetic-Scale Line Graphs

- Show patterns or trends over some variable, usually time.
- In epidemiology these graphs are often used to show the history of incidence of a disease over time.
- Arithmetic-scale line graphs are also good for comparing two or more sets of data.

Here are a few simple rules for constructing arithmetic-scale line graphs:

- Mark off each axis at equal intervals.
- Use a scale on the x-axis that matches the intervals used when collecting the data (for example, days, weeks, months or years). If very small intervals were used, combine them into larger ones.
- Make the y-axis shorter than the x-axis, so the graph appears horizontal.
- Always start the y-axis with 0.
- Pick a range of values for the y-axis that is slightly higher than the largest number you will be plotting.
- Select an interval size for the y-axis that will give you enough intervals to show the data in enough detail for your purposes.

A **histogram** is another type of graph that is very important in field epidemiology. We will learn about histograms later in this module.

IV. INTERPRETING EPIDEMIOLOGIC DATA: TIME, PLACE AND PERSON VARIABLES

As we look at tables, graphs and charts to draw inferences and form hypotheses, we often make comparisons between:

- Different **time** periods,
- **Places**, and
- Groups of **people**.

It is very important to use comparable data when making such comparisons. Be sure the data are comparable with respect to:

- Case definitions
- Level of effort in case-finding and data collection
- Time periods (compare weeks with weeks, months with months etc.)
- Populations (if two populations differ in age distribution or density, this should be taken into account)

Apparent differences in disease incidence can be influenced by any of these factors.

A. Time

1. In General

Variations over time in the frequency of a disease can tell us a lot about the determinants of that disease in a given population.

We may look at trends over many years (called secular trends), or seasonal variations, or variations over the short time period of an outbreak.

- As we look at disease data over time, we should ask these questions:
 - What is the pattern?
 - What factors might explain it?
 - What is the most likely future pattern?
- Variations over time may result from:
 - True increases or decreases--actual changes in the frequency of the disease in that population
 - Changes in sensitivity or specificity of the surveillance system
 - Mistakes made in collecting or organizing the data
 - Changes in the perceptions of the public or the health care community about the importance of diagnosing and reporting that particular disease
- A long-term increase in incidence of a disease may reflect changes such as:
 - Introduction of a new disease agent into a population
 - Decreasing effectiveness of control measures
 - Changes in the environment (for example, climatic conditions affecting the tick population)
 - Changes in societal practices (for example, urbanization leading to greater population density)

- Many diseases are subject to **cyclic changes** over time. Diseases that are strongly influenced by environmental factors may show seasonal variation.
- Another form of cyclic change is called **secular trends**. These are marked changes over long time periods that are caused by changes in environmental factors or host susceptibility.

Example of secular trend:

- Hepatitis A incidence has historically cycled up and down over periods of ten or more years.
 - When the disease is widespread, the number of susceptible people (especially children) goes down, so it begins to wane.
 - When enough susceptibles build up, it increases again.
 - This pattern may be changing now that an effective vaccine is available to prevent hepatitis A.
- Finally, time trends for a disease may simply show erratic change due to chance variations, sometimes called “noise.” This is true of many diseases with low endemic levels. It is also a common pattern when looking at very localized data, since the number of cases may be too low to exhibit a strong pattern.

2. Time during an outbreak or epidemic period

- When investigating an outbreak, monitoring time trends becomes critical.
 - An outbreak or epidemic period is the time during which the number of cases of a disease exceeds the expected number.
 - Time is usually measured in hours, days, weeks or months.
- A special kind of graph, called a **histogram**, is used to plot cases according to the time of onset of symptoms. This is called an **epidemic curve**. Interpretation of epidemic curves will be covered in more detail in the workshop portion of this course.
 - o A histogram looks somewhat like a bar graph, but with several important differences. In an epidemic curve:
 - The x-axis is always made up of equal time intervals, and should begin just before the outbreak
 - The time intervals should be appropriate to the disease in question (hours, days, weeks, months)
 - The y-axis is the number of cases
 - Each case is represented by one square, and all squares are of equal size
 - There are no spaces between the columns

- There may or may not be horizontal lines between the squares
 - You may show a second variable in a histogram through the use of shading.
 - For example, you may want to look at the distribution of hepatitis cases who are county residents vs. visitors to the county.
 - This can be done either by shading the squares representing visitors, or by constructing two separate histograms.
 - The most common spreadsheet software, such as Excel, does not generate proper histograms. They may be constructed by hand, or using specialized software such as Epi Info, which was produced by CDC.
- A **frequency polygon** may be used instead of a histogram to show an epidemic curve. In this type of graph, the squares are replaced by a line. However, a frequency polygon is not the same as a line graph, as shown in the next example.

