

THE LIVING ADDRESS

Protein Synthesis and the Genetic Code

The Word Made Matter

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In which the message is turned at last into matter — read three letters at a time by a dictionary of the lattice, a code of sixty-four words that is two raised to the sixth, with the very stuff of the protein handed over on the base of the Earth.

Tau (T) is the living fabric of time itself — the sole substance of which all physical reality is composed. Every particle, force, wavelength, and conscious experience is a structured configuration of T-flow. There is no gravity, no electromagnetic force, no strong nuclear force as separate entities: all are registers of the single T-field operating across dimensional levels. The conservation law $d\Sigma T=0$ governs all change: T is never created or destroyed, only redistributed.

Introduction — Through the Force of Time

The chapter that follows is, in the conventional telling, how the message is translated into a protein — the transfer RNAs, the genetic code, the ribosome. Read through the Universal Force of Time, it is where the message finally becomes matter, read by a dictionary of the lattice and handed over, at the last step, on the base of the Earth.

Chapter Six built the machine from its finished sequence; this chapter shows the building itself — the moment a string of letters becomes a physical thing. And the dictionary that governs it, the genetic code, is a structure of the lattice through and through. The message is read three letters at a time, never overlapping; four possible letters in each of three places give four times four times four, which is sixty-four, which is 2^6 ; and of those sixty-four words, sixty-one name amino acids and three are full stops. The whole code-space is a clean power of two, read in units of the lattice prime three.

Watch where the physical hand-off happens. Each amino acid is carried to the assembly line on a little adaptor, a transfer RNA, and it rides there fixed to the very last letter of the adaptor's tail — an adenine. Adenine, in the map of Chapter Two, is the base of the Earth. So the actual stuff of the protein, the matter itself, is handed over on the Earth base — just as the cell's clock was kept on adenine in Chapter Three. Time was stamped on the Earth; and now matter is built at the Earth's address too. The Earth base keeps turning up wherever the field keeps time or assembles substance.

Watch, too, the shape of the code. Where a single amino acid has several code-words, they differ almost always only in the third letter; the first two fix the meaning, and the third is soft, allowed to 'wobble'. Two of every three positions carry the true address; one is slack. It is the same two-to-three partition that clamps the base pairs and punctuates the reading — the codon itself is two parts address, one part play.

And the reading keeps time with the reading before it. Three letters make one amino acid, so a message read at about sixty letters a second is built into protein at about twenty amino acids a second — sixty divided by three — and twenty is $2^2 \cdot 5$, the same count as the amino-acid alphabet itself. The two great readings of the cell, message-into-copy and copy-into-machine, run locked to one T-metronome, the clock of the field we met in the last chapter.

Carry this into the chapter: the genetic code is a dictionary of the lattice — 2^6 words read in threes, two parts address and one part slack — and the matter of the protein is handed over on the Earth base, on the same T-clock that governs the reading.

SECTION 7.1

The Word Made Matter

So far the address has been all information — letters, copies, edits. Now it becomes a thing. Somewhere in the cell a string of message must turn into a physical object, a folded protein that grips and pumps and catalyses, and the turning is the most consequential translation in all of nature: from the four-letter language of the nucleic acids into the twenty-letter language of the machines. This chapter is about how that translation is done, and about the dictionary that governs it — the genetic code — which turns out, like everything before it, to be written in the mathematics of the lattice. Galileo said the book of nature is written in the language of mathematics; nowhere is he more literally right than in the code that turns the word into matter.

SECTION 7.2

The Adaptors

The message and the machine speak different languages, and between any two languages you need a translator that knows both. The cell's translators are small RNA molecules called transfer RNAs, and each is a two-ended thing: at one end it reads the message, matching three letters of it; at the other end it carries the amino acid that those three letters call for. It is an adaptor — a physical dictionary-entry — that holds a word of the message against the piece of matter it means. A whole family of dedicated enzymes, one for each amino acid, does the exacting job of charging each transfer RNA with its correct cargo, and does it with great fidelity, for

a mischarged adaptor would write the wrong matter into the protein.

