

The Genetic Code and the I Ching

Jan Krämer

Theoretical Computer Science Dept. of Computer Science
RWTH Aachen University

Helping Donald Knuth Seminar 2005

Outline

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Background

- The I Ching
- The Genetic Code
- Question 7.2.1.7-2(a)
- Q&A

2

The Exercise

- The Question
- Algebraical Approach
- Empirical Approach
- Q&A

3

Recomendation

- My Recomendation
- Final Recomendation

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I Ching, the Bible of Changes

I Ching

- One of the five classics of confucian wisdom
- Contains 64 chapters
- Each chapter is marked by an hexagram
(for example ☰☷)

I Ching, the Bible of Changes

King Wen's ordering

- The original ordering of the chapters in the *I Ching*
- This implies a certain ordering of the hexagrams
- (The *I Ching* has been ascribed to King Wen c. 1100 B.C.)

Example

1: ☰, 2: ☷, 3: ☱, 4: ☲, ..., 63: ☴, 64: ☵

DNA and Proteins

DNA for Computer Scientists

- DNA sequence strings over 4 letter alphabet $\{T, C, A, G\}$ (nucleotides)
- Triplet of nucleotides form a codon
- Each codon maps to an amino acid $\{A, C, D, E, F, G, H, I, K, L, M, N, P, Q, R, S, T, V, W, Y\}$
- Proteins are strings of amino acids

Mapping DNA to proteins

Algorithm

- Chop DNA sequence into codons (x, y, z)
- Map $T \rightarrow 1, C \rightarrow 2, A \rightarrow 3, G \rightarrow 4$
- Lookup amino acid in matrices below
 $(x, y, z) \rightarrow$ matrix x , row y , column z
- Keep going until you hit separator '-'

Amino Acid Lookup Matrix

$\begin{pmatrix} F & S & Y & C \\ F & S & Y & C \\ L & S & - & - \\ L & S & - & W \end{pmatrix}$	$\begin{pmatrix} L & P & H & R \\ L & P & H & R \\ L & P & Q & R \\ L & P & Q & R \end{pmatrix}$	$\begin{pmatrix} I & T & N & S \\ I & T & N & S \\ I & T & K & R \\ M & T & K & R \end{pmatrix}$	$\begin{pmatrix} V & A & D & G \\ V & A & D & G \\ V & A & E & G \\ V & A & E & G \end{pmatrix}$
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Mapping Hexagrams to Amino Acids

Question

Show that there is a simple way to map each codon into a hexagram of the I Ching, with the property that the 21 possible outcomes $\{A, C, D, \dots, W, Y, -\}$ correspond to 21 consecutive hexagrams of the King Wen ordering.

Question 7.2.1.7-2(a)

Mapping Hexagrams to Amino Acids

Solution

- Represent (T, C, A, G) by
 - $\left(\begin{smallmatrix} \vdots & \vdots & \vdots & \vdots \\ \hline \hline \end{smallmatrix}, \begin{smallmatrix} \vdots & \vdots & \vdots & \vdots \\ \hline \hline \end{smallmatrix}, \begin{smallmatrix} \vdots & \vdots & \vdots & \vdots \\ \hline \hline \end{smallmatrix}, \begin{smallmatrix} \vdots & \vdots & \vdots & \vdots \\ \hline \hline \end{smallmatrix} \right)$ in first nucleotide
 - $\left(\begin{smallmatrix} \vdots & \vdots & \vdots & \vdots \\ \hline \hline \end{smallmatrix}, \begin{smallmatrix} \vdots & \vdots & \vdots & \vdots \\ \hline \hline \end{smallmatrix}, \begin{smallmatrix} \vdots & \vdots & \vdots & \vdots \\ \hline \hline \end{smallmatrix}, \begin{smallmatrix} \vdots & \vdots & \vdots & \vdots \\ \hline \hline \end{smallmatrix} \right)$ in second nucleotide
 - $\left(\begin{smallmatrix} \vdots & \vdots & \vdots & \vdots \\ \hline \hline \end{smallmatrix}, \begin{smallmatrix} \vdots & \vdots & \vdots & \vdots \\ \hline \hline \end{smallmatrix}, \begin{smallmatrix} \vdots & \vdots & \vdots & \vdots \\ \hline \hline \end{smallmatrix}, \begin{smallmatrix} \vdots & \vdots & \vdots & \vdots \\ \hline \hline \end{smallmatrix} \right)$ in third nucleotide