B. Place

- Place is a specific geographic area that can be described by latitude, longitude and altitude. As used in epidemiology, place:
 - May be a street address, city, state, region, or country, or
 - May be expressed as a dichotomous, “either-or” variable such as
 - urban/rural
 - domestic/foreign
 - institutional/non-institutional
 - lower vs. higher socioeconomic areas
- The association of a disease with a place implies that the most important causative factors are in the environment or people of that place. For example:
 - Population density (urban vs. rural) can affect how rapidly an airborne disease is transmitted, or determine risk of exposure to a vectorborne disease.
 - The incidence of many diseases increases as socioeconomic status decreases, due to the effects on immune status, quality of the environment, overcrowding, etc.
 - Regional variations can reflect the specific ecologic requirements of a disease agent or vector

- Country of origin or of exposure can be important, because diseases may be imported from endemic areas (for example, malaria)
- Rates, not counts, must be used to compare disease incidence in different places. Otherwise the difference in population size would make it impossible to interpret any differences.
- Place comparisons are most useful if we look at the data over time. Remember, no one point in time can give us all the information we need.

C. Person

As we learned in Module I, people can be described in terms of many inherited or acquired characteristics such as:

- Age
- Sex
- Race
- Immune status
- Marital status
- Educational level

They may also be described in terms of their activities, such as:

- Occupation
- Recreational activities
- Religious practices
- Customs

Or, they may be described by the circumstances in which they live, such as:

- Social conditions, for example housing
- Economic status
- Environmental conditions

These variables are important since they determine, to a large degree, who is at the greatest risk of acquiring specific infections.

Age is the single most important personal characteristic. To a large extent, it determines:

- The physiologic activity of the disease organism
- The level of immunity or resistance, and
- The potential for exposure to a disease agent

Our behavior, and therefore our risk of exposure, differs markedly at different life stages. Examples are the mouthing behavior of toddlers and increased sexual activity during adolescence and young adulthood.

Sex can also influence the risk of disease.

For many diseases, both sexes have about the same level of risk. However, if there is a gender difference in a particular disease it usually means either males or females had a greater opportunity for exposure. This could be due to differences in:

- occupation (for example, child care, agricultural work)
- recreational activities (for example, hunting), or
- social behaviors (for example, intravenous drug use)

Race and/or ethnicity can also be a factor in disease risk.

- Such disparities most often result from differences in exposure or immunity status (for example, immunization rates).
- With most diseases, racial and ethnic differences are a consequence of socio-economic and cultural differences, rather than physical differences.

Summary

Epidemiologic variables are characteristics that can be observed and/or measured. They may be characteristics of

- **time,**
- **place,** or
- **person.**

Tables, charts and graphs are good tools for organizing epidemiologic data. They make it possible to identify, explore, understand and present data distributions, trends and relationships.

After we organize the data, we can look at epidemiologic variables to:

- identify characteristics that might be important, and
- form hypotheses about the source, causative agent, and mode of transmission of illness.

PRINCIPLES OF INFECTIOUS DISEASE EPIDEMIOLOGY

MODULE VI – INTERVIEWING 101

I. INTRODUCTION

- Interviewing people to elicit information is a key skill for any field epidemiologist.
- When investigating a disease case or outbreak, we must help people “open up” and give us accurate information.
- Success in disease control often depends on the ability to develop trust and rapport, and to aid people in recalling what has happened.

Because interviewing is a skill that requires practice, this module will present only the “basics.” You will have the opportunity to practice during the Workshop portion of this course.

Objectives: Module VI is designed to prepare public health workers to meet the following objectives:

- Define the purpose and goals of epidemiologic interviewing
- Describe methods used to assure confidentiality
- Identify the three components of effective communication
- Describe several techniques for maintaining objectivity and eliciting accurate information

II. WHO DOES AN EPIDEMIOLOGIST INTERVIEW?

Field epidemiologists may interview a wide variety of people in the course of their work, such as:

- Health professionals reporting a disease case or outbreak
- People diagnosed with, or suspected to have, an infectious disease
- People who may have come into contact with someone with an infectious disease
- People who may have been exposed to a disease source (for example, a food establishment, specific food product, or water supply)
- Workers involved in a suspected disease outbreak, such as food service, health care or childcare workers

Interviews may be done in person or by telephone and may take place in all kinds of settings, such as:

- homes
- hospitals
- workplaces
- schools or childcare centers, or even
- “on the street” - anywhere a disease investigation takes us.

Example:

A hospital laboratory reports a case of *Salmonella* in a young woman who is hospitalized.

Interview #1:

The field epidemiologist calls the hospital's records department to get information to complete the initial CD-1 report.

