The Kodonian Project

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Welcome

i Purpose & Goal

The Kodonian Project seeks to demonstrate that the Universal Genetic Code contains an embedded signal—an intentional, structured message—suggesting a deeper informational or even extraterrestrial origin behind the blueprint of life on Earth.

Preface

here we go... $\,$

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FAQ

Find here answers to common questions around the topic of our Genetic Code Signal research.

i Are we serious?

Yes

Who is the main contributor of this project?

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i Has research in this area been conducted elsewhere?

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i What are the strongest arguments for a signal in the Genetic Code?

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What are the strongest arguments against a signal in the Genetic Code?

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i Evolution did it, didn't it?

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1 Demo Site

1.1 MathJax - mhchem

1.2 R

1+1

1.3 Python

Quarto enables you to weave together content and executable code into a finished document. To learn more about Quarto see https://quarto.org.

```
db_table(
   'SELECT name, symbol3, symbol1, nuclsum_b, nuclsum_s, nuclsum_t FROM amino_acids ORDER BY nu
   headers=['Name', 'Short Name', 'Symbol', 'Block Sum', 'Side Chain Sum', 'Total Sum'])
```

| Name | Short Name | Symbol | Block Sum | Side Chain Sum | Total Sum |
|------------|----------------------|--------------|-----------|----------------|-----------|
| STOP | Stop | * | 0 | 0 | 0 |
| Glycine | Gly | G | 74 | 1 | 75 |
| Alanine | Ala | A | 74 | 15 | 89 |
| Serine | Ser | \mathbf{S} | 74 | 31 | 105 |
| Proline | Pro | P | 74 | 41 | 115 |
| Valine | Val | V | 74 | 43 | 117 |
| Threonine | Thr | ${ m T}$ | 74 | 45 | 119 |
| Cysteine | Cys | \mathbf{C} | 74 | 47 | 121 |
| Leucine | Leu | L | 74 | 57 | 131 |
| Isoleucine | Ile | I | 74 | 57 | 131 |
| Asparagine | Asn | N | 74 | 58 | 132 |

| Name | Short Name | Symbol | Block Sum | Side Chain Sum | Total Sum |
|------------------|----------------------|--------------|-----------|----------------|-----------|
| Aspartic Acid | Asp | D | 74 | 59 | 133 |
| Lysine | Lys | K | 74 | 72 | 146 |
| Glutamine | Gln | Q | 74 | 72 | 146 |
| Glutamic Acid | Glu | ${ m E}$ | 74 | 73 | 147 |
| Methionine/START | Met | M | 74 | 75 | 149 |
| Histidine | His | Η | 74 | 81 | 155 |
| Phenylalanine | Phe | \mathbf{F} | 74 | 91 | 165 |
| Arginine | Arg | \mathbf{R} | 74 | 100 | 174 |
| Tyrosine | Tyr | Y | 74 | 107 | 181 |
| Tryptophane | Trp | W | 74 | 130 | 204 |

i Expand for a list of Genetic Code variants

Source

| Id | Species |
|--------------------------|--|
| $\overline{\text{GC01}}$ | Standard Code |
| GC02 | Vertebrate Mitochondrial Code |
| GC03 | Yeast Mitochondrial Code |
| GC04 | Mold, Protozoan, Coelenterate Mitochondrial Code |
| GC05 | Invertebrate Mitochondrial Code |
| GC06 | Ciliate, Dasycladacean and Hexamita Nuclear Code |
| GC07 | Deleted Kinetoplast Code |
| GC09 | Echinoderm and Flatworm Mitochondrial Code |
| GC10 | Euplotid Nuclear Code |
| GC11 | Bacterial, Archaeal and Plant Plastid Code |
| GC12 | Alternative Yeast Nuclear Code |
| GC13 | Ascidian Mitochondrial Code |
| GC14 | Alternative Flatworm Mitochondrial Code |
| GC15 | Blepharisma Nuclear Code |
| GC16 | Chlorophycean Mitochondrial Code |
| GC21 | Trematode Mitochondrial Code |
| GC22 | Scenedesmus Obliquus Mitochondrial Code |
| GC23 | Thraustochytrium Mitochondrial Code |
| GC24 | Pterobranchia Mitochondrial Code |
| GC25 | Candidate Division SR1 and Gracilibacteria Code |
| GC26 | Pachysolen Tannophilus Nuclear Code |

