



Ciências
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da Universidade
de Lisboa

Experimental design and data analysis

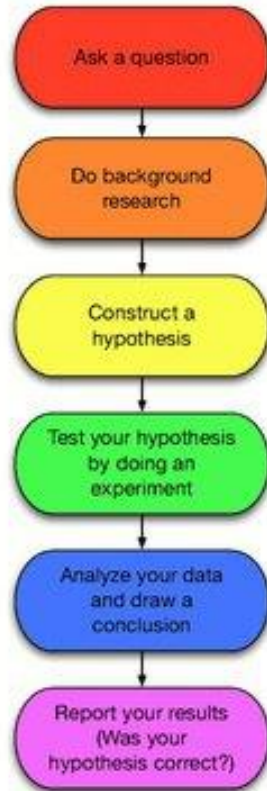
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Scientific Method

The Scientific Method



1. Ask a question: What is the role of a protein “Z” in plant innate immunity?

While asking your question you also need to decide an approach. The approach sets the theme of how you are going to address the question. Are you going for a top-down or bottom-up design?

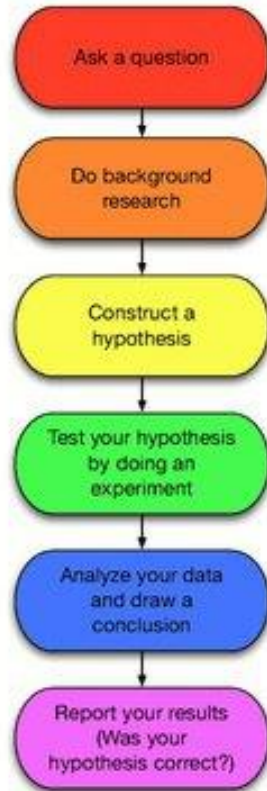
(it is always advisable to break down your question into multiple small parts, and answer them individually as you move ahead)

2. Do background research: Search for research and review articles on PubMed, Google Scholar etc using keywords like “plant innate immunity”, “proteins involved immunity”, “immunity regulation in plants”

3. Construct a hypothesis: protein “Z” is a membrane receptor or interacts with a membrane receptor and participates in pathogen molecules recognition.

Scientific Method

The Scientific Method



4. Test hypothesis: This step involves designing multiple experiments to address each part of your question. This includes from basic experiments like checking for the presence of “z” in plant membranes or complex experiments like protein-protein interaction. Include positive and negative controls to avoid any false positive or false negatives in your experiment.

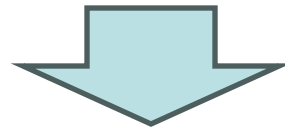
5. Data analysis: make use of statistical tools to make sense of the data you have obtained so far. Make relevant inferences, apply statistical tests and draw conclusions.

6. Results: report your conclusions in an understandable format which includes gel images, quantitative analysis, graphs etc. If “Z” shows a role in plant immunity, you can report you results in peer-reviewed journals, if not: either change the hypothesis, or check for errors in your analysis or try to report your findings as negative results.

Scientific Method

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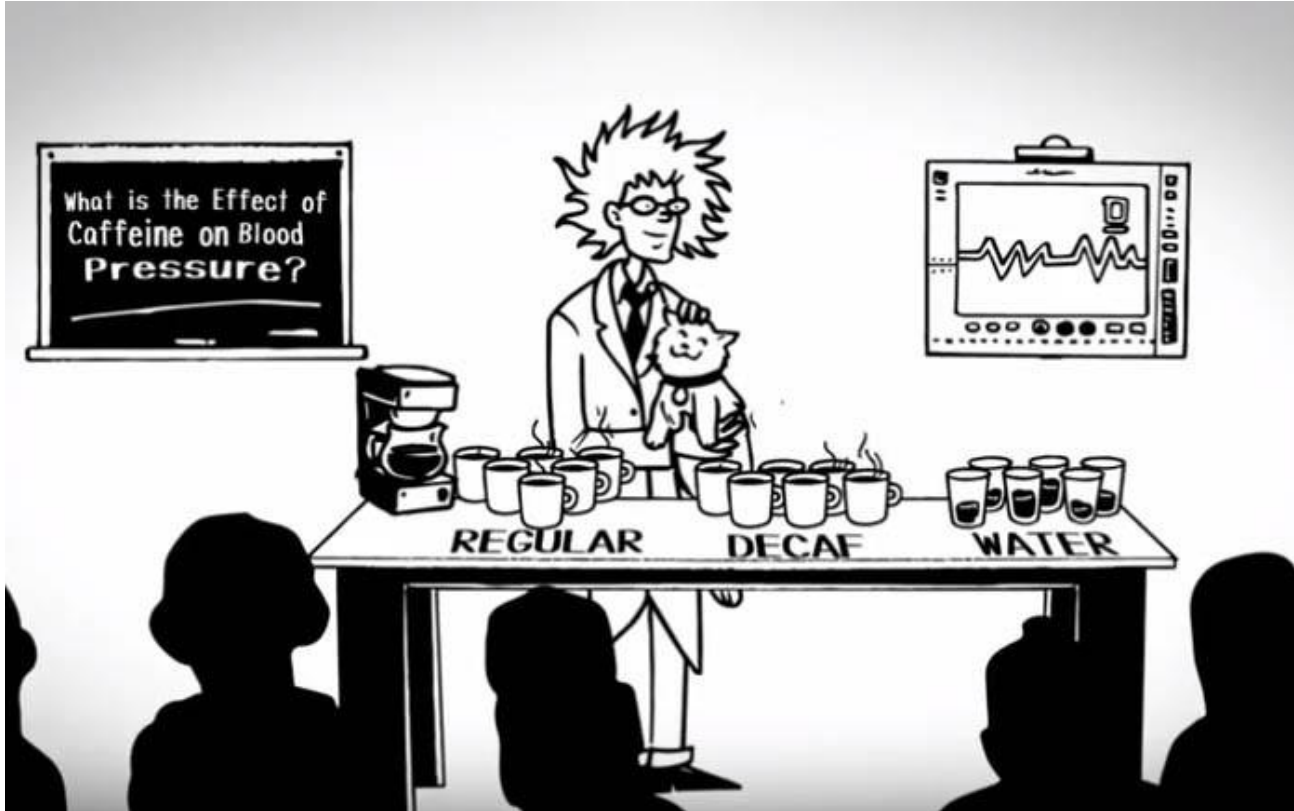


Experimental Design?



"There's a flaw in your experimental design.
All the mice are scorpions." AllPosters

Experimental design



Experimental design includes both:

- Strategies for organizing data collection
- Because of variability: Data analysis procedures matched to those data collection strategies
- We wouldn't need a science of experimental design:
 - If all units (for e.g. organisms) were identical
 - If all units responded identically to treatments
- We need experimental design to control variability so that treatment effects can be identified

Experimental design

- The idea of controlling variability through design has a long history. One of the first examples is the work in scruvy by Sir James Lind's (1747)

Without stating what method of allocation he used, Lind allocated two men to each of six different daily treatments for a period of fourteen days. The six treatments were:

- 1- 1.1 litres of cider;
- 2- twenty-five millilitres of elixir vitriol (dilute sulphuric acid);
- 3- 18 millilitres of vinegar three times throughout the day before meals;
- 4- half a pint of sea water;
- 5- two oranges and one lemon continued for six days only (when the supply was exhausted);
- 6- medicinal paste made up of garlic, mustard seed, dried radish root and gum myrrh.



Experimental design

“The most sudden and visible good effects were perceived from the use of oranges and lemons; one of those who had taken them being at the end of six days fit for duty ... The other was the best recovered of any in his condition; and being now deemed pretty well, was appointed nurse to the rest of the sick.”



Experimental design

- Later on.. Studies in crop variation (1921 – 1929) by statistician named Fisher was hired at Rothamsted agricultural station
- They had a lot of observational data on crop yields and hoped a statistician could analyze it to find effects of various treatments
- All he had to do was sort out the effects of confounding variables
- Fisher did regression analyses—lots of them—to study (and get rid of) the effects of confounders:
 - soil fertility gradients
 - drainage
 - effects of rainfall
 - effects of temperature and weather, etc.



Conclusion: The effects of confounders are typically larger than those of the systematic effects under study

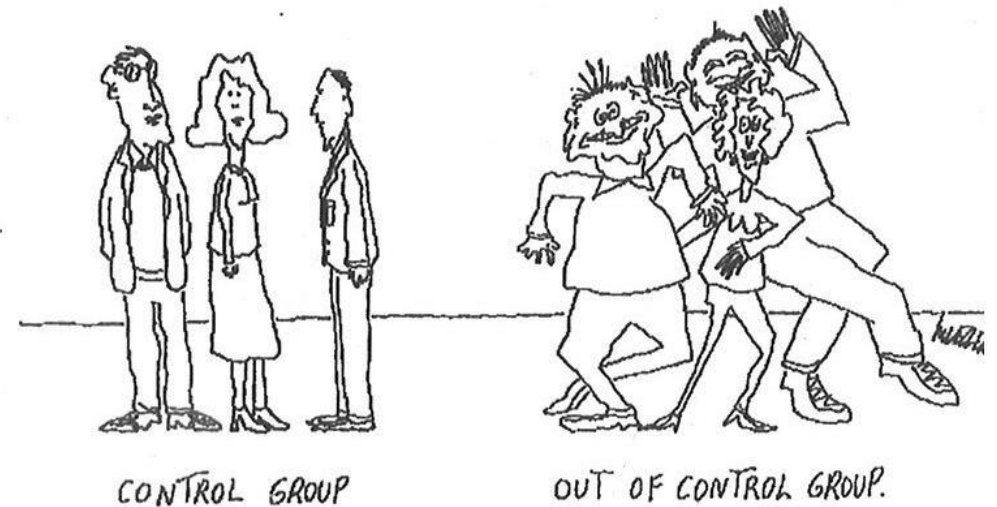
Experimental design

- **Fisher invents**
 - Basic principles of experimental design
 - Control of variation by randomization
 - Analysis of variance
- **Studies in Crop variation IV (1927)**
 - Fisher invents analysis of covariance to combine statistical control and control by randomization
- **Studies in crop variation VI (1929)**
 - Fisher refines the theory of experimental design, introducing most other key concepts known today



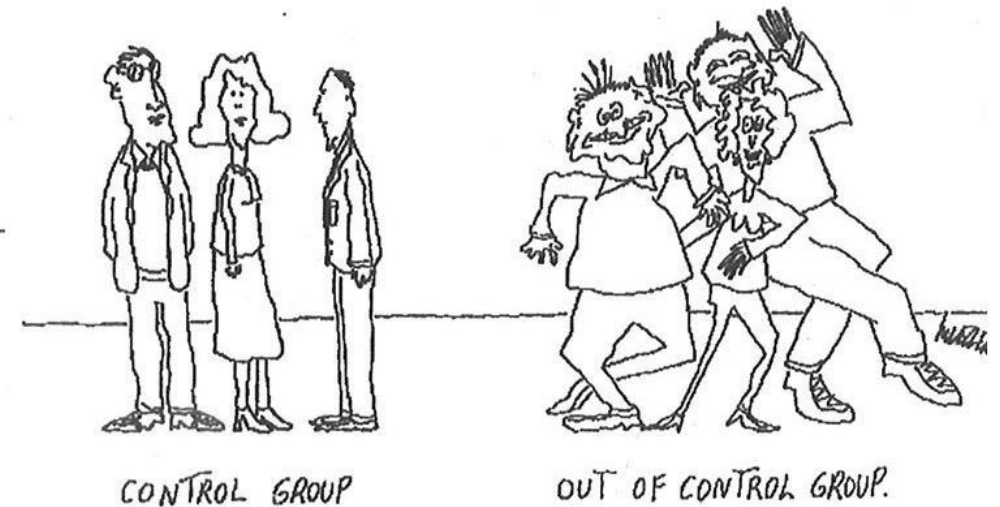
Experimental design

- **Experimental design controls background variability so that systematic effects of treatments can be observed**
- **Three basic principles**
 - **Control by matching (replication)**
 - **Control by randomization**
 - **Control by statistical adjustment**
- **Their importance is in that order**



