An Interactive Introduction to Organismal and Molecular Biology

AN INTERACTIVE INTRODUCTION TO ORGANISMAL AND MOLECULAR BIOLOGY

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CONTENTS

Unit II. Organismal Biology

INTRODUCTION

Andrea Bierema

This "textbook" is interactive, meaning that although each chapter has text, they also have interactive HTML5 content, such as quizzes, simulations, interactive videos, and images with clickable hotspots. Students receive instant feedback when they complete the interactive content, and therefore, can learn and check their understanding all in one place. I still consider this textbook to be fairly text-heavy and will continue to make it even more interactive content!

The image on the cover and above represents the creation of this book. I pulled most of the content from open resources, modified them, added questions, and now offer them for you to use!

I chose the content to align with two courses that I teach: environmental and organismal applications and biomedical applications. Unit 1 introduces students to science, which both courses use. Unit 2 covers content necessary for understanding conservation implications (the underlying theme of the course is de-extinction), and Unit 3 focuses on proteins so that students can understand the implications of modifying DNA (the underlying theme is CRISPR).

Please use this book as you see fit for your classes. I look forward to hearing how to make this book even more useful in the future!

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UNIT I

INTRODUCTION TO SCIENCE

1.

NATURE OF SCIENCE

Nature of Science

Andrea Bierema

Learning Objectives

- Identify aspects and misconceptions regarding the nature of science and scientific inquiry.
- Describe the processes of science.
- Given a description of an investigation, determine the type of study it is, the research question, and control and experimental variables, when appropriate.

An Introduction to the Nature of Science

To understand what ***science*** is, just look around you. What do you see? Perhaps your hand on the mouse, a computer screen, papers, ballpoint pens, the family cat, the sun shining through the window, etc. Science is, in one sense, our knowledge of all that: all the stuff that is in the universe from the tiniest subatomic particles in a single atom of the metal in your computer's circuits, to the nuclear reactions that formed the immense ball of gas that is our sun, to the complex chemical interactions and electrical fluctuations within your own body that allow you to read and understand these words. But just as importantly, science is also a reliable process by which we learn about all that stuff in the universe. However, science is different from many other ways of learning because of the way it is done. Science relies on ***testing*** ideas with ***evidence*** gathered from the ***natural world***.

Given the way that science is often taught—memorizing facts from a thick textbook based on research done decades ago and completing lab activities in which there is one known answer—many students have ***misconceptions*** about what science is and how it works. Complete the following interactive to learn more about the real side of science!

Exercise

Before beginning the interactive element below, try this!

- Please see this interactive tool that contains a list of statements regarding the nature of science; some correct and some not. Click on a statement and hold down to move the statements around.
- The first thing to do on the website is to put the statements into three groups:
    - Agree: Statements that you agree with
    - Disagree: Statements that you disagree with
    - In between: Statements that you believe to be true under some conditions, but not others.
- Second, once the statements are put into three groups, order the statements from those that you most agree to those that you least agree with. Discuss your list with your peers.

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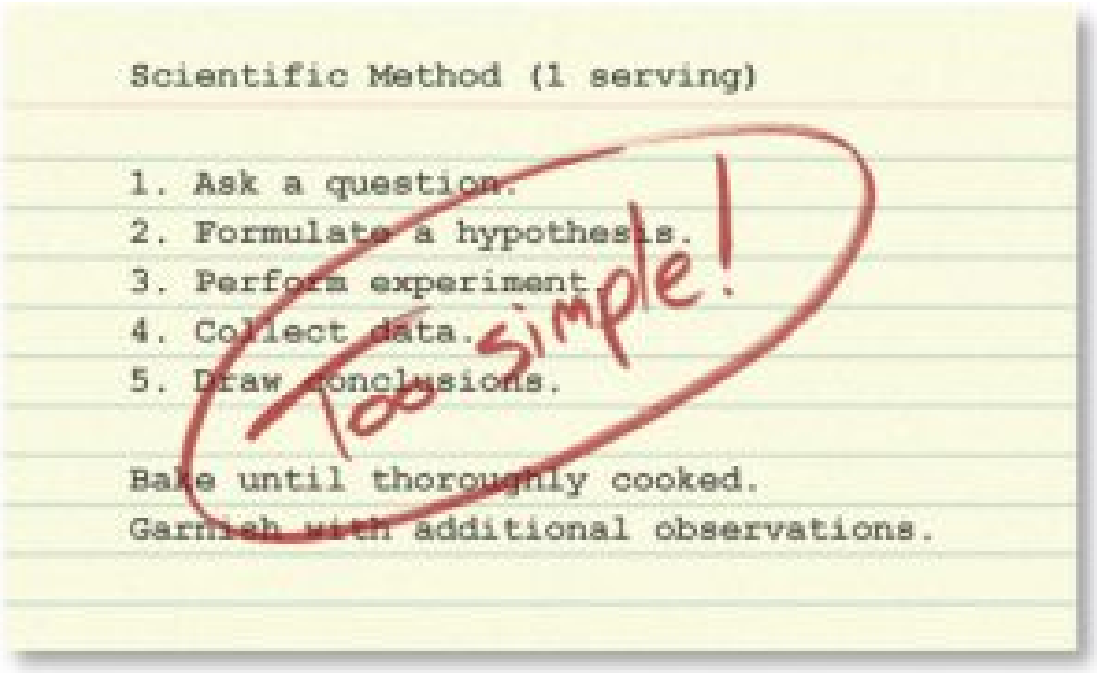
Scientific Practices

The scientific method is too simple to capture the various ways that science occurs, and therefore, we refer to the way science occurs as scientific practices. This section provides more detail on scientific practices, including how it can be explained with a flow chart, that scientific investigations are based on testable research questions and include observations and inferences, and how scientific ideas are tested.



Science Flowchart

As you learned from the activity above, scientists do not follow one Scientific Method. Rather, science is complex. Notice in the flowchart, for instance, that it is non-linear; every study is unpredictable and follows a different path. Moreover, the research is not "done" after one investigation. Results often lead to new questions or new ways to investigate a similar question. Also notice in the flowchart that one of the main elements is "Community Analysis and Feedback." Science is a social endeavor and scientists talk to each other about their research before, during, and after an investigation is done and even published.

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Exercise

At first, the science process, as demonstrated in the science flowchart, might seem overwhelming. Even within the scope of a single investigation, science may involve many different people engaged in all sorts of different activities in different orders and at different points in time—it is simply much more dynamic, flexible, unpredictable, and rich than many other representations. Let's break it down by looking at an example. The video below explains how an investigation on past climate change fits into the elements of the science flow chart.

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Testable Research Questions

Only ***testable ideas*** are within the purview of science. For an idea to be testable, it must logically generate specific expectations—in other words, a set of observations that we could expect to make if the idea were true and a set of observations that would be inconsistent with the idea and lead you to believe that it is not true.

Example

Some topics initially appear to be scientific, but actually are not. For example, the Intelligent Design movement promotes the idea that many aspects of life are too complex to have evolved without the intervention of an intelligent cause—assumed by most proponents to be a **supernatural** being, like God. Promoters of this idea are interested in explaining what we observe in the natural world (the features of living things), which does align well with the aims of science. However, because Intelligent Design relies on the action of an unspecified "intelligent cause," it is not a **testable idea**. The *Understanding Science* website has more information.

A related aspect to being testable is that the question is in regards to the ***natural world***. This includes the components of the physical universe around us like atoms, plants, ecosystems, people, societies, and galaxies, as well as the natural forces at work on those things. In contrast, science cannot study ***supernatural forces*** and explanations. For example, the idea that a supernatural afterlife exists is not a part of science because this afterlife operates outside the rules that govern the natural world.

For example:

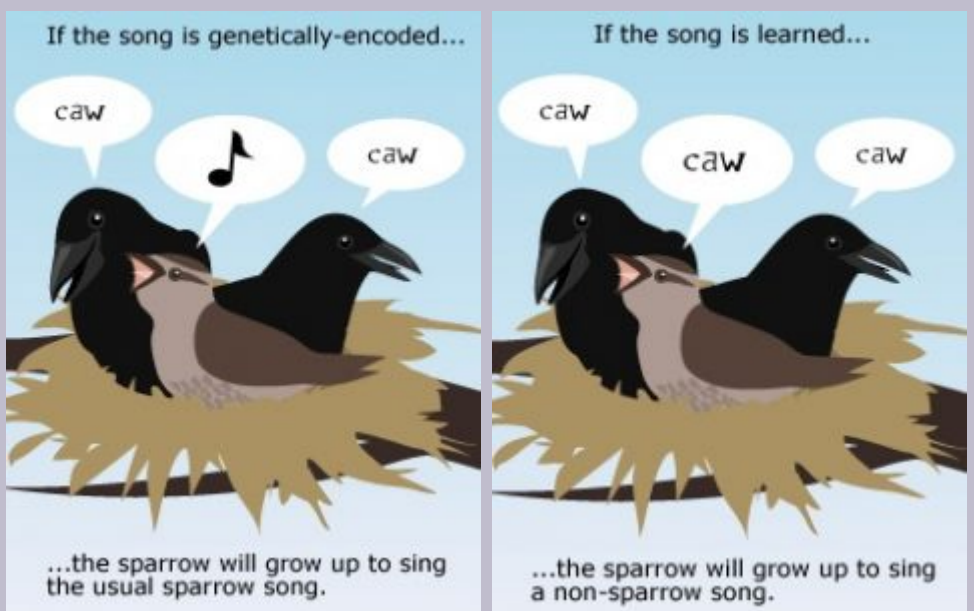
- When did the oldest rocks on earth form?
- Through what chemical reactions do fungi get energy from the nutrients they absorb?
- What causes Jupiter's red spot?
- How does smog move through the atmosphere?

Example of a Scientific Question

Question: Is a sparrow's song genetically-encoded or learned?

Test: See what happens when a sparrow is raised in the nest of another species.

Hypotheses:



Observations and Inferences

We typically think of observations as having been seen "with our own eyes," but in science, ***observations*** can take many forms. Of course we can make observations directly by seeing, feeling, hearing, and smelling, but we can also extend and refine our basic senses with tools: thermometers, microscopes, telescopes, radar, radiation sensors, X-ray crystallography, mass spectroscopy, etc. These tools do a better job of observing than we can! Further, humans cannot directly sense many of the phenomena that science investigates (no amount of staring at this computer screen will ever let you see the atoms that make it up or the UV radiation that it emits) and, in such cases, we must rely on indirect observations facilitated by tools. Through these tools, we can make many more observations much more precisely than those our basic senses are equipped to handle.

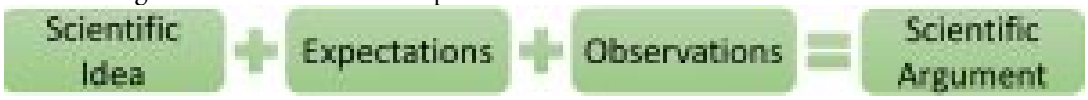
Scientific Arguments

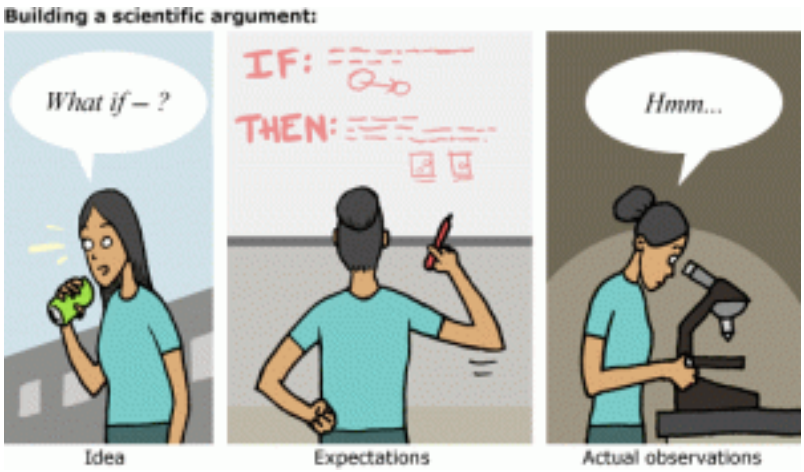
Taken together, the expectations generated by a scientific idea and the actual observations relevant to those expectations form what we'll call a ***scientific argument***.

In this case, the term *argument* refers not to a disagreement between two people, but to an evidence-based line of reasoning; so scientific arguments are more like the closing argument in a court case (a logical description of what we think and why we think it) than they are like the fights you may have had with siblings.

A scientific argument uses evidence to make a case for whether a scientific idea is accurate or inaccurate. For example, the idea that illness in new mothers can be caused by doctors' dirty hands ***generates*** the expectation that illness rates should go down when doctors are required to wash their hands before attending births. When this test was actually performed in the 1800s, the results matched the expectations, forming a strong scientific argument in support of the idea—and hand-washing!

Scientific arguments involve three components:





These components are always related in the same logical way:

image

1. What would we expect to see if this idea were true (i.e., what is our expected observation, frequently called predictions)?

2. What do we actually observe that is relevant to those expectations (to be used as evidence)?

3. Do our expectations match our observations?

Tests typically generate what scientists think of as raw data—unaltered observations, descriptions, or measurements—but those must be analyzed and interpreted. Data become evidence only when they have been interpreted in a way that reflects on the accuracy or inaccuracy of a scientific idea.

For example, an investigation of the evolutionary relationships among crustaceans, insects, millipedes, spiders, and their relatives might tell us the genetic sequence of a particular gene for each organism. This is raw data, but what does it mean? A long series of the As, Ts, Gs, and Cs that make up genetic sequences don't, by themselves, tell us whether insects are more closely related to crustaceans or to spiders. Instead, that data must be analyzed through statistical calculations, tabulations, and/or visual representations. In this case, a biologist might begin to analyze the genetic data by aligning the different sequences, highlighting similarities and differences, and performing calculations to compare the different sequences. Only then can she interpret the results and figure out whether or not they support the hypothesis that insects are more closely related to crustaceans than to spiders.

Furthermore, the same data may be interpreted in different ways. So another scientist could analyze the same genetic data in a new way and come to a different conclusion about the relationships between insects, crustaceans, and spiders. Ultimately, the scientific community will come to a consensus about how a set of data should be interpreted, but this process may take some time and usually involves additional lines of evidence.

Testing Ideas

Ultimately, scientific ideas must not only be testable but must actually be tested—preferably with many different lines of **evidence** by many different people. This characteristic is at the heart of all science. Now that we have learned about the complexity of science as a process. let's explore some of the common ways in which scientists test ideas.

Experimental Studies

An ***experiment*** is a test that involves manipulating a variable in a system. In an experiment, some subjects or individuals have the manipulated variable (called an ***experimental group***) and other subjects have the original form of the variable. (called a ***control group***).

Ideally, experiments also involve keeping as many other factors as ***constant*** as possible to isolate the cause of the experimental results. Experiments can be quite simple tests set up in a lab—like rolling a ball down different inclines to see how the angle affects the rolling time. But large-scale experiments can also be performed out in the real world. For example, classic experiments in ecology involved removing a species of barnacles from intertidal rocks on the Scottish coast to see how that would affect other barnacle species over time. But whether they are

large- or small-scale, performed in the lab or in the field, and require years or mere milliseconds to complete, experiments are distinguished from other sorts of tests by their reliance on the intentional manipulation of some factors and, ideally, keeping others constant.

Observational Studies

Experiments are one way to test some sorts of ideas, but science doesn't live on experiments alone. For many ideas in science, testing via experiment is impossible, inappropriate, or only part of the picture. In those cases, testing is often a matter of making the right ***observations***. For example, we can't actually experiment on distant stars in order to test ideas about which nuclear reactions occur within them, but we can test those ideas by building sensors that allow us to observe what forms of radiation the stars emit. Similarly, we can't perform experiments to test ideas about what *T. rex* ate, but we can test those ideas by making detailed observations of their fossilized teeth and comparing those to the teeth of modern organisms that eat different foods.

Modeling Studies

In science, the term ***model*** can mean several different things, such as an idea about how something works, a physical model of a system that can be used for testing or demonstrative purposes, or a mathematical model (a set of equations that indirectly represents a real system). These equations are based on relevant information about the system and on sets of hypotheses about how the system works.

Given a set of parameters, a model can generate expectations about how the phenomenon will behave in a particular situation. A model and the hypotheses it is based upon are supported when the model generates expectations that match the behavior of its real-world counterpart. In other words, the hypotheses are the potential ***causes*** of an observed phenomenon—the phenomenon being the resulting ***effects*** of that cause. Therefore, models test cause and effect relationships. Modeling often involves idealizing the system in some way, leaving some aspects of the real system out of the model to isolate particular factors or to make the model easier to work with computationally.

Are Model Organisms Scientific Models?

A ***model organism*** is a non-human species that has been widely studied, usually because it is easy to maintain and breed in a laboratory setting and has particular experimental advantages. For example, they may have particularly robust embryos that are easily studied and manipulated in the lab, which is useful for scientists studying development. Or, they may occupy a pivotal position in the evolutionary tree; this is useful for scientists studying evolution.

If a study uses a model organism, is it a modeling study? The answer may seem like an obvious "yes", but consider how modeling organisms are used. For instance, what if the modeling organism is used in an experimental design in which some individuals are part of an ***experimental group*** and others are in a ***control group***. The purpose of doing the research may be ultimately to predict how the treatment may affect humans, but what is the research question? The research question is "how does treatment x affect this model species?" Scientists then use that information to ***infer*** how the treatment might affect humans, but it cannot by itself

actually answer the question of how humans will be impacted. Instead, it is *inferred* how humans may be impacted based on prior research that connects physiological similarities and differences between the model species and humans.

Therefore, when asking if a study is a modeling study, do not assume it is because it is using a model species. Rather, consider what the research question is, not just the general purpose of doing the research.

Exercise

Now try applying what you have learned about scientific research to an investigation!

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Science Checklist

To conclude this introduction to the nature of science, we'll look at a checklist that summarizes key characteristics of science and compare it to an "atypical" example of science to see how they measure up and which characteristics they share.

This checklist provides a guide for which sorts of activities are encompassed by science, but since the boundaries of science are not clearly defined, the list should not be interpreted as all-or-nothing. Some of these characteristics are particularly important to science (e.g., all of science must ultimately rely on evidence), but others are less central. For example, some perfectly scientific investigations may run into a dead-end and not lead to ongoing research. Use this checklist as a reminder of the usual features of science. If something doesn't meet most of these characteristics, it shouldn't be treated as science.

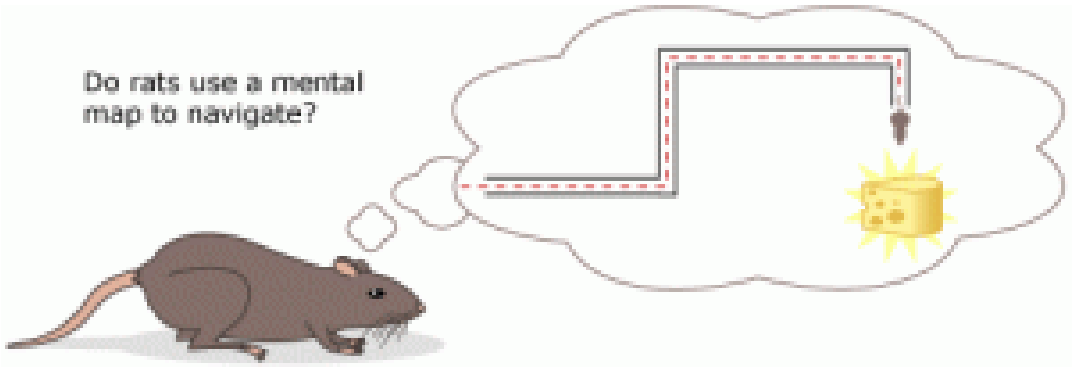
Science Checklist

How scientific is it?

- Focuses on the natural world
- Aims to explain the natural world
- Uses testable ideas
- Relies on evidence
- Involves the scientific community
- Leads to ongoing research
- Benefits from scientific behavior

Example Case Study:

Most of us have probably wondered how other animals think and experience the world (e.g., is Fido really happy to see me or does he just want a treat), but can that curiosity be satisfied by science? After all, how could we ever test an idea about how another animal thinks? In the 1940s, psychologist Edward Tolman investigated a related question using the methods of science. He wanted to know how rats successfully navigate their surroundings—for example, a maze containing a hidden reward. Tolman suspected that rats would build mental maps of the maze as they investigated it (forming a mental picture of the layout of the maze), but many of his colleagues thought that rats would learn to navigate the maze through stimulus-response, associating particular cues with particular outcomes (e.g., taking this tunnel means I get a piece of cheese) without forming any big picture of the maze.



Here's how Tolman's investigation measures up against our checklist:

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So is it science? Though less stereotypically scientific than splitting atoms, this psychological research is very much within the realm of science.

Attributions

This chapter is a modified derivative of the following articles:

Understanding Science. 2020. University of California Museum of Paleontology. 11 June 2020 <http://www.understandingscience.org>.

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2.

SCIENTIFIC CONTROVERSIES

Andrea Bierema

Learning Objectives

- Define and describe "scientific controversy."
- Identify if a science topic is a scientific controversy.
- Distinguish between scientific hypotheses, theories, and laws.

What is a Scientific Controversy?

The Merriam-Webster Dictionary defines *controversy* as "a discussion marked especially by the expression of opposing views." So, what does it mean for something to be a "scientific" controversy?

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So, what is a scientific controversy?

Exercise

The following activity is a series of scenarios. Determine if each one represents a scientific controversy or not. Each scenario is followed by which aspect of the science flowchart it represents. Below the interactive is a simplified flowchart (see the previous chapter for an in-depth version of the flowchart).

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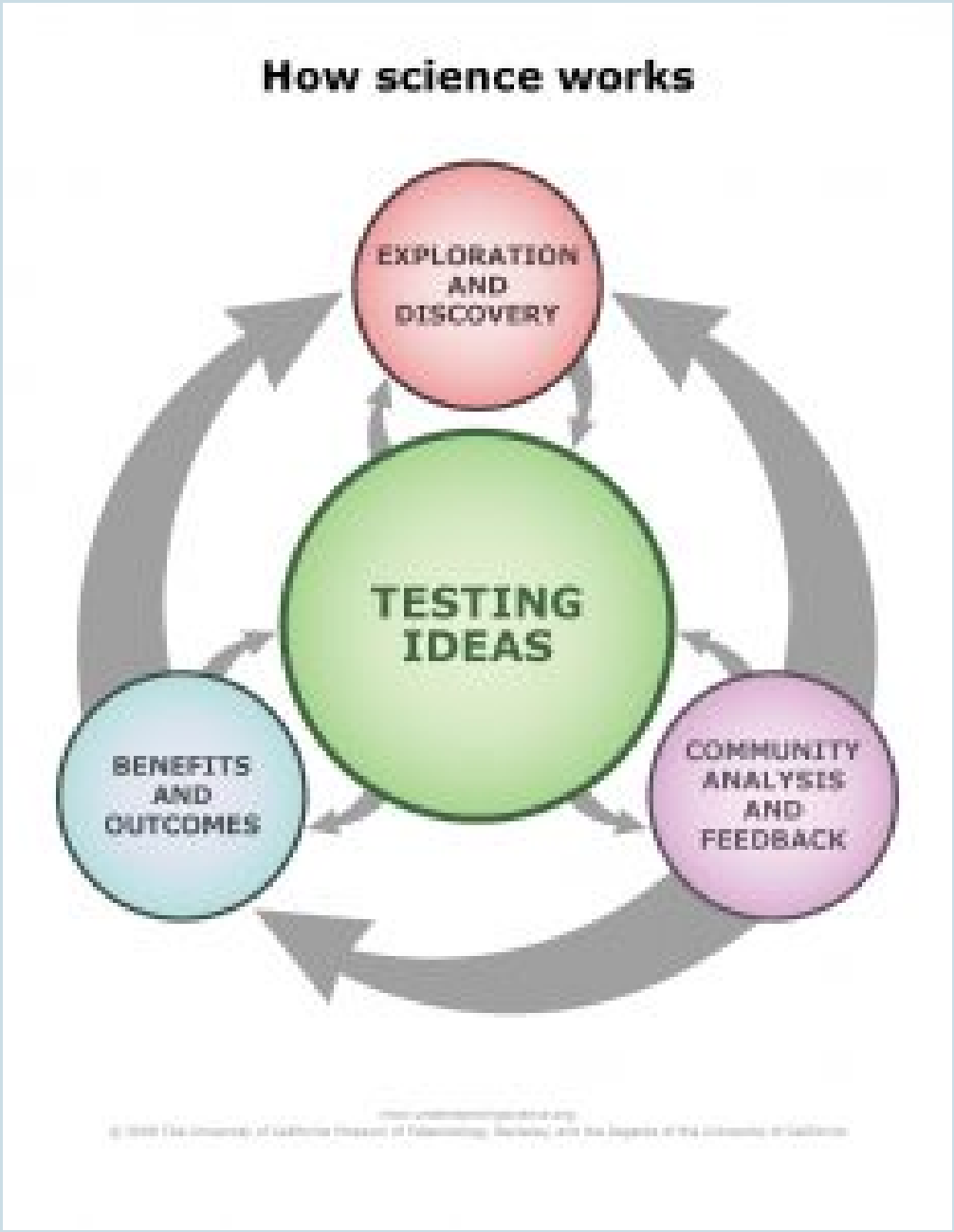

As you saw in the exercise above, *controversies* can take many forms, but many are not considered to be *scientific controversies*. For instance, ethical concerns are essential when society creates policy, but ethics do not use scientific ***evidence*** and so are not scientific controversies (reminder that science cannot answer all questions nor solve all of our problems).

Also, having unexpected results is very common in science, and the fact that things may turn out different than what you expected does not mean that it is a controversy. Even having a single published article contradicting a widely accepted concept does not result in controversy. However, once the scientific community continues to research the concept and is identifying mounting evidence that counters the accepted conclusion, then the concept may be a scientific controversy.

Also seen in the exercise, some ideas may appear to be controversial in science but really are not. This may happen unintentionally. For instance, balanced reporting is generally considered good journalism, and balance does have its virtues. The public should be able to get information on all sides of an issue, but that doesn't mean that all sides of the issue deserve equal weight. Science works by carefully examining the evidence supporting different hypotheses and building on those that have the most support. Journalism and policies that falsely grant all viewpoints the same scientific legitimacy effectively undo one of the main aims of science: to weigh the evidence.

Theories, Hypotheses, and Laws

To really understand scientific controversies, it is essential to know the relationship between laws, theories, and hypotheses.

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Some introductory science courses treat hypotheses as "things we're not sure about yet" and only explore established and accepted theories. In fact, hypotheses, theories, and laws are rather like apples, oranges, and kumquats: one cannot grow into another, no matter how much fertilizer and water are offered. Hypotheses, theories, and laws are all scientific explanations that differ in breadth, not in the level of support. **Hypotheses** are explanations that are limited in scope, applying to a fairly narrow range of phenomena. The term **law** is sometimes used to refer to an idea about how observable phenomena are related—the term is also used in other ways within science. **Theories** are deep explanations that apply to a broad range of phenomena and that may integrate many hypotheses and laws.

Theories

Theories are broad explanations for a wide range of phenomena. They are concise (i.e., generally don't have a long list of exceptions and special rules), coherent, systematic, predictive, and broadly applicable. In fact, theories often integrate and generalize many hypotheses. For example, the theory of **natural selection** broadly applies to all populations with some form of inheritance, variation, and differential reproductive success—whether that population is composed of alpine butterflies, fruit flies on a tropical island, a new form of life discovered on Mars, or even bits in a computer's memory. This theory helps us understand a wide range of observations (from the rise of antibiotic-resistant bacteria to the physical match between pollinators and their preferred flowers), makes predictions in new situations (e.g., that treating AIDS patients with a cocktail of medications should slow the evolution of the virus), and has proven itself time and time again in thousands of experiments and observational studies.

Hypotheses

Hypotheses are proposed explanations for a fairly narrow set of phenomena. These reasoned explanations are not guesses—of the wild or educated variety. When scientists formulate new hypotheses, they are usually based on prior experience, scientific background knowledge, preliminary observations, and logic. For example, scientists observed that alpine butterflies exhibit characteristics intermediate between two species that live at lower elevations. Based on these observations and their understanding of speciation, the scientists hypothesized that this species of alpine butterfly evolved as the result of hybridization between the two other species living at lower elevations.

Laws

In everyday language, a law is a rule that must be abided or something that can be relied upon to occur in a particular situation. Scientific laws, on the other hand, are less rigid. They may have exceptions, and, like other scientific knowledge, may be modified or rejected based on new evidence and perspectives. In science, the term *law* usually refers to a generalization about data and is a compact way of describing what we'd expect to happen in a particular situation.

Some laws are non-mechanistic statements about the relationship between observable phenomena. For example, the ideal gas law describes how the pressure, volume, and temperature of a particular amount of gas are related to one another. It does not describe how gases must behave; we know that gases do not precisely conform to the ideal gas law.

Other laws deal with phenomena that are not directly observable. For example, the second law of thermodynamics deals with entropy, which is not directly observable in the same way that volume and pressure are. Still, other laws offer more mechanistic explanations of phenomena. For example, Mendel's first law offers a model of how genes are distributed to gametes and offspring that help us make predictions about the outcomes of genetic crosses.

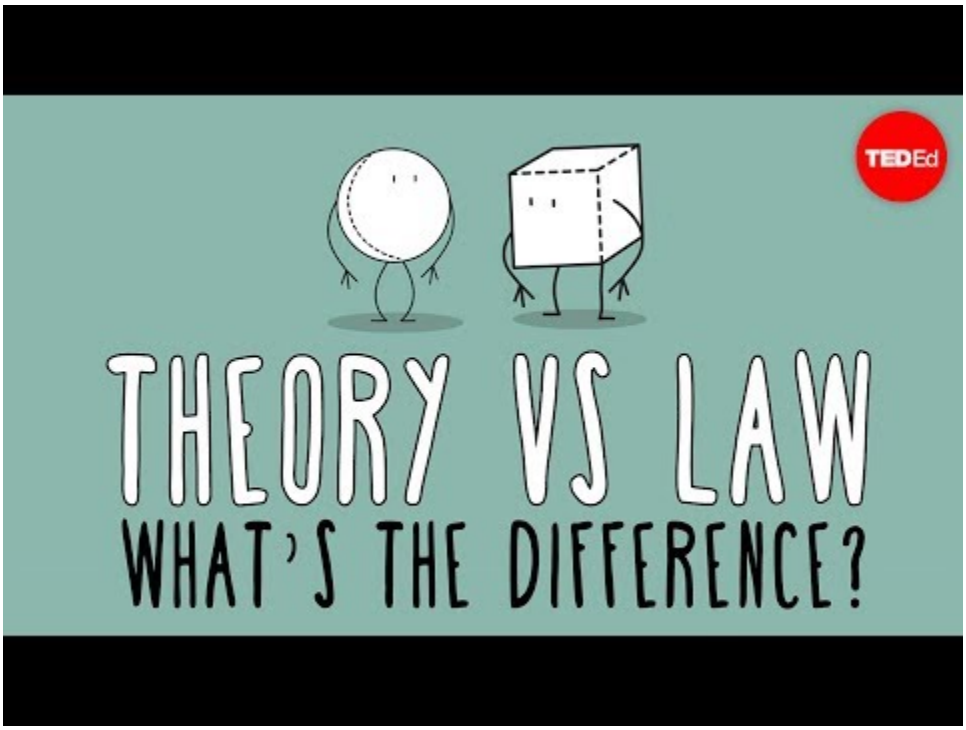
The term "law" may be used to describe many different forms of scientific knowledge, and whether or not a particular idea is called a law has much to do with its discipline and the time period in which it was first developed.

