Computational Molecular Biology

Lecture Seven: A Mathematical Diversion

More on longest common subsequences

Semester I, 2009-10

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Permutations

There are 24 permutations of the four letters ACGT. These are:

```
AGCT TGCA TACG TAGC
GACT GTCA ATCG ATGC
GCAT GCTA ACTG AGTC
CGAT CGTA CATG GATC
CAGT CTGA CTAG GTAC
ACGT TCGA TCAG TGAC
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```

There are 20!=2432902008176640000 permutations of the 20 amino acids. One such permutation is

 $\label{lem:conditional} Phe-Ser-Tyr-Sys-Leu-Trp-Pro-Hys-Arg-Glu-Ile-Thr-Asn-Met-Lys-Val-Ala-Asp-Gly-Glu$

Longest Common Subsequences of Permutations

The problem of finding longest common subsequences between permutations, say

$$\pi = \mathsf{ACGT}$$
 and $\pi' = \mathsf{CGAT}$,

can always be translated into a problem of finding longest common subsequences between permutations of integers, in this case

$$\tau = 1 \ 2 \ 3 \ 4 \ \text{and} \ \tau' = 2 \ 3 \ 1 \ 4.$$

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The first letter of π' is the 2nd letter of π , the second leter of π' is the 3rd letter of π , the thrid letter of π' is the 1st letter of π , and the fourth letter of π' is the 4th letter of π .

Longest Increasing Subsequence of a Permutation

The problem of finding longest common subsequences between the integer permutations

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is equivalent to the problem of finding a longest increasing subsequence of $\tau^\prime.$

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The subsequence 3 1 4 is not increasing.

Problem

Find the length of a longest common subsequence between the permutations

 $\pi = \mathsf{Phe}\text{-}\mathsf{Ser}\text{-}\mathsf{Tyr}\text{-}\mathsf{Sys}\text{-}\mathsf{Leu}\text{-}\mathsf{Trp}\text{-}\mathsf{Pro}\text{-}\mathsf{Hys}$

and

 $\pi' = \mathsf{Leu}\text{-}\mathsf{Phe}\text{-}\mathsf{Trp}\text{-}\mathsf{Ser}\text{-}\mathsf{Tyr}\text{-}\mathsf{Pro}\text{-}\mathsf{Sys}\text{-}\mathsf{Hys}$

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Solution

Just find a longest increasing subsequence of

$$\tau'$$
=5 1 6 2 3 7 4 8

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Solution

Just find a longest increasing subsequence of

$$\tau'=51623748$$

A longest increasing subsequence is $1\ 2\ 3\ 7\ 8$. So the length of a LCS is 5.



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In order to explain this KRS algorithm we need to know about partitions and Young tableaux.

Partitions

A partition of a positive integer n is a sequence of integers

$$\lambda_1 \ge \lambda_2 \ge \cdots \ge \lambda_I$$

with

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We write

$$(\lambda_1, \lambda_2, \dots, \lambda_l) \vdash n.$$

Partitions of n = 4

$$(4) \vdash 4$$

$$(3,1) \vdash 4$$

$$(2,2)\vdash 4$$

$$(2,1,1)\vdash 4$$

$$(1,1,1,1) \vdash 4$$

Shapes of partitions

The shape of $\lambda = (\lambda_1, \lambda_2, \cdots, \lambda_l) \vdash n$ is an array of l left-justified rows with each row i containing λ_i cells (or boxes).

Shapes of partitions of n = 4

$$(3,1) \vdash 4$$

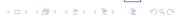






 $(2,1,1)\vdash 4$

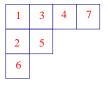




A Young tableau of shape $\lambda \vdash n$ is the shape with the numbers $1, 2, \cdots, n$ entered in the cells.

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If each row is increasing from left to right, and each column is increasing from top to bottom, then the Young tableau is said to be a standard Young tableau.

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The above example is a standard Yound tableau.