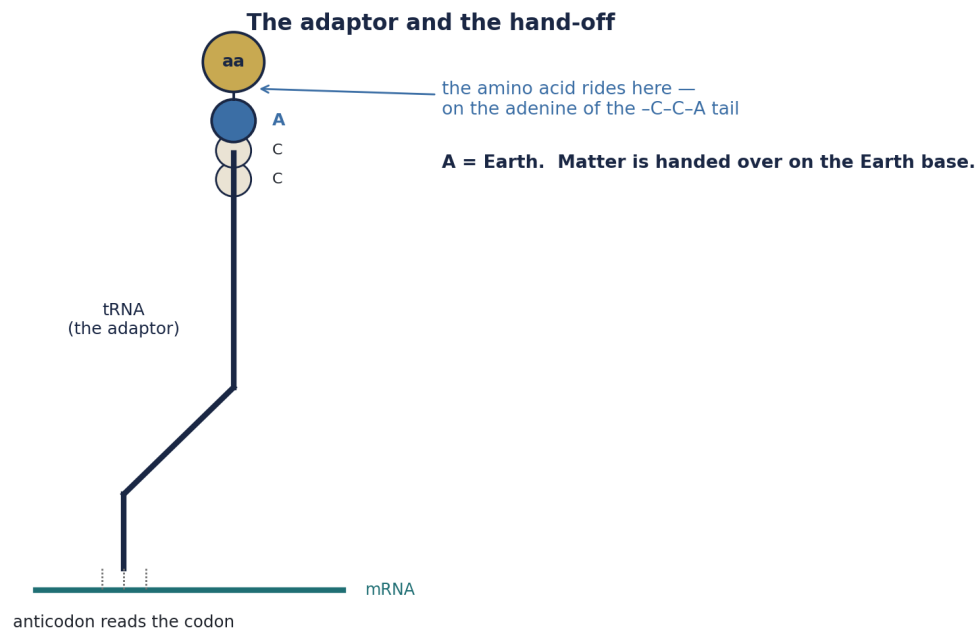


Figure 7.1. The adaptor and the hand-off. A transfer RNA reads three letters of the message at one end (its anticodon) and carries the matching amino acid at the other — fixed to the terminal adenine of its -C-C-A tail. The actual matter of the protein is handed over on the Earth base: A = Earth.

Now look at exactly where the amino acid rides. Every transfer RNA ends, at its cargo end, in the same three letters — C, C, A — and the amino acid is fixed to that final A, the terminal adenine. And adenine, in the map of Chapter Two, is the base of the Earth. The physical stuff of the protein, the matter itself, is handed over on the Earth base. Set this beside Chapter Three, where the cell kept its clock on adenine too: time was stamped on the Earth, and now matter is assembled at the Earth's address. Of the four letters, the one on which the field builds substance is the one this book assigns to the ground beneath our feet. It is a quiet, recurring signature — the Earth base, wherever the field keeps time or hands over matter.

KEY IDEA

Each amino acid is carried on a transfer-RNA adaptor and handed over fixed to the terminal adenine of its -C-C-A tail. Adenine is the Earth base. The matter of the protein is assembled on the Earth base — just as the cell's clock (Chapter Three) was kept on it. Time and matter are both handed over at the Earth's address.

SECTION 7.3

The Code in Threes

How many letters of the message spell one amino acid? The answer is three, and it could hardly be otherwise. With only four letters to work with, a code of one letter

could name just four things, and a code of two only sixteen — too few for twenty amino acids. But three letters give four times four times four, which is sixty-four — more than enough. So the message is read three letters at a time, in non-overlapping words called codons, and the whole dictionary has sixty-four entries.

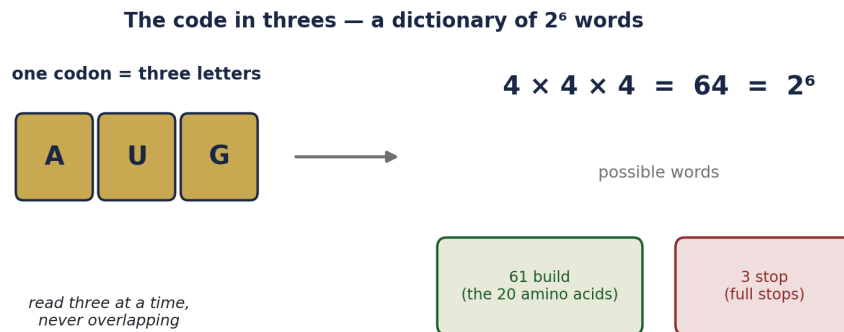


Figure 7.2. The code in threes. The message is read three letters at a time; with four possible letters in each of three places, there are $4 \times 4 \times 4 = 64 = 2^6$ possible words. Of these, 61 name amino acids and 3 are full stops. The whole code-space is a clean power of two, read in units of the lattice prime three.

