

## **B2- Mathematics**

**B-MAT-150** 

## 106bombyx

Bombyx booming bylaw





## Bombyx

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binary name: : 106bombyx repository name: : 106bombyx repository rights: : ramassage-tek

language: : C, C++, perl 5, python 3.4, ruby 2.1, php 5.6, bash 4

group size: : 1-2

compilation (when necessary): : via Makefile, includig re, clean and fclean rules



• Your repository shall contain the totality of your source files, but no useless file (binary, test files, obj files,...).

• All the bonus files (including a specific Makefile) shall be in a directory named bonus.

• Error messages have to be written on the error output, and the program should then exit with the 84 error code (O if there is no error).

In the 70's, chaos theory opened the way for a better understanding of some animal species. Butterflys for instance. Let's say... bombyx.

If a generation is crowded, chances are that the next generation will be crowded too, regarding the natural rules of reproduction. But resources may lack for this new generation, so it may not be able to develop. Therefore, the evolution of the number of bombyx from a generation to the next one is not trivial.

Let's call  $x_i$  the number of the  $i^{th}$  generation of butterflys. Here is a model for the evolution of  $x_i$ :

$$\begin{cases} x_1 = n & \text{where } n \text{ is the number of first generation individuals} \\ x_{i+1} = k.x_i.\frac{1000 - x_i}{1000}, & \text{for } i \geq 1, k \text{ being the } \textit{growth rate}, \text{ from 1 to 4}. \end{cases}$$

In order to study this evolution, you are asked to plot two things:

- the curve representing the number of individuals in relation to the generation (varying from 1 to 100)
- a synthetic scheme summing all the results for a given n; it consists in plotting every value of  $x_i$  (between two given bounds), in relation to k (k varying from 1 to 4 by 0.01 steps).

In both cases, your program shall print on the standard outuput the values to be entered into *gnuplot* to draw the graphes.

```
Terminal + X

~/B-MAT-150> ./106bombyx -h

USAGE

./106bombyx n [k | i0 i1]

DESCRIPTION

n number of first generation individuals
k growth rate from 1 to 4
i0 initial generation (included)
i1 final generation (included)
```







Your program output has to be strictly identical to the one below.

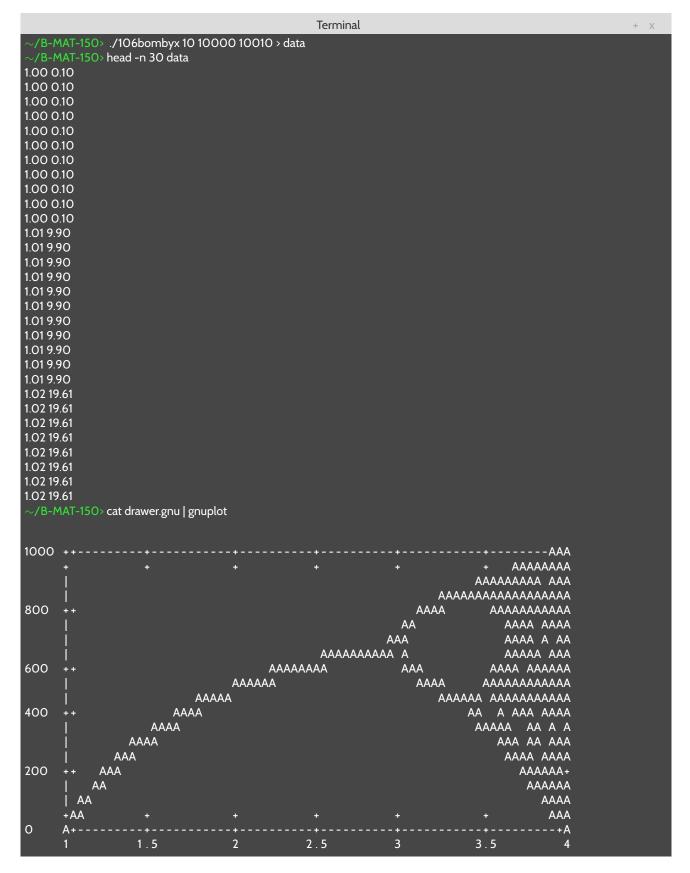
Terminal

+ X

~/B-MAT-150> cat drawer.gnu set terminal dumb set nokey plot "data"











Terminal ~/B-MAT-150> ./106bombyx 10 3.3 > data ~/B-MAT-150> head data 110.00 2 32.67 3 104.29 4 308.26 5 703.68 6 688.10 7 708.24 8 681.89 9 715.82 10 671.29  $\sim$ /B-MAT-150> tail data 91 823.60 92 479.43 93 823.60 94 479.43 95 823.60 96 479.43 97 823.60 98 479.43 99 823.60 100 479.43

