



# Are Irrational Numbers (Like the Square Root of the Number Seven) Applicable to Genetic Sequences?

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## Abstract

According to Quantum Perspective Model, this article researches whether there is a link between the square root of **seven** numbers and the genetic sequences. At first, the square root digits of the number seven after the comma are added respectively. Secondly, the resulting sum corresponds to the nucleotide bases, the results obtained in this way are expressed as nucleotide bases. (**A, T, C, G, and U**). (**A**) Adenine, (**T**) Thymine, (**C**) Cytosine, (**G**) Guanine, (**U**) **Uracil**. From this point of view, when the first four hundred and fifty digits of the square root of the number seven after the comma are calculated, the gene sequence is obtained as follows:

[**GATTUAAAGUTAATATTAUTAGTTTGATT**]. Thirdly, after researching this sequence at NCBI (National Biotechnology Information Center), the search result is similar to **bony fishes**, especially *Danio rerio* (zebrafish). Fourthly, the genetic codes of zebrafishes were found to be similar to human genetic codes. Lastly, some repetitions were detected exactly like this: as “**GAT**” and “**UTA**”. In summary, the connection between these results and the square root of the seven in mathematical science and the genetic codes in biochemistry shed light on explaining irrational numbers.

## Subject Areas

Biochemistry, Mathematics, Number Theory

## Keywords

Quantum Perspective Model, Biochemistry, *Danio rerio*, *Danio aesculapii*, Timema, the Square Root of Seven and NCBI (National Biotechnology Information Center)

# 1. Introduction and Calculation of the Square Root of Seven numbers

## 1.1. Introduction

The relationships between numbers and genetic codes are not only research with the square root of the number two [1] and but also researched with the square root of the number three [2] and the square root of the number five [3], too from the Quantum Perspective Model [4], this paper attempts to search the relationships between the square root of the number seven and the genetic codes.

## 1.2. Calculating the Square Root of Seven

The square root of seven =

2.645751311064590590501615753639260425710259183082450180368334459201068823;  
 2302836277603928864745436106150645783384974630957435298886272147844273905;  
 558801077227171507297283238922996895948650872607009780542037238280237159;  
 4110034193911600157852559630594574103515239680271640737379907404158151990440347431;  
 945367139973059700505139969223754561609711902737815499163328828770400065757;  
 0674651963497752083793818114613090876473786595624330579947981281632307054 [5].

## 2. Methods and Discussion

### 2.1. Methods

The chemical structures of bases include Carbon (C), Nitrogen (N), Oxygen (O), and Hydrogen (H). **Calculation of bases with chemical atoms** (See also **Table 1**) (Ölmez T, 2020) [6].

The atomic numbers of them: Carbon (C): 6, Nitrogen (N): 7, Oxygen (O): 8, Hydrogen (H): 1 (Wieser E M *et al.*, 2013) [7]. The chemical structures of bases (A, T, C, G, and U) are shown below (Ölmez T, 2020) [6].

**Table 1.** Representation of nucleotide bases (A, T, C, G and U) in chemical atoms.

ATOMS/NUCLEOTIDE BASES	C = 6	H = 1	O = 8	N = 7	SUM
ADENINE: C <sub>5</sub> H <sub>5</sub> N <sub>5</sub>	5	5	-	5	70
THYMINE: C <sub>5</sub> H <sub>6</sub> N <sub>2</sub> O <sub>2</sub>	5	6	2	2	66
CYTOSINE: C <sub>4</sub> H <sub>5</sub> N <sub>3</sub> O <sub>1</sub>	4	5	1	3	58
GUANINE: C <sub>5</sub> H <sub>5</sub> N <sub>5</sub> O <sub>1</sub>	5	5	1	5	78
URACIL: C <sub>4</sub> H <sub>4</sub> N <sub>2</sub> O <sub>2</sub>	4	4	2	2	58

(A) Adenine: C<sub>5</sub>H<sub>5</sub>N<sub>5</sub>: **70**; (T) Thymine: C<sub>5</sub>H<sub>6</sub>N<sub>2</sub>O<sub>2</sub>: **66**, (C) Cytosine: C<sub>4</sub>H<sub>5</sub>N<sub>3</sub>O<sub>1</sub>: **58**, (G) Guanine: C<sub>5</sub>H<sub>5</sub>N<sub>5</sub>O<sub>1</sub>: **78**, and (U) Uracil: C<sub>4</sub>H<sub>4</sub>N<sub>2</sub>O<sub>2</sub>: **58** (Lodish H *et al.*, 2018) [8].

