



Pre-breeding strategies for obtaining new resilient and added value berries

From genomes to New Breeding Techniques to obtain new resilient and quality cultivars.

Prof. Bruno Mezzetti

Dip. di Scienze Agrarie, Alimentari ed Ambientali

Università Politecnica delle Marche

b.mezzetti@staff.univpm.it



The History of Genetic Modification in CROPS

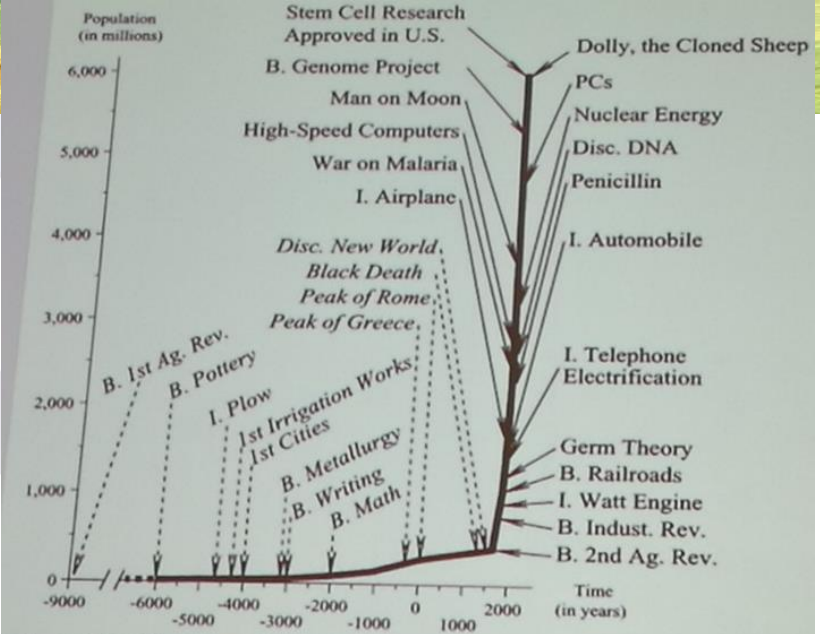
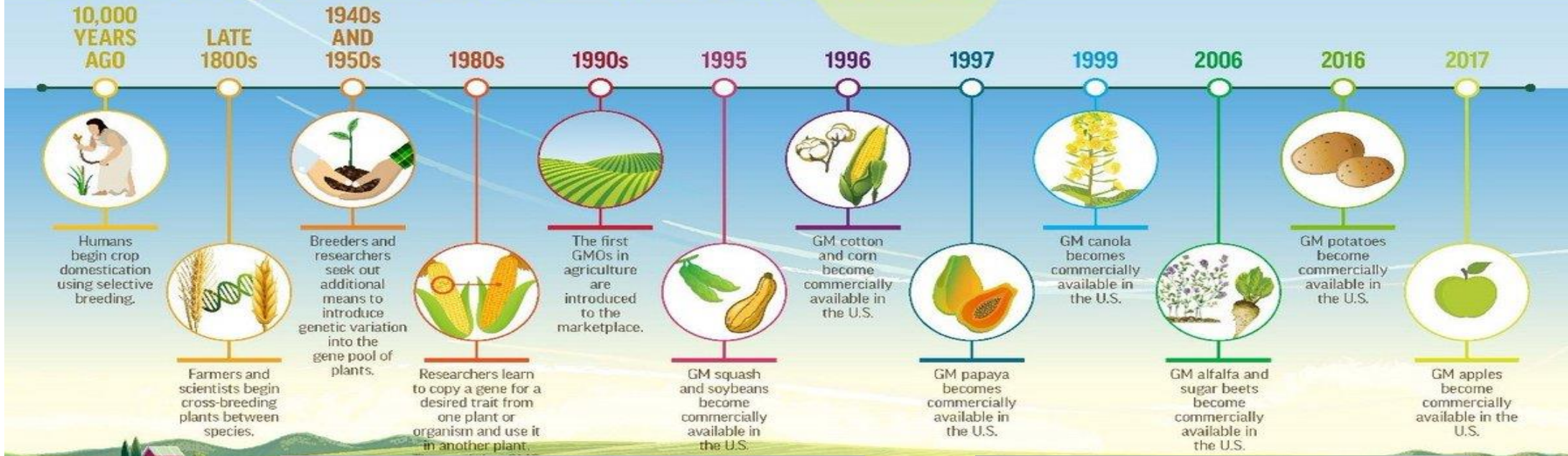
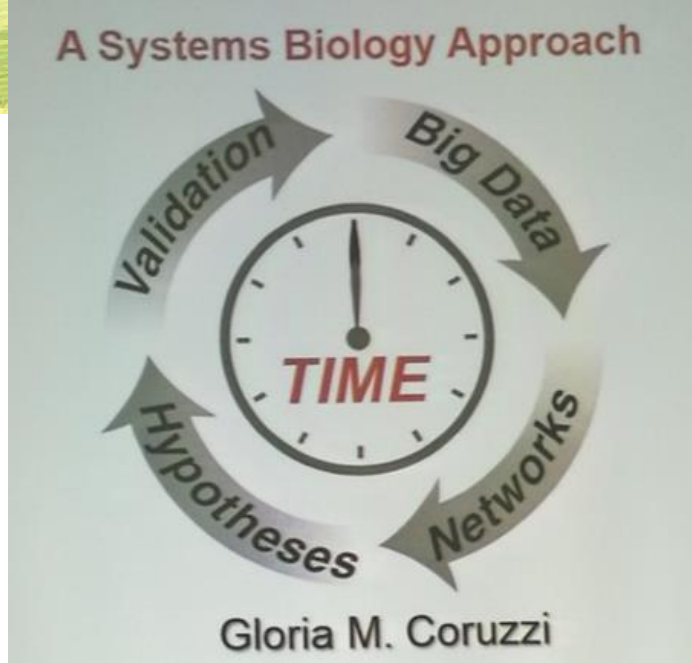
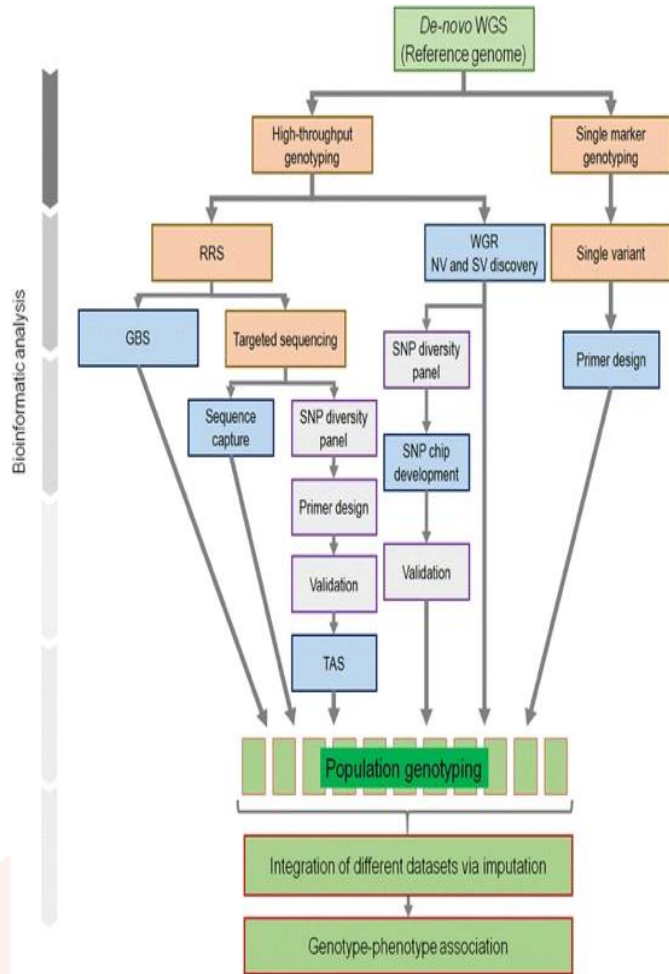


