Abstract

Today there are two major theoretical frameworks in biology. One is the ‘chemical paradigm’, the idea that life is an extremely complex form of chemistry. The other is the ‘information paradigm’, the view that life is not just ‘chemistry’ but ‘chemistry-plus-information’. This implies the existence of a fundamental difference between information and chemistry, a conclusion that is strongly supported by the fact that information and information-based-processes like heredity and natural selection simply do not exist in the world of chemistry. Against this conclusion, the supporters of the chemical paradigm have pointed out that information processes are no different from chemical processes because they are both described by the same physical quantities. They may appear different, but this is only because they take place in extremely complex systems. According to the chemical paradigm, in other words, biological information is but a shortcut term that we use to avoid long descriptions of countless chemical reactions. It is intuitively appealing, but it does not represent a new ontological entity. It is merely a derived construct, a linguistic metaphor. The supporters of the information paradigm insist that information is a real and fundamental entity of Nature, but have not been able to prove this point. The result is that the chemical view has not been abandoned and the two paradigms are both coexisting today. Here it is shown that an alternative does exist and is a third theoretical framework that is referred to as the ‘code paradigm’. The key point is that we need to introduce in biology not only the concept of information but also that of meaning because any code is based on meaning and a genetic code does exist in every cell. The third paradigm is the view that organic information and organic meaning exist in every living system because they are the inevitable results of the processes of copying and coding that produce genes and proteins. Their true nature has eluded us for a long time because they are nominable entities, i.e., objective and reproducible observables that can be described only by naming their components in their natural order. They have also eluded us because nominable entities exist only in artifacts and biologists have not yet come to terms with the idea that life is artifact making. This is the idea that life arose from matter and yet it is fundamentally different from it because inanimate matter is made of spontaneous structures whereas life is made of manufactured objects. It will be shown, furthermore, that the existence of information and meaning in living systems is documented by the standard procedures of science. We do not have to abandon the scientific method in order to introduce meaning in biology. All we need is a science that becomes fully aware of the existence of organic codes in Nature.

Keywords – Information, meaning, organic codes, mechanism, observables, ontology.

Introduction

From time immemorial it has been taken for granted that life is fundamentally different from matter, but in the last few centuries this belief has been seriously challenged by the view that ‘life is chemistry’. The idea that life had a natural origin on the primitive Earth suggests that the first cells came into being from previous chemical systems by spontaneous chemical reactions, and this is equivalent to saying that there is no fundamental divide between life and matter.

This chemical paradigm is very popular, today, and is often considered a complement of the Darwinian paradigm but this is not the case. The reason is that natural selection, the cornerstone of Darwinian evolution,
does not exist in inanimate matter. In the 1950s and 60s, furthermore, molecular biology has uncovered two fundamental entities of life - biological information and the genetic code - that are totally absent in the inorganic world, and this again suggests that a deep divide does exist between life and matter.

Ernst Mayr, one of the architects of the Modern Synthesis, has been one of the most outspoken supporters of the view that life is fundamentally different from inanimate matter. In *The Growth of Biological Thought* (1982), he made this point in no uncertain terms:

“… The discovery of the genetic code was a breakthrough of the first order. It showed why organisms are fundamentally different from any kind of nonliving material. There is nothing in the inanimate world that has a genetic program which stores information with a history of three thousand million years!” (p. 124)

“… Except for the twilight zone of the origin of life, the possession of a genetic program provides for an absolute difference between organisms and inanimate matter.” (p. 56)

The discoveries of molecular biology, in short, appear in contrast with the chemical paradigm, and this raises formidable problems. On the one hand it is an experimental fact that natural selection, biological information and the genetic code do not exist in inanimate matter. On the other hand, we seem unable to accept that life evolved from inanimate matter and yet it is fundamentally different from it. How can something give rise to something fundamentally different from itself? How could the physical world produce life if there is an absolute discontinuity between them?

The aim of this paper is to show that a solution to these problems does exist, but it is not provided by the paradigms that are based respectively on chemistry and information. It is provided instead by a third approach that here is referred to as the ‘code paradigm’ because it is based on the organic codes of life. To this purpose the paper has been divided into two parts. The first is dedicated to the two present paradigms of modern biology and the other to the new theoretical framework.

**PART 1**

**Chemistry versus Information**

**1-1 The Chemical Paradigm**

Ever since the scientific revolution, physics has been the ‘queen’ science, and biologists have been split into opposite camps, one in favour and one against adopting its method, an approach which has become known as *mechanism*. In biology, the first version of mechanism was the Cartesian doctrine that “the body is a machine” and that the clock is its model: “A healthy man is like a well functioning clock, and an ill man is like a clock that needs repairing” (Descartes, 1637).

The mechanical concept of nature spread very quickly in 17th century Europe, but not without conflict. Opposition came particularly from a new science that was slowly emerging from alchemy and that regarded the human body essentially as a seat of chemical reactions. The heirs of the alchemists were determined to leave magic behind but had no intention of accepting the ‘mechanical’ view of nature, and one of chemistry’s founding fathers, Georg Ernst Stahl (1659-1731), launched an open challenge to mechanism. He claimed that organisms cannot be machines because what is taking place inside them are real transmutations of substances and not movements of wheels, belts and pulleys.

The arguments of the chemists did have an impact, and eventually forced mechanists to change their model. In the course of the 18th century, the view that organisms are mechanical machines, gradually turned into the idea that they are chemical machines. This change went hand in hand with the development of the steam engine, and that machine became the new model of biology. In the 19th century, furthermore, the study of the steam engine was pushed all the way up to the highest level of theoretical formalism, and culminated with the discovery of the first two laws of thermodynamics. The result is that any living system came to be seen as a thermodynamic machine, i.e., as a chemical machine that must be continuously active in order to obey the laws of thermodynamics.

The old opposition between physics and chemistry came to an end, and the two sciences together gave origin to a unified framework that is often referred to as the ‘chemical paradigm’, the idea that life is an extremely complex form of chemistry. This is equivalent to saying that all biological processes are chemical transformations of matter and energy, and are completely described, in principle, by physical quantities.
The chemical paradigm has underlined time and again - against all forms of vitalism - that living systems are subject to the laws of thermodynamics, but it is by no means limited to this principle. It is a paradigm which has steadily grown by adding new arguments to its thesis. The non-equilibrium thermodynamics of Ilya Prigogine, the phase-transitions of Stuart Kauffman, chaos theory and complexity theory, are all descriptions of natural processes that rightly belong to the framework of the chemical paradigm. The same is true for the idea that life is shaped by physical forces and by mathematical principles, a recurrent theme in the history of science, from Goethe and D’Arcy Thomson to modern structuralists like René Thom and Brian Goodwin. The chemical paradigm, in short, is the view that the laws of physics and chemistry and the principles of mathematics are all that we need to account for the presence of life in the universe.

1-2 The Information Paradigm

At the beginning of the 20th century, the rediscovery of the laws of Mendel led Wilhelm Johannsen to make a sharp distinction between the visible part of an organism (the phenotype) and the invisible part that carries its hereditary instructions (the genotype). Johannsen (1909) proposed that every living being is a dual entity, a synthesis of two complementary realities. This idea was largely ignored, at first, but a few decades later the computer made it immediately comprehensible. The phenotype-genotype duality was a hardware-software distinction, and became the prototype description of any organism. The model of the living system changed again and became the computer.

