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► **To cite this version:**

| Jean-Yves Boulay. Genetic code, quantum physics and the $3/2$ ratio. 2020. hal-02902700v1

HAL Id: hal-02902700

<https://hal.science/hal-02902700v1>

Preprint submitted on 20 Jul 2020 (v1), last revised 25 Oct 2020 (v4)

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Genetic code, quantum physics and the 3/2 ratio

Quantum analysis of the atoms constituting the genetic code

Jean-Yves Boulay

Abstract. The analysis of the quantum structure of the five atomic elements composing the coded twenty amino acids and the four coding nucleotides of DNA working in the organization of the genetic code reveals an opposition of their respective constituents in always an arithmetic ratio of value $3/2$ according to the parity of the number of their quantum shells. Also, the quantum analysis of the amino acid Glycine, the smallest component of peptides that can be confused with saturated base, reveals the same arithmetic oppositions of $3/2$ value of its components by the differentiation, operated according to their number of protons, of its five chemical groups.

1. Introduction

The genetic code is organized into two main entities including a coding structure, DNA (and/or RNA), made up of nucleotides and a coded structure, peptides, chains of amino acids. These two structures each consist of only five different atoms. Thus, Hydrogen, Nitrogen, Carbon, Oxygen and Phosphorus are the only elements of DNA (and RNA) the coding structure of the genetic code. All of the twenty amino acids that make up the peptides, the coded structure, are made up of Hydrogen, Nitrogen, Carbon, Oxygen and Sulphur. These two biological structures therefore each use three atoms with an even number of electron shells (C, N and O) versus two atoms with an odd number of quantum shells (H and P in DNA or H and S in amino acids). These two groups of atomic elements are opposed in various $3/2$ value ratios according to almost all of their own quantum criteria.

2. Differentiation of atoms according to the parity of the number of electron shells.

Only five atoms make up the twenty genetically encoded amino acids. These five different atoms distribute their electrons over one, two and three quantum shells. According to these physico-chemical criteria, chart Figure 1, these five atoms are opposed in two groups in a duality of three versus two atoms: Carbon, Nitrogen and Oxygen are with even number of quantum shells; Hydrogen and Sulphur have an odd number of quantum shells. Still in a $3/2$ ratio duality, the three atoms with an even number of electron shells total six layers ($2 + 2 + 2 = 6$ shells) versus four ($1 + 3 = 4$ shells) for the two atoms with odd number of quantum shells.

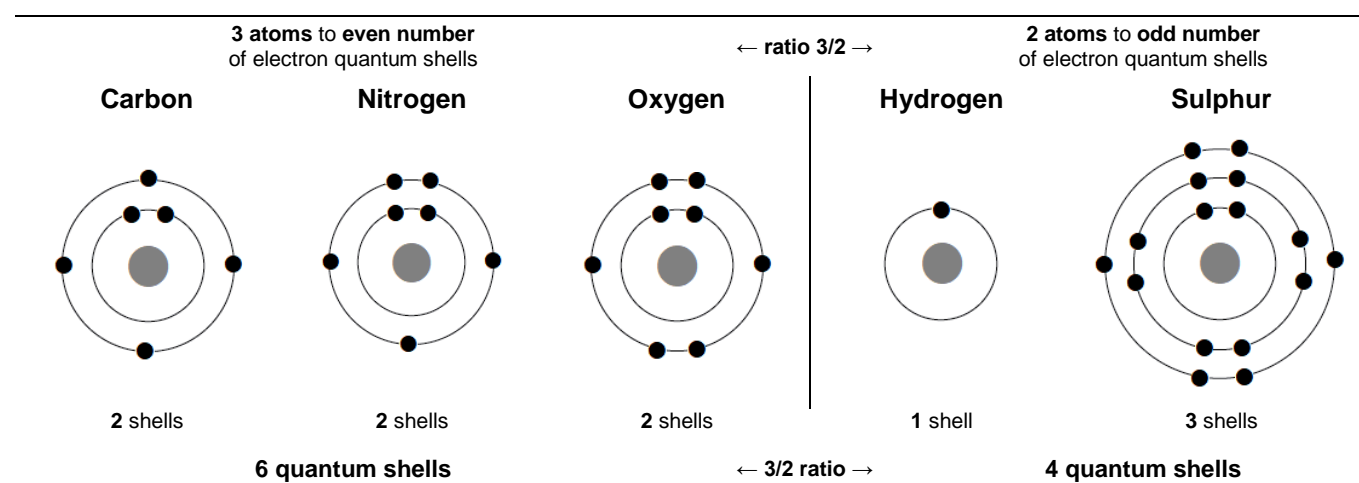


Fig. 1 Differentiation of the 5 atoms constituting 20 amino acids into 2 groups of 3 and 2 atoms according to the parity of their number of electron quantum shells.

3. Quantum structure

By studying the quantum structure of these five atoms, a multitude of $3/2$ ratios is revealed, opposing the three atoms with an even number of electronic shells to the two atoms with an odd number of electronic shells. DNA is also made up of the same five different qualities of atoms except that Phosphorus* replaces Sulphur. However, these last two atoms have the same number of electron shells and the same electronic structure in their saturated state (inside molecules) with the same maximum number of electrons that can orbit their nucleus. This means that the same $3/2$ ratio dualities also operate in DNA.

* Phosphorus and Sulphur having the same saturated quantum configuration, these two elements can be confused in some demonstrations.

The chart in Figure 2 describes the quantum shells and subshells of electrons of the five atoms constituting the twenty amino acids as well as those of the Phosphorus for DNA. Also detailed are the values of the three quantum numbers n , l and m^{**} as well as the numbers of orbitals. The description of the atoms is that in their saturated state, that is to say with their full electron shells such as they are inside amino acids or nucleotides (DNA).

** Here it is the quantum number m_l which is the subject of study. For graphic simplification this value is simply noted m in the demonstrations.

Shells value of n	1 (K) $n1$		2 (L) $n2$			3 (M) $n3$			
Subshells value of l	1s $l0$	2s $l0$	2p $l1$			3s $l0$	3p $l1$		
H	[••] $m0$								
C	[••] $m0$	[••] $m0$	[••] $m-1$	[••] $m0$	[••] $m1$				
N	[••] $m0$	[••] $m0$	[••] $m-1$	[••] $m0$	[••] $m1$				
O	[••] $m0$	[••] $m0$	[••] $m-1$	[••] $m0$	[••] $m1$				
P	[••] $m0$	[••] $m0$	[••] $m-1$	[••] $m0$	[••] $m1$	[••] $m0$	[••] $m-1$	[••] $m0$	[••] $m1$
S	[••] $m0$	[••] $m0$	[••] $m-1$	[••] $m0$	[••] $m1$	[••] $m0$	[••] $m-1$	[••] $m0$	[••] $m1$

Fig. 2 Maximum number of electron shells, subshells and orbitals of the atoms constituting the 20 amino acids and the four DNA bases. **Own electrons** (•) and **external electrons** (◐). Value of quantum numbers n , l and m . ($m \rightarrow m_l$)

The opposition of the values of Carbon, Nitrogen and Oxygen to those of Hydrogen and Sulphur (Phosphorus for nucleotides in DNA), always generates an arithmetic ratio of value 3/2 according to multiple criteria studied.

The table in Figure 3 lists the impressive series of quantum situations in which this remarkable duality takes place between sets of $3x$ entities versus $2x$ entities. Thus, the ratio for the numbers of electron subshells (1s, 2s, 2p, 3s, 3p) is 3/2. It is still 3/2 if we detail the subshells of those where the quantum number $l = 0$ of those where the quantum number $l = 1$.

Also, the ratio for the numbers of orbitals is 3/2. It is still on 3/2 if we detail the orbitals of those where the quantum number $m = 0$, of those where the quantum number $m = -1$ and those where the quantum number $m = 1$. This ratio is always 3/2 if we detail the orbitals of those where the quantum number $l = 0$ of those where the quantum number $l = 1$. Also, the maximum number of electrons that can orbit inside all of the electronic shells of these two groups of atoms is still in a ratio of 3/2: thirty electrons can orbit inside the electronic shells of Carbon, Nitrogen and Oxygen versus twenty on the electron shells of Hydrogen and Sulphur (Phosphorus for DNA bases).

For this last criterion, the distinction of the electrons which can orbit either on the first internal shell (2 electrons for each of the five atoms) or on the set of the other (external) shells always opposes the different values in ratios 3/2: 6 versus 4 electrons for the inner shell and 24 versus 16 for the other shells.

Quantum criteria:	Atoms to even number of electron quantum shells			Atoms to odd number of electron quantum shells	
	Carbon 1	Nitrogen 1	Oxygen 1	Hydrogen 1	Sulphur* 1
Number of atoms	3 atoms			2 atoms	
	$\leftarrow 3/2 \text{ ratio} \rightarrow$				
Number of electron shells (K, L, M)	Carbon 2	Nitrogen 2	Oxygen 2	Hydrogen 1	Sulphur* 3
	6 electron shells			4 electron shells	
	$\leftarrow 3/2 \text{ ratio} \rightarrow$				
Number of subshells (1s, 2s, 2p, 3s, 3p)	Carbon 3	Nitrogen 3	Oxygen 3	Hydrogen 1	Sulphur* 5
	9 subshells			6 subshells	
	$\leftarrow 3/2 \text{ ratio} \rightarrow$				
Number of subshells where the quantum number $l = 0$	Carbon 2	Nitrogen 2	Oxygen 2	Hydrogen 1	Sulphur* 3
where the quantum number $l = 1$	1	1	1	0	2
	6 subshells where $l = 0$			4 subshells where $l = 0$	
	3 subshells where $l = 1$			2 subshells where $l = 1$	
	$\leftarrow 3/2 \text{ ratio} \rightarrow$				
Maximum number of orbitals	Carbon 5	Nitrogen 5	Oxygen 5	Hydrogène 1	Soufre* 9
	15 orbitals			10 orbitals	
	$\leftarrow 3/2 \text{ ratio} \rightarrow$				
Number of orbitals where the quantum number $m = 0$	Carbon 3	Nitrogen 3	Oxygen 3	Hydrogen 1	Sulphur* 5
where the quantum number $m = -1$	1	1	1	0	2
where the quantum number $m = 1$	1	1	1	0	2
	9 orbitals where $m = 0$			6 orbitals where $m = 0$	
	3 orbitals where $m = -1$			2 orbitals where $m = -1$	
	3 orbitals where $m = +1$			2 orbitals where $m = +1$	
	$\leftarrow 3/2 \text{ ratio} \rightarrow$				
	$\leftarrow 3/2 \text{ ratio} \rightarrow$				
	$\leftarrow 3/2 \text{ ratio} \rightarrow$				
number of orbitals where the quantum number $l = 0$	Carbon 2	Nitrogen 2	Oxygen 2	Hydrogen 1	Sulphur* 3
where the quantum number $l = 1$	3	3	3	0	6
	6 orbitals where $l = 0$			4 orbitals where $l = 0$	
	9 orbitals where $l = 1$			6 orbitals where $l = 1$	
	$\leftarrow 3/2 \text{ ratio} \rightarrow$				
	$\leftarrow 3/2 \text{ ratio} \rightarrow$				
Maximum number of electrons orbiting on quantum shells	Carbon 10	Nitrogen 10	Oxygen 10	Hydrogen 2	Sulphur* 18
of which the first shell (internal)	2	2	2	2	2
of which the outer shell (s)	8	8	8	-	8+8
	30 electrons			20 electrons	
	6 electrons			4 electrons	
	24 electrons			16 electrons	
	$\leftarrow 3/2 \text{ ratio} \rightarrow$				
	$\leftarrow 3/2 \text{ ratio} \rightarrow$				
	$\leftarrow 3/2 \text{ ratio} \rightarrow$				

Fig. 3 3/2 ratio of the electron shells and subshells, orbitals and maximum numbers of electrons according to the parity of the number of electron shells of the five atoms constituting the twenty amino acids (* Or Phosphorus for DNA). Other 3/2 ratios generated in relation to the values of the different quantum numbers of the electrons. See Fig. 1 and 2.

