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BIOCHEMISTRY

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THE MANGA GUIDE™ TO BIOCHEMISTRY



THE MANGA GUIDE" TO BIOCHEMISTRY

MASAHARU TAKEMURA, KIKUYARO, AND OFFICE SAWA





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PREFACE

This book introduces the world of biochemistry in an approachable comic format. Biochemistry is a synthesis of biology and chemistry, which together elucidate the processes of life at the most basic level. It is the study of the molecules that constitute our bodies and those of other living organisms, and the chemical reactions that occur within cells. In recent years, the field of biochemistry has been growing at an unprecedented rate. From the end of the 19th century and into the 20th, scientists have conducted chemical research on phenomena in the fields of medicine, nutritional science, agriculture, biology, and many other subjects, and this research has led to some incredible discoveries.

When you consider the diversity of the fields listed above, biochemistry may seem like a disjointed collection of different sciences. But even though the objectives differ, the concepts on which they are based are the same—the chemical elucidation of life phenomena. Therefore, the fundamentals of biochemistry must be learned by anyone who intends to participate in any field that deals with the human body or life phenomena to any extent, such as medicine, dentistry, pharmacology, agriculture, nutritional science, and nursing.

This book explains the most important points in biochemistry in an easy-to-understand format. It can be used as a reference book or supplementary reader for a biochemistry course, or for a course in medical science or nutritional science. You can use this book as a quick refresher or to gain a better understanding of this fascinating science. Even a high school student would certainly be able to comprehend this material.

The organization of this book differs somewhat from other existing biochemistry books. For example, although the major cellular chemical components (substances that are present in all living things: saccharides, lipids, nucleic acids, and proteins) are usually described first in an ordinary biochemistry textbook, discussions of each of these substances are incorporated organically, rather than in an independent chapter. I did this because I believe that introducing these substances in context makes them easier to understand and remember.

In addition, I've included information about biochemistry in our everyday lives in Chapter 3 to highlight the significance of biochemistry by showing how it applies to subjects that most people are familiar with.

The protagonist of this book is a high school girl named Kumi who is very concerned with dieting. I chose this story because it relates to my own educational background as a member of a nutritional science division in an agricultural sciences department. These days, when people talk about biochemistry, the discussion often centers around nutrition and health. Many people are concerned with the phenomena that make up *metabolic syndrome*, a general name for the risk factors of an increasingly-common collection of disorders: type 2 diabetes, coronary artery disease, and stroke.

When I was writing this book, I had the entire text checked at both the manuscript and scenario stages by Professor Yukio Furuichi (emeritus professor at Mie University and currently a professor at Nagoya Women's University), whose specialty is lipid biochemistry, and Professor Shonen Yoshida (emeritus professor at Nagoya University and currently a consultant at the Cancer Immunotherapy Center of Nagoya Kyoritsu Hospital), whose specialties are biochemistry and molecular biology. Professor Furuichi provided guidance for my graduate thesis, and Professor Yoshida provided guidance for my PhD thesis. I would like to take this opportunity to express my deep gratitude to both of them for taking time from their busy schedules to proofread this manuscript.

I would also like to take this opportunity to thank Professor Kazuo Kamemura, my mentor during my graduate student days, and his graduate student, Mitsutaka Ogawa, both of Nagahama Institute of Bio-Science and Technology. Specifically, I would like to thank them for the lectin blotting data that they provided. I would also like to thank everyone at the Ohmsha Development Bureau for their ongoing help on my previous work, *The Manga Guide to Molecular Biology*; Sawako Sawada of Office Sawa; the manga artist Kikuyaro, who created the delightful scenario and drawings; and, above all, you for choosing to read this book.

MASAHARU TAKEMURA JANUARY 2009



























1. Cell Structure









2. What Happens Inside a Cell?
















3. A Cell Is the Location of Many Chemical Reactions





















4. Fundamental Biochemistry Knowledge

In this section, we'll explain some technical terms that you need to know to study biochemistry.

CARBON

First, we'll examine an extremely important chemical element in biochemistry—carbon.

Carbon is the element identified by chemical symbol C and possessing the atomic number 6 and an atomic weight of 12.0107. It's the primary component of all known life, which is why people sometimes refer to Earth's organisms as "carbon-based life." Carbon is the backbone of all organic compounds, and the bodies of living organisms are made almost entirely out of these compounds. Carbon is ideal as a backbone for complex organic molecules such as biopolymers, because it forms four stable bonds, which is an unusually high number for an element. Proteins, lipids, saccharides, nucleic acids, and vitamins are all built with carbon as a framework.

Although carbon is common on Earth—in the biosphere, lithosphere, atmosphere, and hydrosphere—there is a finite amount of it, so it's recycled and reused. Over time, a carbon atom passes through air, soil, rocks, and living creatures via biogeochemical cycles. The carbon in your body today may have once been inside a dinosaur!

CHEMICAL BONDS

When carbon combines with other elements, such as oxygen, hydrogen, or nitrogen, different chemical compounds are produced. Except for certain gases, like helium and argon, almost all chemical substances are composed of *molecules*, two or more atoms attached via a *chemical bond*. For example, a water molecule (H_2O) is created when two hydrogen atoms (H) and one oxygen atom (O) join together.

There are several different types of chemical bonds. Some examples include: *covalent bonds*, in which electrons are shared between a pair of atoms, *ionic bonds*, in which oppositely-charged atoms are attracted to one another, and *metallic bonds*, in which a pool of electrons swirl around numerous metal atoms.

The four stable bonds that carbon forms are all covalent bonds.

BIOPOLYMERS

Biopolymers are extremely important molecules to the study of biochemistry.

Biopolymer is a generic term for large, modular organic molecules. Modular means "assembled from repeating units," like the beads of a necklace. Proteins, lipids, nucleic acid, and polysaccharides are all biopolymers. Because they tend to be especially large molecules, biopolymers can form complex structures, which makes them very useful in advanced systems such as cells.

Biopolymers can form these complex chains because they're more than simple beads. Let's consider proteins, for example. Imagine a protein as a necklace made from a variety of different LEGO blocks that can all connect to one another. Since you can twist the necklace easily, it doesn't matter whether the blocks are close together or far apart, but the individual properties of each block result in some connecting better than others. If this necklace was a mile long, imagine the many strange and complex forms you could build. This isn't precisely how proteins function, but you get the idea.

ENZYMES

Since biochemistry explains life from a chemical point of view, it is vital to understand how chemical reactions work, and *enzymes* are essential to these reactions. Enzymes are proteins that act as catalysts—that is, they increase the rate of chemical reactions. An enzyme catalyzes nearly every chemical reaction that occurs in an organism.

In a chemical reaction catalyzed by an enzyme, the substance that the enzyme acts upon is called the *substrate*. The new substance that's formed during the reaction is called the *product*. The activity of an enzyme is affected by the environment inside the organism (temperature, pH, and other factors), the availability of the substrate, and, in some cases, the concentration of the product.

Although almost all enzymes are proteins, it has recently been discovered that a special type of ribonucleic acid (RNA) can act as a catalyst in certain chemical reactions. This is called an RNA enzyme, or a *ribozyme*.

OXIDATION-REDUCTION

Enzymes are broadly classified into six types, which will be introduced in detail in Chapter 4. *Oxidation-reduction* is one of the most important enzyme reactions, in which electrons are exchanged between two substances. If electrons are lost, the substance is *oxidized*, and if electrons are gained, the substance is *reduced*. Normally, when one substance is oxidized, another substance is reduced, so oxidation and reduction are said to occur simultaneously.

The movement of hydrogen ions (H^+ , aka protons) often accompanies the exchange of electrons in an organism, and NADPH, NADH, and similar compounds (which we'll discuss in Chapter 2) work as *reducing agents* on other substances.

RESPIRATION

In Chapter 2, we will examine *respiration*. In the broadest sense, respiration is the process of obtaining energy by breaking down large compounds, but this only gives us a vague sense of the meaning.

More specifically, when respiration occurs, an organic substance (for example, the carbohydrates that make up spaghetti) is broken down into simple, inorganic components, like carbon dioxide (CO_2) and water (H_2O). Energy is produced when electrons are transferred between molecules (oxidation-reduction), along a sort of factory line, until they reach oxygen (O_2). This process is known as *internal respiration* or *cellular respiration*.

The oxygen we mentioned above is very important in respiration. It comes from the air that we breathe, and carbon dioxide is produced as a waste product of cellular respiration. When we use our lungs to inhale oxygen and exhale carbon dioxide, it's known as *external respiration*.

METABOLISM

The processes that alter an organism's chemical substances are called *metabolism*. Broadly speaking, metabolism can be divided into *substance metabolism* and *energy metabolism*. However, since these two types occur together during metabolism, the distinction isn't very clear. In this book, when we refer to metabolism, you may assume that we mean substance metabolism.

Substance metabolism This refers to the changes to substances that occur in an organism, including the chemical reactions that are catalyzed by enzymes. More specifically, a reaction that breaks down a complex substance into simpler substances is called *catabolism*, and, conversely, a reaction that synthesizes a more complex substance is called *anabolism*.

Energy metabolism This refers to the energy that's gained or lost through anabolic and catabolic processes within an organism, including reactions in which the energy created via respiration or photosynthesis is stored as ATP and other high-energy intermediates.



1. Ecosystems and Cycles











Since Kumi eats beef, the carbon can also move from the cow to Kumi.



Mmm...l love a tasty cheeseburger! So carbon really moves around a lot, doesn't it? Is this movement the biogeochemical cycle?



Yep! The entire Earth is carrying out this cycle on a gigantic scale.





Wow! So the rice, potatoes, and apples that I eat were originally related to carbon dixiode that somebody else exhaled.



And it's not just carbon. Hydrogen (H), oxygen (O), nitrogen (N), and sulfur (S) cycle around the Earth as well, going from organism to organism, getting emitted into the atmosphere, dissolved into the ocean, or buried deep underground. When this cycle works smoothly, the ecosystem and the global environment are healthy.

CARBON CYCLE



Next I'll explain the carbon cycle in a little more detail.



Umm, carbon...? I know we've talked a little about it, but I still don't understand what it is.

No problem! Nemoto, can you review some basic facts about carbon for Kumi?



Sure. Carbon is one of the most important chemical elements to living creatures like us.

Carbon is at the center of amino acids, which are the building blocks of proteins. It's also the element that creates the framework of saccharides and lipids, and it's a vital part of DNA.





You're right. It looks like we'd be in real trouble without any C. It really is important!



When carbon leaves an organism's body, it can bond with two oxygen atoms to become carbon dioxide (CO₂) or bond with four hydrogen atoms to become methane (CH₄).





Also, it can accumulate deep underground and, over long periods of time, it will become crude oil, coal, and, in some cases, even diamonds.