Question 7.2.1-7-2(a)

Mapping Hexagrams to Amino Acids

Solution

- Represent (*T*, *C*, *A*, *G*) by
 - $\left(\begin{smallmatrix} \vdots & \vdots & \vdots & \vdots \\ \hline \text{---} & \text{---} & \text{---} & \text{---} \end{smallmatrix} \right)$ in first nucleotide
 - $\left(\begin{smallmatrix} \vdots & \vdots & \vdots & \vdots \\ \hline \text{---} & \text{---} & \text{---} & \text{---} \end{smallmatrix} \right)$ in second nucleotide
 - $\left(\begin{smallmatrix} \vdots & \vdots & \vdots & \vdots \\ \hline \text{---} & \text{---} & \text{---} & \text{---} \end{smallmatrix} \right)$ in third nucleotide

Example

- $\begin{smallmatrix} \text{---} \\ \text{---} \\ \text{---} \\ \text{---} \\ \text{---} \\ \text{---} \end{smallmatrix} = \begin{smallmatrix} \vdots & \vdots & \vdots & \vdots \\ \hline \text{---} & \text{---} & \text{---} & \text{---} \end{smallmatrix} + \begin{smallmatrix} \vdots & \vdots & \vdots & \vdots \\ \hline \text{---} & \text{---} & \text{---} & \text{---} \end{smallmatrix} + \begin{smallmatrix} \vdots & \vdots & \vdots & \vdots \\ \hline \text{---} & \text{---} & \text{---} & \text{---} \end{smallmatrix}$
- $\Leftrightarrow$ codon *TTC*
- $\Leftrightarrow$ amino acid *F*

Question 7.2.1.7-2(a)

Mapping Hexagrams to Amino Acids

Solution

- Represent (T, C, A, G) by
 - $\left(\begin{smallmatrix} \vdots & \vdots & \vdots & \vdots \\ \hline \hline \end{smallmatrix}, \begin{smallmatrix} \vdots & \vdots & \vdots & \vdots \\ \hline \hline \end{smallmatrix}, \begin{smallmatrix} \vdots & \vdots & \vdots & \vdots \\ \hline \hline \end{smallmatrix}, \begin{smallmatrix} \vdots & \vdots & \vdots & \vdots \\ \hline \hline \end{smallmatrix} \right)$ in first nucleotide
 - $\left(\begin{smallmatrix} \vdots & \vdots & \vdots & \vdots \\ \hline \hline \end{smallmatrix}, \begin{smallmatrix} \vdots & \vdots & \vdots & \vdots \\ \hline \hline \end{smallmatrix}, \begin{smallmatrix} \vdots & \vdots & \vdots & \vdots \\ \hline \hline \end{smallmatrix}, \begin{smallmatrix} \vdots & \vdots & \vdots & \vdots \\ \hline \hline \end{smallmatrix} \right)$ in second nucleotide
 - $\left(\begin{smallmatrix} \vdots & \vdots & \vdots & \vdots \\ \hline \hline \end{smallmatrix}, \begin{smallmatrix} \vdots & \vdots & \vdots & \vdots \\ \hline \hline \end{smallmatrix}, \begin{smallmatrix} \vdots & \vdots & \vdots & \vdots \\ \hline \hline \end{smallmatrix}, \begin{smallmatrix} \vdots & \vdots & \vdots & \vdots \\ \hline \hline \end{smallmatrix} \right)$ in third nucleotide
- Hexagrams 34 - 54 mapped to 21 possible amino acids

Q&A Session 1

Q&A

Any Questions?