Thus, fourteen different quantum criteria oppose, in a duality of ratio 3/2, the five atoms constituting the twenty amino acids (and also constituting the four DNA bases with the Phosphorus in place of Sulphur). The fact that the genetic code is organized only with these five different atoms in this duality is therefore not random. The perfect complementarity of the quantum characteristics of Hydrogen and Sulphur (Phosphorus in DNA) is particularly remarkable. These last two atoms have indeed very different quantum characteristics (in contrast to Carbon, Nitrogen and Oxygen with common characteristics) which however complement each other perfectly to always oppose in a 3/2 ratio to three other atoms, constituents of amino acids (and DNA bases). For example, Sulphur has a maximum number of nine orbitals versus only one for Hydrogen. These two very different values nevertheless complement each other (10 orbitals) to oppose in a duality of ratio 3/2 to the three times five quantum orbitals of Carbon, Nitrogen and Oxygen (15 orbitals).

Thus, the 3/2 ratio is revealed at the bottomest of the subatomic structure of the constituents of the twenty amino acids that are on the one hand the three atoms of Carbon, Nitrogen and Oxygen and on the other hand the two atoms of Hydrogen and Sulphur. It is therefore remarkable to note that these same phenomena are found in DNA, another mechanical component of the genetic code, where the quantum properties of the Phosphorus mimic those of Sulphur.

Also, Figure 4, these six atoms constituting the entire mechanism of the genetic code therefore oppose three to three depending on the parity of their number of electron shells. In a ratio of 3/2, the Hydrogen - Phosphorus - Sulphur group totals 63 (3 times 21) nucleons versus 42 (2 times 21) for the Carbon - Nitrogen - Oxygen group. These same two groups are inversely opposed in the 3/2 ratio with respectively nine valences for C, N and O versus six valences for H, P and S.

Physical entities:	Atoms to even number of electron quantum shells			Atoms to odd number of electron quantum shells		
nucleons	Hydrogen 1	Phosphorus 30	Sulphur 32	Carbon 12	Nitrogen 14	Oxygen 16
	63 nucléons			42 nucléons		
	← 3/2 ratio →					
valences	Hydrogen 1	Phosphorus 3	Sulphur 2	Carbon 4	Nitrogen 3	Oxygen 2
	6 valences			9 valences		
	← 2/3 ratio →					

Fig. 4 The six atoms constituting the genetic code: 3/2 ratio opposing the 3 atoms with an even number of electron shells of the 3 with an odd number according to their global number of nucleons and a 3/2 ratio according to their number of valences.

4 Quantum analysis

4.1 New quantum chart

This quantum study of the genetic code is an opportunity to propose a new type of table describing the quantum organization of atoms. In this chart, illustrated in Figure 5, the different quantum shells and subshells are presented in the form of chevrons. At the top end of each rafter are indicated the names of the different shells and subshells; at the left end of these chevrons, the numbers of orbitals and electrons of these different shells and quantum subshells are indicated. At each chevron vertex is the orbital where the quantum number $m = 0$. The orbitals with positive quantum number m are progressively positioned towards the top of these chevron vertices and the orbitals with negative quantum number m are progressively positioned towards the outside left of these chevron vertices.

In the appendix, the same type of table is presented describing the quantum organization of the shells and subshells up to the 5th shell (O) and 15th subshell ($5g$). This innovative presentation, more explicit in describing the quantum structure of the atomic elements, will be used in various tables of this quantum study of the constituents of the genetic code.

amount of orbitals		amount of electrons •		shells and subshells					
by shell:	by subshell:			1(K)		2(L)		3(M)	
				1s $l=0$	2s $l=0$	2p $l=1$	3s $l=0$	3p $l=1$	3d $l=2$
1	2	1	2	□ $m=0$					
4	8	1	2		□ $m=0$	□ $m=+1$			
		3	6		□ $m=-1$	□ $m=0$			
9	18	1	2				□ $m=0$	□ $m=+1$	□ $m=+2$
		3	6				□ $m=-1$	□ $m=0$	□ $m=+1$
		5	10				□ $m=-2$	□ $m=-1$	□ $m=0$

Fig. 5 Quantum distribution of orbitals and electrons in the first three shells and the first six subshells. Chart in chevron form, see appendix Fig. 26.

4.2 Quantum structure of atoms

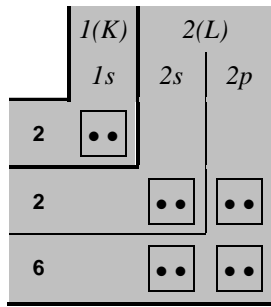
Figure 6 illustrates the quantum structure of the five atoms working in the genetic code. As stated above, the Phosphorus, working in DNA and the Sulphur, involved in peptides, are confused in this analysis. Also, the three atoms Carbon, Nitrogen and Oxygen with even number of quantum shells present the same quantum configuration in their saturated state.

As already introduced in Figure 3, it appears more explicitly in this type of chevron form chart that, in a 3/2 value ratio, the 30 electrons (10 + 10 + 10) orbiting in the three atoms with an even number of quantum shells oppose the 20 electrons (2 + 18) orbiting in the two atomic elements with an odd number of quantum shells.

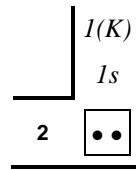
Atoms to **even number** of quantum shells

Atoms to **odd number** of quantum shells

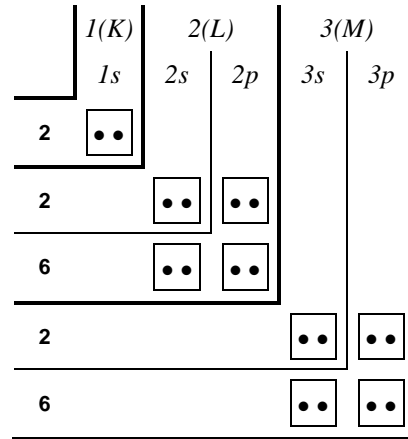
Carbon, Nitrogen, Oxygen



Hydrogen



Sulphur (or Phosphorus for DNA)



Maximum number of electrons orbiting in the quantum shells:

Carbon 10	Nitrogen 10	Oxygen 10
---------------------	-----------------------	---------------------

Hydrogen 2

Sulphur* 18

30 orbiting electrons

← 3/2 ratio →

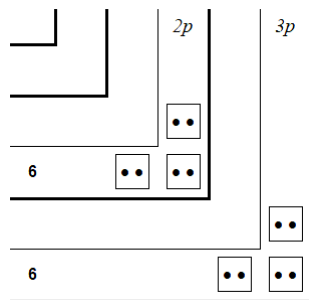
20 orbiting electrons

Fig. 6 Quantum structure of the 5 elements constituting the genetic code. *Or Phosphorus for DNA. From the chevron form chart introduced in Figure 5.

4.3 Azimuthal quantum number

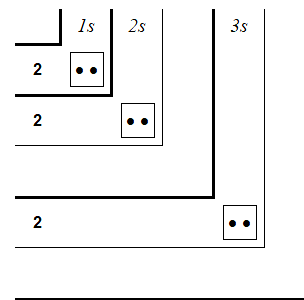
Figure 7 details the distribution of electrons according to the value of the azimuthal quantum number. It appears that according to this criterion and the parity of the number of quantum shells, the distribution of the electrons of these three and two elements is organized into numerous ratios of 3/2 value including ratios transcendent according to the criteria considered. This arithmetic transcendence is directly related to the remarkable identity $(a + b)^2 = a^2 + 2ab + b^2$ where a and b have the respective values 3 and 2. This relationship to the remarkable identity which operates in several of the next tables is illustrated and explained in Chapter 4.7.

Azimuthal quantum number $l = 1$
external subshells **2p** and **3p**



30 electrons orbiting on external subshells **2p** and **3p**

Azimuthal quantum number $l = 0$
internal subshells **1s** **2s** and **3s**



20 electrons orbiting on internal subshells **1s** **2s** and **3s**

← 3/2 ratio →

C	N	O
6	6	6
-	-	-

← 2p →
← 3p →

H	S*
-	6
-	6

C	N	O
2	2	2
2	2	2
-	-	-

← 1s →
← 2s →
← 3s →

H	S*
2	2
-	2
-	2

18 orbiting electrons

← 3/2 ratio →

12 orbiting electrons

← 3/2 ratio →

12 orbiting electrons

← 3/2 ratio →

8 orbiting electrons

← 3/2 ratio →

Fig. 7 Count of the electrons orbiting on the external (azimuthal quantum number $l = 1$) and internal subshells (azimuthal quantum number $l = 0$) in the 5 elements H, C, N, O and S (* or P in DNA) constituting amino acids. See Fig. 5 and 6.

4.5 Magnetic quantum number

Illustrated in Figure 8, the distinction between electrons with a magnetic number $m = 0$ and those with a magnetic number $m = -1$ or $m = +1$ generates exactly the same transcendent arithmetic ratios of $3/2$ value. It is essential to emphasize that despite different individual values for the elements H and S, the same global values are found in these counts from two different quantum criteria: the azimuthal quantum number (l) and the magnetic number (m).