Experimental design

- **Experimental design controls background variability so that systematic effects of treatments can be observed**
- **Three basic principles**
 - **Control by matching (replication)**
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Experimental design

- Known sources of variation may be eliminated by matching (replication)
- However **matching is limited**
 - matching is only possible on observable characteristics
 - perfect matching is not always possible
 - matching inherently limits generalizability by removing (possibly desired) variation
- **Matching ensures that groups compared are alike on specific known and observable characteristics** (in principle, everything we have thought of)

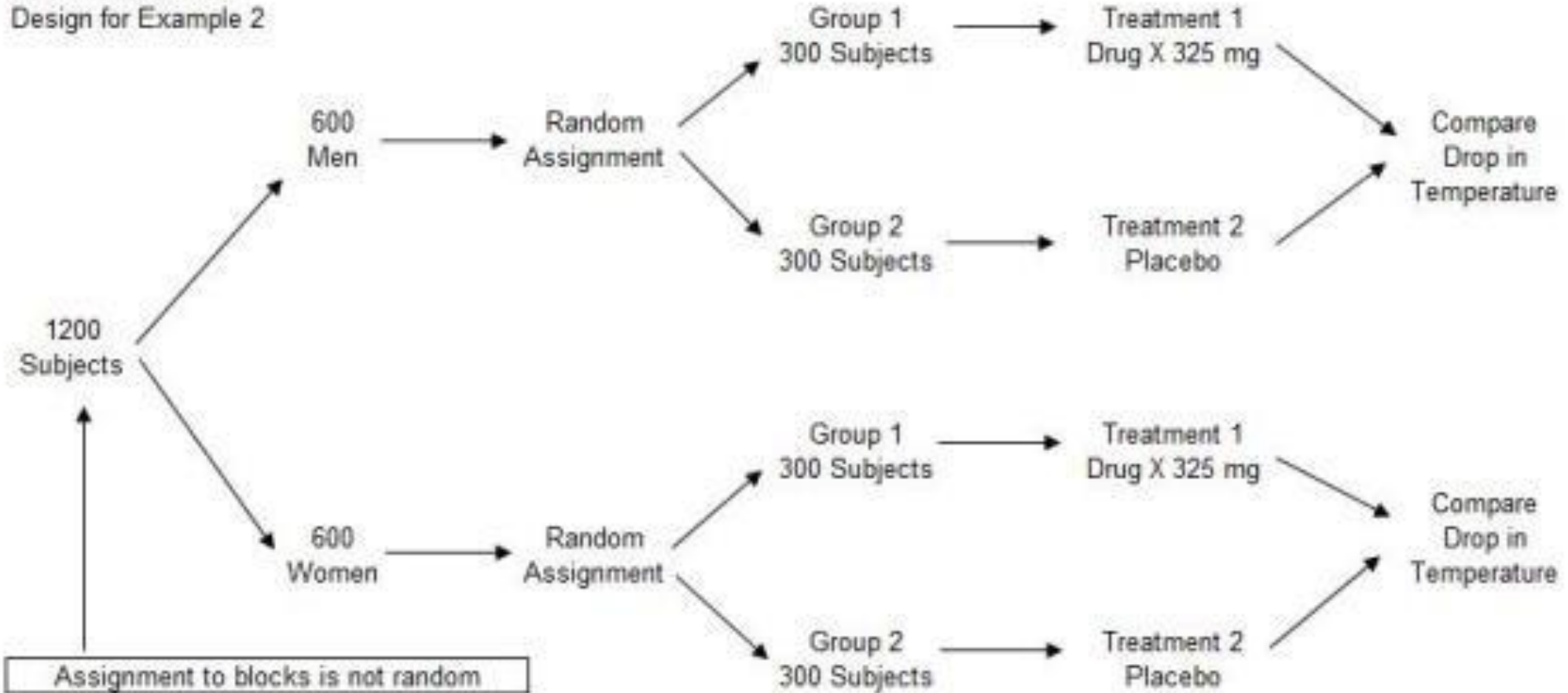


Wouldn't it be great if there were a method of making groups alike on not only everything we have thought of, but everything we didn't think of too?

Experimental design

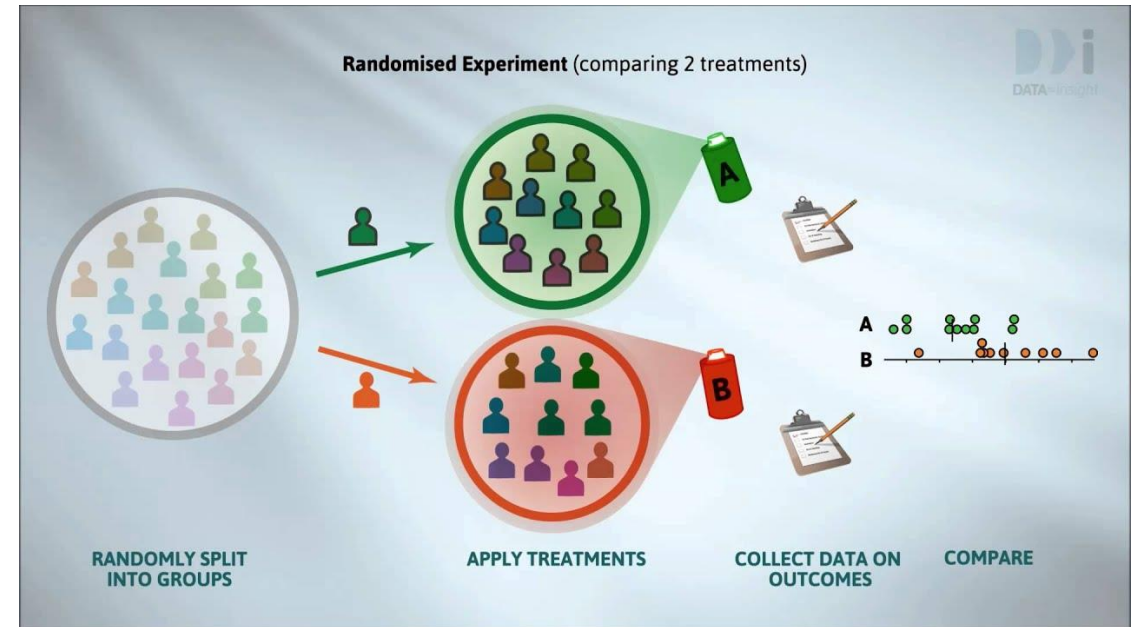
Figure 3

Randomized Block
Design for Example 2



Experimental design

- Randomization controls for the effects all (observable or non-observable, known or unknown) characteristics
- Randomization makes groups equivalent (on average) on all variables (known and unknown, observable or not)
- Randomization also gives us a way to assess whether differences after treatment are larger than would be expected due to chance.
- Random assignment is not assignment with no particular rule. It is a purposeful process



Experimental design

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Experiments are used to study causal relationships - you later independent variables and measure the alteration effect on the dependent variables

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Set of procedures to systematically test a hypothesis. A good experimental design requires a strong understanding of the system you are studying.

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Set of procedures to systematically test a hypothesis. A good experimental design requires a strong understanding of the system you are studying.

Where to start?

1- define your variables

2- define your hypothesis

3- design experimental treatments to manipulate the independent variables

4- Assign subjects to groups, either between-subjects or within-subjects (matching and randomization)

5- Think of how to measure the effects in the dependent variables

Experimental design

Examples:

Phone use and sleep

Question?

How does phone use before bedtime affects sleep patterns.

Can you further specify your question?

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Define your variables!

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Independent variable - minutes of phone use before sleep

Dependent variable - hours of sleep per night

| Research question | Independent variable | Dependent variable |
|---------------------|-----------------------------------|--------------------------|
| Phone use and sleep | Minutes of phone use before sleep | Hours of sleep per night |

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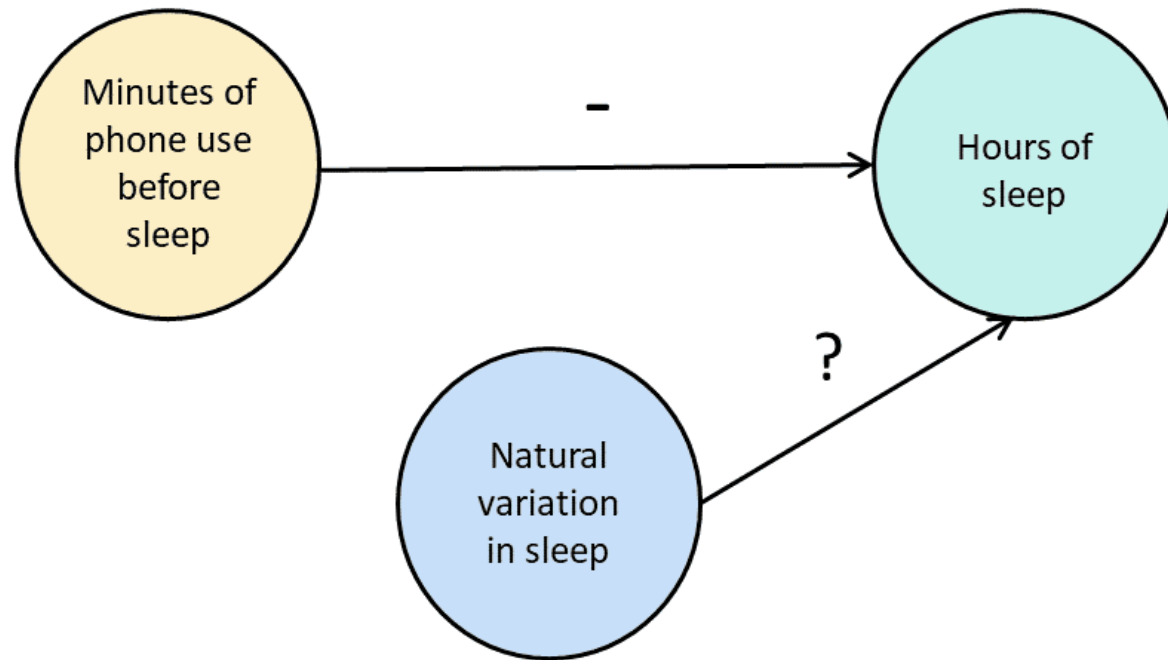
Dependent variable - hours of sleep per night

What are the external factors influencing your experiment?

Individual sleep patterns - how to overcome this?

-measure the difference between sleep with phone use and sleep without phone use in the groups (rather than the average amount of sleep per treatment group)

Experimental design



Experimental design

Write your hypothesis:

Experimental design

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H0: phone use before sleep does not correlate with the amount of sleep a person gets

H1: Increasing phone use before sleep leads to a decrease in sleep.

Define a controlled experimental approach

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Design your experimental treatments and groups

Use the phone as a:

- *categorical variable: either as binary (yes/no) or as levels of a factor (no phone use, low phone use, high phone use).*
- *continuous variable (minutes of phone use measured every night).*

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- How many individuals? Control group? How to define a “control”*
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Experimental design

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- Randomized design or randomized block design?

- Between subjects design or within subjects design?

Experimental design

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-How many individuals? Control group? How to define a “control”

-Statistical power?

- Randomized design or randomized block design?