In 1953, James Watson and Francis Crick pointed out that the sequence of nucleotides represents the information carried by a gene. A few years later, the mechanism of protein synthesis was discovered and it was found that the sequence of nucleotides in genes determines the sequence of amino acids in proteins, with a process that amounts to a transfer of linear information from genes to proteins. In both types of molecules, therefore, biological information was identified with, and defined by, the specific sequence of their subunits.

These discoveries gave origin to the ‘information paradigm’, the second great theoretical framework of modern biology. It is the idea that living systems are information-processing machines, and that life is based not only on chemistry (energy and matter) but also, and above all, on information (Maynard-Smith, 2000). In this framework, chemistry accounts for the ‘hardware’ of living systems, whereas information provides the software. The view that ‘life is chemistry’ was replaced in this way by the idea that ‘life is chemistry+information’.

This, in turn, led to the concept of the ‘genetic programme’, the idea that the genome is for the cell what a programme is for a computer. The logical separation that exists between programme and machine implies that something similar exists between the genome and the cell, and such a biological separation has in fact been documented by an outstanding number of experimental results (Danchin, 2009). Many genes, for example, have been transplanted from one organism to another and have turned out to be fully functional inside the new cells. Many bacteria now produce human proteins, and the very existence of viruses can be explained by the transmission of independent genetic strings, thus confirming that genes are separable from the cell machine. It has even been possible to transplant an entire genome from one species to another, thus proving that a genome does have a substantial degree of autonomy (Lartigue et al., 2007).

This informational view of life, has been immediately accepted into the Modern Synthesis because the concept of information goes hand in hand with the processes of heredity and natural selection. Heredity is precisely the transmission of genetic information from one generation to the next, the short-term result of molecular copying. The long-term repetition of copying, on the other hand, is inevitably accompanied by errors, and in a world of limited resources not all copies can survive and a selection is bound to take place. That is how natural selection came into existence. It is the long-term result of molecular copying, and can exist only in a world of molecules that carry information.

Today, in other words, heredity and natural selection are both squarely based on information, and the information paradigm has become, to all effects, the modern version of the Darwinian paradigm, a view of life which is in conflict with the chemical paradigm, because information, heredity and natural selection simply do not exist in the world of chemistry.

1-3 Shannon’s Information Theory

The concept of information has been introduced in science in two very different ways. In biology, as we have seen, the information of genes and proteins is defined by their sequences and is referred to as biological
information. In engineering, on the other hand, the information of a message is defined by an entropy-like formula introduced by Claude Shannon in 1948, and is referred to as statistical information.

Shannon was particularly interested in telephone transmissions, and described any communication system as a combination of a source (that produces signals), a destination (that receives them) and a channel in between. He conceived information as an entity that is generated when uncertainty is reduced, so he proposed to measure information by measuring changes in uncertainty (Shannon, 1948). To this purpose he described the state of all communication systems with a probability function, and was able to prove a number of theorems on their ability to transmit information. Shannon established in this way an entirely new field of research which has become known as ‘Information Theory’. Perhaps the most important result of this field was the demonstration that reliable communication is possible over unreliable channels, a result which opened the way to the tremendous expansion and success of the communication technologies.

Shannon underlined that our messages are digital entities, because they are made of discreet units, and it is precisely their digitality that allows us to associate a probability function to each of them. This measures the statistical information of the sequence in terms of digital units called bits (or shannons, according to the International Standards Organization).

Sequences, on the other hand, are made of units which are not only discreet but are also arranged in a specific order, and in genes and proteins this order represents the biological information, or the specificity, of the sequence. Digitality, in short, is associated with the statistical information of a sequence, whereas specificity represents its biological information. The important point is that these two types of information deal with different but equally fundamental processes. Statistical information is concerned with the faithful transmission of messages irrespective of their meaning, whereas biological information is concerned with their unique meanings.

The goal of communication is the reliable transmission of all messages, whatever is their meaning, and this is why in communication technology information has been sharply separated from meaning. In his seminal papers, Shannon expressed this concept in no uncertain terms:

“The fundamental problem of communication is that of reproducing at one point, exactly or approximately, a message selected at another point. Frequently the messages have meaning; that is they refer to or are correlated according to some system with certain physical or conceptual entities. These semantic aspects of communication are irrelevant to the engineering problem.”

In the life sciences, on the other hand, no such clear distinction has been made. Modern biology has accepted the concept of information but not the concept of meaning, with the result that meaning has either been ignored or it has not been regarded as an entity in its own right. This is a major unresolved problem in the information paradigm, a problem that arises from the lack of a clear distinction between information and meaning, in sharp contrast with the lesson that comes from Shannon’s theory.

1-4 Digital and analogue

Shannon’s statistical information is totally different from the biological information of molecular sequences, but they do have something in common. Both of them are totally absent in chemical reactions and are therefore in conflict with the view that ‘life is chemistry’.

Perhaps the strongest criticism of the chemical paradigm has come in fact from the Information Theory camp, and in particular from Hubert Yockey, one of the organizers of the first congress dedicated to the introduction of Shannon’s Information in Biology (Yockey et al, 1958). In a long series of articles and books, Yockey (1974, 1992, 2000, 2005) has underlined that heredity is transmitted by factors that are “segregated, linear and digital” whereas the compounds of chemistry are “blended, three-dimensional and analog”.

“Chemical reactions in non-living systems are not controlled by a message. If the genetic processes were purely chemical, the law of mass action and thermodynamics would govern the placement of amino acids in the protein sequences according to their concentrations … There is nothing in the physico-chemical world that remotely resembles reactions being determined by a sequence and codes between sequences” (Yockey, 1992)

Yockey has tirelessly pointed out that no amount of chemical evolution can cross the barrier that divides the analog world of chemistry from the digital world of life, and concluded from this that the origin of life cannot have been the result of chemical evolution. At the same time, however, Yockey did not invoke an extraterrestrial origin or Intelligent Design. He claimed instead that the origin of life is unknowable, in the
same sense that there are propositions of logic that are undecidable. The problem, with this argument, is that
the existence of undecidable propositions has been proven in logic, whereas the conclusion that the origin of
life is unknowable is just an assumption. It may be a legitimate assumption, in principle, but in no way it is
complicable to Godel’s theorem and certainly it does not carry the same weight.

It is important however to recognize that Yockey’s distinction between analog and digital entities cannot
be ignored. He was absolutely right in saying that the spontaneous reactions of chemistry cannot produce
molecules with linear, digital and specific properties, and this is a point that must be taken into account by
any scientific theory on the origin of life.

The information paradigm is based on the experimental fact that heredity and natural selection do not
exist in the inanimate world, and the discovery that they are both based on information leads to the
conclusion that ‘life is chemistry-plus-information’. At the same time, however, the information paradigm
maintains that information is fully compatible with the laws of physics and chemistry. But how? How can we
prove that information is distinct from chemistry and yet it is perfectly compatible with the laws of physics
and chemistry? This is the classical problem of understanding how it is possible that life evolved from matter
and yet it is fundamentally different from it, and the information paradigm has not been able to solve it.

1-5 The claim of Physicalism

The view that ‘life is chemistry’ was proposed for the first time by Jan Baptist van Helmont (1648), and has
been re-proposed countless times ever since. One of the most recent formulations has been given by Günther
Wächtershäuser (1997) in these terms “If we could ever trace the historic process backwards far enough in
time, we would wind up with an origin of life in purely chemical processes”.