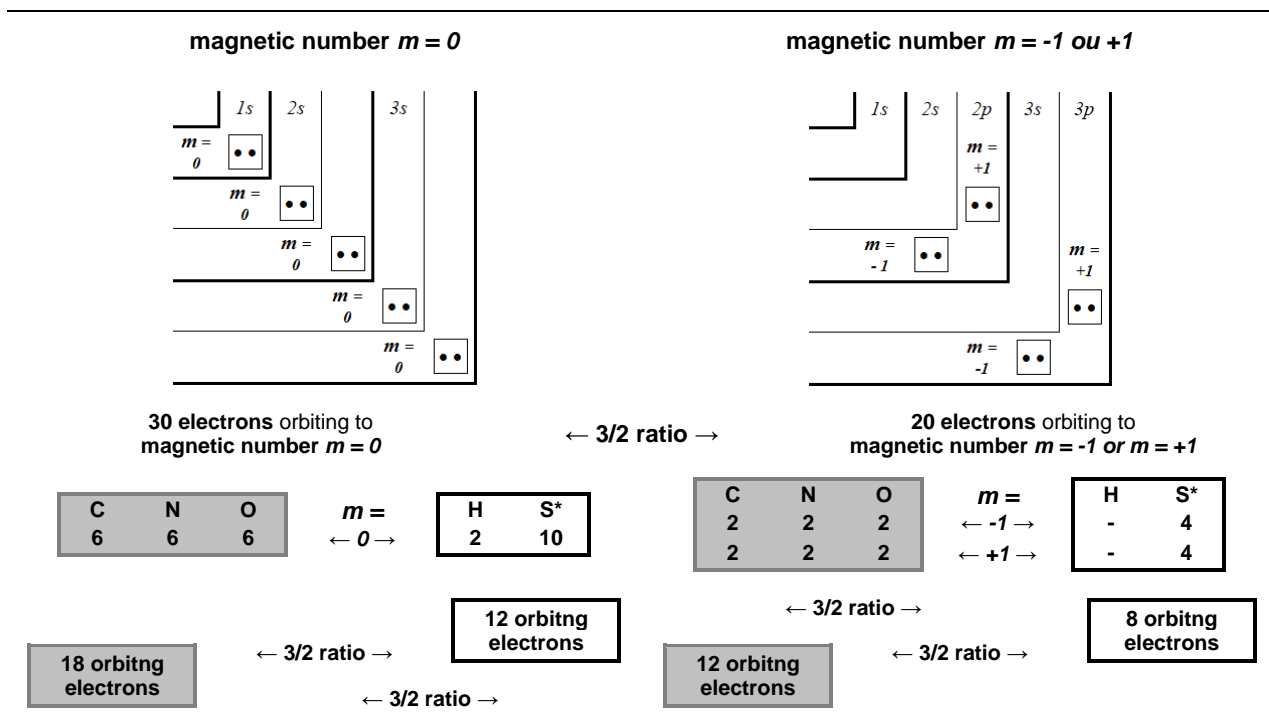


Fig. 8 Count of electrons orbiting with a magnetic number $m = 0$ and electrons orbiting with a magnetic number $m = -1$ or $m = +1$ in the 5 elements H, C, N, O and S (* or P in DNA) constituting the amino acids. See Fig. 5, 6 and 10 (Recall: $m \rightarrow m_l$).

4.6 Quantum shells

As illustrated in Figure. 9, the individual number of subshells of the three atomic elements C, N and O (with even number of shells) is equal to $3/2$ of their number of quantum shells. The values of these same ratios are to $1/1$ for Hydrogen and $5/3$ to Sulphur. However, the global values of these two elements with an odd number of shells complement each other perfectly to also generate a $3/2$ ratio between their number of subshells and shells but also with the global values of the three atoms with an even number of quantum shells.

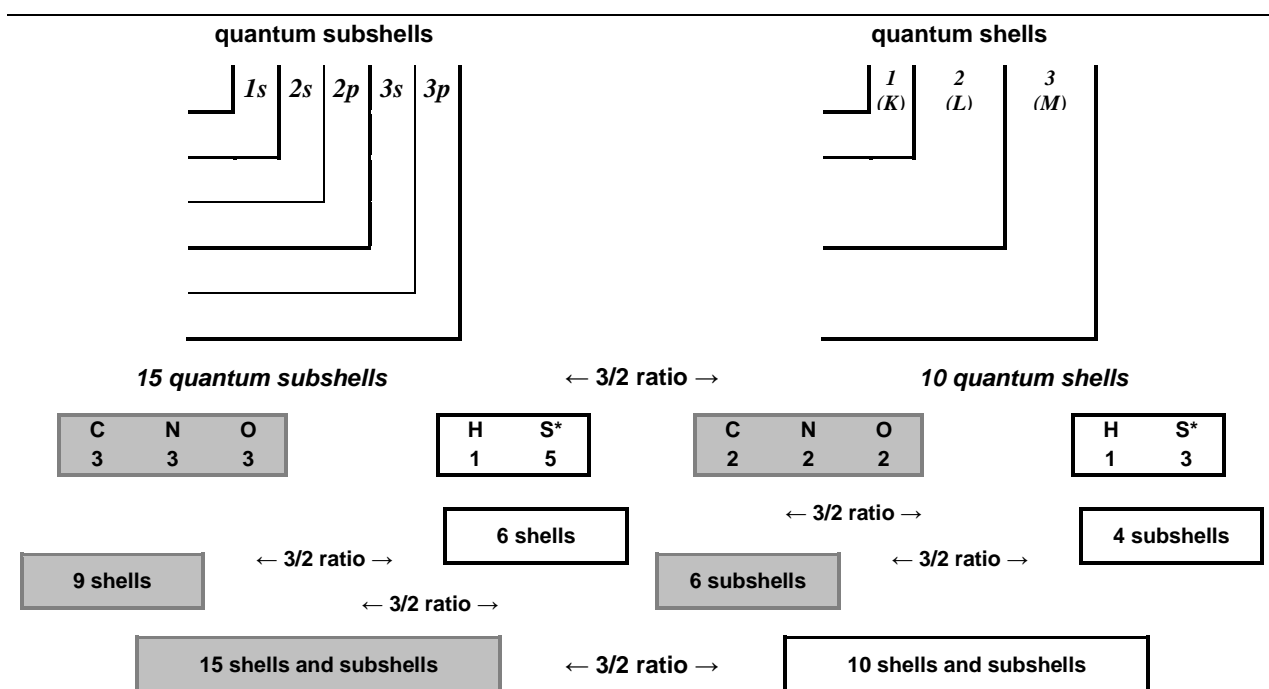


Fig. 9 Count of subshells and quantum shells of the 5 elements H, C, N, O and S (* or P in DNA) constituting the amino acids. See Fig. 5, 6 and 10.

4.7 Remarkable identity

Thus, these various ratios opposing the subshells and shells and transversely, the two categories of atoms previously defined according to the parity of their number of quantum shells, are organized in the remarkable identity $(a + b)^2 = a^2 + 2ab + b^2$ where a and b have the respective values 3 and 2. Figure 10 explains this arithmetic organization operating in the quantum structure of the five elements working within the genetic code.

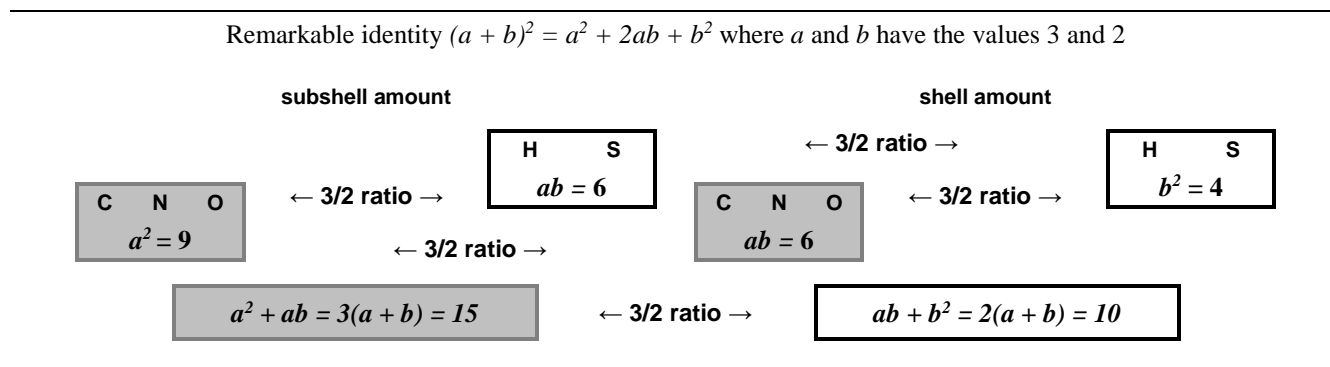


Fig. 10 Remarkable identity revealed in the count of subshells and quantum shells of the five elements H, C, N, O and S (P in DNA). See Fig. 9.

Thus, the quantity of subshells in C, N and O corresponds to the value a^2 of the remarkable identity and the quantity of subshells in H and S corresponds to the value ab . The quantity of quantum shells in C, N and O also corresponds to the value ab and that in H and S corresponds to the value b^2 . These different values therefore transcend into these equal ratios:

$$(a^2/ab) = (ab/b^2) = (a^2+ab)/(ab+b^2)$$

$$(3^2/6) = (6/2^2) = (3^2+6)/(6+2^2)$$

$$(9/6) = (6/4) = (15)/(10)$$

As it was previously revealed, this remarkable identity therefore also operates in the counts of electrons according to their azimuthal quantum number (Figure 8) and according to their magnetic number (Figure 9). In these electron counts, the values are just double and, for a and b at the root values 3 and 2, the respective and transcendent values are equal to:

$$2a^2 \rightarrow 2ab \rightarrow 2ab \rightarrow 2b^2$$

$$18 \rightarrow 12 \rightarrow 12 \rightarrow 8$$

5 Anatomy of Glycine

Within the mechanism of the genetic code and therefore among the twenty amino acids, Glycine is distinguished by its absence of radical. Its radical is reduced to a simple hydrogen atom which in a way simply closes the "base" structure common to each amino acid. The quantum study of this *glycined base*, identifying with Glycine, reveals singular arithmetic arrangements of its different components.

5.1 Modules of Petoukhov

The notion of modules is an original system proposed by Sergei Petoukhov [1 and 2] to describe the structure of biological molecules. In the appendix is introduced this concept of modular structure and detailed two sets of amino acids with the number of protons equal or not equal to eight times their respective number of Petoukhov modules.

