# A forest of sigmoidalistic monsters: Part 1: phenotypic description

## Plomion C<sup>1</sup>

**Abstract :** This paper describes a series of genetic monsters<sup>1</sup> created by crossing straight and sigmoidal maritime pine trees.

#### Introduction

This is the first report of such phenomenon in trees...to our best knowledge of course!!

#### **Materials and Methods**

Simple genetic manipulation....cannot say more...

#### Results

image speaks for itself.



<sup>&</sup>lt;sup>1</sup>: Bio-gecho, Landes pampa

<sup>&</sup>lt;sup>1</sup> no transgenics were produced in this experiment...so José don't cut my trees!!





**Figure 1:** Genetically manipulated forest (somewhere...in the Landes pampa). See the remarkable segregation obtained by this genetic manipulation

The basic statistics for stem straightness (ST, deviation from verticality) are indicated in supp file 1. Although skewed, the distribution of the trait clearly shows the presence of transgressive genotypes, so called "sigmoid phenotype". Data for two growth traits are given for comparison purpose. Note the extremely high CVp for ST! We are aware that such high value will probably make Dr. S. Delzon Jealous. Indeed compared to P50, ST presents a 10-fold increase in CVp making it a perfect trait for breeding for ornamental varieties in the year to come. Resistance to the pine wood nematode will have to be checked to evaluated the sustainability of the sigmoid forest. According to preliminary studies and feedback from users (Hugo Plomion, 6 yo) the ecological services provided by Sigmoid forest are expected to be high...Their impact on the forestry wood chain remains to be assessed.

#### Discussion

you wonder of course about the hydraulic properties of these trees...Part 2 (after the evaluation of the EFPA division....) will provide some answers to this key question....suspense...

This being said, we urge the reader to carefully check the following references (can be retrieved at http://herve.cochard.free.fr/) that will provide essential background to understand the next chapter of this story.

Cochard et al 1992, Cochard et al 1993, Cochard et 1994, Cochard et al 1995, Cochard et al 2013 (new insights in tree monsters: a comment on the paper by Plomion, JSPH 0002).

The double inversed signmoidal phenotype, also called in the corridor of the cavitron facility the double S phenotype, is inspiring and deserve more in depth analysis. Annabel Porté, a well known specialist of the double S phenotype will provide some clues about this issue in the next issue of the Journal.

#### Acknowledgements

Two non anonymous reviewers, A Porté and G Le Provost, for certifying the uniqueness of these trees...and for supporting tons of mosquitoes . Franck Zappa for kepping me awake at night.

#### References

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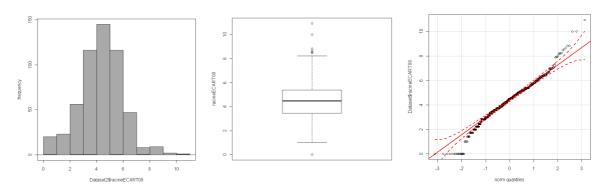
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Supp file 1:

### Stem straightness (Root-squared transformed data)



|                                    | CIRC10                    | HT10                        | ECART08                      | logECART08                  | rtsqECART08                  |
|------------------------------------|---------------------------|-----------------------------|------------------------------|-----------------------------|------------------------------|
| Kolmogorov-Smirnov:<br>D (p-value) | 1 (<2.2e <sup>-16</sup> ) | 1 (<2.2e <sup>-16</sup> )   | 0.95 (<2.2e <sup>-16</sup> ) | 0.7 (<2.2e <sup>-16</sup> ) | 0.9(<2.2e <sup>-16</sup> )   |
| Shapiro-Wilk:<br>W (p-value)       | 0.99 (0.0001)             | 0.96 (6.5e <sup>-06</sup> ) | 0.88 (<2.2e <sup>-16</sup> ) | $0.94(1.24e^{-13})$         | 0.98 (1.35e <sup>-07</sup> ) |

<sup>\*:</sup> p-value > 0.05: Ho accepted- the trait display normal distribution

### Classical statistics for three traits

#### Standard Deviation Traits Mean (M) (SD) CVp Min Max ECART08 14.9941956 0.68235402 21.97421 0 543 120 HT10 722.86654 98.9859463 0.1369353 370 1020 532 CIRC10 464.27941 103.782013 0.22353352 700 157 544

Pearson correlations for 3 traits (below diagonal) and associated p-values (above diagonal).

|         | Circ10 | ECART08 | HT.10  |
|---------|--------|---------|--------|
| CIRC10  |        | 0       | 0      |
| ECART08 | 0.18   |         | 0.2506 |
| HT.10   | 0.82   | 0.05    |        |

Correlation circle in the main plane (F1xF2) of the principal component analysis for 3 traits.

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