Interview #2:

- The field epidemiologist:
 - visits the hospital and interviews the patient, using a standard enteric disease investigation form.
 - finds out, through this interview, that the patient had lunch with a friend two days before she became ill.
 - finds out that the friend has been mildly ill with nausea, diarrhea, and fever.

Interview #3:

- The field epidemiologist:
 - calls the friend and completes an enteric disease investigation interview.
 - finds out that the friend works in a childcare center, and has continued to work throughout her illness.
 - arranges for the friend to be tested for salmonellosis.

Interviews #4, #5 and so on:

From here, the investigation may take two different directions, each of which will require more interviews.

- First, the field epidemiologist will go to the restaurant where the two women had lunch and interview the manager (and perhaps staff) to find out whether any of the staff are ill, whether complaints of other illnesses have been received, etc.
- Second, if the friend's lab test is positive for *Salmonella*, interviews will need to be conducted at the childcare center where the friend works.

III. WHY CONDUCT INTERVIEWS?

The basic purpose of an epidemiologic interview is to get information that can help prevent the spread of disease. Interviews may:

- Obtain complete data for disease reporting and analysis
- Provide "clues" that lead to hypothesis development
- Identify the source and/or connections between disease cases or outbreaks
- Help prevent the development of disease in those potentially exposed
- Help prevent the complications of untreated disease in those already infected

The goals of an interview vary with the type of interview. Some common goals are:

- To get honest, complete and accurate information

- To educate the person being interviewed about the disease and its treatment
- To motivate the person to assist in identifying additional people at risk and preventing the spread of infection
- To reduce the risk of the person spreading the disease

IV. TRUST AND CONFIDENTIALITY

Trust

To gain the trust of the interview subject, **rappport** must be established by

- discussing common interests.
- showing the person that this is not just a job, and that you truly care about their health and the health of those around them.
- explaining the goal of the interview and how it will benefit the person and others.

Confidentiality

- Confidentiality is a very important principle in public health work.
- Interviewing is a voluntary process, which requires the acceptance and cooperation of the person being interviewed.
- If the person has concerns about whether the information will be kept confidential, they may be reluctant to provide information.
- The preservation of confidentiality is established by health department policy and practice, and by state and federal statutes. The interviewer should be familiar with the policies and procedures of his/her agency.
- Missouri law (Revised Statutes of MO 192.067) allows the Department of Health and Senior Services to obtain information from medical records, for purposes of conducting epidemiological studies to be used in promoting and safeguarding the health of the citizens of Missouri. This authority extends to the local public health agencies through departmental rules and contracts. See Module III, Public Health Surveillance, for more information about public health's legal authority to collect information.
- In most epidemiologic investigations, the person giving information must be reassured that they will not be identified publicly. There may be important exceptions, however. If bioterrorism or other intentional exposure is suspected, a criminal investigation may be necessary. In that case confidentiality could not be guaranteed.
- What does confidentiality mean to the person being interviewed?
 - Contact names must only be used for field investigation and notification.

- A contact must never be told the name or identity of the person who named them as a contact (or given other information that could lead to identifying the source, such as specific times, places, etc.).
- Information must be shared only with other health professionals who have a need to know in order to perform the investigation.
- Methods for maintaining confidentiality:
 - Avoid revealing any identifying information about other cases or contacts—not only name, but also age, race, gender, location of index patient, etc.
 - When using the telephone, ensure that you are talking with the correct person. Verify that he/she is in a private setting.
 - Follow special procedures when interviewing persons or other sexually transmitted diseases (more information on this will be shared in the Workshop).
- If an interview is not about personal medical information, then confidentiality safeguards may not apply. For example, information collected from a foodhandler about the process used in food preparation is not considered medical information, and may be an open record after an outbreak investigation is completed.

V. COMMUNICATION: SOME BASIC COMPONENTS

Effective interviewing requires good communication skills. There are three components of effective communication:

1. Non-verbal communication
2. Verbal communication
3. Effective listening

1. Non-verbal communication is extremely important - some say it is more important than the actual words we use. Here are some tips for good non-verbal communication:

- Eye contact: Good eye contact conveys confidence in yourself, your message, or both. Too much eye contact can be seen as aggressive.
- Facial expressions: Be aware that any facial expressions can affect the interaction. It is best to approach the person in a non-threatening manner.
- Body orientation: Fully face the person while maintaining appropriate eye contact. The optimum distance from the other person is between 18 inches and four feet.
- Posture: Body positioning can have a great impact on communication. Lean forward and listen intently to show interest. Don't appear too relaxed (by pulling back or putting your feet up) - this tells the person you don't care.
- Physical environment: Ideally, the interview should be held in a private location with few distractions.

When interviews are done by phone, of course, non-verbal communication is less important. However, your tone of voice and the pace of your speech can convey some of the same messages that non-verbal cues such as posture convey in person.