5.2 Detailed structure of Glycine

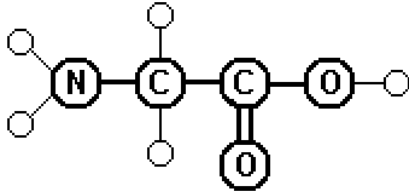
Figure 11 describes the structure of Glycine (or saturated base called *glycined base*) according to many criteria including its chemical composition, modular, but also atomic. It turns out that Glycine consists of 40 protons, either $5x$ protons or $(3 + 2)x$ protons. This glycined base also consists of 5 groups or modules, i.e. $(3 + 2)x$ chemical groups. In Glycine, the number of protons is therefore an exact multiple of 8 (5 times 8 protons) and it turns out that the average number of protons per chemical group (or Petoukhov module) is therefore 8. For two groups (CH_2 and O), the amount of protons is exactly 8 whereas for the other three groups, these proton amounts are 9 or 6 ($\text{NH}_2 \rightarrow 9$, $\text{OH} \rightarrow 9$ and $\text{C} \rightarrow 6$).

The differentiation of these two types of modules, made up or not made up of 8 protons reveals a multitude of oppositions of the different natures of the components of Glycine (glycined base) in always an arithmetical ratio of 3/2 value. As described in the appendix, the multiplicity of protons/modules within an 8/1 ratio of amino acids is not random, but concerns exactly 50% of the twenty amino acids used in the genetic code, i.e. 10 amino acids out of 20.

Chemical structure of a saturated base (glycined) identifying with Glycine

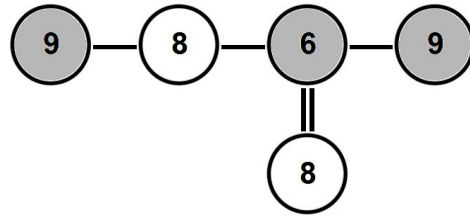
molecular structure: 10 atoms

5 atoms without neutron and to odd number of quantum shell: H → 1 quantum shell



5 atoms with neutrons and to even number of quantum shells: C N O → 2 quantum shells

modular structure: 5 modules

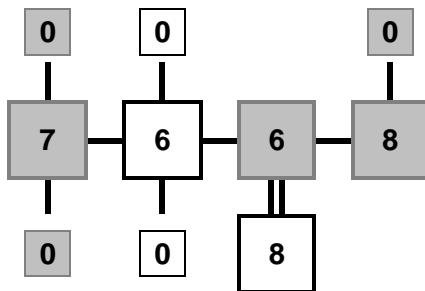


3 modules not composed of 8 protons

← 3/2 ratio →

2 modules composed of 8 protons

atomic structure: 35 neutrons

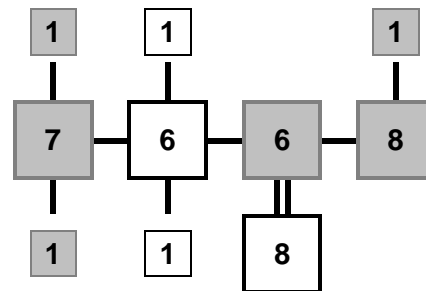


21 neutrons

← 3/2 ratio →

14 neutrons

atomic structure: 40 protons



3 + 21 = 24 protons

← 3/2 ratio →

2 + 14 = 16 protons

Fig. 11 Chemical, modular and atomic structure of a saturated base identified with the amino acid Glycine: 5 modules, 10 atoms, 40 protons and 35 neutrons. See also Fig. 12.

Glycine is made up of a multitude of entities whose numbers are all multiples of five. Thus the glycined base consists of five modules, two times five atoms, five of which have one electron shell (H) and five at two shells (C, N and O). Also Glycine consists of 5 times 15 nucleons (75) including 5 times 7 (35) neutrons and 5 times 8 (40) protons. The covalent bonds between these different components are also in numbers which are multiple of 5.

Physico-chemical entities:	proton number modules not equal to 8		proton number modules equal to 8	
	9	6	8	8
modules amount (chemical groups): 5	3	← 3/2 ratio →	2	
Full atoms amount: 10	6	← 3/2 ratio →	4	
Atoms amount to 1 electron shell: 5 (satellite atoms: H)	3	← 3/2 ratio →	2	
Atoms amount to 3 electron shells: 5 (master atoms: C, N and O)	3	← 3/2 ratio →	2	
Nucleons amount: 75	45	← 3/2 ratio →	30	
Protons amount: 40	24	← 3/2 ratio →	16	
whose master atoms (in C, N and O): 35	21	← 3/2 ratio →	14	
whose satellite atoms: (in H): 5	3	← 3/2 ratio →	2	
Neutrons amount: 35	21	← 3/2 ratio →	14	
Number of covalent bonds*: 20	12	← 3/2 ratio →	8	
whose satellites/masters bonds: 10	6	← 3/2 ratio →	4	
whose masters/masters bonds: 10	6	← 3/2 ratio →	4	

Fig. 12 Count of the different chemical, modular and atomic entities of a saturated base identifying with the amino acid Glycine. *Cumulative bonds (valences) by atom. See Fig.11 also.

Also, it therefore appears, [Figures 11 and 12](#), that the different constituents of Glycine, always $5x$ in number, are always at 3 same x entities in the set of three modules (chemical groups) with number of protons not equal to 8 and always of amount at 2 same x entities in the set of two modules whose number of protons is equal to 8.

5.3 Quantum analysis of the glycined base

In the next [Figures 13 and 17](#), the detailed quantum structure of Glycine (shown as glycined base) is illustrated. This graphic representation, more explicit than a classic version, is inspired by *the concept of chart in chevron* introduced in [Chapter 4](#).

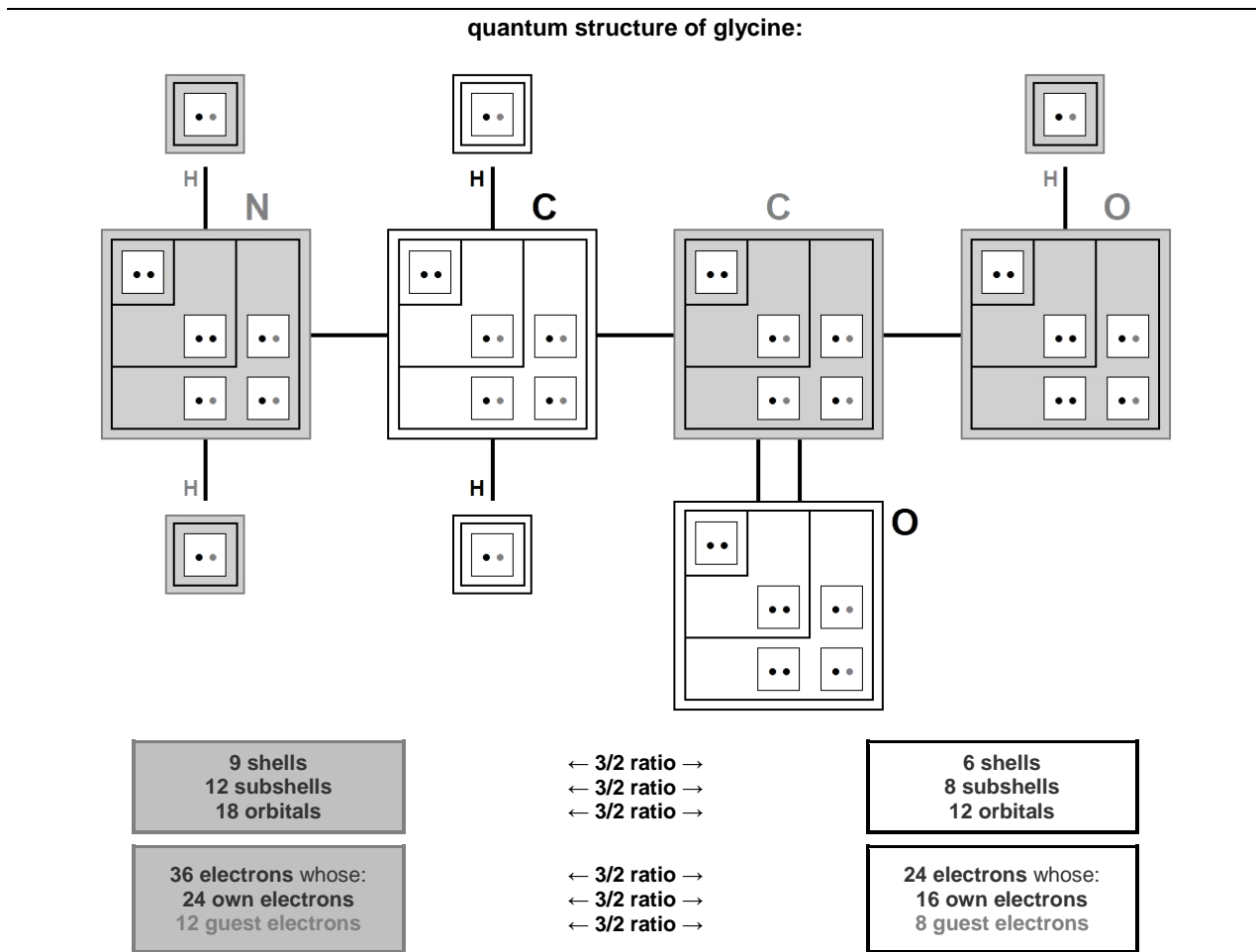


Fig. 13 Quantum structure of Glycine: 30 orbitals, 20 subshells, 15 electron shells, 60 electrons whose 40 own electrons (•) and 20 guest electrons (*). See [Fig. 5](#) and [9](#).

In its quantum structure, Glycine (glycined base) is therefore also always made up of $5x$ entities. Its ten atoms total 15 ($5x \rightarrow x = 3$) quantum shells and 20 ($5x \rightarrow x = 4$) electron subshells. These 20 subshells total 30 ($5x \rightarrow x = 6$) orbitals where 60 ($5x \rightarrow x = 12$) electrons can evolve, including 40 ($5x \rightarrow x = 8$) individually own to these ten atoms and 20 ($5x \rightarrow x = 4$) covalent electrons (20 shared electrons).

5.3.1 Orbitals, shells, and quantum subshells

Depending on whether they are in the three modules with a number of protons not equal to 8 or in the two with a number of protons equal to 8, these various entities are always with the respective numbers of $3x$ and $2x$. Thus, in these two groups of modules we can oppose, in a ratio of $3/2$, 18 orbitals to 12 others, 9 quantum shells to 6 others and 12 subshells to 8 other subshells.

Also, as illustrated in [Figure 14](#), the values of the orbitals and of the quantum subshells oppose in transcendent ratios of value $3/2$. So more, this arithmetic transcendence is organized from the remarkable identity $(a + b)^2 = a^2 + 2ab + b^2$ where a and b have the respective values 3 and 2.