Oil! Diamonds! Wow, carbon is really valuable stuff!



That's true, but remember: It's not all about excitement and riches. The way carbon circulates is extremely important. Disrupting this balance could make carbon dioxide concentration on Earth steadily increase...



Which would be a serious problem, wouldn't it?



Oh man, that sounds like it would be a total bummer...



Well, think of it this way: If you can understand the Earth as a single circulating system and your own body as a system as well, then you'll realize the importance of a healthy balance and the dangers of disrupting that balance. This knowledge can lead to insights into the pursuit of health and beauty!



The pursuit of health and beauty? Now you've got my attention!



(Wow, Professor Kurosaka is really good at motivating Kumi...)



Let's look at this figure again.



Now look at @ on the right. This is the flow of carbon in which those saccharides are used by a living creature, and through this process, they become carbon dioxide once more and are returned to the atmosphere via respiration.

Let's talk more about these two flows. Now that we're finished with lunch, we can really sink our teeth into the subject!



Super cool!

2. Let's Talk Photosynthesis

THE IMPORTANCE OF PLANTS

At the base of the ecosystem, plants, algae, and some bacteria supply food to all living creatures. The majority of them use a process called *photosynthesis*, in which saccharides and oxygen (O_2) are created from carbon dioxide by splitting atoms of water, using energy from the Sun. Saccharides are also known as *carbohydrates* and are vital to living creatures like us.



That's why plants are called *producers*. In contrast, we animals are called *consumers*.

Photosynthesis is important not just because it creates saccharides. It also maintains a steady, balanced concentration of carbon dioxide in the atmosphere, and it produces oxygen, which living animals require to survive.



As you can imagine, deforestation by humans greatly reduces the number of "producers," upsetting the delicate balance of the biogeochemical cycle and potentially threatening "consumers" (like us!).

Now, let's take a closer look at the way in which plants use sunlight to create saccharides.



CHLOROPLAST STRUCTURE

This image (courtesy of RoboCat) shows green sacks, called *chloroplasts*, within a plant cell. Inside a chloroplast, structures shaped like very thin pouches are stacked in multiple layers to form peculiar structures. Each of these flat pouches is called a *thylakoid*, and a stack of multiple thylakoids is called a *granum*.



The thylakoid membrane is a bilayer composed primarily of phospholipids, just like in cell membranes.

Now let's look at the surface of the thylakoid membrane.



Do you see the groups of tiny grains that appear to be embedded in the surface of the thylakoid? Each of these is a collection of molecules called *chlorophyll*, as well as various proteins that aid in photosynthesis.

Chlorophyll molecules absorb sunlight. However, they don't absorb the entire spectrum; they reflect the green portion of sunlight back outside, which is why plants appear green to us.




















3. Respiration









SACCHARIDES AND THE "-OSE" SUFFIX

You've probably noticed that most of the saccharides we've discussed so far, like glucose and galactose, end with the suffix "-ose." There are standardized rules for naming saccharides, so these names usually end with "-ose."

For instance, glucose is the saccharide that's the basis of energy production, and it's the sugar referred to when we talk about *blood sugar*. Common table sugar is technically called *sucrose*. Milk contains a saccharide called *milk sugar*, which is known as *lactose*, and the sugar contained in fruit is called *fructose*. It's important to remember that there are several kinds of saccharides in the natural world, and the structures of sucrose, lactose, glucose, galactose, and fructose actually differ somewhat. Also, the starch found in rice, potatoes, and other starchy foods is made from *amylose* and *amylopectin*.

In Chapter 3, we'll examine the structures of these saccharides in detail.

WHY DO MONOSACCHARIDES TAKE A CYCLIC STRUCTURE?

Why do monosaccharides take a cyclic structure more often than an open-chain structure? The secret is in an OH that's bonded to a carbon in the molecule.

Alcohol is a good example of this: All types of alcohol are represented in the form R-OH (where R is the variable group). Alcohol can bind with an aldehyde group or a ketone group to create a substance called *hemiacetal*. Since an OH of a monosaccharide also has this property, the monosaccharide ends up reacting with an aldehyde group or a ketone group within the molecule, and a cyclic structure is formed as the result.













* LACTATE IS CREATED FROM PYRUVATE WHEN OXYGEN ISN'T AVAILABLE.









EACH MOLECULE OF GLUCOSE IS CONVERTED INTO TWO MOLECULES OF PYRUVATE, SO TWO MOLECULES OF ATP, EIGH MOLECULES OF NADH, AND TWO MOLECULES OF FADH₂ ARE CREATED IN THE CITRIC ACID CYCLE.



STAGE 3: MASS PRODUCTION OF ENERGY BY THE ELECTRON TRANSPORT CHAIN





* IN OTHER WORDS, NAD* OR FAD IS "REDUCED," AND NADH OR FADH_ IS CREATED. (SEE PAGE 37 FOR AN EXPLANATION OF REDUCTION.)







78 CHAPTER 2







4. ATP—The Common Currency of Energy

Plants and animals use cellular respiration to turn the sugar created by photosynthesis into potential energy, mainly in the form of adenosine triphosphate (ATP). ATP is often called the "common currency" of energy because it's used by almost every living thing: bacteria, plants—even complex organisms like Tom Cruise. However, individual molecules of ATP are *not* exchanged between organisms, so how does it operate like currency?

As you can see below, ATP has three phosphate groups attached to adenosine. When the outermost phosphate group is detached and becomes adenosine diphosphate (ADP) and inorganic phosphate (Pi), 7.3 kcal (31 kJ) per mole of energy is released. If ATP is hydrolyzed in a test tube, the surrounding water is warmed by this energy, but in an actual cell that energy is used when an enzyme catalyzes a chemical reaction, a muscle moves, or a neural signal is transmitted.



5. Types of Monosaccharides

ALDOSES AND KETOSES

We already saw that one of the basic forms of a saccharide (monosaccharide) has its first carbon forming an aldehyde group (see page 61 for more details). Another form of mono-saccharide has its second carbon forming a ketone group.

Monosaccharides that have an aldehyde group are called *aldoses*, and monosaccharides with a ketone group are called *ketoses*.

Some examples of aldoses are glucose and galactose, and the most well-known ketose is fructose. (For more details about fructose, see Chapter 3.)



PYRANOSE AND FURANOSE

Earlier we learned that when certain monosaccharides, like glucose, take a cyclic structure, they resemble a hexagon. A monosaccharide that takes this form of a six-membered ring, made up of five carbons and one oxygen, is called a *pyranose*. However, there are some cases in which a monosaccharide will resemble a pentagon, made up of four carbons and one oxygen. This is called a *furanose*.

Although glucose normally takes the form of pyranose, in extremely rare cases it becomes furanose. To distinguish the two, the former is called *glucopyranose*, and the latter is called *glucofuranose*.

Fructose can also become a pyranose or furanose when it takes a cyclic structure, and these forms are called *fructopyranose* and *fructofuranose*, respectively.



D-FORM AND L-FORM

Monosaccharides, such as glucose, can exist as *D*-form or *L*-form isomers. All monosaccharides that appear in this book are in D-form.

To tell the difference between the two forms, first find the asymmetric carbon. This is the carbon for which the four bonded substances all differ—in glucose, it's the fifth one. When the OH connected to this asymmetric carbon is on the *right* side in the structural formula of the open-chain form, this monosaccharide is the D-form. When the OH is on the *left* side, it's the L-form. Using this as a basis, the H and OH that are bonded to the carbons at the second through fifth positions of the D-form for glucose are all reversed in the L-form.

The fact that the H and OH are reversed for all carbons from the second to the fifth positions is important. If, for example, only the H and OH at the fourth position of glucose are reversed, a different monosaccharide called galactose is formed (see page 62).

Note that most monosaccharides that exist in the natural world are known to be of the D-form.



6. What Is CoA?

When pyruvate, which is created by glycolysis, enters into the citric acid cycle, it becomes a substance called acetyl-CoA. But what does CoA mean?

CoA stands for *coenzyme* A. Its structural formula is shown below. CoA is a substance in which two phosphates in a row are bonded to the fifth carbon of adenosine triphosphate, and a vitamin called pantothenic acid, as well as 2-mercaptoethylamine, are also bonded. Within the CoA, shown in the shaded part of the figure below, is the *phosphopantetheine group*, which works as a "carrier" because it transports the acetyl group (aka the carbohydrate chain of fatty acids). Acetyl-CoA has an acetyl group bonded at the front of this intimidating-looking molecule.



Another protein that works similarly to CoA is ACP (acylcarrier protein). ACP, which we'll examine more thoroughly in Chapter 3, is also a "carrier." And like CoA, it has a phosphopantetheine group but at a different location. The phosphopantetheine group is bonded to the serine (a type of amino acid) of ACP rather than to adenosine triphosphate.

Since it's called a coenzyme, CoA plays the role of assisting in chemical reactions necessary in the procession of a metabolic pathway.











THERE ARE EXCEPTIONS, THOUGH: SOME GLYCOLIPIDS DC DISSOLVE IN WATER.






FATTY ACIDS



Fatty acids are a source of energy, and they can become phospholipids, which are a raw material for creating cell membranes. If there were no fatty acids, humans could not live.



Wow, really? That's surprising! And I thought they were the enemy.



First, let's look at the structure of fatty acids. Although they can be constructed by connecting a few to several dozen carbon (C) atoms together, the fatty acids that are in our bodies always contain 12 to 20 carbon atoms.





At the farthest end of that long chain (which is called a *hydrocarbon chain*) is a structure called a *carboxyl group* (-COOH).

Since only hydrogen (H) atoms are attached to each of the carbon atoms, the fatty acid does **not** mix easily with water. It lacks hydroxyl groups (OH), which saccharides have (see page 61 for information about saccharides).



Oh, I get it, like oil and water. So tough to mix!



Some fatty acids are made in our bodies. For instance, excess carbohydrates are converted into palmitic acid. Two fatty acids, linoleic and α -linolenic acid, are *essential*, which means that they are necessary for good health and cannot be synthesized by humans. On the other hand, stearic acid and arachidonic acid cannot be synthesized but are not essential to good health either. All these fatty acids contain over 16 Cs!



Yikes! So many Cs...it's like a nightmare report card!

Palmitic acid	CH ₃ (CH ₂) ₁₄ COOH	
Stearic acid	CH ₃ (CH ₂) ₁₆ COOH	
Linoleic acid	CH ₃ (CH ₂) ₄ (CH=CHCH ₂) ₂ (CH ₂) ₆ COOH	When there is a double bond inside the molecule, we write it like this.
lpha-linolenic acid	CH ₃ CH ₂ (CH=CHCH ₂) ₃ (CH ₂) ₆ COOH	
Arachidonic acid	CH ₃ (CH ₂) ₄ (CH=CHCH ₂) ₄ (CH ₂) ₂ COOH	



Certain fatty acids have double bonds between the carbon atoms in the middle of the molecule, as you can see in this figure.