- Between subjects design or within subjects design?

| | Completely randomized design | Randomized block design |
|----------------------------|--|---|
| Phone use and sleep | Subjects are all randomly assigned a level of phone use using a random number generator. | Subjects are first grouped by age, and then phone use treatments are randomly assigned within these groups. |

Experimental design

Measure your dependent variable

- *How to do this? How to operationalize it?*
- *How to get robust data?*

| | Between-subjects (independent measures) design | Within-subjects (repeated measures) design |
|----------------------------|--|--|
| Phone use and sleep | Subjects are randomly assigned a level of phone use (none, low, or high) and follow that level of phone use throughout the experiment. | Subjects are assigned consecutively to zero, low, and high levels of phone use throughout the experiment, and the order in which they follow these treatments is randomized. |

Experimental design

Measure your dependent variable

- *How to do this? How to operationalize it?*
- *How to get reliable and valid measurements that minimize research bias or error?*
- *How to get robust data?*

suggestions:

ask participants to record the time they go to sleep and get up,

Ask participants to wear a sleep tracker

Challenge

To define an experimental design

Copyright 2006 by Randy Glasbergen. www.glasbergen.com



**“My team has created a very innovative solution,
but we’re still looking for a problem to go with it.”**



Speaking the language of lipids in grapevine-*Plasmopara viticola* interaction



BioISI
[integrating sciences]



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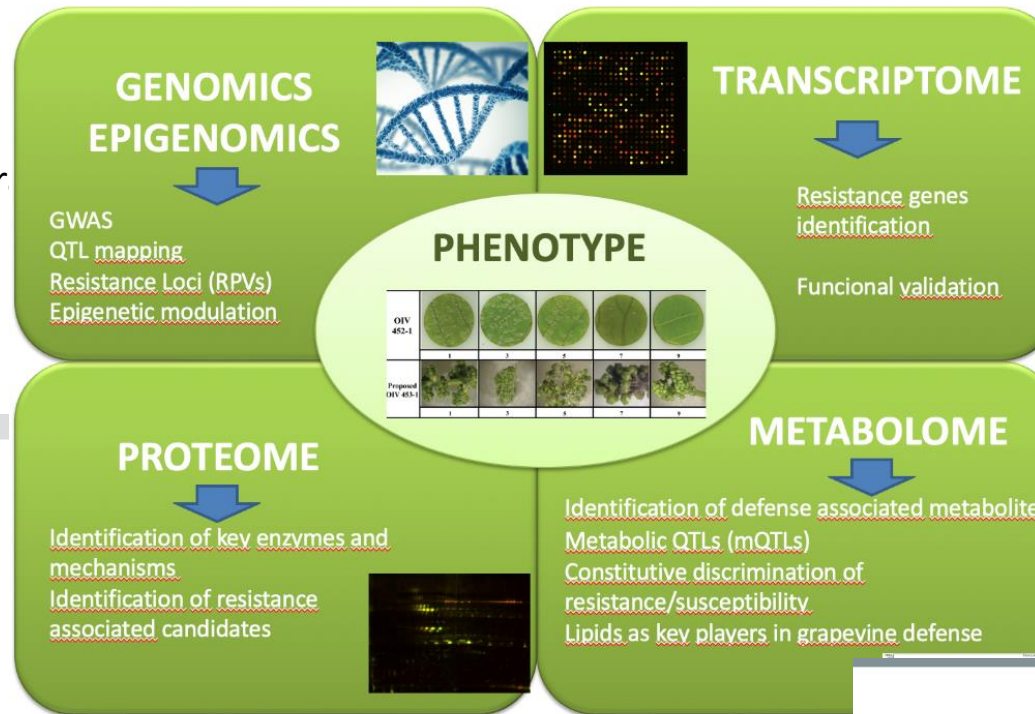
Grapevine downy mildew research

Fifteen years of molecular approaches for disease and resistance mechanisms disclosure

Vitis vinifera

2007 - Grapevine first genome draft

2014 - Genome re-annotation



Plasmopara viticola

Obligate biotroph

2012 - Sequencing projects – INRA

2016 - First genome draft

2019 - Genome available

Vol 449 | 27 September 2007 | doi:10.1038/nature06148

nature

LETTERS

The grapevine genome sequence suggests ancestral hexaploidization in major angiosperm phyla

The French–Italian Public Consortium for Grapevine Genome Characterization*

The analysis of the first plant genomes provided unexpected evidence for genome duplication events in species that had previously been considered as true diploids on the basis of their genetics^{1,2}. These whole-genome duplications may have had important roles in the evolution of plants.

All grapevine varieties are highly heterozygous; preliminary data showed that there was as much as a 13% sequence divergence between alleles, which would hinder reliable contig assembly when a whole-genome shotgun strategy was used for sequencing. Our comparison

OPEN ACCESS Freely available online

PLOS one

A High Quality Draft Consensus Sequence of the Genome of a Heterozygous Grapevine Variety

Ricardo Velasco^{1,2*}, Andrey Zharkikh^{3,4*}, Michela Troggio^{1,5*}, Dustin A. Cartwright^{1,2*}, Alessandro Cestaro¹, Dmitry Pruss², Massimo Pindo¹, Lisa M. FitzGerald⁶, Silvia Vezzulli¹, Julia Reid⁶, Giulia Malacarne¹, Diana Iliev⁷, Giuseppina Coppola¹, Bryan Wardell⁸, Diego Micheletti¹, Teresita Macalima⁹, Marco Facci¹, Jeff T. Mitchell¹⁰, Michele Perazzoli¹¹, Glenn Eldredge¹², Pamela Gatto¹³, Rozan Ozyerski¹⁴, Marco Moretto¹⁵, Natalia Gutin¹⁶, Marco Stefanini¹⁷, Yang Chen¹⁸, Cinzia Segala¹⁹, Christine Davenport²⁰, Lorenzo Demattè¹, Amy Mraz²¹, Juri Battilana¹, Keith Stormo²², Fabrizio Costa²³, Quanzhou Tao²⁴, Azeddine Si-Ammour²⁵, Tim Harkins²⁶, Angie Lackey²⁷, Clotilde Perbot²⁸, Bruce Tallon²⁹, Alessandra Stella³⁰, Victor Sokoyev³¹, Jeffrey A. Faucett³², Lieven Sterck³³, Kees Vandepoel³⁴, Stella M. Grando³⁵, Stefano Toppo³⁶, Claudio Moser³⁷, Jerry Lanchbury³⁸, Robert Rogden³⁹, Mark Skolnick⁴⁰, Vittorio Sgaramella⁴¹, Satish K. Bhatnagar⁴², Paolo Fontana⁴³, Alexander Gutin⁴⁴, Yves Van de Peer⁴⁵, Francesco Salami⁴⁶, Roberto Viola⁴⁷

¹IASMA Research Center, San Michele all'Adige, Trento, Italy, ²Myriad Genetics Inc, Salt Lake City, Utah, United States of America, ³Life Sciences Corporation, Branford, Connecticut, United States of America, ⁴Roche Diagnostics Corporation, Roche Applied Science, Indianapolis, Indiana, United States of America, ⁵Amplicon Express Inc., Pullman, Washington, United States of America, ⁶Technology Park Lodi, Lodi, Italy, ⁷Department of Plant Systems Biology, VIB, Ghent University, Ghent, Belgium, ⁸Department of Biological Chemistry, Padova University, Padova, Italy, ⁹Department of Computer Science, Royal Holloway, University of London, Egham, Surrey, United Kingdom



genome announcements



Draft Genome Sequence of *Plasmopara viticola*, the Grapevine Downy Mildew Pathogen

Yann Dussert,^a Jérôme Gouzy,^b Sylvie Richart-Cervera,^a Isabelle D. Mazet,^a Laurent Delière,^a Carole Couture,^a Ludovic Legrand,^b Marie-Christine Piron,^a Pere Mestre,^a François Delmotte^a
^aSAVE, Bordeaux Sciences Agro, INRA, Villenave d'Ornon, France; ^bLIPM, Université de Toulouse, INRA, CNRS, Castanet-Tolosan, France; ^cSVOV, INRA, Université de Strasbourg, Colmar, France

www.nature.com/scientificreports

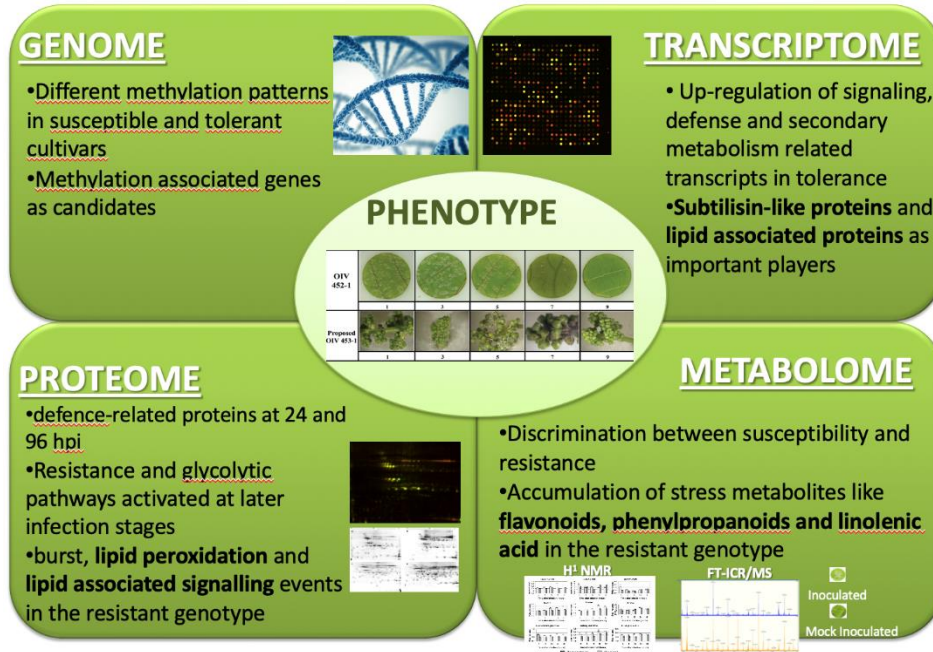
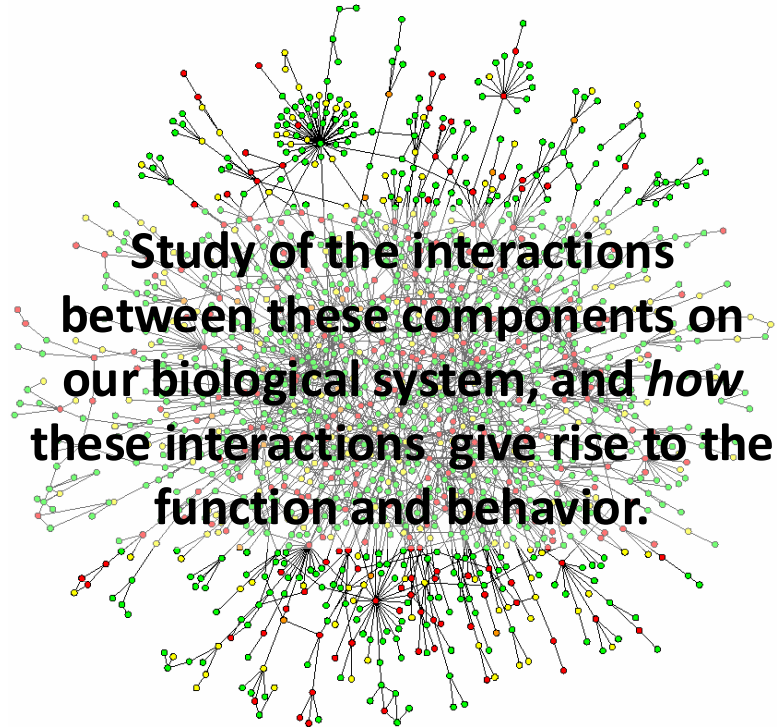
SCIENTIFIC REPORTS