Figure 2.1 The Growth of World Population and Some Major Events in the History of Technology.

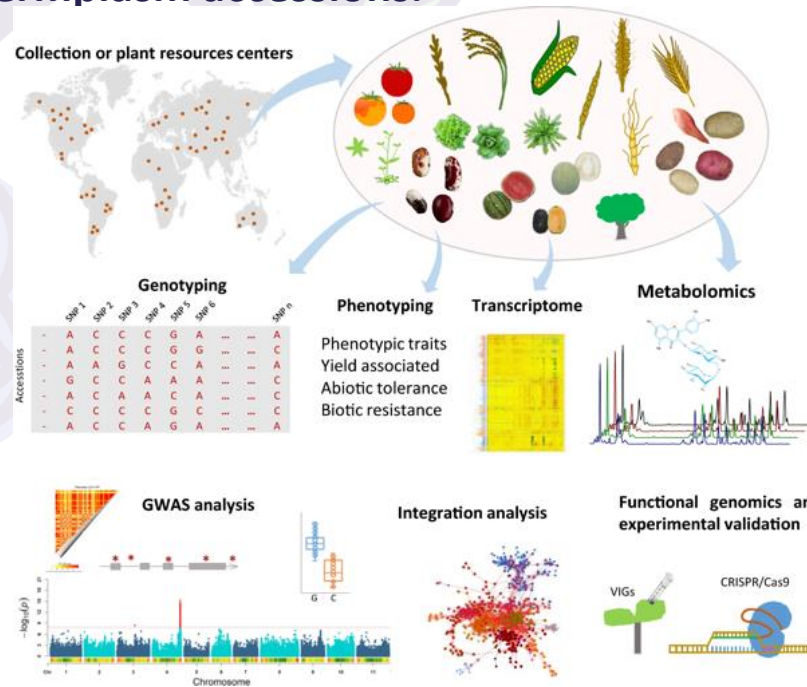


Bioinformatic tools and sequencing strategies

GENOME WIDE ASSOCIATION STUDIES - GWAS

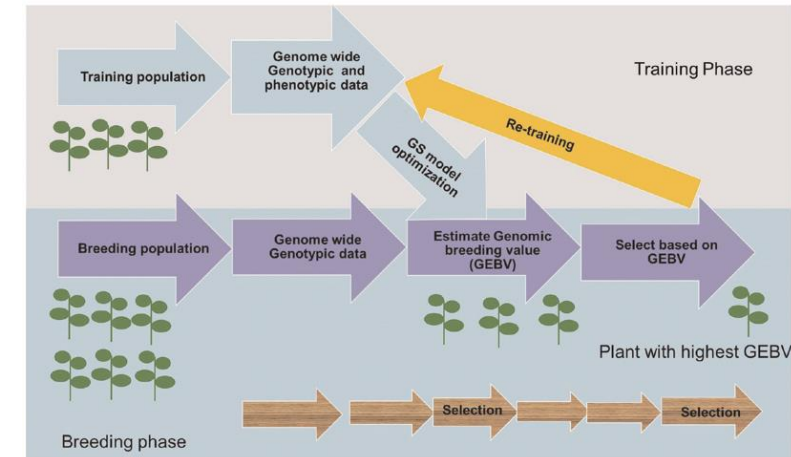


Genome-wide association studies (GWAS) enable the detection and identification of quantitative trait loci (QTLs) and genes controlling phenotypic variations in a collection of cultivars and germplasm accessions.

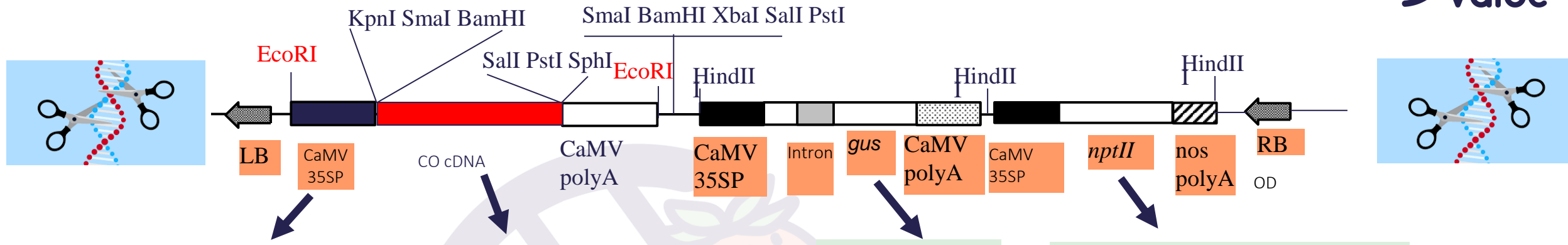


Genomic selection

Genomic selection (GS) enables the selection of superior genotypes based on genomic estimated breeding values (GEBV) derived from the information of genome-wide DNA polymorphisms.