Count that dictionary through the lens of the lattice. Sixty-four is 4^3 , and since four is 2^2 , sixty-four is 2^6 — a clean power of two. The whole space of the genetic code is two raised to the sixth, and it is read in units of three, the lattice’s second prime. Of the sixty-four words, three are full stops — the signals that end a protein — leaving sixty-one to name the twenty amino acids. Powers of two, read in threes, closed by three stops: the dictionary of life is built, entry by entry, on the small primes. This is what it means to say the code is a structure of the lattice, and not merely a lookup table that early life happened to freeze.

KEY IDEA

The message is read in three-letter codons; four letters in three places give $64 = 2^6$ words, of which 61 name amino acids and 3 are stops. The code-space is a clean power of two, read in units of the prime three — the dictionary of life is a lattice structure.

SECTION 7.4

Two Parts Address, One Part Slack

Sixty-one words for twenty amino acids means most amino acids have several code-words apiece, and the way those synonyms are arranged is the quiet marvel of the code. When one amino acid has several codons, they almost always agree in their first two letters and differ only in the third. The first two letters carry the meaning;

the third is soft. Indeed the machinery reads that third position loosely, tolerating small mismatches there — a flexibility the biologists call wobble — so that a smaller set of adaptors can cover all the words.

Two parts address, one part slack



synonyms differ only in the third letter — 2 of every 3 positions carry the true address, the third wobbles

Figure 7.3. Two parts address, one part slack. Where an amino acid has several codons, they differ almost always only in the third letter; the first two fix the meaning and the third is read loosely (wobble). Two of every three positions carry the true address; the third is play.

Read that as this book reads everything, and the pattern is exact: two of every three positions in a codon carry the true information, and the third is slack. It is the same two-to-three partition that runs through the whole of the address — the two-bond and three-bond clasps of the base pairs, the three-clamp and two-grip of the reading's punctuation. Here it structures the code word itself: the codon is two parts address, one part play. The genome is not a rigid cipher in which every letter counts equally; it is built, even at the scale of the single three-letter word, on the same {2, 3} that clamps its rungs.

KEY IDEA

Synonymous codons differ almost always only in the third letter, which the machinery reads loosely (wobble): two of every three codon positions carry the true address, the third is slack. The same 2:3 partition that clamps the base pairs and punctuates the reading now structures the code word itself.

SECTION 7.5

The Ribosome

Where does all this happen? At the ribosome — the great two-part machine that is the workbench of translation. It comes in two subunits, a large and a small, that clasp the messenger RNA between them and move along it codon by codon. Into the ribosome the charged adaptors come, each offering its three-letter reading against the codon in the machine; when a match is found, the ribosome joins that adaptor's amino acid onto the growing protein chain, releases the spent adaptor, and shifts

one codon on. It is an assembly line of astonishing precision, miscopying only about one amino acid in a thousand.