## 2.2. Discussion

Every cell in our body contains genes, but only a certain subset of them is “turned on” or expressed in every cell. The amount of protein in each cell is controlled by a series of different processes. Now, if we analyze the genetic languages with the lowest level of *information processing*, it is the language of DNA and RNA with four nucleotide base alphabets and the language of proteins with twenty amino acid alphabets (Jonathan S, 2005) [9]. *Genetic information* is stored in DNA, the mobile messenger nucleic acid is transcribed into RNA and translated into amino acid sequences, which are then folded into proteins. Now, if we look at the *Quantum Perspective Model* starting from the amino acid class in the genetic code; The universal triplet has significant degeneration in genetic codes, with 64 codons (Penrose R *et al.*, 2008) [10]. The reason it’s called the QUANTUM PERSPECTIVE MODEL, which evokes quantum mechanics, implies that the universe is effectively computing not only at the most microscopic level but also at larger scales (Penrose R *et al.*, 2008) [10].

## 3. Calculation of the Square Root of Seven Numbers and Genetic Codes

The first four hundred and fifty digits of the square root of seven after the comma are here:

2.6457513110645905905016157536392604257102591830824501803683344592  
01068823;  
230283627760392886474543610615064578338497463095743529888627214784  
4273905;  
558801077227171507297283238922996895948650872607009780542037238280  
237159;  
411003419391160015785255963059457410351523968027164073737990740415  
8151990440347431;  
945367139973059700505139969223754561609711902737815499163328828770  
400065757;  
067465196349775208379381811461309087647378659562433057994798128163  
2307054 [5].

At first, the first group of the square root numbers of seven after comma was taken. For example, 6, 4, 5, 7, 5, 1, 3, 1, 1, 0, 6, 4, 5, 9, 0, 5, 9, 0, 5, 0, 1, ...and so on. Secondly, all decimal numbers are subjected to the addition process, respectively.  $(6+4+5+7+5+1+3+1+1+0+6+4+5+9+0+5+9+0+5+0+1 = 77)$ . The sum of the first group of the root square numbers of seven after comma is “77”. Just like as in (G) Guanine: 78 (See also **Table 1**).

The **first** group of the root square numbers of seven after comma:

$6+4+5+7+5+1+3+1+1+0+6+4+5+9+0+5+9+0+5+0+1 = 77$  (**G**) Guanine: **78**

The **second** group of the root square numbers of seven after comma:

$6+1+5+7+5+3+6+3+9+2+6+0+4+2+5+7 = 70$  (**A**) Adenine: **70**

The **third** group of the root square numbers of seven after comma:

$1+0+2+5+9+1+8+3+0+8+2+4+5+0+1+8+0+3+6 = 66$  (**T**) Thymine: **66**

The **fourth** group of the root square numbers of seven after comma:

$8+3+3+4+4+5+9+2+0+1+0+6+8+8+2+3 = 66$  (**T**) Thymine: **66**

The **fifth** group of the root square numbers of seven after comma:

$2+3+0+2+8+3+6+2+7+7+6+0+3+9 = 58$  (**U**) Uracil: **58**

The **sixth** group of the root square numbers of seven after comma:

$2+8+8+6+4+7+4+5+4+3+6+1+0+6+1+5 = 70$  (**A**) Adenine: **70**

The **seventh** group of the root square numbers of seven after comma:

$0+6+4+5+7+8+3+3+8+4+9+7+4 = 68$  (**A**) Adenine: **70**

The **eighth** group of the root square numbers of seven after comma:

$6+3+0+9+5+7+4+3+5+2+9+8+8 = 69$  (**A**) Adenine: **70**

The **ninth** group of the square numbers of seven after comma:

$8+6+2+7+2+1+4+7+8+4+4+2+7+3+9+0+5 = 79$  (**G**) Guanine: **78**

The **tenth** group of the root square numbers of seven after comma:

$5+5+8+8+0+1+0+7+7+2+2+7+1+7 = 60$  (**U**) Uracil: **58**

The **eleventh** group of the root square numbers of seven after comma:

$1+5+0+7+2+9+7+2+8+3+2+3+8+9 = 66$  (**T**) Thymine: **66**

The **twelfth** group of the root square numbers of seven after comma:

$2+2+9+9+6+8+9+5+9+4+8 = 71$  (**A**) Adenine: **70**

The **thirteenth** group of the root square numbers of seven after comma:

$6+5+0+8+7+2+6+0+7+0+0+9+7+8+0+5 = 70$  (**A**) Adenine: **70**

The **fourteenth** group of the root square numbers of seven after comma:

$4+2+0+3+7+2+3+8+2+8+0+2+3+7+1+5+9 = 66$  (**T**) Thymine: **66**

The **fifteenth** group of the root square numbers of seven after comma:

$4+1+1+0+0+3+4+1+9+3+9+1+1+6+0+0+1+5+7+8+5+2 = 71$  (**A**) Adenine: **70**

The **sixteenth** group of the root square numbers of seven after comma:

$5+5+9+6+3+0+5+9+4+5+7+4+1+0+3 = 66$  (**T**) Thymine: **66**

The **seventh** group of the root square numbers of seven after comma:

$5+1+5+2+3+9+6+8+0+2+7+1+6+4+0+7 = 66$  (**T**) Thymine: **66**

The **eighteenth** group of the root square numbers of seven after comma:

$3+7+3+7+9+9+0+7+4+0+4+1+5+8+1 = 68$  (**A**) Adenine: **70**

The **nineteenth** group of the root square numbers of seven after comma:

$5+1+9+9+0+4+4+0+3+4+7+4+3+1 = 58$  (**U**) Uracil: **58**

The **twentieth** group of the root square numbers of seven after comma:

$9+4+5+3+6+7+1+3+9+9+7+3 = 66$  (**T**) Thymine: **66**

The **twenty-first** group of the root square numbers of seven after comma:

$0+5+9+7+0+0+5+0+5+1+3+9+9+6+9+2 = 70$  (**A**) Adenine: **70**

The **twenty-second** group of the root square numbers of seven after comma:

$2+3+7+5+4+5+6+1+6+0+9+7+1+1+9+0+2+7+3 = 78$  (**G**) Guanine: **78**

The **twenty-third** group of the root square numbers of seven after comma:

$7+8+1+5+4+9+9+1+6+3+3+2+8 = 66$  (T) Thymine: **66**

The **twenty-fourth** group of the root square numbers of seven after comma:

$8+2+8+7+7+0+4+0+0+0+6+5+7+5+7 = 66$  (T) Thymine: **66**

The **twenty-fifth** group of the root square numbers of seven after comma:

$0+6+7+4+6+5+1+9+6+3+4+9+7 = 67$  (T) Thymine: **66**

The **twenty-sixth** group of the root square numbers of seven after comma:

$7+5+2+0+8+3+7+9+3+8+1+8+1+1+4+6+1+3 = 77$  (G) Guanine: **78**

The **twenty-seventh** group of the root square numbers of seven after comma:

$0+9+0+8+7+6+4+7+3+7+8+6+5 = 70$  (A) Adenine: **70**

The **twenty-eighth** group of the root square numbers of seven after comma:

$9+5+6+2+4+3+3+0+5+7+9+9+4 = 66$  (T) Thymine: **66**

The **twenty-ninth** group of the root square numbers of seven after comma:

$7+9+8+1+2+8+1+6+3+2+3+0+7+0+5+4 = 66$  (T) Thymine: **66**

This sequence can be shown as

[GATTUAAGUTAATATTAUTAGTTTGATT]. Let me try to explain this sequence with the “Quantum Perspective Model”. For example, The **first** group of the square root of seven after comma is equal to Guanine (G): 77 with the lack of one “1” Hydrogen bond (H:1) (Remember, See **Table 1**; Guanine (G): 78). This result may mean the sequence of the square root of seven in groups [GATTUAAGUTAATATTAUTAGTTTGATT]. The **fifteenth** group of the square root of seven after the comma is regarded as Adenine (A) with one more Hydrogen bond (H:1); Adenine: 71. (Remember, See **Table 1**; (A) Adenine:70) (Because the **deviations** in the calculation of the square root of seven numbers can be derived from the Cytosine (C)—Guanine (G) Hydrogen bonds because of Cytosine (C)—Guanine (G) pairs with by *three* hydrogen bonds. Adenine (A) pairs with Thymine (T) by *two* hydrogen bonds [6]. The reason for the lack/more of hydrogen bonds: **Hydrogen bonding** is a very versatile attraction (Ölmez T, 2020) [6]. Hydrogen bonds are relatively *weak and easily broken* by increasing hardness (Farrell R E, 2010) [11].