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The Question

Question 7.2.1.7-2(b)

Question

Is that a sensational discovery?

The Question

Question 7.2.1.7-2(b)

Question

Is that a sensational discovery?

Knuth's Answer

No

Approach

Note

- The mapping between hexagramms and amino acids, can be reduced to a serialization of the amino acid matrices.
- The number of the hexagram is the position of the mapped amino acid in the string

Approach

Note

- The mapping between hexagramms and amino acids, can be reduced to a serialization of the amino acid matrices.
- The number of the hexagram is the position of the mapped amino acid in the string

Question to Solve

How many permutation of the aminoacid matrices exists that contain a run of 21 distinct consecutive elements.

Number of Total Possible Permutations

Knuth States

$$\binom{64}{6,6,6,4,4,4,4,4,3,3,2,2,2,2,2,2,2,2,1,1} \approx 2.3 \times 10^{69}$$

Where Does That Formula Come From?

Multiset: $\{n_1 * x_1, \dots, n_r * x_r\}$, $n_1 + \dots + n_r = n$

$\binom{n}{n_1, \dots, n_r}$: multinomial coefficient

You draw the positions for:

- The k_1 elements of type x_1
- ...
- The k_r elements of type x_r

$$\binom{n}{n_1, \dots, n_r} = \binom{n}{n_1} \binom{n-n_1}{n_2} \dots \binom{n-n_1-\dots-n_{r-1}}{n_r} = \frac{n!}{n_1! \dots n_r!}$$

Number of Total Possible Permutations

Knuth States

$$\binom{64}{6,6,6,4,4,4,4,4,3,3,2,2,2,2,2,2,2,2,1,1} \approx 2.3 \times 10^{69}$$

Exact Solution

$$\binom{64}{6,6,6,4,4,4,4,4,3,3,2,2,2,2,2,2,2,2,1,1} =$$

$$2316278295874198383354290667060208126761409719904245070233600000000000 \approx 2.3 \times 10^{69}$$

Algebraical Approach

Multiset Permutations

with Run of r Distinct Elements

—

Explanations

Principle of inclusion and exclusion:

Permutations with 21 consecutive elements – multiples

Definitions

Multiset $= \{(n_1 + 1) * x_1, (n_2 + 1) * x_2, \dots, (n_r + 1) * x_r\}$

$$n = n_1 + n_2 + \dots + n_r$$

Algebraical Approach

Multiset Permutations

with Run of r Distinct Elements

$$r! -$$

Explanations

Number of permutations of the 21 consecutive elements

Definitions

$$\text{Multiset} = \{(n_1 + 1) * x_1, (n_2 + 1) * x_2, \dots, (n_r + 1) * x_r\}$$

$$n = n_1 + n_2 + \dots + n_r$$

Algebraical Approach

Multiset Permutations with Run of r Distinct Elements

$$\binom{n}{n_1, \dots, n_r} r! -$$

Explanations

Number of permutations of the rest of the elements

Definitions

Multiset $= \{(n_1 + 1) * x_1, (n_2 + 1) * x_2, \dots, (n_r + 1) * x_r\}$

$$n = n_1 + n_2 + \dots + n_r$$

Algebraical Approach

Multiset Permutations

with Run of r Distinct Elements

$$(n+1) \binom{n}{n_1, \dots, n_r} r! -$$

Explanations

Number of places where the consecutive elements can be inserted

Definitions

Multiset $= \{(n_1 + 1) * x_1, (n_2 + 1) * x_2, \dots, (n_r + 1) * x_r\}$

$$n = n_1 + n_2 + \dots + n_r$$

Algebraical Approach

Multiset Permutations with Run of r Distinct Elements

$$(n+1) \binom{n}{n_1, \dots, n_r} r! -$$

Where Do the Multiples Come From?