DNA AND GENES: Transgenes, Cisgenes, RNAi, gene editing



Promotor
Transgene/Cisgene

Regulator Gene

Constitutive
 Site specific
 Stress induced

Gene

Transgene

Cis/Intragenic

RNAi

CRISPR

Protein/enzyme/RNAi/CAS9

TRAIT

GUS/GFP Reporter

Transgene TO AVOID

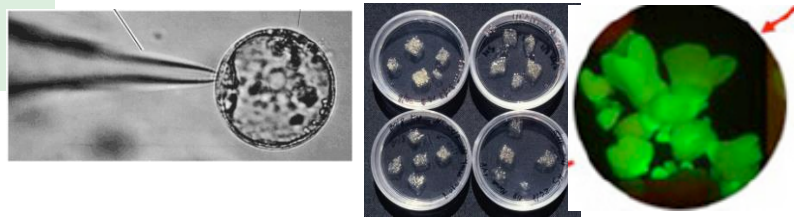
nptII

Antibiotic marker - to be avoided

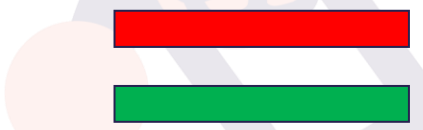
Use new cisgenic markers such as the MYB gene

Problems in selection efficiency and chimerism

Clean Gene editing: transient expression in protoplast



Or plant segregation of Agro mediated transformed plant



Stable strawberry intragenic lines expressing *FaWRKY1*-RNAi and *FaNPR3.1*-RNAi driven by *FaAAT2* and *FaDOF2* promoters

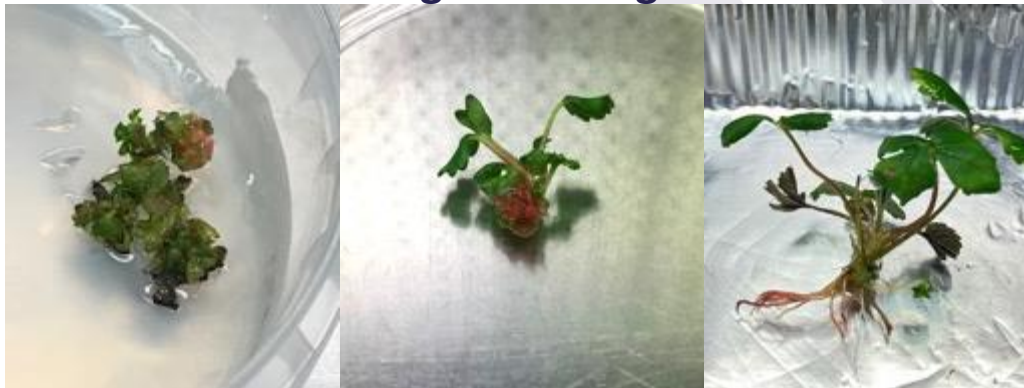


6 weeks after infection

15 weeks after infection

18 weeks after infection

Putative elongated intragenic lines



- Without using selectable marker genes, the possibility of obtaining a complete transformed shoot is very low
- High percentage of chimeric shoots

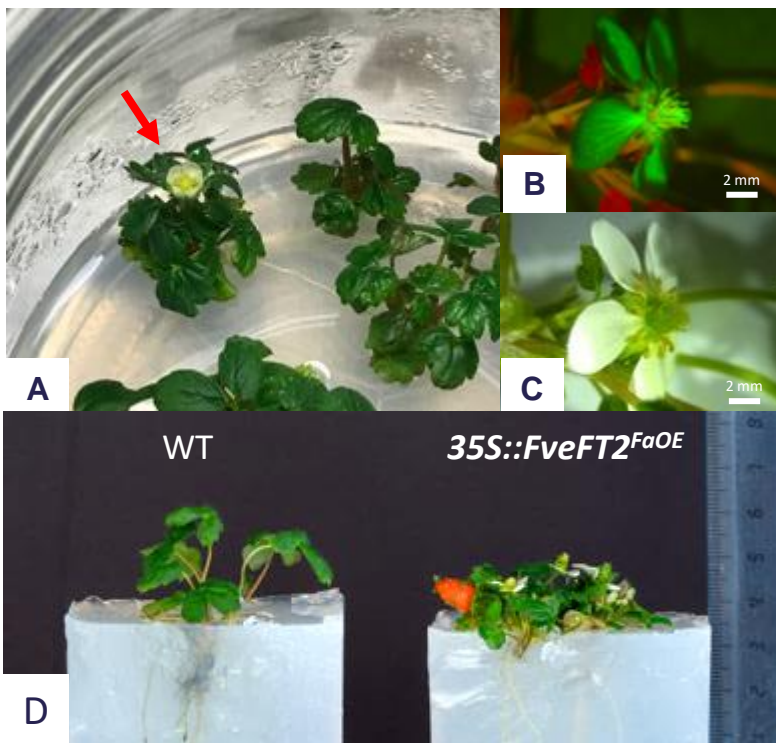


Alternative strategies are necessary

Transformation efficiency of four apple varieties without the use of selectable marker genes (through MYB expression only) (Transformation efficiency of apple with kanamycin is around 0.22 e 13.8%).