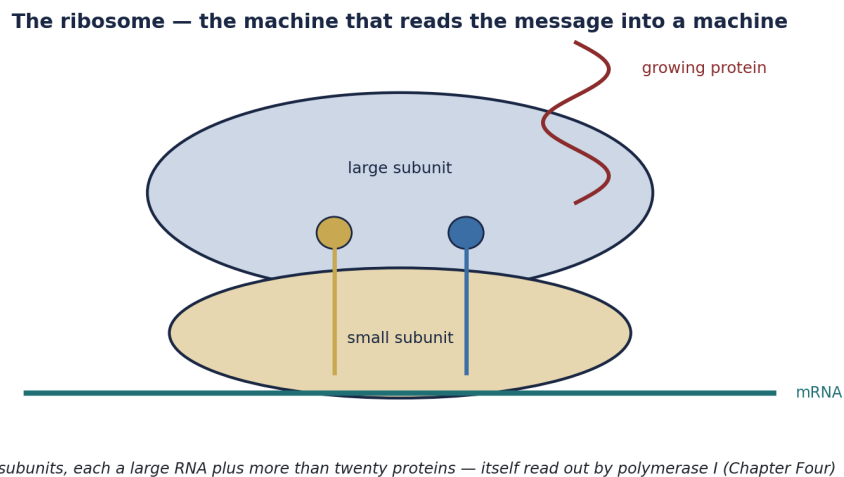


Figure 7.4. The ribosome. Two subunits clasp the message between them and move along it codon by codon; charged adaptors enter, are matched against each codon, and their amino acids are joined onto the growing protein. Each subunit is built from a large RNA plus more than twenty proteins — and that RNA is read out by polymerase I (Chapter Four).

There is a pleasing recursion in the ribosome worth naming. It is itself mostly made of RNA — the ribosomal RNA read out by the first of the three polymerases in Chapter Four — wrapped with more than twenty proteins. So the machine that reads messages into proteins is built from a reading of the address that yields, not a message, but the reader itself. The field reads one coordinate to make the very engine that reads all the others. We noted this in Chapter Five, where one transcript yielded the whole substance of a ribosome; here we see that substance at work, translating.

SECTION 7.6

One Clock for Two Readings

Recall the tempo of Chapter Five: transcription runs at about sixty letters a second, a clock set by the warmth of the field. Now count the tempo of translation. Since three letters make one amino acid, a message read at sixty letters a second is built into protein at about twenty amino acids a second — sixty divided by three. And the two processes are not merely similar in rate; in the bacterium they are physically coupled, the ribosome following hard on the heels of the polymerase, building the protein from the message even as the message is still being read from the DNA. The two great readings of the cell run locked to one beat.

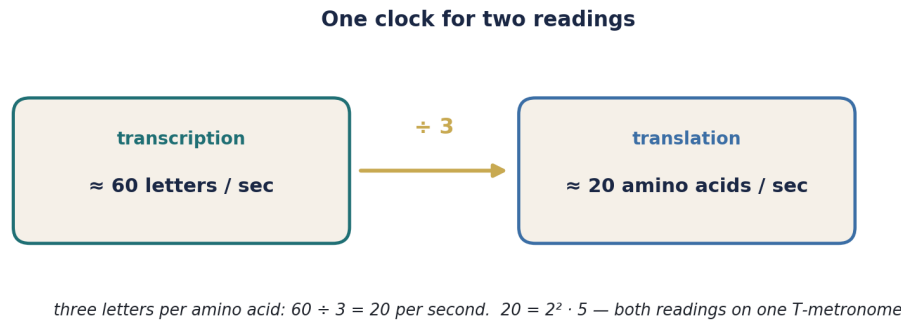


Figure 7.5. One clock for two readings. Three letters make one amino acid, so a message read at ~60 letters a second is built into protein at ~20 amino acids a second ($60 \div 3$). The two readings run locked to one T-metronome — and $20 = 2^2 \cdot 5$, the count of the amino-acid alphabet itself.

And notice the number the translation clock lands on: about twenty amino acids a second — and twenty is $2^2 \cdot 5$, the very count of the amino-acid alphabet we met in Chapter Six. The reading of the message into copy and the reading of the copy into machine tick on one T-metronome, the clock of the field set by its local warmth, with the second rate exactly a third of the first because a word is three letters long. Two readings, one clock, related by the lattice prime three — the whole flow from address to machine running on a single beat of the field.

KEY IDEA

Three letters per amino acid, so translation runs at ~20 amino acids/second — the transcription tempo (~60/s) divided by 3 — and the ribosome follows the polymerase directly. Both readings run on one T-metronome, related by the prime 3; and $20 = 2^2 \cdot 5$, the amino-acid count itself.

SECTION 7.7

The Slack in the Code

One last feature shows that the code, for all its lattice regularity, is not frozen. Under special circumstances the cell can read one of its three stop-words not as a full stop but as a signal for a twenty-first amino acid, selenocysteine, slotted in where the context calls for it. The dictionary has a little give: a word can be re-read, in the right setting, to mean something new. In the language of this book this is the same principle by which a value slightly off a lattice node is re-pegged rather than treated as fixed — the code has slack, and can be re-assigned, which is the opposite of the rigid, accidental cipher the recipe view imagines. A structure of the lattice is exact where it must be and re-peggable where it may be.