2. Verbal communication refers to the way we organize and present the words we use. Some tips for good verbal communication include:

- **Brevity:** Be brief and to the point.
- **Primacy:** Say the most important things first, to help the person remember them.
- **Organization:** Ask your questions and present your messages in a logical and sequential manner. It is always best to use a prepared format for an interview.
- **Appropriate educational level:** Remember that many people cannot understand technical terms or complicated sentences. The average reading level among the public is 4th to 6th grade. Use familiar terms and avoid “talking down.”
- **Clarification:** Always give the person the opportunity to ask clarifying questions. If they don’t understand what you are asking, they cannot give accurate information.
- **Repetition:** Repeat important topics to help the person understand and remember the message. You may also ask the person to repeat the information.
- **Specificity:** Be explicit. Do not raise irrelevant points or “beat around the bush,” or the message will be lost.

3. Effective Listening is a vital part of communication. Effective listening includes non-verbal and verbal feedback to the person who is talking.

- Non-verbal feedback conveys encouragement, and may include nodding, smiling, and other appropriate gestures.
- Verbal feedback means paraphrasing what the person said, or responding to the content. This conveys you have heard and understood what was said.
- Avoid selective listening. Listen to the entire message being communicated, not just what you want to hear. You may miss something very important!

VI. OBJECTIVITY

There is one other key ingredient of a good interview. The interviewer must remain objective and try to elicit accurate information. To do this:

- Don’t anticipate the answers - let the person speak for him- or herself.
- Go back over any responses that seem inconsistent. For example, when reconstructing an exposure history, if there are gaps, contradictions or the timing seems “off,” gently guide the person through the sequence of events again. It can be hard to remember things that happened days or weeks ago!
- Never “lead” a person to a particular answer, even if you have a strong suspicion what the answer “should be.” You will only harm the investigation by influencing the responses.

- If you sense that the person is hesitant to share some information, concentrate on making them comfortable. You may need to “back off” and come back to that subject again later.
- Take some time at the end of the interview to go over your notes and ask any needed follow-up questions.

Interviewing is an important but challenging part of field epidemiology. In the Workshop portion of this course, you will have the opportunity to practice with other students.

Remember, a good interviewer:

- is well organized.
- establishes rapport with the person being interviewed.
- assures confidentiality (when appropriate).
- uses good verbal and non-verbal communication skills.
- always remains objective.

MODULE VII

Outbreak Investigations

A. Overview

- Outbreak investigations should be a collaborative effort, since several tasks requiring different skills must be done at the same time.
- **Steps in Disease Investigation**

While every outbreak is unique, the investigative process generally follows the sequence outlined below:

 1. Obtain the initial report.
 2. Determine the extent of illness. Are there other associated cases?
 3. Learn about the suspected agent—is it transmissible from person to person? Is it transmissible through the environment, such as by food or water?
 4. Plan the investigation.
 5. Conduct the investigation.
 6. Formulate a case definition for analytical purposes.
 7. Analyze the cases and characterize by time, place, and person.
 8. Evaluate the hypothesis and formulate conclusions.
 9. Select, implement, and evaluate control measures.
 10. Prepare investigation report.
 11. Distribute the approved final report to all agencies that contributed to the investigation effort.
 12. Conduct after-action evaluation.
- This list is a summary of the things that need to be considered in any investigation. In real life, several of these steps may go on at the same time. Their order will vary, and several of the steps may occur more than once. However, all of these things are necessary to the successful resolution of an outbreak.

B. Preparations for an outbreak must begin before the outbreak occurs.

1. Each agency should establish a multidisciplinary investigative team and assign responsibilities. Members should include:
 - nursing
 - communicable disease
 - environmental
 - support staff
 - laboratory
 - public information, and
 - computer information

2. Staff should receive training, including Introduction to Epidemiology, Principles of Epidemiology and other disease specific courses on investigative procedures.
3. Assemble materials: laboratory kits, forms, reference materials, personal protective equipment such as gloves and masks.
4. Maintain a current phone directory, including e-mail and Internet addresses, home addresses and phone numbers of team participants, and key contact personnel outside the Local Public Health Agency.
5. Maintain adequate local surveillance systems for the early detection of increased disease incidence.

C. Definitions

An *outbreak or epidemic* is the occurrence in a community or region of an illness(es) of similar nature, clearly in excess of normal expectancy, and derived from a common or a propagated source (19 CSR 20-20.010).

Acute gastroenteritis is an illness with sudden onset characterized by symptoms such as diarrhea, vomiting, fever, or abdominal cramping.

NOTE: Always consider the possibility of intentional contamination when investigating an outbreak. If a bioterrorism event is suspected, notify your Regional Communicable Disease Coordinator and appropriate law enforcement officials immediately.