In these counts, the respective and transcendent values are equal to two times the root values of this identity, that is:

$$2a^2 \rightarrow 2ab \rightarrow 2ab \rightarrow 2b^2$$

$$18 \rightarrow 12 \rightarrow 12 \rightarrow 8$$

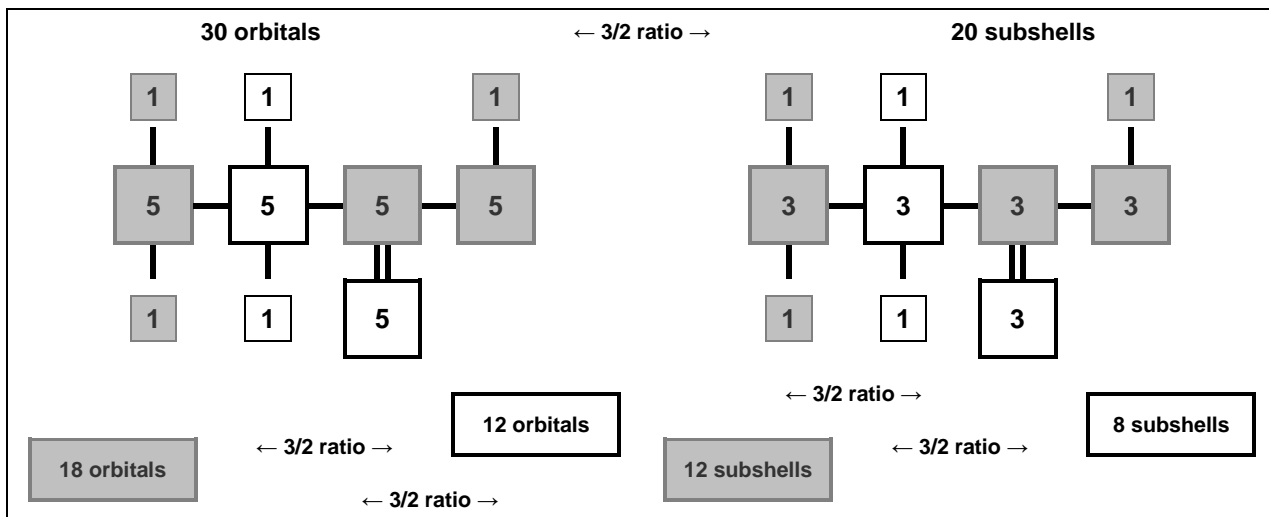


Fig. 14 Counting the orbitals and quantum subshells of Glycine. See Fig. 13.

5.3.2 Orbiting electrons and own electrons

As illustrated in Figure 15, depending on whether they are specific to each atom or orbiting (own + invited), and according to their membership of one or the other type of modules (chemical groups previously defined), the electrons of Glycine s oppose in transcendent ratios on $3/2$ value. Also, this arithmetic transcendence is organized from the remarkable $(a + b)^2 = a^2 + 2ab + b^2$ where a and b are to respective values 3 and 2.

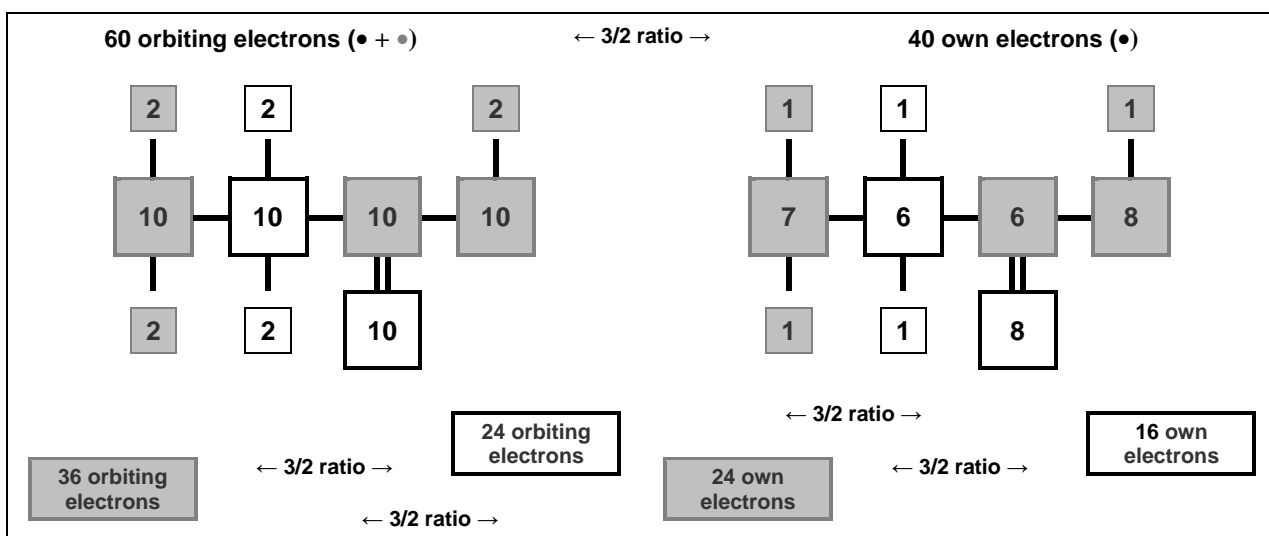


Fig. 15 Count of the orbiting electrons (• + •) and the own electrons (•) of the Glycine. See Fig. 13 also.

In these counts, the respective and transcendent values are equal to four times the root values of this identity, that is:

$$4a^2 \rightarrow 4ab \rightarrow 4ab \rightarrow 4b^2$$

$$36 \rightarrow 24 \rightarrow 24 \rightarrow 16$$

5.3.2.1 Electric charges and $3/2$ ratio

Since the numbers of clean electrons correspond to those of the numbers of protons, the negative and positive electric charges are therefore also opposed in a ratio of value $3/2$ with, for the whole of Glycine, 60 electrons (orbiting) charge negative (-e) versus 40 protons of positive charge (e). The same opposition of electric charges is observed in the two groups of modules previously defined with, for one and the other group, 36 negative charges versus 24 positive and 24 negative charges versus 16 positive charges.

5.3.3 Own electrons and electron sharing

In Glycine, the number of semi-full orbitals is double the number of full orbitals. Also, as illustrated in Figure 16, the distribution of these two types of orbital is organized in $3/2$ value ratios between the two groups of modules previously defined according to their proton number equal or not equal to 8.

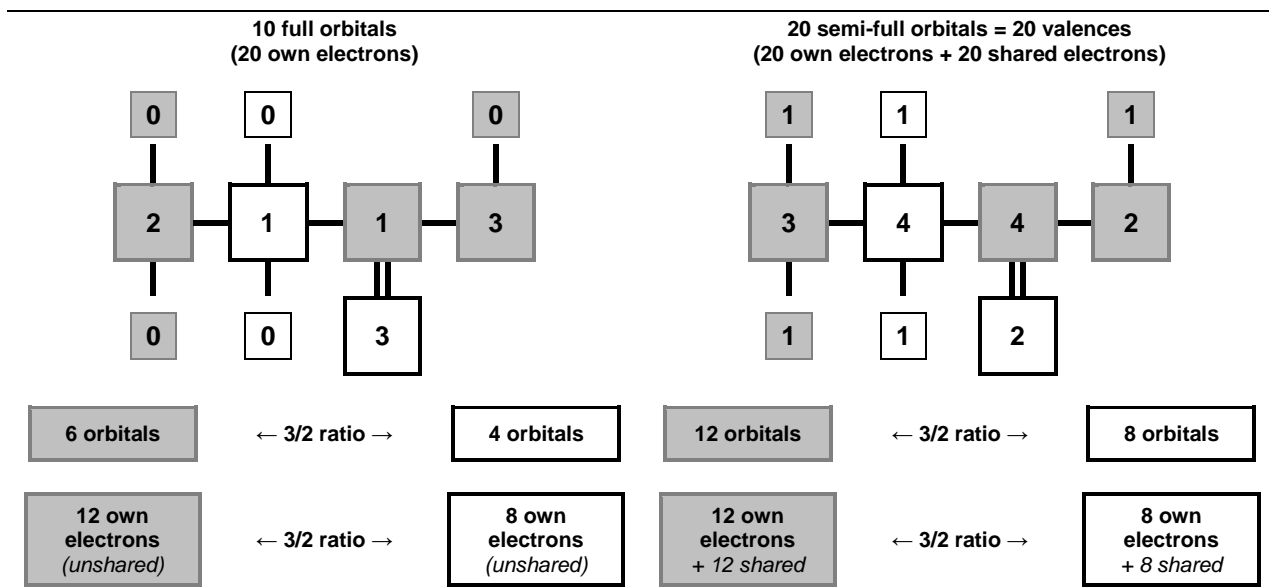


Fig. 16 Counting of the full and semi-full orbitals of the atoms constituting the Glycine. Count of own and shared electrons from these two types of orbital. See also Fig. 13, 15 and 17.

5.3.4 Jumps in quantum shells

Figure 17 describes, for each atom of Glycine, the amplitude of subshell jumps of electrons which are shared with another atom. Among the 20 shared electrons, it turns out that 10 change (jump) subshell level and 10 do not change level (see also Figure 20).