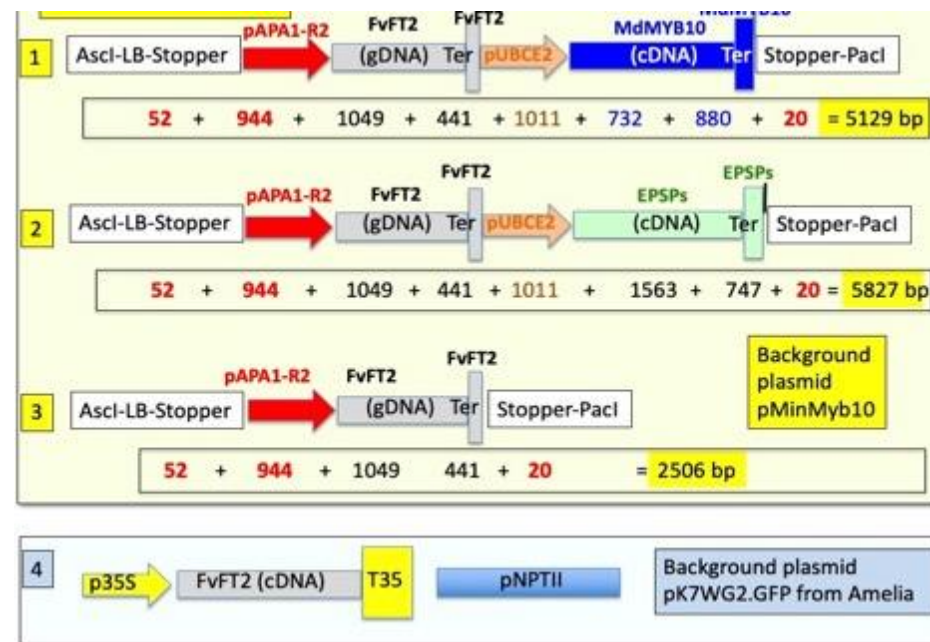
In vitro phenotype



In vivo phenotype



Gaston et al., 2021



Intragenic constructs with Myb or EPSP genes from strawberry

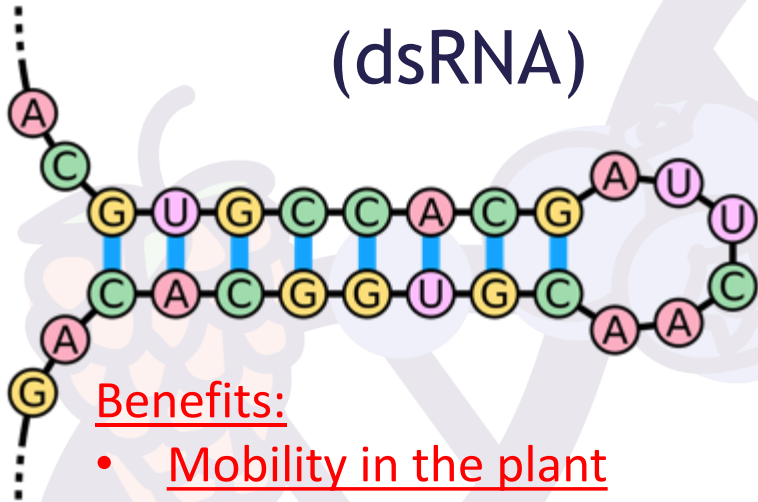


RNAi versus CRISPR/Cas9

GENE EDITING/CRISPR



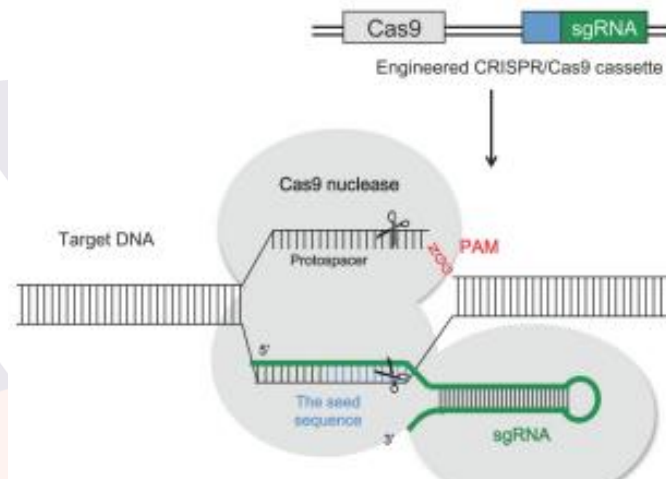
RNAi inhibits gene expression in a sequence-specific manner induced by double-stranded RNA (dsRNA)



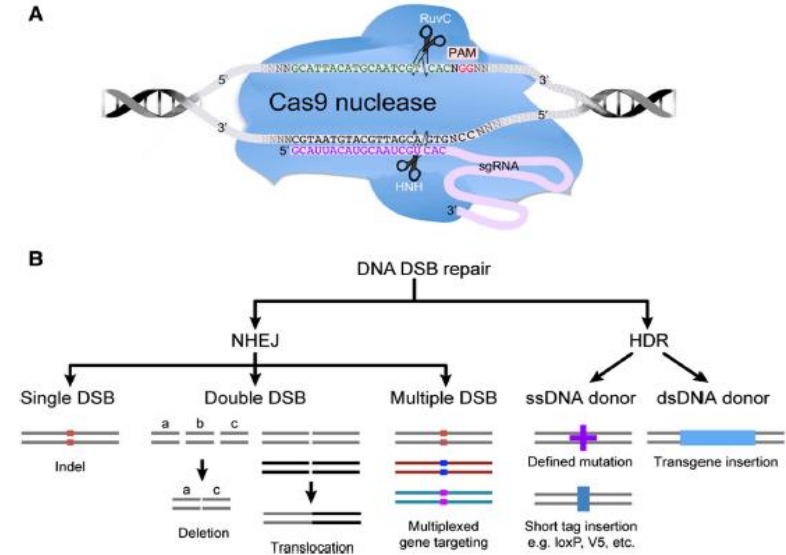
Benefits:

- Mobility in the plant
- Mobility between plant and target organisms
- Stable expression in planta
- Direct application - spray

Fire e Mello - Nobel prize for Medicine in 2006



Genome editing allows the integration, deletion, and/or mutation of genes of interest. In theory, replacing RNAi but only for cutting endogenous plant genes.



CAS9 protein is the tool for induce gene editing and it must be inserted in the DNA or in the cell and cuts the DNA. For this reason, according to the legislation, it is still a GMO.

Charpentier e Doudna Nobel prize for Chemistry in 2020



New Biotechnological Tools for the Genetic Improvement of Major Woody Fruit Species

Cecilia Limera¹, Silvia Sabbadini¹, Jeremy B. Sweet² and Bruno Mezzetti^{1*}

¹ Department of Agricultural, Food and Environmental Sciences, Università Politecnica delle Marche, Ancona, Italy; ² J. T. Environmental Consultants Ltd., Cambridge, United Kingdom