OPEN Genome sequence of *Plasmopara viticola* and insight into the pathogenic mechanism

Ling Yin^{1,2,3}, Yunhe An^{1,3,4}, Junjie Qu⁵, Xinlong Li², Yali Zhang¹, Ian Dry⁶, Huijuan Wu¹ & Jiang Lu^{1,3}

Received: 03 November 2016
 Accepted: 22 March 2017
 Published: 18 April 2017

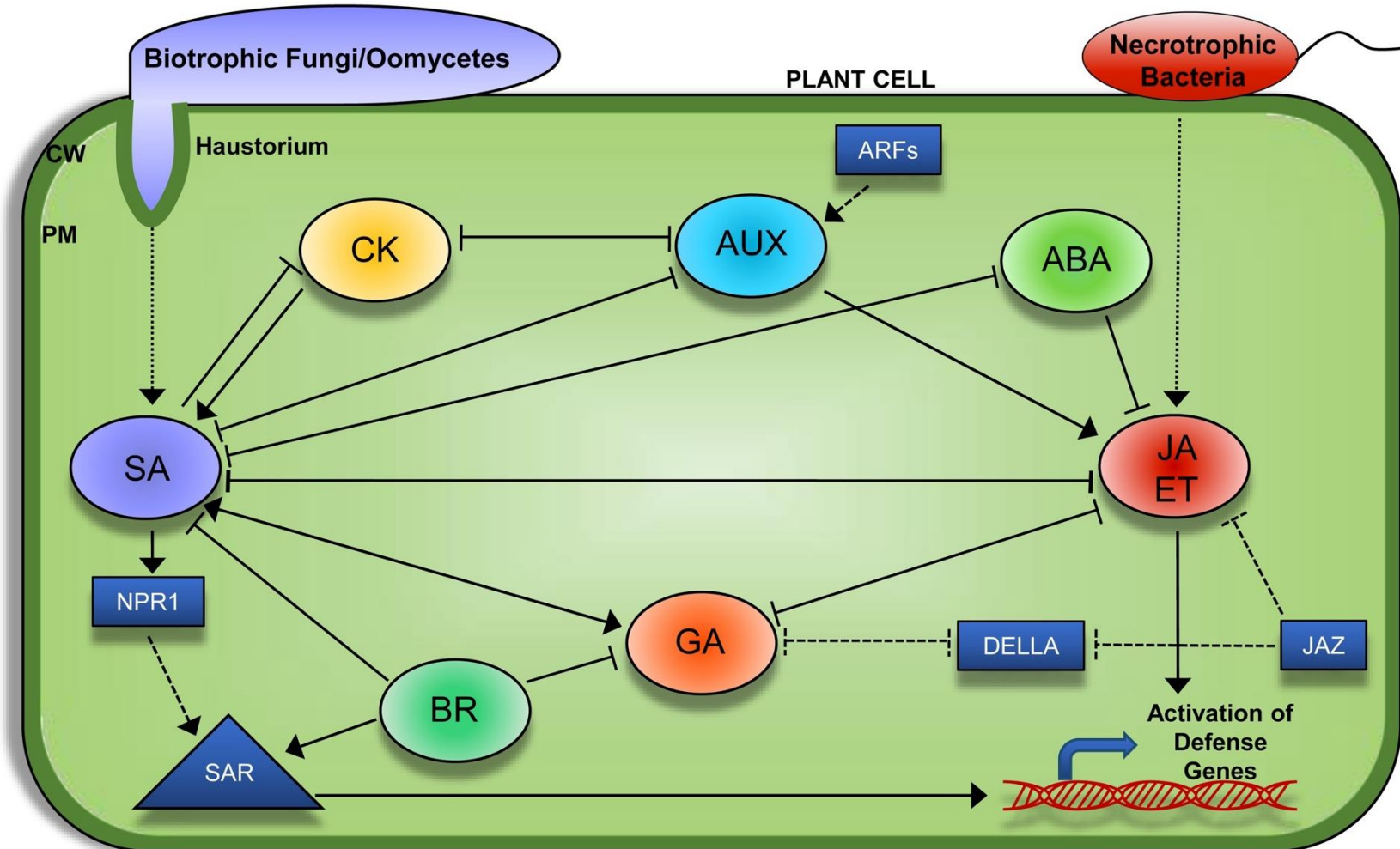
Systems biology – something about networks



Jasmonic acid signaling? Lipids? Fatty acids?

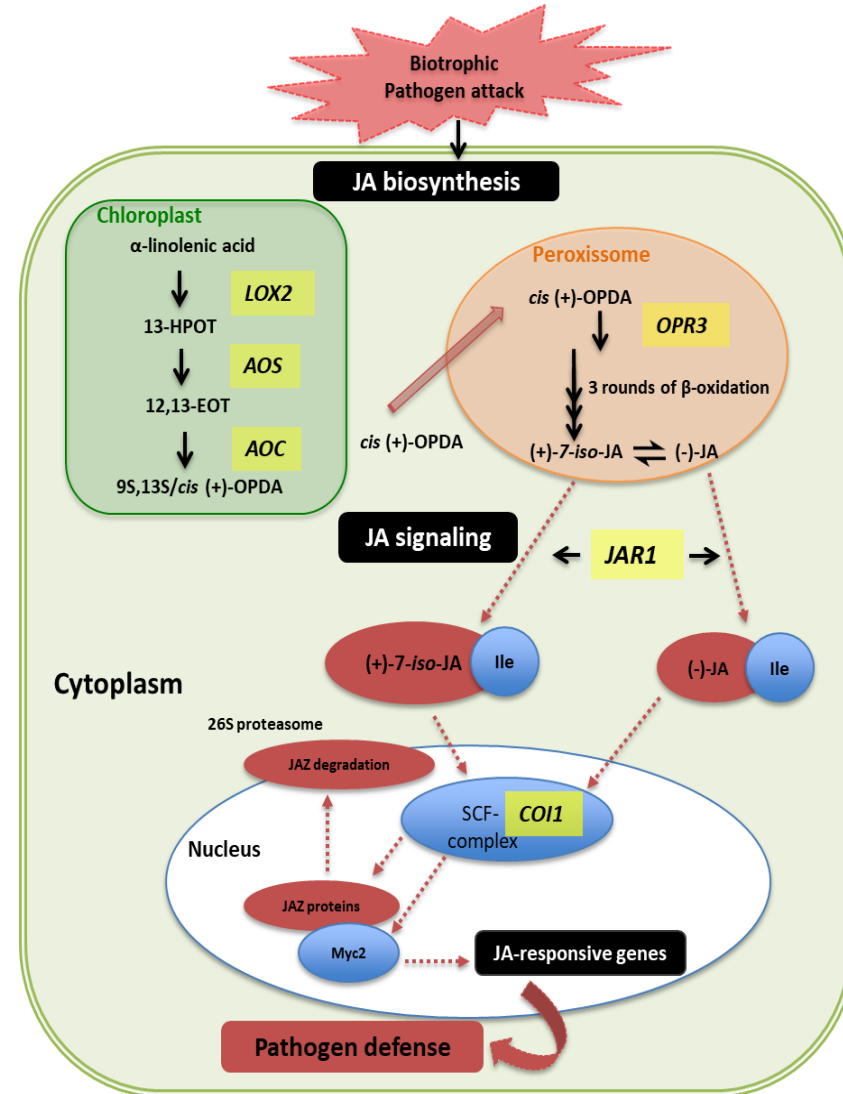
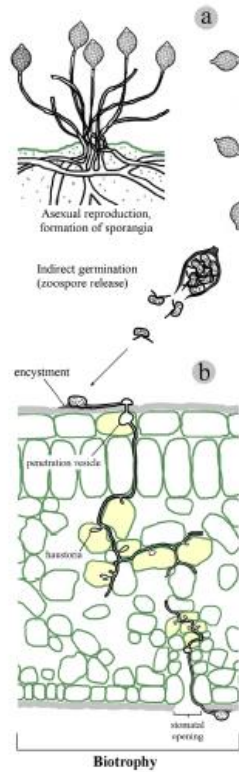
Hormone signaling in plant-pathogen interactions

What was the basic knowledge on hormone signaling in plant-pathogen interactions?



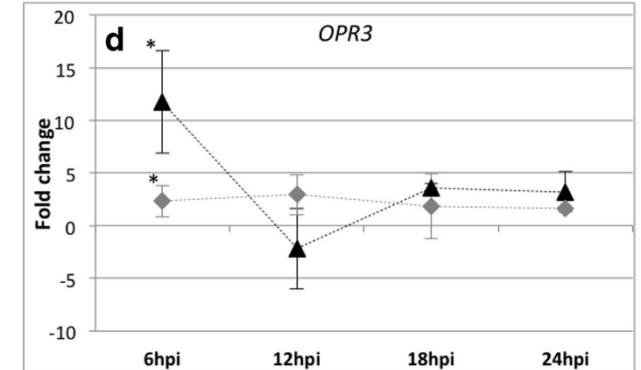
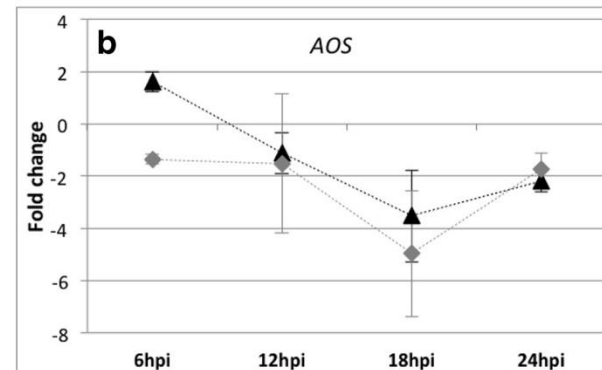
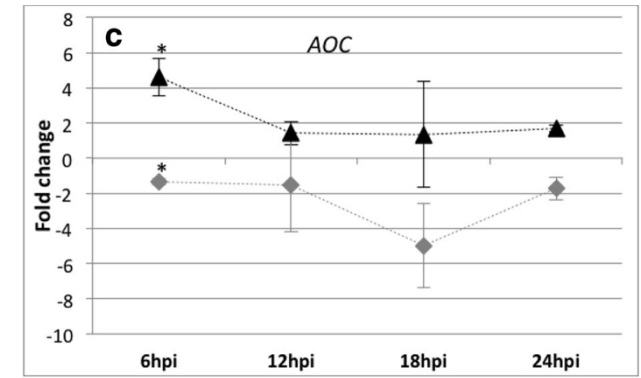
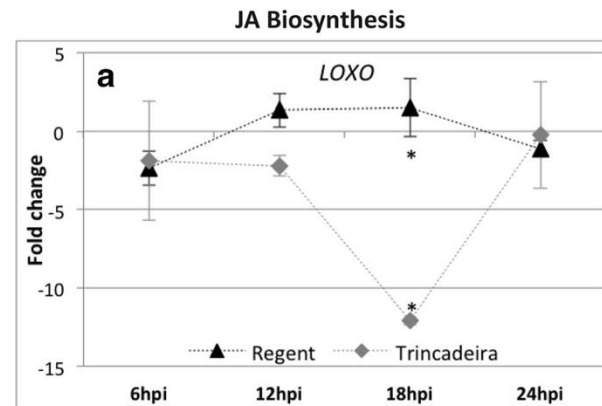
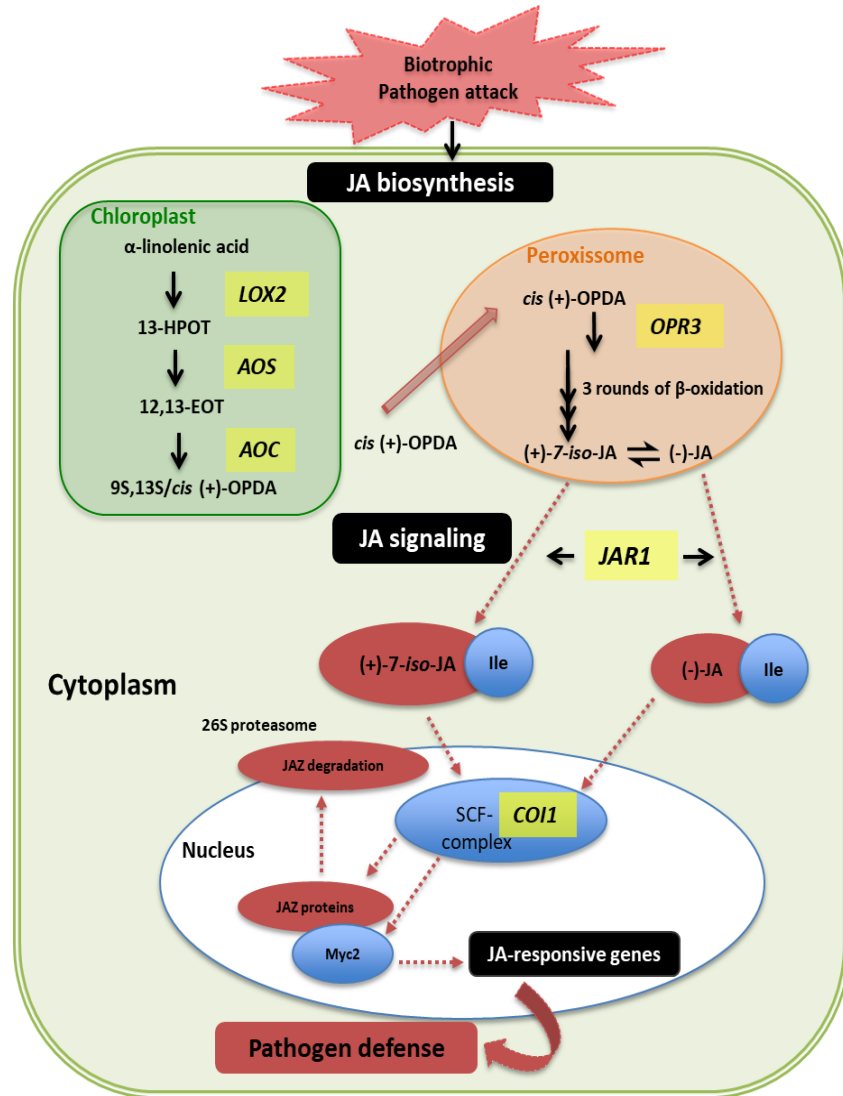
Hormone signaling in grapevine-*P. viticola* interaction