Carbons that are double bonded are called *unsaturated carbons*, and fatty acids that have unsaturated carbons are called *unsaturated fatty acids*.

Unsaturated fatty acids don't solidify as easily, remaining as liquid at lower temperatures than saturated fatty acids, so they are often included as a component of cell membranes (that is, phospholipids) for which flexibility is important.



So... if there are a lot of double bonds, fatty acids are harder to solidify?



That's right. Double bonds can create kinks, which prevent the unsaturated fatty acids from forming a stable solid. This means that **the melting point of fatty acids differs significantly depending on the number of carbon atoms and the number of double bonds between those carbon atoms**.





CHOLESTEROL'S JOB



I understand that cholesterol is a type of steroid, but what exactly **is** a steroid? All I know is that bodybuilders use them to get super buff.



Well, when we talk about "steroids," it often brings to mind images of pharmaceuticals and intimidating pectoral muscles, but there are actually a number of steroids that exist in our bodies normally, like cholesterol.

For example, there is a type of hormone called a *steroid hormone*. The most famous of these are the sex hormones—the indispensible hormones that make males males and females females.



Sex hormones, eh? Whatever they are, I'm just glad they made me a girl. Boys are totally gross!



Um...whatever you say, Kumi.

Anyway, *testosterone*, which is a sex hormone produced mainly in a male's testicles, is actually created with cholesterol as a raw material.

Progesterone, another sex hormone, is produced in a female's ovaries or placenta and is also created from cholesterol.





PROGESTERONE



Vitamin D is also a type of steroid created from cholesterol. Since Vitamin D is produced when ultraviolet rays strike our skin, it's very important for human beings to be exposed to the sun.



In addition, cholesterol plays other very important roles, such as being a raw material of bile acid,* which is required for digestion and the absorption of fat in the small intestine.



Wow! Cholesterol has a ton of important jobs! I had no idea...



That's right. When people who pay attention to their health (like your father) talk about cholesterol, usually a negative image, like arteriosclerosis or obesity, comes to mind, but cholesterol is actually an extremely important substance for our bodies.



Jeez, now I'm really confused. Why does such a vital substance make us think of illness, unhealthiness, or being overweight?



Don't worry, we'll unravel that mystery soon enough...

* Bile acid is created in the liver, stored in the gallbladder, and secreted to the duodenum.









WHAT IS ARTERIOSCLEROSIS?



Now that you have a better understanding of cholesterol, imagine what would happen if the density of LDL (bad) in the blood increases while the density of HDL (good) decreases.



Hmmm, I guess cholesterol would be transported into the body's tissues faster than it could be removed, and it would pile up in the blood vessels.



Right. Cholesterol will accumulate on the walls of blood vessels, the lumens of the blood vessels will get narrower, and the flow of blood will be obstructed. This is called *arteriosclerosis*.



The blood vessels get thicker and harder, and as the symptoms advance, it can lead to sickness or even death.



Cholesterol can lead to **death**?! Not to be taken lightly, I guess...



Let's look at a typical type of arteriosclerosis called *atherosclerosis*. Scientists believe the process goes something like this: First, LDL cholesterol is deposited on the inner wall of a damaged blood vessel. This cholesterol is then eaten by phagocytes,* such as *macrophages*, and bulging cells filled with fat, called *foam cells*, accumulate. When this happens, the smooth muscle cells that create the walls of a blood vessel also end up changing significantly, getting harder and thicker.

* A *phagocyte* is a gluttonous kind of white blood cell that eats just about anything. They're an essential part of the immune system.







2. Biochemistry of Obesity—Why Is Fat Stored?



* MORE ACCURATELY, SINCE SOME OF THESE PARTICLES OF FOOD ARE DIRECTLY EXCRETED AFTER THEY ARE EATEN, INGESTED ENERGY IS THE TOTAL ENERGY THAT IS CREATED FROM THE SACCHARIDES, LIPIDS, AND PROTEINS THAT WERE ACTUALLY ABSORBED BY THE BODY.





ANIMALS PRESERVE FAT



Animals have a mechanism for preserving a fixed level of fat in the body. If they overeat and fat accumulates, that triggers a signal that tells the brain to reduce the amount that is eaten. Conversely, if the amout of fat is reduced, causing the animal to get hungry, it will eat until the fat level returns to its original amount.



Interesting! I guess that instinct must come from animals living in the wild, where life is severe. If it gets too fat, it won't be able to catch food, and if it gets too skinny, it could starve and die.



That's right. For example, the protein hormone *insulin* causes glucose in the blood to be absorbed into muscle or adipose tissue and stored as glycogen or fat. The result is that it lowers the blood sugar level. (Insulin is also used to treat diabetes.)





Oh, so "blood sugar level" is just the concentration of glucose in the blood! And if that glucose is absorbed into muscle or fat, the concentration in the blood will drop.



Also, the cell membranes of nerve cells in the hypothalamus (a special part of the brain) contain a protein called an *insulin receptor*.*

* Insulin receptors actually exist in the cell membranes of many types of cells in the body, not just nerve cells.



So insulin regulates the eating behavior and fat level of an animal via the hypothalamus.





The urge to eat is suppressed by the nervous system when insulin binds with the insulin receptors in the hypothalamus.

The experiments scientists have done to prove this may surprise you—for instance, lab mice that were prevented from producing insulin receptors ended up becoming really obese.



Wow! Instructions go to my brain to keep me from eating too much? That's crazy it's like my own body is playing a trick on me!



There's also a protein called *leptin*, which is only created in adipose tissue, that can notify the brain of an accumulation of fat via leptin receptors in the hypothalamus, much like insulin receptors. This also suppresses hunger.



I see. So both insulin and leptin are important proteins for controlling how much we want to eat.



That's right. And scientists have done experiments with leptin too—mice with deformed leptin genes end up just as tubby as the mice without insulin receptors.



That's awful! I wonder why scientists are always picking on mice, anyway...



If we compare the leptin concentration in the blood of nonobese people versus obese people, they both have a value that is proportional to the amount of fat they carry. The amount of leptin created by people who overeat is no different than that of people who eat normally. Although the creation of leptin is linked with the suppression of eating in nonobese people, this is not the case in obese people. In





What a nightmare! If our insulin or leptin is ineffective, we'll eat and eat, and our appetite will never be satisfied...



There are also a number of other substances that help balance our fat levels and appetite, which we'll talk about next!





112 CHAPTER 3

* HYDROLYSIS IS THE DECOMPOSITION OF A SUBSTANCE BY AN ENZYME USING WATER. FOR DETAILS, SEE PAGE 172.











WHEN FAT IS USED AS AN ENERGY SOURCE



Okay, now I understand how fat is produced, but what I really want to know is how to get rid of the fat I'm producing.



To lose weight, you just have to steadily use up your stored fat.

To do this, you need to remember one important thing: **When both saccharides** and lipids are present, the saccharides will be used as an energy source first.

If you eat foods containing lots of saccharides and lipids, your blood sugar level will rise. The saccharides are used first for energy production, and the lipids are stored in adipose tissue.



However, when the saccharides are used up and your blood sugar level decreases, the stored lipids will gradually start getting metabolized. So when your hungry stomach growls, lipids are actively being consumed.

> ടറ INGR

> > NOW IT'S

FAT

OUR TURN





Hmm, I guess that means I've got to exercise for guite a while before those pounds start to burn off.



Now let's talk about how fat is metabolized.

First, the fat in adipose tissue, **triacylglycerol**, is broken down into **fatty acid** and **glycerol** by an enzyme called *hydrolase*. Hydrolase is a hormone-sensitive lipase—note that this is different from a lipoprotein lipase, which we talked about on page 112.





Fatty acid is released into the blood and transported to the various organs and muscles of the body, where it undergoes a chemical reaction called β -oxidation (that's beta oxidation).







You got it!

First, the fatty acid is activated by the attachment of CoA. This fatty acid-CoA compound is called *acyl-CoA*. A carnitine is then added to form *acyl-carnitine*, which is shuttled into the cell. It then enters the mitochondrion and is broken down into acetvl-CoA.

Fatty acids are long carbon chains (10+), but acetyl-CoA has only two carbon atoms.

The fatty acid is broken down by β -oxidation so that one molecule of acetyl-CoA (two carbon atoms) is detached each time. The process is called β -oxidation because the CoA is attached to the second-to-last carbon atom of the fatty acid (the β carbon).

Ultimately, all carbon atoms in the fatty acid will become acetyl-CoA. The following figure shows the β -oxidation of palmitic acid, which has 16 carbon atoms.



PALMITIC ACID WITH 16 CARBON ATOMS



The $\beta\mbox{-}oxidation$ cycle is repeated seven times to make eight molecules of acetyl-CoA!



Right, and since they're already in the mitochondrion, they can directly enter the citric acid cycle there to create ATP.



So when you're burning fat and dieting, this crazy process is breaking down fatty acids inside your body. Wow!



Earlier, I told you how palmitic acid is produced from malonyl-CoA during fatty acid synthesis. When this palmitic acid is broken down, a total of 129 molecules of ATP are produced through the TCA cycle and electron transport chain.



Wait...if I remember correctly, weren't 38 molecules of ATP the most that could be created from 1 molecule of glucose? 129 is an awful lot, isn't it?





Yup, that's true. Fatty acid is a very efficient storage material.



Hmmm. That must be why it takes so much energy to burn it away! Biochemically speaking, dieting is a lot of work...





BIOCHEMISTRY IN OUR EVERYDAY LIVES 123

3. What Is Blood Type?

BLOOD TYPE



The third mystery is: What is blood type? This should be fun!



Even if we totally avoid the science of blood types, they can still be an interesting way to classify people. For example, in Japan it's common to tell someone's fortune based on their blood type.



That's right! A Type A person is often said to be calm, composed, and serious. Incidentally, I am an unconventional, do-it-my-own-way Type B!



My father is a power-hungry Type O, and my mother is a sensitive and moody Type AB.



Hmm, that makes sense. Children of parents with Type O and Type AB should be either Type A or Type B but not Type O or Type AB. So if you were to have a sibling, he or she would be Type A or Type B.





If I had a younger brother who was Type A, my family would have people of all four blood types.

But it's a little strange, isn't it? Members of one family can have different blood types, but people from completely different families can have the same blood type! What exactly **is** blood type, anyway?

HOW IS BLOOD TYPE DETERMINED?



Kumi, you've probably learned about red blood cells in school, right?



Yeah! They're the cells that give blood its color. They look something like this:





You got it. Blood type is determined by the type of saccharide molecules that protrude from the surface of red blood cells. The surface of many cells, including red blood cells, is covered by a *sugar coating* (*glycocalyx*) made up of saccharides.