Controversies

Fundamental scientific controversy: scientists disagreeing about a central hypothesis or theory. If you imagine scientific knowledge as a web of interconnected ideas, theories and hypotheses are at the center of the web and are connected to many, many other ideas—so, a controversy over one of these principal ideas has the potential to shake up the state of scientific knowledge. For example, physicists are currently in disagreement over the basic validity of string theory, the set of key ideas that have been billed as the next big leap forward in theoretical physics. This is a fundamental scientific controversy.

Secondary scientific controversy: scientists disagreeing about a less central aspect of a scientific idea. For example, evolutionary biologists have different views on the importance of punctuated equilibrium (a pattern of evolutionary change, characterized by rapid evolution interrupted by many years of constancy). This controversy focuses on an important aspect of the mode and rate of evolutionary change, but a change in scientists' acceptance of punctuated equilibrium would not shake evolutionary biology to its core. Scientists on both sides of the punctuated equilibrium issue accept the same basic tenets of evolutionary theory.

For an overview of the relationship between theories and laws, watch the following video:



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Honesty is the Best Policy

In science, honesty really is the best policy—even if that means publicizing a slip-up. Geoffrey Chang, a professor at the Scripps Research Institute, has made a successful career working out the physical structures of proteins used in cell membranes. His work was published in top journals and cited by other scientists many times. Then, in 2006, he found a mistake. Prompted by conflicting results from other researchers, Chang discovered that, for the past five years, he had been analyzing his data with a flawed computer program, leading to incorrect results. So what did he do? Exactly what the culture of science expected of him: he published letters retracting his previous work, offered an apology, and then started the work of reanalyzing his data in order to correct his results.

What happens when someone within the community doesn't meet those expectations? In science, not playing by the rules amounts to scientific misconduct, or at least scientific misbehavior. Serious misconduct is rare, but nevertheless, because scientists are people and have human frailties, it does happen. Perhaps a chemist is asked to review the paper of a personal friend and chooses to overlook a flaw in the research; thus, failing to fairly scrutinize the work. Perhaps a physicist performs an experiment and chooses only to report results that fit

with his or her favorite hypothesis; thus, failing to be fully honest. Perhaps a biologist writes a research article but doesn't cite a previous study that inspired the work; thus, failing to assign credit fairly. Or perhaps a psychologist studies a group of students' problem-solving skills but circumvents a few guidelines about how the participants should be recruited; thus, failing to work within the ethical guidelines established by the scientific community. Such behavior works against one of science's main goals: to build accurate knowledge about how the world works in ways that are ethical and humane.

Now that you have learned about scientific controversies, let's look at an example that is commonly referred to as a controversy, but it actually is not. We will look at the history of how the connection of vaccination and autism was created and how this led to further investigation and eventually, an article being retracted—an extremely rare event.

Because it undermines science, scientists take misconduct very seriously. In response to misconduct, the scientific community may withhold esteem, job offers, and funding; effectively preventing the offender from participating in science. For example, a scientist found to have plagiarized parts of a grant application to the National Institutes of Health will likely be prevented from participating in federally funded grants for a period of time—a tough punishment for someone whose salary may be partly dependent on such grants. Some types of misconduct are even punishable by law. For example, because he faked data in funding applications and journal articles, medical researcher Eric Poehlman received a $180,000 fine, a year in prison, and a lifetime ban on receiving federal research funds!

Serious and damaging cases of scientific misconduct are almost invariably found out. That's because science is designed to get at how the world really works. Any fraudulent results that paint a false picture of the world will be uncovered as science proceeds and zooms in on the true picture. For example, in the early 1900s, the influential physiologist Emil Abderhalden claimed to have shown that humans produced protective enzymes that could be used in many practical ways—foremost among them, detecting pregnancy. The only problem? Such enzymes don't actually exist. So, of course, Abderhalden's fraud was eventually found out by other scientists who could not reproduce his test results and found that his pregnancy test simply didn't work.

Science's system of scrutiny, peer review, and checks and balances help accelerate the process of discovering and weeding out occasional cases of fraud. For example, the world was first clued into Woo Suk Hwang's fraudulent claims regarding stem cells when other scientists scrutinizing his work drew attention to an anomaly: some of his data looked too good to be true. DNA fingerprint graphs purportedly representing DNA from different samples showed peaks that seemed to be exact duplicates of one another—more likely the result of image manipulation than actual DNA fingerprinting analysis. The ensuing investigation revealed that the copycat graph peaks were only the tip of the iceberg. In fact, Hwang's basic claim—that his lab had cloned human embryos and collected stem cells from them—turned out to be entirely fabricated!

Such flagrant examples of fraud can be disturbing and should lead to the indictment of offenders, but they should not lead to the indictment of science. Science has many safeguards in place to prevent fraud, and when fraud does happen, science has mechanisms for detecting it. Scientific misconduct may temporarily lead science towards incorrect conclusions, but the ongoing processes of science regularly correct such diversions.

Examples

Let's look at some examples of science topics to determine if and to what extent they are scientific controversies.

Vaccines & Autism

Does vaccination cause autism, and is this a scientific controversy?

As emphasized by the World Health Organization and the Centers for Disease Control and Prevention, there is no evidence supporting that vaccination causes autism.

The idea of this possible link began by Wakefield and his research group when they published a paper in "The Lancet" journal in 1998 titled "Ileal-lymphoid-nodular hyperplasia, non-specific colitis, and pervasive developmental disorder in children" (see the original article, which has since been retracted). In the article, Wakefield et al. (1998) claimed that "In eight children, the onset of behavioural problems had been linked, either by the parents or by the child's physician, with measles, mumps, and rubella vaccination."

After years of investigation and additional research studies—all of which found no link between autism and vaccination (Eggertson, 2010)—the General Medical Council held a hearing in 2010. During this hearing, Wakefield "admitted and found proved" that the research was funded by Mr. Barr, who "had the benefit of public funding from the Legal Aid Board in relation to the pursuit of litigation against manufacturers of the MMR vaccine" (p. 4).

Additionally, the children in the study were not randomly selected for the study. Rather, according to the General Medical Council (2018), each child in the study was carefully selected after conversations with the children's parents and doctors. For instance, the report described procedures for the selection of each child, and Child 3 was referred by the child's general practitioner for having "behavioural problems of an autistic nature, severe constipation and learning difficulties all associated by his parents with his MMR vaccination" (p. 18). "In reaching its decision, the Panel concluded that [Wakefield's] description of the referral process as "routine", when it was not, was irresponsible and misleading and contrary to [Wakefield's] duty as a senior author" (p. 46).

The General Medical Council (2018) also found other irresponsible measures such as telling assistants to increase the amount of medication without reporting it to the doctor and taking blood samples from children at his son's birthday party (documented on pages 50-56).

Wakefield and the so-called link between vaccination and autism are both discredited, and Wakefield is no longer a practicing physician. In 2016, Wakefield directed a propaganda movie called "Vaxxed: From Cover-Up to Catastrophe." The movie was supposed to air at the Tribeca Film Festival but was pulled (Ryzik, 2016). Robert De Niro, one of the founders of the festival, originally supported the movie but later denounced it (Ryzik, 2016). Although, as of August 2020, a video clip of him supporting it is still on the Vaxxed website.

So, is this a scientific controversy? The Wakefield article caused scientists to investigate the potential for a relationship. But, no additional evidence was found showing a link between vaccination and autism, making the only "evidence collected" based on a retracted article written by a discredited physician. Therefore, the idea of any link between vaccines and autism is not a scientific controversy.

More Information

To learn more about vaccination and immunity, check out *OpenStax*'s chapter on Vaccines.

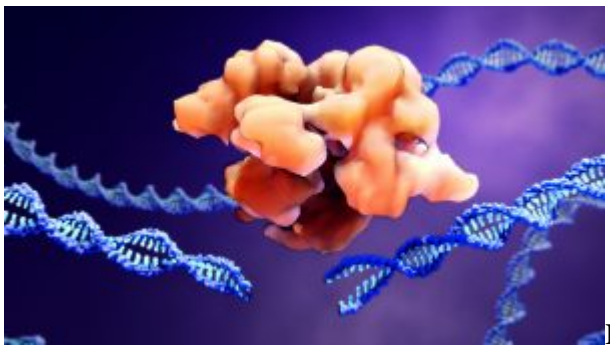
CRISPR

Genome editing technologies enable scientists to make changes to DNA, leading to changes in physical traits like eye color and disease risk. Scientists use different technologies to do this. These technologies act like scissors, cutting the DNA at a specific spot. Then scientists can remove, add, or replace the DNA where it was cut.

The first genome editing technologies were developed in the late 1900s. More recently, a new genome-editing tool called **CRISPR**, invented in 2009, has made it easier than ever to edit DNA. CRISPR is simpler, faster, cheaper, and more accurate than older genome editing methods. Many scientists who perform genome editing now use CRISPR.

One way that scientists use genome editing is to investigate different diseases that affect humans. They edit the genomes of animals, like mice and zebrafish, because animals have many of the same genes as humans. For example, mice and humans share about 85 percent of their genes! By changing a single gene or multiple genes in a mouse, scientists can observe how these changes affect the mouse's health and predict how similar changes in human genomes might affect human health.

Scientists also are developing gene therapies—treatments involving genome editing—to prevent and treat diseases in humans. Genome editing tools have the potential to help treat diseases with a genomic basis, like cystic fibrosis and diabetes. There are two different categories of gene therapies: germline therapy and somatic therapy. Germline therapies change DNA in reproductive cells (like sperm and eggs). Changes to the DNA of reproductive cells are passed down from generation to generation. Somatic therapies, on the other hand, target non-reproductive cells, and changes made in these cells affect only the person who receives the gene therapy.

 Even though CRISPR improved upon older genome editing technologies, it is not perfect. For example, sometimes genome editing tools cut in the wrong spot. Scientists are not yet sure how these errors might affect patients. Assessing the safety of gene therapies and improving genome editing technologies are critical steps to ensure that this technology is ready for use in patients.

There are also several ethical concerns that can emerge with genome editing, including safety. First and foremost, genome editing must be safe before it is used to treat patients. Some other ethical questions that scientists and society must consider are:

- Is it okay to use gene therapy on an embryo when it is impossible to get permission from the embryo for treatment? Is getting permission from the parents enough?
- What if gene therapies are too expensive and only wealthy people can access and afford them? That could worsen existing health inequalities between the rich and poor.
- Will some people use genome editing for traits not important for health, such as athletic ability or height? Is that okay?
- Should scientists ever be able to edit germline cells? Edits in the germline would be passed down through generations.

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Most people agree that scientists should not edit the genomes of germline cells at this time because the scientific communities across the world are approaching germline therapy research with caution since edits to a germline cell would be passed down through generations. Many countries and organizations have strict regulations to prevent germline editing for this reason. The NIH, for example, does not fund research to edit human embryos.

There are many applications of CRISPR, including medical treatments (similar to gene therapy), agriculture (genetically modified organisms), and conservation (adding genetic diversity to species or de-extincting species—described in the next section). Many of these applications have non-CRISPR alternatives that have been around longer. For instance, scientists are researching how CRISPR can be used to treat cancers rather

than chemotherapy. Given the novelty of CRISPR, its effectiveness and safety are continually studied and compared to traditional approaches. Depending on the findings of the research, some of these applications may be a scientific controversy and will take time to solve.

De-extinction

One of the more interesting conservation debates to have emerged in recent years involve efforts to reverse extinction. This field, known as de-extinction or resurrection biology aims to revive extinct species, and eventually to reintroduce viable populations to their original locations (Seddon, 2017).

One possible method, called "breeding back", aims to produce individuals genetically similar to an extinct species by selective breeding of extant species that carry the genetic material of their extinct relatives. This is the main method currently being used to revive the aurochs (*Bos primigenius*), the ancestor of today's domestic cattle (Stokstad, 2015). Other "breeding back" projects place less emphasis on genetics and more on morphology, by selectively breeding individuals with certain traits to produce individuals that visually appear similar to the extinct species. Such is the case at The Quagga Project, where selectively breeding of plains zebras (*Equus quagga*) with quagga-like characteristics (reduced striping and brown hues) are resulting in animals that look increasingly like extinct quaggas (Harley et al., 2009).

The second popular method used for de-extinction is cloning. This involves the transfer of viable genetic material from an extinct species to the eggs (or embryo) of a closely related surrogate mother, who will hopefully give birth to an individual of the extinct species. Cloning has been used in selective breeding of livestock for many years, and plans are also currently underway to use cloning to prevent the extinction of highly threatened species such as the northern white rhinoceros.

Spanish ibex

Despite the promise that cloning offers for reviving extant and recently extinct species, cloning species that went extinct many years ago has been more challenging. So far, attempts to clone Spain's Pyrenean ibex (*Capra pyrenaica pyrenaica*) and Australia's gastric-brooding frog (*Rheobatrachus silus*) have produced individuals that lived for only a few minutes (Ogden, 2014).

Despite the progress made, de-extinction is one of the most controversial and polarising debates to emerge among conservation biologists in recent years. Proponents of de-extinction hope that the early work described above paves the way for the resurrection of extinct species once the threats that drove them to extinction have been managed. Many de-extinction biologists have even started establishing banks where the genetic material of threatened species is cryopreserved for future use.

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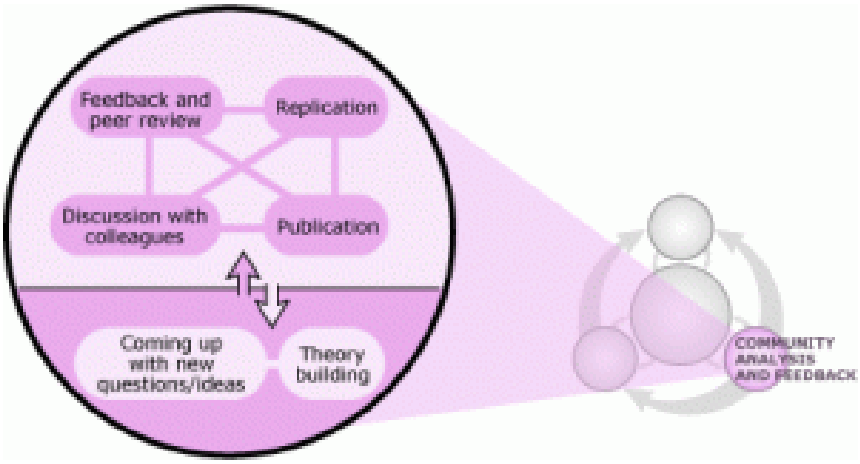
3.

INFORMATION COMMUNICATION

Andrea Bierema and Sara Miller

Learning Objectives

- Recognize that a given scholarly work may not represent the only, or even the majority, perspective on the issue.
- Recognize that information may be perceived differently based on the format in which it is packaged.
- Recognize issues of access or lack of access to scientific information sources.
- Recognize that one's personal information and online interactions affect the information one receives and the information one produces or disseminates online.
- Evaluate an information source.



Community Analysis and Feedback

One of the main elements of the science flowchart is "community analysis and feedback." Members of the scientific community play several essential and direct roles:

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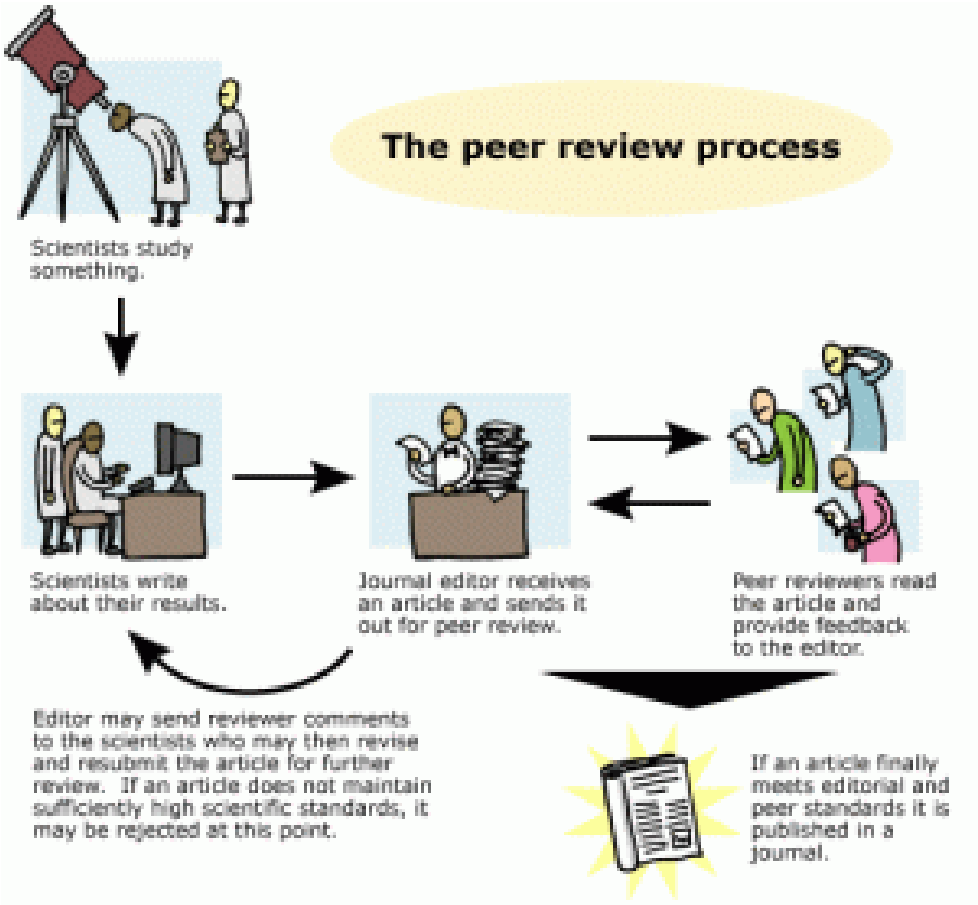

Learn more about the peer review process of scientific articles!

The scientific community provides a system of checks and balances that ensures the quality of scientific work, double-checks arguments, and makes sure that ideas are evaluated fairly. This scrutiny can serve a few different functions from fact-checking to whistleblowing.

The community evaluates evidence and ideas. Scientists describe their work at conferences, in journal articles, and in books. By disseminating their ideas, study methods, and test results in these ways, scientists allow other community members to check their work, both by reviewing what has been done and trying to replicate all or part of it. This helps to ensure that evidence meets high standards, that all relevant lines of evidence are explored, that judgments are not based on flawed reasoning, and hence, that science moves in the direction of more and more accurate explanations. For example, in 1989, when two scientists claimed to have produced nuclear fusion at temperatures lower than was thought possible, the scientific community took a close look at their methods and results. Community members found several ways to improve the experiments and several tests that the original researchers had failed to perform. Meanwhile, other scientists got started on trying to replicate the experimental results and discovered that they could not be consistently reproduced. The scientific community ultimately found that the evidence was not compelling enough to warrant accepting the researchers' claims.

Scientists are people, too. They come from different backgrounds, have different personal beliefs, and favor different hypotheses and theories which can result in unintentional biases even when scientists strive to remain objective. Luckily, the scientific community is diverse, and for every scientist who looks at a result through rose-

tinted glasses, there is another who peers at it through her own blue-tinted ones. Because of the community's diversity, individual biases are balanced out and the community as a whole can evaluate scientific ideas fairly.

The community helps identify and eliminate fraud. Though fraud is rare in science, it sometimes happens. These occasional cases of fraud are identified through the scrutiny of the scientific community. For example, a recent case in which medical researcher Jon Sudbø faked data on 900 Norwegian patients was discovered by another scientist familiar with the group of patients with whom he claimed to be working. Because they build upon the work of others, scientists take fraud very seriously. No one wants to build their own work on a shaky foundation supported by fraudulent ideas.

Science depends on its community in many ways: from the specific (e.g., catching a mistake in an article) to the general (e.g., dividing up the enormous amount of work that keeps science moving forward). Being part of that community means meeting some expectations.

Article Types

Scientists distribute information about their ideas in many ways (informally communicating with colleagues, making presentations at conferences, writing books, etc.), but among these different modes of communication, peer-reviewed journal articles are especially important.

A journal article is a formal, souped-up version of the standard high school lab report. In journal articles, scientists (usually a group of collaborators) describe a study and report any details one might need to evaluate that study: background information, data, statistical results, graphs, maps, explanations of how the study was performed and how the researchers drew their conclusions, etc. These articles are published in scientific journals, either in print or on the internet. Print journals look much like any magazine, except that they are chock full of firsthand reports of scientific research. Journals distribute scientific information to researchers all around the world so that they can keep current in their fields and evaluate the work of their peers. Journal articles neaten up the messy process of science by presenting ideas, evidence, and reasoning in a way that's easy to understand—in contrast to the often circuitous

Example

Checks and Balances

Before the 1970s, the field of primatology was dominated by men. Male scientists observed and recorded primate behavior in the wild, male scientists developed explanations to understand those behaviors, and male scientists read and evaluated each others' work. And at that time, observations suggested that primate social life was largely controlled by males, with females playing a more passive role. But that changed when women scientists began to work in the field in the 1970s. Because of their own gender experiences, these women paid more attention to subtleties in the female primates' behavior, and revealed that female primates actually have elaborate sex lives and manipulate male behavior in many ways. So in this case, a diverse assemblage of scientists counterbalanced each others' biases, leading to a more complete

and accurate understanding of primate societies.

(and sometimes tedious) process of science. Check out the article below to learn more about the different ways in which scientific research is published.

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Headlines

- Trans-fat free!
- Ethanol production: an eco-nightmare?
- Cancer researchers discover new hope.
- Major petroleum company acknowledges the reality of global warming.
- Clinically proven to reduce the appearance of wrinkles!

The above statements aren't exactly the headlines you'd find in a scientific journal, but they are examples of the sorts of scientific messages that one might encounter every day. Because science is so critical to our lives, we are regularly targeted by media messages about science in the form of advertising or reporting from newspapers, magazines, the internet, TV, or radio. Similarly, our everyday lives are affected by all sorts of science-related policies from what additives are allowed (or required) to be mixed in with gasoline, to where homes can be built, to how milk is processed, but you don't have to take these media messages and science policies at face value. Understanding the nature of science can help you uncover the real meaning of media messages about science and evaluate the science behind policies.

Exercise

Let's practice!

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Source Evaluation

Every day, we are bombarded with messages based on science: the nightly news reports on the health effects of cholesterol in eggs, a shampoo advertisement claims that it has been scientifically proven to strengthen hair, or the newspaper reports on the senate's vote to restrict carbon dioxide emissions based on their impact on global warming. Media representations of science and science-related policy are essential for quickly communicating scientific messages to the broad public. However, some important parts of the scientific message can easily get lost or garbled in translation. Understanding the nature of science can make you a better-informed consumer of those messages and policies. It can help you:

- separate science from spin
- identify misrepresentations of science, and
- find trustworthy sources for further information.

Moreover, an original piece of scientific research may be interpreted many times over before it reaches you. First, the researchers will write up the research for a scientific journal article, which may then be adapted into a simplified press release, which will be read by reporters and translated yet again into a newspaper, magazine, or internet article and so on. Just as in a game of telephone, errors and exaggerations can sneak in with each adaptation.

Exercise

Before getting into the details on how to analyze information sources, let's consider the following hypothetical article. What are some things to consider while evaluating this article? In the image, click on the exclamation point icons to learn about some concerns regarding this article.

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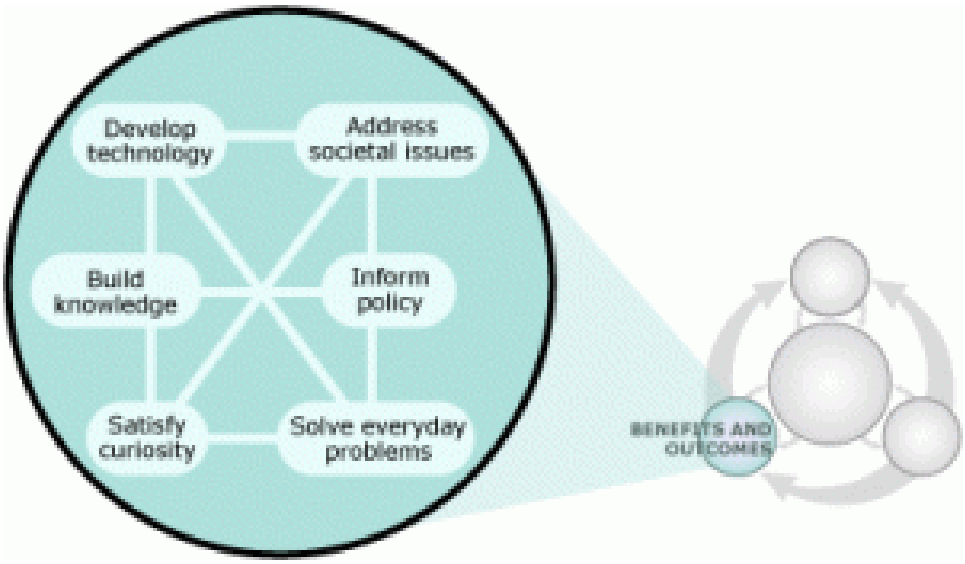
4.

STAKEHOLDERS AND AUTHORITY

Andrea Bierema

Learning Objectives

- Identify different types of authority, such as subject expertise, societal position, or individual experience.
- Recognize that a given information source may not give voice to all—or even the majority— of stakeholders involved in and impacted by an issue, such as conservation.
- Identify and explain the roles of primary, secondary, opposition, and marginalized stakeholders in a given case study.



Scientific studies are used to address societal issues and inform policy, but who is impacted by these implications? This chapter addresses this question.

What is a Stakeholder?

Note

Issues can affect more than just stakeholders. For instance, in conservation, policies include the impact on wildlife. Wildlife are part of the ecosystem rather than a stakeholder. Stakeholders are only people or representations for people

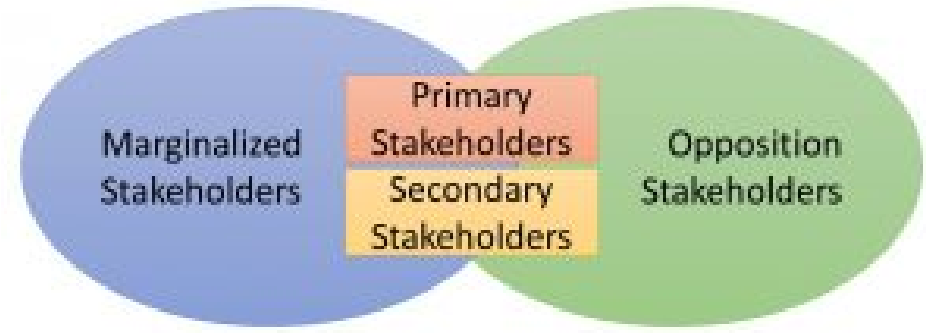
Stakeholders are people, institutions, or social groups that are affected by, and/or involved in, a particular issue, such as the creation of policies. While this definition is seemingly straightforward, it is often difficult to answer fundamental questions such as:

- Who are "the people?"
- What does "institution" mean?
- What are the limits of a "social group?"

Yet, these questions must be answered if the right **stakeholders** are to be identified and mobilized.

Types of Stakeholders

There are stakeholders who directly influence, or are influenced by, outcomes (called "primary stakeholders") and others that indirectly affect, or are affected by, outcomes (called "secondary stakeholders"). Some of these stakeholders may be marginalized stakeholders or opposition stakeholders. See below for definitions.



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Exercise

Answer the following questions after reading through the different types of stakeholders.

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Stakeholder Influence on Decision-Making

In some cases, stakeholders are easy to identify. In other cases, a more in-depth understanding of the issues involved in a particular situation is needed to determine who should be included in the early phases of the process. For a variety of reasons, stakeholders may disagree about who should be included in the dialogue. It may be that a group is perceived as too combative, or is not thought to have the appropriate skills to participate effectively. These objections may or may not be justifiable and may often be the result of historic biases held by stakeholder groups. While assumptions are inevitable, it is important that the initial stakeholder identification process avoids reaching premature conclusions about which stakeholders should, or should not, be involved.

Initially, dialogue should be as open and participatory as possible, encouraging stakeholders from a variety of backgrounds and perspectives to contribute to the identification and framing of collaboration goals and objectives. If the process is not participatory, there is a risk that it will quickly become dominated by the strongest, loudest, or best-resourced groups who seek to shape the process for their own objectives. Over time, it may be determined that additional interests must be brought into the dialogue and the process needs to be open enough to facilitate this.

Given the challenges that "open participation" in a collaboration process brings (in terms of multiple, often conflicting perspectives and interests), many groups choose to promote collaboration more gradually. In these cases, "start-up" involves bringing together like-minded groups and allies. Steering committees can be established by these groups to formulate shared goals and objectives and assess and strengthen capacities before a wider collaboration process is initiated.

In some instances, initial dialogue may lead to consortiums, alliances, or coalitions. While this approach can provide for a strong and coherent voice, there are associated risks. These include the premature establishment of partnerships before issues, opportunities, and appropriate stakeholder roles and responsibilities have been

fully defined. When alliances with only like-minded groups are formed, the risk of generating negative reactions among other stakeholders can increase due to perceived "exclusivity." Effective information sharing, communication, and public education can help alleviate these risks. Development of a strategic plan for progressively bringing in other key stakeholders—primary, secondary, or opposition—will also be essential.

The number of parties engaged in the collaboration process is also an important consideration. All stakeholders do not need to participate all the time, or to the same degree. A review of who is participating in a conservation initiative should be made on a regular basis, and participation revised as needed.

Differences in Power

Power differentials exist in all forms of social organizations and between social groups. The source of these differences may be based on the heredity rights that leaders enjoy in certain cultural settings, or the power differences earned through channels that economic and political opportunity afford individuals and groups.

Two power issues are particularly relevant to facilitators of stakeholder collaboration: addressing power inequalities between key parties so that adequate representation and collaboration can be achieved, and reaching agreements among parties on how a disagreement over issues will be resolved.

Conflicts involving core group values and identity are difficult to resolve. For example, less politically powerful stakeholders may fear that a powerful outsider will impose its views on a process. This may provoke them to withdraw from a given negotiation process even if they stand to benefit from staying involved. Similarly, distrust can make it particularly difficult for the parties to begin constructive talks. Facilitators need to understand the source of this distrust and determine what, if anything, can be done to remove it.