Jasmonic acid signaling activated in the first hours of infection with a biotroph?



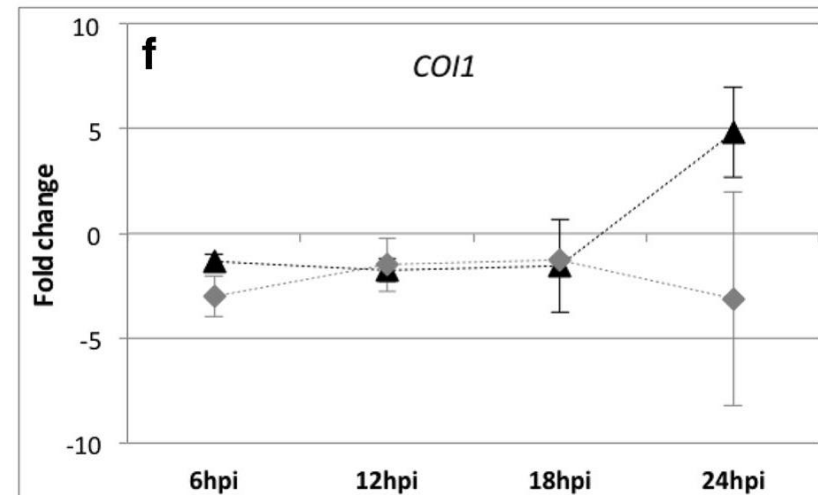
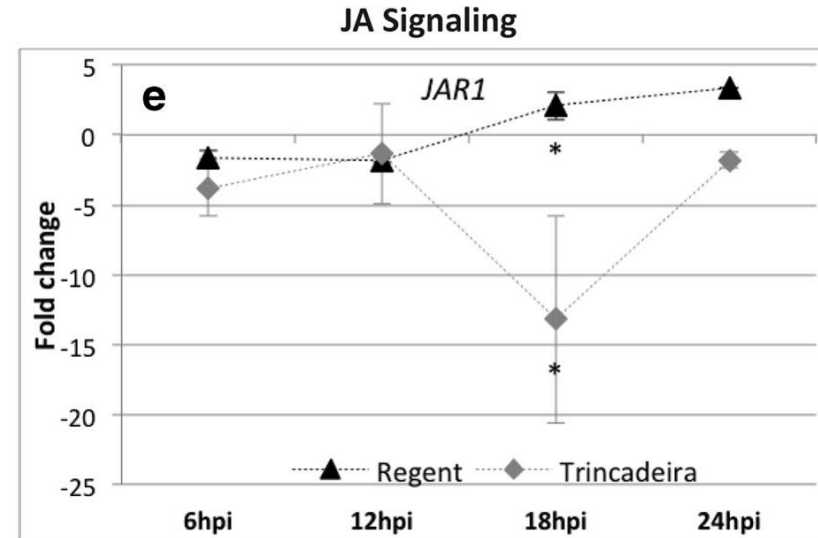
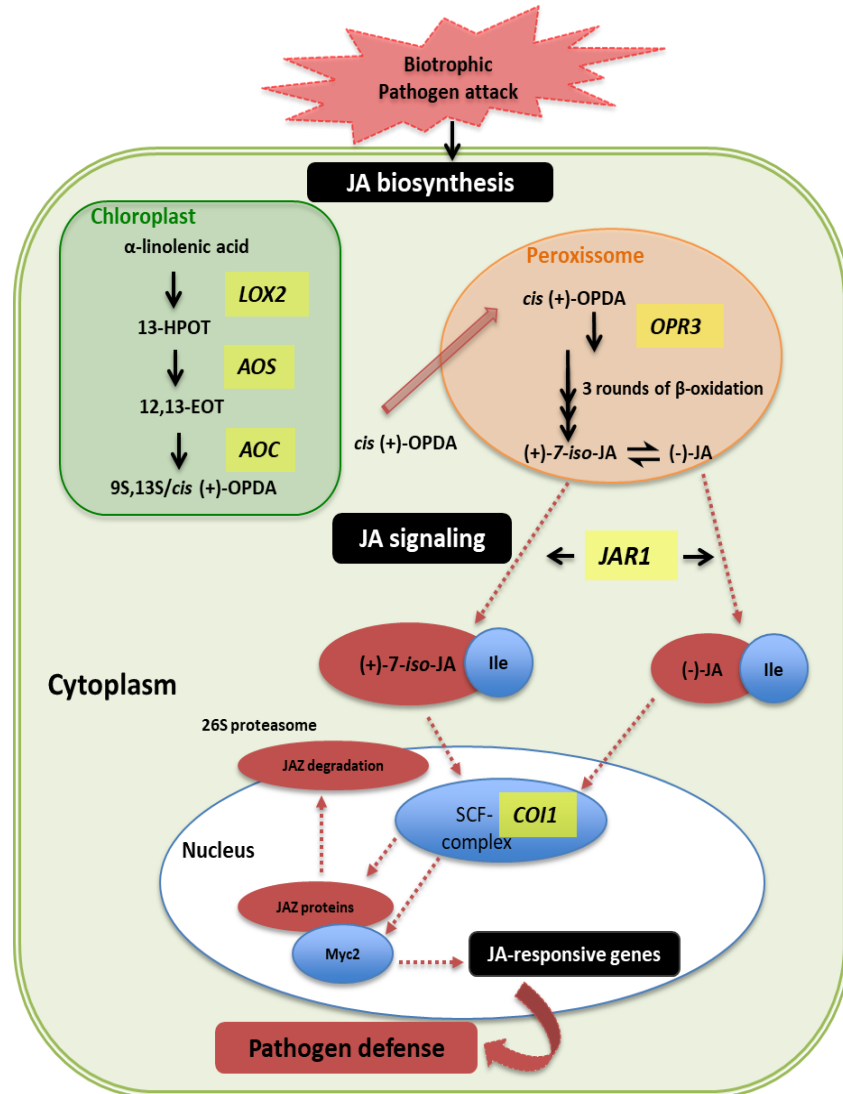
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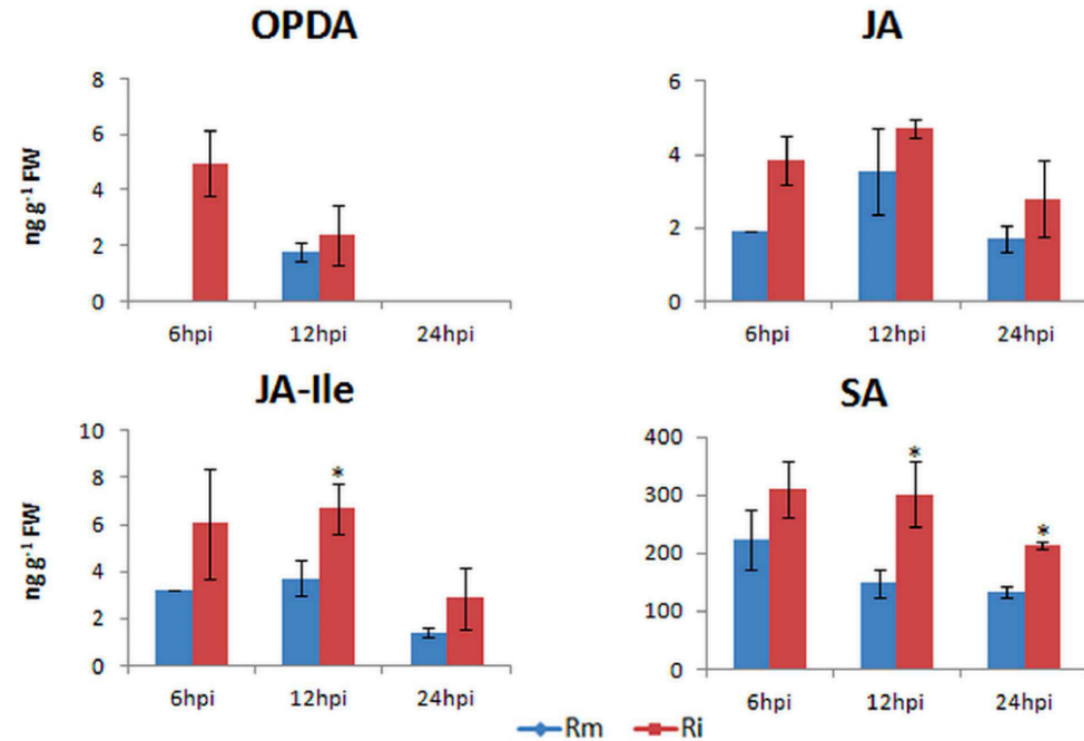
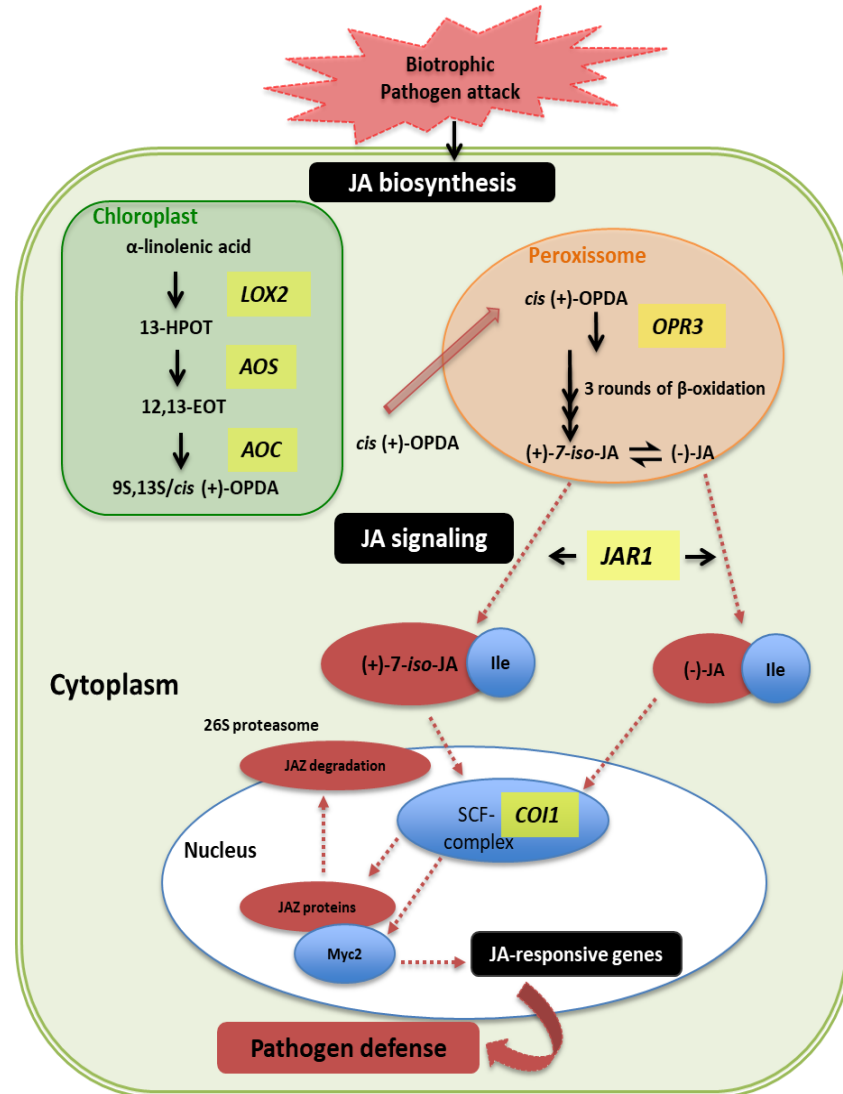
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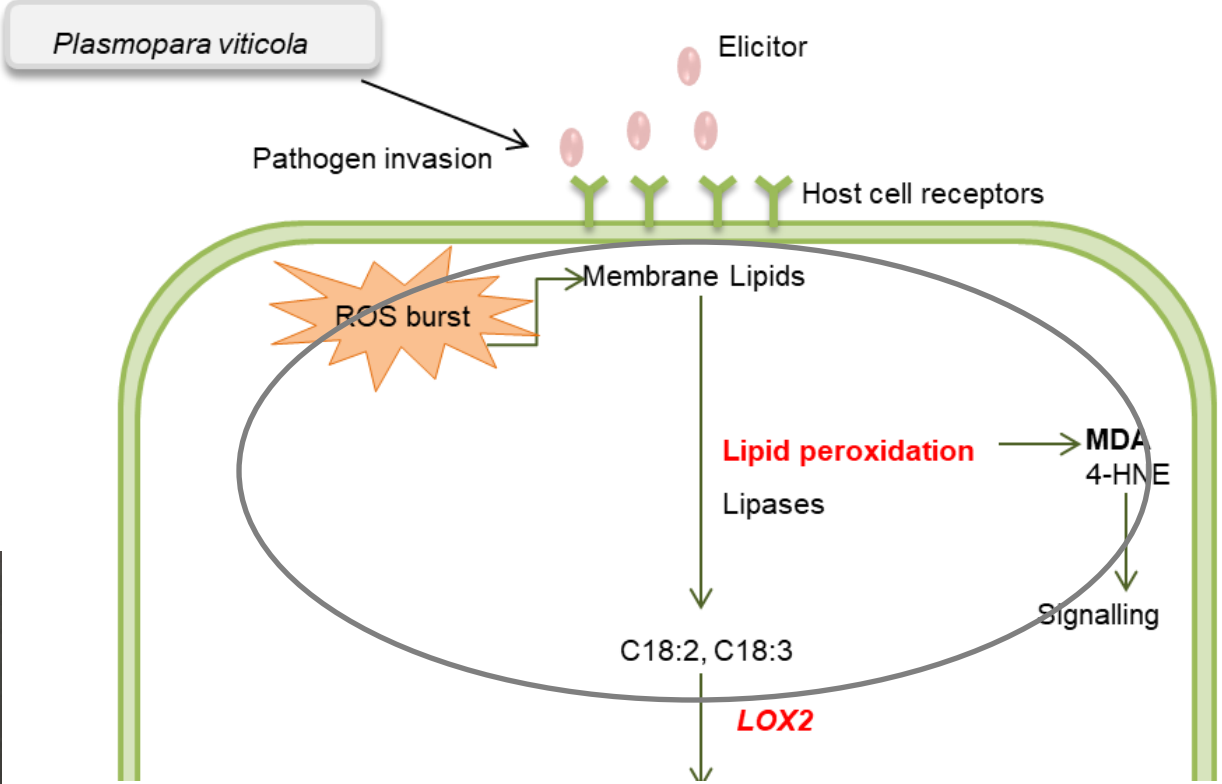


