

Week 8, LEC02 - November 9th  
Discrete Mathematics, Fall 2018

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# CSCA67 - Lecture Notes

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Current Instructor: Dr. Richard Pancer



## Instructors:

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| <b>Office Hours:</b> | Monday 12:10 - 1:30                           | Monday 11:10 - 12:30      |
|                      | Wednesday 1:10 - 2:00                         | Friday 1:30 - 3:00        |
|                      | Friday 1:10 - 2:00 (will change after week 6) |                           |

$S(n)$  : Given  $n$  lines in the plane such that no two are parallel, and no three intersect at a single point, we have  $\binom{n}{3}$  Triangles.

Claim:  $\forall n \in \mathbb{N}, n \geq 3, S(n)$

Base  $n = 3$ , by observation (inspection), given three lines only one triangle can be formed.

$\binom{3}{3} = 1, S(n)$  for  $n = 3$ .

Inductive Hypothesis:  $k \in \mathbb{N}, k \geq 3, S(k)$

Inductive Step:  $S(k) \rightarrow S(k+1)$

With  $k$  lines in the plane, we have  $\binom{k}{3}$  triangles, by Inductive Hypothesis.

When the  $(k+1)^{st}$  line is added, it must cross every existing pair of lines in the plane.

When this happens, a new triangle is formed.

we have  $\binom{k}{2}$  ways of selecting 2 lines from  $k$  lines.

$\therefore$  Total number of Triangles after the  $(k+1)^{st}$  line is added is  $\binom{k}{3}(\text{IH}) + \binom{k}{2}(\text{New triangle from } (k+1)^{st} \text{ line})$ .  
 $= \frac{k!}{(k-3)!3!} + \frac{k!}{(k-2)!2!} = \frac{(k+1)!}{(k+1-3)!3!} = \binom{k+1}{3} \cdot S(k+1)$

Conclusion: Since  $k$  is arbitrary,  $\forall n \in \mathbb{N}, n \geq 3, S(n)$

This question will re-appear on assignment 2!

## 1 Counting with Repetitions

The genetic code of an organism stored in DNA molecules consist of 4 nucleotides:

Adenine, Cytosine, Guanine and Thymine

- It is possible to sequece short strings of molecules.
- One way to sequence the nucleotides of a longer string of DNA is to split the string into shorter sequences.
- A C-enzyme will split a DNA-sequence at each C. This means that each fragment will end at a C except possibly the last fragment.
- Similarly for A-enzymes, G-enzymes and T-enzymes.
- If the original nucleotide is split on each of C, A, G and T then it can be sequenced as it is most likely a unique sequence that can be constructed by each of the four sets of fragments.

**Example.** Given a 20-nucleotide string split at the Cs, one might have the fragments:

AC, AC, AAATC, C, C, C, TATA, TGGC

**Q.** How many different 20-nucleotide strings could have given rise to the above set of fragments? In other words, how many different arrangements are there of these fragments?

**A.**