Jeez, we're back to saccharides again? Though I do like the sound of this "sugar coating"...



Remember the lipid bilayer of the cell membrane? Proteins are embedded in various spots in that bilayer, and saccharide molecules are often attached to their outer surfaces. Collectively, these saccharide molecules make up the cell's sugar coating.





How strange! They look almost like little sea anemones or something, reaching out from the bottom of the ocean. And they're all over our blood cells?





That's right. There are actually over 100 different antigens that can coat red blood cells and be used to classify blood types, so many different *blood group systems* are possible. The most famous is the *ABO blood group system*, which was discovered in 1900 by the Austrian immunologist Karl Landsteiner.



That's the system that we use now, right? With Type A, Type B, Type AB, and Type O?



Right again! The ABO blood group system is based on **three types of saccharide molecules** present on the surface of red blood cells. Each of these has a structure called a *sugar chain*, which consists of several monosaccharides connected together.





Three types of "anemones" waving around down there, huh?



If we look at the left side of the diagrams below, we can see that the tip for each type of sugar chain is different.





But wait...why does one person become Type A while another becomes Type B? What decides which blood type a person will have?



Well, blood type is determined by a certain gene, and an enzyme is created by that gene! (We'll come back to this later, but you can skip ahead to page 169 for more details, if you'd like.)



Again with the enzymes! You mentioned them when we discussed fat, and now again when we're talking about blood types? They must be really important.



Definitely. We'll have to get the professor to tell us more about enzymes when we see her again later.



Yeah! By the way, is it possible that the differences in these sugar chains could actually be related to a person's personality? Can you really tell a person's fortune from their blood type?



Hmm, good question. I guess if you went out on a limb, you could say that the gene that determines blood type might also have some kind of influence on nerve cells. But claiming that that might actually influence one's personality is a pretty big leap of logic, and there's no scientific evidence to support it.



That's a relief. I'd hate to think that my entire personality is controlled by a bunch of dinky little sugar chains.



I don't care what fortune-tellers say—there's no one in the world with a personality like yours, Kumi...





4. Why Does Fruit Get Sweeter as It Ripens?

WHAT TYPES OF SUGAR ARE IN FRUIT?



Time for the fourth mystery: **Why does fruit get sweeter as it ripens?** Speaking of fruit, we just got some fresh pears at my house...

Mmmm, they are ripe, sweet, and delicious.





Actually, that's true for many different kinds of fruit. Mandarin oranges, grapes, cantaloupes, and watermelons all get tastier as they ripen.



You've got that right. When mandarin oranges are still a little greenish, they can be way too acidic, but ripe ones are really sweet. And the melon that you brought over earlier was in season and perfectly ripe!



Yep, but what does "ripe" mean, biochemically speaking? We know that ripe fruit is sweeter, but why?

The reason is that three types of sugars are contained in large quantities in fruits and berries: sucrose (table sugar), fructose (fruit sugar), and glucose (grape sugar*).



Weird! I would have expected them to have only fruit sugar. We are talking about fruit, after all...

* Despite the name, "grape" sugar is found in many fruits and berries, not just grapes.
MONOSACCHARIDES, OLIGOSACCHARIDES, AND POLYSACCHARIDES



Earlier we said that sucrose, glucose, and fructose all have different structures. (See page 63 for details.)



Right! There are a bunch of different kinds of saccharides, and I love eating all of them.



Well, now it's time to learn more about those different saccharides! The basic unit of saccharides is called a *monosaccharide*, which is formed by at least three carbon atoms connected together.

Glucose and fructose are monosaccharides that consist of six carbon atoms. If two or more monosaccharides are attached, they become an *oligosaccharide*. Although sucrose is an oligosaccharide, since it's formed by connecting only two

monosaccharides, it's called a *disaccharide*. Here's what these saccharides look like:





Oh, so sucrose is made up of one glucose and one fructose connected together!



That's right! There are also saccharides made of many monosaccharides, which form extremely long molecules or complex branching structures. These are called *polysaccharides*.

Can you think of any polysaccharides that we might be familiar with?



Oh! We talked about potatoes and rice earlier, and they were chock-full of saccharides. So...starch is a polysaccharide, right?



That's right. Starch consists of many glucose monosaccharides connected together. Plants store glucose in this form after it's created during photosynthesis.



Our bodies (and those of animals) also contain a "storage material" like starch. It's called glycogen, and it's produced mainly by the liver or muscles by connecting together excess glucose molecules to store them for later.



Oh yeah, I remember you mentioning glycogen earlier, but now it actually makes sense!





Other types of polysaccharides include cellulose and chitin. Cellulose is the main component of plant cell walls, and chitin is the main component of the hard shells of crustaceans, like shrimp and crabs. Mushrooms also use chitin as a structural material.

HOW FRUITS BECOME SWEET



Now, let's get back to fruit. Fruits, such as mandarin oranges and melons, get sweeter and more delicious as they ripen. Why is that?



Hmm. When I was shopping for strawberries and mandarin oranges at the supermarket earlier, there was a sign that said, "Sugar Content: 11–12%."

I suppose if fruit gets sweeter as it ripens, that just means that the sugar content has increased, right? Biochemically speaking, we must be talking about a change in the saccharides.



You got it. So...let's talk about saccharides!

Take a look at the graphs below. Before they're ripe, citrus fruits, such as mandarin oranges, contain roughly an equal amount of glucose, fructose, and sucrose. But as they ripen, the relative amount of sucrose steadily increases. In a Japanese pear, all three types increase.



Source: Saburo Ito, Editor, Science of Fruit, Asakura Publishing Co., Ltd. (1991)



When fruit ripens, polysaccharides, like starch, are broken down into monosaccharides, like glucose, and the activity of *sucrose-phosphate synthase*, an enzyme that synthesizes sucrose in fruit, increases while the activity of *invertase*, which breaks down sucrose, decreases.







* ADDITIONALLY, SUBSTANCES SUCH AS SORBITOL, XYLOSE, AND ORGANIC ACIDS ARE ALSO RELATED TO THE SWEETNESS OR ACIDITY OF FRUIT.

5. Why Are Mochi Rice Cakes Springy?

DIFFERENCES BETWEEN NORMAL RICE AND MOCHI RICE



The final mystery is: Why are mochi rice cakes springy? I love mochi, so this is my kind of

number of the second se



First, let's talk about how mochi is made. Did you know that mochi is made from special rice, which is springier than normal rice?



Of course I know that! I've got serious mochi-making experience. I've even participated in a traditional "mochipounding" ceremony!

But what makes mochi rice springier than normal rice?





It's all because of the structural differences of the starches that make up the rice. Rice is 75% starch, so slight variances in its starch composition can seriously affect its physical properties.

As you can see in the figure below, "normal," non-glutinous rice contains the two starches amylose and amylopectin. Amylose makes up about 17 to 22% of the starch, and the rest is amylopectin.





Okay, the graph makes perfect sense, but what does that actually mean?



Just hang on, it should become clear in a second. So...amylose and amylopectin are both made up of glucose molecules connected together to form polysaccharides.





In other words, they're both made up of the same raw materials!



That's right. The only real difference is the way that their glucose molecules are connected together.

So the "mystery of springy mochi" is **actually** the "mystery of monosaccharides connecting in different ways to form polysaccharides and oligosaccharides." I guess the name's not quite as catchy, but it's still a very interesting subject.



Aha! So the secret of mochi's springiness is hidden in how the glucose connects together!

THE DIFFERENCE BETWEEN AMYLOSE AND AMYLOPECTIN



So we know that the difference between these two starches is the way in which their glucose molecules are connected. Specifically, the shape of amylose is **straight**, and the shape of amylopectin is **branched**.



Straight? Branched? I have no idea what you're talking about...



Amylose is formed when glucose molecules are connected in a straight line via a method called the $\alpha(1\rightarrow 4)$ glycosidic bond.





Oh, the glucose molecules are lined up in a row. That makes sense.



However, in amylopectin, there are places where glucose molecules are connected via another process called the $\alpha(1\rightarrow 6)$ glycosidic bond.





Oh, I see. That bond connects the molecules vertically! That must be what causes amylopectin to branch off in different directions.



You got it, Kumi. These two types of bonds are what give amylopectin its more complex, branched structure and amylose its straightforward, linked structure.



I totally understand! Man, I thought this stuff was supposed to be difficult.



Because of this branched structure, when amylopectin is viewed from a distance, it has a distinct "fringed" shape.



Mochi rice, which is mostly made up of amylopectin, becomes very viscous when cooked because the branches stick together. In other words, it feels very springy and stretchy.



So since this "fringed" kind of starch is more elastic than the straight kind, mochi made with this special rice is springy and delicious!



By the way, even in normal, non-glutinous rice, the stickiness varies according to the amount of amylose it contains.

WHAT DO THE NUMBERS MEAN IN α (1 \rightarrow 4) AND α (1 \rightarrow 6)?



Since we've come this far, let's take this opportunity to learn what the numbers mean in the two types of bonds we discussed: $\alpha(1\rightarrow 4)$ and $\alpha(1\rightarrow 6)$.



Yeah, I was wondering about that, but I was kind of afraid to ask...



This may seem a little out of the blue, but...do you like baseball?



Um, I guess so. My dad is a big fan, so I see a lot of games on TV.



Numbering the bases on the field makes the game much easier to understand, right?



If they weren't numbered, it would make describing a game a lot trickier. For instance, we'd have to call third base "the base to the catcher's left." And I don't even want to **think** about how we'd talk about a triple play!



It would make the announcer's life difficult, that's for sure.



We learned that glucose and fructose have six carbon atoms each, right? Numbers have been assigned to each of those atoms just like the bases in baseball!



Look carefully at the carbon atoms (C) in the following figure. This is the ring structure of glucose, and the numbers 1 through 6 have been assigned as shown.





Wait, I think I get it! The numbers in the $\alpha(1\rightarrow 4)$ glycosidic bond and the $\alpha(1\rightarrow 6)$ glycosidic bond refer to those numbers!



Right you are! The $\alpha(1 \rightarrow 4)$ glycosidic bond means that the first carbon of one glucose is attached via a glycosidic bond to the fourth carbon of the neighboring glucose.*





Okay, so the $\alpha(1\rightarrow 6)$ glycosidic bond means that the first carbon of one glucose is attached to the sixth carbon of the next glucose in exactly the same way, right?



Yup, but when the carbon atoms at the first and sixth positions are attached, the glucose molecules can't be side-by-side in a straight line.

* The glucose molecules are simplified throughout this section in order to highlight the relevant bonds.



They can connect only as you see in this diagram—vertically, rather than horizontally.