Hormone signaling in grapevine-*P. viticola* interaction

Jasmonic acid signaling activated in the first hours of infection with a biotroph?



Which events lead to JA precursor accumulation?



Tolerant cultivar:

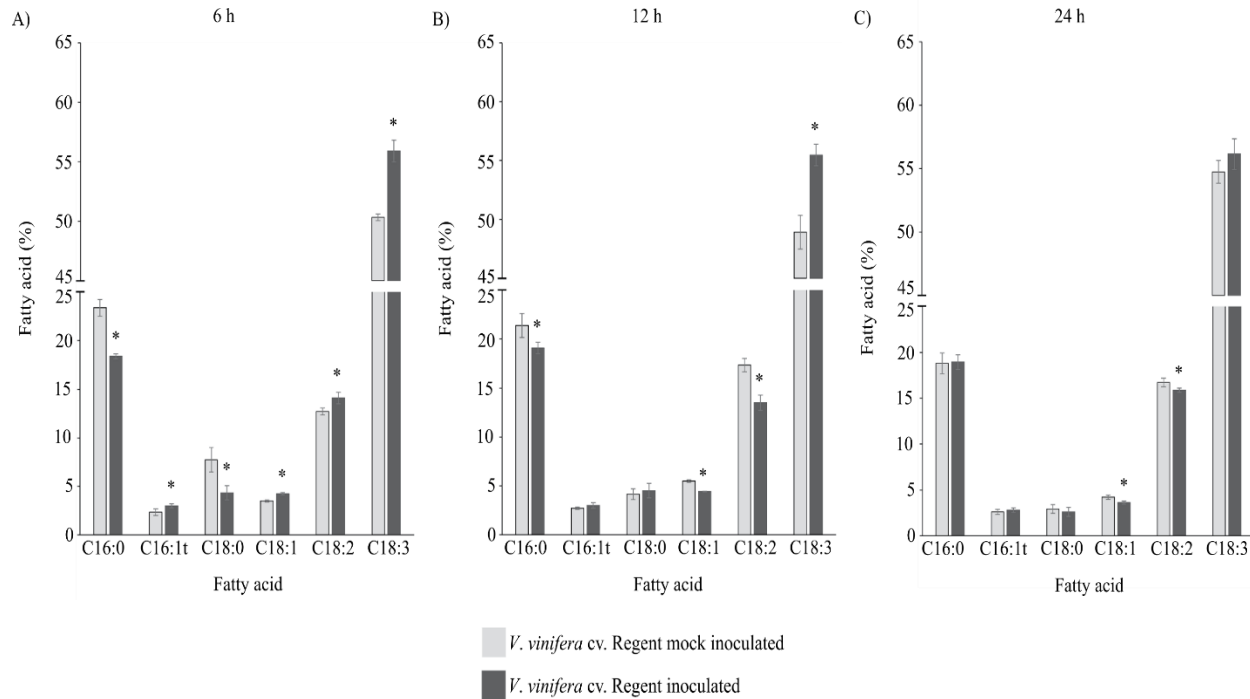
↑ JA biosynthesis

↑ Accumulation of the precursor C18:3



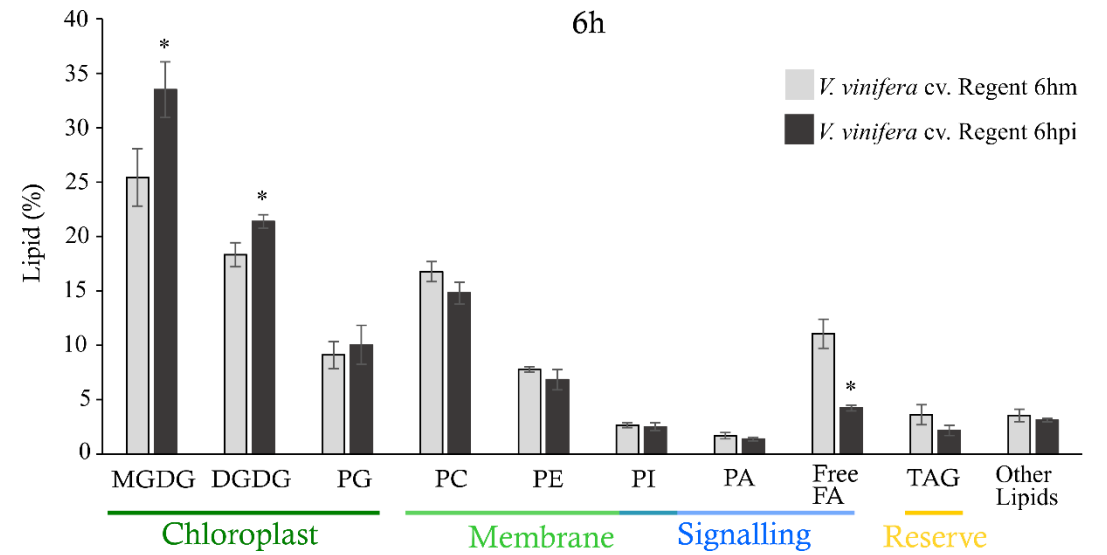
Why and how tolerant cultivars present a higher content of free C18:3 in the chloroplast?

Which events lead to JA precursor accumulation?