Select x, y, v, w, α so that $|xvy| = n$, $w = \sigma(v)$ and $|v\alpha| = r$ distinct elements.
 $xv\alpha wy$ can be created by:

- αw in xvy , or
- $v\alpha$ in xwy

Algebraical Approach

Multiset Permutations with Run of r Distinct Elements

$$(n+1) \binom{n}{n_1, \dots, n_r} r! -$$

A Special Case Makes Our Life Easier

$xv\alpha w$ of length $n+r$ has been multiply inserted

$w = \sigma(v)$ but both $v\alpha$ and αw must contain all possible elements exactly once

$\Rightarrow \alpha$ must contain all elements that only occur once

Since we have such an element (w) in the sequence of amino acids

$\Rightarrow$ This case is indeed the only of multiple insertions we must consider

Algebraical Approach

Multiset Permutations

with Run of r Distinct Elements

$$(n+1) \binom{n}{n_1, \dots, n_r} r! - \sum_{k=1}^r$$

Explanations

Vary over $|v|$ in $xv\alpha wy$

Definitions

Multiset $= \{(n_1 + 1) * x_1, (n_2 + 1) * x_2, \dots, (n_r + 1) * x_r\}$

$n = n_1 + n_2 + \dots + n_r$

$n_r = 0$

Algebraical Approach

Multiset Permutations

with Run of r Distinct Elements

$$(n+1) \binom{n}{n_1, \dots, n_r} r! - \sum_{k=1}^r \binom{n-k}{n_1, \dots, n_r}$$

Explanations

Generate number of possible words xy for $xv_\alpha wy$ with $|v| = k$

Definitions

Multiset $= \{(n_1 + 1) * x_1, (n_2 + 1) * x_2, \dots, (n_r + 1) * x_r\}$

$n = n_1 + n_2 + \dots + n_r$

$n_r = 0$

Algebraical Approach

Multiset Permutations

with Run of r Distinct Elements

$$(n+1) \binom{n}{n_1, \dots, n_r} r! - \sum_{k=1}^r \sum_{\substack{0 \leq d_1, \dots, d_r \leq 1 \\ d_1 + \dots + d_r = k}} \binom{n-k}{n_1 - d_1, \dots, n_r - d_r}$$

Explanations

Generate number of possible words xy for $xv\alpha wy$ with $|v| = k$ but take into account the possible elements that can make up v

Definitions

Multiset = $\{(n_1 + 1) * x_1, (n_2 + 1) * x_2, \dots, (n_r + 1) * x_r\}$

$n = n_1 + n_2 + \dots + n_r$

$n_r = 0$

Algebraical Approach

Multiset Permutations with Run of r Distinct Elements

$$(n+1) \binom{n}{n_1, \dots, n_r} r! - \sum_{k=1}^r$$

$$\sum_{\substack{0 \leq d_1, \dots, d_r \leq 1 \\ d_1 + \dots + d_r = k}} \binom{n-k}{n_1 - d_1, \dots, n_r - d_r}$$

Explanations

This defines not only the number of xys but the specific elements contained in v , w and α

Definitions

Multiset = $\{(n_1 + 1) * x_1, (n_2 + 1) * x_2, \dots, (n_r + 1) * x_r\}$

$n = n_1 + n_2 + \dots + n_r$

$n_r = 0$

Algebraical Approach

Multiset Permutations with Run of r Distinct Elements

$$(n+1) \binom{n}{n_1, \dots, n_r} r! - \sum_{k=1}^r (n+1-k)$$

$$\sum_{\substack{0 \leq d_1, \dots, d_r \leq 1 \\ d_1 + \dots + d_r = k}} \binom{n-k}{n_1 - d_1, \dots, n_r - d_r}$$

Explanations

Number of places where we can split xy

Definitions

Multiset = $\{(n_1 + 1) * x_1, (n_2 + 1) * x_2, \dots, (n_r + 1) * x_r\}$

$n = n_1 + n_2 + \dots + n_r$

$n_r = 0$

Algebraical Approach

Multiset Permutations

with Run of r Distinct Elements

$$(n+1) \binom{n}{n_1, \dots, n_r} r! - \sum_{k=1}^r (n+1-k) a_k \sum_{\substack{0 \leq d_1, \dots, d_r \leq 1 \\ d_1 + \dots + d_r = k}} \binom{n-k}{n_1 - d_1, \dots, n_r - d_r}$$