Oh! So that's why a branch happens at that point, and that's what makes it springy. We're really getting to the bottom of things! \varGamma



By the way, there's another type of bond that connects carbons in glucose. It's called the $\beta(1\rightarrow 4)$ glycosidic bond.



Beta? What's the difference?



Beta means no starch! When glucose molecules are connected via $\beta(1 \rightarrow 4)$ glycosidic bonds, the polysaccharide *cellulose* is formed rather than starch. This is the main component for creating cell walls in plants, and it's also a type of dietary fiber.



Oh yeah! I read about dietary fiber in the latest issue of *Dieter's Digest*. It's difficult to digest, so it just passes right through the body. Heh heh.



That's correct. But dietary fiber also includes substances that easily dissolve in water, such as hemicellulose or pectin, and these are easily digested.



Weird! Are they energy sources for us too?



They are, but let's get back to cellulose, okay? There's an enzyme contained in our saliva called α -amylase, which can break down starches, like rice, into pieces, but α -amylase cannot break down cellulose.



Why not?



Well, look at the $\beta(1\rightarrow 4)$ glycosidic bond shown in the following figure.



 β (1) \rightarrow 4) GLYCOSIDIC BOND



Hey, the parts that are connected are different, aren't they? They have a strange shape, almost like the letter N.



The $\beta(1\rightarrow 4)$ glycosidic bond differs from the $\alpha(1\rightarrow 4)$ glycosidic bond in that the positions of the hydrogen atom (H) and the hydroxyl group (OH) are flipped around their carbon. This creates a bond that is N-shaped rather than U-shaped, like the $\alpha(1\rightarrow 4)$ glycosidic bond.



Yeah, no offense to the β type, but its connection seems twisted and totally weird!





The Greek letters α and β represent the position of the hydroxyl group (OH) on carbon 1. When OH is on the bottom, as shown on the left in the above figure, it is the α type. When it's at the top, it is the β type.



Because of this difference in structure, α -amylase, which can break down $\alpha(1\rightarrow 4)$ glycosidic bonds, is unable to break down $\beta(1\rightarrow 4)$ glycosidic bonds.





Wow! That twisted connection really makes a big difference!



So cellulose, which is formed by $\beta(1 \rightarrow 4)$ glycosidic bonds, is not broken down in our digestive system. This makes it very effective as dietary fiber. It...you know... *keeps you regular.*

Well. Moving right along! Remember when we were talking about sucrose in fruit? We discovered that sucrose is made up of one glucose and one fructose connected together. The carbon at position 1 of glucose is connected to the carbon at position 2 of fructose, as you can see below.





I get it! So sucrose is formed with an $\alpha(1\rightarrow 2)$ glycosidic bond, right?



Exactly! And now that you know how individual monosaccharides are connected together, you should have a much better understanding of why these substances have their unique physical properties.























AMINO ACIDS





PRIMARY STRUCTURE OF A PROTEIN

First, a long amino acid chain is formed by connecting amino acids one at a time with *peptide bonds*. Remember earlier when we said that amino acids are joined together via chemical reactions to form proteins? This chemical reaction, which creates the peptide bonds, is called the *peptidyl transfer*.

This reaction occurs inside *ribosomes*, which are protein synthesis apparatuses contained in the cytoplasm and attached to the rough endoplasmic reticulum. We'll talk more about ribosomes in Chapter 5.

The long amino acid chain connected together by peptide bonds is called a *polypeptide chain*. This amino acid string is called the *primary structure* of the protein.



SECONDARY STRUCTURE OF A PROTEIN

Each of the 20 amino acids has a characteristic part (indicated by [R] on page 155) that is unique. This is called the *side chain*.

Since a number of difference forces can act on these side chains, such as hydrogen bonds and hydrophobic and electrostatic interactions, neighboring amino acids are attracted to or repelled by each other in particular ways, which results in a characteristic, local threedimensional structure. This is called the *secondary structure* of the protein.

There are a number of different secondary structures, such as an α -helix, in which a part of the polypeptide chain forms a spiral, and a β -sheet, which takes a folded planar shape.





TERTIARY STRUCTURE OF A PROTEIN

Even when the polypeptide chain takes its secondary structure, it's not yet a fully folded, functional protein.

To become functional, the polypeptide chain has to take on a specific three-dimensional shape, which is determined by the interactions of the amino acid side chains. This shape is called the *tertiary structure* of the protein.

For example, myoglobin, which is shown in the following figure, is a type of protein that exists in the muscles of animals. This protein is formed from eight α -helices surrounding an iron-containing *heme*, which binds oxygen.



MYOGLOBIN CONSISTS OF THE EIGHT $\alpha\text{-Helices}$ indicated by the letters a through H.

TERTIARY STRUCTURE

QUATERNARY STRUCTURE OF A PROTEIN AND SUBUNITS

Many proteins are able to operate as a protein or as an enzyme at the tertiary structure stage. However, some proteins create an aggregation in which multiple polypeptide chains that have taken tertiary structures are assembled together into a large functional unit.

For example, our red blood cells contain many iron-binding proteins called *hemoglobin*, which are used to transport oxygen. Hemoglobin is formed by assembling four polypeptide chains called *globin* (two each of two types, α and β , which are indicated below as α_1 , α_2 , β_1 , and β_2). An enzyme such as RNA polymerase II, which creates RNA in our cells, is formed by assembling 12 polypeptide chains.

This state is called the *quaternary structure*, and each of the polypeptide chains used to create the quaternary structure is called a *subunit*.



SINCE THE SUBUNITS OF HEMOGLOBIN ARE VERY SIMILAR IN STRUCTURE TO THOSE IN MYOGLOBIN, HEMOGLOBIN IN THIS FIGURE HAS BEEN DRAWN WITH THE SAME STRUCTURE AS ON PAGE 160. HOWEVER, THE STRUCTURES ARE ACTUALLY SOMEWHAT DIFFERENT.

QUATERNARY STRUCTURE

2. An Enzyme's Job





STRICT ENZYME? RELAXED ENZYME?





Some enzymes are "strict," which means they only act on very specific substrates. Other enzymes are more "relaxed" and act on a broad range of substrates.



Strict and relaxed? Sounds like the difference between my mom and my dad...



There are enzymes that can act on substances that are similar or closely related to their substrates. Many examples of these enzymes are seen in the digestive system—for example, the *protein catabolism enzymes*.



Remember, there are a huge number of different proteins. If enzymes were *too* specific, there would have to be a separate enzyme to break down every single kind of protein! Things would get pretty unwieldy.





Since proteins are so complex, the enzymes that break them down had to become a bit more flexible.



That's right! Protein catabolism enzymes (the enzymes that break down proteins) often have a certain degree of leeway in the substrates they can interact with.



For example, one of the protein catabolism enzymes secreted from the pancreas, carboxypeptidase, detaches amino acids sequentially from the end of a protein.

Carboxypeptidase comes in various types, including A, B, C, and Y. Carboxypeptidase A, for example, can detach almost any amino acid from the C-terminal end of a protein, but it doesn't work well on amino acids with bulky or aromatic R-groups, like arginine, lysine, and proline.





There are 20 types of amino acids that make up proteins, right? So even though carboxypeptidase A can't deal with arginine, lysine, and proline, there are still 17 types it *can* deal with. It seems pretty flexible!

That's right. It has some serious leeway in the substrates it can interact with. However, there are also some "strict" protein catabolism enzymes. For example, trypsin cuts only through the C-terminal side of arginine and lysine.





* A NEWLY DISCOVERED ENZYME IS ASSIGNED AN EC NUMBER AND SYSTEM NAME AS WELL AS A RECOMMENDED NAME ACCORDING TO STANDARDS DETERMINED BY THE ENZYME COMMISSION OF THE INTERNATIONAL UNION OF BIOCHEMISTRY AND MOLECULAR BIOLOGY (IUBMB).




GLUCOSYLTRANSFERASE DETERMINES BLOOD TYPE



Remember when we solved the blood type mystery (in Chapter 3)? When we were trying to figure out what determines blood type, we discovered that it was the work of an enzyme.



I remember!

The ABO blood group system is classified according to the differences between the "sugar chains" on the surface of red blood cells. A sugar chain is a collection of monosaccharides connected together!



And the differences between those chains are the monosaccharides at the tips, right?



Yeah! For people with Type A blood, that monosaccharide is *N*-acetylgalactosamine (GalNAc); for people with Type B blood, it's galactose (Gal); and for people with Type O blood, there is no monosaccharide at the tip.

(After writing such a detailed report for the professor, I'll probably remember that for the rest of my life...)



That's correct. The particular monosaccharide that's attached to the tip of a sugar chain (or the lack of one) is determined by a certain *gene*.

Genes are like the "blueprints" for proteins. (Remember: An enzyme is a type of protein.)

So what really determines blood type is the gene that creates *glycosyltransferase*, which is the enzyme that attaches a particular monosaccharide to the tip of a sugar chain found on the surface of the red blood cell.



So glycosyltransferase attaches a certain monosaccharide, and the type of that monosaccharide determines blood type?



That's right. Look again at the structure of the sugar chain of each blood type.





Let's review it again. The only difference in these three types is the monosaccharide at the very tip.

For people with **Type A** blood, it's *N*-acetylgalactosamine. For people with **Type B** blood, it's galactose. For people with **Type O** blood, it's nothing!



Yeah, yeah, I've got it.



So, the sugar chain that Type O people have is the "prototype," or the minimal sugar chain. People with A and B types have the O type chain plus something extra. If someone has the transferase gene that attaches *N*-acetylgalactosamine,

his or her blood type will be Type A! But if someone has the transferase gene that attaches galactose, his or her blood type will be Type B!





That all makes sense, but I still don't quite understand what's going on with people with Type O blood. Why don't they get monosaccharides on the end of their sugar chains? Seems unfair!



Type 0 people also have the genes associated with glycosyltransferases, but since there is no enzyme activity for the proteins created by those genes, no saccharide is attached at the tips. The activity may have been lost because of a genetic mutation during the evolutionary process.



A mutation? Cool!



So to wrap it all up: The ABO blood group system is nothing but the result of differences in glycosyltransferase genes.



Nemoto, you must have a genetic mutation that gives you an oversized brain. How else could you possibly know so much?!



Gyuh...



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3. Using Graphs to Understand Enzymes



WHY ARE ENZYMES IMPORTANT FOR CHEMICAL REACTIONS?

A substance that expedites a chemical reaction is called a *catalyst*. An enzyme, which is a type of catalyst, is also known as a *biological catalyst*.

Although an enzyme is required to make a chemical reaction advance efficiently, it is not neccesarily required for the chemical reaction to occur.

Most chemical reactions in the body will occur eventually even without enzymes, but many would take an overwhelmingly long time. Complex chemical reactions, like glycolysis, may never reach completion without a catalyst.