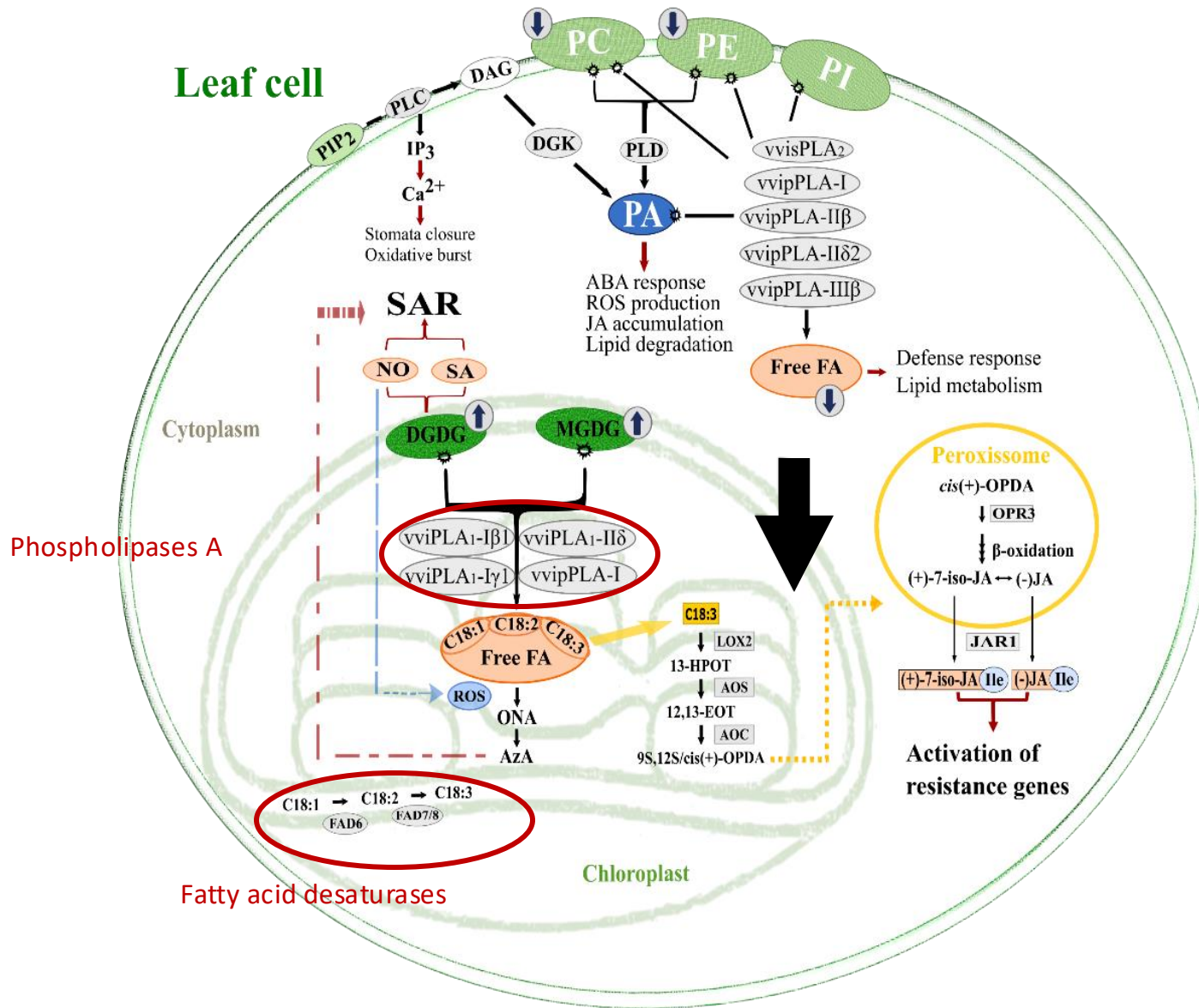


FA modulation after *P. viticola* inoculation **only occurs in the tolerant cultivar**

Increase of C18 desaturation;
C18:3 accumulation;
Increase of PUFAs.



P. viticola inoculation triggers FA and lipid content alteration



Players

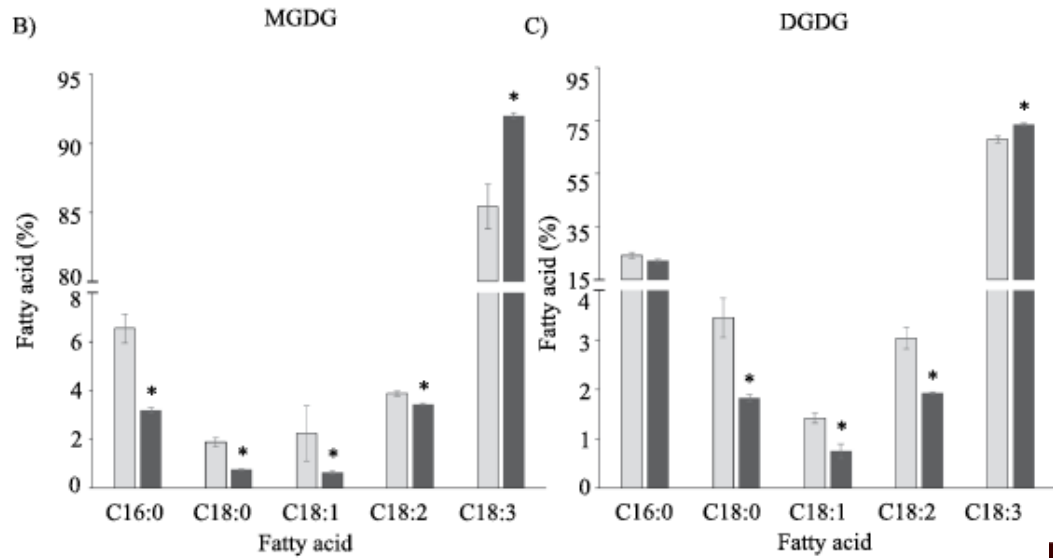
Phospholipases

Catalyze the hydrolysis membrane lipids into lysophospholipids and fatty acids

Fatty acid desaturases

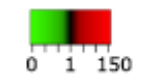
Introduce double bonds in fatty acids present in membrane lipids leading to the synthesis of unsaturated FA

P. viticola inoculation triggers FA and lipid content alteration



V. vinifera cv. Regent 6hm
 V. vinifera cv. Regent 6hpi

Activation of chloroplastidial *Fatty Acid Desaturases* (FAD6, FAD7, FAD8)



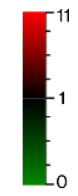
| | | | |
|---------------|----------------|--------------|---------------------|
| 27.96±19.90 * | 114.87±42.36 * | 18.43±5.56 * | <i>VviPLA1-β1</i> |
| 2.79±1.73 | 18.57±3.16 * | 29.28±0.69 * | <i>VviPLA1-γ1</i> |
| 4.27±0.15 * | 0.71±0.58 | 1.01±0.14 | <i>VviPLA1-IIIδ</i> |

Activation of chloroplastidial Phospholipases A



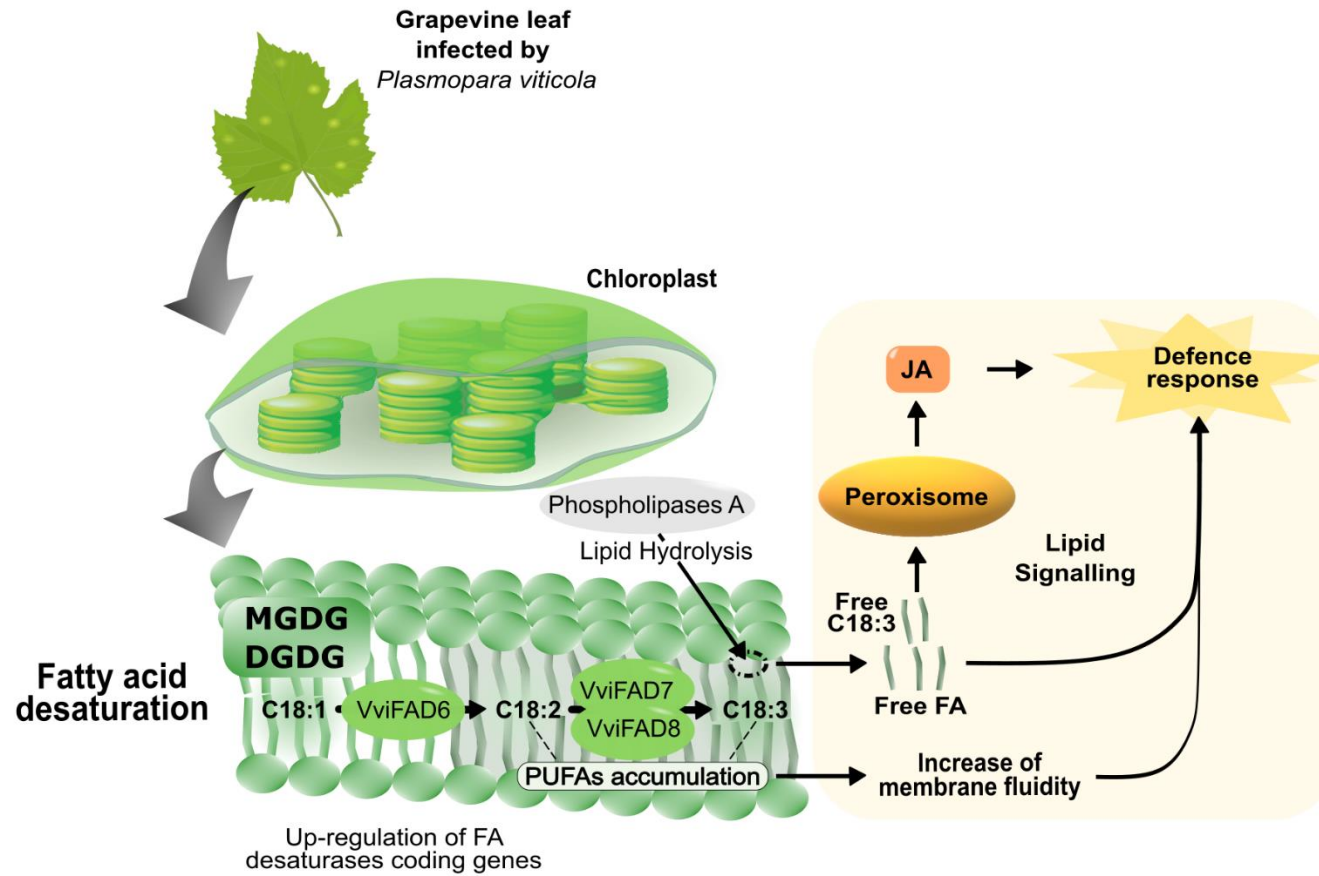
Increase Free FA pool

| | | | |
|----------------|---------------|---------------|------------------|
| 3.25 ± 0.65 * | 1.08 ± 0.38 | 1.53 ± 0.20 | <i>VviFAD2-1</i> |
| 3.17 ± 1.09 * | 0.06 ± 0.04 | 0.21 ± 0.01 | <i>VviFAD2-2</i> |
| 0.03 ± 0.01 * | 0.34 ± 0.30 * | 1.51 ± 0.53 | <i>VviFAD3-1</i> |
| 2.66 ± 0.61 * | 1.13 ± 0.41 * | 2.8 ± 0.11 * | <i>VviFAD4</i> |
| 2.01 ± 0.75 * | 1.11 ± 0.61 | 2.25 ± 0.24 * | <i>VviADS</i> |
| 2.78 ± 0.72 * | 1.17 ± 0.14 | 1.76 ± 0.24 | <i>VviFAD6</i> |
| 1.14 ± 0.26 * | 1.12 ± 0.43 | 1.91 ± 0.19 * | <i>VviFAD7</i> |
| 10.87 ± 4.51 * | 1.13 ± 0.91 | 2.44 ± 0.21 * | <i>VviFAD8</i> |
| 2.21 ± 0.52 * | 1.38 ± 0.11 | 1.92 ± 0.01 * | <i>VviSAD-1</i> |
| 2.96 ± 0.72 * | 1.4 ± 0.18 | 1.56 ± 0.05 * | <i>VviSAD-2</i> |
| 6hpi | 24hpi | 48hpi | |



Increase FA desaturation

P. viticola inoculation triggers FA and lipid content alteration



Increase of membrane fluidity leads to the protection of photosynthetic apparatus