iPlanta

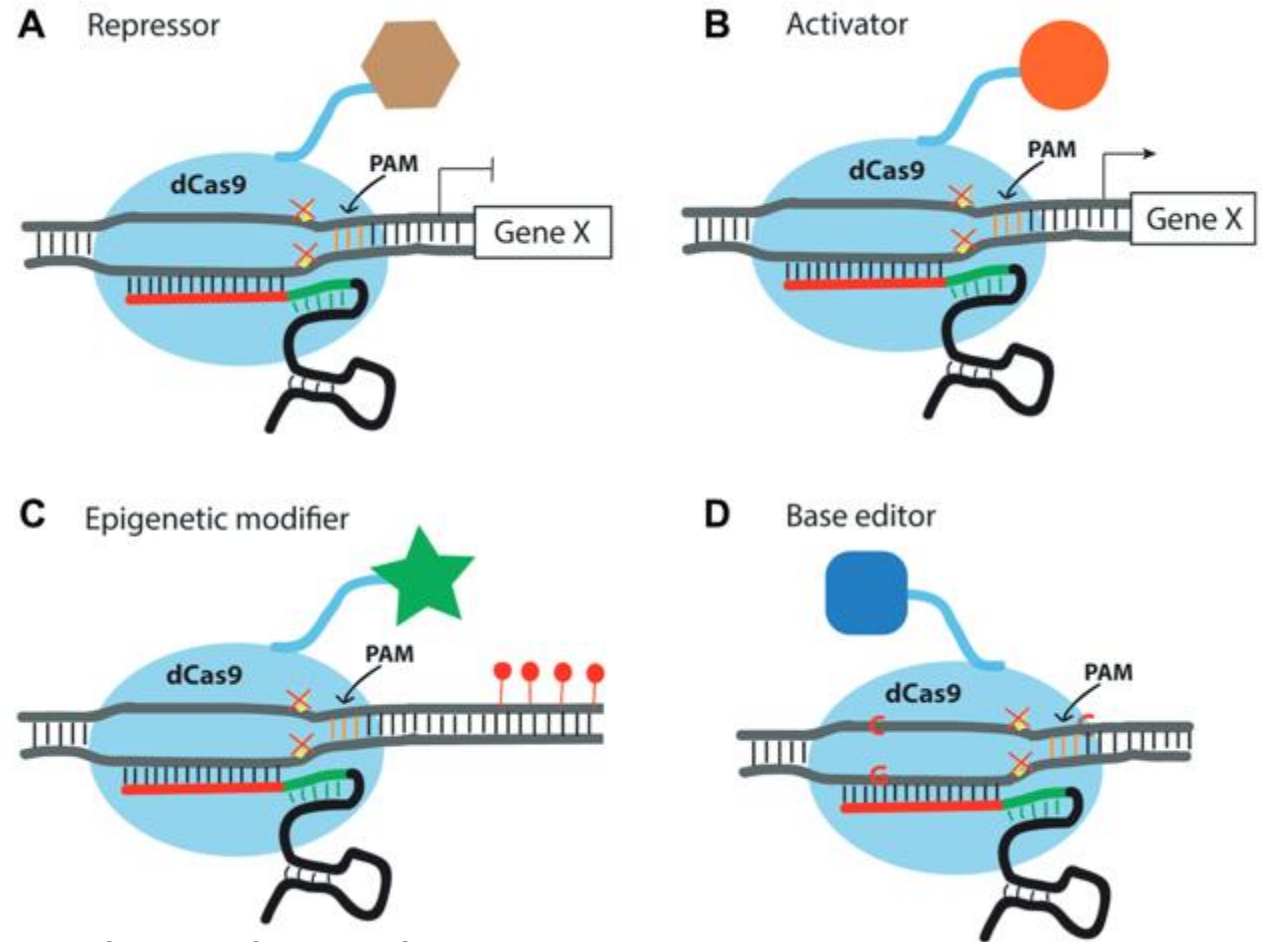
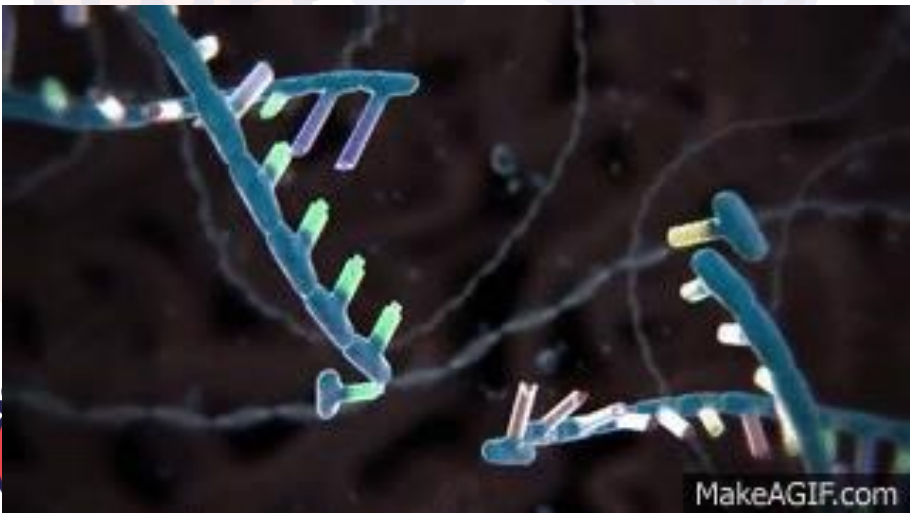


CRISPR/Cas9 mechanism

sgRNA that activates and guides Cas9 nuclease



DSBs mutation



Cas9 can be engineered:

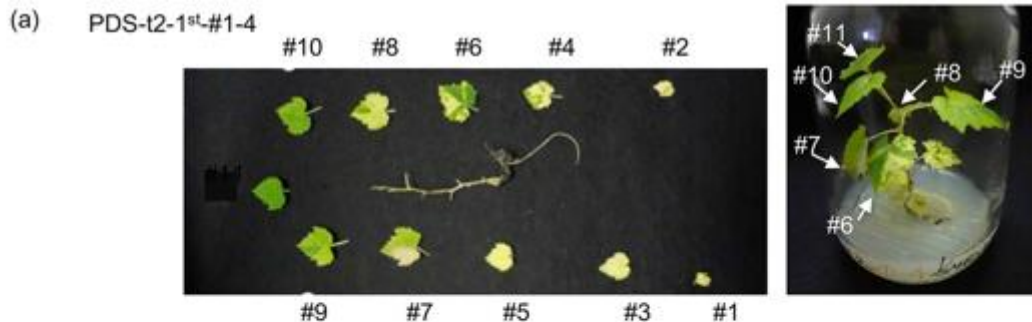
- By linking it to a transcription activator or repressor
- To convert one base pair to another without performing a DSBs (base editing)
- Epigenetic modifier

CRISPR/Cas9 applied to fruit tree species

Reviewed by Fizikova et al., 2021

a) Proof of concept

PDS mutation = Chlorophyll-deficient variegated plant



grapevine

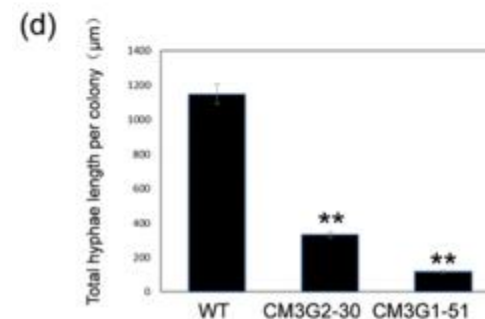
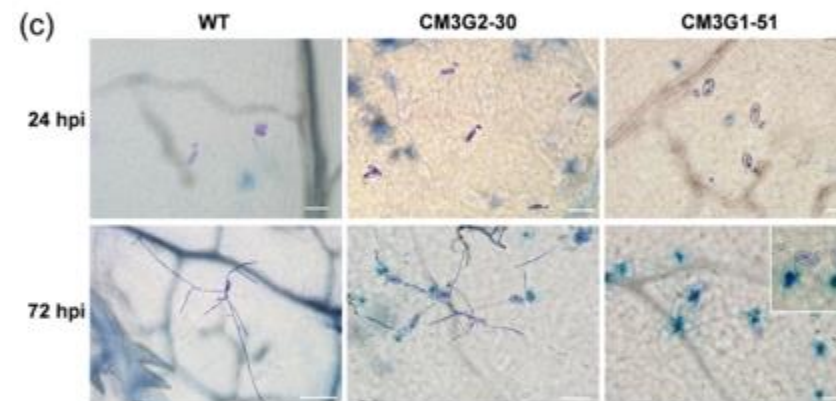
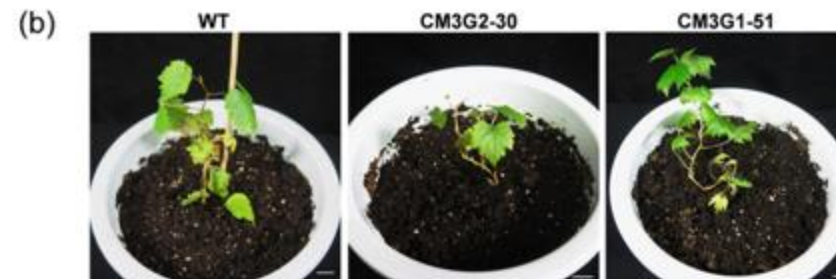
(b)