He added that “The science of chemistry, however, is an ahistoric science striving for universal laws... so
this is the challenge of the origin of life: to reduce the historic process of biological evolution to a universal
chemical law of evolution”. The difficulty of this task, he pointed out, is due to the fact that “Chemistry is
mechanistic and history teleological, and the life sciences are the arena where mechanistic explanations and
teleological understanding come into close encounter.”

Wächtershäuser claimed that “information is a teleological concept”, and gave a specific example of the
conflict between mechanism and teleology: “On the level of nucleic acid sequences it is quite convenient to
use the information metaphor … and apply teleological notions such as ‘function’ or ‘information’… but in
the course of the process of retrodiction the teleological notions, whence we started, fade away. And what
remains is purely chemical mechanism”. This amounts to saying that biological information, the most basic
concept of molecular biology, does not really belong to science.

The same thesis has been expressed by the supporters of physicalism, the view that all natural processes
are completely described, in principle, by physical quantities. The crucial point is that a sequence cannot be
measured and this means that biological information, or biological specificity (as some prefer to call it) is not
a physical quantity. So, what is it? A similar problem arises with the genetic code. The rules of a code cannot
be measured and cannot be reduced to physical quantities. So what are they?

According to physicalism, biological information and the genetic code are mere metaphors. They are
linguistic expressions that we conveniently use as shortcuts in order to avoid repeating every time all the
details of long chains of chemical reactions. But behind those terms there are only chemical reactions and
nothing else. They are like those computer programs that allow us to write our instructions in English, thus
saving us the trouble to write them with the binary digits of the machine language. Ultimately, however,
there are only binary digits in the machine language of the computer, and in the same way, it is argued, there
are only physical quantities at the most fundamental level of Nature.

This conclusion, known as the physicalist thesis, has been proposed in various ways by a number of
scientists and philosophers (Chargaff, 1963; Sarkar, 1996; Mahner and Bunge, 1997; Griffiths and
Knight, 1998; Griffith, 2001, Boniolo, 2003), and it is equivalent to the thesis that ‘life is chemistry’.

This is one of the most deeply dividing issues of modern science. Many biologists are convinced that
biological information and the genetic code are real and fundamental components of life, but physicalists
insist that they are real only in a very superficial sense and that there is nothing fundamental about them
because they must be reducible, in principle, to physical quantities.
1-6 Two ontological problems

The discovery of biological information was the event that transformed biochemistry into molecular biology, and the paradigm that ‘life is chemistry’ into the new paradigm that ‘life is chemistry-plus-information’. Surprisingly, however, the old regime has not been deposed. The idea that life is an extremely complex form of chemistry is still very popular, today, because it is widely accepted (1) that life evolved spontaneously on our planet from primitive chemical systems and (2) that all biological processes are completely described, in principle, by physical quantities.

These are the two key points that lie at the heart of the chemical paradigm, and we can go beyond that paradigm only if we replace them with more general concepts.

The idea that life is ‘chemistry-plus-information’, implies that information is ontologically different from chemistry. But can we prove it? Ontology is the study of being and saying what an entity is amounts to defining it. Ontology, in short, is concerned with the definition of entities at the most basic level. In order to prove that life is ‘chemistry-plus-information’, therefore, we need to prove that there is an ontological difference between information and chemistry. More precisely, we need to prove that the above two pillars of the chemical paradigm are both wrong, and to this purpose we must show (1) that it was not spontaneous chemical reactions that gave origin to the first cells and (2) that in addition to physical quantities we need other fundamental entities to describe what goes on in living systems.

These are the two great problems that we have before us. Is there an ontological difference between life and matter? Is there an ontological difference between information and chemistry? The rest of the paper is dedicated precisely to these two problems. The first is addressed in the remaining sections of Part 1, whereas the whole of Part 2 is dedicated to the ontological definitions of organic information and organic meaning.

1-7 The idea that “Life is artifact-making”

According to the chemical paradigm, the first cells evolved from chemical systems by spontaneous chemical reactions that are all fully described, in principle, by physical quantities. No other entities are required to explain the origin of life by chemical evolution, and this is why physicalism concludes that biological information and the genetic code are purely metaphorical terms.

It must be underlined that the physicalist thesis would be absolutely correct if genes and proteins were spontaneous molecules because there is no doubt that all spontaneous reactions are completely accounted for by physical quantities. This, however, is precisely the point that molecular biology has proved wrong. Genes and proteins are not produced by spontaneous processes in living systems. They are produced by molecular machines that physically stick their subunits together according to sequences and codes and are therefore manufactured molecules, i.e., molecular artifacts. This in turn means that all biological structures are manufactured, and therefore that the whole of life is artifact-making (Barbieri, 2004, 2006, 2008). This conclusion may appear paradoxical, at first, but let us take a closer look.

All chemical reactions are either spontaneous or catalyzed processes, and biochemistry has clearly shown that virtually all reactions that take place in living systems are catalyzed processes. What molecular biology has discovered is that the production of genes and proteins requires not only catalysts but also templates. The catalysts join the subunits together by chemical bonds, and the templates provide the order in which the subunits are assembled. It is precisely that order that determines biological specificity, the most important characteristic of life, and that order comes from a molecule that is outside the assembled molecule.

This is precisely the characteristic that divides spontaneous objects from artifacts. In spontaneous and in catalyzed processes, the order of the components comes from within the molecules, i.e., is determined by internal factors, whereas in genes and proteins it comes from without, from an external template.

The difference between spontaneous and manufactured objects, in short, does not exists only at the macroscopic level of culture. It exists also at the molecular level, because it is an experimental fact that genes and proteins are manufactured molecules. It is also an experimental fact that they are template-dependent molecules, and this means that they are molecular artifacts.

Let us now look at the difference between the processes that manufacture genes and proteins. They both require catalysts and templates, but in addition to that, proteins also require a set of coding rules (in the form of molecular adaptors). This is because genes are nucleic acids that are formed by copying a template, whereas proteins cannot be copied. Their order must still come from nucleic acids (because only these
molecules can be inherited) but a sequence of nucleic acid has to be translated into a sequence of amino acids
and this is achieved, in protein synthesis, by the rules of the genetic code.

We realize in this way that there are two distinct processes at the basis of life: the copying of genes and
the coding of proteins. Genes are manufactured by molecular machines that can be referred to as copymakers
and proteins by molecular machines that can be called codemakers. Copying and coding, on the other hand,
are both artifact-making processes and life as we know it requires both of them. We can truly say therefore
that life is artifact-making, or, more precisely, that life is artifact-making by copying and coding.

This makes us realize that the physicalist thesis is wrong because it is only spontaneous processes, not all
processes, that are completely described by physical quantities. Manufacturing processes require additional
entities, like sequences and coding rules, that are not physical quantities, because they cannot be measured,
but which are absolutely essential to the description of all living systems.

1-8 A useful metaphor

We find it difficult to accept that life evolved from matter and, at the same time, that it is fundamentally
different from it. How can something give origin to something fundamentally different from itself? The way
out of this dilemma, as we have seen, is the idea that life is artifact-making, i.e., that the fundamental
properties of life did not arise spontaneously from inanimate matter but were brought into existence by
molecular machines. This idea, however, does not seem intuitively appealing, so it may be useful to illustrate
it with a metaphor. It is a sort of cartoon, if you like, but if used consistently it is as rigorous as a technical
argument.

