## Supplementary Material for: On the origin of degeneracy in the genetic code

Diego Luis Gonzalez Simone Giannerini Rodolfo Rosa

September 4, 2019

## A Chemical nucleotide transformations and symmetric groups

The genetic alphabet

$$\mathcal{B} := \{U(T), C, A, G\}$$

is composed of four letter or nucleotides: Uracil (Thymine), Cytosine, Adenine, and Guanine. The set of all possible bijective transformations over  $\mathcal{B}$  has 24 elements and is defined as

$$S_{\mathcal{B}} = \{\pi : \mathcal{B} \to \mathcal{B} \mid \pi \text{ is bijective}\}$$

 $S_{\mathcal{B}}$ , together with the usual group operation given by composition as functions (denoted by  $\circ$ ), forms a symmetric group on  $\mathcal{B}$ . The group  $S_{\mathcal{B}}$  is isomorphic to the symmetric group  $S_4$  on four elements (see ? for more details). Hence, we write  $\pi : (A, T, C, G) \to (G, A, T, C)$  if  $\pi$  satisfies  $\pi(A) = G, \pi(T) = A, \pi(C) =$ T, and  $\pi(G) = C$ . Bijective mappings  $\pi : \mathcal{B} \to \mathcal{B}$  can be applied componentwise to  $x \in \mathcal{B}^3$ , the set of codons, and thus induce a bijective map  $\mathcal{B}^3 \to \mathcal{B}^3$  which we denote also by  $\pi$ .

As shown in ?, there are 4 bijective transformations that are invariant with respect to the chemical characters of the nucleotides. These are: Identity:

$$I: (A, T, C, G) \to (A, T, C, G);$$

Strong/Weak (SW) or complementary transformation:

$$SW: (A, T, C, G) \rightarrow (T, A, G, C);$$

Pyrimidine/Purine (YR) or parity transformation:

 $\mathrm{YR}: (A, T, C, G) \to (G, C, T, A);$ 

and Keto/Amino (KM) or Rumer's transformation:

$$\mathrm{KM}: (A, T, C, G) \to (C, G, A, T).$$

In turn, these 4 transformations, together with the operation  $\circ$  defined above, form a group on  $\mathcal{B}^3$ , which is isomorphic to the Klein V group and whose composition table is the following:

0	I	SW	$\mathbf{K}\mathbf{M}$	YR
I	Ι	SW	KM	YR
SW	SW	Ι	$\mathbf{YR}$	$\mathbf{K}\mathbf{M}$
KM	$\mathbf{K}\mathbf{M}$	$\mathbf{YR}$	Ι	SW
YR	YR	$\mathbf{K}\mathbf{M}$	SW	Ι

For instance, given the codon ACT, we can apply the two transformations SW and KM: SW(ACT) = TGA, KM(TGA) = GTC. Hence,  $(SW \circ KM)(ACT) = GTC = YR(ACT)$ .

## B Jukes' code in a random world of tetracodons.

Jukes' code has 16 amino acids and 1 stop signal. Hence, we assume we have 17 objects/amino acids to be assigned randomly to  $4^4 = 256$  tetracodons partitioned into 32 symmetric codons (with degeneracy 2) and 224 asymmetric codons (with degeneracy 4). Under this hypothesis, we compute the probability of observing a given degeneracy distribution. This corresponds to the hypergeometric probability of extracting without replacement k symmetric codons out of 17 objects, where the probability of extracting a symmetric codon is p = 32/256 = 1/8. We obtain:

$$P(X = k) = \frac{\left(\frac{32}{k}\right) \left(\frac{224}{17-k}\right)}{\left(\frac{256}{17}\right)}.$$
(1)



Figure 1: Hypergeometric probability function associated to choosing at random k symmetric elements/codons from 17. The mode of the distribution k = 2corresponds to the degeneracy of Jukes' code and indicates that such code has maximal probability of being chosen if we assume a random assignation of tetracodons to amino acids.

The hypergeometric probability function above is presented in Figure 1. Clearly, the mode k = 2 corresponds to the degeneracy distribution of Jukes' code, that is, 2 elements of degeneracy 2 and 15 elements with degeneracy 4.