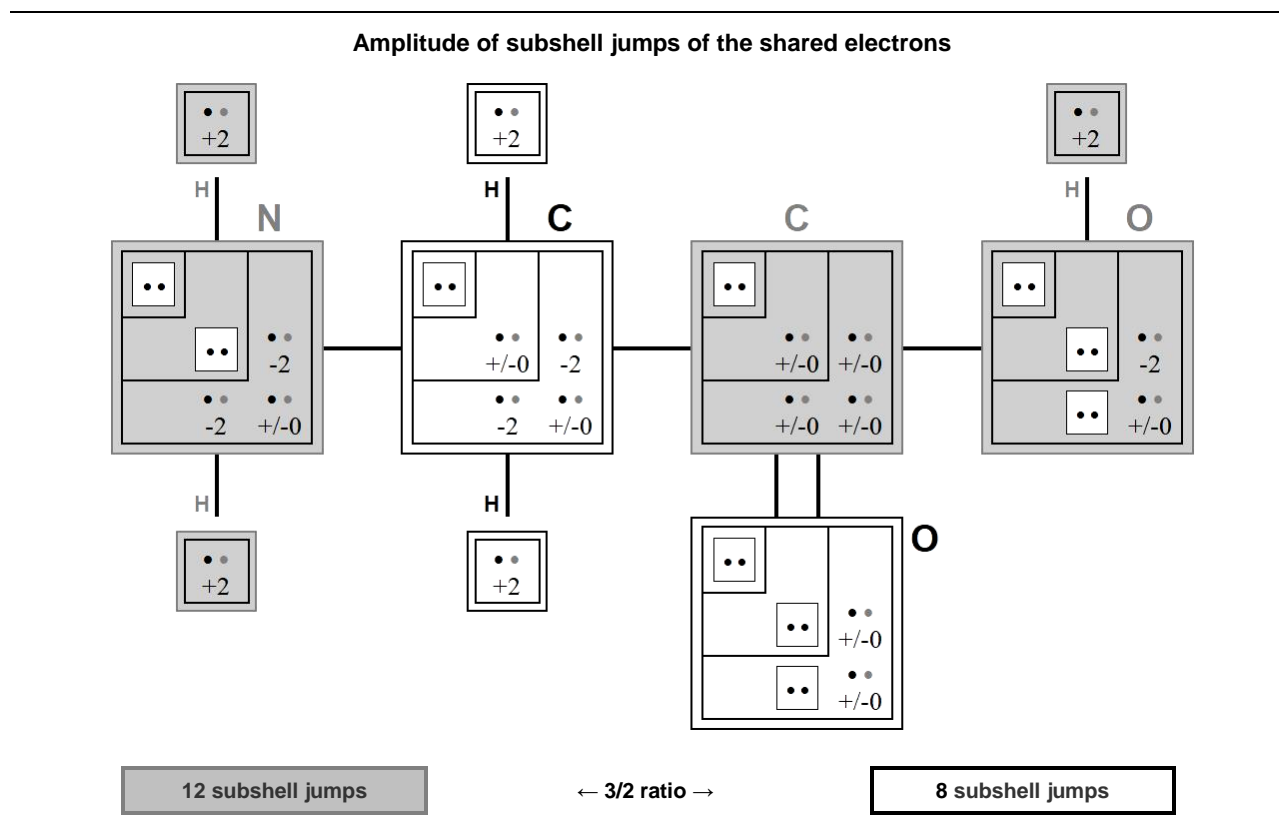


Fig. 17 Subshell jumps of the 20 shared electrons of the atoms constituting Glycine. See Fig. 16, 18 and 19 also.

For example (Figure 18), in the NH₂ group (module), the own electron of the Hydrogen atom evolves on its original level 1 subshell and on the level 3 subshell of the Nitrogen atom.

Subshell jump and evolving of an electron from of a satellite Hydrogen of the NH₂ group of Glycine

the electron of H jumps from a subshell 1 level (in H) to a subshell 3 level (in N), that is a jump of value +2

the electron of H evolves in subshell of level 1 (in H) and in subshell of level 3 (in N), that is a cumulative levels of subshell evolution of value 4

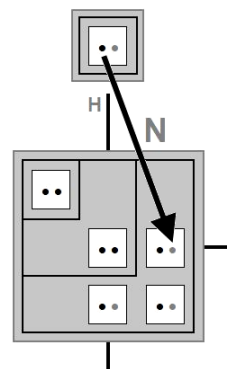


Fig. 18 Example of a subshell jump and of evolution of an electron from a satellite Hydrogen of the NH₂ group (module) of Glycine. See Fig. 16 and 17.

5.3.4.1 Jumps and levels of quantum subshells

As it appears Figures 17 and 19, the counting of these subshell jumps always registers in a ratio of 3/2 according to the membership of the electrons to one or the other of the two groups of modules (chemical groups) differentiated according to their number of protons. Also, for each of the shared electrons, the distribution of the cumulative levels of the subshells where these electrons orbit are still organized in a 3/2 value ratio according to these same criteria of differentiation of the modules considered.

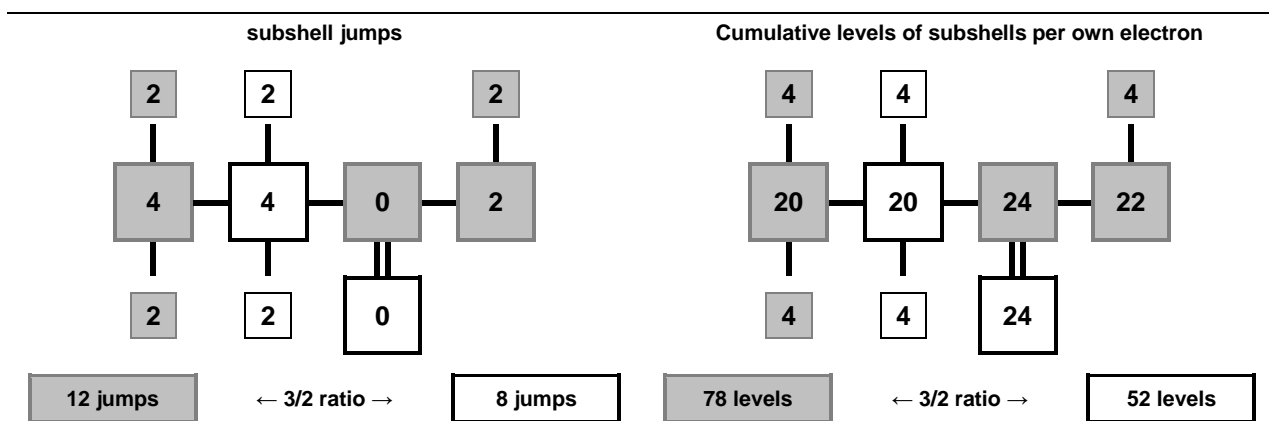


Fig. 19 Subshell jumps and cumulative levels of subshells by own electrons of the atoms constituting Glycine. See Fig. 15, 16 and 17 also.

5.3.4.2 Electrons shared and subshells

As shown in Figure 20 and previously Figure 17, in their covalent migration, 50% of the shared electrons change of subshell level and 50% are found on the same quantum level. The count of these two types of migration in one and the other group of modules previously defined according to their number of protons is written in arithmetic ratios of value 3/2 with, for each type of migration, six electrons versus four.

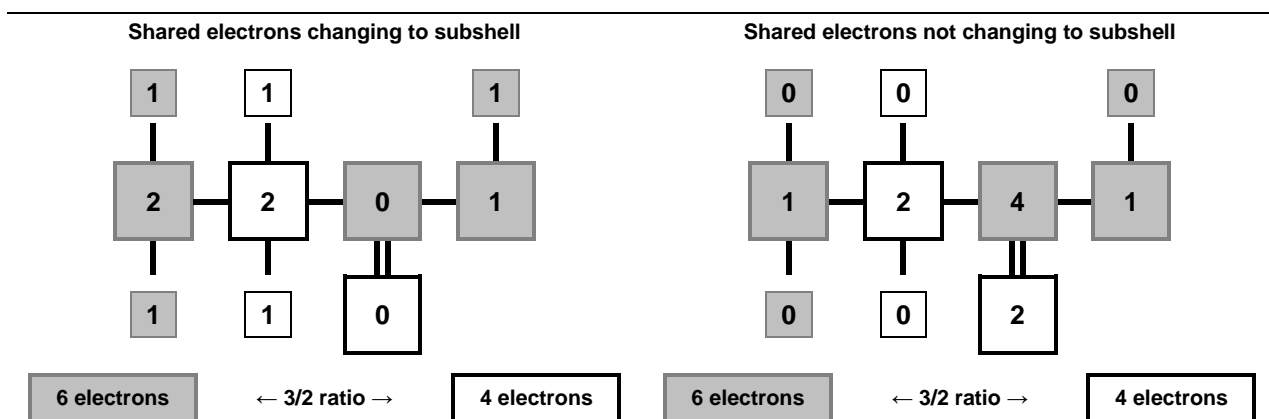


Fig. 20 Count of shared electrons changing or not changing to subshell of the atoms constituting Glycine. See also Fig. 15, 17 and 18.

5.4 Bonds (valences) and modules

According to the concept of modular structure proposed by Sergei Petoukhov, concept introduced in [Chapter 5.1](#) and detailed in the [appendix](#), it is possible to differentiate two types of covalent bonds:

- the module↔module bonds operating between two non-hydrogen atoms, bonds which can be qualified as *master-master*,
- the module↔Hydrogen bonds operating between a Hydrogen and a non-hydrogen atom, bonds which can be qualified as *master-satellite*.

As demonstrated in [Figure 21](#) and more explicitly in [Figures 13](#) and [17](#) where the modular and quantum structures of the atoms are illustrated, the cumulative numbers of *master-master* bonds and of *master-satellite* bonds are identical in Glycine (glycined base), i.e. ten bonds (cumulated per atom) for these two categories of covalent bonds.

Also, the distribution of these two types of bonds is organized into 3/2 value ratios between the two groups of modules previously differentiated according to their number of protons which can be equal or not equal to 8.

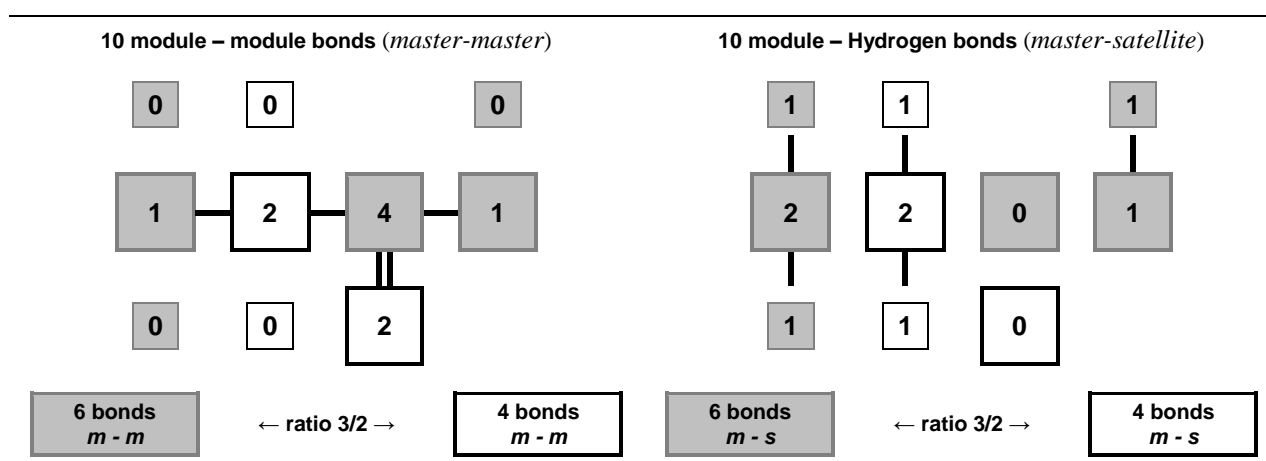


Fig. 21 Count of the module - module bonds (*master - master*) and the module - Hydrogen bonds (*master - satellite*) of the atoms constituting Glycine (cumulative bonds per atom). See [Fig. 13](#) and [17](#) also.