The metaphor consists in saying that all spontaneous molecules are ‘grey’ (all shades of grey between
white and black), whereas all manufactured molecules are ‘coloured’ (all colours of the rainbow). With this
terminology, the concept that life is artifact-making amounts to saying that the world of life is coloured
whereas the world of inanimate matter is grey, and this gives us a new way of formulating the problem of the
origins. Earth was a lifeless planet, at the beginning, and all its molecules were grey, so how did coloured
molecules appear out of grey matter?

Spontaneous genes and spontaneous proteins did appear on the primitive Earth but they did not evolve
into the first cells, because spontaneous processes do not have biological specificity. They gave origin to
molecular machines and it was these machines and their products that evolved into the first cells. The
simplest molecular machines that could appear spontaneously on the primitive Earth were molecules that
could stick monomers together at random (bondmakers) or in the order provided by a template (copymakers).
These molecules started manufacturing polymers such as polypeptides, polynucleotides and polysaccharides,
and had the potential to produce them indefinitely, thus increasing dramatically their presence on the
primitive Earth. The unlimited repetition of copying, furthermore, is inevitably accompanied by errors, and
in a world of limited resources a selection is bound to take place. That is how natural selection came into
being, and that is why there is no natural selection in the spontaneous reactions of chemistry.

It must be underlined that the origin of molecular copying does require extremely improbable events. In a
primitive environment where chemical evolution had already accumulated many varieties of organic
molecules, the appearance of bondmakers and copymakers was as likely as that of any other average-size
structure. The origin of proteins, on the other hand, was a much more complex affair, because proteins
cannot be copied and their reproduction required the evolution of supramolecular systems that developed a
code and which can therefore be referred to as codemakers. The evolution of the molecular machines, in
short, started with bondmakers, went on to copymakers and finally gave rise to codemakers.

If we translate all this in the terminology of grey and coloured molecules, we can say that the first
molecular machines were grey (because they appeared spontaneously) and that they started producing
coloured molecules (because manufactured molecules are coloured). The first molecular machines were
therefore a special type of grey molecules, and we may call them ‘silver’ molecules. The machines that came
after them, however, could incorporate also coloured molecules, and eventually these replaced all grey
elements in them. The silver molecular machines evolved into coloured machines and we can illustrate this
transformation by saying that they became ‘golden’ molecular machines. At this stage, the divide between
life and matter became complete, because all the components of life, molecules and molecular machines,
were coloured, whereas all the components of inanimate matter were grey.
1-9 Linear, digital and specific objects

The existence of linear, digital and specific entities in life is a fact, an experimental fact, and all biologists acknowledge it. It is equally a fact that digital and specific sequences and codes do not exist in the inanimate world, so it is beyond dispute that a divide does exist between life and matter. It is the divide between the analog world of chemistry and the digital world of life, and it is not a fiction. The problem is the origin of that divide, not its existence.

Hubert Yockey has underlined that spontaneous reactions cannot produce a living cell, and that, let us repeat it, is formally correct. The real answer to Yockey is not a denial of this point, but the argument that it does not apply to living cells because spontaneous reactions simply do not exist in them. The evidence shows that genes and proteins are manufactured by molecular machines in all present cells, and the most logical conclusion we can draw is that this has been true also for all the cells of the past, including the first cells.

Yockey’s critique of chemical evolution is justified only if we assume that chemical evolution was but a sequence of spontaneous reactions, because linear, digital and specific properties do not exist in spontaneous processes. But they do exist in all manufacturing processes, including those that take place at the molecular level. The answer to Yockey’s argument, in short, is that genes and proteins are molecular artifacts, that life itself is artifact-making (Barbieri, 2003, 2008).

When a copymaker scans a nucleic acid and makes a copy of that molecule, what is happening is precisely an operation that brings into existence a linear, digital and specific copy of a pre-existing molecule. It was molecular copying, the simplest form of artifact-making, that started manufacturing biological objects on the primitive Earth, and that is what started the process that we call life.

That simple beginning is all that was needed to start the odyssey of life on Earth, and we don’t have to rely on extremely complex or extremely unlikely events. But it was a real beginning and what it produced was an absolute novelty in the history of the Universe.

There was a time when atoms did not exist. They came into being within giant stars, and were scattered all over the place when those stars exploded. There was a time when molecules did not exist. They originated from the interaction of atoms in many different places such as comets and planets. There was a time when the world was inhabited only by spontaneously formed molecules, but that period did not last forever. At a certain point molecular machines appeared and the world became also inhabited by manufactured molecules.

By natural artifact

That was the beginning of life, and that is why life arose from matter and yet it is fundamentally different from it. The idea that life is artifact-making is the only logical alternative to the chemical view of life. The divide between life and matter is real because inanimate matter is made of spontaneous structures and life is made of manufactured objects.

1-10 What is Mechanism?

The model of the chemical paradigm is the steam-engine whereas the model of the information paradigm is the computer. Each of them is very different from the clock-model of Descartes, but they are all mechanistic models of life, so we need to ask ourselves ‘what is mechanism?’

One of the expressions that best catches the spirit of mechanism is John Maynard Smith’s statement that “We understand biological phenomena only when we have invented machines with similar properties” (Maynard Smith, 1986).

In fact, ‘understanding’ something means explaining it with a model that we are familiar with, and a machine gives us an immediate sense of familiarity. When we see it working before our eyes, we feel that we ‘know’ it. Actually, we do not even need to build a machine to get this feeling. A description is enough, and so a machine is often just a model, or even an algorithm. One of the most famous machines of all times was built by Turing with just pencil and paper.

A model, furthermore, does not necessarily have a mathematical form. Natural selection, for example, is a mechanistic model which is entirely expressed in words. The important point is that the model has the logic of a machine (i.e. that it delivers the same sense of familiarity that we get from a real functioning machine). Mechanism, in short, is the view that scientific knowledge is obtained by building machine-like models of what we observe in nature. Let us briefly summarize it.

(1) Mechanism is not reductionism, because a machine is a machine not when it is reduced to pieces but when it is put together into a working whole.
(2) Mechanisms is not determinism, because it is more general than classical physics (quantum theory is mechanism, and so is non-equilibrium thermodynamics, chaos theory, complexity theory and the like).

(3) Mechanism is not physicalism, because it is not limited to physical quantities (natural selection, the Turing machine and Godel’s theorem are mechanistic models that are not based on physical quantities).

(4) Finally, and most importantly, mechanism is made of models and models do not coincide with reality (“the map is not the territory”), which means that mechanism is intrinsically incomplete and continuously evolving.

Mechanism, in short, is virtually equivalent to the scientific method. The difference is that the hypotheses of the scientific method are replaced by models, i.e., by descriptions of fully functional working systems. Mechanism, in other words, is ‘scientific modelling’.

Ever since it first appearance, at the beginning of the scientific revolution, mechanism has been highly effective in accounting for particular aspects of Nature, and at the same time it has shown an extraordinary ability to change in the face of adversity. The first mechanistic model of the body was the clock-machine, then came the steam-engine-machine, and after that the computer-machine. Which amounts to saying that mechanism has introduced in biology first mechanical energy, then chemical energy, and finally information.

