# Project 1: Genomics



## Cancer hallmarks orthologues in Arabidopsis thaliana suggests partial overlap and potentiality as a research model for the disease

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### Introduction

- Cancer is a prevalent disease worldwide.
- Cancer hallmarks were described as common traits for rational drug design for the disease.
- Although cancer study is important, animal model organism are expensive.

- Phenologs (phenotypes based on orthologs) has been poorly used to find alternatives to animal models.
- Plant models are cheap and they are not limited by informed consent or bioethics rules applied to animals.



- (Knijnenburg et al., 2015 + Kiefer et al, 2017)
- Categorization in hallmarks using GO terms.

6476 genes annotated using Entrez IDs

- Gene Ontology enrichment analysis using A. thaliana as target.
- Non-metric multidimensional scaling (NMDS).
- Bar plots of Jaccard index, and Szymkiewicz–Simpson coefficient.
- Visualization using ggplot2



- 1793 orthologues were found in A.thaliana while 63% genes were unique in humans.
- A partial overlap between orthologues suggested a possible potential of A. thaliana as a study model (>23% shared genes) for the following hallmarks:
  - 1. Deregulating Cellular Energetics



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