5.5 Radius and electronegativity

The study of the different main physical values of the atomic elements constituting Glycine* and listed in the table in [Figure 22](#) reveals some like arithmetic phenomena opposing the three modules with a number of protons not equal to 8 to the two with number of protons equal to 8.

atomic element:	H	C	N	O
radius of covalence (pm):	38	77	75	73
atomic radius (pm):	25	70	65	60
Van der Waals radius (pm):	120	170	155	152
electronegativity (Allred):	2,2	2,5	3,07	3,5
electronegativity (Pauling):	2,2	2,55	3,04	3,44
electronegativity (Mulliken):	3,01	2,67	3,08	3,22

Fig. 22 Radius and electronegativity [3] of the four atomic elements constituting Glycine. See [Fig. 23](#) and [24](#).

5.5.1 Atomic radii

As illustrated in [Figure 23](#), the cumulation of the individual values of the covalent radii, the atomic radii and the Van der Waals radii of the atomic elements constituting the Glycine also register in arithmetic ratios of 3/2 value by the opposition of the two groups of modules previously defined according to their number of protons.

These ratios are not all exactly equal to the 3/2 arithmetic value but approach more than 99%. These differences are justified by the lack of precision (rounding of values) of the source data.

*Note: the values considered are those of the elements in their primordial state.

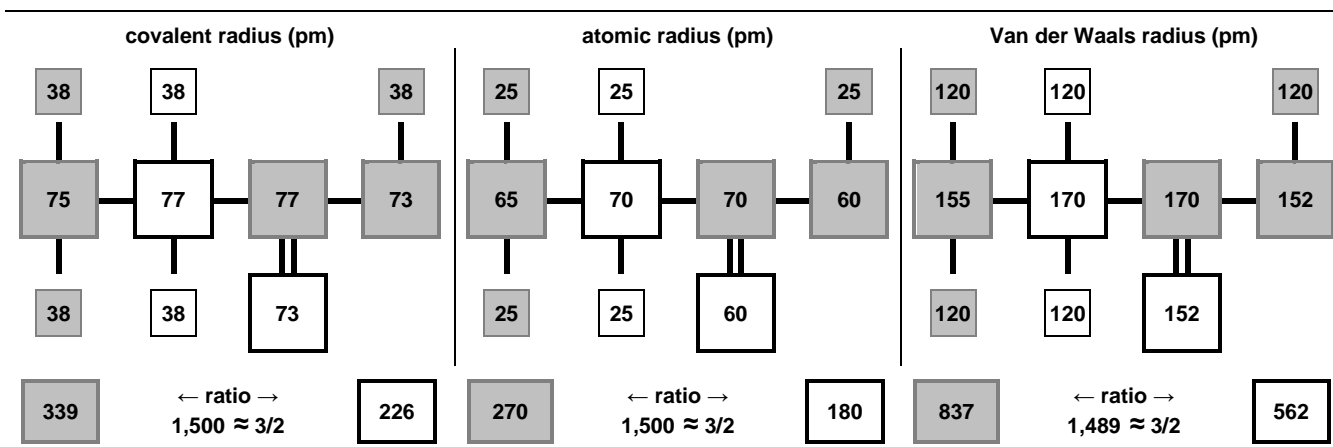


Fig. 23 Counts of the individual values of the covalent radii, atomic radii and Van der Waals radii of the atomic elements constituting Glycine.

5.5.2 Electronegativity

As illustrated in Figure 24, the cumulative individual values of the electronegativity of the atomic elements constituting Glycine also fall into arithmetic ratios of 3/2 by the opposition of the two groups of modules previously defined according to their number of protons. This is true from the Allred scale, from the Pauling scale and also from the Mulliken scale with slight oscillations of the ideal ratio of 3/2.

Again, these ratios approach at more than 99% of the ideal 3/2 value ratio and these differences are justified by the variability and relative imprecision of the raw source data on these different scales.

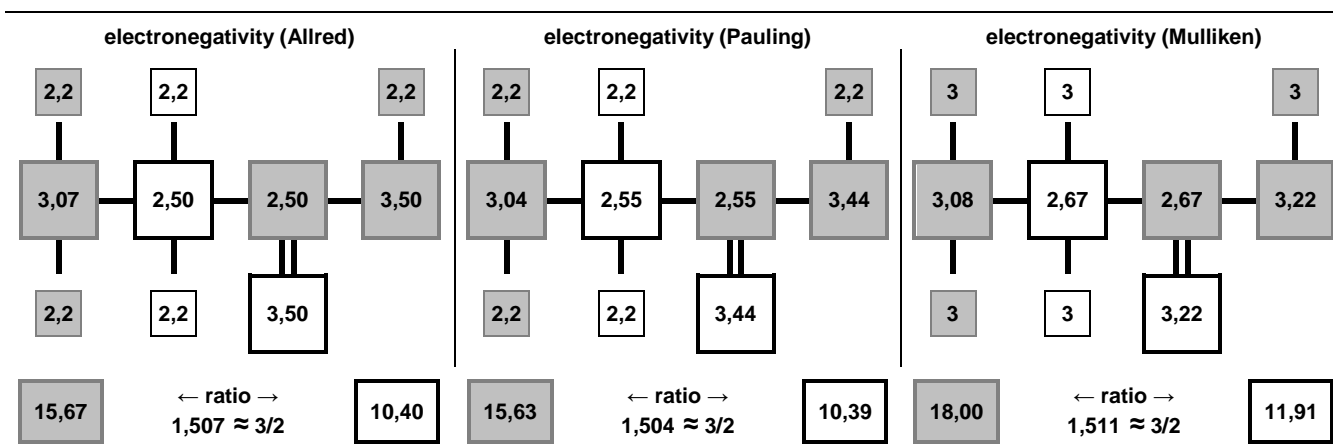


Fig. 24 Counts of individual electronegativity values of the chemical elements constituting Glycine according to the Allred, Pauling and Mulliken scales.

Thus, although different individually, these different physical values of atomic dimension and electronegativity are nevertheless and always organized in arithmetic ratios of value 3/2 by the opposition of the modules (chemical groups) with equal or unequal proton number to the value 8.

Discussion et Conclusion

Analysis of the quantum organization of the atomic elements working in the constituents of the genetic code reveals a systematic opposition of their different components in a 3/2 value ratio depending on whether these atoms have an even or odd number of quantum layers. The multitude of these singular arithmetic arrangements, always identical in their final ratios, prohibits the idea of any random interaction of these different constituents.

These arithmetic configurations are of a highly structured level, often even organized around the remarkable identity $(a + b)^2 = a^2 + 2ab + b^2$ where a and b have precisely the respective values 3 and 2. The arithmetic mechanics arising of this identity thus allows the different values considered to be organized and opposed in a triple ratio of 3/2 transcendent values and in arithmetic form: $(a^2/ab) = (ab/b^2) = (a^2+ab)/(ab+b^2)$.

As highlighted in some of his various works including those referenced in [1] and [2], Sergei Petoukhov draws attention to the organization of the components of the genetic code around the values 3 and 2. For example, and this is of great importance, there are two types of DNA base associations, one with three hydrogen bonds between the bases Guanine and Cytosine and the other with two hydrogen bonds between the bases Adenine and Thymine.

The fact that this 3/2 arithmetic ratio also operates between the components of Glycine, the primary amino acid identified to a glycined base, supports the idea of an arithmetically non-random organization of the mechanics of the genetic code. The latest investigations studying the different values of radii and electronegativity of the components of this glycined base greatly reinforce this analysis.

In this organization of living matter, its modular structure, the multiplicity (or not multiplicity) protons/modules and more broadly the multiplicity (or not multiplicity) of the numbers of protons of the twenty amino acids by the value 8 appear to be factors to consider with the utmost care. Also, we suggest, without discussing it further here, that this criterion of multiplicity by 8 of the number of protons in amino acids is related to the byte rule, another quantum constraint operating in the atomic elements used in these amino acids.

To conclude this quantum study of the components of living matter working in the organization of the genetic code, we advance the idea that this matter known as "living" is only the prolongation of a general organization of matter since its atomic structure towards its molecular structure. Indeed, as it is revealed in this study, the elements working within living matter are not randomly organized according to arithmetic criteria which depend on their primordial quantum structure.

This 3/2 fractional value arithmetic ratio is very similar to the fractional values of the electric charges of different quarks, which are ratios of whole numbers (2/3 and -1/3). Thus these phenomena, operating in the most complex organization of matter, depend on its most basic structure. By the amplitude of the phenomena presented here, it is therefore not possible to imagine a non-relation between this primary structure of matter and its highly organized structure as it appears in the structural mechanics of the genetic code.

Annexe

A1. Quantum chart in chevron form

A1.1 Depiction

Figure 26 shows the chevron form quantum table of the first 15 electronic shells. This graphic concept is introduced in Chapter 4.1 and illustrated in Figure 5. We suggest that this new graphic type be favoured for the description of the quantum organization of the different chemical elements.

In this table, the different quantum shells and subshells are presented in the form of chevrons. At the top end of each rafter are indicated the names of the different shells and subshells; at the left end of these chevrons, the numbers of orbits and electrons of these different shells and quantum subshells are indicated. At each chevron vertex is the orbital where the quantum number $m = 0$. The orbitals with positive quantum number m are progressively positioned towards the top of these chevron vertices and the orbitals with negative quantum number m are progressively positioned towards the outside left of these chevron vertices.

This new graphic design is more explicit in describing the quantum structure of atomic elements than any other usual linear chart. Very visually, this chevron configuration clearly highlights the arithmetic progression of the orbital numbers of the different quantum layers in square powers of the level of these electronic shells:

- 1st shell → $1^2 = 1$ orbital,
- 2nd shell → $2^2 = 4$ orbitals,
- 3rd shell → $3^2 = 9$ orbitals,
- 4th shell → $4^2 = 16$ orbitals, etc.

In this table, and more generally in this quantum study of the atomic elements, constituents of the genetic code, the electronic spin is not detailed (by ascending or descending arrows). In return, it is the migratory or non-migratory nature of the electrons that is the subject of study. Thus, for example, the representation of the nitrogen atom such as that illustrated in Figure. 25 (and in various other figures of this quantum study) is favoured.

chevron form quantum representation of the atomic element 7 (N):

2 quantum shells
 3 subshells
 5 orbitals ($1^2 + 2^2$)
 10 orbiting electrons whose
 7 own (●) + 3 guest (◦)

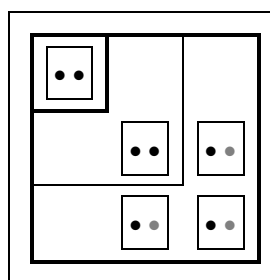


Fig. 25 Graphical quantum representation of Nitrogen in chevron form design (in its saturated state). See also Fig. 5 and 13.