AN OUTBREAK CASE STUDY: Foodborne Outbreak Associated with a Wedding, Southwest Missouri, June 2002

Introduction

In this case study you will investigate a disease outbreak following the **Steps in Disease Investigation** above. For purposes of this exercise, you will be the head of the investigation team for the Washaw County Health Department (WCHD), and you will work on each of the steps in order. For each step, you will be given some information describing the current situation and then asked to describe how you would handle that step. The correct answers will be provided after each step.

STEP 1. Obtain the initial report.

Situation: On June 7, 2002, the WCHD nurse received a call from a person who had attended a wedding and reception on June 1, 2002 and soon afterward became ill with acute gastroenteritis. The caller said he knew of several other people who were also ill.

Question: What additional information should you get from the caller?

STEP 2. Determine the extent of the illness. Are there other associated cases?

Situation: You have contacted the local hospital. The Laboratory Director said they had recently sent five *Salmonella* isolates from stool cultures to the State Public Health Laboratory (SPHL) for additional testing. Within the next three days, the SPHL reported that the five isolates were all *Salmonella infantis*. One of these patients was the caller who attended the wedding.

You have telephoned the Southwest Regional Communicable Disease Coordinator, who reviewed the surveillance data in the MOHSIS computer system. No other cases of *Salmonella infantis* had been reported in the Southwest Region in all of 2002.

a. Question: Should this episode be considered an outbreak? If so, what other activities should you start?

b. Question: What definition of an outbreak related case would best serve the investigation at this point?

STEP 3. Learn about the suspected agent—is it transmissible from person to person? Through the environment (including food or water)?

Situation: Consult the Communicable Disease Investigation Reference Manual (CDIRM) or other current references such as *Control of Communicable Diseases Manual* or the *American Academy of Pediatrics Red Book* to answer the questions in Steps 3 and 4 and get other information about the natural history of the disease. In this case, the agent is known to be *Salmonella infantis*, which is transmissible from person to person and through the environment. (See CDIRM manual section on salmonellosis at <http://www.dhss.mo.gov/CDManual/Salm.pdf>)

a. Question: Considering salmonellosis is transmissible from person to person, what additional steps should be taken?

b. Question: Considering that salmonellosis is transmissible through the environment, what additional steps should be taken?

STEP 4. Plan the investigation.

Situation: By carrying out the activities in the previous steps, the following information was obtained:

- There were three meals associated with the wedding celebration:

May 31, 2002	Rehearsal Dinner
June 01, 2002	Bridal Brunch (morning)
June 01, 2002	Wedding Reception (7:00 p.m.)
- The bridal brunch was held at a private residence, and was attended by 12 women.
- The rehearsal dinner and the wedding reception were catered by a local catering firm and held at a convention center. The catering firm is operated out of a home with a separate kitchen devoted to the business.
- About 30 people attended the rehearsal dinner and 300 attended the wedding reception.
- All of the first five identified cases attended the reception, but only one was at the rehearsal dinner and none attended the bridal brunch.

a. Question: At this early stage, what would be a reasonable tentative hypothesis about what may have caused the outbreak?

Clues:

- Formulate a tentative hypothesis based on the time, place, and person associations you have found so far. This hypothesis will form a basis for the investigation. It is very

important not to be too narrow in your focus, thereby excluding potentially important cases or events.

- Develop the hypothesis by interpreting available data to determine:
 - Identity of most likely agent(s)
 - Most likely source(s) of agent
 - Most likely mode by which agent was transmitted.

b. Question: The purpose of the detailed investigation is to gather the information needed to test your tentative hypothesis. What are the required tests of each component of the hypothesis, and what information is needed for each?

c. Question: You have developed your hypothesis and determined what information you will need to gather. How will you gather the information and test your hypothesis?

STEP 5. Conduct the investigation

Situation: Your team is assembled and the questionnaires are developed.

Question: How will you proceed with the investigation?

STEP 6. Formulate a case definition for analytical purposes.

Situation: You've gathered a lot of information. Now you need to develop a more refined case definition so you can clearly identify the relevant cases and analyze the data. The goal is to create a case definition that is sufficiently "tight" to include only the people whom you are reasonably sure had *Salmonella* infections related to the outbreak.

Question: What elements will you use to formulate your case definition? What is your new, refined case definition?

STEP 7. Analyze the cases and characterize by time, place, and person.

Situation: The investigation was carried out as planned. With the assistance of the Regional Communicable Disease Coordinator, the following steps were taken:

- The case definition was used to identify which people were considered to be "cases."
- An epidemiologic curve (histogram) was created.
- A case-control study was conducted and attack rates were calculated.

Following are the major findings, grouped again by the three major parts of the hypothesis.