#### 4. Results

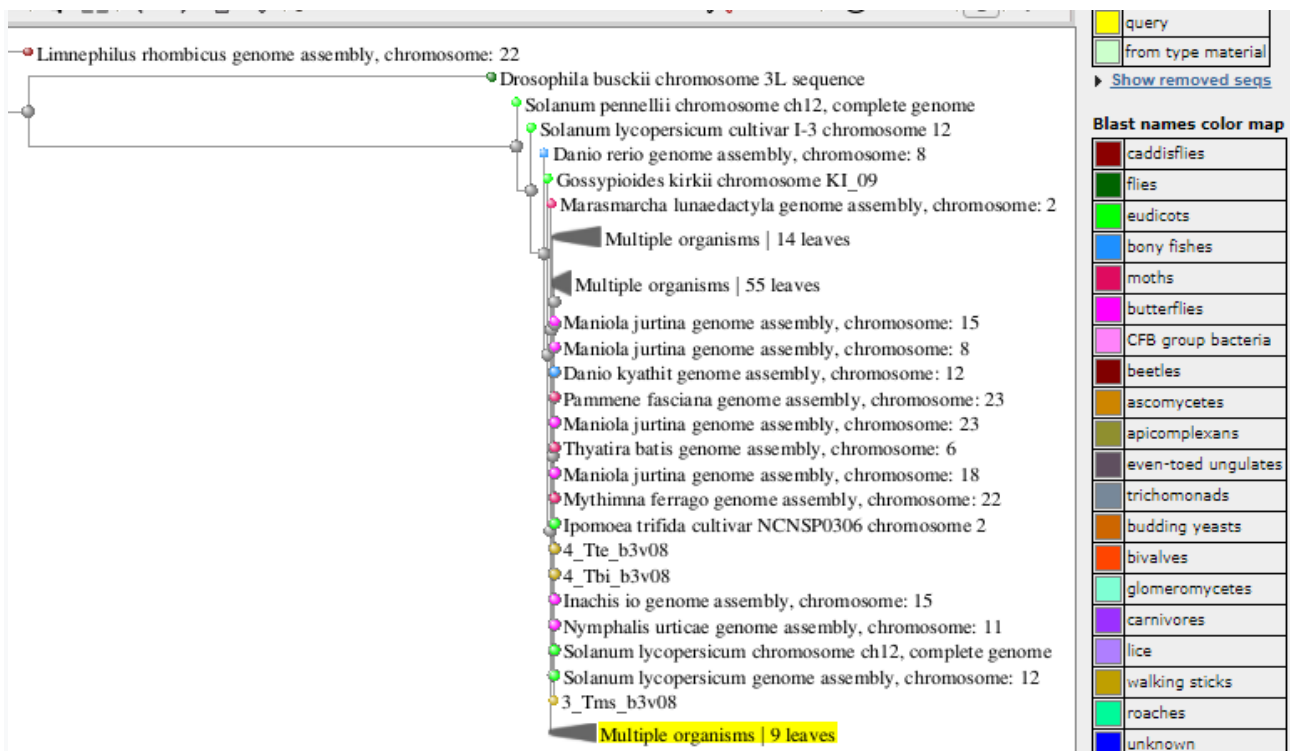
After searching the square root of the number seven with the National Biotechnology Information Center (NCBI) databases [12], some conceptual relationships with **bony fishes** can ultimately be found. Types of **bony fishes** are based on *Danio rerio* (zebrafish) (See **Figure 1**). Types of other living creatures are eudicots, walking sticks, *Danio aesculapii*, flies beetles, moths, butterflies, lice, bivalves, roaches, hawks and eagles, caddisflies, *Timema tahoe*, *Timema bartmani* and carnivores. Another interesting result of NCBI is *Timema shepardii*, an animal that has not engaged in sexual reproduction for many years and has the longest known asexual period [11] (See **Figure 2**).