In situations that become polarized around opposing values or identities, facilitators of a collaboration process need to be resourceful in sensitizing participants to the validity of different stakeholder perspectives. Instrumental techniques such as adapting the meeting structure and process to better enable constructive cross-stakeholder learning can be useful.

Stakeholder Scenarios

The following is a series of scenarios that illustrate the stakeholder concepts described above.

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5.

BIBLIOGRAPHIES

Andrea Bierema and Sara Miller

Learning Objectives

- Give credit to the original ideas of others through paper attribution and citation.
- Identify interested parties, such as scholars, organizations, governments, and industries, who might produce information about a topic and then determine how to access that information.
- Match information needs and search strategies to appropriate search tools.
- Manage searching processes and results effectively.
- Correctly cite an information source.

Citing Sources

Citation is something that we do almost every day, whether or not we're aware of it. We may think of citation as a requirement for papers and assignments, which is one function of citation, but it has several purposes. In the sciences, what does it really mean to **cite** your sources and why bother?

Locating Information

Whenever we share a story or link on social media, we're citing—including information about the source—such as where it can be found online. Very simply, this information allows others to find the original source and identify where it came from. If someone is curious about the story you shared, they can read the original post and possibly follow links in that story (citations) which will lead to other discoveries. Citation helps us to find information. If you're reading a news article about a scientific study, check to see if the article provides a link to

the scholarly journal article where the study was published. This link is a form of citation that will help lead you to the original information.

Scientific Conversation

Citation also serves to show a record of how other sources impacted the current source. Scientific research articles published in **scientific journals** always provide a list of citations, which show where the ideas, techniques, and studies that were built upon by the current research came from. This reference creates a sort of paper trail that helps other scientists better evaluate the new study and see how it fits with previous research. By providing a list of references, an author invites other scientists to see for themselves if the ideas the author cites are supported by evidence, if the assumptions he or she makes are justified, and if the techniques described by others have been properly implemented. In this way, citation functions as a record of a conversation: how other scientists' work speaks to and informs new work.

Copyright Infringement and Credit

Another important function of citation is to identify the original creators of information and to give them credit. In science, credit matters. A magazine or newspaper article only sometimes acknowledges the sources of its arguments—the books the author read or the interviews conducted. Science, on the other hand, is scrupulous about giving credit where credit is due. The bibliography or list of citations that you find in scientific research articles serves to credit other scientists for ideas, techniques, and studies that were built upon by the current research.

Legally and ethically, It's important to not give the direct or indirect impression that someone else's work or ideas were written or created by you. When you hear the term plagiarism, it refers to this phenomenon. For example, some of the content in this chapter was created by people other than the listed authors, and to avoid plagiarism, we've given credit to the original authors through citation at the end of the chapter in the "References" section. It's crucial to use proper citation to indicate where that source material or idea originated. If you use copyrighted work in your own creations without citation, it's a copyright infringement—a legal issue—in addition to the ethical issue of plagiarism.

Exercise

Take this quiz from Turnitin to learn more about plagiarism and copyright infringement!

Finding Sources

There are many ways to find both popular and scholarly sources, including Google and Google Scholar. You also have the option of beginning your search through your school's Library website (e.g., MSU Libraries). There are two main advantages to using the Libraries' search as your starting point, especially for scholarly or journal articles. The Libraries' search can make it easier to narrow and sort your searches by type of article or subject. You will also have automatic access to articles for which the library subscribes. Often when you're using Google or Google Scholar, you may be asked to pay or log in to view the whole article (full-text). If you start at the Libraries' page, you typically just need to enter your school ID and password to get access.

Exercise

Watch this linked video about searching the MSU Libraries and answer the questions.

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Citation Guidelines

When citing your sources, your instructor will specify which citation style they would like you to use. For each source that you use, you will want to keep track of certain types of information to include in your citations.

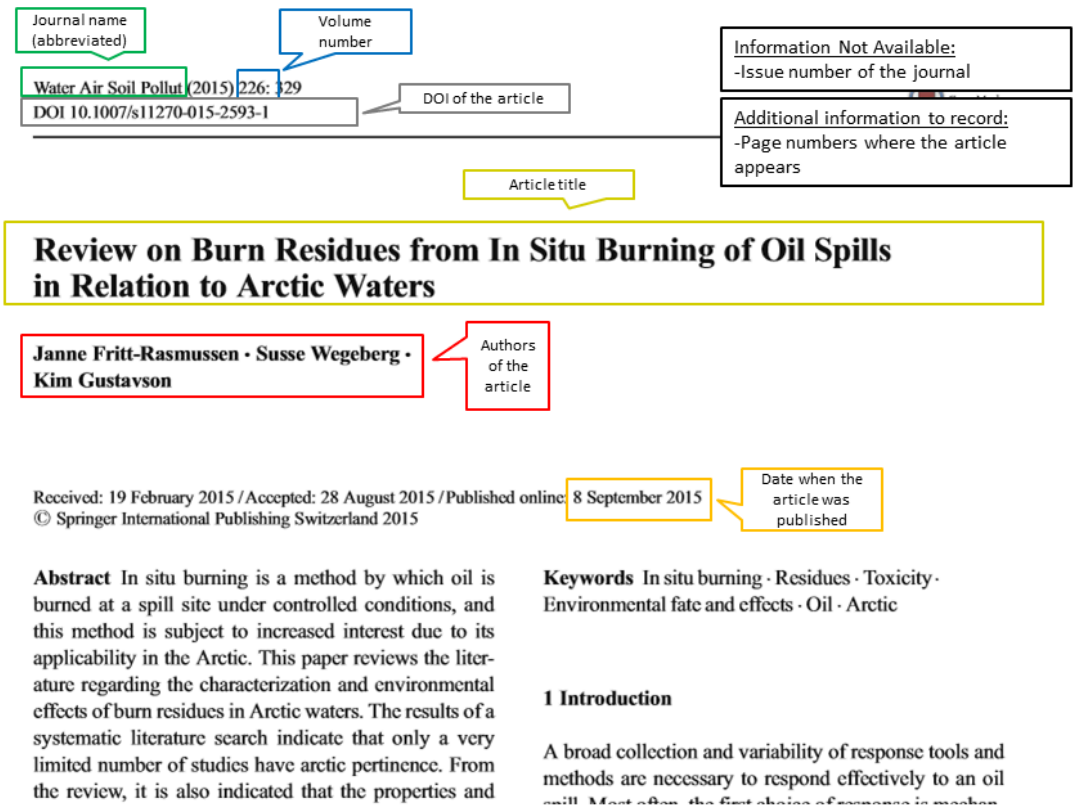
The following is an example of a scholarly journal article with the pieces of information that you'll need to create the citation listed and highlighted. The example comes from Marcyk and Moll's Information Literacy Badges.

Scholarly Journal Article

To cite a scholarly journal article, you'll need to collect the following pieces of information:

- Author(s)
- Date when the article was published
- Title of the article
- Title of the journal
- Volume and issue number
- Page numbers where the article appears
- DOI (alphanumeric code) or URL (website address) of the article

These pieces of information are highlighted in the following image, and the completed APA-style citation is listed beneath the image.



Completed APA Style Citation:

Fritt-Rasmussen, J., Wegeberg, S., & Gustavson, K. (2015). Review on burn residues from in situ burning of oil spills in relation to arctic waters. Water, Air and Soil Pollution, 226(329), n.p. https://doi-org.proxy1.cl.msu.edu/10.1007/s11270-015-2593-1

For more practice with citations, you can go through the Information Literacy Badges' self-guided lesson.

Exercise

For the following citation, match the different parts to their descriptor.

Colla, S. R., & Packer, L. (2008). Evidence for decline in eastern North American bumblebees (hymenoptera: Apidae), with special focus on *bombus affinis* cresson. *Biodiversity and Conservation, 17*(6), 1379-1391. doi:10.1007/s10531-008-9340-5

An interactive or media element has been excluded from this version of the text. You can view it online here:
https://openbooks.lib.msu.edu/isb202/?p=76

References

Marcyk, E. & Moll, E. *Citation as conversation*. Information Literacy Badges. https://informationliteracybadges.org/

Attribution

This chapter is a modified derivative of the following article:

Understanding Science. 2020. University of California Museum of Paleontology. 11 June 2020 <http://www.understandingscience.org>.

Media Attributions

- apa-journal-2

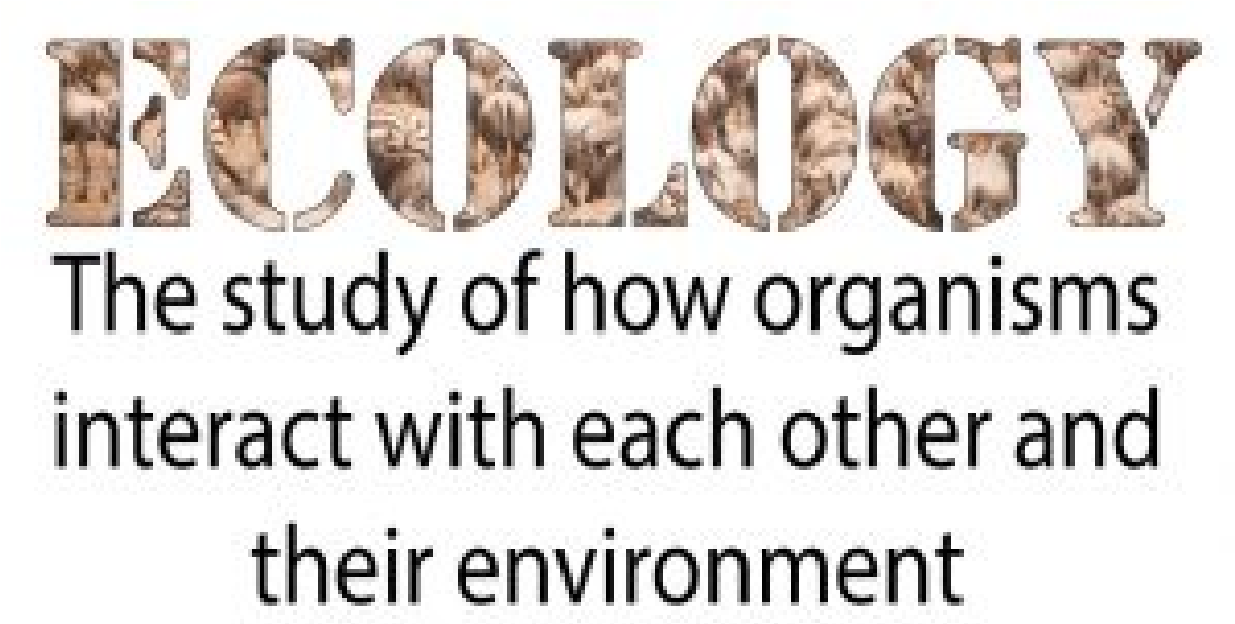
UNIT II

ORGANISMAL BIOLOGY

6.

INTRODUCTION TO ECOLOGY

Andrea Bierema



Ecological Levels

Ecology is studied at different levels. For instance, it can focus on the interaction of individuals within the same species (population ecology) or individuals of different species (community ecology). See the figure below for all of the levels and click on the plus hotspots for definitions and more information.

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In this unit, we will examine ecology at these various levels. First, we examine biodiversity, which addresses all of the ecological levels. Next, we consider the broad levels of biosphere and ecosystem ecology by learning about nutrient cycling and climate change. Then we focus on community ecology, learning about the various ways that species interact with one another. Later, we consider population ecology, focusing on how population sizes change over time and the factors that influence those changes. Lastly, we learn about evolution and modeling evolution using phylogenetic trees, which evolution occurs at the population level but is often influenced by biotic and abiotic factors.

Media Attributions

- Ecology © Andrea Bierema

7.

BIODIVERSITY

Andrea Bierema

Figure 1. From left to right: Row 1: frogfish, moss, sea urchin, woodland crocus, and a flatworm; Row 2: alien lizard, archean cell infected with a virus, monarch butterfly, and platypus; Row 3: weevil, salmonella, hornbill, garden spider, and mushroom.

Learning Objectives

Students will be able to:

- Define biodiversity.
- Use visual models to characterize the scope of biodiversity on earth.
- Describe efforts to conserve threatened and endangered species.
- Explain the Red List of Threatened Species.
- Recognize types of protected areas.
- Describe the benefits of biodiversity.
- Characterize the threats to biodiversity.
- Explain ways in which organizations are working to save biodiversity.

Introduction to Biodiversity

Earth's biodiversity includes the entire range of living species, including single-celled and multicellular organisms. View the following interactive video to learn more!

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The previous video showed how to measure biodiversity and examples of how the amount of biodiversity varies throughout the world. Another aspect of biodiversity is that it is studied at different levels:

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The relationship between species, genetic, and ecosystem diversities is complex and interdependent. That is, a species cannot exist without genetic diversity or ecosystem diversity and vice versa. For that reason, it is virtually impossible to affect one aspect of diversity without affecting the other. We can, therefore, think of species, genetic, and ecosystem diversities simply as different ways to measure the variety of life.

Let's dive deeper into ecosystem biodiversity:

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Benefits of Biodiversity

How do humans benefit from biodiversity? Watch the interactive video to find out!

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Threats to Biodiversity

Despite humans obtaining several benefits from high biodiversity, there are several things that we are doing that limit biodiversity.

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Saving Biodiversity

Although many of our actions threaten biodiversity, there are things that people are doing to minimize our impact and save biodiversity.

Further Information

Curious what you can do to help save biodiversity? Check out the World Wildlife Fund's guide!

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Species Richness and Threat Level Comparison

The following is a series of maps illustrating global species richness as well as the number of threatened species for various taxa. Slide the white vertical line to see the threatened species map and to compare the two maps.

Further Information

If you would like to see maps for additional species, check out the Biodiversity Mapping website!

Terrestrial Species

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Marine Species

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Discussion Question

How do the species-richness and threatened-species maps compare? Are there consistent trends across the different taxa? Why do you think these patterns or lack of patterns exist?

Attributions

The maps in this chapter are from "Biodiversity Mapping" by Jenkins, C. N. Used with permission.

This chapter is a modified derivative of "What is Biodiversity?" by Wilson, J. W., & Primack, R. B., Conservation Biology in Sub-Saharan Africa, Cambridge, UK, Open Book Publishers, 2019.

Figure 1 Attributions

(in order from left to right, starting in row 1)

"Antennarius pictus- Painted frogfish" by Steve Childs is licensed under CC BY 2.0.

"Aloe-moss (Polytrichaceae) at Philip Edward Island in Killarney, Ontario, Canada" by Ryan Hodnett is licensed under CC BY-SA 4.0.

"Tripneustes ventricosus (West Indian sea egg-top) and Echinometra viridis (reef urchin – bottom)" by Nick Hobgood is licensed under CC BY-SA 3.0.

"Flowering woodland crocus in the garden reserve Jonkervallei, Joure, Netherlands" by Dominicus Johannes Bergsma is licensed under CC BY-SA 4.0.

"Lembeh, Indonesia – flatworm – Eurylepta sp." by Rickard Zerpe is licensed under CC BY-SA 2.0.

"The alien lizard" by Kalablink is licensed under CC BY-SA 4.0.

"Sulfolobus infected with the DNA virus STSV1" by Xiangyux is licensed under Public Domain.

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8.

SYSTEMS THINKING AND THE CARBON CYCLE

Andrea Bierema

Learning Objectives

Students will be able to:

- Describe systems thinking.
- Identify and describe a system that is observed in the natural world.
- Apply systems terminology (e.g., closed and open system, reservoir, and flux) to the structure of a systems model, including the carbon cycle.
- Identify the characteristics of positive (reinforcing) and negative (balancing) feedbacks in systems model output.
- Convert information into a carbon cycle model to illustrate the relationships of components within a system.
- Use a carbon cycling system model to predict how a carbon molecular travels through the earth's four spheres.
- Estimate residence time for a system in equilibrium.

Systems Thinking

A system is a set of components that are linked through interconnections and functions to create an outcome. The interconnections of ***components*** and their ***interactions*** create ***system*** behavior. This is a broad definition

and it describes systems in biology (like the circulatory system or nutrient cycling), a game system (like chess, cards, or football), and even a social system (such as 4H or Girl Scouts).

Example

For example, a football team is a system with:

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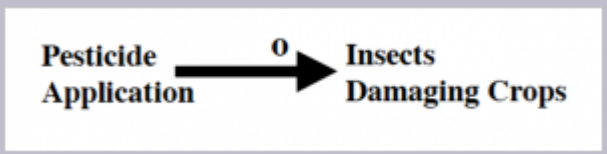
Systems thinking is one way of cognitively framing how we examine the world around us. In contrast to the more common mechanistic or deterministic model that analyzes and understands the whole as the sum of its parts, systems thinking moves the focus of analysis away from the parts themselves and instead concentrates on understanding how the different parts interact with each other.

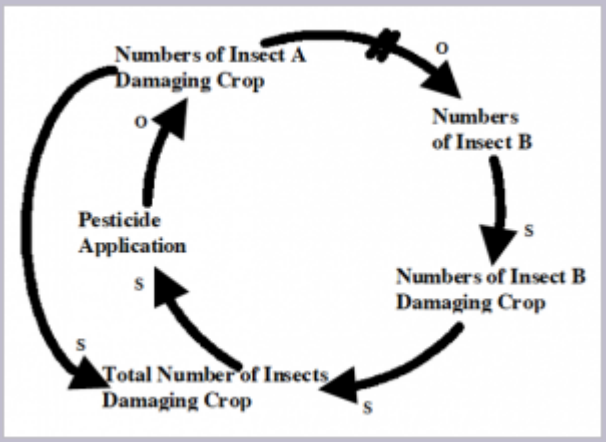
This is a fundamental difference from traditional analysis (analysis means to break into constituent parts); systems thinking works "by expanding its view to take into account larger and larger numbers of interactions" as a system is being studied (Aronson 1996).

Example

Here is a classic example to show the difference between traditional analysis and systems thinking.



"If you apply pesticide, it will kill insect A which is damaging the crops. If you get more insects, apply more pesticide..."



If it were that simple, as the top figure to the left suggests, we'd have no crop losses. Looking at the problem from a broader perspective, we can see one reason why the application of more pesticide doesn't have the expected outcome. As seen in the second figure, the total numbers of insect A are competing with insect B and keeping the population in check. When Insect A is exterminated, insect B's population explodes, and they fill the niche of insect A.

Systems thinking is a modality of thinking that keeps a focus on interactions between parts, with special vigilance to identify unintended consequences of changes that take place in a system because of these interactions.

The components and interactions in a system are referred to by the scientific terms used in system analysis

Now to introduce some important system science concepts. Scientists describe the "container" that holds an Earth material as a reservoir and the interactions as fluxes.

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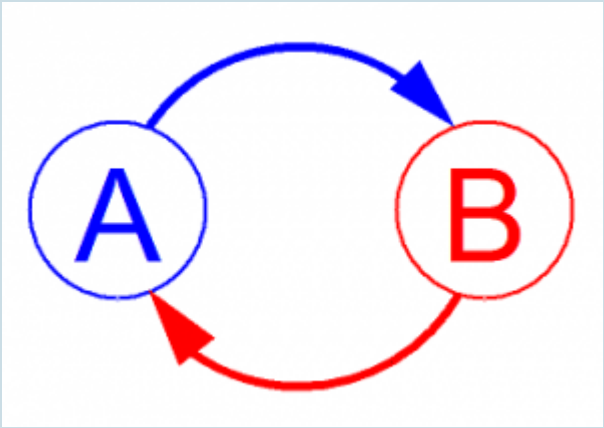
Exercise

A feedback loop exists between two components/reservoirs of a system (labeled “A” and “B” below) when each affects the other.



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Let's look at this concept closer with an example. The following is a slideshow—move the slides with the arrows at the bottom of the slideshow.

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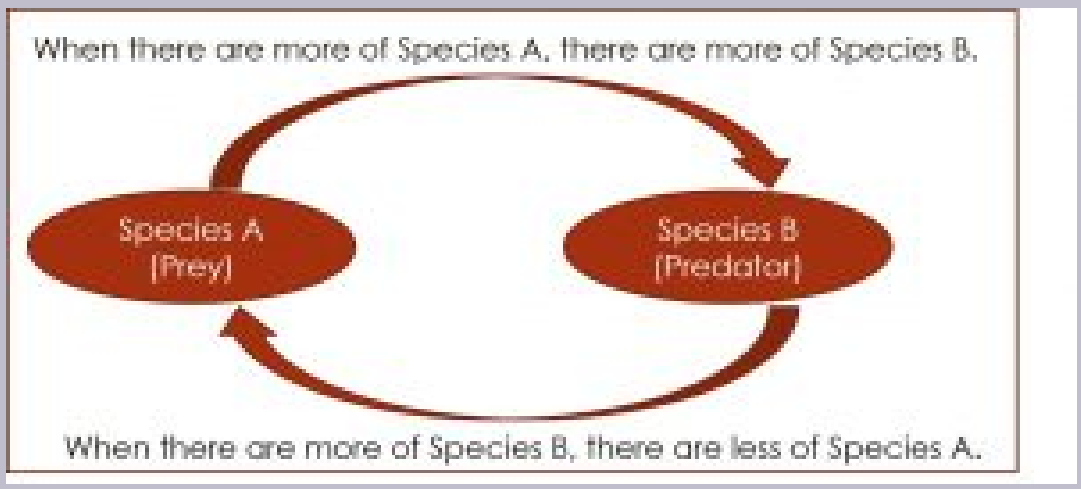
Example



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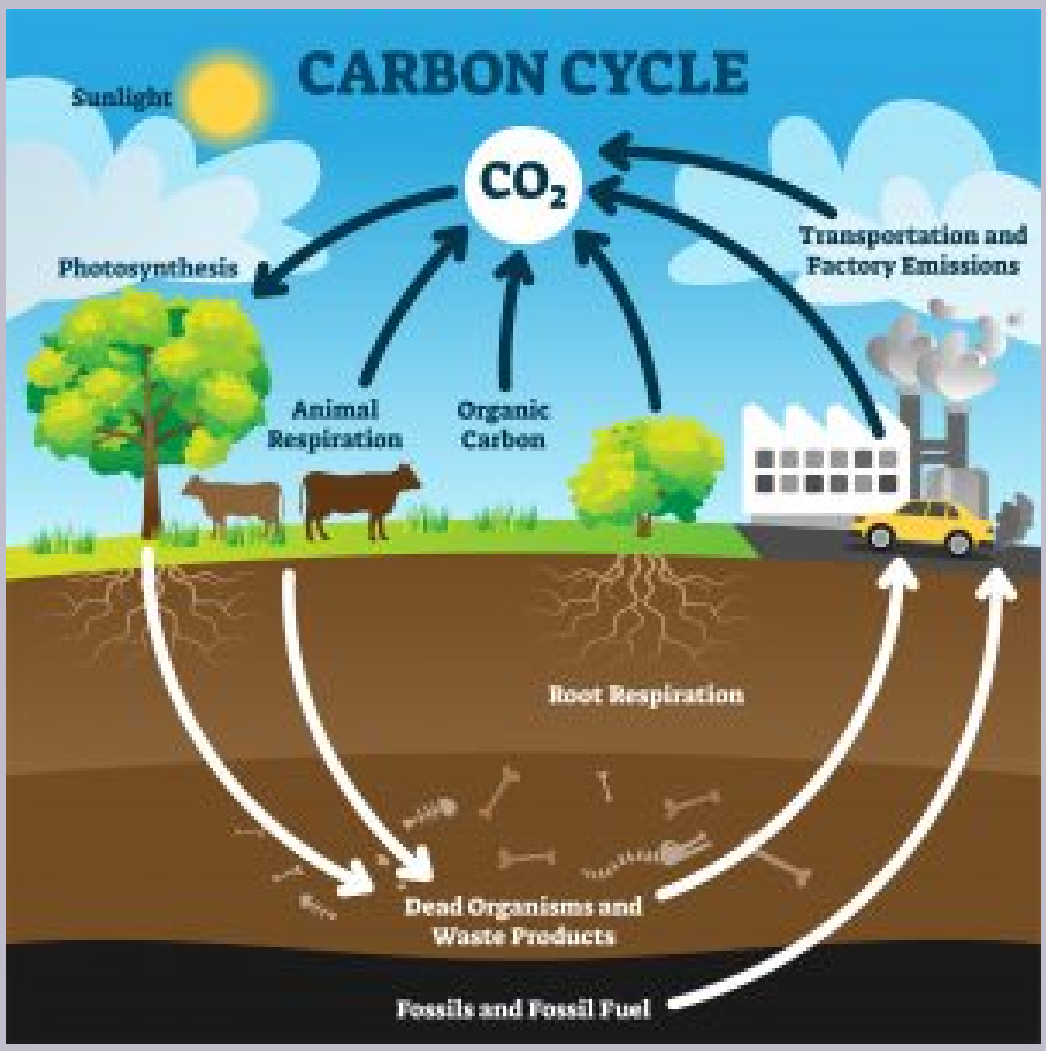
The Carbon Cycle

According to the ***Law of Conservation of Mass***, matter can be neither created nor destroyed. Therefore, when nutrients (i.e., matter) are melted, burned, etc., they aren't destroyed. Also, when more nutrients appear in one part of the earth, it was not created. Rather, nutrients cycle through different parts of the earth. ***Nutrient cycling is a system with fluxes and reservoirs.***

Example

Here is an example of a typical carbon cycle. Apply the concepts learned in the previous section to the carbon cycle using the following diagram.



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The diagram in the example box above is similar to how carbon cycles are drawn in biology textbooks. It is technically a system, but does not explicitly illustrate how carbon moves through Earth's four spheres, depicted below.

The four spheres of Earth include the atmosphere (the air), biosphere (anything living, including organisms living in water), lithosphere (rocks, minerals, sand), and hydrosphere (marine and freshwater).

The following activity will end with a carbon cycle that focuses on these four spheres.

Exercise

The following is a list of processes describing how carbon moves (i.e., the ***fluxes*** of the carbon cycle). Refer to the definitions of each of these processes while performing the activity below.

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Using the definitions of each carbon flux above, develop a carbon cycle model. In the model below, each corner is one of the spheres (e.g., atmosphere), and each arrow represents a flux, which is the movement of carbon from one sphere to another sphere. Notice that one of the arrows (between the atmosphere and hydrosphere) is a double arrow; one of the fluxes represents carbon moving back and forth between the atmosphere and hydrosphere.

To complete the model below, drag each carbon flux (on the right side of the activity) to its appropriate arrow. The two red, dotted-line arrows are human activities, and the black solid-line arrows are natural processes.

Alternatively, feel free to complete the activity below the carbon cycle model, which addresses the same information, but in sentence form.

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This activity meets the same objectives as above, but just in a different format. Feel free to do this one instead or do it before creating the visual model above.

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Potential Carbon Molecule Paths

As seen in the above activity, there are many ways in which carbon moves through the earth's spheres. To combine the ideas of the first carbon diagram shown earlier and the created carbon model, consider what are the more specific reservoirs of carbon in each sphere. For instance, the first model shows trees and cows, which are in the biosphere, and fossil fuels used by factories and cars, in which fossil fuels are part of the lithosphere.

Exercises

Here's one example of how a carbon molecule may travel through the cycle:

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Here's another example; this time moving through all four spheres:

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Residence Time

Residence time is the average amount of time that matter remains in a reservoir. For carbon, this varies rather dramatically from sphere to sphere.

How to calculate residence time, assuming equilibrium in fluxes adding to, and fluxes removing, carbon from a given sphere:

Residence Time = [Amount of substance in sphere] / [total flux in OR out]

For example, if the total amount of carbon in a sphere is 8 billion tons and total flux in or out of the sphere is 4 billion tons per year:

[8 billion tons of carbon] / [4 billion tons per year] = 2 years

The conclusion to be made from this residence time example is that one atom of carbon will be in the given sphere for an average of 2 years before moving naturally to another sphere in the carbon system. Remember that this is an average; some atoms will spend much less time, and some will spend much more, in the sphere.

Exercise

Here is the total amount of carbon in each sphere:

- Biosphere: 605 billion tons
- Atmosphere: 760 billion tons
- Hydrosphere: 39,000 billion tons
- Lithosphere: 40,005,600 billion tons

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Reference

Aronson, D. (1996). Overview of systems thinking. Retrieved from http://www.thinking.net/Systems_Thinking/OverviewSTarticle.pdf

Attributions

This chapter is a modified derivative of the following articles:

"Development and Evaluation of an Inquiry-Based Unit for Teaching about Paleoclimate and Climate Change" by Barone, S., Master's Theses, 113, 2019, Used with permission.

"Introductory system slides." by Gilbert, L. *In* Systems Thinking, InTeGrate, CC BY-NC-SA 3.0.

"Unit 2 pre-class homework tutorial." by Low, R. *In* The Wicket Problem of Global Food Security, InTeGrate, CC BY-NC-SA

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9.

CLIMATE CHANGE

Andrea Bierema

Learning Objectives

Students will be able to:

- Distinguish between climate and weather.
- Explain past, present, and future climate.
- Describe examples that illustrate that life on earth depends on, is shaped by, and affects climate.
- Describe the Intergovernmental Panel on Climate Change.
- Describe examples of how humans influence the climate system.
- Describe how system climate models are developed and analyzed.

How is climate change portrayed in television comedies? Watch the following video to find out!

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Weather vs. Climate

Although these words are often used interchangeably, they have very different meanings. View the following video to learn how they differ.

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Exercise

What are your local weather and climate like? Check out *Weather Underground's* Historical Weather Data to find out! Once on the website, type in your location and then click "view." The data charts represent the current day's temperature, precipitation, and wind speed. Scroll down to the charts, which also show the day's patterns as well as the average. Change the setting toward the top of the page to "month" and scroll further down to see the daily observations. If viewing this at the beginning of the month, then change the month to the previous one.

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The Atmospheric Blanket and its Warming Effect

The gasses in the atmosphere act like a blanket, warming the earth and resulting in the natural greenhouse gas effect.

The Atmosphere

The Earth's atmosphere is an extremely thin shell compared with the size of our planet. The primary gases in the atmosphere by volume are nitrogen (78.1%), oxygen (20.9%), and argon (0.9%). These figures don't include water vapor, which varies significantly with location and altitude but averages about 0.4% of the atmosphere globally. Other naturally occurring gases include carbon dioxide (designated by chemists as CO_2), ozone, and methane, which all occur in trace amounts. Although CO_2, methane, and ozone occur naturally, human activities are increasing their concentrations. This blanket of atmosphere sustains life in many fundamental ways, as shown in the figure below.

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The Natural Greenhouse Gas Effect

On the image below, click on the "question mark" hotspots to learn about solar radiation.

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Not all of the emitted heat energy can escape to space. The greenhouse gases in the intervening atmosphere absorb (or trap) some of this heat energy. As a result, the heat energy leaving the planet is reduced by the intervening atmosphere. It is this trapping of heat energy that otherwise would have escaped to space through the atmosphere that is referred to as the ***greenhouse effect***.