1. "This is an outbreak of *Salmonella infantis*..."

- Ten stool specimens from outbreak-related cases were confirmed positive for *Salmonella infantis*. The earliest specimen was collected on 6/4/02 and the last one on 7/2/02. No other enteric pathogens were isolated.
- One person with a stool specimen positive for *Salmonella infantis* did not meet the case definition and was excluded from the analysis. She was the mother of the bride (who did not handle the food). Her symptoms started on 5/31/02 (the day before the wedding) and therefore she did not meet the definition. It is possible that she had

“nervous diarrhea” before the wedding, and her *Salmonella* infection actually began later.

- Sadly, the bride and groom were also reported to be ill. They were not included in the study because they had left on their honeymoon anyway.
- The common signs and symptoms were diarrhea (100%), cramps (84%), nausea (52%), fever (52%), chills (44%), and headache (36%).
- There were 24 cases who reported onset time. The mean onset time was 23 hours after the wedding reception began; the range was 6 to 63 hours.

2. “...caused by the ingestion of contaminated foods (or beverages)...”

- Exposure histories of ill and well attendees:
 - The case-control study began on June 12, 2002. Fifty-one persons were interviewed, all of whom ate at the reception. Twenty-five of the 51 were ill and met the case definition. Three more ill people were interviewed but did not meet the case definition and so were excluded from the analysis. The controls were 23 well guests identified during interviews with wedding attendees.
 - All cases interviewed were over 18 years old. The mean age of the cases was 36 and the mean age of the controls was 29.
 - The menu consisted of: turkey, ham, roast beef, potato salad, pasta salad, raw vegetables and dip, raw fruit and sauce, chips, bread, condiments, a variety of cakes, iced tea, soda and beer.
 - Two-by-two tables were constructed for each menu item served at the wedding reception. Two foods were found to be statistically significantly associated with illness:

Food Item	Odds Ratio	95% Confidence Interval	Uncorrected “p” Value
Turkey	5.45	1.19 – 26.99	0.01
Potato Salad	5.2	1.22 – 23.52	0.01

- Environmental evaluation
 - No food remained for testing.
 - On June 10, 2002, the convention center in which the reception was held was inspected and revealed the following:
 - The tables used for serving food had no cold holding capacity.
 - Serving tables were not provided with sneeze shields.
 - The kitchen area was adequately equipped and clean.
 - The inspection and evaluation of the caterer’s facility on June 10 revealed the following:
 - Foodhandling equipment appeared to be in good working order.
 - The operation did not have a three-vat sink for proper dishwashing.
 - All foods served at the wedding reception were to be served cold.
 - The caterer received uncooked boneless turkey breasts at approximately 10:00 p.m., Wednesday, May 29. They were delivered frozen by the bride’s family from Smallville, individually vacuum packaged. The caterer immediately placed the breasts in a tub of water. The caterer could not remember if the breasts were placed in refrigeration or left on the counter at room temperature to thaw. The thawed breasts were cooked in the original vacuum packaging Thursday afternoon, May 30,

to a temperature of 170°F using a meat thermometer to check cooking temperature. They were removed from the electric roaster oven and cooled at room temperature for 1½ to 2 hours. The breasts were then placed in refrigeration. They were sliced at the caterer's on a commercial meat slicer Friday afternoon, May 31. Old food debris was found on the slicer on June 10, the day of inspection.

- The potato salad was prepared on Thursday, May 30, at the caterer's with the following ingredients: potatoes, Miracle Whip salad dressing, mustard, commercially prepared pickle relish, celery, sugar, salt and pepper.
- The pasta salad was prepared at the caterer's on Thursday, May 30, with the following ingredients: commercially prepackaged noodles, oil, vinegar and mustard.
- Pre-cooked boneless hams were served, which were shaved and packaged at two large grocery stores in a nearby town. Unannounced visits to both stores on June 11, 2002 revealed that the hams were sliced in the meat cutting departments. Only a single meat slicer was present in each meat department. During the day, raw and cooked products were being sliced. The meat slicers were not thoroughly cleaned and sanitized between the slicing of raw and cooked products. Raw beef particles were present on both slicers at the time of inspection. The ambient air temperatures in the meat cutting rooms were in the mid to upper 70s.
- Pre-cooked Hormel brand roast beef was sliced and packaged at a grocery in a small town in the southern part of the county. The roast beef was picked up the morning of the dinner and delivered in coolers to the convention center where the dinner was served. A visit to the grocery on June 11, 2002, revealed the following: The meat slicer was used only for precooked prepackaged deli meats. The meat slicer was clean. The walk-in meat cooler used for storage was 40°F and also clean.
- On June 10, 2002, a sample of the water supply at the catering establishment was obtained and analyzed based on Department of Health and Senior Services standards for drinking water. The water was determined to be unsatisfactory, with bacteria too numerous to count with coliforms. It was also noted that the well that supplies the water was located within 50 feet of a hog lot.