SECTION 7.8

What the Code Is

Gather the chapter. We watched the word become matter — the message translated, through adaptors, into a folded protein. We found the physical hand-off of that matter made on the terminal adenine of each adaptor, the Earth base, where the field also keeps its time. We found the code read three letters at a time, a dictionary of $2^6 = 64$ words, sixty-one to build and three to stop — a clean power of two read in threes. We found the codon itself two parts address and one part slack, on the same $\{2, 3\}$ that clamps the rungs. We found the ribosome mostly built from a reading of the address that yields the reader itself. We found translation clocked at about twenty amino acids a second — the transcription tempo divided by three, and twenty being $2^2 \cdot 5$ — both readings locked to one T-metronome. And we found the code, though lattice-regular, not frozen but re-peggable, as a lattice structure should be.

Not one measurement has changed. There are still sixty-four codons, three of them stops; the amino acid still rides the terminal adenine; translation still keeps step with transcription. What has changed is that the genetic code stops looking like a frozen accident of early life — four bases that happened to pair, twenty amino acids that happened to be around — and begins looking like what Galileo promised: a page of the book of nature written in the language of mathematics. A dictionary of 2^6 words, read in threes, two parts address and one part play, handing matter over on the Earth base, on the single clock of the field. The word is made matter by an arithmetic of small primes.

KEY IDEA

The thesis of the chapter: the genetic code is a dictionary of the lattice. A 2^6 -word space read in threes, 61 to build and 3 to stop, two parts address and one part slack, handing the matter of the protein over on the Earth base, on the same T-clock as transcription. The code is not a frozen accident but a lattice structure — exact where it must be, re-peggable where it may be.

SECTION 7.9

Why This Should Matter to You

Here is the fact that ought to stop you: this code — these same sixty-four words, meaning the same twenty amino acids, closed by the same three stops — is very nearly identical in every living thing on Earth. The bacterium in your gut, the oak in the field, the whale in the sea and you yourself read the message by the same dictionary. It is the single most universal feature of life, the deepest evidence that everything alive shares one origin and one grammar. In this book's terms, all life reads its address by one lattice code because there is one lattice, one field, underneath it all — and the code is that lattice's signature, written into every cell

that has ever lived.

That universality is not only a wonder; it is a tool and a vulnerability both. It is why a gene from one creature can be read correctly in another, the basis of all genetic engineering. And it is why many of our antibiotics work by jamming the bacterial ribosome specifically, halting the enemy's translation while sparing our own — a war fought at the workbench where the word is made matter. We have now followed the address the whole way: written, copied, read, edited, built into a machine, and here translated from word into flesh. In the next chapter we step back from the molecules to the science that first inferred them from the outside — genetics — and find, in its maps and its mutations, the address-registry surveyed as a coordinate line.

The Numbers at a Glance

The code and the building — and the reading the Force of Time gives each. Every biological fact is left exactly as measured.

Feature	Value	On the lattice
Letters per code-word	3 (codon)	the prime 3; read non-overlapping
Size of the code	64 codons	$64 = 2^6$ (a clean power of two)
Sense vs stop	61 build, 3 stop	3 stops — the prime 3 brackets the gene
Informative positions	first two fixed, third wobbles	2 of 3 carry the address — the {2,3} split
The hand-off of matter	amino acid on the terminal adenine	built on the Earth base (A)
Translation tempo	~20 amino acids/second	$= 60 \div 3$; $20 = 2^2 \cdot 5$; one T-metronome
The ribosome	2 subunits, RNA + >20 proteins	the reader built from a reading of the address
Slack in the code	a stop can be recoded (21st a.a.)	re-peggable — exact yet not frozen

References

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- [1] S. Daubney, The Universal Force of Time — Master Compendium v5, The Daubney Foundation (2026).
 - [2] R. Schleif, Genetics and Molecular Biology, 2nd ed., Johns Hopkins University Press, Baltimore (1993).
 - [3] S. Daubney, The Force of Time — Where It Departs From Current Science, The Daubney Foundation (2026).
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