Enzymes are important because, for an organism that must maintain specific conditions and produce energy to stay alive, the chemical reactions that occur inside its body must be as efficient as possible. For living organisms as a whole, it would be disastrous if reactions took too long.



Now we'll use some simple graphs and formulas to study the essential qualities of chemical reactions that rely on enzymes, and we'll learn why enzymes are so meaningful to those reactions.









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You're right. As the substrate concentration increases, the reaction rate approaches its *limit*, V_{max} , which is the maximum rate.

If we keep taking measurements like this, we'll be able to figure out V_{max} , 1/2 V_{max} , and K_m , right?



In theory, that's true—we could just take a bunch of measurements, and eventually we'd approach V_{max} , but in practice, it's a bit more difficult than that.



Huh? But why?





But we can't make calculations using infinity, right? So what can we do?



It's no problem. We'll just use a little trick. We can take advantage of the fact that **the reciprocal of infinity is zero**. In other words, when we take the reciprocal, we know that the *y*-axis value is V_{max} when the *x*-axis is zero!

Although, actually, the numerical value of the *y*-axis at this time is $\frac{1}{V_{max}}$.

It looks a little something like this:



 V_{max} V_{max} V_{max}

Since infinity is too abstract for our calculations, we need to replace it with something that makes sense in the real world, so we just take the reciprocals of the values for both the x- and y-axes.

Step 2

When the *x*-axis value is $\frac{1}{\infty}$, the *y*-axis value is $\frac{1}{V_{\text{max}}}$.

Luckily, we know that the reciprocal of infinity is zero.

Step 3

When the x-axis value is 0, the y-axis value is $\frac{1}{V}$.

If we create a graph for this, we can get a definite numerical value for V_{max} .



I can't believe it, but...that actually makes sense! Maybe graphs aren't quite as evil as I thought they were...



Now that you understand the theory behind the calculations, don't you feel like your life is richer? Like you understand the universe just a little bit better? Now let's return to the straight line graph you created earlier and try to obtain the value of V_{max} .



Yeah, let's do it! Graphs and reciprocals are child's play for the likes of me!



** REMEMBER, THE DNA POLYMERASE REACTION TIME WAS 60 MINUTES.

LET'S USE THE MICHAELIS-MENTEN EQUATION TO FIGURE OUT KM AND VMAX!





In other words, we can draw the graph of the straight line y = ax + b, where $y : \frac{1}{y}$





4. Enzymes and Inhibitors

Why in the world do we have to use equations and graphs—which are no fun at all—to get the values of V_{max} and K_m ?

One reason is to demonstrate that enzyme reactions are extremely precise and predictable. They proceed in strict accordance with chemical and mathematical laws.

More importantly, for scholars doing enzyme research or research concerning the world surrounding enzymes, the values of V_{max} and K_m are very useful.

One of these areas of research deals with the relationship between enzymes and inhibitors. An *inhibitor* is a substance that affects the binding of an enzyme and substrate, or affects the enzyme itself. As a result, it inhibits the enzyme's activity.

Many inhibitors are created artificially and used for enzyme research. Since they inhibit enzymes, they are often toxic to living organisms. However, they can also be used in positive ways—to kill cancer cells, for example.

Inhibitors also exist in the natural world. In that case, they're known as "cellular" enzyme inhibitors rather than "medicinal" inhibitors. For example, some inhibitors are created directly in an organism's cells and play important roles in regulating enzyme reactions. Plants can create inhibitors as well. The seeds of certain legumes, for example, contain cellular enzyme inhibitors such as α -amylase inhibitors or trypsin inhibitors. These are known as *anti-nutritional factors*. Scientists believe that these inhibitors might be part of the plant's natural defense system, which helps protect them against predation.

If the structure of an inhibitor is similar to that of the substrate, it can bond quite easily with the enzyme. However, since the shape of an inhibitor is subtly different from that of the substrate, no reaction will occur, and once the enzyme has already bonded with another substance, it isn't able to bond with its substrate. Actually, there are several different types of enzyme inhibition, and we can tell which is occuring by using the Michaelis-Menten equation.

Two of the types of enzyme inhibition are competitive inhibition and non-competitive inhibition.

Competitive inhibition occurs when a substance that is very similar to the substrate bonds with the enzyme, inhibiting the enzyme reaction.



* THE ACTIVE SITE IS THE PART WHERE THE SUBSTRATE BONDS TO THE ENZYME IN ORDER TO UNDERGO A CATALYTIC ACTION. Since the enzymes remain active, the maximum reaction rate V_{max} is not affected, but the K_m value will rise, since some of the enzymes accidentally bind to inhibitors. With inhibitors getting in the way, it takes longer for enzymes to bind to substrates, producing the same result as if a lower substrate concentration was present.

If a competitive inhibitor is added, the slope of the straight line will be greater on a Lineweaver-Burk reciprocal plot, as shown below. Note that the intersection with the *y*-axis, $1/V_{max}$ doesn't change.



Although the intersection with the *y*-axis doesn't change, the intersection with the *x*-axis does. This means that if you add an unknown inhibitor to an enzyme reaction, take measurements, graph the result, and see that $1/K_m$ has increased, you know it's a competitive inhibitor.

Non-competitive inhibition is when an inhibitor bonds at a part of the enzyme unrelated to the substrate bonding site to inhibit the enzyme reaction.



In this case, the inhibitor doesn't block the substrate—it takes the enzymes out of the game. K_m is unaffected because it depends on substrate concentration. Instead, the maximum reaction rate will steadily decrease as the amount of inhibitor increases. The same thing would happen if the enzyme concentration decreased.

This means that when a non-competitive inhibitor is added, the slope of the straight line will again be greater on a Lineweaver-Burk reciprocal plot, as shown below. But unlike competitive inhibition, the intersection with the *y*-axis will change, while the intersection with the *x*-axis will stay the same.



Since inhibitors significantly affect enzyme reactions, they are often used in research for elucidating enzyme structures or reaction mechanisms. They are also used in cancer research by applying small amounts to inhibit enzymes that benefit the cancer cells.



ALLOSTERIC ENZYMES

Although we've introduced enzyme reactions for which the Michaelis-Menten equation applies, there are also many enzymes that exhibit activity for which the Michaelis-Menten equation does not apply.

One of these is a type of enzyme called an *allosteric enzyme*, which can respond to environmental signals with changes in activity. This is referred to as an *allosteric effect*. For example, a substrate that bonds to one subunit may bring about a change in the three-dimensional structure of the enzyme, which causes the substrate to bond more easily with another subunit.

In a case like that, the curve representing the relationship between the substrate concentration and the reaction speed will be an S-shaped *sigmoid curve* rather than the hyperbola that is typical of the Michaelis-Menten equation.

We'll skip the details here, but not all enzyme reactions are simple ones that occur according to certain fixed equations—in fact, in the body, most are extremely complex and diverse.









MOLECULAR BIOLOGY AND THE BIOCHEMISTRY OF NUCLEIC ACIDS







1. What Is Nucleic Acid?



* GENES ALSO CONTAIN THE "BLUEPRINTS" FOR RNA.




THE DISCOVERY OF NUCLEIN

The Swiss biochemist Friedrich Miescher (1844–1895) successfully isolated a new substance from a sample of white blood cells. He extracted the cells from used bandages that he received from a nearby hospital.



White blood cells adhering to used bandages? Ugh, so we must be talking about...

That's right! We're talking about **pus**!



Friedrich Miescher



He studied pus?! Gross! I guess biochemistry isn't all glamour and diets...



To extract white blood cells, Miescher processed the pus by adding an enzyme called *protease*, which breaks down protein, and then he used a method called *ether extraction* to remove the lipids.





The substance that was obtained from the white blood cells showed strong acidity. Miescher named it *nuclein* since it was found inside the **nucleus** of the white blood cells. Miescher also succeeded in extracting nuclein from salmon sperm. Nuclein later became known as *nucleic acid*, and in the first half of the 20th century, scientists determined that there two types of nucleic acid: DNA and RNA.













Right. In other words, a "pair" is created via hydrogen bonds between two bases, so that the polynucleotides form a double strand. The pairings are always faithful: A pairs with T (with two hydrogen bonds), and G pairs with C (with three hydrogen bonds).





This pairing configuration is called *complementarity*. Because of this complementarity, DNA can be *replicated*—in other words, the two-stranded polynucleotide chain can be separated into individual strands, and those strands can be used as *templates* to build new strands. The result of this replication is two DNA molecules that have the same arrangement of bases.





A template is like the mold that you pour batter into to make cupcakes.



I get it! If you have a mold, they're easy to replicate.

If you know what the sequence of the template strand is, you'll know the sequence of a new strand that uses that tempate as a mold!



That's right. The biggest difference between DNA and other biopolymers (proteins, saccharides, and lipids) is that ability to replicate easily.

DNA REPLICATION AND THE ENZYME DNA POLYMERASE



So why is DNA replicated? Since DNA contains our genes, it must be replicated whenever cells divide and multiply.

Genes are passed on from parents to children, as well as from cell to cell. When a cell divides, the same genes must be inherited by the two new cells, so the DNA must be replicated before the cell splits.



CLEANLY



That's...totally cool!

The first person to isolate an enzyme involved in DNA replication—called *DNA polymerase I*—was the American biochemist Arthur Kornberg (1918–2007).

Yeah! That's the enzyme that replicates DNA?

Enzymes really do a lot, don't they?

DNA polymerase was the enzyme we used when we obtained V_{max} and K_m . Do you remember?



Arthur Kornberg



In 1956, Kornberg extracted an active enzyme used for synthesizing DNA from a liquid in which *E. coli* was pulverized. This enzyme could be used to artificially synthesize DNA in a test tube, which created a great sensation in the scientific world at the time.



That discovery was seriously important.



It sure was! In fact, Kornberg was awarded the Nobel Prize in Physiology or Medicine in 1959.

DNA polymerase is officially called "DNA-dependent DNA polymerase." This means that a DNA polymerase enzyme requires the presence of an existing DNA template molecule in order to create a new DNA molecule.





I get it. Since the pairs are complementary, DNA polymerase steadily creates the chain by referring to the template.



The chemical reaction that DNA polymerase catalyzes is shown below.





In other words, DNA polymerase catalyzes the attachment of the next deoxyribonucleotide phosphate to the 3' end of a new DNA strand. This attachment is called a *phosphodiester bond*.

The nucleic acids that belong together are joined firmly by the "matchmaker" DNA polymerase.



That's right! Ha ha ha...





On the other hand, RNA usually exists as a single strand. The enzyme RNA polymerase actually uses DNA as a template when it binds ribonucleotides together into a new RNA strand.







Oh, that reminds me...earlier you said that RNA is an extremely important substance, but you never said why!