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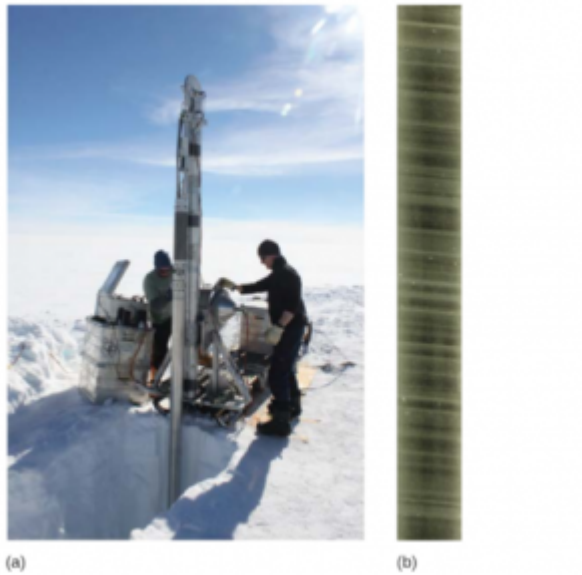
Past Climate

Because scientists cannot go back in time to directly measure climatic variables, such as average temperature and precipitation, they must instead indirectly measure temperature. To do this, scientists rely on historical evidence of Earth's past climate.

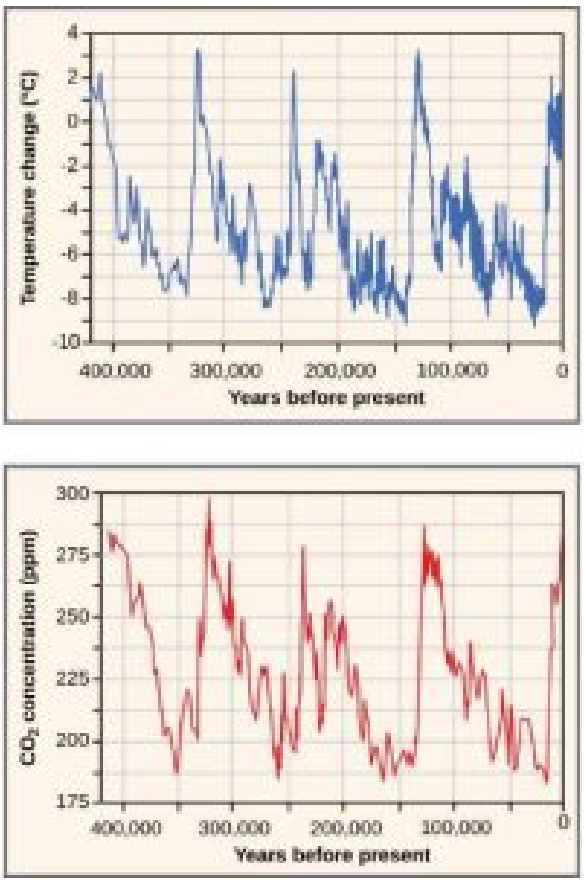
Antarctic ice cores are a key example of such evidence for climate change. These ice cores are samples of polar ice obtained by means of drills that reach thousands of meters into ice sheets or high mountain glaciers. Viewing the ice cores is like traveling backward through time; the deeper the sample, the earlier the time period. Trapped within the ice are air bubbles and other biological evidence that can reveal temperature and carbon dioxide data. Antarctic ice cores have been collected and analyzed to indirectly estimate the temperature of the Earth over the past 400,000 years.



Scientists drill for ice cores in polar regions. The ice contains air bubbles and biological substances that provide important information for researchers. (credit: a: Helle Astrid Kjær; b: National Ice Core Laboratory, USGS)

The 0°C on the graph below represents the long-term average. Temperatures that are greater than 0°C exceed Earth's long-term average temperature. Conversely, temperatures that are less than 0°C are less than Earth's average temperature. This figure shows that there have been periodic cycles of increasing and decreasing temperature.

Before the late 1800s, the Earth has been as much as 9°C cooler and about 3°C warmer. Note that the second graph below shows that the atmospheric concentration of carbon dioxide has also risen and fallen in periodic cycles. Also, note the relationship between carbon dioxide concentration and temperature. The graph shows that carbon dioxide levels in the atmosphere have historically cycled between 180 and 300 parts per million (ppm) by volume.



Ice at the Russian Vostok station in East Antarctica was laid down over the course of 420,000 years and reached a depth of over 3,000 m. By measuring the amount of CO2 trapped in the ice, scientists have determined past atmospheric CO2 concentrations. Temperatures relative to modern-day were determined from the amount of deuterium (an isotope of hydrogen) present.

The figure above does not show the last 2,000 years with enough detail to compare the changes in Earth's temperature during the last 400,000 years with the temperature change that has occurred in the more recent past. Two significant temperature anomalies, or irregularities, have occurred in the last 2,000 years. These are the Medieval Climate Anomaly (or the Medieval Warm Period) and the Little Ice Age. A third temperature anomaly aligns with the Industrial Era. The Medieval Climate Anomaly occurred between 900 and 1300 AD. During this time period, many climate scientists think that slightly warmer weather conditions prevailed in many parts of the world; the higher-than-average temperature changes varied between 0.10°C and 0.20°C above the norm. Although 0.10°C does not seem large enough to produce any noticeable change, it did free seas of ice. Because of this warming, the Vikings were able to colonize Greenland.

The Little Ice Age was a cold period that occurred between 1550 AD and 1850 AD. During this time, a slight cooling of a little less than 1°C was observed in North America, Europe, and possibly other areas of the Earth.

This 1°C change in global temperature is a seemingly small deviation in temperature (as was observed during the Medieval Climate Anomaly); however, it also resulted in noticeable climatic changes. Historical accounts reveal a time of exceptionally harsh winters with much snow and frost.

The Industrial Revolution, which began around 1750, was characterized by changes in much of human society. Advances in agriculture increased the food supply, which improved the standard of living for people in Europe and the United States. New technologies were invented that provided jobs and cheaper goods. These new technologies were powered using fossil fuels; especially coal. The Industrial Revolution starting in the early nineteenth century ushered in the beginning of the Industrial Era. When fossil fuel is burned, carbon dioxide is released. With the beginning of the Industrial Era, atmospheric carbon dioxide began to rise.

Activity

What proportion of Earth's history have humans been on Earth? Check out Learn.Genetic's Geologic Time Scale to find out! Note: Once on the website, don't give up scrolling down.

Climate Models

How We Use Models

Models help us to work through complicated problems and understand complex systems. They also allow us to test theories and solutions. From models as simple as toy cars and kitchens to complex representations such as flight simulators and virtual globes, we use models throughout our lives to explore and understand how things work.

Climate Models, and How They Work

Climate models are based on well-documented physical processes to simulate the transfer of energy and materials through the climate system. Climate models, also known as general circulation models or GCMs, use mathematical equations to characterize how energy and matter interact in different parts of the ocean, atmosphere, and land. Building and running a climate model is a complex process of identifying and quantifying Earth system processes, representing them with mathematical equations, setting variables to represent initial conditions and subsequent changes in climate forcing, and repeatedly solving the equations using powerful supercomputers.

Climate Model Resolution

Climate models separate Earth's surface into a three-dimensional grid of cells. The results of processes modeled in each cell are passed to neighboring cells to model the exchange of matter and energy over time. Grid cell size defines the resolution of the model: the smaller the size of the grid cells, the higher the level of detail in the model. More detailed models have more grid cells, so they need more computing power.

See an animation showing different grid sizes »

Explore information about supercomputer systems used to run global climate models »

Climate models also include the element of time, called a time step. Time steps can be in minutes, hours, days, or years. Like grid cell size, the smaller the time step, the more detailed the results will be. However, this higher temporal resolution requires additional computing power.

How are Climate Models Tested?

Once a climate model is set up, it can be tested via a process known as "hind-casting." This process runs the model from the present time back into the past. The model results are then compared with observed climate and weather conditions to see how well they match. This testing allows scientists to check the accuracy of the models and, if needed, revise their equations. Science teams around the world test and compare their model outputs to observations and results from other models.

Using Scenarios to Predict Future Climate

Once a climate model can perform well in hind-casting tests, its results for simulating future climate are also assumed to be valid. To project climate into the future, the climate forcing is set to change according to a possible future scenario. Scenarios are possible stories about how quickly the human population will grow, how land will be used, how economies will evolve, and the atmospheric conditions (and therefore, climate forcing) that would result in each storyline.

In 2000, the Intergovernmental Panel on Climate Change (IPCC) issued its Special Report on Emissions Scenarios (SRES), describing four scenario families to describe a range of possible future conditions. Referred to by letter-number combinations such as A1, A2, B1, and B2, each scenario was based on a complex relationship between the socioeconomic forces driving greenhouse gas and aerosol emissions and the levels to which those emissions would climb during the 21st century. The SRES scenarios have been in use for more than a decade, so many climate model results describe their inputs using the letter-number combinations.

In 2013, climate scientists agreed upon a new set of scenarios that focused on the level of greenhouse gases in the atmosphere in 2100. Collectively, these scenarios are known as Representative Concentration Pathways or RCPs. Each RCP indicates the amount of climate forcing, expressed in Watts per square meter, that would result from greenhouse gases in the atmosphere in 2100. The rate and trajectory of the forcing is the pathway. Like their predecessors, these values are used in setting up climate models.

Learn more about RCPs »

Results of Current Climate Models

Around the world, different teams of scientists have built and run models to project future climate conditions under various scenarios for the next century. The model results project that global temperature will continue to increase, but show that human decisions and behavior we choose today will determine how dramatically climate will change in the future.

How are Climate Models Different From Weather Prediction Models?

Unlike weather forecasts, which describe a detailed picture of the expected daily sequence of conditions starting from the present, climate models are probabilistic, indicating areas with higher chances to be warmer or cooler and wetter or drier than usual. Climate models are based on global patterns in the ocean and atmosphere, and records of the types of weather that occurred under similar patterns in the past.

View maps showing short-term climate forecasts »

IPCC (Intergovernmental Panel on Climate Change)

The Intergovernmental Panel on Climate Change (IPCC) is the most prominent international scientific body for assessing climate change. It was formed in 1988 by the World Meteorological Organization (WMO) and the United Nations Environment Programme (UNEP). There are currently 195 member-countries in the IPCC, and membership is open to all countries in the WMO and UN. The IPCC is responsible for reviewing and evaluating scientific, technical, and socioeconomic information related to climate change. While the IPCC neither conducts research nor monitors any climate change data directly, it provides policymakers with the most comprehensive picture of the scientific consensus.

Who are the people in this panel?

An interactive or media element has been excluded from this version of the text. You can view it online here:
https://openbooks.lib.msu.edu/isb202/?p=155

To learn more about some of the scientists involved, check out the IPCC's playlist "25 Years of the IPCC- Individual Videos."

IPCC's Climate Assessment Report

The IPCC is currently working on its 6th assessment report, which is scheduled to be released in 2022. The following video is a summary of the 5th assessment report.

An interactive or media element has been excluded from this version of the text. You can view it online here:
https://openbooks.lib.msu.edu/isb202/?p=155

If you would like to learn more about these reports, then view the videos for each working group:

- Working Group 1: Physical Science Basis
  - View in English, or
  - Choose another language from this playlist
- Working Group 2: Impacts, Adaptation, and Vulnerability
  - View in English, or
  - Choose another language from this playlist
- Working Group 3: Mitigation of Climate Change
  - View in English, or
  - Choose another language from this playlist

View the synthesis report.

So what do the models look like? The following are links to the IPCC 5th Assessment Report, "Summary for Policymakers" figures per working group.

Working Group 1: Physical Science Basis

Working Group 1 investigates past, present, and future climate.

- Figure SPM.1 Temperature: a) The observed annual and decadal average temperature from 1850 to 2012,

compared to the average for 1961 to 1990. Negative values indicate average temperatures were below the 1961-1990 average and positive values indicate above that average. b) World map of the observed change in surface temperature from 1901 to 2012.

- Figure SPM.2 Precipitation: World maps illustrating the change in precipitation from 1901 to 2010 and 1951 to 2010.
- Figure SPM.3 Cryosphere (snow and ice): For the 20th century, a) the amount of spring snow cover in the Northern Hemisphere, b) the extent of Arctic summer sea ice, c) change in global average upper ocean heat content relative to 1970, and d) change in global average sea level relative to 1900-1905.
- Figure SPM.4 Carbon Dioxide: a) Atmospheric concentration of carbon dioxide and b) surface ocean carbon dioxide and pH (the lower the number, the more acidic).
- Figure SPM.5 Drivers of Climate Change: Anthropogenic and natural climate change drivers and their impacts in 2011 compared to 1750.
- Figure SPM.6 Natural and Anthropogenic Impact: Global and regional graphs illustrating the impact of natural forcings and combined natural and anthropogenic forcings.
- Figure SPM.7 Predicted Temperature: Predicted temperature changes relative to 1986-2005 using four models (each model represents a different scenario (from cutting all carbon emissions today to not changing carbon emissions at all).
- Figure SPM.8 Predicted Climate: World maps showing predicted changes in climate for 2081-2100 relative to 1986-2005 for two different scenarios for a) temperature, b) precipitation, c) Northern Hemisphere September sea ice extent, and d) ocean surface pH (lower values mean more acidic).
- Figure SMP.9 Predicted Sea Level Rise: Graph depicting predicted sea-level rise using four models (each model represents a different scenario (from cutting all carbon emissions today to not changing carbon emissions at all).
- Figure SMP.10 Carbon Dioxide and Temperature: Global surface temperature correlated with temperature change relative to 1861-1880.
- See the WG1 SMP figures in context with full captions.

Working Group 2: Impacts, Adaptation, and Vulnerability

Working Group 2 investigates how changes in climate impact the environment and society.

- Figure SPM.2 Widespread Impacts: A world map illustrating how different regions are impacted by climate change.
- Figure SPM.5 Species Migration: The maximum speeds that species can move across landscapes compared to how quickly temperatures are expected to change.
- Figure SPM.6 Fisheries: Predicted climate change impact on a) catch potential and b) change in pH and which taxa are most impacted.
- Figure SPM.7 Crop Yield: Predicted climate change impact on future crop yields.
- Figure SPM.8 Solution Model: A model illustrating risks and mitigations.
- See the WG2 SMP figures in context with full captions.

Working Group 3: Mitigation of Climate Change

Working Group 3 investigates how we may reduce future climate change and the impact of its effects.

- Figure SPM.1 Greenhouse Gases: Total annual greenhouse gases by type: fossil fuel and industrial processes, forestry and other land use (FOLU), nitrous oxide (N_2O), and fluorinated gases covered under the Kyoto Protocol (F-gases).
- Figure SPM.2 Economic Sectors: Direct and indirect carbon dioxide emissions by economic sector.
- Figure SPM.4 Baseline and Mitigation Scenarios: Global greenhouse gas emissions for four scenarios: baseline (not changing our emissions path), two scenarios with varying emission levels, and one scenario for if we stopped carbon emissions.
- Figure SPM.9 Investment: Changes in annual investment flows projected for 2010-2029 based on keeping carbon dioxide emissions in a range of about 430 to 530 parts per million.
- See the WG3 SMP figures in context with full captions.

Attributions

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"Climate Change" by Ramanathan, V., *Bending the Curve: Climate Change Solutions Digital Textbook*, 2019, CC-BY-NC-SA.

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10.

SPECIES INTERACTIONS

Andrea Bierema

Learning Objectives

Students will be able to:

- Define niche.
- Describe types of species interactions.
- Define competitive exclusion and resource partitioning principles.
- Use food webs to infer examples of species interaction within a community.
- Use ecological models to appropriately predict how an abundance of species may impact other species within a community.

Community Ecology

Ecology is studied at different scales and species interactions are part of the "community ecology" scale.

An interactive or media element has been excluded from this version of the text. You can view it online here:
https://openbooks.lib.msu.edu/isb202/?p=199

Two-Species Interactions

Community ecology includes the ways in which species interact. Research sometimes focuses on two species of a complex community and the general ways those species interact with one another can be classified by whether the species are positively, negatively, or neutrally impacted. The following video describes the main types of species interactions, with examples. The table below summarizes these types.

An interactive or media element has been excluded from this version of the text. You can view it online here:
https://openbooks.lib.msu.edu/isb202/?p=199

Interaction	Definition	Species A	Species B
Mutualism*	A long-term, close association between two species in which both partners benefit	+	+
Commensalism*	Organisms of two species use the same limited resource and have a negative impact on each other	+	N
Consumption · Parasitism* · Parasitoidism* · Predation · Herbivory	One species eats all or part of the body of a member of another species. The consumed species is either killed from the interaction (e.g., predation) or harmed but likely survives (e.g., parasitism)	+	–
Competition	Two species "fight" over a resource. Although one species may "win," it is still negatively impacted by taking part in the competition	–	–

*These interactions may also be labeled as symbiotic interactions (i.e., symbiosis) if the two species live and interact closely together.

Exercise

Test your understanding of species interactions!

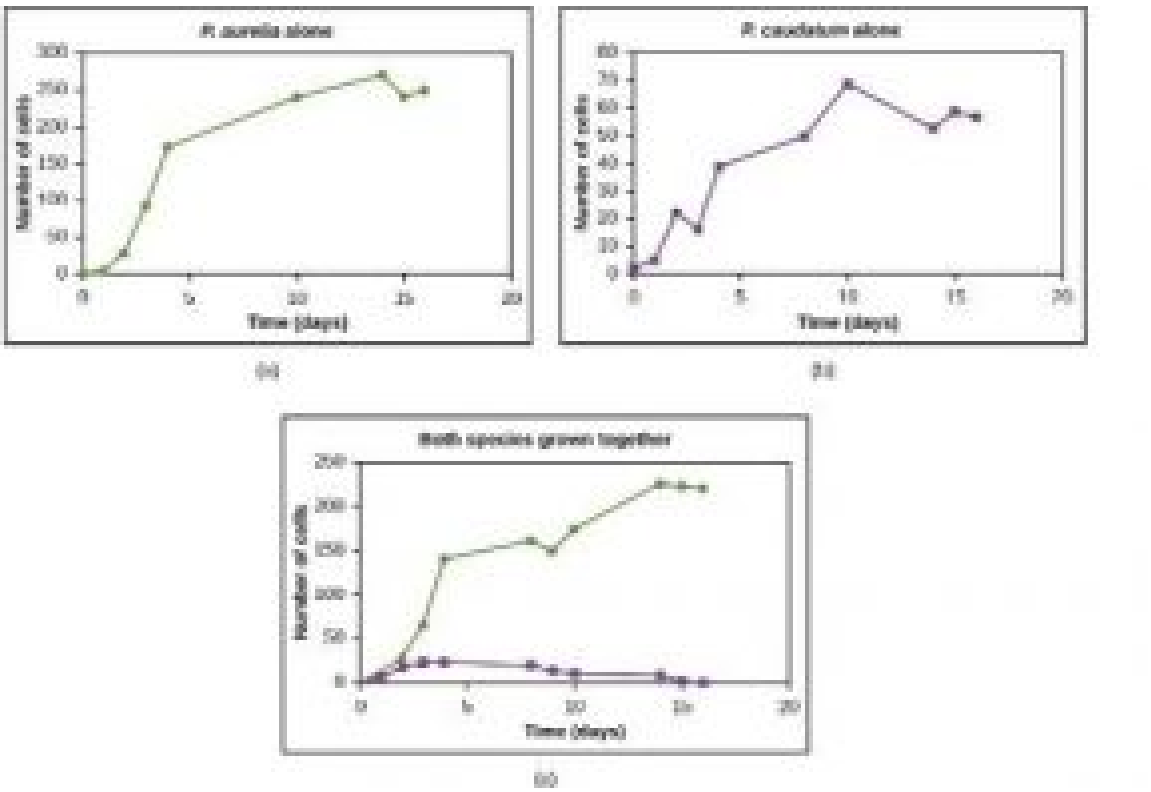
An interactive or media element has been excluded from this version of the text. You can view it online here:
https://openbooks.lib.msu.edu/isb202/?p=199

Learn more about "aquatic cleaning stations"!

Niches and How They Influence Competition

Resources are often limited within a habitat and multiple species may compete to obtain them. All species have an ***ecological niche*** in the ecosystem, which describes how they acquire the resources they need and how they interact with other species in the community. The ***competitive exclusion principle*** states that two species cannot occupy the same niche in a habitat. In other words, different species cannot coexist in a community if they are competing for all the same resources.

An example of this principle is shown below with two protozoan species: *Paramecium aurelia* and *Paramecium caudatum*. When grown individually in the laboratory, they both thrive. When they are placed together in the same test tube (habitat), *P. aurelia* outcompetes *P. caudatum* for food, leading to the latter's eventual extinction.



Paramecium aurelia (graph a) and *Paramecium caudatum* (graph b) grow well individually, but when they compete for the same resources, the *P. aurelia* outcompetes the *P. caudatum*. In graph c, the top growth curve is *P. aurelia* and the bottom growth curve is *P. caudatum*.

Exercise

Does competition between two species that share a similar niche always result in one dying off? Try out the Virtual Biology Lab's simulation of barnacle competition. Once on the website, read through the background information and the tutorial. Then run the experiment- feel free to keep the variables consistent or see what happens when you change them.

An interactive or media element has been excluded from this version of the text. You can view it online here:
https://openbooks.lib.msu.edu/isb202/?p=199

Multi-Species Species Interactions

This section refers to food webs. Please see Khan Academy's Food Chains and Food Webs for a review.

In community ecology, we can examine the interaction of two species or we can think at a larger, more complex scale and examine how many species interact. One way to do so is to start with a food web to identify some of the interactions that are occurring within a community. The following video describes this in more detail using the example of a coastal food web.

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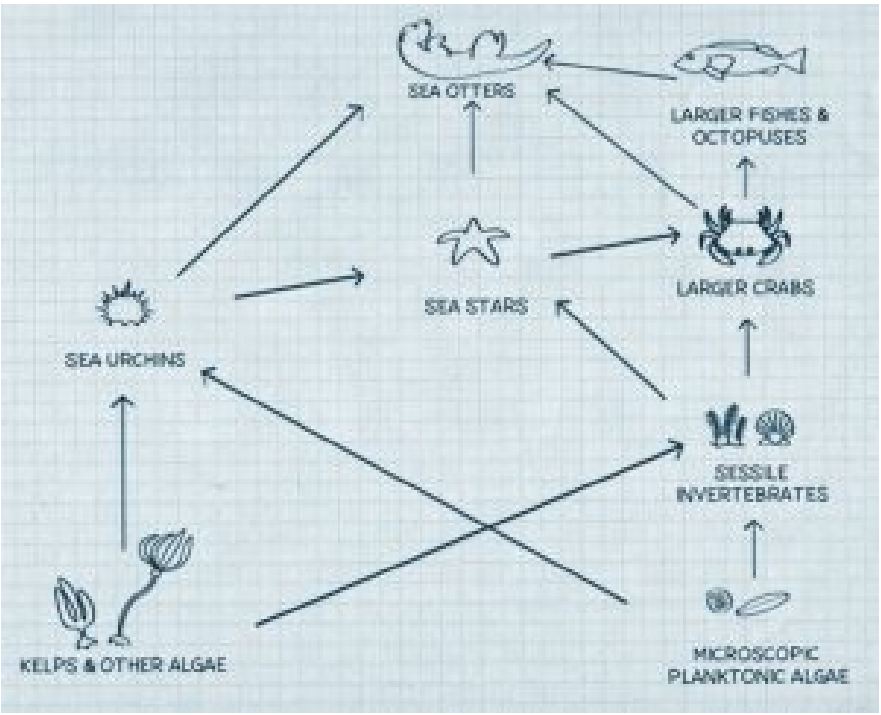
https://openbooks.lib.msu.edu/isb202/?p=199



This is the food web shown in the video above by Khan Academy.

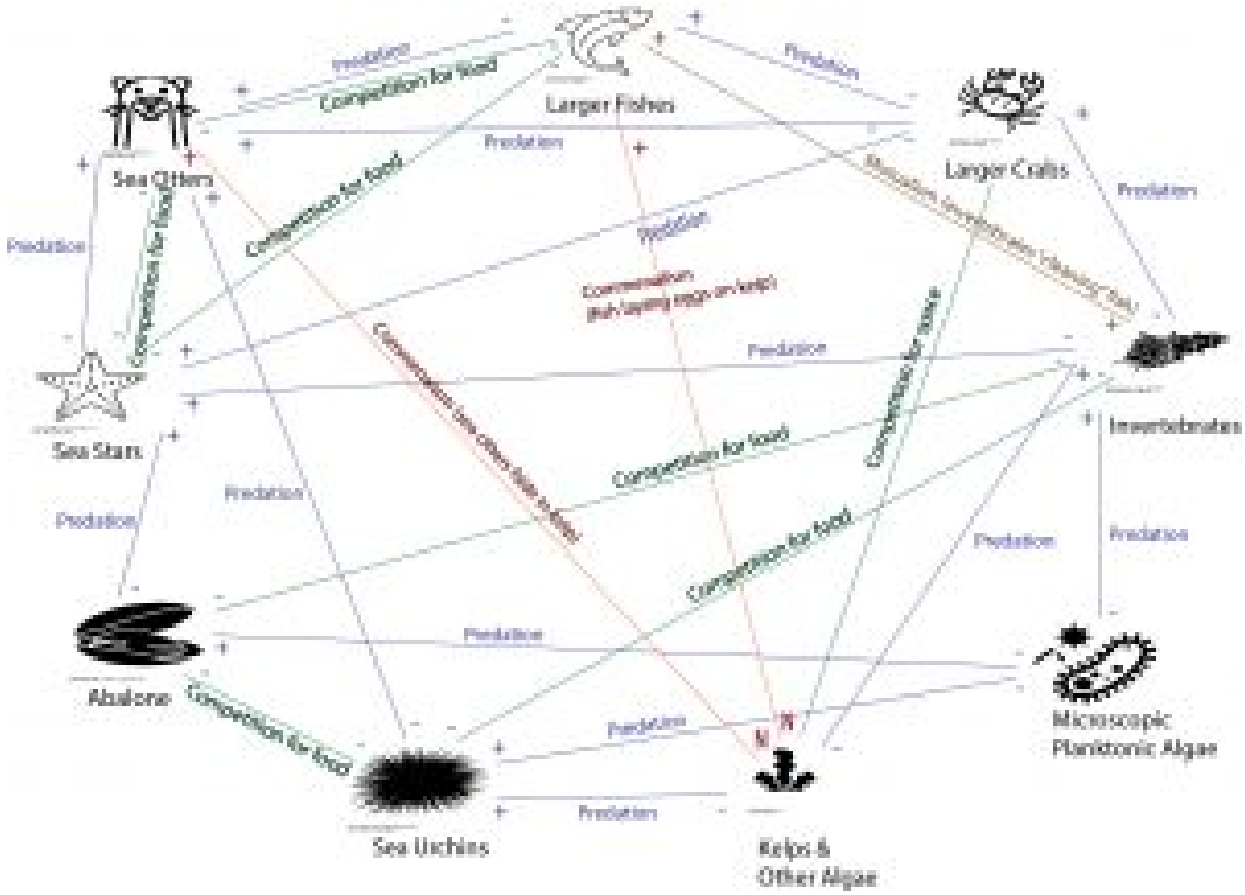
An interactive or media element has been excluded from this version of the text. You can view it online here:
https://openbooks.lib.msu.edu/isb202/?p=199

Species Interaction Model

The food web is a model that illustrates how energy moves. Models can also be created to show how species interact with one another (beyond predator/prey and competition for food). The following model was introduced in the video. Rather than having each line represent energy movement, they can be labeled with different species interactions such as mutualism. This is a visual way to explain the complexity of species interactions within a community.

For instance, the following model is a complete species interaction model using the species from the video's Pacific coast food web. Notice how species are now connected with lines rather than arrows. Interactions influence both species involved, while an arrow in a food web indicates the direction that energy moves. Each end of a line is also labeled with a positive (+), negative (-), and neutral (N), indicating how the species is impacted by the interaction. These symbols align with the symbols used in the interactions table shown toward the beginning of the chapter. For instance, in looking at the lines/interactions labeled as "competition for food," both species are labeled as being negatively impacted by placing a negative on each side of the line.

The "predation" and "competition for food" interactions align with the food web. Food webs allude to predation interactions and if species are eating the same species, then they are likely competing for the food source. The rest of the interactions labeled are from outside research, rather than from studying the food web.



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11.

POPULATION GROWTH

Andrea Bierema

Learning Objectives

Students will be able to:

- Define population.
- Describe common population growth models.
- Define carrying capacity.
- Provide accurate explanations of population growth graphs, including carrying capacity.
- Define and identify examples of density-independent and density-dependent factors.
- Explain how population growth data plays a role in conservation.
- Draw connections between human population growth and social/political and economic conditions.

Population Ecology

Ecology is studied at different scales, and population growth is part of the "population ecology" scale.

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Graph Interpretation

Exercise: Graph Interpretation

Population growth, a common topic in population ecology, plots population size over time on a graph. Before learning about population growth, let's test your graph reading skills.

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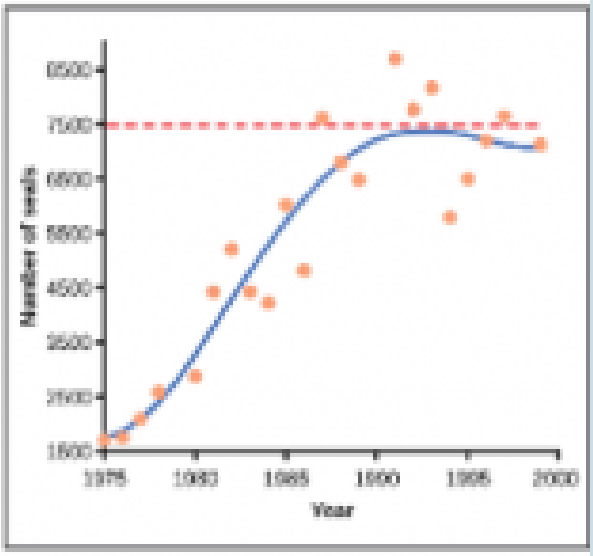
Exercise: Population Growth Graph

Now, let's read a population growth graph:



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Population Growth

There are two main models used to describe how population size changes over time: exponential growth and logistic growth. Click on the information hotspots (labeled as "i") in the figure below to learn more.

Exercise

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Although several populations in nature follow logistic and exponential growth patterns, population growth can be much more complicated. For instance, many insects undergo brief exponential growth, followed by periods of mass death (or drop in population).