Now we face a new challenge, and once again we hear that mechanism is not enough, that we need something completely different. Which could be true, of course, but mechanism remains our best chance to find out what makes living systems tick. Mechanism may well be able to change again and introduce in biology not only the concepts of energy and information, but also the last frontier, the concept of meaning.

PART 2

The Code paradigm

2-1 Schrödinger’s prophecy

In 1944, Erwin Schrödinger wrote “What is Life?”, a little book that inspired generations of physicists and biologists and became a landmark in the history of molecular biology. There were two seminal ideas in that book: one was that the genetic material is like an “aperiodic crystal”, the other was that “the chromosomes contain a code-script for the entire organism”. The metaphor of the aperiodic crystal was used by Schrödinger to convey the idea that the atoms of the genetic material must be arranged in a unique pattern in every individual organism, an idea that later was referred to as biological specificity. The metaphor of the code-script was used to express the concept that there must be a miniature code in the hereditary substance, a code that Schrödinger compared to “a Morse code with many characters”, and that was supposed to carry “the highly complicated plan of development of the entire organism” (Schrödinger, 1944). That was the very first time that the word ‘code’ was associated with a biological structure and was given a role in organic life.

The existence of specificity and code at the heart of life led Schrödinger to a third seminal conclusion, an idea that he expressed in the form of a prophecy: “Living matter, while not eluding the ‘laws of physics’ as established up to date, is likely to involve hitherto unknown ‘other laws of physics’, which, however, once they have been revealed, will form just an integral part of this science as the former”. Schrödinger regarded this prophecy as his greatest contribution to biology, indeed he wrote that it was “my only motive for writing this book”, and yet that is the one idea that even according to his strongest supporters did not stand up to scrutiny. Some 30 years later, Gunther Stent (1978) gave up the struggle and concluded that “No ‘other laws of physics’ turned up along the way. Instead, the making and breaking of hydrogen bonds seems to be all there is to understanding the workings of the hereditary substance”.

Schrödinger’s prophecy of new laws of physics appears to have been shipwrecked in a sea of hydrogen bonds, but in reality that is true only in a superficial sense. The essence of the prophecy was the idea that the two basic features of life - specificity and the genetic code - require new fundamental entities of Nature that are “hitherto unknown”, and in that form it is still valid. The fact that Schrödinger invoked new laws of physics should not have obscured the substance of the prophecy, which can be expressed in this way: in order to understand life we need to discover something fundamentally new, something that is still not part of physical theory.

Let us turn therefore to this generalized version of Schrödinger’s prophecy. He anticipated the concept of biological specificity (what today we call biological sequences, or biological information), and announced
that there must be a ‘code-script’ in every living cell. Both ideas were truly prophetic, at the time, and both turned out to be true. That should be enough for us to take a new look at the essence of his prophecy: is it true that we need something fundamentally new in order to explain biological information and the genetic code?

2-2 The ‘special constraints’ solution

In the 1960s, Howard Pattee pointed out that the genetic code is fully compatible with the theory developed by John von Neumann on self-replicating machines. Von Neumann had shown that a self-replicating system capable of open-ended evolution must necessarily contain a description of itself, and such a description must be categorically different from the controlled system (“the map is not the territory”). The description of a system, on the other hand, is necessarily made of entities that represent, or ‘stand for’, its material components, and function therefore as signs or symbols. According to von Neumann, in short, an evolvable self-replicating system must be a physical system controlled by symbols, or, more precisely, by a programme, by the rules of a code (von Neumann, 1951, 1958, 1966).

This was enough, according to Pattee, to prove that every living cell is controlled by a real code, and he set out to find out how physical theory can account for the existence of the genetic code without resorting to the Schrödinger solution of “new laws of physics”. To this purpose, Pattee focussed on the idea that physical theory does not consist only of physical laws, but of laws plus initial conditions and boundary conditions, both of which are often referred to as constraints.

This had been known since Newton’s time, of course, but physicists had consistently assumed that laws are fundamental whereas constraints have only an accessory role. The reality, however, turned out to be very different. Murray Gell-Mann (1994) has underlined that “the effective complexity of the universe receives only a small contribution from the fundamental laws. The rest comes from ‘frozen accidents’, which are precisely the result of constraints. All planets, for example, are formed according to universal physical laws, and yet they are all different. Their individual features are due to the particular constraints of their development, and the distinction between laws and constraints is so important that Eugene Wigner (1964) called it “Newton’s greatest discovery”.

In this novel theoretical framework where laws and constraints have equally fundamental roles, Pattee argued that information and codes are perfectly compatible with physical theory because they have precisely the defining features of constraints. The rules of a code, for example, are limitations that drastically reduce the number of possibilities and can be regarded therefore as true natural constraints. In a similar way, Claude Shannon underlined that information is obtained whenever uncertainty is reduced, and concluded from this that the notions of information and constraint are interchangeable (Shannon, 1948).

The solution proposed by Pattee, in short, is that information and codes do not require new laws of physics, because they are a special type of constraints and constraints are an integral part of physical theory (Pattee, 1968, 1972, 1980, 1995, 2001, 2008). This is the ‘special constraint’ solution to the problem of the genetic code, a solution that is developed in three logical steps: (1) life requires self-replication (a biological principle), (2) evolution requires symbolic control of self-replication (von Neumann), and (3) physics requires that symbols and codes are special types of constraints (Pattee).

Such a conclusion, however, is not entirely satisfactory. It is certainly true that sequences and codes have the defining characteristics of constraints, but not all constraints lead to life, far from it, and it is not enough to say that they must be ‘special’ constraints. What is it that makes them special? What is it that distinguishes the special constraints of information from the special constraints of the genetic code, and what is it that distinguish both of them from the countless constraints of inanimate matter?

2-3 The new observables

Howard Pattee has pointed out that biology does not need new laws of physics because physical theory is based on laws and constraints, and entities like symbols and codes can be regarded as special types of constraints. This is undoubtedly true, but it is not the whole truth. Physical theory starts with the definition of fundamental entities, or observables (time, space, mass etc), and then looks for relationships between them which are referred to as laws and constraints. The basic components of physical theory, in short, are not two but three: laws, constraints, and observables.
The important point here is that the history of physics has not been made only by the discovery of new laws and new constraints, but also by the discovery of new observables. In Newton’s physics, for example, the fundamental observables were time, space and mass, but then electricity required the addition of electric charge and thermodynamics required the addition of temperature.

If we assume a priori that life does not need new observables, we can limit ourselves to laws and constraints, but this is precisely the point that we cannot take for granted. Life is based on the copying of genes and on the coding of proteins and these processes require entities, like biological sequences and the rules of a code, that have all the defining characteristics of new observables. This is because the role of observables is to allow us to describe the world and we simply cannot describe living systems without sequences and codes. But what kind of entities are these new observables?

A biological sequence is a linear chain of units that represents organic information, and a biological code is a set of rules that associate an organic meaning to each unit of information. Sequences and codes, in short, are carriers respectively of organic information and organic meaning, and our problem is to understand the nature of these entities.