#### 5. Conclusions

At first, the result of this research can be summarized as the expression of root

<a href="#">Danio rerio genome assembly, chromosome: 8</a>	<a href="#">Danio rerio</a>	42.1	235	100%	0.98	100.00%	55941223	<a href="#">LR812045.1</a>
<a href="#">Danio rerio genome assembly, chromosome: 8</a>	<a href="#">Danio rerio</a>	42.1	237	85%	0.98	100.00%	51589891	<a href="#">LR812601.1</a>
<a href="#">Danio rerio strain Nadia (NA) genome assembly, chromosome: 8</a>	<a href="#">Danio rerio</a>	42.1	271	85%	0.98	100.00%	53615831	<a href="#">LR812576.1</a>
<a href="#">Danio rerio strain Cooch Behar (CB) genome assembly, chromosome: 8</a>	<a href="#">Danio rerio</a>	42.1	241	85%	0.98	100.00%	53882377	<a href="#">LR812551.1</a>
<a href="#">Diptera punctata clone 19Y milk protein mRNA, partial cds</a>	<a href="#">Diptera pun...</a>	42.1	42.1	89%	0.98	96.00%	649	<a href="#">AY448005.1</a>
<a href="#">Accipiter gentilis genome assembly, chromosome: 17</a>	<a href="#">Accipiter gentilis</a>	40.1	108	100%	3.9	100.00%	31036983	<a href="#">OV839378.1</a>
<a href="#">Limnephilus rhombicus genome assembly, chromosome: 22</a>	<a href="#">Limnephilus rh...</a>	40.1	345	100%	3.9	100.00%	46446577	<a href="#">OV815292.1</a>
<a href="#">Macaria notata genome assembly, chromosome: 27</a>	<a href="#">Macaria notata</a>	40.1	72.4	85%	3.9	95.83%	9005937	<a href="#">OV656751.1</a>
<a href="#">Brassicogethes aeneus genome assembly, chromosome: 4</a>	<a href="#">Brassicogethe...</a>	40.1	736	100%	3.9	100.00%	61642216	<a href="#">OV121135.1</a>
<a href="#">Sphingobacterium sp. SRCM116780 chromosome, complete genome</a>	<a href="#">Sphingobacteri...</a>	40.1	40.1	71%	3.9	100.00%	4282846	<a href="#">CP090446.1</a>
<a href="#">Meles meles genome assembly, chromosome: 13</a>	<a href="#">Meles meles</a>	40.1	74.3	85%	3.9	100.00%	87850455	<a href="#">OV277454.1</a>
<a href="#">Notodonta ziczac genome assembly, chromosome: 4</a>	<a href="#">Notodonta ziczac</a>	40.1	40.1	71%	3.9	100.00%	14050355	<a href="#">OU974041.1</a>
<a href="#">Calonectria ilicicola strain GDYJ169 chromosome 10</a>	<a href="#">Calonectria ilic...</a>	40.1	110	85%	3.9	95.83%	4267126	<a href="#">CP085834.1</a>
<a href="#">Rhizophagus irregularis strain A1 chromosome 4</a>	<a href="#">Rhizophagus ir...</a>	40.1	176	82%	3.9	100.00%	6036189	<a href="#">CP080923.1</a>
<a href="#">Rhizophagus irregularis strain C2 chromosome 4</a>	<a href="#">Rhizophagus ir...</a>	40.1	142	82%	3.9	100.00%	6439466	<a href="#">CP080888.1</a>
<a href="#">Rhizophagus irregularis strain B3 chromosome 4</a>	<a href="#">Rhizophagus ir...</a>	40.1	142	82%	3.9	100.00%	5903266	<a href="#">CP080856.1</a>
<a href="#">Melinaea marsaeus rileyi genome assembly, chromosome: 13</a>	<a href="#">Melinaea mars...</a>	40.1	406	89%	3.9	100.00%	25081093	<a href="#">OU911094.1</a>
<a href="#">Synanthedon vespiformis genome assembly, chromosome: 3</a>	<a href="#">Synanthedon v...</a>	40.1	40.1	71%	3.9	100.00%	12113659	<a href="#">OU906948.1</a>
<a href="#">Lasiommata megera genome assembly, chromosome: 14</a>	<a href="#">Lasiommata m...</a>	40.1	205	96%	3.9	100.00%	17193394	<a href="#">OV743321.1</a>