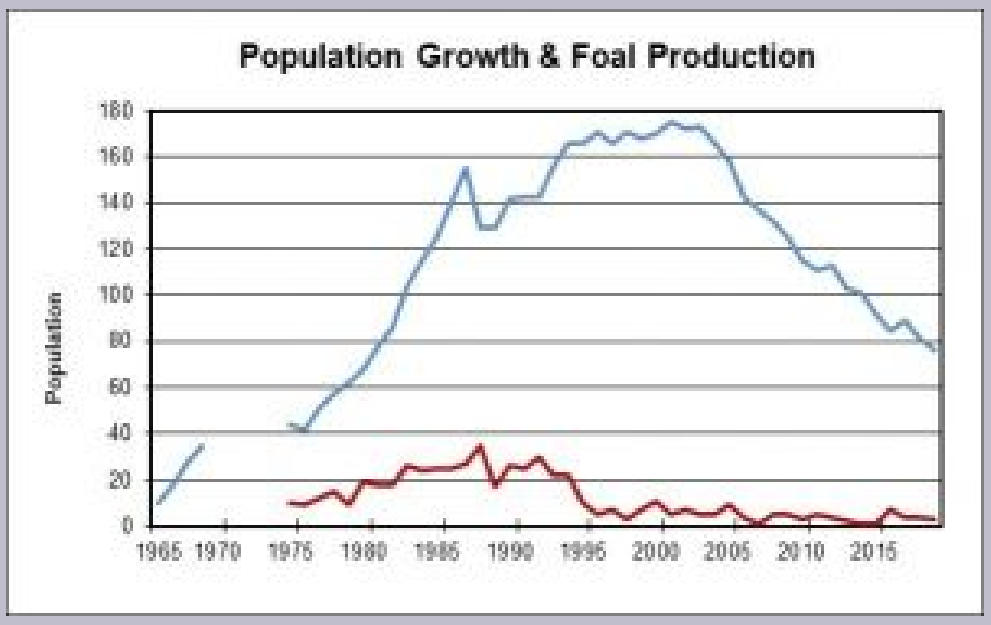
Learn more about population growth modeling through this simulation!

Example

Population growth may not look like either exponential or logistic growth models. For instance, the horse population on Assateague Island in Maryland (pictured below) experienced increase and stabilization—similar to logistic growth—but then underwent a stable decline.

To complicate the interpretation of this graph further, the stabilization and decline were due to human management of the population. Studies showed that the environment would not sustain such a large population over time, and many of the plant-species' population growth would drop dramatically—eventually causing a crash in the horse's population size. Managers injected female horses with a vaccine that caused immune cells to attack sperm cells.

Mare and foal grazing on Assateague Island.



Horse population growth and foal production on Assateague Island National Seashore, 1965-2015. Blue line – horse population, red line – foal births, blank space – unavailable data.

Check out the following article to learn more!

National Park Service. (n.d.). Resource management brief – horses. Retrieved June 24, 2020 from https://www.nps.gov/asis/learn/nature/resource-management-brief-horses.htm

Practice

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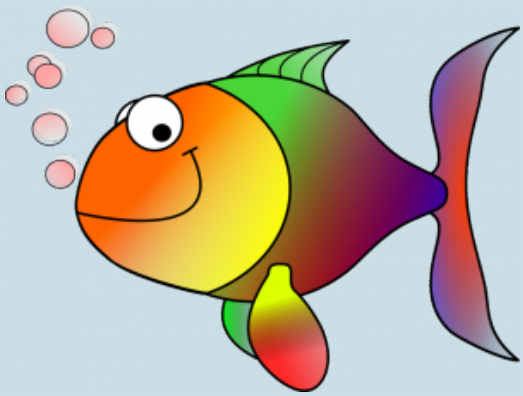

Can you identify the carrying capacity in these populations of the fictitious smiling rainbow fish?

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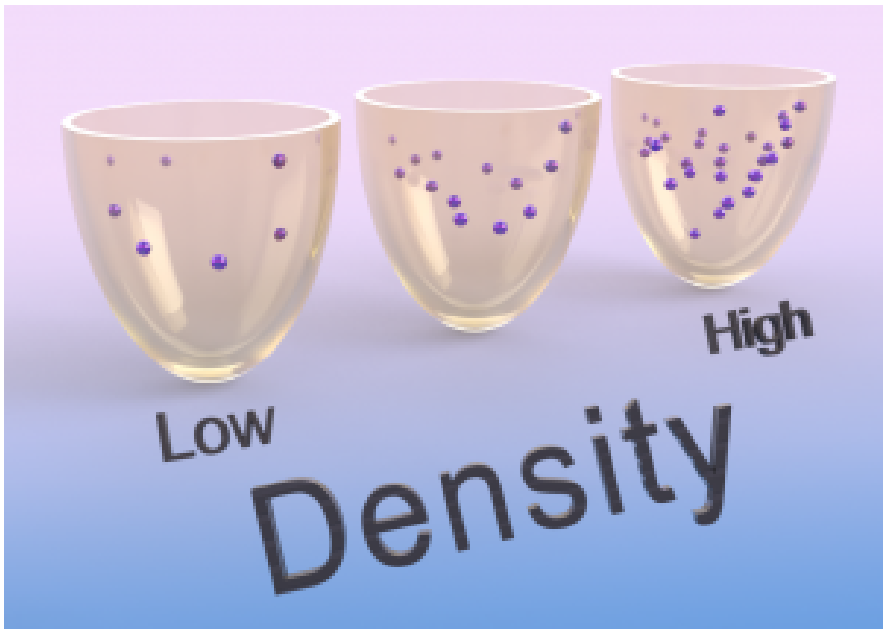
Population Density

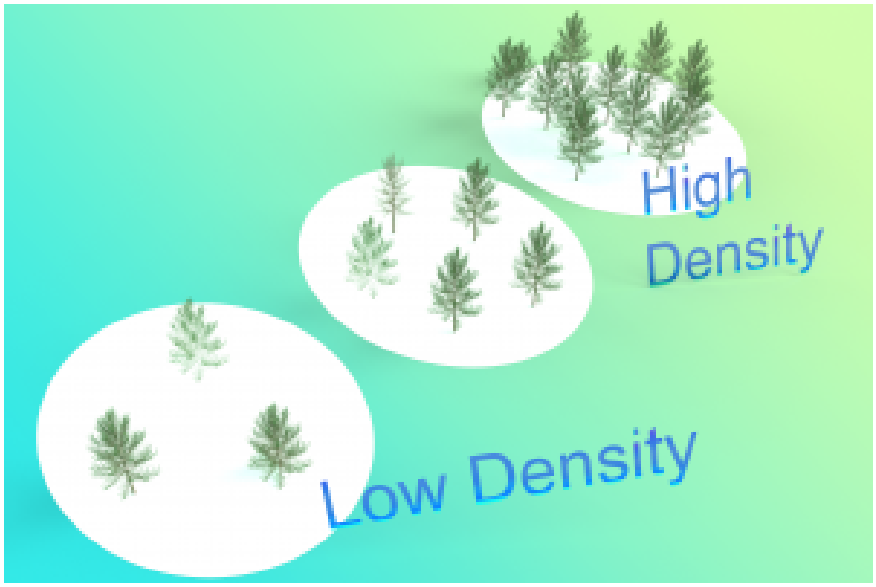
Although examining how the size of the population changes over time is informative, it neglects to take into account how much space the population is occupying. How dense a population is can impact survival and be influenced by a number of factors.

Density

You may have heard of density in a chemistry or physics class before. In those cases, density typically refers to how dense an object is (as seen in the figure below). This is the amount of matter in a given volume (calculated by mass/volume).



The diagram illustrates that higher density means that the particles are closer together.



Density is also used to explain populations. In this image, the density of pine trees in each "environment" varies from low density to high density.

Population density is similar to object density, except it refers to the number of individuals (or organisms) instead of the number of particles.

Population Growth Factors

Factors that influence population size, such as a drought or drop in prey, are categorized into one of two types:

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Based on those definitions, identify each factor as density-dependent or independent.

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Population Growth and Conservation

Population growth is used to inform conservation—whether it be to determine if conservation efforts are needed (i.e., a population is dropping in size) or testing to see if a conservation strategy is working.

Exercises

View the following interactive video to learn about lion conservation in Gorogonsa National Park and practice some of the population growth concepts!

An interactive or media element has been excluded from this version of the text. You can view it online here:
https://openbooks.lib.msu.edu/isb202/?p=157

Need more practice and information? Go through the *Connecting Concepts* interactive lesson on population dynamics, which describes population growth trends of zebra mussels and elephants!

Human Population

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The previous video explored why our population is increasing in size, but when did our population start increasing and what does it look like now? The following video does an excellent job of showing how our rate of increase has changed over time. Note: It's going to seem like not a lot is going for a while, but keep watching!

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More Information

How do your personal habitats and way of life impact the planet? How many Earths would we need if everyone lived a similar way as you? Find out by taking this ecological footprint quiz!

Next, find out your water footprint. You'll be surprised by how many things that we do use water!

Reference

National Park Service. (n.d.). Resource management brief – horses. Retrieved June 24, 2020 from https://www.nps.gov/asis/learn/nature/resource-management-brief-horses.htm

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12.

EVOLUTION

Andrea Bierema

Learning Objectives

Students will be able to:

- Define evolution.
- Describe different mechanisms of evolution, including natural selection, genetic drift, gene flow, and mutation.
- Recognize that new heritable traits result from random mutations.
- Use data and physical traits to predict which evolutionary mechanisms and potential selective agents influence a population.
- Distinguish between microevolution and macroevolution.
- Describe possible mechanisms of speciation.

Definition of Evolution

Biological evolution, simply put, is ***descent with modification***.

Biological evolution is not simply a matter of change over time. Lots of things change over time: trees lose their leaves and mountain ranges rise and erode, but they aren't examples of biological evolution because they don't involve descent through genetic inheritance.

The central idea of biological evolution is that ***all life on Earth shares a common ancestor***, just as you and your cousins share a common grandmother.

Through the process of descent with modification, the common ancestor of life on Earth gave rise to the fantastic diversity that we see documented in the fossil record and around us today. Evolution means that we're all distant cousins: humans and oak trees; hummingbirds and whales.

Evolution occurs at different scales:

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https://openbooks.lib.msu.edu/isb202/?p=211

Mechanisms of Evolution

Although the term "evolution" is often used synonymously with "natural selection," they are actually referring to different concepts. Evolution is an observable phenomenon in which gene frequencies change over time, but it does not explain *why* a population is undergoing evolution. This is where natural selection—and other mechanisms—come into play. They explain "the why."

There are four main mechanisms of evolution:

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https://openbooks.lib.msu.edu/isb202/?p=211

Exercise

Let's look at some examples of evolution and see if we can identify which evolutionary mechanism is at play.

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https://openbooks.lib.msu.edu/isb202/?p=211

Once you complete that, here is another game. This is a matching game using the images from the above quiz. As you match two images, recall which type of mechanism it is and why!

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https://openbooks.lib.msu.edu/isb202/?p=211

All of these mechanisms can cause changes in the frequencies of genes in populations, so all of them are mechanisms of evolutionary change. However, natural selection and genetic drift cannot operate unless there is genetic variation—that is, unless some individuals are genetically different from others. If the population of beetles was 100% green, selection and drift would not have any effect because their genetic make-up could not change.

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Important Notes Regarding Mutations

Mutations are random

Mutations can be beneficial, neutral, or harmful for the organism, but mutations do not "try" to supply what the organism "needs." In this respect, mutations are random—whether a particular mutation happens or not is unrelated to how useful that mutation would be.

Not all mutations matter to evolution

Because all cells in our body contain DNA, there are lots of places for mutations to occur; however, not all mutations matter for evolution. Mutations that occur in non-reproductive cells won't be passed onto offspring.

For instance, if a skin cell has a mutation that causes uncontrollable cell division (i.e., cancer), that mutation is not passed to the next generation.

See the "Protein Structure and Function" chapter in this textbook for more information.

Evolution Examples

Learn more about the mechanisms of evolution via the following examples.

Exercises

Learn more about evolution through this example:

An interactive or media element has been excluded from this version of the text. You can view it online here:
https://openbooks.lib.msu.edu/isb202/?p=211

To continue learning how predation impacts the natural selection of traits, check out *Connected Bio's* simulation on the evolution of fur color in deer mice and *PhET*'s simulation (below) on the natural selection of rabbits!

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Exercise

Check out Virtual Biology Lab's Fish Pond simulation. In this simulation, change the migration rate (to test genetic flow), mutation rate, or genotype relative fitness (to test natural selection). You can also change the initial population size.

How does having a small population size affect genotype frequency, by chance (i.e., genetic drift)?

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Macroevolution

Before examining speciation, let's first learn about what it means to be a species.

Species Concept

A species is often defined as a group of individuals that actually or potentially interbreed in nature. In this sense, a species is the biggest gene pool possible under natural conditions.

That definition of a species might seem cut and dried, but it is not—in nature, there are lots of places where it is difficult to apply this definition. For example, many bacteria reproduce mainly asexually through binary fission. The definition of a species as a group of interbreeding individuals cannot be easily applied to organisms that reproduce only or mainly asexually.

Also, many plants and some animals form hybrids in nature. Hooded crows and carrion crows look different, and largely mate within their own groups, but in some areas they hybridize. Should they be considered the same species or separate species?

Hooded crow

Carrion crow

There are lots of other places where the boundary of a species is blurred. It's not so surprising that these blurry places exist—after all, the idea of a species is something that we humans invented for our own convenience!

Speciation

Speciation is a lineage-splitting event that produces two or more separate species. Let's learn about speciation in the birds of paradise in the following interactive video.

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Exercise

Go through additional case studies on evolution, including speciation in *Connecting Concepts*' Natural Selection interactive lesson and Speciation interactive lesson.

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- Carrion crow © Bob is licensed under a CC BY (Attribution) license

13.

PHYLOGENETIC TREES: MODELING EVOLUTION

Andrea Bierema

Learning Objectives

Students will be able to:

- Provide accurate explanations of a phylogenetic tree, which is a scientific model that explains how species are evolutionarily related to each other.
- Convert relevant information into a phylogenetic tree.
- Identify common ancestors on a phylogenetic tree.
- Label shared derived characteristics on a phylogenetic tree.

The History of Life: Looking at the Patterns

The central ideas of evolution are that life has a history—it has changed over time—and that different species share common ancestors.

Here, you can explore how evolutionary change and evolutionary relationships are represented in "family trees," how these trees are constructed, and how this knowledge affects biological classification. You will also find a timeline of evolutionary history and information on some specific events in the history of life: human evolution and the origin of life.

The Family Tree

The process of evolution produces a pattern of relationships between species. As lineages evolve and split and modifications are inherited, their evolutionary paths diverge. This produces a branching pattern of evolutionary relationships.

By studying inherited species' characteristics and other historical evidence, we can reconstruct evolutionary relationships and represent them on a "family tree," called a phylogeny. The phylogeny you see below represents the basic relationships that tie all life on Earth together.

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The Three Domains

This tree, like all phylogenetic trees, is a hypothesis about the relationships among organisms. It illustrates the idea that all of life is related and can be divided into three major clades, often referred to as the three domains: Archaea, Bacteria, and Eukaryota. We can zoom in on particular branches of the tree to explore the phylogeny of particular lineages, such as Animalia (outlined in red). Then we can zoom in even further to examine some of the major lineages within Vertebrata. Just click the button in the center of the image below.

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The tree is supported by many lines of evidence, but it is probably not flawless. Scientists constantly reevaluate hypotheses and compare them to new evidence. As scientists gather even more data, they may revise these particular hypotheses, rearranging some of the branches on the tree. For example, evidence discovered in the last 50 years suggests that birds are dinosaurs, which required adjustment to several "vertebrate twigs."

Understanding Phylogenies

The following diagram describes the different components of phylogenetic trees. Click on the Information tab in each box to learn more!

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For additional help on understanding phylogenies, see Learn.Genetic's "Tree Diagrams" video and "Tree Diagrams" interactive.

Exercise

Let's see what you learned!

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You can also go back to the HHMI Biointeractive's Lizard Evolution lab. Once you launch the interactive, click on the module tab to the left, and select "Module 2: Phylogeny."

Want even more practice? Try Nova Lab's The Evolution Lab!

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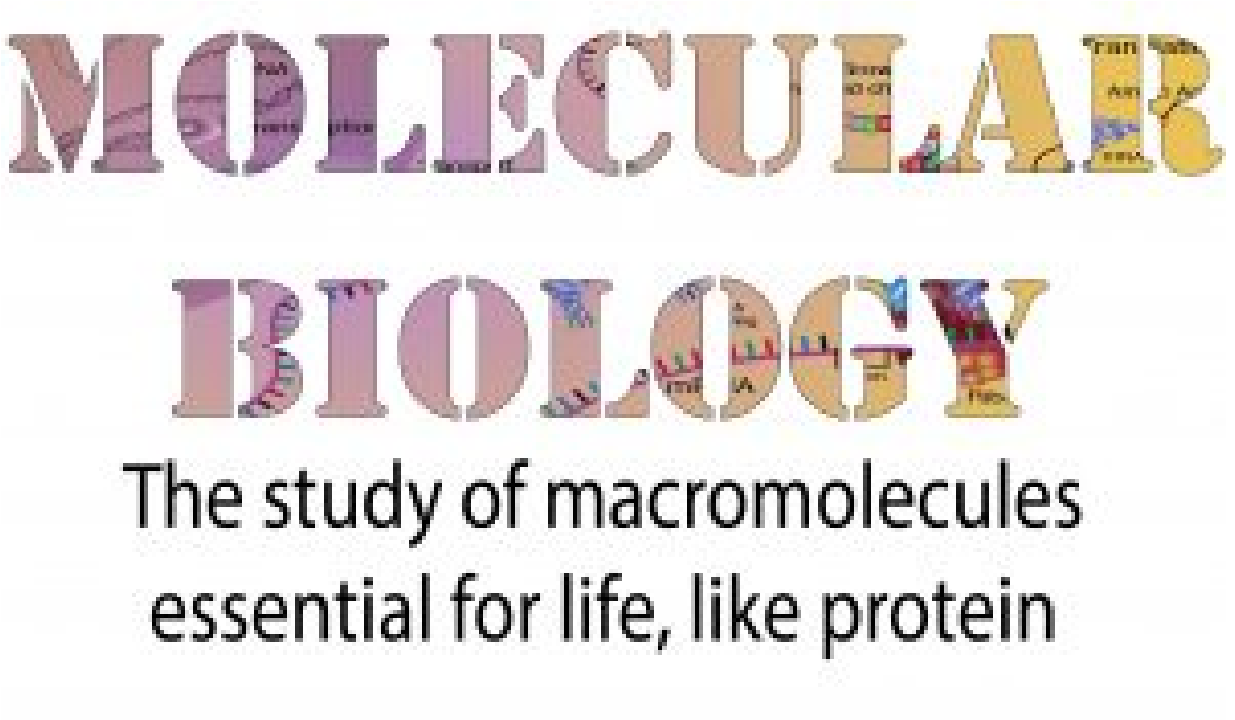
UNIT III

MOLECULAR BIOLOGY

14.

INTRODUCTION TO MOLECULAR BIOLOGY

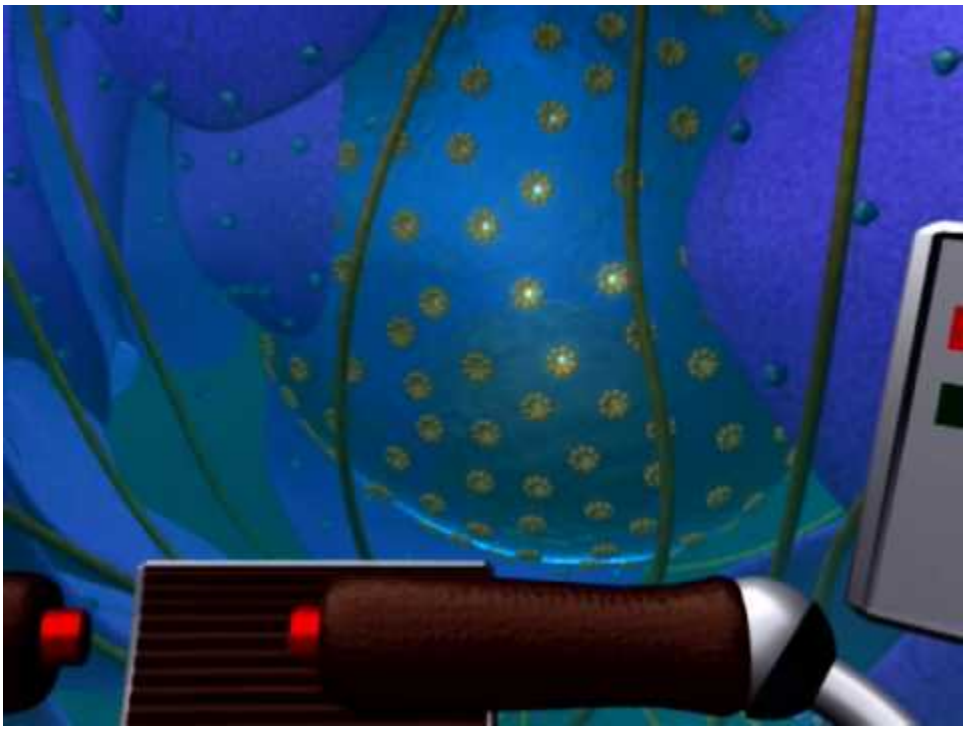
Andrea Bierema



This unit addresses how DNA is read to create protein and that proteins determine an organism's characteristics. It ends with a discussion on how we as humans can modify and edit DNA—which then affects protein synthesis—via genetic engineering.

The Cell

Review the main structures of a cell before continuing on to the next chapters. The following video gives a tour of the cell:



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Additional Information

To take a closer look into the organelles of a cell, check out Genome Unlocking Life's Code animations!

Molecular Biology

We cover a span of topics in this unit. To find out some of the basics of molecular biology and how they relate to one's life, watch the following video. It's an excellent animation!

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15.

PROTEIN STRUCTURE AND FUNCTION

Andrea Bierema

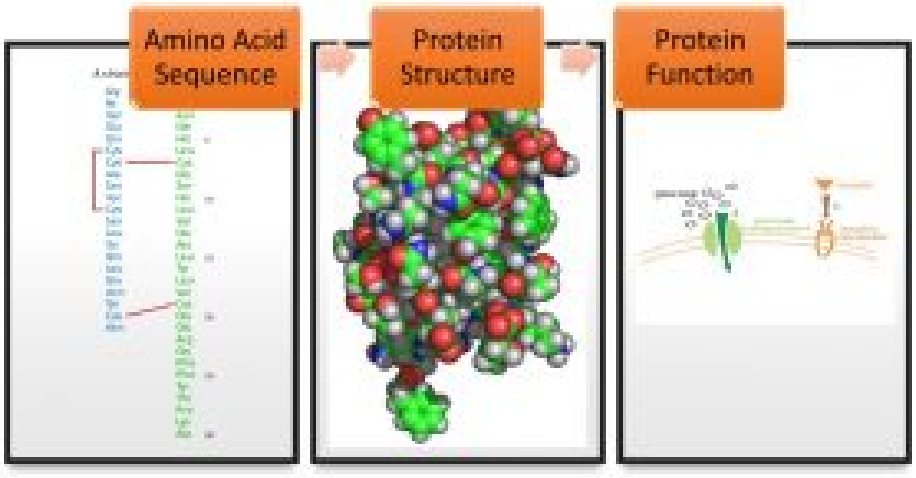
Learning Objectives

Students will be able to:

- Explain the relationship between amino acids and proteins.
- Identify examples of proteins.
- Recognize that molecular structure determines molecular interactions and relates to the cellular functions of proteins.
- Describe how protein structure influences its function.
- Describe the relationship between mutation and evolution.

Overview

This chapter is titled “protein structure and function” because protein structure heavily influences its function. The structure of a protein is caused by the chemical properties of its amino acids.



Insulin, a protein, has a specific amino acid sequence, which then causes a specific structure. This structure influences its function, which is to aid glucose in entering cells.

Proteins

Proteins are one of the most abundant organic molecules in living systems and have the most diverse range of functions of all macromolecules. Proteins may be structural, regulatory, contractile, or protective. They may serve in transport, storage, or membranes; or they may be toxins or enzymes. Each cell in a living system may contain thousands of proteins, each with a unique function. Their structures, like their functions, vary greatly. They are all, however, amino acid polymers arranged in a linear sequence (also referred to as a "peptide").

Protein types and functions:

Genetic Diseases

Learn more about protein function by checking out Learn.Genetic's "Examples of Single Gene Disorders", which describes how

proteins are involved in various gene disorders.

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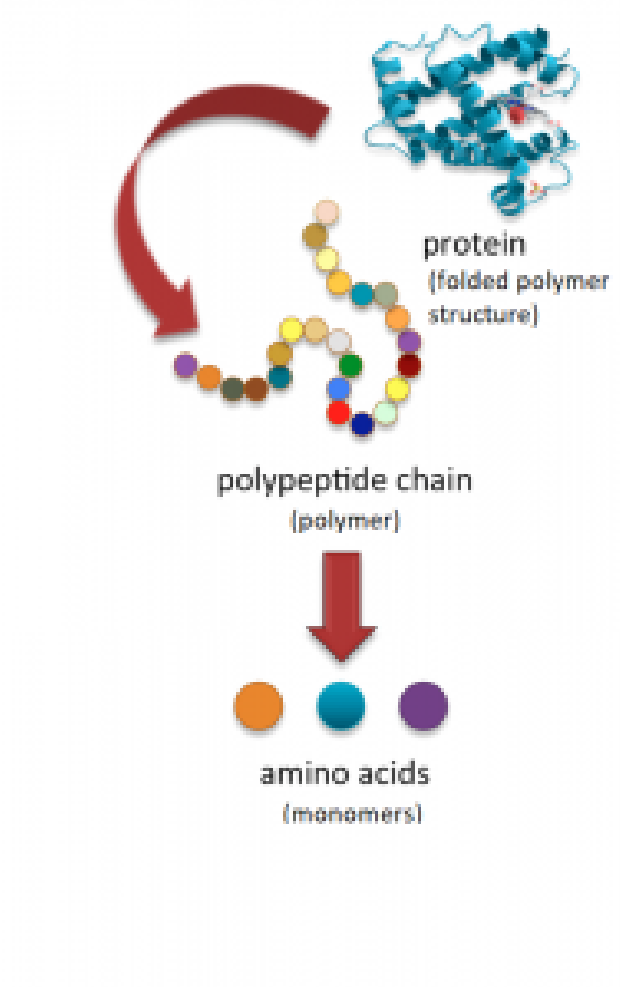
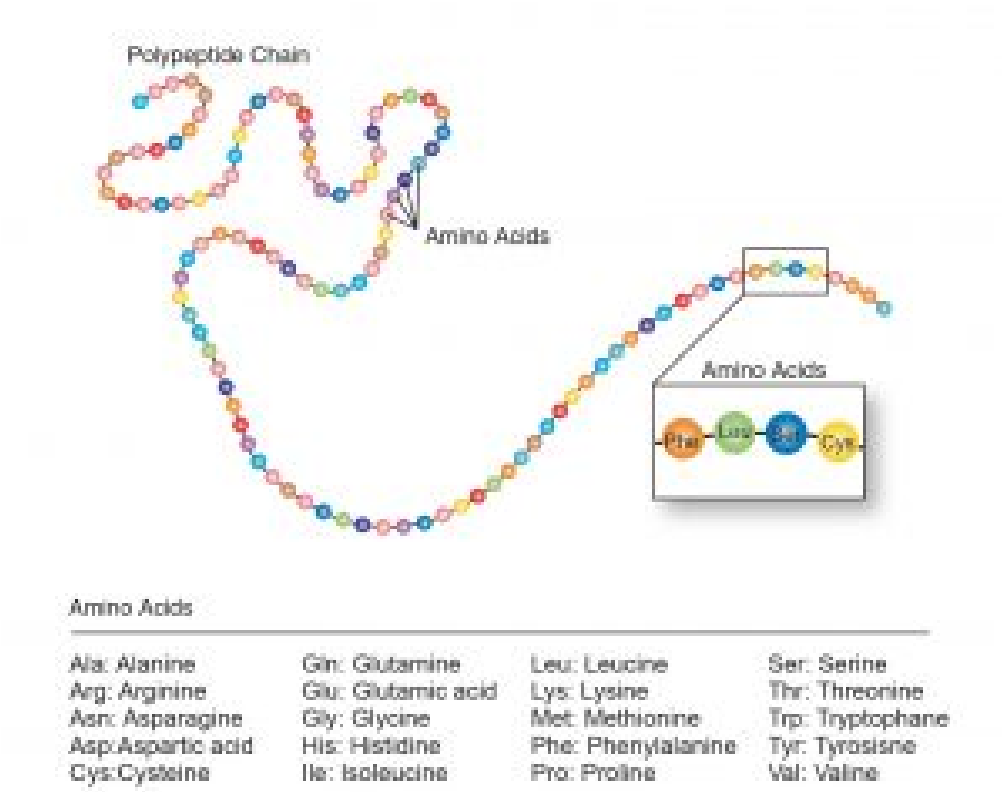
Monomers and Polymers

Monomers are molecules that can bind into long chains—these long chains are called "polymers." In other words, a polymer ("poly" = many) are made of monomers ("mono" meaning "one").

Amino acids are the monomers that comprise polypeptides (polypeptides being the polymers). A polypeptide folds into a 3D structure called a protein. Scientists use the name "amino acid" because these acids contain both amino group and carboxyl-acid-group in their basic structure. As we mentioned, there are 20 common amino acids present in proteins. Nine of these are essential amino acids in humans because the human body cannot produce them and we obtain them from our diet. Below are two illustrations depicting the relationship between amino acids and polypeptides.



A protein is composed of polypeptide chain(s) and a polypeptide chain is made of amino acids.



A polypeptide chain is a chain composed of amino acids. There are 20 amino acids commonly found in organisms.

Protein Structure

Example

For an interactive illustration of the protein structure levels, check out the protein folding simulation by LabXchange, which uses

As mentioned above, a protein's shape is critical to its function. For example, an enzyme can bind to a specific substrate at an active site. If this active site is altered because of local changes or changes in overall protein structure, the enzyme may be unable to bind to the substrate. To understand how the protein gets its final shape or conformation, we need to understand the four levels of protein structure: primary, secondary, tertiary, and quaternary. See the image below and click on the information hotspots (labeled with an "i") for explanations.

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hemoglobin as an example and describes the molecular structure in more detail.

As seen in the image above, a strand of amino acids folds on itself, creating a unique shape in the tertiary structure of the protein. This is caused by the chemical properties of the amino acids. The chemical properties of the amino acids determine how this shape occurs. For instance, each amino acid is negatively (-), positively (+), or neutrally (N) charged. Negatively charged amino acids bind with positively charged amino acids (neutrally charged amino acids are not affected). Also, the amino acid called cysteine contains sulfur and sulfurs easily bind with each other, creating a "disulfide bond." Because of this, cysteines bind with other cysteines. See the table below for a list of all 20 amino acids and their charges. There are other properties that also influence a protein's shape, such as the amino acid's polarity. Note that these bonds are not as strong as what is created between amino acids when an amino acid chain is created, but these bonds are strong enough to hold the shape in the protein.