According to a long tradition, natural entities are divided into quantities and qualities. Quantities can be measured and are objective, whereas qualities are subjective and cannot be measured. In the case of organic information and organic meaning, however, this scheme breaks down. Organic information, for example, is not a quantity because a specific sequence cannot be measured. But it is not a quality either, because linear specificity is a feature that we find in organic molecules, and is therefore an objective feature of the world, not a subjective one. The same is true for organic meaning. This too cannot be measured, so it is not a quantity, but it is not a quality either because the rules of the genetic code are the same for all observers in all living systems.

A scheme based on quantities and qualities alone, in short, is not enough to describe the world. In addition to quantities (objective and measurable) and qualities (subjective and not-measurable) we must recognize the existence in Nature of a third type of entities (objective but not-measurable).

Organic information and organic meaning belong precisely to that new type of entities, and we can also give them a suitable name. Since organic information and organic meaning can be described only by naming their components, we can say that they are nominable entities, or that they belongs to the class of the nominable entities of Nature (Barbieri, 2004, 2006, 2008).

It must be underlined that the existence of new observables in living systems is perfectly compatible with physics, because observables are an integral part of physical theory and the discovery of new observable has gone on throughout the history of science. Let us take therefore a closer look at these new natural entities and see if we can learn something more about them.

2-4 Names and ‘nominable’ entities

Physical theory consists of laws, constraints and observables, but in addition to these three components there is also a fourth one that should be taken into account, and that is names. Science is always expressed in words and we need therefore to give names to the objects and the processes that we observe in Nature. Names (including those that we call ‘numbers’) are necessarily a fourth essential component of physical theory, but are different from the first three because they change from one language to another. Laws, constraints and observables, in other words, do not depend upon the language that is employed to express them, whereas names are totally language-dependent. This is because names (or nominal entities, to use a classical term) in general have nothing to do with the intrinsic features of the named objects, and are therefore mere labels that we attach to them.

The deep divide that exists between ‘names’ and ‘objects’ has been at the centre of many controversies in the past, in particular of the celebrated medieval dispute over ‘nominal entities’ and ‘real entities’. It has also had a long history in the philosophy of mathematics, where some have argued that numbers are ‘invented’ by the human mind, and others that they are ‘discovered’, a conclusion which implies that they have an existence of their own in some abstract Platonic world.

The relationship between names and objects is also a crucial issue in science, but here it has taken on a new form. Let us underline that all names are sequences of characters (alphabetic, numerical or alpha-numerical) and that each sequence is unique. Names, in other words, have specificity. In general, the specificity of a name has nothing to do with the characteristics of the named object, and in these cases we
can truly say that names are mere labels. Science, however, has invented a new type of names where the sequence of characters does represent an order that is objectively present in the named objects.

The chemical formula of a molecule, for example, describes an objective sequence of atoms, and any atom can be described by the objective sequence of its quantum numbers. In these cases, the names are no longer arbitrary labels but true ‘observables’ because they describe characteristics that we observe in Nature.

This shows that there are two distinct types of names in science: labels and observables.

In the case of the observables, furthermore, there is another distinction that must be considered. When a molecule is formed spontaneously, its final sequence is due to the interactions between its own components, and in most cases it is completely determined by them. In the case of a protein, however, all its different amino acids interact by the same peptide bonds and a spontaneous assembly would produce a completely random order (which is incompatible with life). In this case, a specific sequence can be obtained only if the amino acids are put together by a molecular machine according to the order provided by a template that is external to the protein itself. We need therefore to distinguish between two different types of observables.

The sequence of quantum numbers in an atom, or the sequence of atoms in inorganic molecules, is determined from without, by external templates. In the first case the sequence is a physically computable entity, in the sense that it is the automatic result of physical forces, whereas in the second case it can only be described by ‘naming’ its components, and is therefore a nominable entity (this term should not be confused with the classical concept of nominal entity, which applies to all names). A nominable entity is not a label but an observable, and more precisely a non-computable observable.

All names, in conclusion, are specific sequences of characters, and in science they can be divided into two great classes: labels and observables. The observables, in turn, can be divided into computable entities and nominable entities. The important point is that physics and chemistry deal exclusively with computable entities (physical quantities), whereas nominable entities (information and coding rules) exist only in living systems. We need therefore to pay a special attention to these new observables, and make sure that they truly are fundamental entities of Nature.

2-5 Organic information

In genes and proteins, biological, or organic, information has been defined as the specific sequence of their subunits. This definition however is not entirely satisfactory because it gives the impression that information is a static property, something that molecules have simply because they have a sequence. In reality, there are countless molecules which have a sequence but only in a few cases this becomes information. That happens only when copymakers use it as a guideline for copying. Even copymakers, however, do not account, by themselves, for information. Copymakers can stick subunits together and produce sequences, but without a template they would produce only random sequences, not specific ones. Sequences alone or copymakers alone, in other words, have nothing to do with information. It is only when a sequence provides a guideline to a copymaker that it becomes information for it. It is only an act of copying, in other words, that brings organic information into existence.

This tells us that organic information is not just the specific sequence of a molecule, but the specific sequence produced by a copying process. This definition underlines the fact that organic information is not a thing or a property, but the result of a process. It is, more precisely, an 'operative' definition, because information is defined by the process that brings it into existence. We realize in this way that organic information is as real as the copying process that generates it.

We have also seen that organic information is neither a quantity (because a specific sequence cannot be measured), nor a quality (because it is an objective feature of all copied molecules), and belongs instead to a third class of objects that have been referred to as nominable entities (Barbieri, 2004, 2006, 2008).

We conclude that organic information is a new type of objects, and that it is essential to describe the organic molecules of Nature. To this purpose, in fact, it is no less essential than the physical quantities, and this means that organic information has the same scientific 'status' as a physical quantity. They both belong to the class of objective and reproducible entities that allow us to describe the world.

This conclusion, however, raises immediately a new problem, because there are two distinct groups of physical quantities: a small group of fundamental quantities (space, time, mass, charge and temperature) and a much larger group of derived quantities. That distinction applies to all objective entities, so we need to find out whether organic information belongs to the first or to the second group.
Luckily, this problem has a straightforward solution because the sequences of genes and proteins have two very special characteristics. One is that a change in a single component of a biological sequence may produce a sequence which has entirely new properties. This means that although a biological sequence can be said to have ‘components’, it is at the same time a single indivisible whole. The second outstanding feature is that from the knowledge of n elements of a biological sequence we cannot predict the element (n+1). This is equivalent to saying that a specific sequence cannot be described by anything simpler than itself, so it cannot be a derived entity.

We conclude that organic information has the same scientific status as the physical quantities, because it is an objective and reproducible entity. But we also conclude that it does not have the status of a derived physical quantity because it cannot be expressed by anything simpler than itself. This means that organic information has the same scientific status as the fundamental quantities of physics, and is therefore a new irreducible entity of Nature, i.e., a new fundamental observable.

2.6 Organic meaning

A code is a set of rules which establish a correspondence between the objects of two independent worlds. The Morse code, for example, is a correspondence between groups of dots and dashes with the letters of the alphabet, and in the same way the genetic code is a correspondence between groups of nucleotides and amino acids. Let us notice now that establishing a correspondence between, say, object 1 and object 2, is equivalent to saying that object 2 is the meaning of object 1. In the Morse code, for example, the rule that ‘dot-dash’ corresponds to the letter ‘A’, is equivalent to saying that letter ‘A’ is the meaning of ‘dot-dash’. In the code of the English language, the mental object of the sound ‘apple’ is associated to the mental object of the fruit ‘apple’, and this is equivalent to saying that that fruit is the meaning of that sound.