3. "... served at the wedding reception on June 1."

- Thorough questioning of the ill and well persons included in the case-control study revealed no other activities or food or beverage sources in common in the week preceding the outbreak.

a. Question: Was the planned investigation adequately conducted?

b. Question: Referring to the epidemic curve (the histogram you previously downloaded), is the distribution of the cases compatible with a common exposure at the wedding reception? Describe how you arrived at your conclusion.

c. Question: Will the information obtained allow an adequate test of each element of the hypothesis?

STEP 8. Evaluate the hypothesis and formulate conclusions.

Situation: You now have all available information from the statistical analysis, along with laboratory data, environmental inspection findings, and other relevant information with which to evaluate the hypothesis and formulate conclusions.

a. Question: Is the first part of the hypothesis (“This is an outbreak of *Salmonella infantis*. . .”) supported well enough by the data that you accept it as true?

b. Question: Is the second part of the hypothesis (“ . . . caused by the ingestion of contaminated foods (or beverages)...”) supported well enough by the data that you accept it as true?

c. Question: Is the third part of the hypothesis (“ . . . served at the wedding reception on June 1”) supported well enough by the data that you accept it as true?

STEP 9. Select, implement, and evaluate control measures.

Situation: The outbreak was caused by an organism, *Salmonella infantis*, which causes gastrointestinal symptoms.

a. Question: What control measures have you selected and implemented?

b. Question: How can you determine whether the control measures were effective?

STEP 10. Prepare the report

You will learn how to prepare an outbreak report in Module VIII and you will be able to view a report for this outbreak of *Salmonella infantis*.

STEP 11. Distribute the report

The outbreak report should be submitted to the WCHD Administrator and to DHSS for approval. Once approved, it should be distributed according to agency guidelines. At a minimum, it should be shared with each agency involved in the investigation, and with any other entities who have made a formal request for it.

STEP 12. Conduct after-action evaluation

Every outbreak investigation provides an opportunity for learning. The team(s) involved in the investigation should be pulled together and a discussion held about what went well, what did not go so well, and what changes can be put in place to make it easier and better the next time.

PRINCIPLES OF INFECTIOUS DISEASE EPIDEMIOLOGY

MODULE VIII – PREPARING AN OUTBREAK INVESTIGATION REPORT

Note: You will need to access the course to view the examples associated with this module, as they are not included in this outline.

INTRODUCTION

Module VIII is designed to prepare public health workers to meet the following objectives:

1. Correctly identify the necessary elements of an outbreak investigation report
2. Correctly sequence epidemiologic information within an outbreak report
3. Identify the appropriate recipients of an outbreak report

Reports about an outbreak investigation may take many forms. During an investigation, **interim reports** help to transmit information about:

- what has happened
- what is happening
- what progress is being made

Preparing an interim report can help clarify the investigator's understanding and provide new insights.

Appropriate interim reports during the investigation may include:

- Verbal reports to the outbreak coordination team
- Verbal and written progress reports to administrators
- Information briefings for the media/public

This module will focus on the preparation of a formal, **written report** after the conclusion of the outbreak investigation. Preparing and distributing a report assures that the experience gained and the

discoveries made during the investigation are not lost. They can be used to design and implement improvements in the surveillance system and prevention/control measures. Ultimately, this knowledge can help reduce the risk of similar situations occurring in the future.

A final report should be prepared within 90 days after the outbreak investigation, while the information is still fresh and the findings will have the most impact.

Let's look at each of the report components, in order.

I. TITLE

The title should contain, at a minimum, the type of outbreak, location, and date.

II. SUMMARY

The Summary section should contain all of the key facts that describe what happened. It should be brief and concise. The information can be explained and elaborated in other sections of the report.

The Summary should include the following information:

- Date and place of outbreak
- Number exposed
- Number interviewed
- Numbers of suspect, probable and confirmed cases
- Number hospitalized
- Number of deaths
- Key statistics (see below)
- Causative organism
- Control measures
- Recommendations

Key statistics about the outbreak include:

- Attack rate (if available)
- Hospitalization rate
- Death rate
- Frequency distribution of symptoms
- Median date of exposure
- Median date of onset
- Average incubation period
- Average duration of illness
- Average duration of hospitalization

A reader should be able to read through the summary to gain a basic understanding of how many people were ill, how badly they were affected, what agent caused the problem, how the outbreak was controlled, and any recommendations for preventing future outbreaks.