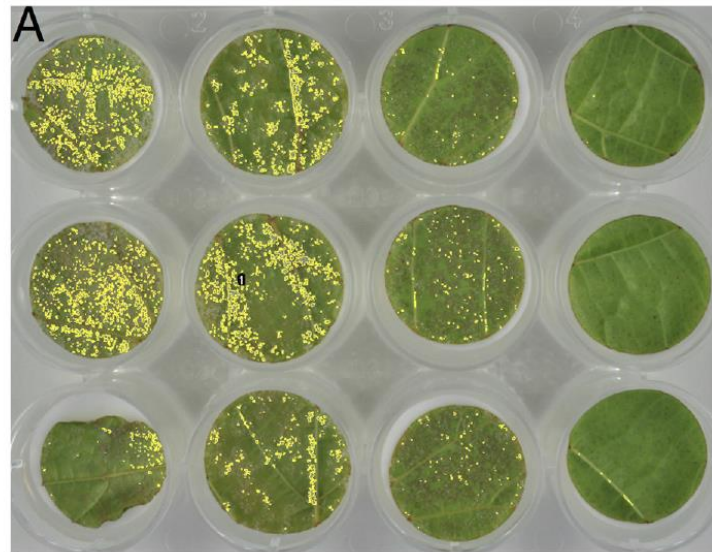
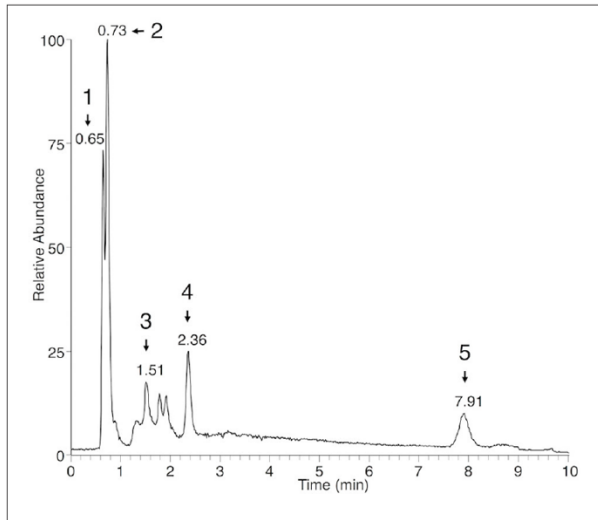
Higher availability of bounded and free C18:3, the Jasmonic Acid precursor



lipid signaling activation

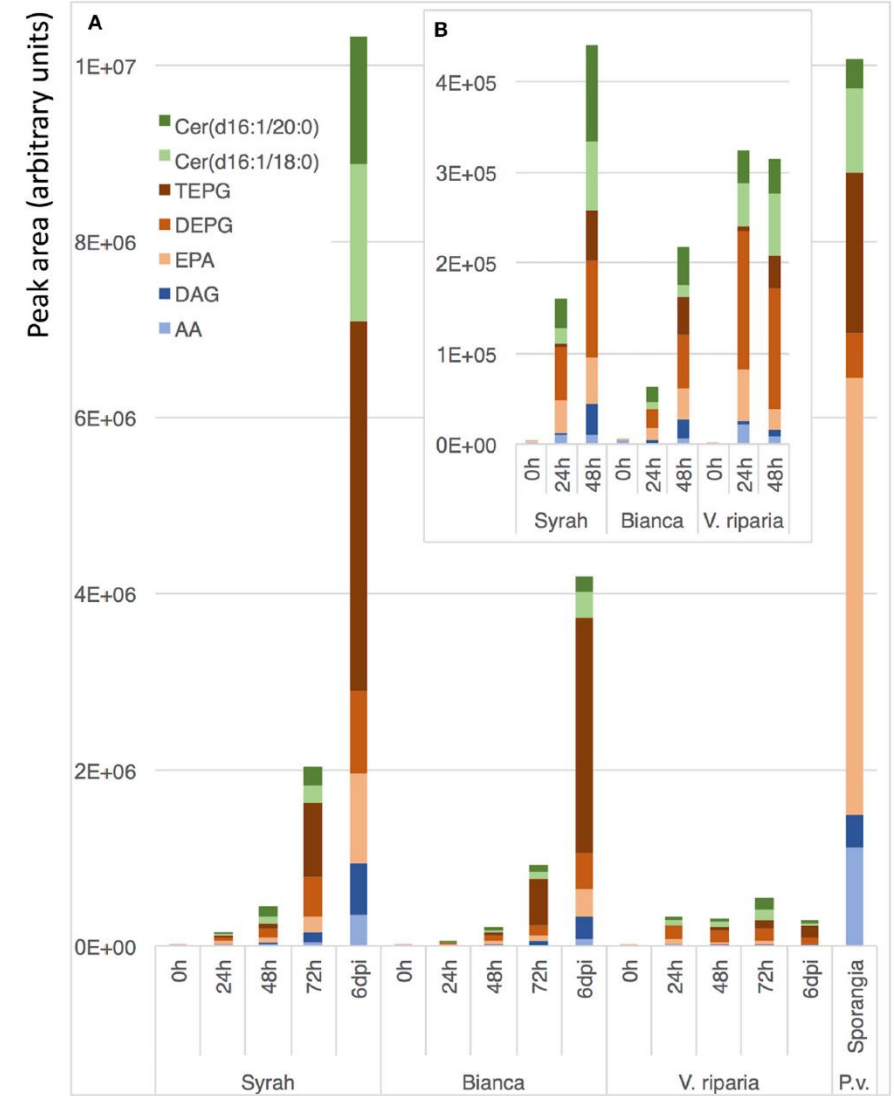
How about *Plasmopara viticola* lipids?

Are there pathogen specific lipids accumulated during the interaction?



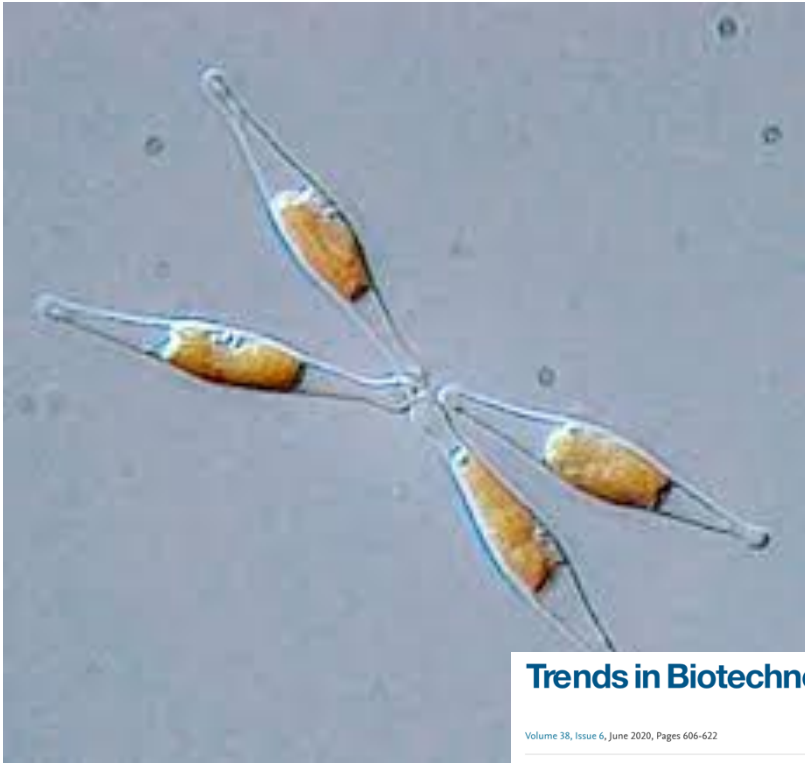
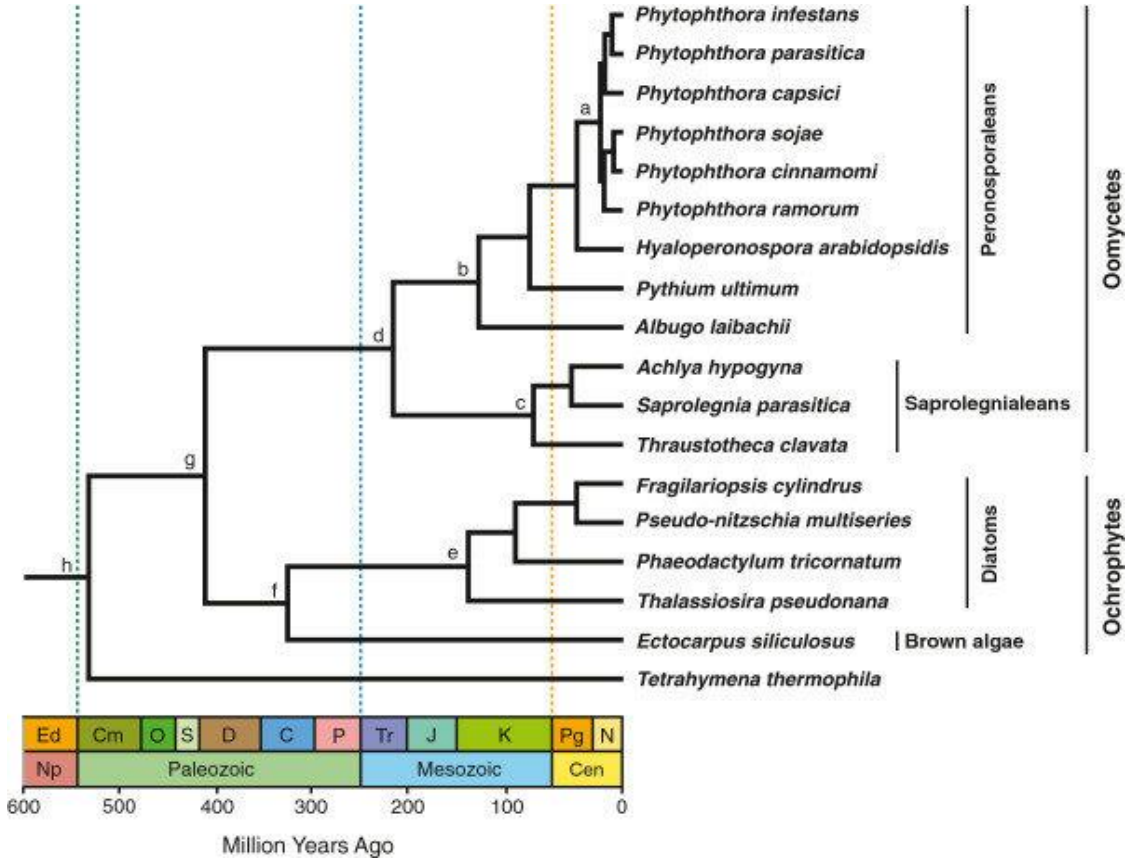
B

| Metabolite | Pearson's correlation coefficient | P-value |
|-----------------|-----------------------------------|-----------|
| EPA | 0.849 | 1.37 e-18 |
| AA | 0.848 | 1.91 e-18 |
| Cer(d16:1/18:0) | 0.840 | 7.56 e-18 |
| Cer(d16:1/22:0) | 0.775 | 8.82 e-14 |
| DAG | 0.741 | 3.71 e-12 |
| Cer(d16:1/16:0) | 0.728 | 1.38 e-11 |
| Cer(d16:1/20:0) | 0.709 | 8.17 e-11 |
| TEPG | 0.702 | 1.47 e-10 |
| AG | 0.693 | 3.02 e-10 |
| EPG | 0.691 | 3.58 e-10 |
| DEPG | 0.689 | 4.18 e-10 |



How about *Plasmopara viticola* lipids?

Plasmopara viticola lipids that accumulate during infection and sporulation- do not exist in grapevine but there are a shared trait with diatoms.



Trends in Biotechnology



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Review

Phaeodactylum tricornutum: A Diatom Cell Factory

Thomas Butler,¹ Rahul Vijay Kapoor,^{1,2} Seetharaman Vaidyanathan,¹

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**“My team has created a very innovative solution,
but we’re still looking for a problem to go with it.”**