Explanations

Number of possible permutation for v

(2134) is not an interesting permutation for v

as that would have been covered in $k = 3 \Rightarrow |v| = 3$

Definitions

Multiset = $\{(n_1 + 1) * x_1, (n_2 + 1) * x_2, \dots, (n_r + 1) * x_r\}$

$n = n_1 + n_2 + \dots + n_r$

$n_r = 0$

a_k = number of indecomposable permutations with k elements

Algebraical Approach

Multiset Permutations with Run of r Distinct Elements

$$(n+1) \binom{n}{n_1, \dots, n_r} r! - \sum_{k=1}^r (n+1-k) k! a_k \sum_{\substack{0 \leq d_1, \dots, d_r \leq 1 \\ d_1 + \dots + d_r = k}} \binom{n-k}{n_1 - d_1, \dots, n_r - d_r}$$

Explanations

Number of permutations of v we can insert as w

Definitions

Multiset = $\{(n_1 + 1) * x_1, (n_2 + 1) * x_2, \dots, (n_r + 1) * x_r\}$

$n = n_1 + n_2 + \dots + n_r$

$n_r = 0$

a_k = number of indecomposable permutations with k elements

Algebraical Approach

Multiset Permutations with Run of r Distinct Elements

$$(n+1) \binom{n}{n_1, \dots, n_r} r! - \sum_{k=1}^r (n+1-k) k! (r-k)! a_k \sum_{\substack{0 \leq d_1, \dots, d_r \leq 1 \\ d_1 + \dots + d_r = k}} \binom{n-k}{n_1 - d_1, \dots, n_r - d_r}$$

Explanations

Number of permutations of α we can insert

Definitions

Multiset $= \{(n_1 + 1) * x_1, (n_2 + 1) * x_2, \dots, (n_r + 1) * x_r\}$

$n = n_1 + n_2 + \dots + n_r$

$n_r = 0$

a_k = number of indecomposable permutations with k elements

Using the Formula

Knuth States

2402880402175789790003993681964551328451668718750185553920000000 $\approx 2.4 \times 10^{63}$

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2402880402175789790003993681964551328451668718750185553920000000 $\approx 2.4 \times 10^{63}$

Result of Verification

2402880402175789790003993681964551328451668718750185553920000000 $\approx 2.4 \times 10^{63}$

Bringing it Together

It would seem that only one of every million permutations contains a run of 21 distinctive elements.

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But

There are $4!^3 \binom{6}{2,2,2} = 1244160$ (Verified) ways to represent codons as in part (a) and effectively change the ordering of the hexagrams.

Bringing it Together

It would seem that only one of every million permutations contains a run of 21 distinctive elements.

But

There are $4!^3 \binom{6}{2,2,2} = 1244160$ (Verified) ways to represent codons as in part (a) and effectively change the ordering of the hexagrams.

Thus

The one in a million chance cannot be seen as a proof that the authors of the I Ching must have foreseen the Genetic code.

Empirical Test

Knuth States

About 31% of all permutations turn out to have a suitable codon mapping.

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Knuth States

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Method of Verification

- I developed a C++ application that
 - Randomly generates permutation of hexagrams
- Searches for codon mappings

Empirical Test

Knuth States

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Result of Verification

After 1100 tested permutations, 347 were found to have mappings

Empirical Test

Knuth States

About 31% of all permutations turn out to have a suitable codon mapping.

Result of Verification

After 1100 tested permutations, 347 were found to have mappings

⇒ 31%

Q&A Session 2

Q&A

Any Questions?

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My Recommendation

My Recommendation

Knuths provided answer is correct, albeit extremely slow to retrace due to the long, underdocumented formula.

Final Recommendation

Let the Group Decide

What do you think?