By the same token, the rule of the genetic code that a group of three nucleotides (a codon) corresponds to an amino acid is equivalent to saying that that amino acid is the organic meaning of that codon. Anywhere there is a code, be it in the mental or in the organic world, there is meaning. We can say, therefore, that meaning is an entity which is related to another entity by a code, and that organic meaning exists whenever an organic code exists (Barbieri, 2003, 2008).

The existence of meaning in the organic world may seem strange, at first, but in reality it is no more strange than the existence of a code because they are the two sides of the same coin. To say that a code establishes a correspondence between two entities is equivalent to saying that one entity is the meaning of the other, so we cannot have codes without meaning or meaning without codes. All we need to keep in mind is that meaning is a mental entity when the code is between mental objects, but it is an organic entity when the code is between organic molecules.

Modern biology has readily accepted the concept of information but has carefully avoided the concept of meaning, and yet organic information and organic meaning are both the result of natural processes. Just as it is an act of copying that creates organic information, so it is an act of coding that creates organic meaning. Copying and coding are the processes; copymakers and codemakers are their agents; organic information and organic meaning are their results.

But the parallel goes even further. We have seen that organic information cannot be measured, and the same is true for organic meaning. We have seen that organic information is an objective entity, because it is defined by the same sequence for any number of observers, and that is also true for organic meaning, which is defined by coding rules that are the same for all observers. Finally, we have seen that organic information is an irreducible entity, because it cannot be described by anything simpler than its sequence, and the same is true for organic meaning, which cannot be defined by anything simpler than its coding rules.

Organic information and organic meaning, in short, belong to the same class of entities because they have the same defining characteristics: they both are objective-but-not-measurable entities, they both are fundamental entities because they cannot be reduced to anything simpler, and they both are nominable entities because we can describe them only by naming their components (Barbieri, 2004, 2008).

Finally, let us underline that they are the twin pillars of life because organic information comes from the copying process that produces genes, while organic meaning comes from the coding process that generates proteins.
Physical quantities have three fundamental properties: (1) they are objective, (2) they are reproducible, and (3) they are defined by operative procedures. This last property is particularly important because it has provided the solution to one of the most controversial issues of physics. The controversy was about the theoretical possibility that the entity which is measured may not be the same entity which has been defined. This led to the idea that there should be no difference between what is measured and what is defined, i.e., to the concept of operative (or operational) definition: a physical quantity is defined by the operations that are carried out in order to measure it.

It was this operational approach that solved the definition problem in physics, and it is worth noticing that we can easily generalize it. Rather than saying that a natural entity is defined by the operations that measure it, we can say that a natural entity is defined by the operations that evaluate it. The advantage of this generalized formulation is that it applies to all objective entities, so it can be used not only in physics, but in biology as well. To this purpose, we only need to notice that a measurement is an objective and reproducible description of a physical quantity, just as the naming of a specific sequence is an objective and reproducible description of organic information, and just as the naming of a coded entity is an objective and reproducible description of organic meaning.

Whereas the physical quantities are evaluated by measuring, sequences and codes are evaluated by naming their components, but in both cases the entities in question are defined by the operations that evaluate them, and this is the essence of the operative approach. We may add that organic information and organic meaning can also be defined by the processes of copying and coding that bring them into existence, and that too amounts to an operative definition (Barbieri, 2003, 2008).

We conclude that organic information and organic meaning can be defined by generalized operative procedures that are as reliable as the operative procedures of physics. This means that the definitions of information and meaning should no longer be at the mercy of endless debates on terminology as they have been in the past. The operative definitions are scientific tools which are justified by their own prescriptions, so there is no point in asking whether they are right or wrong. All we can ask of them is whether they contribute or not to our description and to our understanding of Nature.

At this point, we can summarize all the above arguments with the following concepts:

1. The sequence used by a copymaker during a copying process is organic information.
2. The sequence produced by a codemaker during a coding process is an organic meaning.
3. Organic information and organic meaning are neither quantities nor qualities. They are a new kind of natural entities that are referred to as nominable entities.
4. Organic information and organic meanings have the same scientific status as the quantities of physics because they are objective and reproducible entities that can be defined by operative procedures.
5. Organic information and organic meanings have the same scientific status as the fundamental quantities of physics because they cannot be reduced to, or derived from, simpler entities.

The discoveries of the double helix and of the genetic code are the two pillars of modern biology, but there is a strange discrepancy between them. The first brought biological information to light and that concept was immediately accepted into modern biology. The genetic code revealed the existence of biological meaning - because any code is a correspondence between signs and meanings - but that concept has been completely ignored by modern biology.

It is often said that the concept of meaning has also been kept out of Information Theory, but that is not exactly the case. Information theory has certainly made a clear separation between information and meaning, but has not ignored meaning. On the contrary, the mobile telephone, to name just one example, would not even exist without the introduction of error-correcting codes (Battail, 2007), and almost all applications of Information Theory are heavily dependent on such codes. Information theory, in other words, does deal with codes, and therefore with meaning, but keeps them sharply distinct from information.

In biology, however, no such clear distinction has been made, and meaning has been regarded not as an entity in its own right, but as a ‘qualification’ of information. Rather than talking of information and meaning, many biologists are talking of “meaningful information”, “semantic information”, “functional information” and the like.
In a recent review entitled “Information in Biological Systems” John Collier (2008) has listed at least seven different types of information that apparently form a nested hierarchy: (1) physical information (or “It from bit” information), (2) statistical information (or “negentropy”), (3) expressed information, (4) functional information, (5) meaningful information, (6) intentional information, and (7) social information.

Similar proposals have been made by many other authors with different terminologies, and there seem to be no end in sight to the proliferation of the information categories. But why does this happen? Why do we keep multiplying the types of information in order to account for properties that belong to the category of meaning?

It is high time to acknowledge that in biology too we must face the issue of meaning, and to this purpose we should treasure the example of the communication sciences. We should accept that information and meaning are two distinct entities and stop trying to reduce one to the other.

The important point, at any rate, is that a genetic code exists in every cell, a fact which tells us that there are two distinct fundamental processes at the basis of life. The coding of proteins is as essential as the copying of genes and this implies that biological meaning is as necessary as biological information in living systems. This conclusion is nothing less than a new theoretical framework, and we have therefore, three distinct paradigms in modern biology.

In addition to the idea that ‘life is chemistry’, and to the idea that ‘life is chemistry-plus-information’, we have a third paradigm which states that ‘life is chemistry-plus-information-plus-codes’. This is the Code paradigm, the idea that life is based on copying and coding, that we need to introduce in biology not only the concept of biological information but also the concept of biological meaning.

2.9 The discovery of new worlds

The history of physics tells us that scientific discoveries require three logical steps. First we look at the world and choose a certain number of entities to describe it, entities that are called observables (space, time, mass, etc.) precisely because they represent what we observe. Then we look for relationships between observables and obtain models of the observed phenomena (regularities, equations, laws, etc.). Finally we use our models to make predictions that test them (we predict, for example, the nest eclipse of the moon etc.).

The choice of the observables is the first step in the procedure and the most critical. The movements of planets and stars, for example, can be described with only two observables - space and time - and in that case we get either a Ptolemaic model or a Copernican system. By introducing a third observable - mass - we obtain the laws of motion, universal gravitation and the Newton model of the world.