III. INTRODUCTION

The Introduction can be brief, and should set the scene for the investigation. It should include:

- Date of initial report
- Agency that received the initial report
- Place and date of the outbreak
- Name and official title of the person submitting the report

IV. BACKGROUND

The Background section should provide relevant information about the community and population in which the outbreak occurred, such as:

- A map of the community (or, if the outbreak occurred in an institution, a map of the facility)
- A description of the relevant demographics (size and composition of the population affected, any recent demographic changes, etc.)

V. METHODS

The Methods section should answer the reader's questions about what was done, how and by whom. It should include:

- **What population** was considered to be at risk?
- **What** and **how much** data was collected?
- **From whom** and **from how many people** were data collected?
- **By whom** were data collected?
- **How were case definition(s)** developed and used?
- **How was the well comparison group selected, and how many** people were in it?
- **How were data collected and analyzed?**
 - Records reviewed
 - People interviewed
 - Questionnaires developed and distributed
 - Questionnaire reliability and validity
- **How were laboratory specimens** collected and analyzed?
- **What laboratory standards** were used?
- **What hypotheses** were developed (including tentative ones)?
- **Where, by whom and how were environmental inspections done?** (including the standards used for the inspections, for example, the 2000 city ordinance or 1999 state food code)

The Methods section is usually one of the longer sections of a report.

VI. RESULTS

The Results section should present all of the results from all of the methods used, including laboratory testing, interviews and environmental inspections. The information included in the Summary section can be presented and explained in more detail here (except for the control measures and recommendations).

This section should also include:

- the epidemic curve (histogram) showing illness onset dates

- a summary of the exposure histories of the persons interviewed (for example, their food histories)
- results of statistical probability testing
- test results from the environmental samples
- test results from the human specimens

The Results section is also pretty lengthy.

VII. ANALYSIS

This is the place to present what you have learned from the investigation. It should show your conclusions and interpretations regarding the:

- source of infection
- agent
- reservoir
- mode of transmission
- the group at highest risk

VIII. CONTROL MEASURES

This section should answer the reader's questions about the measures taken to control the outbreak:

- **What methods** were used for outbreak control?
- **How** were they implemented?
- **Where, when and by whom** were they implemented?
- **How was their effectiveness measured?**
- **How effective were they?**

IX. RECOMMENDATIONS

What recommendations can be made, based on what has been learned? This is the place to suggest changes in policies, procedures, and/or educational efforts in order to:

- prevent future outbreaks
- improve surveillance and detection of outbreaks
- improve the process of outbreak investigation and control

X. OTHER OUTCOMES

This is the place to describe what the outbreak, and the efforts to control it, have done to the population at risk. First, what impact did the outbreak itself have, including both health and economic consequences?

- Did any individuals have serious complications that will cause long-term health problems? Was the health care system adversely impacted by the outbreak, for example by a surge in hospital admissions?
- Were businesses or institutions affected economically (for example, by adverse publicity)?

Just as important is the impact of the control measures on the:

- Population—was their way of life affected? Did their immune status change (because of an immunization effort, for example)?
- Reservoirs—if the reservoir was animal or environmental, how did the control measures change it? Was there a change in the abundance or distribution of the disease agent (because of sanitizing, spraying, trapping or other interventions, for example)?
- Vectors—if there was an animal vector, how was it impacted by the control measures? Are there now fewer vector animals? Are they distributed differently in the environment?

Finally, this is the place to share your other discoveries. Did you learn something new to science in the course of the investigation? Often, outbreaks yield new knowledge that needs to be shared with other public health workers. Examples: new agent, reservoir, vector, temperature range, novel mode of transmission, unusual symptoms or complications, etc.

Outbreaks in MO have contributed to new discoveries, such as:

- Transmission of *E. coli* O157:H7 via drinking water contaminated with sewage (first waterborne outbreak of this agent)
- Discovery of erlichiosis as a new tickborne disease in humans

- Transmission of *Salmonella* via public drinking water. The water was cross-contaminated from an abandoned industrial water tower with birds roosting in it. This led to a new water tower inspection program in the Department of Natural Resources.

XI. DISTRIBUTING THE REPORT

At a minimum, the final report of an outbreak investigation should go to the:

- Investigation team members
- Administrator of the investigating health agency
- DHSS (Regional Communicable Disease Coordinator and state office)

Others may request a copy of the report (for example, an affected business, the press, and/or complainants in a lawsuit related to the outbreak). Final outbreak reports are subject to the state's Open Records (Sunshine) law, so they may be released in some circumstances. The field epidemiologist should refer any such requests to his/her supervisor or administrator.

XII. OTHER REPORTING REQUIREMENTS

Depending upon the type of outbreak, there may be other requirements for filing specific forms in addition to the narrative report. Please consult the Communicable Disease Investigation Reference Manual (CDIRM) for details. <http://www.dhss.mo.gov/CDManual/CDsec30.pdf>