PDS-t2	GGCTGGTTTGTCTACTGCAAAAATATTGGCAGATGCAGGTCACAA	WT
#1	GGCTGGTTTGTCTACTGCAAA--ATTGGCAGATGCAGGTCACAA	(-2) 1/14
	GGCTGGTTTGTCTACTGCAA---ATTGGCAGATGCAGGTCACAA	(-3) 1/14
	GGCTGGTTTGTCTACTGCA----ATTGGCAGATGCAGGTCACAA	(-4) 2/14
#2	GGCTGGTTTGTCTACTGCAAA--ATTGGCAGATGCAGGTCACAA	(-2) 2/14
	GGCTGGTTTGTCTACTGC-----ATTGGCAGATGCAGGTCACAA	(-5) 1/14
	GGCTGGTTTGTCTACTG-----TTGGCAGATGCAGGTCACAA	(-7) 1/14
	GGCTGGTTTGTCTACT-----GGCAGATGCAGGTCACAA	(-11) 1/14
	GGCTGGTTTGTCTACT-----GCAGATGCAGGTCACAA	(-12) 1/14
	-----	(-135) 1/14



apple

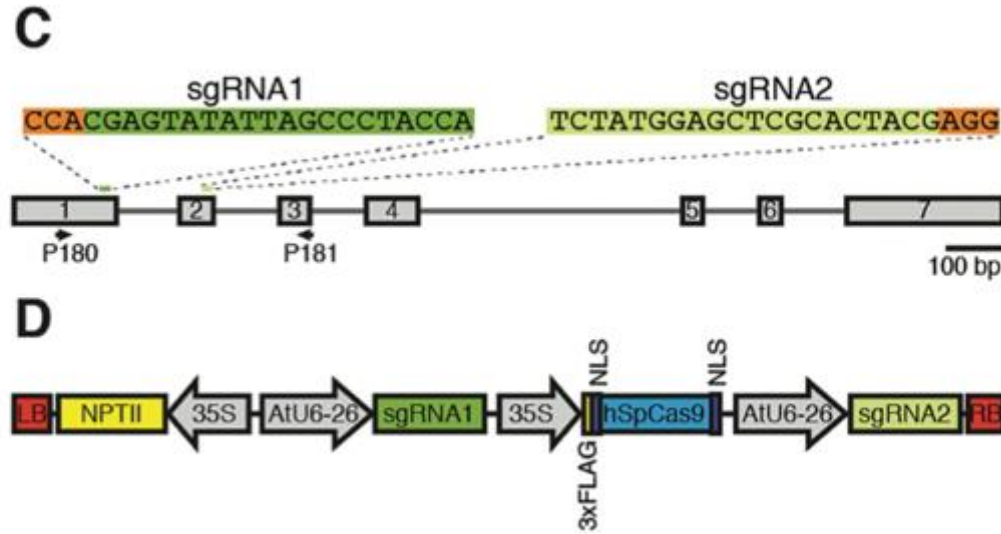
b) Pathogen resistance



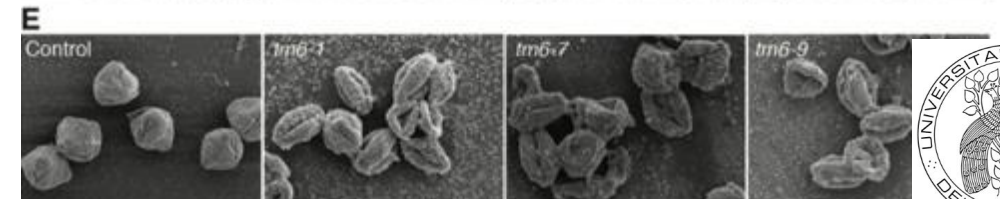
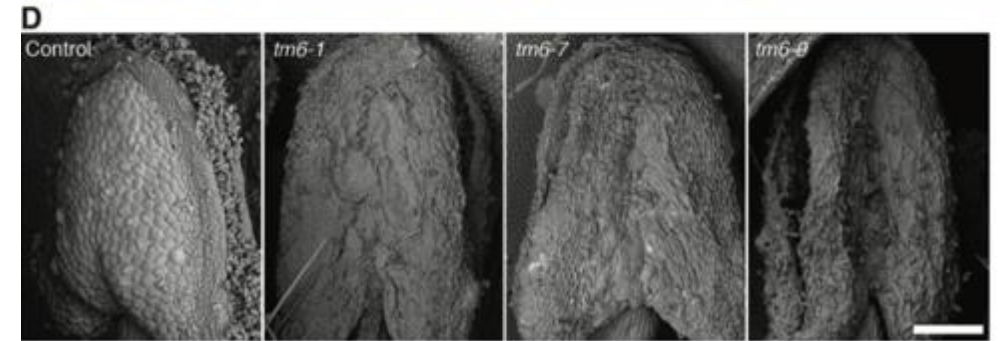
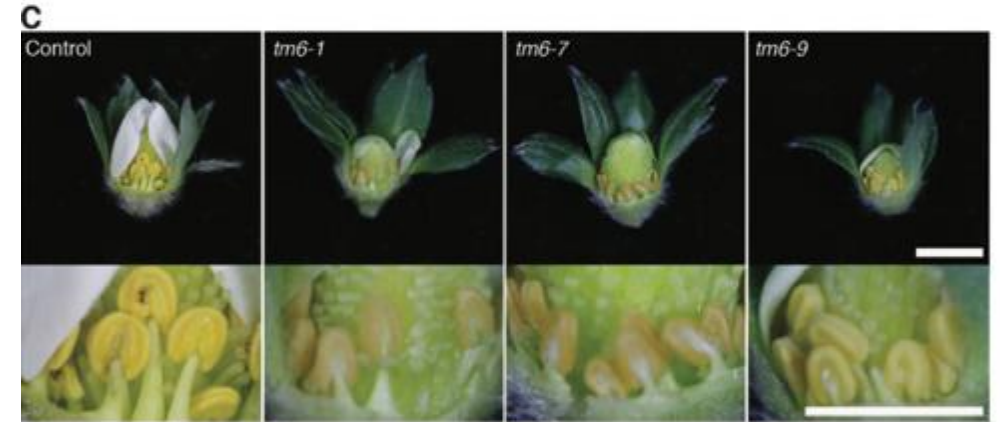
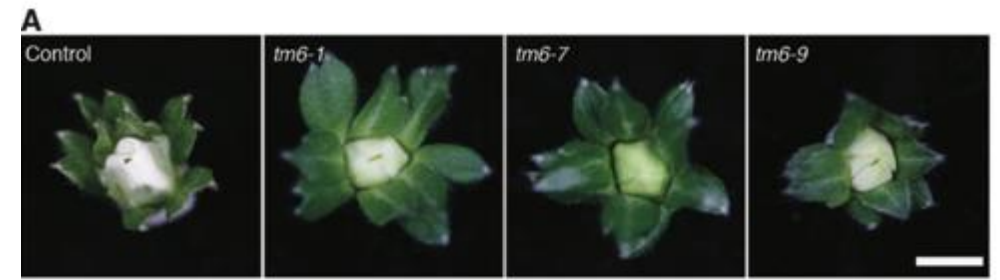
Knockout VvMLO3 results in enhanced resistance to powdery mildew in grapevine (*Vitis vinifera*)

c) Functional genomic studies

APETALA3 (AP3)
Mutation in *Fragaria x ananassa*



Phenotypic characterization of the mutant lines indicated that FaTM6 plays a key role in anther development in strawberry.