**Figure 1.** The NCBI (National Center for Biotechnology Information) result for nucleotide sequence “GATTUAAGUTAATATTAUTAGTTTGATT” [13]. ([https://blast.ncbi.nlm.nih.gov/Blast.cgi?PROGRAM=blastn&PAGE\\_TYPE=BlastSearch&LINK\\_LOC=blasthome](https://blast.ncbi.nlm.nih.gov/Blast.cgi?PROGRAM=blastn&PAGE_TYPE=BlastSearch&LINK_LOC=blasthome))



**Figure 2.** The NCBI (National Center for Biotechnology Information) result blast tree view widget for “GATTUAAGUTAATATTAUTAGTTTGATT” nucleotide sequence [14]. ([https://www.ncbi.nlm.nih.gov/blast/treeview/treeView.cgi?request=page&blastRID=32WBVGR8016&queryID=lcl|Query\\_51663&entrezLim=&ex=&exl=&exh=&ns=100](https://www.ncbi.nlm.nih.gov/blast/treeview/treeView.cgi?request=page&blastRID=32WBVGR8016&queryID=lcl|Query_51663&entrezLim=&ex=&exl=&exh=&ns=100))

seven numbers after commas by genetic codes. Secondly, digits after the comma were thought to be an indicator of genetic codes. As a common feature of Bio-

chemistry and Mathematical sciences, NCBI blasts are obtained [12]. Because these results include both bony fishes and especially *Danio Rerio* (zebrafish) (See **Figure 1**), some genetic stem cells have also been studied in zebrafish (*Danio rerio*) in studies with functional macrophages (Takahashi K, 2001) [15]. Zebrafish not only have muscarinic in their nervous system but also, receptor isoforms not previously identified in other species are present in zebrafish (Nuckels R J, 2006) [16] and (Abrams P *et al.* 2006) [17]. Although there are no specific subtypes of muscarinic acetylcholine receptors, detected in the zebrafish retina. An enzyme was studied in the developing zebrafish brain and retina. But more work may be needed to confirm the studies and experiment with zebrafish (Nuckels R J, 2006).

In biology-related experiments and gene sequencing, it is an excellent favorite example [18]. The common feature of pi numbers [4] and Euler numbers [19] [20] along with the square of the speed of light is that the NCBI results are bony fish [21] [22]. Let alone this result, NCBI result for CAAT Box also consists of **bony fishes**, too [6]. Thirdly, the genetic codes of the square root of two [1] and the genetic codes of the square root of three consist of the same NCBI blast results as the same as bony fishes [2]. Fourthly, although there is no periodic sequence of irrational numbers, in this paper a periodic sequence is obtained in terms of genetic sequences, just as in “GAT” and “UTA”. Remember, this sequence can be shown as [GATTUAAGUTAATATTAUTAGTTTGATT]. Finally, the results of calculating the square root of seven numbers with genetic codes can be expressed by chemical formulas of nucleotide bases. It is stated by Steward that life is between genes and Mathematics (Stewart I, 1999) [23]. Indeed, the relationships between *Mathematics* and *Genetics* have been described by a molecular-genetic alphabet matrix. According to the quantum perspective model, just like this matrix, is it possible to define the square root of the number seven as genetic sequences? (Petoukhov S V, 2011) [24].

## Conflicts of Interest

The author declares no conflicts of interest.

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- [14] [https://www.ncbi.nlm.nih.gov/blast/treewiew/treeView.cgi?request=page&blastRID=32WBVGR8016&queryID=lcl|Query\\_51663&entrezLim=&ex=&exl=&exh=&ns=100](https://www.ncbi.nlm.nih.gov/blast/treewiew/treeView.cgi?request=page&blastRID=32WBVGR8016&queryID=lcl|Query_51663&entrezLim=&ex=&exl=&exh=&ns=100)
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