Just hold your horses, Kumi. We'll get into that soon (on page 220, to be exact).



Got it!



Unlike DNA, RNA usually breaks down soon after it's produced. Kumi, do you remember what we said about deoxyribose?



It's the pentose that's in DNA but not RNA, right?



That's right! In RNA, a hydroxyl group (2-OH) is attached to the carbon at the 2' position on each ribose, as shown in the following figure. In DNA, only a hydrogen is attached, hence the name *deoxy*ribose.

This hydroxyl group in RNA is actually a real troublemaker. The H could even be considered a biochemical playboy.





Because of the existence of this hydroxyl group, RNA is known to undergo *autolysis* (a self-inflicted decomposition) due to a phenomenon known as *base catalysis*.





If you think of the O and the H of a hydroxyl group as a couple, then H is the "playboy" that can be tempted by bases that exist around the RNA.

It's important to note that these aren't the A, U, C, and G bases we've been talking about so far. In this context, a *base* is a substance that can accept a proton (H^+), such as a hydroxide ion (OH^-). Bases can be thought of as the chemical opposite of *acids*, because they neutralize each other.



So the 2'-OH proton is extracted by a base?



That's right. Then the lone oxygen goes looking for a new partner to bond with.



Actually, after the proton is extracted, the oxygen (O^{-}), which carries a negative charge, will bond with the phosphate (P) in the phosphodiester bond at the neighboring 3' position (that is, the important bond connecting two ribonucleotides).





Since RNA frequently undergoes this kind of autolysis, it's considered "unstable" and is not very suitable for storing genetic information. DNA is far more stable than RNA since it does not contain the 2'-OH. Therefore, trustworthy DNA is the best for this purpose. A different but still very important role is assigned to RNA instead.



We'll learn about that role next!





* IN THIS BOOK, WE'LL GENERALLY SAY THAT GENES ARE "WRITTEN" (IN THE FORM OF A BASE SEQUENCE) IN DNA. ** ACTUALLY, THE EXON PART OF GENES IS CONSIDERED TO BE 1.5% OF THE WHOLE. (SEE PAGE 227.)

*** WE'LL EXAMINE INTRONS AND EXONS FURTHER ON PAGE 227.





MRNA



The enzyme called *RNA polymerase* enables genes that are written in the DNA—that is, their base sequences—to be used as templates to synthesize RNA with complementary base sequences.

This process is called *transcription* (it can also be referred to as RNA synthesis), and the synthesized RNA is called *messenger RNA* (*mRNA*). For example, take a look at the gene shown in the following figure.







DNA is double stranded, and, for each gene, only one side of the DNA is meaningful (that is, its base sequence becomes the blueprint). This is the *sense strand* for that gene. The other strand, which is complementary to the sense strand, is called the *antisense strand*.



That makes...sense.



Since mRNA is synthesized using the antisense strand as a template, the base sequence of the RNA that's produced is the same as that of the sense strand. However, where the base is T (thymine) in the sense strand, there will be a U (uracil) in the mRNA.



The synthesized mRNA is called *precursor mRNA*. It undergoes a series of chemical reactions and other processing* to become mature mRNA, which is then transported from the nucleus to the cytoplasm and is soon delivered to a ribosome.





So you can see why we call it a "messenger": it's transporting vital information out of the stronghold of the nucleus.

* This process includes *splicing*, which removes intron parts (see page 227), and the addition of a 5' "cap" and a poly-A tail.

RRNA AND TRNA



A ribosome is like a big protein-making machine consisting of several types of RNA called *ribosomal RNA* (*rRNA*) and several dozen types of ribosomal proteins.



They're not as large as mitochondria or chloroplasts, though.



They may not look like much, but ribosomes are pretty darned impressive!





Another type of RNA, which transports amino acids, is called *transfer RNA* (*tRNA*). The tRNA matches the base sequences of the delivered mRNA with the amino acids needed to build the coded proteins.



It should be pretty obvious where transfer RNA gets its name: it **transfers** amino acids.



The mRNA base sequence is a code for an amino acid sequence. More specifically, a three-base sequence forms the code for one amino acid. This three-base sequence is called a *codon*.





There are unique tRNA molecules for each codon. In other words, only a tRNA with the proper base sequence—called an *anticodon*—that is complementary to the mRNA's codon can adhere at the site of the ribosome.*

The role of rRNA is to connect the amino acids that were transported via tRNA into long chains. Amino acids are connected according to the mRNA base sequence (that is, the codon sequence) to create proteins in accordance with the base sequence (the genes, or "blueprint") of the original DNA.





Here's what a single rRNA looks like:





Huh? It looks like a maze.

If you look carefully, you can see that it's actually just **one RNA strand folded** in a complex way. The ends are marked 3' and 5'.

Oh! You're right!



When an RNA strand folds, its bases pair with each other. In the image above, those pairings are represented by the transverse lines, or "ladder rungs." Base pairing within a single strand is a feature of RNA not found in DNA: **It can fold into a variety of forms according to differences in base sequences**.









Wow, cool! They look like little line drawings.



RIBOZYMES



To summarize, RNA breaks down quickly, but it can do things that DNA can't. RNA exists both in the nucleus and the cytoplasm, and it can take various forms depending on the base sequence.



It's a very flexible molecule, isn't it?



Definitely. RNA researchers have long believed that RNA probably has other important jobs besides copying genes and interacting with ribosomes.

Ribozymes, which were independently discovered at the beginning of the 1980s by the American microbiologist Thomas Cech and the Canadian molecular biologist Sidney Altman, foretold later developments in RNA research. You learned earlier that one very important job of proteins is their work as enzymes (see page 153), but Cech and Altman actually discovered that RNA has the flexibility to work as an enzyme as well.



Huh!



So they combined the words "RNA (ribonucleic acid)" and "enzyme" to coin the term *ribozyme*.





Many of the ribozymes that have been discovered or artificially created so far are involved in building or cleaving RNA or DNA.

Cech discovered that part of RNA itself catalyzes the chemical reaction to remove sections that are meaningless as protein blueprints (introns) and to join the sections that are meaningful (exons). This is referred to as *self-splicing** and is fairly rare.

Usually, splicing involves an enzyme known as *spliceosome*, which is a large complex made of RNA and proteins.



Splicing involves several hydroxyl groups (-OH) of RNA, such as the 2'-OH or the 3'-OH.





Hey, it made a loop!



The discovery of ribozymes caused RNA to be viewed as a multifaceted molecule that does a variety of tasks, and RNA research advanced rapidly. As the 21st century began, this research showed that RNA has other various roles and that many different forms of RNA exist in cells. RNA research is truly in full bloom now, and there are high expectations for future research.

* The genes of eukaryotes (like humans) are divided into several exons by non-coding base sequences called introns. The introns must be removed from the mRNA before translation, in a reaction known as splicing.

3. Biochemistry and Molecular Biology

THE DIRTY JOB OF A BIOCHEMIST

Nowadays—especially in cities—there are fewer and fewer opportunities for exploring nature and really getting your hands dirty. In a world where a mustard-stained tie is the greatest potential tragedy, most people can't even imagine trekking out into the wilderness to really dig through the (often filthy!) wonders of the natural world.

But humans, of course, are products of nature. Long before the miracles of instant spot remover and nectarine-scented hand sanitizer, we lived like other animals, surrounded by the great outdoors.

Although biochemistry is a discipline that attempts to view the incredible phenomena of life through the lens of science, the raw materials of that research are the living organisms themselves—again, products of nature.

Once, during my graduate student days, I went to the mountains in order to collect certain plants that contain large amounts of the proteins that I was researching.

I drove with a younger graduate student to some nearby mountains, and we proceeded along a road that was thickly overgrown with plants. When we got to a place where the car could no longer pass, we continued on foot to find the plant called *Phytolacca Americana* (American pokeweed).

We brought the pokeweed back to the laboratory, washed off the dirt, and used kitchen knives and scissors to cut them up so that we could extract the target proteins from them.

I would also routinely visit a nearby meatpacking plant to obtain an organ called the *thymus* from recently slaughtered cows. (The thymus is usually discarded, so they were happy to give them away free of charge.) I would use scissors to cut up the organ and store the little pieces in the laboratory's refrigerator as experimental samples. This was necessary to extract the research target protein (DNA polymerase).

In this way, biochemistry originally developed based on a methodology of extracting (isolating, purifying, etc.) chemical substances from organic raw materials and checking their chemical properties.

In contrast, molecular biology is a discipline that tries to elucidate the phenomena of life by using *biopolymers* such as DNA and proteins.

Simply speaking, if molecular biology just deals with DNA and RNA or prepares an environment for artificially creating proteins (by using *E. coli*, for example), it is unnecessary to use flesh-and-blood, organic raw materials such as a cow's thymus or plant materials.

In a manner of speaking, the data obtained in these artificial lab environments is "digital." Molecular biology is practiced using high-tech equipment and cutting-edge technologies, and molecular biologists are rarely caked in mud or smeared with animal blood...

As a result, some biochemists refer to their own research as "dirty work." I think that when they do this, it's somehow self-deprecating.

However, it is an undeniable fact that the accumulation of knowledge from that kind of dirty work built the foundation of molecular biology in the first place. Although many young molecular biologists have never once used flesh-and-blood, organic materials other than *E. coli*, cell cultures, or experimental animals, biochemistry and molecular biology have long been bound together and remain so even today. This fact should never be forgotten.

EARLY BIOCHEMISTRY AND MOLECULAR BIOLOGY

In 1897, German biochemist Eduard Buchner made the revolutionary discovery that *fermentation* occurred in an extract created from yeast cells, which contained yeast protein but no living yeast cells.

Prior to that, people believed that the chemical reaction called fermentation, which is characteristic of living organisms, could not happen without the presence of living cells. However, this idea was utterly obliterated by Buchner's discovery.

Because of this, the theory of *vitalism*, in which the phenomena of life occur only because of a characteristic force (spirit or life energy) of living organisms, practically disappeared, and the foundation was built for research on chemical reactions occurring in living organisms in test tubes. In other words, biochemistry was born.

Since Buchner found that an actual living organism need not be present, it would not be an exaggeration to say that this was the discovery that paved the way for the arrival of molecular biology.

As a clearer understanding of the chemistry of life developed, it became apparent that certain mechanisms common in all living organisms form the basic foundation of life. Some of these mechanisms, for example, are that DNA is used as the genetic language, that the basic theory (central dogma) behind the reading of genes to create proteins is consistent, and that the same proteins often do the same kinds of work.

Because these mechanisms are universal among living organisms, it was important to develop methodologies for studying DNA, which serves as the blueprint for proteins, and for elucidating the function of proteins.