Amino Acid	3-Letter Abbrev.	1-Letter Abbrev.	Charge	Di-sulfide Bond Formation?
Alanine	Ala	A	Neutral	
Arginine	Arg	R	(+)	
Asparagine	Asn	N	Neutral	
Aspartate (Aspartic acid)	Asp	D	(-)	
Cysteine	Cys	C	Neutral	Yes
Glutamine	Gln	Q	Neutral	
Glutamate (Glutamic acid)	Glu	E	(-)	
Glycine	Gly	G	Neutral	
Histidine	His	H	(+)	
Isoleucine	Ile	I	Neutral	
Leucine	Leu	L	Neutral	
Lysine	Lys	K	(+)	
Methionine	Met	M	Neutral	
Phenylalanine	Phe	F	Neutral	
Proline	Pro	P	Neutral	
Serine	Ser	S	Neutral	
Threonine	Thr	T	Neutral	
Tryptophan	Trp	W	Neutral	
Tyrosine	Tyr	Y	Neutral	
Valine	Val	V	Neutral	

Exercise

Use the chart above to determine which amino acids may bond together to form the tertiary structure.

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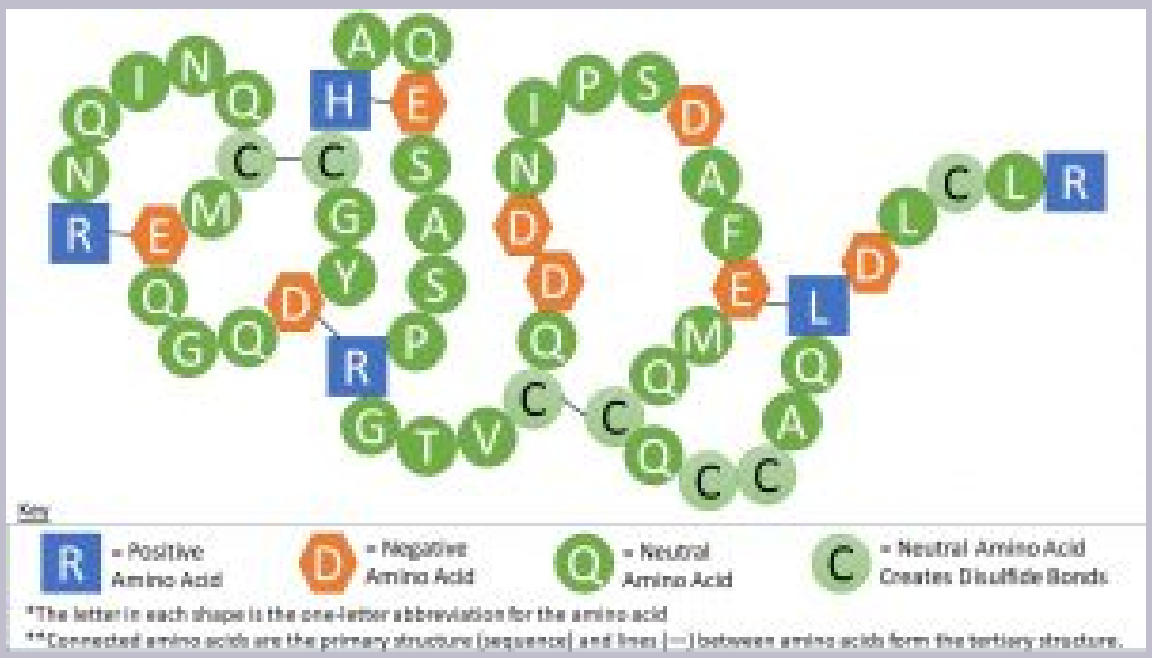
Example

Here is an example of a polypeptide model depicting how charges influence the tertiary structure. The first and second images are the same, except the second image has hotspots with additional information marked with a question mark (?). The key at the bottom of the image is necessary for interpreting the image.



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Mutations

Mutations can impact protein synthesis and amino acid sequence. If these mutations are heritable, then they may influence the evolution of a species. Therefore, this chapter includes information on mutations and evolution.

What Are Mutations?

Mutation is a change in DNA, the hereditary material of life. An organism's DNA codes for the production of proteins, which affects how it looks, how it behaves, and its physiology—all aspects of its life. So, a change in an organism's DNA can cause changes in all aspects of its life.

The gene encoding the protein ultimately determines the unique sequence for every protein. A change in the nucleotide sequence of the gene's coding region may lead to adding a different amino acid to the growing polypeptide chain, causing a change in protein structure and function. In sickle cell anemia, the hemoglobin β chain has a single amino acid substitution, causing a change in protein structure and function. Specifically, valine in the β chain substitutes the amino acid glutamic. What is most remarkable to consider is that a hemoglobin molecule is comprised of two alpha and two beta chains that each consist of about 150 amino acids. The molecule, therefore, has about 600 amino acids. The structural difference between a normal hemoglobin molecule and a sickle cell molecule—which dramatically decreases life expectancy—is a single amino acid out of the total 600. What is even more remarkable is that three nucleotides each encode those 600 amino acids and a single base change (point mutation)—1 in 1800 bases—causes the mutation.

This change to one amino acid in the chain causes hemoglobin molecules to form long fibers that distort the biconcave, or disc-shaped, red blood cells and causes them to assume a crescent, or "sickle," shape that clogs blood vessels. This can lead to a myriad of serious health problems such as breathlessness, dizziness, headaches, and abdominal pain for those affected by this disease.

Mutation Examples

See Learn.Genetics' "The Outcome of Mutation" for descriptions of how specific traits are influenced by mutation.

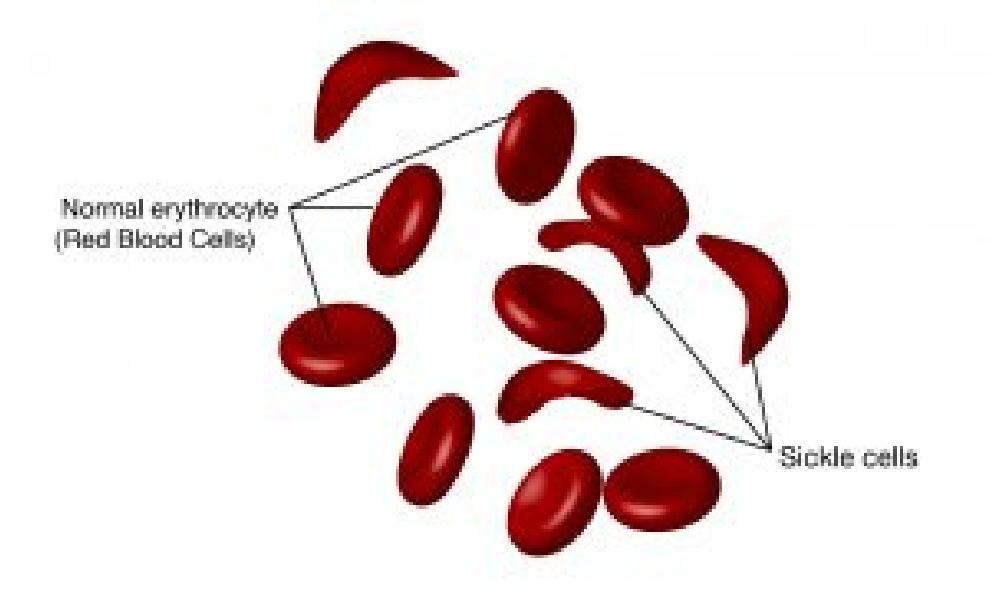
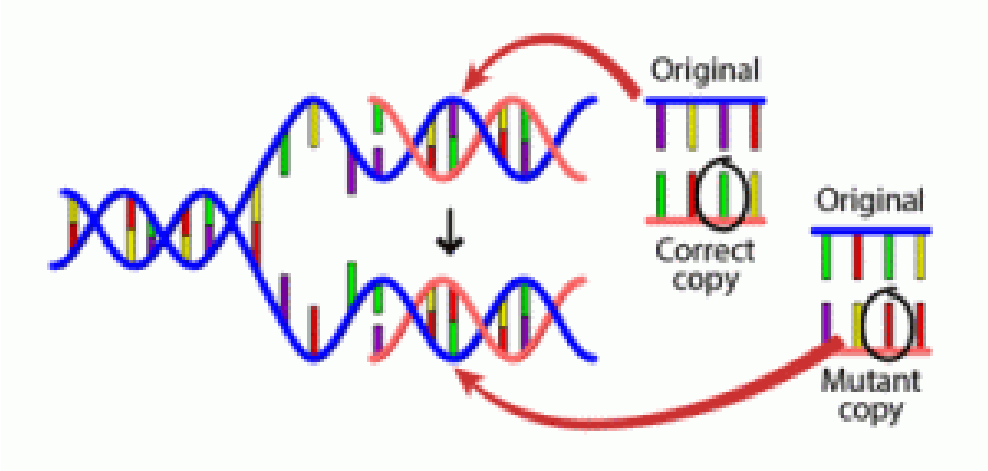


Illustration of normal and sickle cells.

The Causes of Mutations

Mutations happen for several reasons:

- DNA fails to copy accurately: Most of the mutations that we think matter to evolution are "naturally occurring." For example, when a cell divides, it makes a copy of its DNA and sometimes that copy is not quite perfect. That small difference from the original DNA sequence is a mutation.



Mutation can occur during DNA replication.

- External influences can create mutations: Mutations can also be caused by exposure to specific chemicals or radiation. These agents cause the DNA to break down. This is not necessarily unnatural—even in the most isolated and pristine environments, DNA breaks down. Nevertheless, when the cell repairs the DNA, it might not do a perfect job of the repair. So, the cell would end up with DNA slightly different than the original DNA and hence, a mutation.

Evolution

Biological evolution, simply put, is descent with modification. This definition encompasses small-scale evolution (changes in gene—or, more precisely and technically, allele—frequency in a population from one generation to the next) and large-scale evolution (the descent of different species from a common ancestor over many generations). Evolution is responsible for both the remarkable similarities we see across all life and the amazing diversity of that life, but how does it work?

For evolutionary mechanisms (such as natural selection) to act, there needs to be genetic variation and mutations, or changes, in the DNA. DNA codes for proteins, and when those proteins are produced, mutations create variation. Mutations can be beneficial, neutral, or harmful for the organism, but mutations do not "try" to supply what the organism "needs." In this respect, mutations are random—whether a particular mutation happens or not is unrelated to how useful that mutation would be.

Because all cells in our body contain DNA, there are lots of places for mutations to occur; however, not all mutations matter for evolution. Somatic mutations occur in non-reproductive cells and won't be passed onto offspring. Mutations can also be caused by exposure to specific chemicals or radiation. These agents cause the DNA to break down. This is not necessarily unnatural—even in the most isolated and pristine environments, DNA breaks down. Nevertheless, when the cell repairs the DNA, it might not do a perfect job of the repair. So the cell would end up with DNA slightly different than the original DNA and hence, a mutation.

A single germline mutation can have a range of effects:

- ***No change occurs in phenotype***: Some mutations don't have any noticeable effect on the phenotype of an organism. This can happen in many situations: perhaps the mutation occurs in a stretch of DNA with no function, or perhaps the mutation occurs in a protein-coding region but ends up not affecting the amino acid sequence of the protein.
- ***Small change occurs in phenotype***: A single mutation caused this cat's ears to curl backward slightly.

- ***Big change occurs in phenotype***: Some really important phenotypic changes, like DDT resistance in insects, are sometimes caused by single mutations. A single mutation can also have strong negative effects for the organism. Mutations that cause the death of an organism are called lethals—and it doesn't get more negative than that.

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There are some sorts of changes that a single mutation, or even a lot of mutations, could not cause. Neither mutations nor wishful thinking will make pigs have wings; only pop culture could have created the Teenage Mutant Ninja Turtles—mutations could not have done it.

See the "Evolution" chapter in this textbook for more information.

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16.

PROTEIN SYNTHESIS OVERVIEW

Andrea Bierema

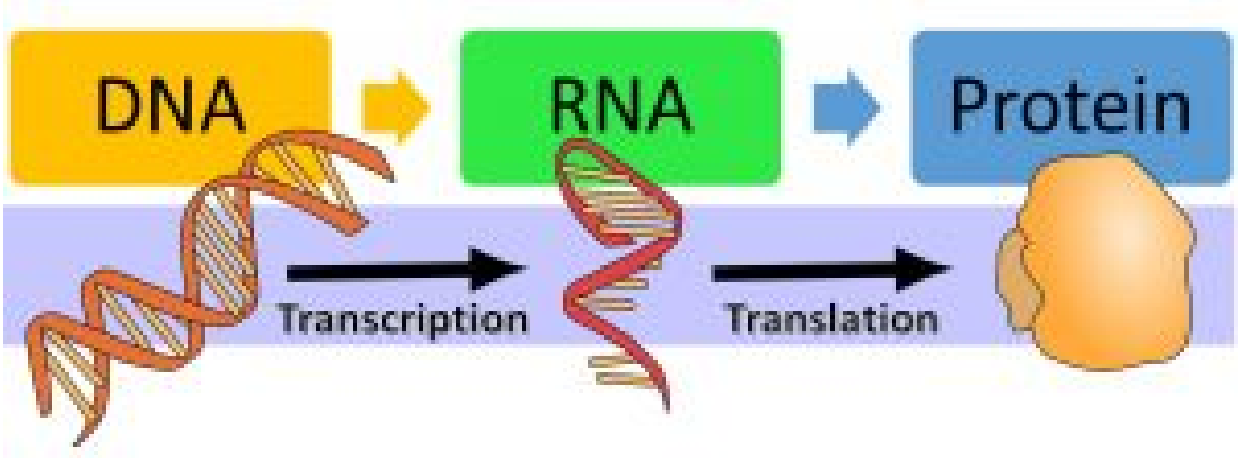
Learning Objectives

Students will be able to:

- Describe the structure and purpose of DNA and RNA.
- Describe the general process of protein synthesis.
- Describe the molecular anatomy of genes and genomes.
- Identify DNA and mRNA bases and binding patterns.
- Interpret a codon-amino acid chart.
- Given a DNA sequence, determine the corresponding mRNA sequence and amino acid sequence.

Central Dogma

The central dogma of molecular biology is that DNA codes for RNA and RNA codes for protein. In addition to DNA coding for RNA, much of the DNA regulates the synthesis of RNA- which ultimately means that it regulates the synthesis of protein. We will learn about protein synthesis regulation in a later chapter.



Protein synthesis consists of two main processes: transcription and translation. During the process of ***transcription***—which occurs in the nucleus—an mRNA molecule is created by reading the DNA. Note that DNA never "becomes" RNA; rather, the DNA is "read" to make an RNA molecule. The mRNA leaves the nucleus and then, through the process of ***translation***, the mRNA is read to create an amino acid sequence which folds into a protein.

Consider what the terms "transcribe" and "translate" mean in relation to language. To "transcribe" something means to rewrite text again in the same language while to "translate" something means to rewrite the text in a different language. Similar to these meanings, in biology, DNA is transcribed into RNA: both DNA and RNA are made of nucleic acid (i.e., the same "language"). With the assistance of proteins, DNA is "read" and transcribed into an mRNA sequence. To read RNA and create protein, though, we refer to it as being translated: RNA is made of nucleic acid and protein is made of amino acid (i.e., different "languages"). Therefore, DNA is transcribed to create an mRNA sequence, and then the mRNA sequence is translated to make a protein.

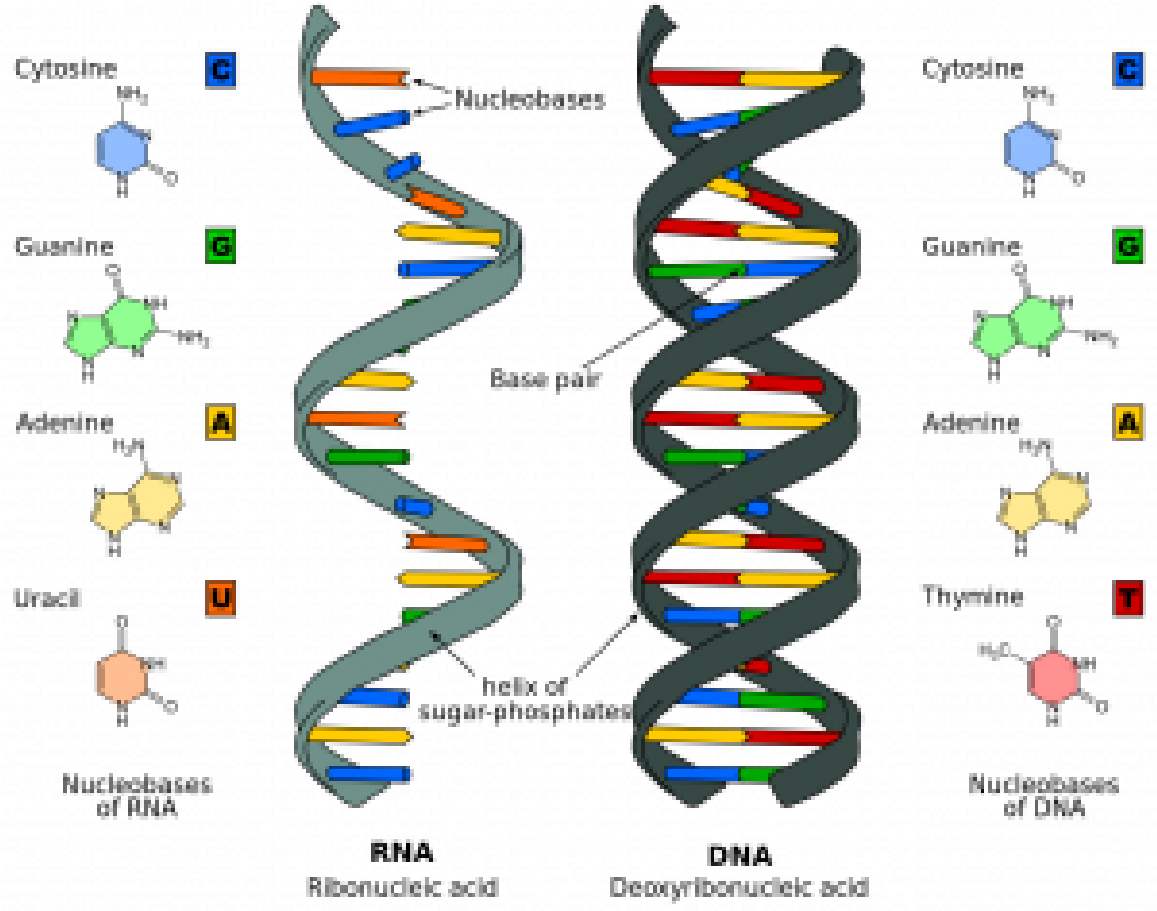
DNA and RNA

The two main types of nucleic acids are deoxyribonucleic acid (DNA) and ribonucleic acid (RNA). DNA is the genetic material in all living organisms, ranging from single-celled bacteria to multicellular mammals. It is in the nucleus of eukaryotes and in the organelles, chloroplasts, and mitochondria. In prokaryotes, the DNA is not enclosed in a membranous envelope.

The cell's entire genetic content is its genome, and the study of genomes is genomics. In eukaryotic cells but not in prokaryotes, a DNA molecule may contain tens of thousands of genes. Many genes contain information to make protein products (e.g., mRNA). Other genes code for RNA products. DNA controls all of the cellular activities by turning the genes "on" or "off."

The other type of nucleic acid, RNA, is mostly involved in protein synthesis. The DNA molecules never leave the nucleus but instead use an intermediary to communicate with the rest of the cell. This intermediary is the messenger RNA (mRNA). Other types of RNA—like rRNA, tRNA, and microRNA—are involved in protein synthesis and its regulation.

DNA and RNA are comprised of ***monomers*** that scientists call ***nucleotides***. The nucleotides combine with each other to form a ***polynucleotide***, DNA or RNA. Three components comprise each nucleotide: a nitrogenous base, a pentose (five-carbon) sugar, and a phosphate group. Each nitrogenous base in a nucleotide is attached to a sugar molecule, which is attached to one or more phosphate groups. Therefore, although the terms "base" and "nucleotide" are sometimes used interchangeably, a nucleotide contains a base as well as part of the sugar-phosphate backbone.



Comparison of RNA (left molecule) and DNA (right molecule). The color of the bases in RNA and DNA aligns with the colored boxes next to each base molecule.

Exercises

Examine the image above and then answer the following questions:

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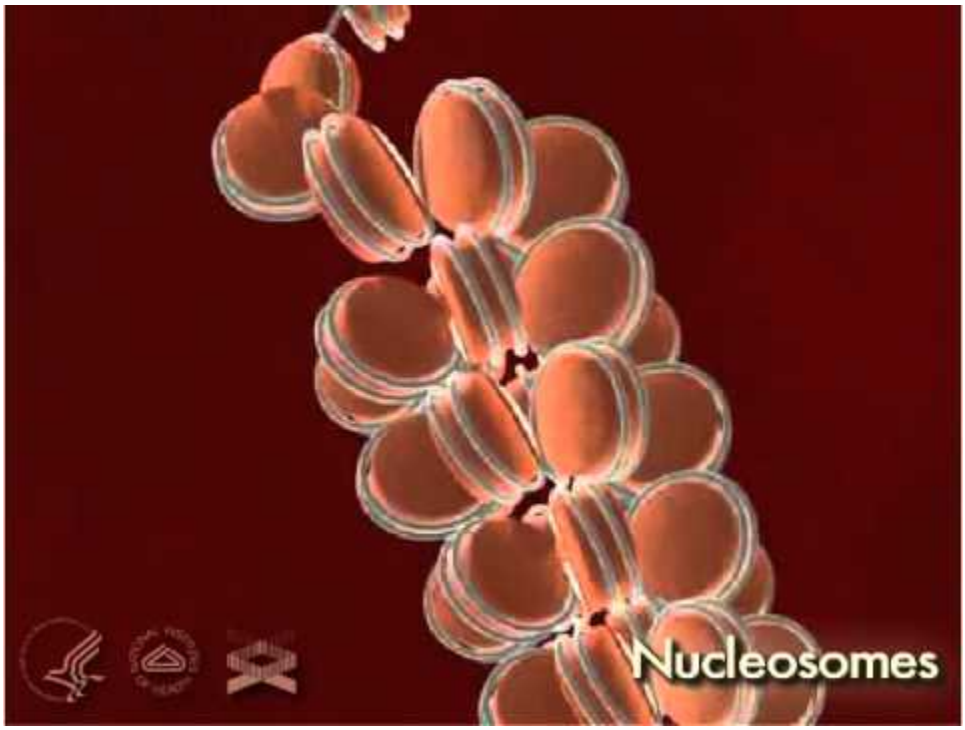
What is a Gene?

The gene is the basic physical unit of inheritance. Genes are passed from parents to offspring and contain the information needed to specify traits. Genes are arranged, one after another, on structures called chromosomes. A chromosome contains a single, long DNA molecule, only a portion of which corresponds to a single gene. Humans have approximately 20,000 genes arranged on their chromosomes. Watch the following video for an animated view on the relationship between chromosomes and genes.



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Protein Synthesis Overview

The two main processes in protein synthesis are transcription and translation. The following is an overview of each of these processes. Each process will be described in more detail in future chapters.

Transcription

A gene is complex: it contains not only the code for the resulting protein but also several regulatory factors that determine if and when the region that codes for a protein are read to create protein. What follows is a diagram of the components of a gene that are used in transcription.

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For this chapter, we focus on the DNA and the ending product of transcription: mRNA.

Exercise

Given a specific DNA strand, what is the sequence of the resulting mRNA molecule? We will learn about how mRNA is created in a later chapter.

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Translation

Translation involves different types of RNA, and we will explain them in more detail in a later chapter: rRNA, tRNA, mRNA, and microRNA.

After an mRNA is created, it leaves the nucleus and is attracted to or attracts a ribosome, which is a molecule made of rRNA and polypeptides. Then, in the ribosome, and with the assistance of tRNAs, the mRNA is read and an amino acid sequence is created.

DNA and mRNA create sequences with just four types of bases; yet, these bases code for 20 unique amino acids (the makeup of protein). How is this possible? Watch the following video to find out!

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The mRNA is read in sets of three bases known as codons. Each codon codes for a single amino acid. In this way, the mRNA is read and the protein product is made.

Below is a chart showing which codons code for which bases. There are two representations; move to the next slide for the second representation.

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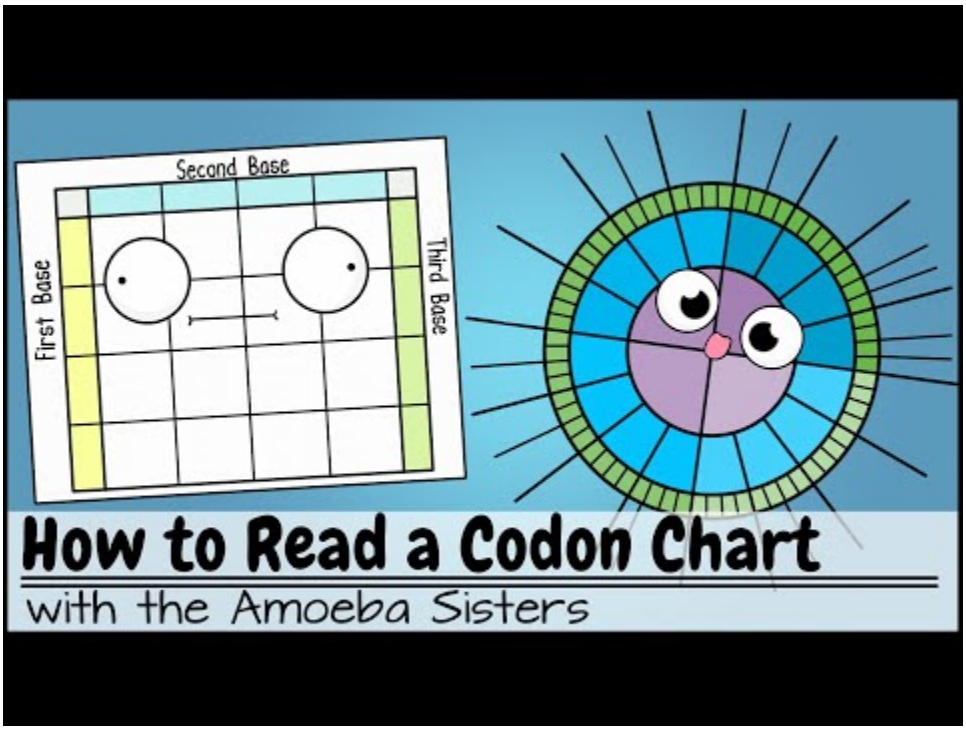
These charts can be a little confusing at first. Watch the following video to learn how to interpret both chart formats.



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Exercise

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Conclusion

This chapter focused on DNA, mRNA, and protein sequences. The next several chapters describe the processes that take place during protein synthesis. Master how sequences are read during protein synthesis (the focus of the current chapter) before moving on to the next chapter. Below are some sources to help further your understanding!

Example

Check out Learn.Genetics' "How a Firefly's Tail Makes Light" video for an overview of protein synthesis!

Exercises

Need a little more practice?

Try out Learn.Genetics' "Transcribe and Translate a Gene" and The Concord Consortium's "DNA to Protein" interactives for further practice!

Attributions

This chapter is a modified derivative of the following articles:

"Gene" by National Human Genome Research Institute, National Institutes of Health, *Talking Glossary of Genetic Terms.*

"Nucleic Acids" by OpenStax College, Biology 2e, CC BY 4.0. Download the original article at https://openstax.org/books/biology-2e/pages/3-5-nucleic-acids

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17.

PROTEIN SYNTHESIS I: TRANSCRIPTION

Andrea Bierema

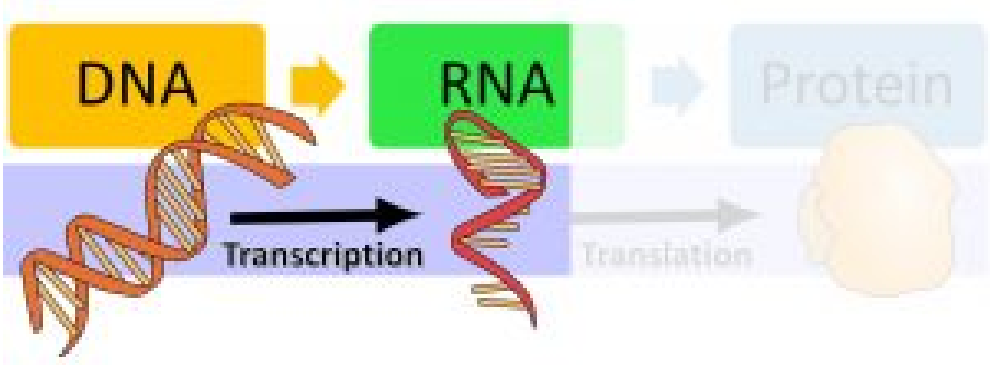
Learning Objectives

Students will be able to:

- Explain the processes necessary for transcription to begin.
- Explain how DNA is transcribed to create an mRNA sequence.
- Describe the role of polymerase in transcription.
- Recognize that protein synthesis regulation (i.e., changes in gene expression) allow cells to respond to changes in the environment.
- Explain which gene-expression regulatory factors are at play for transcription.

Overview

This chapter focuses on how transcription works; that is, how information coded in the DNA molecule is read to create an mRNA sequence. Please see the previous chapter for a general overview of transcription and DNA and RNA bases before continuing to read this chapter.



Transcription is the process of creating an mRNA sequence by "reading" the DNA sequence.

The Process of Transcription: A First Look

Let's first look at a basic overview of what the process of transcription looks like. At the beginning of the following video, you will see that transcription is regulated by a variety of proteins. By "regulation", we mean that certain proteins are needed for transcription to start and some proteins can even prevent transcription from happening. Transcription is happening throughout your body all of the time, but not every gene is constantly being transcribed in every cell; it is regulated by different proteins and depends on which proteins your body needs in which cells.

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Exercise

Now that you have watched a basic overview of transcription, test your knowledge with the following activity in which you will place the following transcription steps in the correct order.

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Role of the Polymerase

The polymerase is an enzyme—and a protein—that aids in the transcription process. The polymerase was depicted in the previous video. Now let's look more closely at what is happening within the polymerase in relation to the steps described previously.

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Transcription Regulation

The overview above depicted components of transcription regulation. Basically, there are proteins that have to bind to the DNA, and each other, before the polymerase can begin transcription.

There are many steps along the way of protein synthesis and gene expression is regulated. Gene expression is when a gene in DNA is "turned on," that is, used to make the protein it specifies. Not all the genes in your body are turned on at the same time or in the same cells or parts of the body.

For many genes, transcription is the key on/off control point: if a gene is not transcribed in a cell, it can't be used to make a protein in that cell.

If a gene does get transcribed, it is likely going to be used to make a protein (i.e. expressed). In general (but not always) the more often a gene is transcribed, the more protein that will be made.

Various factors control how much a gene is transcribed. For instance, how tightly the DNA of the gene is wound around its supporting proteins to form chromatin can affect a gene's availability for transcription.