A1.2 Quantum chart

orbital amount		shells and subshells																
electron amount •		1(K)			2(L)		3(M)			4(N)				5(O)				
by shell:	by subshell:	1s	2s	2p	3s	3p	3d	4s	4p	4d	4f	5s	5p	5d	5f	5g		
1	2	1	2															
4	8	1	2															
		3	6															
9	18	1	2															
		3	6															
		5	10															
16	32	1	2															
		3	6															
		5	10															
		7	14															
25	50	1	2															
		3	6															
		5	10															
		7	14															
		9	18															

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Fig. 26 Chevron form chart representing the first 5 shells and first 15 quantum subshells of the atomic elements. Distribution of orbitals and electrons in these shells and subshells.

A.2 Number of protons and modules of the twenty amino acids

A.2.1 Modules of Petoukhov

The notion of module is an original system proposed by Sergei Petoukhov [1 and 2] to describe the structure of biological molecules. This system represents each non-hydrogen atom by the number of protons in its nucleus. For example, the nitrogen atom is represented by the number 7. If one or more hydrogen atoms are joined to a non-hydrogen atom, the number of hydrogen protons is added to the number of atom protons of non-hydrogen.

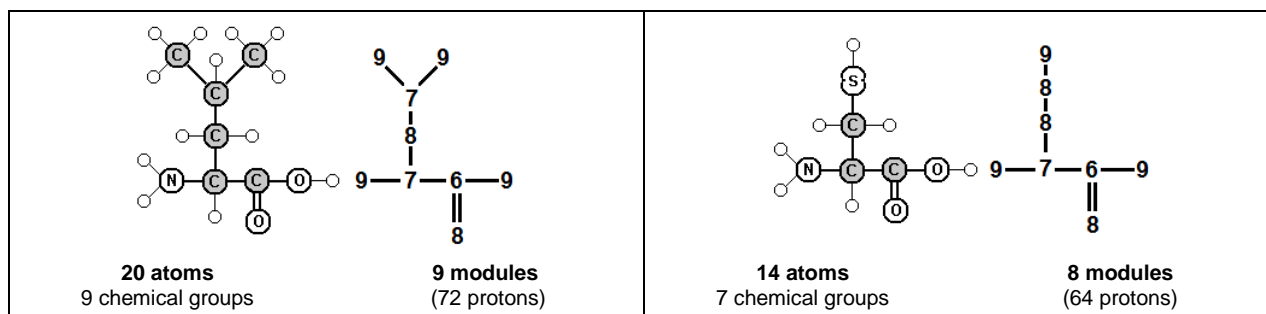


Fig. 27 Molecular and modular structure of amino acids (here Leucine and Cysteine).

Thus, such a group made up of a non-hydrogen atom and its adjoining satellite atoms of Hydrogen is defined by the total sum of protons of this group and this set is called "module". Figure 27 illustrates this modular concept applied to amino acids.

For example, the amide group NH₂ forms a module: it is denoted by the number 9 which is the sum of 7 protons of one nitrogen atom and 2 protons of two satellite atoms of Hydrogen. Sergei Petoukhov provides [1 and 2] modules from 6 to 9 protons according to the configurations. Thus according to this system, the Sulphur atoms (present in *Met* and *Cys*) break down conventionally into two modules of 8 protons (or 9 protons if associated with a Hydrogen in *Cys*).

A.2.2 Protons/modules multiplicity

The genetic code is organized only with 20 amino acids. Figure 28 isolates amino acids with a number of protons multiple of the number of modules from those with a number of protons not multiple of the number of modules. It turns out that these two groups are of equal size since they each consist of 10 entities.

10 amino acids with protons/modules multiplicity	Gly 40*/5**	Ala 48/6	Ser 56/7	Val 64/8	10 amino acids without protons/modules multiplicity
	Cys 64/8	Thr 64/8	Leu 72/9	Ile 72/9	
	Lys 80/10	Met 80/10	Phe 88/12	Tyr 96/13	
	Pro 62/8	Asn 70/9	Asp 70/9	Gln 78/10	
	Glu 78/10	His 82/11	Arg 94/12	Trp 108/15	

Fig.28 Differentiation of the 20 amino acids according to their protons*/modules** multiplicity.

It is remarkable to note that the respective number of protons of each of the ten amino acids of the first group, that whose number of protons is multiple of the number of modules, is always equal to eight times the number of modules.

Also, for this first group of ten amino acids, the total number of modules is equal to 80, so an average of 8 modules into each amino acid. Thus, the total number of protons in this group is therefore equal to 640 protons, an average of 64 (8²) protons by amino acid. Also, for each of the ten other amino acids, the proton/module ratio never represents an integer but a fraction different from $x/1$ (x being an integer).

A.2.3 Other multiplicities

As described in the left part of Figure 29, it turns out that two other amino acids have a number of protons multiple of 8 but not multiple of their respective number of modules: *Phe* with 88 protons for 12 modules and *Tyr* with 96 protons for 13 modules. Thus, by these multiplicity criteria, in a ratio of value 3/2 are opposed twelve amino acids with number of protons multiple of 8 versus eight amino acids with number of protons not multiple of 8.

<table border="1" style="width: 100%; text-align: center;"> <tr><td>Gly 40</td><td>Ala 48</td><td>Ser 56</td><td>Val 64</td></tr> <tr><td>Cys 64</td><td>Thr 64</td><td>Leu 72</td><td>Ile 72</td></tr> <tr><td>Lys 80</td><td>Met 80</td><td>Phe 88</td><td>Tyr 96</td></tr> <tr><td>Pro 62</td><td>Asn 70</td><td>Asp 70</td><td>Gln 78</td></tr> <tr><td>Glu 78</td><td>His 82</td><td>Arg 94</td><td>Trp 108</td></tr> </table>	Gly 40	Ala 48	Ser 56	Val 64	Cys 64	Thr 64	Leu 72	Ile 72	Lys 80	Met 80	Phe 88	Tyr 96	Pro 62	Asn 70	Asp 70	Gln 78	Glu 78	His 82	Arg 94	Trp 108	← 3/2 → ratio	<table border="1" style="width: 100%; text-align: center;"> <tr><td>Gly 40/5</td><td>Ala 48/6</td><td>Ser 56/7</td><td>Val 64/8</td></tr> <tr><td>Thr 64/8</td><td>Leu 72/9</td><td>Ile 72/9</td><td>Lys 80/10</td></tr> <tr><td>Pro 62/8</td><td>Cys 64/7</td><td>Asn 70/9</td><td>Asp 70/9</td></tr> <tr><td>Gln 78/10</td><td>Glu 78/10</td><td>Met 80/9</td><td>His 82/11</td></tr> <tr><td>Phe 88/12</td><td>Arg 94/12</td><td>Tyr 96/13</td><td>Trp 108/15</td></tr> </table>	Gly 40/5	Ala 48/6	Ser 56/7	Val 64/8	Thr 64/8	Leu 72/9	Ile 72/9	Lys 80/10	Pro 62/8	Cys 64/7	Asn 70/9	Asp 70/9	Gln 78/10	Glu 78/10	Met 80/9	His 82/11	Phe 88/12	Arg 94/12	Tyr 96/13	Trp 108/15	← 2/3 → ratio	<table border="1" style="width: 100%; text-align: center;"> <tr><td colspan="4">12 amino acids with number of protons multiple of 8</td></tr> </table>	12 amino acids with number of protons multiple of 8				<table border="1" style="width: 100%; text-align: center;"> <tr><td colspan="4">8 amino acids without number of protons multiple of 8</td></tr> </table>	8 amino acids without number of protons multiple of 8				<table border="1" style="width: 100%; text-align: center;"> <tr><td colspan="4">8 amino acids with protons/groups multiplicity</td></tr> </table>	8 amino acids with protons/groups multiplicity				<table border="1" style="width: 100%; text-align: center;"> <tr><td colspan="4">12 amino acids without protons/groups multiplicity</td></tr> </table>	12 amino acids without protons/groups multiplicity			
Gly 40	Ala 48	Ser 56	Val 64																																																												
Cys 64	Thr 64	Leu 72	Ile 72																																																												
Lys 80	Met 80	Phe 88	Tyr 96																																																												
Pro 62	Asn 70	Asp 70	Gln 78																																																												
Glu 78	His 82	Arg 94	Trp 108																																																												
Gly 40/5	Ala 48/6	Ser 56/7	Val 64/8																																																												
Thr 64/8	Leu 72/9	Ile 72/9	Lys 80/10																																																												
Pro 62/8	Cys 64/7	Asn 70/9	Asp 70/9																																																												
Gln 78/10	Glu 78/10	Met 80/9	His 82/11																																																												
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Fig. 29 Differentiation of the 20 amino acids according to the multiplicity by 8 of their number of protons (ratio 3/2) and according to their multiplicity protons/groups (ratio 2/3).

Also, as described in the right part of Figure 29, in an inverse ratio of 2/3 value, eight amino acids versus twelve have a number of protons multiple of their respective number of chemical groups. As explained above, the nuance between chemical groups and modules is observed for chemical groups composed of a sulphur atom, an atom conventionally made up of two modules (see Figure 27). Thus, the amino acids *Cys* and *Met* have an amount of chemical groups which differs to one value in relation to their number of modules.

A.2.4 Three progressive arithmetical ratios

Thus, illustrated in Figure 30, according to three different criteria for the multiplicity of proton numbers, the ratio between the twenty amino acids operating in the genetic code changes from the value 3/2 to the value 1/1 then to the value of 3/2, inverse to the initial.

proton number multiplicity by 8			protons/modules multiplicity			protons/groups multiplicity		
yes		no	yes		no	yes		no
12 entities	$\leftarrow \frac{3}{2} \rightarrow$ ratio	8 entities	10 entities	$\leftarrow \frac{1}{1} \rightarrow$ ratio	10 entities	8 entities	$\leftarrow \frac{2}{3} \rightarrow$ ratio	12 entities

Fig. 30 Development of the ratio between amino acids according to three criteria of multiplicity of proton numbers. See Fig. 28 and 29.

All these observations confirm the differentiations made in the chemical, quantum and atomic study of Glycine between the modules (chemical groups) with a proton number equal to 8 and from the modules with a proton number not equal to 8.

Références

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3. www.elementschimiques.fr

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