DEVELOPMENT OF RECOMBINANT DNA TECHNIQUES

In 1972, American molecular biologist Paul Berg performed the first successful recombinant DNA experiment in the world by artificially manipulating DNA in a test tube to create a DNA sequence that did not exist in the natural world.

A method for easily decoding the base sequences of DNA was developed by English biochemist Frederick Sanger in 1977, and a method for copying (or *amplifying*) DNA was developed by American molecular biologist Kary Mullis in 1985. After these discoveries, recombinant DNA experimental techniques advanced rapidly.

Recombinant DNA techniques developed after it had been clearly demonstrated that DNA is the genetic language (or, in other words, that the information referred to as genes is written in the form of base sequences in DNA). Molecular biologists imagined that if they only had the DNA base sequences and if they only created an environment in which proteins could be created from those base sequences, they could elucidate the phenomena of life, which can be thought of as the totality of chemical reactions in which proteins are involved.

For example, if genes from an external source are introduced into a simple, easy-tomanipulate organism for which these mechanisms are well understood (such as *E. coli*) to create proteins within that organism, a large quantity of proteins can be obtained all at once without ever having to perform any more "dirty work."

Generally speaking, this idea can be expressed as: Explain the DNA sequences! Explain life!

Molecular biology has developed with this aim. The methodology used to achieve this was based on recombinant DNA techniques.

RETURNING TO BIOCHEMISTRY

After the Human Genome Project (an international cooperative research project aiming to sequence the genetic information of humans) was completed in 2003, researchers turned their attention from DNA back towards proteins and RNA.

The post-genome era, or post-sequencing era, arrived.

No matter how much DNA steals the limelight, no matter how its handling techniques develop, and no matter what anyone says, when life phenomena are viewed as a collection of "chemical reactions," the only things at work there are proteins and RNA.

This is because, even if all of the base sequences of human DNA (that is, the entire genome) are known, the data is meaningless unless the roles of the proteins and RNA that are made from DNA are understood.

Currently, if the amino acid sequences of many proteins are known and the role they play is also understood, then the work of unknown proteins can be predicted to a certain degree based on only amino acid information.

However, the role of unknown proteins must be verified by using biochemical techniques. No matter how many molecular biological techniques (such as DNA recombination) are used to research the activity of proteins to elucidate their roles, the question of whether or not those proteins really do those jobs inside natural cells will remain unanswered. In a manner of speaking, this is the same as the idiomatic expression "you can't see the forest for the trees." As long as the research target is a biological substance, biochemistry remains an absolutely necessary and important academic discipline.

THE ORIGIN OF THE CELL

No doubt you've heard the phrase "origin of life." We're not going to tackle the intimidating question of what life actually is here. Instead, we're concerned with the first living organisms or, in other words, "the origin of cells."

So how did cells originate on Earth?

By now, you should understand that a cell is the location of many vital chemical reactions in living organisms and that cells can divide and multiply. These are two major traits that define life.

Chemical reactions that transform one substance into another substance, which are performed inside of cells, and networks of many of these chemical reactions, which occur across many cells, are collectively referred to as *metabolism*.

Cells must constantly undergo metabolism in order to produce the energy required to maintain their organization and to reproduce.

In addition to metabolism, reproduction, also called *self-replication*, is a major characteristic of living organisms. Unicellular organisms can often simply replicate their cellular contents and split in two; this process is called *binary fission*. For multicellular organisms, reproduction is more complex and involves specialized reproductive cells, which initiate the development of offspring.

(Although the way in which cells replicate themselves can certainly be called selfreplication, the expression is a little awkward. Let's just call it plain old "replication" from now on.)

Cells perform both metabolism and replication, both of which are extremely important when considering the origin of life. But which came first, metabolism or replication?

This is one of the most difficult questions confronting scientists who are conducting research concerning the origin of life. When the "pouch wrapped in a membrane" (the cell) originated, what had been occurring inside that enclosure? Some scholars think that the "pouch" had been performing metabolism after many diverse molecules had collected together. Some time later, the "pouch" absorbed molecules capable of causing replication, so the entire assembly became able to divide and multiply. This idea is known as *metabolism first*.

Other scholars think that the original "pouch" had enclosed replicating molecules and had been dividing. Later, it became able to perform metabolism, and it eventually obtained a more advanced replication method, causing it to evolve into a cell. This idea is known as *replication first*.

Many other scholars think that posing the question of "which came first" is nonsense and that metabolism and replication coevolved cooperatively.

In any case, the biochemical process of metabolism is a very large-scale phenomenon.

4. Conducting Biochemistry Experiments

It was mentioned on page 230 that the functions of proteins can be verified by using biochemical techniques, but what kinds of experiments do biochemists perform?

There are various experimental methods depending on the field of research, and not each and every one will be presented here. Instead, I will introduce several experimental methods that I have used myself.

COLUMN CHROMATOGRAPHY

Column chromatography is an experimental method for separating substances that have the same property from a mixture of substances. For example, we can collect just proteins with certain properties from the liquid extract of American pokeweed, which was introduced earlier, or from the liquid obtained after a cow thymus was blended in a juicer. Special *resins* are packed in a long, narrow tube made of glass or some other material. Some substances adhere to those resins and are trapped, allowing for the collection of only those substances that do not adhere.

Various types of chromatography, such as ion exchange chromatography, gel filtration chromatography, and affinity chromatography, can be used, depending on the type of resin or target protein. As an example, here we will look at a method for purifying the enzyme DNA polymerase α from the thymus of a calf.

As shown below, the cells are smashed by grinding up the calf thymus, and a solution with a high salt (sodium chloride) concentration is used to extract proteins and other molecules. This extract (the liquid in the flask) passes through the large glass tube (called the *column*) in which the *ion exchange resin* has been packed. The proteins are broadly divided into those that are trapped in the column and those that pass through the ion exchange resin.



The substances that pass through the ion exchange resin are collected in test tubes, but the substances that are absorbed are separated from the resin and collected in test tubes later by adding a liquid with a higher salt concentration. DNA polymerase α adheres to the column and can be retrieved by adding a liquid with a salt concentration of 0.5 M (moles per liter). (This is "sample 1.")

The figure below shows the method of purifying DNA polymerase α from this "sample 1."



This technique, called *affinity chromatography*, uses a small glass tube packed with a resin combined with an antibody (a type of protein that's created by the immune system) that can only bond with DNA polymerase α . When sample 1 is passed directly through this resin, it's divided into two parts—one that is absorbed and one that passes through. DNA polymerase α , which is absorbed, can be retrieved by running a solution with an extremely high concentration (3.2 M) of magnesium chloride through the resin. Since the part that's retrieved here is almost 100% DNA polymerase α , it has been "purified" at this time.

In this way, DNA polymerase α can be efficiently purified by using a combination of ion exchange chromatography and affinity chromatography.

ELECTROPHORESIS AND A WESTERN BLOT

This is an experimental method for isolating a specific protein, recognizing what type of protein is in a sample, or checking the size of a target protein. *Electrophoresis* is a method in which a sample is "loaded" on top of a thin gelatinous slab (gel) and a current passes through it to cause the sample to move through the gel. SDS-polyacrylamide gel electrophoresis (left side of the following figure), which separates proteins based on molecular size, is commonly used. After the proteins are separated, they are detected by reacting them with a special reagent, such as a dye or a fluorescent marker.

A western blot (right side of the following figure) is a method of transferring the proteins that are in the gel onto a thin membrane, while retaining the positions that they were in after they were separated, and then detecting a specific protein on the membrane by adding an "antibody" that only reacts with that protein. *Lectin blotting*, which is described next, is an application of a western blot.



LECTIN BLOTTING

Lectins are proteins that can bond specifically to certain sugar chains. Because lectins will bond differently according to the type of sugar chain, lectins can be used to identify the type of sugar chain that is bonded to a protein. A method similar to a western blot can be used after proteins are transferred onto a membrane. Various lectins are mixed with the proteins, and only the lectins that reacted are detected. This lets us identify the types of sugar chains that exist in the proteins that were transferred onto the membrane. This is called lectin blotting.

The following figure shows an experiment in which a lectin called wheat germ agglutinin (WGA) identifies a sugar chain to which the saccharide called *N*-acetylglucosamine (GlcNAc) is attached.

In this lectin blot performed for the rough protein fraction of the starfish oocyte, it is apparent that two large bands are glowing brightly. WGA was the lectin used here.



CENTRIFUGATION

Like column chromatography, *centrifugation* is an experimental method that is performed to separate organelles, proteins, and other biological molecules that have the same density from a mixture of substances with differing densities. The sample solution is placed in a test tube, and the test tube then spins around the centrifuge axis at a very high speed. To handle small molecules, like proteins, ultracentrifugation may also be performed, which spins the test tube at tens of thousands of revolutions per minute. DNA and other polymers can also be isolated in this way.



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ENZYME REACTION MEASUREMENT

There are various methods of measuring enzyme activity, such as using radioisotopes to measure the amount of a reaction product produced or using a change in color to measure the reaction product formed when a substrate is acted on by an enzyme.

I will explain a method of using radioisotopes to measure the enzyme activity of DNA polymerase and a method of using a color reaction to measure the enzyme activity of α -amylase.

DNA POLYMERASE ACTIVITY MEASUREMENT METHOD

First, place the following in a mirco test tube: the solution for measuring activity (with pH optimized and buffered for DNA polymerase), DNA polymerase, the DNA that is to be the template, the nucleotide that is to be the raw material, and the magnesium chloride co-factor. Add the nucleotide that contains the radioisotope, and let it react at 37° C for a fixed interval.

When this is done, the nucleotides that contain the radioisotopes are captured in the new DNA strands that have been synthesized by the DNA polymerase. The unreacted nucleotides are removed, and the synthesized DNA strands are placed in a small radioisotope bottle for measurement in a device called a *liquid scintillation counter*, which measures radioisotopes. Since higher enzyme activity means more radioisotopes were captured in the DNA, higher numeric values indicate higher enzyme activity.



DNA POLYMERASE ACTIVITY MEASUREMENT METHOD

a-AMYLASE ACTIVITY MEASUREMENT METHOD

An α -amylase solution (such as saliva) is added to a solution in which starch has been dissolved inside a test tube. If an iodine solution is immediately added to this, before any of the starch has been broken down, the starch will react with the iodine and produce a blue-violet color. However, as time passes, the starch is broken down by the α -amylase, and the color steadily changes (blue-violet \rightarrow violet \rightarrow red \rightarrow orange \rightarrow pale orange). Eventually, when all the starch has been broken down, the solution will become colorless. The enzyme activity of α -amylase can be measured by using a spectrophotometer to quantify the appearance of its color as a numeric value.






















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ABOUT THE AUTHOR

Masaharu Takemura, PhD, is currently an Associate Professor at the Tokyo University of Science. His specialties are molecular biology and biology education.

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