Plant flowering habitus

1. Isolation of *FT* genes from *F. vesca*, design and preparation of gene constructs (Gaston et al., 2021)



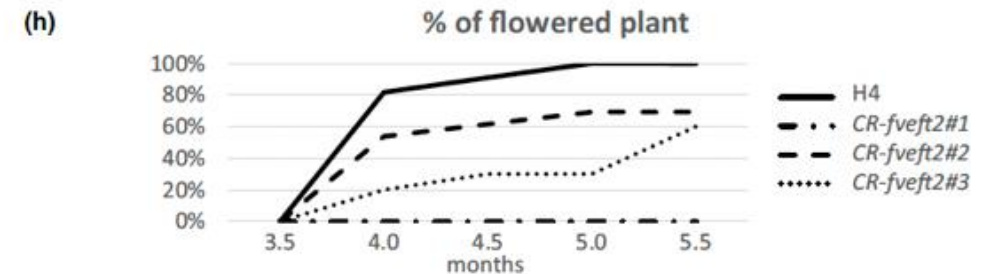
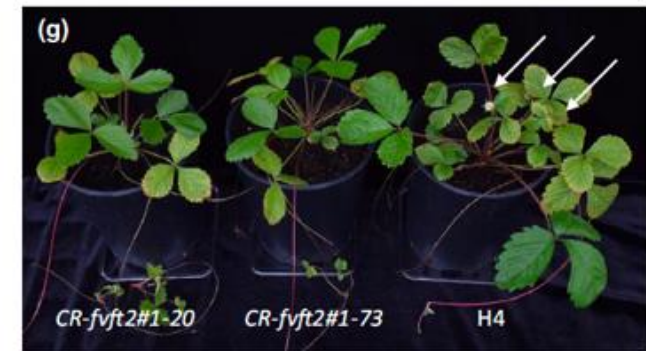
FveFT2 and *FveFT3* genes

2. **Overexpression** of *FveFT2* and *FveFT3* genes in Sveva and *FveFT2* in Romina and phenotypical characterization of transgenic lines in vitro an greenhouse environment

In vitro phenotype



3. **Genome editing** FT2 (Gaston et al. 2021)



RNAi for the control of Botrytis cinerea: more advanced phase for product development

GREENHOUSE/FIELD EXPERIMENTATION OF dsRNA FORMULATES



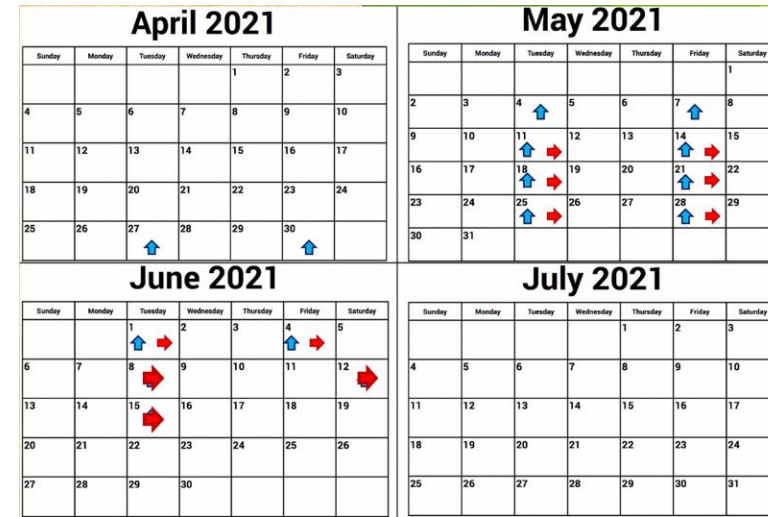
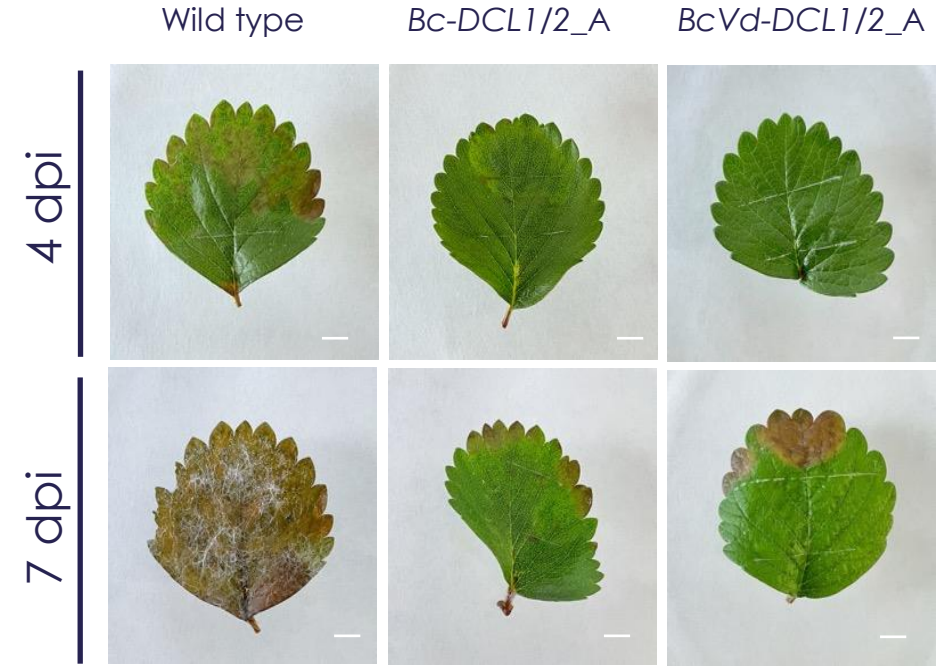
Test dsRNA formulations to identify the most effective ones

Compare dsRNA formulations with chemical and biological pesticides

Identify the level of protection of the dsRNA formulations in pre and post-harvest