| GG | Gly | GA | Asp G l u | AG | Ser Arg | АА | Asn Lys | |
|----|--------------|----|---------------------|------|--------------------|----|-------------|---------|
| | 75 | | 280 | | 279 | | 278 | 1517 |
| GC | A l a | GT | Val | AC | Thr | AT | lle Met | 41 x 37 |
| | 89 | | 117 | | 119 | | 280 | |
| CG | Arg | CA | His G i n | TG | Cys Stop Trp | TA | Tyr Stop | |
| | 174 | | 301 | | 325 | | 181 | 1628 |
| СС | Pro | СТ | Leu | TC | Ser | П | Phe Leu | 44 x 37 |
| | 115 | | 131 | | 105 | | 296 | |
| | | 92 | 22 | 20 | 3145 | | | |
| | | 25 | 60 : | x 37 | 85 x 37 | | | |

Figure 1.1: Matrix

| GC27 | Karyorelict Nuclear Code |
|------|------------------------------------|
| GC28 | Condylostoma Nuclear Code |
| GC29 | Mesodinium Nuclear Code |
| GC30 | Peritrich Nuclear Code |
| GC31 | Blastocrithidia Nuclear Code |
| GC32 | Balanophoraceae Plastid Code |
| GC33 | Cephalodiscidae Mitochondrial Code |
| GC34 | Enterosoma Code |
| GC35 | Peptacetobacter Code |
| GC36 | Anaerococcus and Onthovivens Code |
| GC37 | Absconditabacterales Code |
| | |

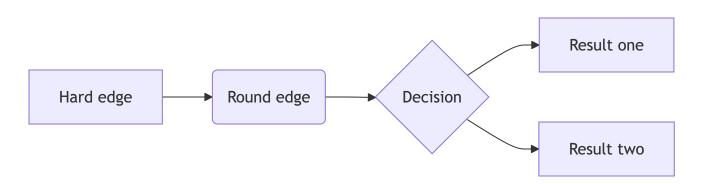
| Codon | Amino Acids | Nucleon BL | Nucleon SC | Table 1.3: Codon assignments for the Nucleon Total Standard Code, contracted by the |
|---------------------|------------------|------------|------------|---|
| \overline{AA} | N, K | 74 | 58 | 132 ^{first} two nucleotide bases. |
| AC | $\mathbf{T}^{'}$ | 74 | 45 | 119 |
| \overline{AG} | S, R | 74 | 31 | 105 |
| AT | I, M | 74 | 57 | 131 |
| CA | H, Q | 74 | 81 | 155 |
| CC | P | 74 | 41 | 115 |
| CG | \mathbf{R} | 74 | 100 | 174 |
| CT | L | 74 | 57 | 131 |
| GA | D, E | 74 | 59 | 133 |
| GC | A | 74 | 15 | 89 |
| GG | G | 74 | 1 | 75 |
| GT | V | 74 | 43 | 117 |
| TA | Y, * | 74 | 107 | 181 |
| TC | S | 74 | 31 | 105 |
| TG | C, *, W | 74 | 47 | 121 |
| TT | F L | 74 | 91 | 165 |

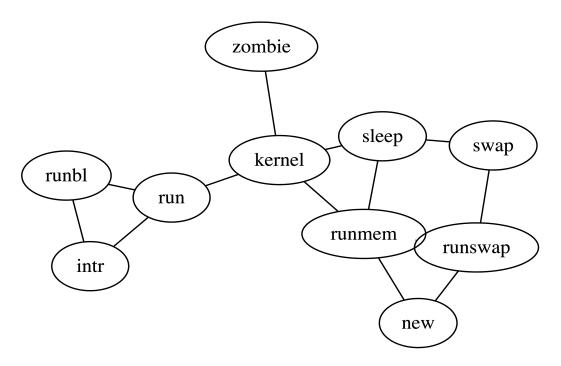
1.4 Running Code

When you click the **Render** button a document will be generated that includes both content and the output of embedded code. You can embed code like this:

1.5 ASCII Table

| | 0 | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 | A | В | С | D | Е | F |
|--------------|---|-------|--------------|--------------|----------------------|--------------|--------------------------|--------------|------------|--------------|---------------|---------------|---------------|---|----------|-----------------|
| 0 | | | | | | | | • | | | | | | 7 | | |
| 1 | | | | !! | \P | § | | | \uparrow | \downarrow | \rightarrow | \leftarrow | | | | |
| 2 | | ! | " | # | \$ | % | & | , | (|) | * | + | , | - | | / |
| 3 | 0 | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 | : | ; | < | = | > | ? |
| 4 | @ | A | В | \mathbf{C} | D | \mathbf{E} | F | G | Η | I | J | K | \mathbf{L} | Μ | Ν | Ο |
| 5 | Ρ | Q | \mathbf{R} | S | Τ | U | V | W | X | Y | \mathbf{Z} | [| \ |] | ^ | _ |
| 6 | 4 | a | b | \mathbf{c} | d | e | f | g | h | i | j | k | l | m | n | О |
| 7 | p | q | r | \mathbf{S} | \mathbf{t} | u | v | w | X | у | \mathbf{Z} | { | | } | ~ | |
| 8 | Ç | ü | é | â | ä | à | $ {a}$ | ç | ê | ë | è | ï | î | ì | Ä | Å |
| 9 | É | æ | Æ | ô | ö | ò | û | ù | ÿ | Ö | Ü | ¢ | £ | ¥ | | f |
| Α | á | í | ó | ú | $\tilde{\mathrm{n}}$ | Ñ | $\underline{\mathbf{a}}$ | \mathbf{O} | į | | \neg | $\frac{1}{2}$ | $\frac{1}{4}$ | i | « | >> |
| В | | | | | | | | | | | | | | | | |
| \mathbf{C} | | | | | | | | | | | | | | | | |
| D | | | | | | | | | | | | | | | | |
| \mathbf{E} | | ß | Γ | | \sum | | μ | | Φ | Θ | Ω | | ∞ | | | |
| F | | \pm | | | | | ÷ | | 0 | | | $\sqrt{}$ | | 2 | | |





A full width figure.

Reach and engagement of campaigns



Low Reach

High Reach

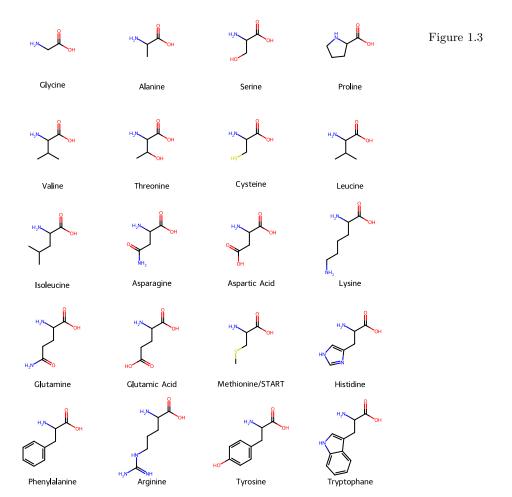
$$\begin{bmatrix} G & A \\ C & T \end{bmatrix} \otimes \begin{bmatrix} G & A \\ C & T \end{bmatrix} = \begin{bmatrix} GG & GA & AG & AA \\ GC & GT & AC & AT \\ CG & CA & TG & TA \\ CC & CT & TC & TT \end{bmatrix}$$

1.6 2D Molecule

As outlined by Crick and Orgel (1973), pp. 16-18, we find this below. Also by referencing this, we end up there¹, as outlined in Figure 1.2. Also refer to Figure 1.1 for more details.

This is a span that has the class Crick, F. H. C., and L. E. Orgel. 1973. aside which places it in the marging): without a tentumbero.1016/00 19-1035(73)90110-3.

¹ Here is the footnote.



- 1.6.1 References
- 1.6.2 Supplementary Material

A hello world

B Appendix 1

here we go...

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