Dubai, 14.02.2022 **Revisiting the number of substitutions in Tandem Repeats**

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ABSTRACT

In this paper, the expected number of substitutions occurring in tandem repeats is investigated.

Single point substitutions happen in the duplication history of tandem repeats. These single substitutions produce a set of variants that we observe at the current time. The number of substitutions that occurred in the past is unknown, a lower bound on the number of substitutions is the length of the Steiner tree of the variants, which can be approximated by the length of the minimum spanning tree of the variants' graphs. However, there might be some parallel substitutions that happened in the duplication history where an existing variant is created again. In this section, we calculate the likelihood P(k,i) that i observed substitutions are the result of $k \ge i$ substitutions on a motif of size n. There are 3n possible substitutions, where n is the length of the motif.

A substitution can either be a new substitution or it can be parallel (it duplicates an existing substitution); in the second case the number of substitutions does not increase. If we observe i substitutions after k substitutions have occurred, then the $k-\frac{th}{substitution}$ produced either a new substitution (with probability $\frac{1}{3n-(i-1)} \frac{3n}{s}$), or reproduced an existing substitution (a parallel substitution with probability $\frac{1}{3n}$). Thus the probability P(k,i) of observing i substitutions after k>0 substitutions can be calculated using the recursive formula

 $P(k,i)=P(k-1,i-1)\times \left(3n-(i-1)\right) (3n)+P(k-1,i)\times \left(3n-(i-1)\right) (3n)+P(k-1,i)\times \left(3n-(i-1)\right) (3n)+P(k-1,i-1)\times \left(3n-(i-1)\right) (3n)+P(k-1,i-1)\times \left(3n-(i-1)\right) (3n)+P(k-1,i-1)\times \left(3n-(i-1)\right) (3n)+P(k-1,i-1)\times \left(3n-(i-1)\right) (3n)+P(k-1,i-1)\times \left(3n-(i-1)\right) (3n)+P(k-1,i)\times \left(3n-(i-1)\right) (3n-(i-1)) (3n-(i$

for k>0, i>0, \$

with initial values

 $P(0,1)=1, P(0,i)=0, P(k,0)=0 \det\{ \text{ for } \} i \le 1.$

Figure 2.2 plots P(k,i) for n=11 and i=19, as is the case for the tandem motifs in <u>NZ1 [1]</u>. This suggest that the number of parallel substitutions in <u>NZ1</u> is more likely to be in the range 5--13.

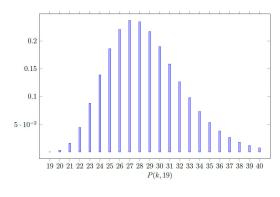


Figure 2.2: The likelihood distribution of the number of substitutions to produce 19 variants of a motif of length 11 bp.

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