Proteins called transcription factors, however, play a particularly central role in regulating transcription. These important proteins help determine which genes are active in each cell of your body.

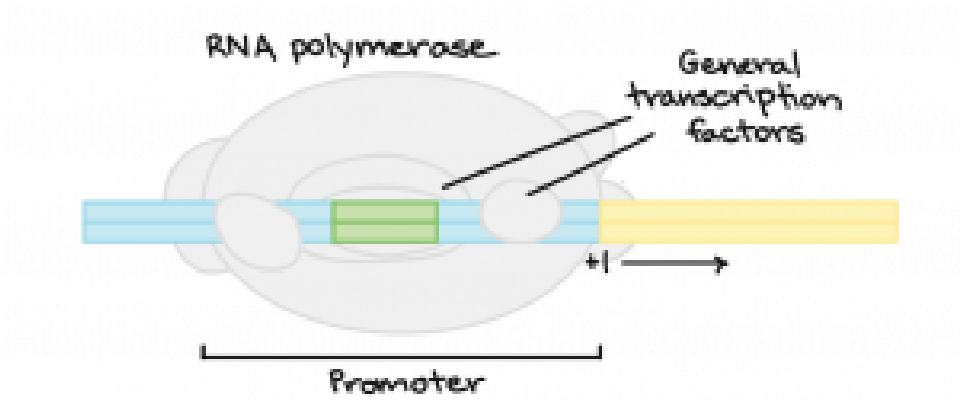
Transcription Factors

What has to happen for a gene to be transcribed? The enzyme **RNA polymerase**, which makes a new RNA molecule from a DNA template, must attach to the DNA of the gene. It attaches to a spot called the **promoter**.

The RNA polymerase can attach to the **promoter** only with the help of proteins called **general transcription factors**. They are part of the cell's core transcription "toolkit," needed for the transcription of any gene.

More information

In bacteria, RNA polymerase attaches right to the DNA of the promoter. You can see how this process works, and how it can be regulated by transcription factors, in the *lac* operon and *trp* operon videos.



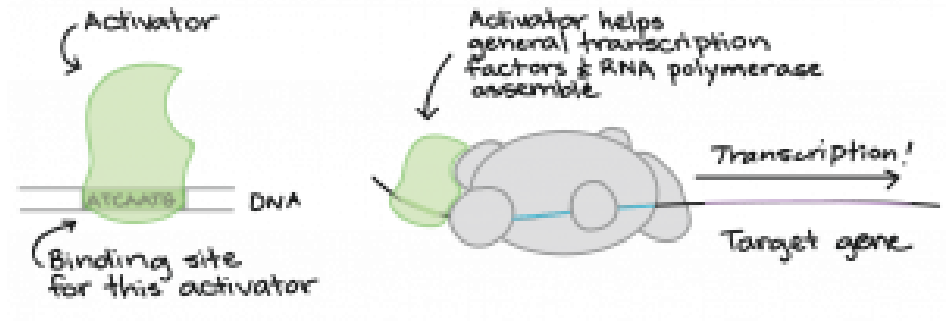
However, many transcription factors (including some of the coolest ones!) are not the general kind. Instead, there is a large class of transcription factors that control the expression of specific, individual genes. For instance, a transcription factor might activate only a set of genes needed in certain neurons.

How do Transcription Factors Work?

A typical transcription factor binds to DNA at a certain target sequence. Once it's bound, the transcription factor makes it either harder or easier for RNA polymerase to bind to the promoter of the gene.

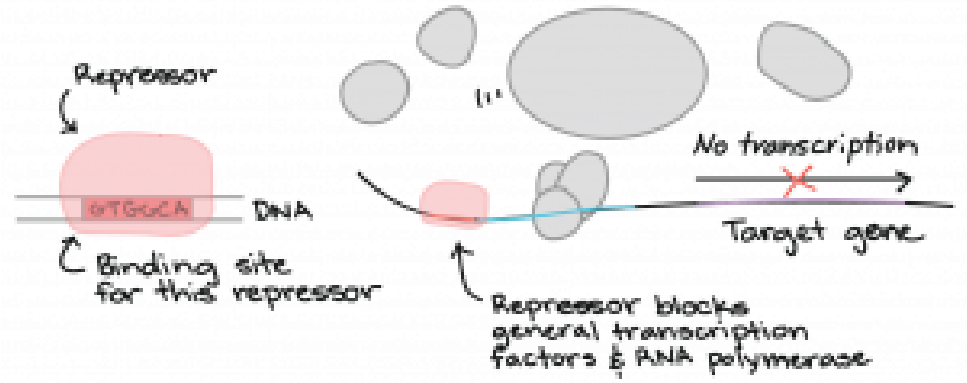
Activators

Some transcription factors **activate** transcription. For instance, they may help the general transcription factors and/or RNA polymerase bind to the promoter, as shown in the diagram below.



Repressors

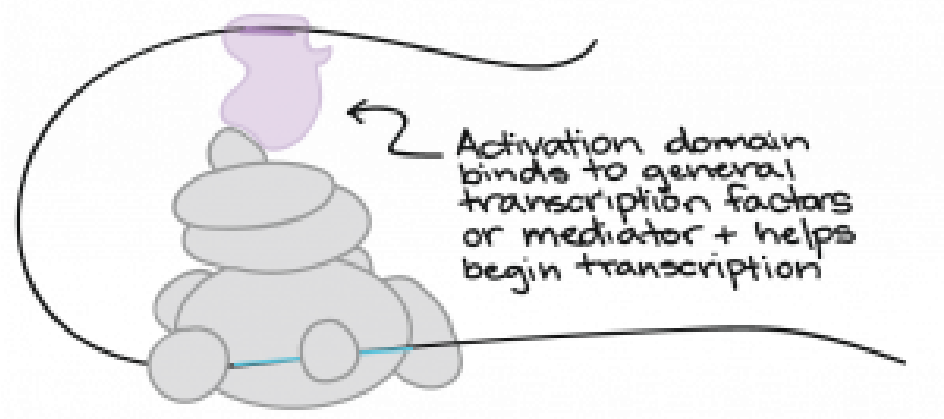
Other transcription factors **repress** transcription. This repression can work in a variety of ways. As one example, a repressor may get in the way of the basal transcription factors or RNA polymerase, making it so they can't bind to the promoter or begin transcription.



Turning Genes on in Specific Body Parts

Some genes need to be expressed in more than one body part or type of cell. For instance, suppose a gene needed to be turned on in your spine, skull, and fingertips, but not in the rest of your body. How can transcription factors make this pattern happen?

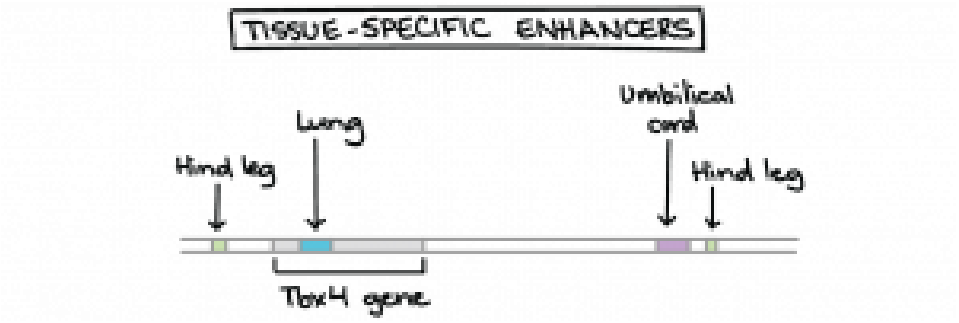
A gene with this type of pattern may have several **enhancers** (far-away clusters of binding sites for activators) or **silencers** (the same thing, but for repressors). Each enhancer or silencer may activate or repress the gene in a certain cell type or body part, binding transcription factors that are made in that part of the body.[1,2]



Example: Modular Mouse

As an example, let's consider a gene found in mice, called *Tbx4*. This gene is important for the development of many different parts of the mouse body, including the blood vessels and hind legs.[3]

During development, several well-defined enhancers drive *Tbx4* expression in different parts of the mouse embryo. The diagram below shows some of the *Tbx4* enhancers, each labeled with the body part where it produces expression.



Not drawn fully to scale. This Image is based on Figure 5 of Menke et al.[3]

Evolution of Development

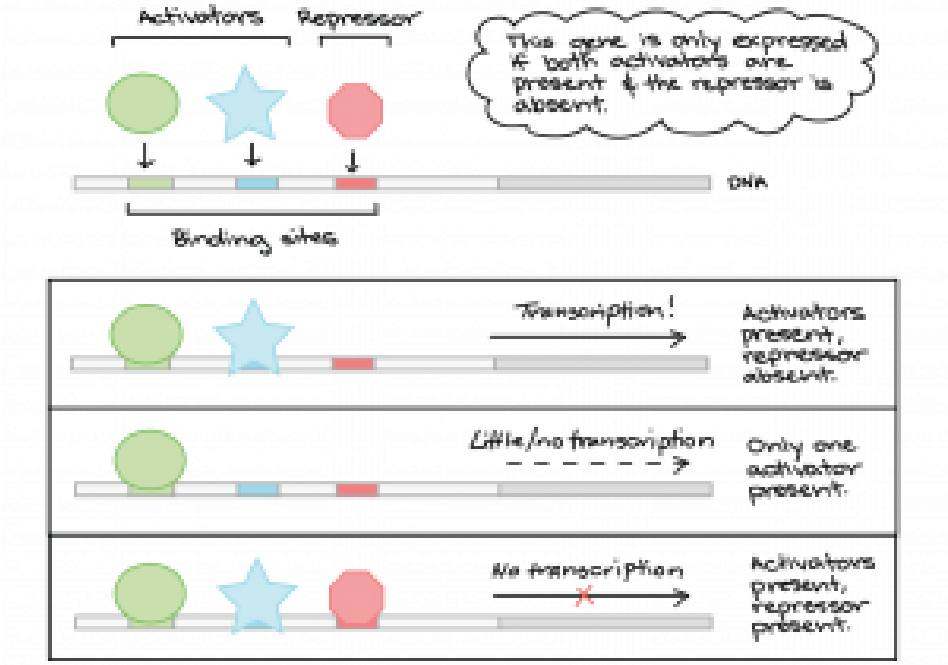
Enhancers like those of the *Tbx4* gene are called tissue-specific enhancers: they control a gene's expression in a certain part of the body. Mutations of tissue-specific enhancers and silencers may play a key role in the evolution of body form.[4]

How could that work? Suppose that a mutation, or change in DNA, happened in the coding sequence of the *Tbx4* gene. The mutation would inactivate the gene everywhere in the body and a mouse without a normal copy would likely die. However, a mutation in an enhancer might just change the expression pattern a bit, leading to a new feature (e.g., a shorter leg) without killing the mouse.

Transcription Factors and Cellular "Logic"

Can cells do logic? Not in the same way as your amazing brain. However, cells can detect information and combine it to determine the correct response—in much the same way that your calculator detects pushed buttons and outputs an answer.

We can see an example of this "molecular logic" when we consider how transcription factors regulate genes. Many genes are controlled by several different transcription factors, with a specific combination needed to turn the gene on; this is particularly true in eukaryotes and is sometimes called combinatorial regulation.[5,6] For instance, a gene may be expressed only if activators A and B are present, and if repressor C is absent.



The use of multiple transcription factors to regulate a gene means that different sources of information can be integrated into a single outcome. For instance, imagine that:

- Activator A is present only in skin cells
- Activator B is active only in cells receiving "divide now!" signals (growth factors) from neighbors
- Repressor C is produced when a cell's DNA is damaged

In this case, the gene would be "turned on" only in skin cells that are receiving division signals and have undamaged, healthy DNA. This pattern of regulation might make sense for a gene involved in cell division in skin cells. In fact, the loss of proteins similar to repressor C can lead to cancer.
Real-life combinatorial regulation can be a bit more complicated than this. For instance, many different transcription factors may be involved, or it may matter exactly how many molecules of a given transcription factor are bound to the DNA.

A Closer Look

After reading through this section, view the following video, which depicts many of the regulatory factors described above.

An interactive or media element has been excluded from this version of the text. You can view it online here:
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Examples

Now that you have learned some of the basics, check out this example that applies what you learned to a specific case study.

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The video above briefly describes the laboratory part of this research. To learn more about what this research looks like, check out the "Stickleback Evolution Virtual Lab."

Lactose Example

If you are still a little unsure of how switches work, then check out this *HMMI Biointeractive* interactive. The ability to digest lactose as an adult is a rare phenomenon in mammals. It evolved twice in humans—in Africa and Europe.

Exercise

Now let's test your understanding of transcription regulation!

Take the quiz below the simulation as you work your way through it. Note that if you are using your mouse to scroll down, it may not work at this point—use the scrolling bar at the right edge of your web browser instead.

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The Process of Transcription: A Detailed Look

This chapter began with an overview of transcription and then focused more deeply on the role of the polymerase and regulatory proteins. Now watch the following video. It is an in-depth version of the first video of this chapter, incorporating aspects described throughout this chapter.

An interactive or media element has been excluded from this version of the text. You can view it online here:
https://openbooks.lib.msu.edu/isb202/?p=274

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1. Gilbert, S. F. (2000). Anatomy of the gene: Promoters and enhancers. In *Developmental biology* (6th ed.). Sunderland, MA: Sinauer Associates. Retrieved from https://www.ncbi.nlm.nih.gov/books/NBK10023/#_A751_.
2. Gilbert, S. F. (2000). Silencers. In *Developmental biology* (6th ed.). Sunderland, MA: Sinauer Associates.

Retrieved from http://www.ncbi.nlm.nih.gov/books/NBK10023/#_A777_.
3. Menke, D. B., Guenther, C., and Kingsley, D. M. (2008). Dual hindlimb control elements in the Tbx4 gene and region-specific control of bone size in vertebrate limbs. *Development*, *135*, 2543-2553. http://dx.doi.org/10.1242/dev.017384.
4. Wray, Gregory A. (2007). The evolutionary significance of *cis*-regulatory mutations. *Nature Reviews Genetics*, *8*, 206-216. http://dx.doi.org/10.1038/nrg2063.
5. Reece, J. B., Urry, L. A., Cain, M. L., Wasserman, S. A., Minorsky, P. V., and Jackson, R. B. (2011). Combinatorial control of gene activation. In *Campbell Biology* (10th ed., pp. 37). San Francisco, CA: Pearson.
6. Reményi, Attila, Hans R. Schöler, and Matthias Wilmanns. (2004). Combinatorial control of gene expression. *Nature Structural & Molecular Biology*, *11*(9), 812. http://dx.doi.org/10.1038/nsmb820. Retrieved from http://www.nature.com/scitable/content/Combinatorial-control-of-gene-expression-16976.

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18.

PROTEIN SYNTHESIS II: RNA PROCESSING

Andrea Bierema

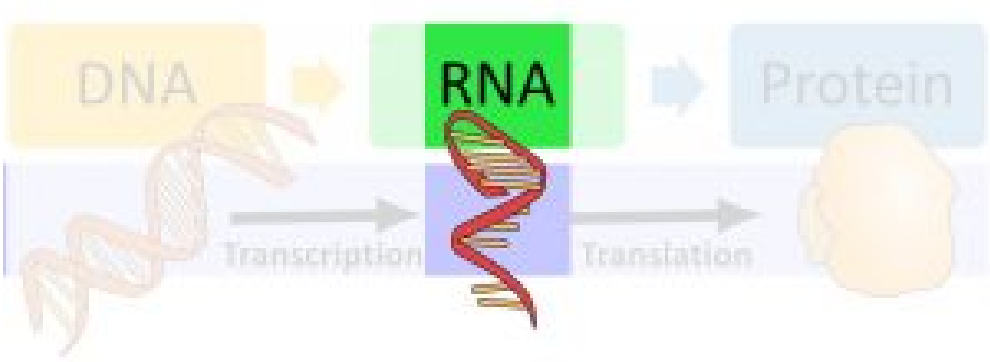
Learning Objectives

Students will be able to:

- Describe the molecular anatomy of genes and genomes.
- Describe exons and introns.
- Explain how splicing occurs during mRNA processing.
- Explain the process of alternative splicing.
- Describe how alternative splicing contributes to cells having different functions.
- Identify additional steps that take place during mRNA processing.

Overview

At the end of transcription, once the polymerase releases the RNA molecule, the RNA is not quite ready for translation and is not technically an "mRNA" yet. It still needs to be "processed," which means that certain nucleotides are removed and others are added—at the end of this, it is an mRNA molecule, but before that, it is a pre-mRNA molecule. This occurs as the RNA molecule is leaving the nucleus. After it is processed, it is then ready for translation, which is covered in a future chapter. This chapter focuses on the different steps that take place during mRNA processing and how some of these steps allow for cells within a single organism to have different functions. Please see the previous chapter for a review of transcription.

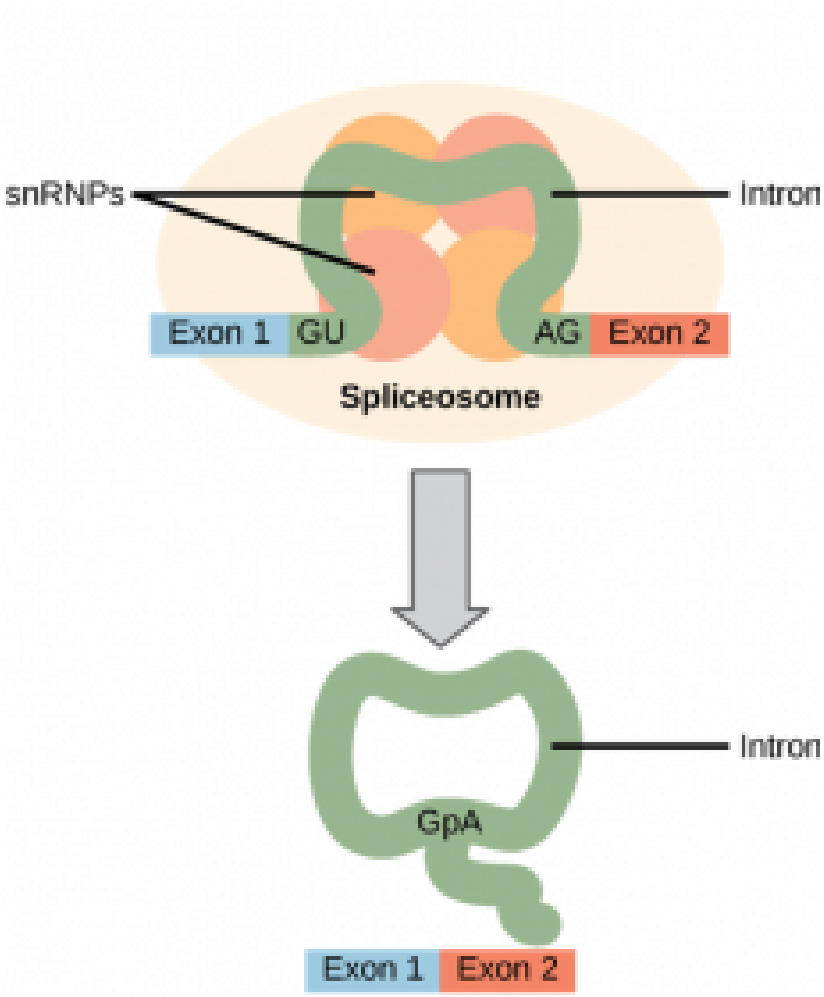


The focus of this chapter is what happens to the RNA produced from transcription before it can be translated.

Splicing

After an mRNA molecule is produced, specific sequences throughout the mRNA molecule are removed. These sequences are called introns. The sequences that remain are called exons. To recall these terms, consider the following: exons correspond to protein-coding sequences ("EX–on" signifies that they are *ex*pressed), and *int*ervening sequences called introns ("IN-tron" denotes their *int*ervening role), which may be involved in gene regulation but are removed from the pre-mRNA during processing. Intron sequences in mRNA do not encode functional proteins.

All of a pre-mRNA's introns must be completely and precisely removed before protein synthesis. If the process errs by even a single nucleotide, the reading frame of the rejoined exons would shift, and the resulting protein would be dysfunctional. The process of removing introns and reconnecting exons is called ***splicing***.



Pre-mRNA splicing involves the precise removal of introns from the pre-mRNA. The splicing process is catalyzed by protein complexes called spliceosomes that are composed of proteins and RNA molecules called small nuclear RNAs (snRNAs).

If we consider that a base sequence is similar to letters on a page, then splicing is removing particular lines of text. View the following video, which follows this analogy, for a brief overview of what splicing does (open the summary dialog at the end of the video to reflect on what you learned).

An interactive or media element has been excluded from this version of the text. You can view it online here:
https://openbooks.lib.msu.edu/isb202/?p=276

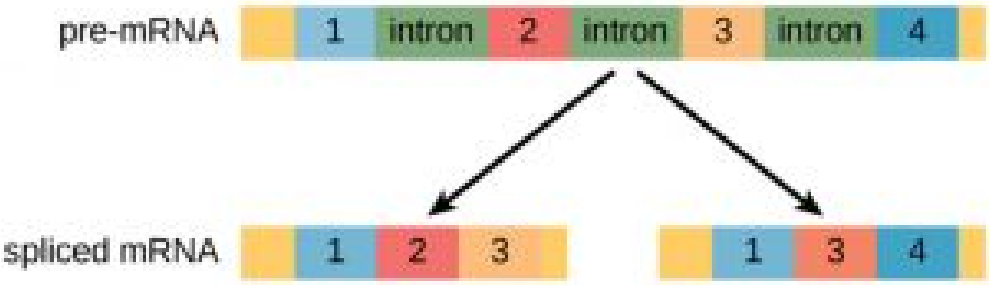
Exercise

Let's test your knowledge! Complete the following image by dragging the parts of the pre-RNA that will be the mature mRNA after mRNA processing. The untranslated regions are already in place.

An interactive or media element has been excluded from this version of the text. You can view it online here:
https://openbooks.lib.msu.edu/isb202/?p=276

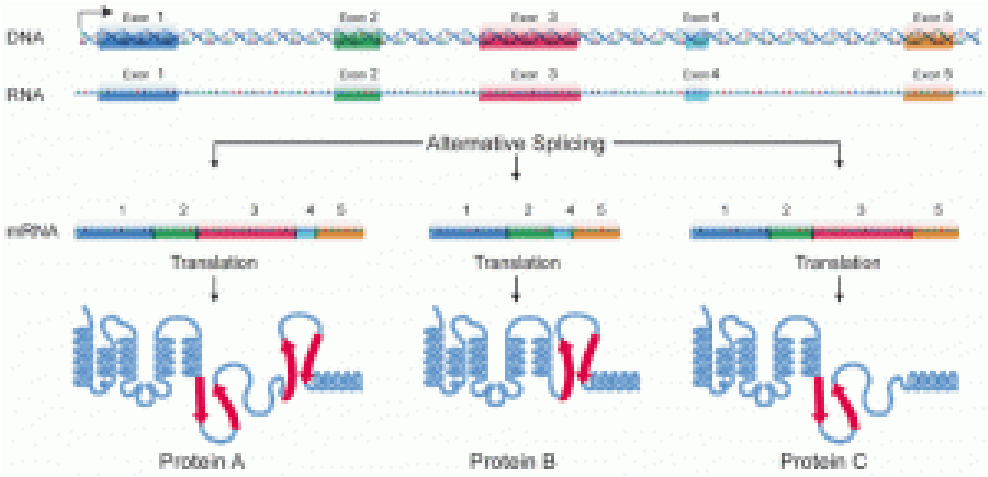
Alternative Splicing

Genes that exhibited alternative RNA splicing were first observed in the 1970s. Alternative RNA splicing is a mechanism that allows different protein products to be produced from one gene when different combinations of exons are combined to form the mRNA. This alternative splicing can be haphazard, but more often it is controlled and acts as a mechanism of *gene regulation*, with the frequency of different splicing alternatives controlled by the cell as a way to control the production of different protein products in different cells or at different stages of development. Alternative splicing is now understood to be a common mechanism of gene regulation in eukaryotes; according to one estimate, 70% of genes in humans are expressed as multiple proteins through alternative splicing. Although there are multiple ways to alternatively splice RNA transcripts, the original 5′-3′ order of the exons is *always conserved*. That is, a transcript with exons 1 2 3 4 5 6 7 might be spliced 1 2 4 5 6 7 or 1 2 3 6 7, but never 1 2 5 4 3 6 7.



An example of alternative splicing. Each number is a different exon, and not all exons are expressed. For example, a pre-mRNA with exons 1, 2, 3, 4 may express exons 1, 2, and 3 in one cell type and exons 1, 3, and 4 in another cell type.

Here is another representation of alternative splicing—this one does not label the introns (assume the molecule segment between the exons are introns), like the previous but it shows how alternative splicing impacts the protein.

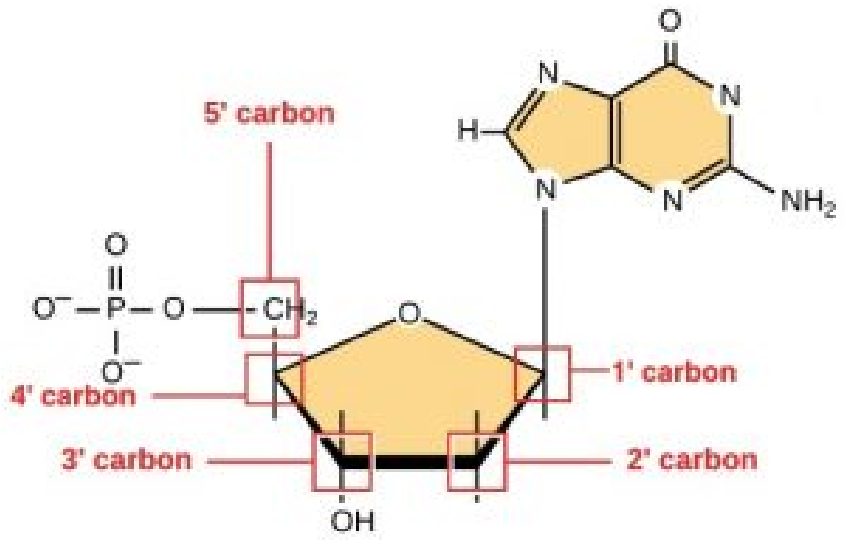


When a pre-mRNA molecule is spliced in different ways, it results in different base sequences, which changes the amino acid sequence and affects the protein's structure and function.

How could alternative splicing evolve? Introns have a beginning—and ending—recognition sequence; it is easy to imagine the failure of the splicing mechanism to identify the end of an intron and instead find the end of the next intron, thus removing two introns and the intervening exon. In fact, there are mechanisms in place to prevent such intron-skipping, but mutations are likely to lead to their failure. Such "mistakes" would more than likely produce a nonfunctional protein. Indeed, the cause of many genetic diseases is abnormal splicing rather than mutations in a coding sequence. However, alternative splicing could possibly create a protein variant without the loss of the original protein, opening up possibilities for adaptation of the new variant to new functions. Gene duplication has played an important role in the evolution of new functions in a similar way by providing genes that may evolve without eliminating the original, functional protein.

Additional Processing

Before the mRNA leaves the nucleus, it is given two protective "caps" that prevent the ends of the strand from degrading during its journey. The two ends of a DNA strand are referred to as 3′ and 5′, which references the position of sugar molecules in the DNA. The 5′ cap is placed on the 5′ end of the mRNA. The poly-A tail, which is attached to the 3′ end, is usually composed of a long chain of adenine (A) nucleotides. These changes protect the two ends of the RNA from being broken down by other enzymes in the cell.



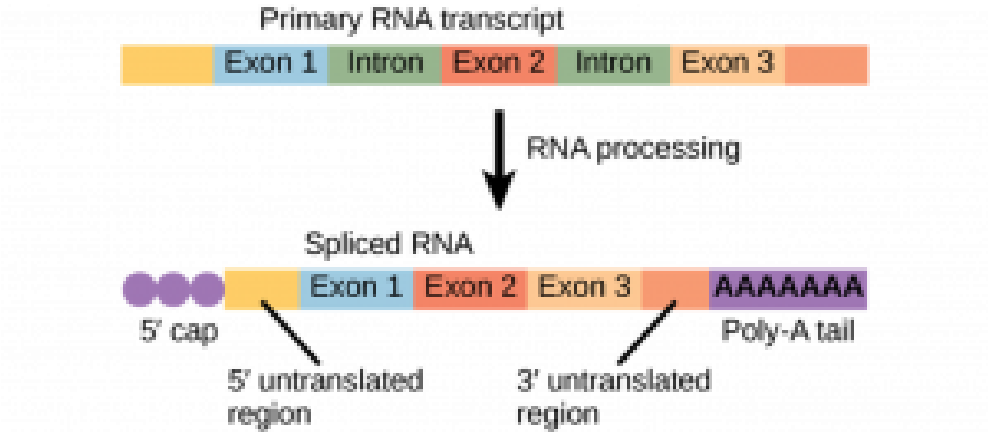
A nucleotide sugar is composed of 5 carbons; each one is numbered (the apostrophe " ' " is called "prime"). Phosphates bind to the 3' carbon and the 5' carbon of each sugar. One end of a DNA or RNA molecule ends with the 3' carbon exposed and the other side ends with a phosphate attached to the last sugar's 5' end. This is why one end is referred to as 3' and the other end is referred to as 5'.

5′ Capping

While the pre-mRNA is still being synthesized, a 7-methylguanosine cap is added to the 5′ end of the growing transcript by a phosphate linkage. This functional group protects the nascent mRNA from degradation. In addition, factors involved in protein synthesis recognize the cap to help initiate translation by ribosomes.

3′ Poly-A Tail

Once elongation is complete, the pre-mRNA is cleaved by an endonuclease between an AAUAAA consensus sequence and a GU-rich sequence, leaving the AAUAAA sequence on the pre-mRNA. An enzyme called poly-A polymerase then adds a string of approximately 200 A residues, called the poly-A tail. This modification further protects the pre-mRNA from degradation and is also the binding site for a protein necessary for exporting the processed mRNA to the cytoplasm. Below is a more complete diagram of mRNA processing.



This image is a more complete representation of RNA processing than the original one introduced earlier in this chapter. In addition to the removal of the introns, the mature mRNA has a cap on one end and poly-A tail on the other end.

Test Your Understanding

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19.