The three basic observables of classical physics can be combined together in different ways and produce many other derived observables (velocity, acceleration, force, energy, power, momentum, etc.), but what defines the whole system is the initial number of fundamental observables. The actual identity of these observables can be changed (space and time, for example, can be replaced by velocity and time, and in that case space becomes a derived entity), but the minimum number of fundamental observables does not change.

That number defines a whole world of phenomena, and we can discover new worlds, i.e., new aspects of reality, only if we discover new fundamental observables. The world of electricity and magnetism, for example, required precisely the introduction of new fundamental observables, and so did the world of thermodynamics, the world of nuclear forces, and the world of elementary particles. All of which takes us to a question: do we need new observables in the world of life or not? This point is crucial, and the different paradigms of biology are nothing less than different ways to answer it.

The chemical paradigm states a priori that we do not need new observables to describe living systems, i.e., that life is completely described, in principle, by the quantities of physics. The information paradigm claims that information is a fundamental entity that exists only in living systems, but it has not been able to contrast the physicalist charge that there is nothing fundamental in it.

We can prove that this charge is wrong only by showing that information is a new observable and this can be done only by showing that information is the result of a manufacturing process by molecular copying. But as soon as we accept the reality of molecular copying we must also accept the reality of molecular coding, and therefore of another fundamental observable. This is the third paradigm of modern biology, the Code view of life, the idea that life is artifact-making by copying and coding.

The crucial point is that the existence of two new observables in living systems is not a hypothesis. It is an experimental fact. We can prove that biological sequences (organic information) and the rules of a code (organic meaning) are fundamental observables with the same procedures that we have used in the case of...
space, time, mass, temperature, etc. The only difference is that sequences and coding rules are non-computable observables, but there is no doubt that observables they are (we do observe them in living systems) and that they are fundamental observables (because we cannot describe living systems without them and because we cannot reduce them to anything else).

The discovery of classical physics, the discovery of thermodynamics, the discoveries of electromagnetism and of elementary particles, were all based on the discoveries of new fundamental observables, and now we realize that this is true also in biology. Life is indeed a new world, a new dimension of reality, because it is the result of copying and coding processes that bring two new fundamental observables into existence.

2-10 The unexpected results of coding

The organic codes may give the impression of being deterministic rules that turn living systems into biological robots, but this far from the truth. They are, in fact, the actual tools that bring creativity into life. It is the rules of grammar, for example, that allow us to create endless combinations of words and generate the universe of language and literature. The key feature of the organic codes is the fact that they bring absolute novelties into existence and in so doing they produce objects that turn out to have totally unexpected properties. This is a crucial point, and in order to illustrate it let us start from the case of those particular human artifacts that we call ‘numbers’.

There is little doubt that numbers were generated by counting and that counting was favoured by natural selection because it has practical advantages. The process of counting, however, produces exclusively natural numbers, but then we have discovered prime numbers, fractional numbers, rational and irrational numbers, real and imaginary numbers, and in so doing we have brought to light an endless stream of mathematical theorems. All these additional entities were not produced by counting, and this is why some mathematicians say that natural numbers were invented by man but all other rules of mathematics had to be discovered, as if they had an existence of their own.

The world of mathematics was generated by the ‘genetic’ rule of counting and then it developed into an increasingly complex world full of additional, or ‘epigenetic’ properties. A world of codified objects, in short, is a world of artifacts, and it is only partially determined by the coding rules that generate the artifacts. In general, it turns out to have unexpected ‘rules of its own’, rules that we call epigenetic because they were not present at the beginning and are brought to light only by processes of exploration and discovery.

This is what we actually find in living systems. In the world of proteins, for example, there is a universal mechanism in every cell that produces linear polypeptides from linear sequences of genes, but then the polypeptides fold themselves up into three-dimensional structures and take up forms that were not written in the genes. That generates a whole new world of objects, and living cells appear to engage in a veritable exploration of the potentialities of the protein universe.

Another outstanding example is the body-plan of animals. It is based on instructions that specify only three essential relationships between the cells of the body (up and down, back and front, left and right) and yet the number of morphological designs that can be built with them is virtually unlimited.

Language, mathematics, proteins and animals are very different worlds, but deep down there is something in common between them. They all have (1) a ‘genetic’ algorithm that produces the objects of a potentially unlimited new world of artifacts (words, numbers, proteins and bodies) and (2) an exploratory procedure that brings into existence additional or ‘epigenetic’ properties of the new world that were not written in the coding rules and were not present at the beginning.

The organic codes, in conclusion, do not explain everything, far from it. They just account for coding. They code for objects that are absolute novelties and which have unpredictable properties. Far from being deterministic rules, the organic codes are the quintessential instruments of creativity and the higher their number the greater is the creative potential of a system. But they account only for the generative rules of life, not for the flesh and blood of history.

Conclusion

The discoveries of the double helix and of the genetic code have been two of the major scientific revolutions of all times and yet the majority view, today, is still the idea that life is an extremely complex form of chemistry. This view is based on the physicalist thesis that all biological processes are reducible, in principle,
to physical quantities, so there is nothing fundamental in genetic information and in the genetic code because they are not physical quantities. They are regarded as metaphorical or teleological terms that we use only because they are intuitively appealing.

This is the great paradox of modern biology. On the one hand, genetic information and the genetic code have become the very basis of biological research, and at the same time we are told that they are little more than linguistic decorations. This paradox is due to the fact that the information paradigm has not been able to offer a convincing alternative to the physicalist thesis.

Here we have seen that such an alternative does exist, because the physicalist thesis is valid only in spontaneous systems, whereas genes and proteins are never formed by spontaneous reactions. They are invariably manufactured by molecular machines, and all manufacturing processes do not require only physical quantities but also additional entities like sequences and coding rules. The alternative to the view that ‘life is chemistry’, in short, is the view that ‘life is artifact-making’.

The charge that information is a teleological concept is simply false, notwithstanding the fact that it is repeated fairly often. The truth is precisely the other way round. Information has all the defining features of a scientific concept because it has been defined in two different ways and in both cases there is nothing teleological about it.

(1) When it is defined by Shannon’s approach, information is actually expressed by a formula, like any other standard physical quantity.

(2) When it is defined by a sequence, information is no longer measurable, but it is still an essential parameter because it is absolutely necessary to the description of a living system.

We simply cannot describe the transmission of genes or the synthesis of proteins without their sequences, and we cannot replace sequences with anything else, which means that using information to describe living systems is perfectly equivalent to using space, time, mass and energy to describe physical systems.

The truth, in other words, is that there is no more teleology in information and in the genetic code than there is in the quantities of physics and chemistry. Sequences (biological information) and the rules of codes (biological meaning) are descriptive entities and are absolutely essential to the scientific study of life.

Unfortunately, the information paradigm has accepted the concept of information but not the concept of meaning, and this is equivalent to saying that genetic information is real but the genetic code is not. What we need, therefore, is a new paradigm that fully accepts the implications of the discovery of the genetic code.

The implication that ‘life is artifact-making’, that life is based on copying and coding. This is the code paradigm, the theoretical framework where biological sequences (organic information) and the rules of a code (organic meaning) are as real and fundamental as the fundamental quantities of physics.

References


von Helmont, J.B. (1648). *Ortus Medicinae*. Amsterdam