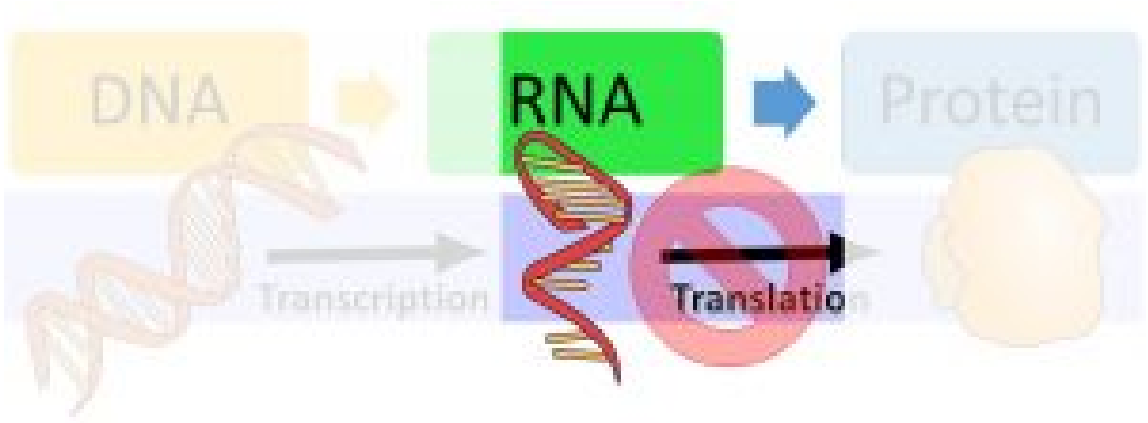
PROTEIN SYNTHESIS III: RNA INTERFERENCE

Andrea Bierema

Learning Objectives

Students will be able to:

- Describe the process of RNA interference.
- Compare the two types of RNA interference: microRNA and siRNA.
- Explain how RNA interference affects gene expression.



RNAi affects protein synthesis after the mRNA is created but before translation begins. It prevents translation from occurring.

Recent Discovery

Not all RNA molecules code for protein. Some RNA control genes in a way that was only discovered recently: a process called RNA interference, or RNAi. Although scientists identified RNAi relatively recently, they now know that organisms have been using this trick for millions of years.

RNAi is a mechanism that organisms use to silence genes when the proteins that they code for are no longer needed. This silencing happens when short RNA molecules bind to stretches of mRNA, preventing translation of the mRNA.

Click on the "plus" hotspots on the figure below to learn more!

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https://openbooks.lib.msu.edu/isb202/?p=278

(credit: modification of work by Robinson, R)

Researchers believe that RNAi arose as a way to reduce the production of a gene's encoded protein for purposes of fine-tuning growth or self-defense. When viruses infect cells, for example, they command their host to produce specialized RNAs that allow the virus to survive and make copies of itself. Researchers believe that RNAi eliminates unwanted viral RNA and some speculate that it may even play a role in human immunity.

Oddly enough, scientists discovered RNAi from a failed experiment! Researchers investigating genes involved in plant growth noticed something strange: when they tried to turn petunia flowers purple by adding an extra "purple" gene, the flowers bloomed white instead.

This result fascinated researchers, who could not understand how adding genetic material could somehow get rid of an inherited trait. The mystery remained unsolved until, a few years later, two geneticists studying development saw a similar thing happening in lab animals.

The researchers, Andrew Z. Fire (then of the Carnegie Institutions of Washington in Baltimore and now at Stanford University) and Craig Mello (of the University of Massachusetts Medical School in Worcester) were trying to block the expression of genes that affect cell growth and tissue formation in roundworms, using a molecular tool called antisense RNA.

To their surprise, Mello and Fire found that their antisense RNA tool wasn't doing much at all. Rather, they determined, a double-stranded contaminant produced during the synthesis of the single-stranded antisense RNA interfered with gene expression. Mello and Fire named the process RNAi (RNA *interference*) and in 2006 were awarded the Nobel Prize in physiology or medicine for their discovery.

Further experiments revealed that the double-stranded RNA gets chopped up inside the cell into much smaller pieces that stick to mRNA and block its action. That is, the mRNA cannot bind to a ribosome and translation cannot occur.

Today, scientists are taking a cue from nature and using RNAi to explore biology. They have learned, for example, that the process is not limited to worms and plants, but operates in humans too.

Medical researchers are currently testing new types of RNAi-based drugs for treating conditions such as macular degeneration (the leading cause of blindness) and various infections (including those caused by HIV and the herpes virus).

RNA Types

There are two main types of RNA that can interfere with mRNA and translation: microRNA and siRNA. This section describes microRNA in more detail. Watch the video in the next section to learn how microRNA and siRNA compare.

Molecules called microRNAs have been found in organisms as diverse as plants, worms, and people. The molecules are truly "micro," consisting of only a few dozen nucleotides, compared to typical human mRNAs that are a few thousand nucleotides long.

What's particularly interesting about microRNAs is that many of them evolved from DNA that used to be considered merely filler material. (14)

How do these small, but important, RNA molecules do their work? They start out much bigger, but get trimmed by cellular enzymes including one aptly named Dicer. Like tiny pieces of Velcro®, microRNAs stick to certain mRNA molecules and stop them from passing on their protein-making instructions.

First discovered in a roundworm model system, some microRNAs help determine the organism's body plan. In their absence, very bad things can happen. For example, worms engineered to lack a microRNA called let-7 develop so abnormally that they often rupture and practically break in half as the worm grows.

Perhaps it is not surprising that because microRNA helps specify the timing of an organism's developmental plan, the appearance of the microRNAs themselves is carefully timed inside a developing organism. Biologists (including Amy Pasquinelli of the University of California, San Diego) are currently figuring out how microRNAs are made and cut to size, as well as how they are produced at the proper time during development.

MicroRNA molecules also have been linked to cancer. For example, Gregory Hannon of the Cold Spring Harbor Laboratory on Long Island in New York, found that certain microRNAs are associated with the severity of the blood cancer B-cell lymphoma in mice.

Since the discovery of microRNAs in the first years of the 21st century, scientists have identified hundreds of them that likely exist as part of a large family with similar nucleotide sequences. New roles for these molecules are still being found.

Mechanisms

Watch the following video to learn about how the two main types of molecules used in RNAi (microRNA and siRNA) differ from one another and how they work to prevent (or interfere with) gene expression.

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https://openbooks.lib.msu.edu/isb202/?p=278

Exercise

After watching the video, answer the following question:

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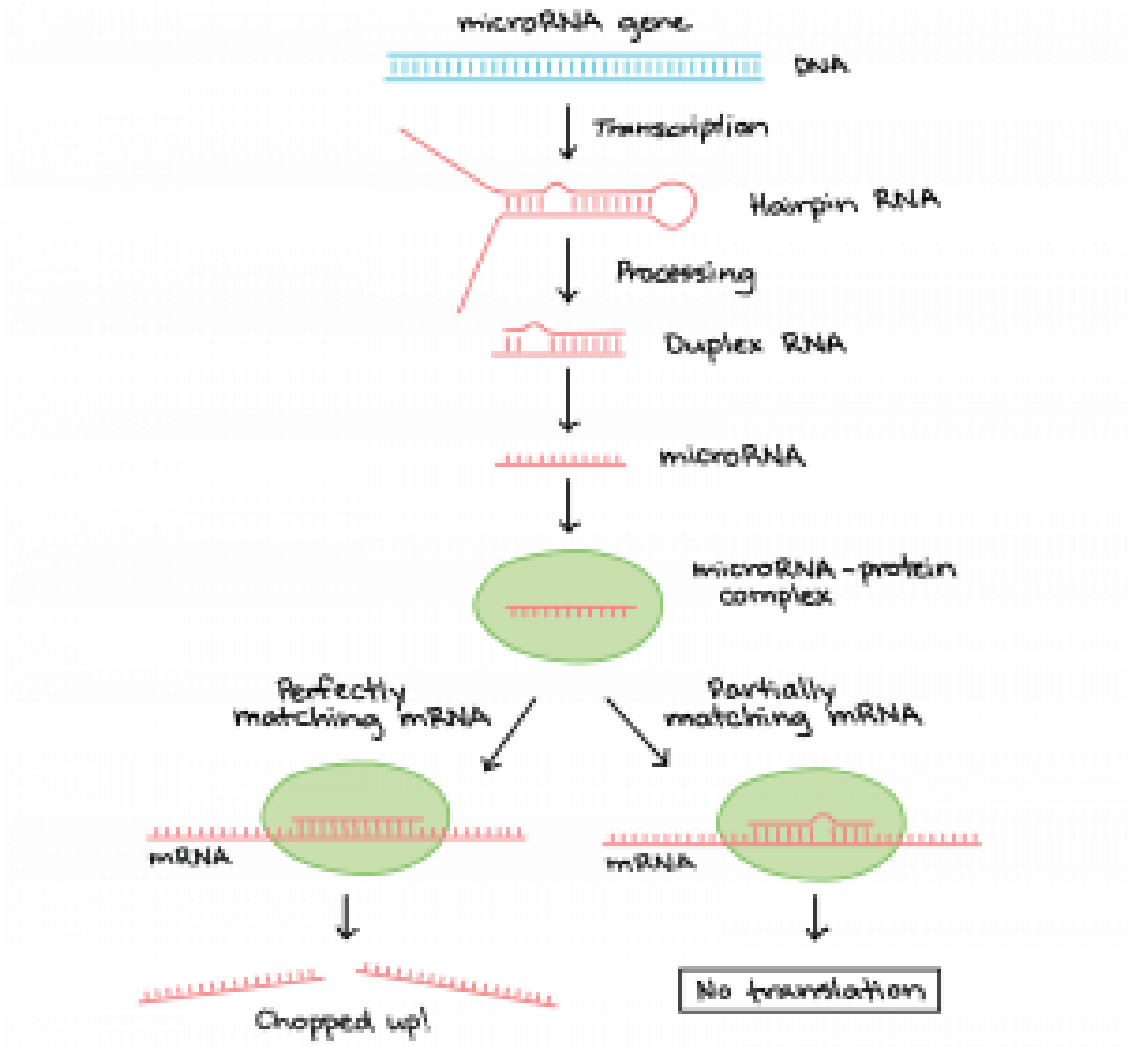
What happens after the microRNA binds to the mRNA?



A microRNA gene is transcribed and processed into a microRNA. The microRNA might perfectly or partially match the mRNA, preventing protein synthesis.

The miRNA directs the protein complex to "matching" mRNA molecules (ones that form base pairs with the miRNA). When the RNA-protein complex binds.[2]

- If the miRNA and its target match perfectly, an enzyme in the RNA-protein complex will typically chop the mRNA in half, leading to its breakdown.
- If the miRNA and its target have some mismatches, the RNA-protein complex may instead bind to the mRNA and keep it from being translated.

These are not the only ways that miRNAs inhibit expression of their targets and scientists are still investigating their many modes of action.[3]

Overview Interactive

For a review of what you learned in this chapter, see HHMI Biointeractive's RNA Interference interactive!

References

1. Carthew, R. W. and Sontheimer, E. J. (2009). Origins and mechanisms of miRNAs and siRNAs. *Cell*, *136*(4), 648. http://dx.doi.org/10.1016/j.cell.2009.01.035.
2. Carthew, R. W. and Sontheimer, E. J. (2009). Origins and mechanisms of miRNAs and siRNAs. *Cell*, *136*(4), 650. http://dx.doi.org/10.1016/j.cell.2009.01.035.
3. Carthew, R. W. and Sontheimer, E. J. (2009). Origins and mechanisms of miRNAs and siRNAs. *Cell*, *136*(4), 652. http://dx.doi.org/10.1016/j.cell.2009.01.035.
4. Sayed, Danish and Abdellatif, Maha. (2011). MicroRNAs in development and disease. *Physiological Reviews* 91(3), 831, 837-839. http://physrev.physiology.org/content/physrev/91/3/827.full.pdf.

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20.

PROTEIN SYNTHESIS IV: TRANSLATION

Andrea Bierema

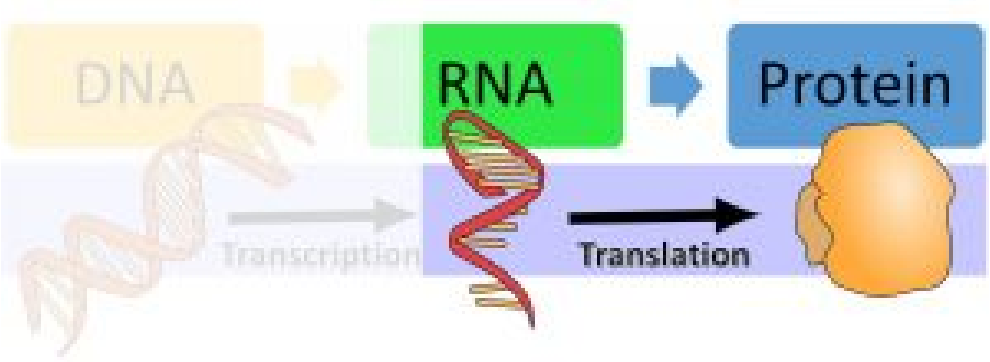
Learning Objectives

Students will be able to:

- Explain how mRNA is translated to synthesize protein.
- Describe the role of the ribosome, the tRNAs, and the mRNA in translation.
- Identify at which point on the mRNA molecule translation begins.
- Explain what happens when a stop codon on the mRNA is reached during translation.

Overview

So far in this book, we have learned how RNA is made, processed, and regulated. This chapter focuses on how translation works; that is, how information coded in the mRNA molecule is read to create an amino acid sequence (i.e., polypeptide), which then folds into a protein. Please see an earlier chapter for a general overview of translation and which codons (i.e., base sequences) code for which amino acids.



Translation is the process of creating a polypeptide (i.e., an amino acid sequence) by "reading" the mRNA sequence and using tRNAs.

The Process of Translation: A First Look

Let's first look at a basic overview of what the process of translation looks like. The video below begins with the mRNA leaving the nucleus and binding with a ribosome. TransferRNAs (tRNAs) move in and out of the ribosome, carrying an amino acid into the ribosome and then leaving without it. An amino acid sequence (i.e., a polypeptide) is produced.

An interactive or media element has been excluded from this version of the text. You can view it online here:
https://openbooks.lib.msu.edu/isb202/?p=280

Exercise

Now that you have watched a basic overview of translation, test your knowledge with the following activity in which you place the following translation steps in the correct order.

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https://openbooks.lib.msu.edu/isb202/?p=280

Molecules

The following video lists the different molecules at play in translation. While watching it, consider how each of these molecules played a role in the first video of this chapter. Watch this before moving on to the mechanism.

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https://openbooks.lib.msu.edu/isb202/?p=280

Mechanism

The first video in this chapter quickly showed what translation looks like. The following video slows down the process and explains in more detail what is happening in the ribosome.

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The Process of Translation: A Detailed Look

This chapter began with an overview of translation and then described in more detail what is happing in the ribosome and how the amino acid chain builds. Now watch the following video, which is an in-depth version of the first video of this chapter, now incorporating aspects described throughout this chapter.

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- Central Dogma (RNA makes protein via translation). © Andrea Bierema

21.

PROTEIN SYNTHESIS V: ADDITIONAL REGULATION

Andrea Bierema

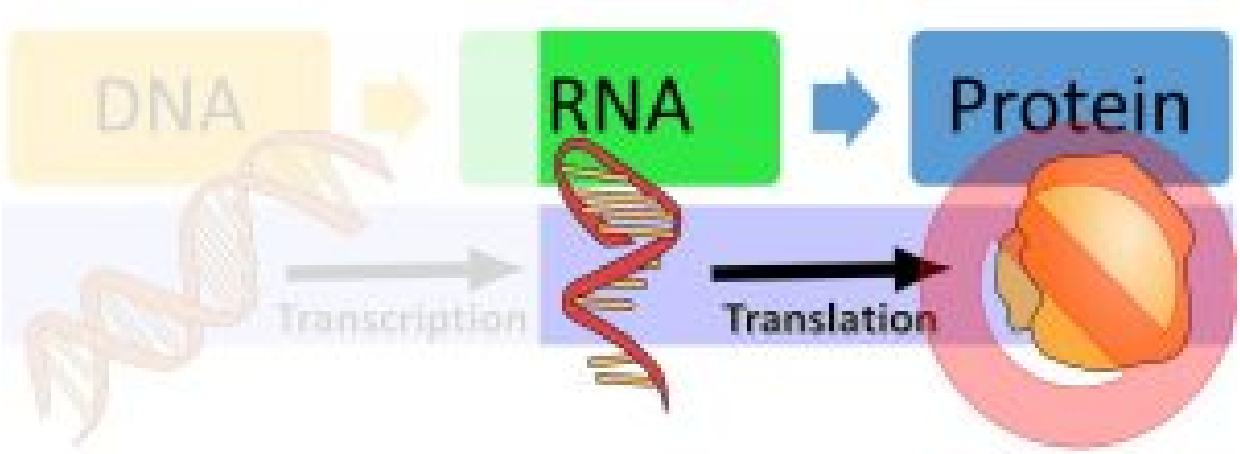
Learning Objective

Students will be able to:

- Identify ways in which gene expression is regulated during and after translation.

Overview

We have learned how we synthesize proteins and how gene expression is regulated before and after transcription, but the regulation can also happen during or after translation.



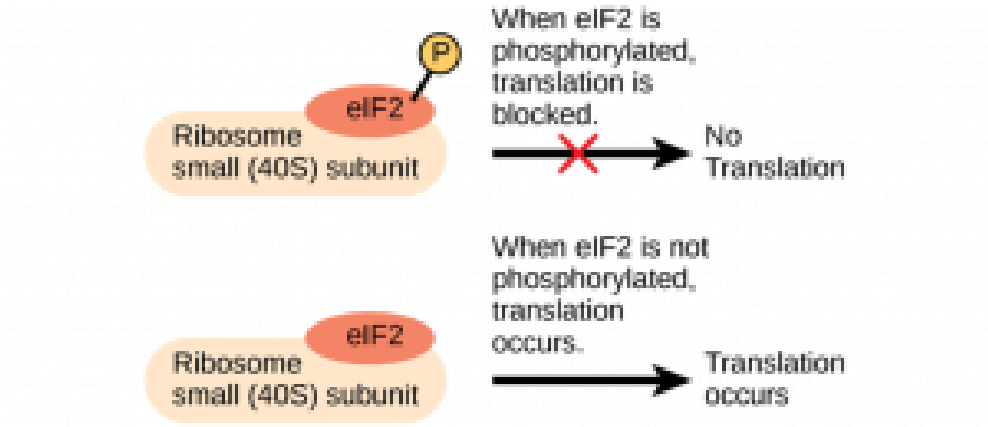
Regulation of Translation

We already saw how miRNAs can inhibit translation, but there are a number of other ways that translation of an mRNA can also be regulated in a cell. One key step for regulation is translation initiation.

In order for translation to begin, the ribosome, an RNA-and-protein complex that houses translation, must assemble on the mRNA. This process involves many "helper" proteins, which make sure the ribosome is correctly positioned. Translation can be regulated globally (for every mRNA in the cell) through changes in the availability or activity of the "helper" proteins.

For example, in order for translation to begin, a protein called eukaryotic initiation factor-2 (eIF-2) must bind to a part of the ribosome called the small subunit. Binding of eIF-2 is controlled by phosphorylation, or addition of a phosphate group to the protein.

When eIF-2 is phosphorylated, it's turned "off"—it undergoes a shape-change and can no longer play its role in initiation, so translation cannot begin. When eIF-2 is not phosphorylated, in contrast, it's "on" and can carry out its role in initiation, allowing translation to proceed.



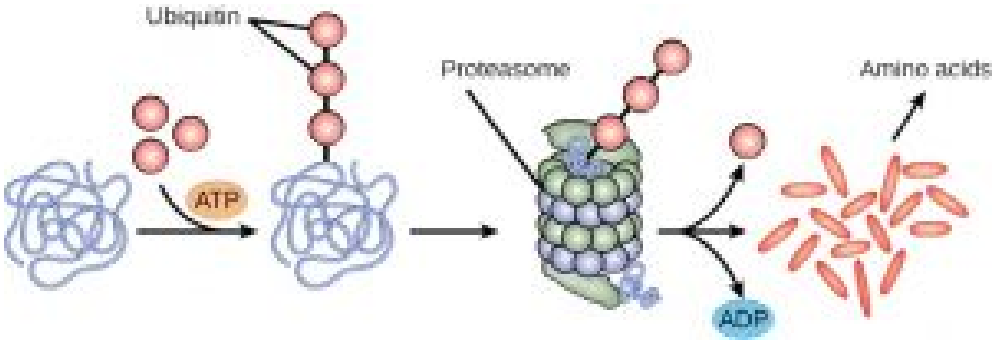
Gene expression can be controlled by factors that bind the translation initiation complex.

Chemical Modifications, Protein Activity, and Longevity

Proteins can be chemically modified with the addition of groups including methyl, phosphate, acetyl, and ubiquitin groups. The addition or removal of these groups from proteins regulates their activity or the length of time they exist in the cell. Sometimes these modifications can regulate where a protein is found in the cell—for example, in the nucleus, in the cytoplasm, or attached to the plasma membrane.

Chemical modifications occur in response to external stimuli such as stress, the lack of nutrients, heat, or ultraviolet light exposure. These changes can alter epigenetic accessibility, transcription, mRNA stability, or translation—all resulting in changes in the expression of various genes. This is an efficient way for the cell to rapidly change the levels of specific proteins in response to the environment. Because proteins are involved in every stage of gene regulation, the phosphorylation of a protein (depending on the protein that is modified) can: alter accessibility to the chromosome; alter translation (by altering transcription factor binding or function); change nuclear shuttling (by influencing modifications to the nuclear pore complex); alter RNA stability (by binding or not binding to the RNA to regulate its stability); modify translation (increase or decrease); or change post-translational modifications (add or remove phosphates or other chemical modifications).

The addition of a ubiquitin group to a protein marks that protein for degradation. Ubiquitin acts like a flag indicating that the protein lifespan is complete. These proteins are moved to the proteasome, an organelle that functions to remove proteins, to be degraded. One way to control gene expression, therefore, is to alter the longevity of the protein.



Proteins with ubiquitin tags are marked for degradation within the proteasome.

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22.

GENETIC ENGINEERING

Andrea Bierema

Learning Objectives

Students will be able to

- Describe examples and mechanisms of genetic engineering.
- Define genetically modified organisms (i.e., GMO).
- Explain how CRISPR is used.
- Describe the role of Cas9 and the guide RNA in CRISPR.
- Identify key steps and their variations in CRISPR.

A variety of biotechnologies exist including cloning organisms, sequencing DNA, and modifying DNA. Although there is a wide range of molecular biotechnologies, this chapter focuses on changes to the genome, which results in changes in proteins or protein synthesis: **genetic engineering**.

Introduction to Genetic Engineering

Genetic engineering refers to the direct manipulation of DNA to alter an organism's characteristics (phenotype) in a particular way. This may mean changing one base pair (A-T or C-G), deleting a whole region of DNA, or introducing an additional copy of a gene. It may also mean extracting DNA from another organism's genome and combining it with the DNA of that individual. It has been used by scientists to enhance or modify the characteristics of an individual organism from a virus to sheep, to possibly humans. For example, genetic engineering can be used to produce plants that have a higher nutritional value or can tolerate exposure to herbicides.

We can change an organism's characteristics by introducing new pieces of DNA into their genomes. This could be:

- DNA from the same species.
- DNA from a different species.
- DNA made synthetically in the lab.

There are several techniques that can be used to modify a genome, including:

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Genetically Modified Organisms (GMOs)

Genetically modified (GM) organisms are organisms that have had their genomes changed in a way that does not happen naturally. By changing an organism's genome, we change the resulting proteins, which change their characteristics. Any organism can be genetically modified, but laws restrict the creation of genetically modified humans, and the production and distribution of other GMOs are tightly regulated.

Exercise

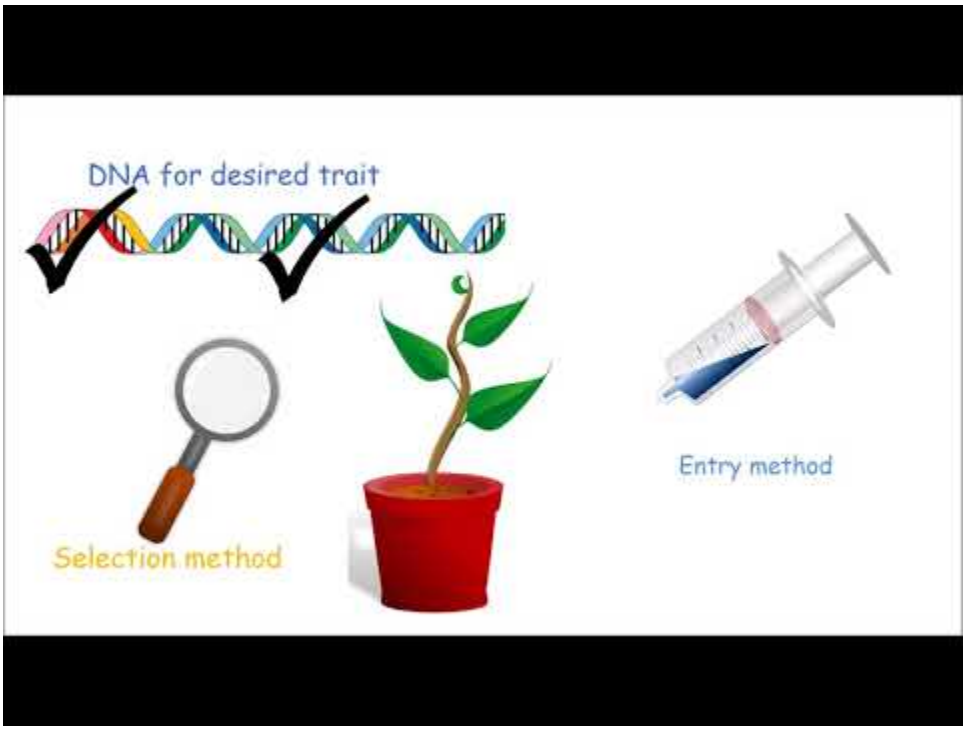
Learn more about the process of creating GMOs by going through *Connecting Concepts'* Interactive lesson on developing a genetically modified canola plant.

The first genetically modified organism was created in 1973 and was a bacterium. Then in 1974, the same techniques were applied to mice. The first genetically-modified foods were made available in 1994.

What is ***not*** a GMO? The genomes of organisms change naturally over time, and these natural changes are not classified as GMO (otherwise, everything would be classified as a GMO). Examples of natural changes include:

- when organisms mate, offspring get bits of DNA from both parents
- mutations arise as a result of mistakes when DNA is copied
- environmental factors like UV radiation can create changes in DNA.

The following video describes how genetically modified plants are made and which qualities may be desired.



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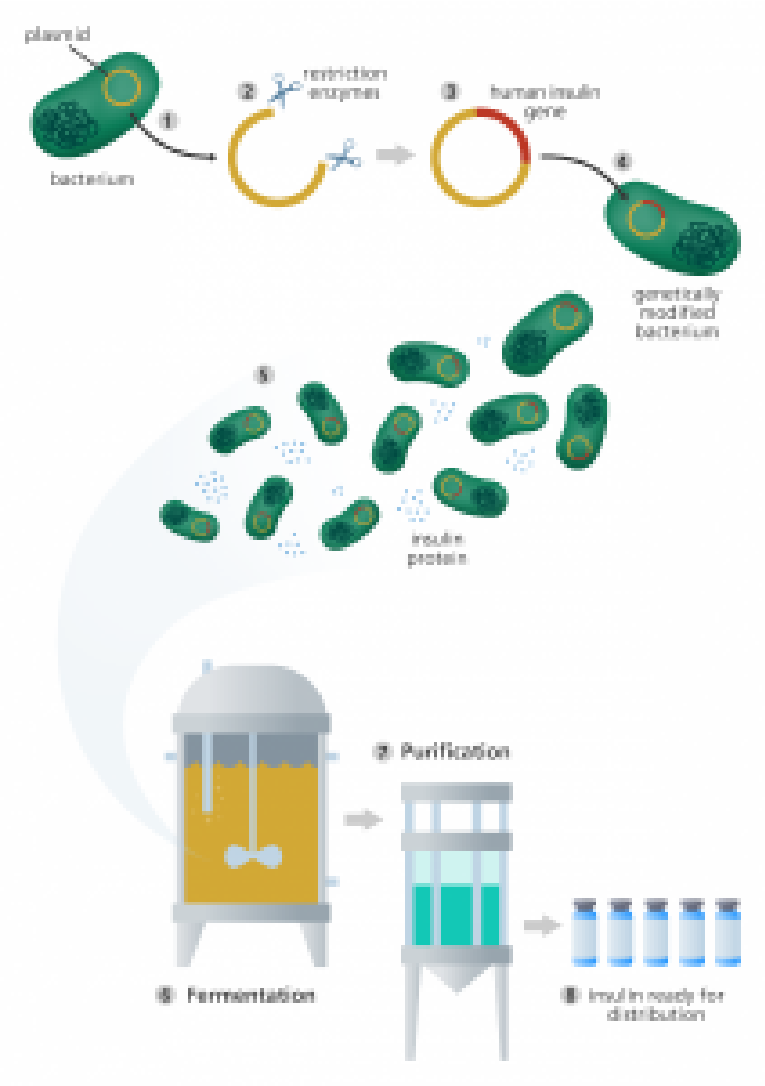
An Example of Genetic Engineering: Insulin Production

Normally, insulin is produced in the pancreas, but in people with type 1 diabetes, there is a problem with insulin production. People with diabetes, therefore, have to inject insulin to control their blood sugar levels. Genetic engineering has been used to produce a type of insulin in yeast and in bacteria like *E. coli* that is very similar to our own. This genetically modified insulin, *Humulin* was licensed for human use in 1982.

To produce genetically-engineered insulin, a small, circular DNA called a plasmid is extracted from the bacteria or yeast cell. A small section is then cut out of the circular plasmid by restriction enzymes that act as "molecular scissors." The gene for human insulin is inserted into the gap in the plasmid, creating a genetically modified plasmid.

This genetically modified plasmid is introduced into a new bacteria or yeast cell. This cell divides rapidly and starts making insulin. To create large amounts of the cells, the genetically modified bacteria or yeast are grown in large fermentation vessels that contain all the nutrients they need. The more the cells divide, the more insulin

is produced. When fermentation is complete, the mixture is filtered to release the insulin. The insulin is then purified and packaged into bottles and insulin pens for distribution to patients with diabetes.



An illustration showing how genetic modification is used to produce insulin in bacteria.

Mosquitos and the Lethal Gene

For another example of genetic engineering, check out *HHMI Biointeractive*'s "Genetically Modified Mosquitos" video.

CRISPR

CRISPR-Cas9 is a genome-editing tool that is creating a buzz in the science world. It is faster, cheaper, and more accurate than previous techniques of editing DNA and has a wide range of potential applications. It can remove, add, and alter sections of the DNA sequence.

The CRISPR-Cas9 system consists of two key molecules that introduce a mutation into the DNA. These are:

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Watch the following videos to learn about the CRISPR-Cas9 process.

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In addition to the videos above, *HHMI Biointeractive* CRISPR Cas-9 Mechanism and Applications interactive is a great illustration!

The following is an overview of the different ways in which DNA is edited:

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Exercise

Check your understanding of the CRISPR process!

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Exercise

Want to learn more about what this looks like in a laboratory? Check out this simulation and this scrolling interactive from LabXchange!

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