Check for unexpected effects of dsRNA formulations

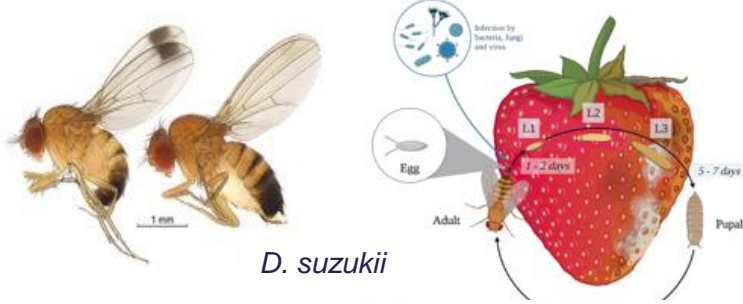
CONFIRMED THE EFFECTIVENESS OF EXPRESSION IN PLANTA: GMOs



RNAi strategy in octoploid strawberry to induce resistance to *Drosophila suzukii*

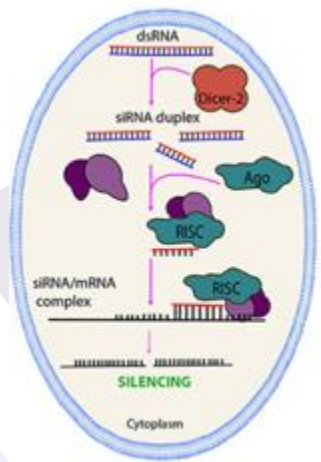


New emergency: *Drosophila suzukii*



D. suzukii

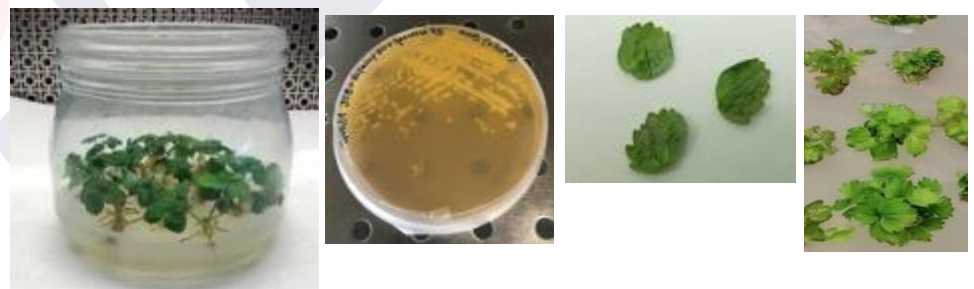
RNAi mechanism



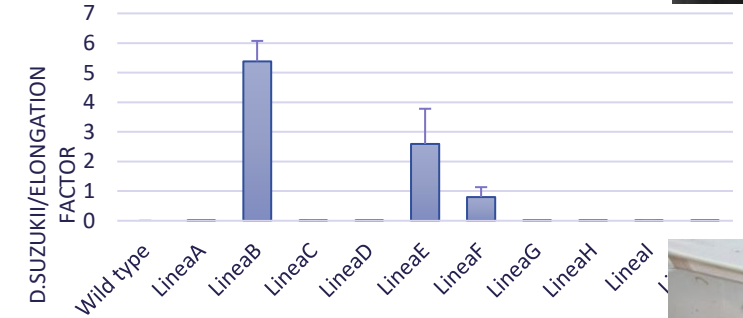
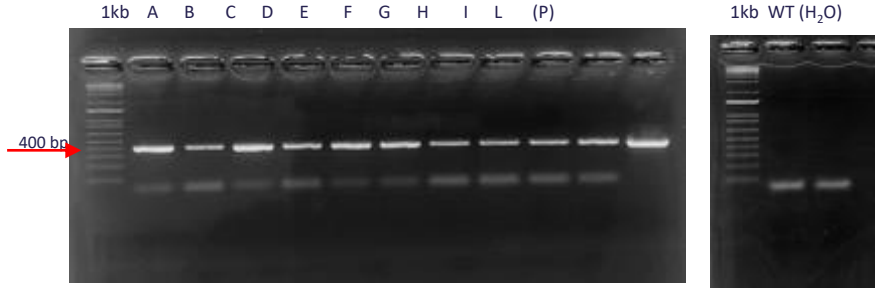
Harpin construct for *D. suzukii* gene target



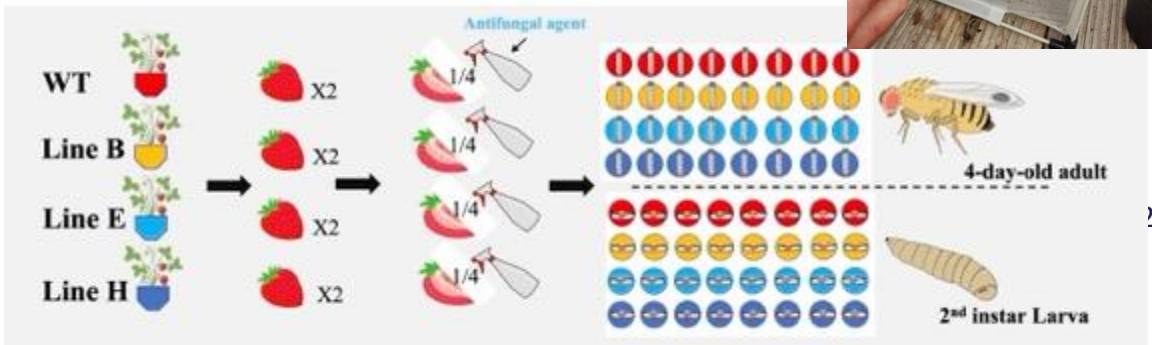
Agrobacterium transformation



Genomic characterization and gene expression



Infections to test the level of tolerance: Preliminary data to be confirmed in the coming season



TO BE OR NOT TO BE GMO

Transformation with exogenous sequences

GMO

Cisgenesis / intragenesis E

GMO EU Court (25/07/2018)

Plant with a GMO intermediate grafting

no GMO

Non GMO cultivar grafted on GMO rootstock

no GMO

Genome editing

GMO EU Court (25/07/2018)

Stable expression of GMO dsRNA in plant

no GMO

DsRNA-based products for direct application

no GMOs

**TO RESOLVE THE OPPOSITION AGAINST BIOTECHNOLOGIES IT IS IMPORTANT TO GIVE
CORRECT INFORMATION AND SHOW THE PLANTS**



Recent decision of the EU parliament

Note that it still needs to get approved by 2/3rds of the European Commission and then implementing regulations need to be written and implemented, so nothing will happen any time soon.

What did pass Parliament divides edited crops into two groups - those with the repair of a single double-strand cut (NGT1) - which are indistinguishable from mutagenesis, and those with more extensive edits (NGT2), such as a fragment deletion.

- NGT1 plants would still need to be labeled and listed (despite the lack of verification ability) and cannot be patented.
- NGT2 plants remain as GMOs, though an expedited review is planned.

Overall, I see continued dysfunction, not enablement, when it comes to getting edited products on the market.

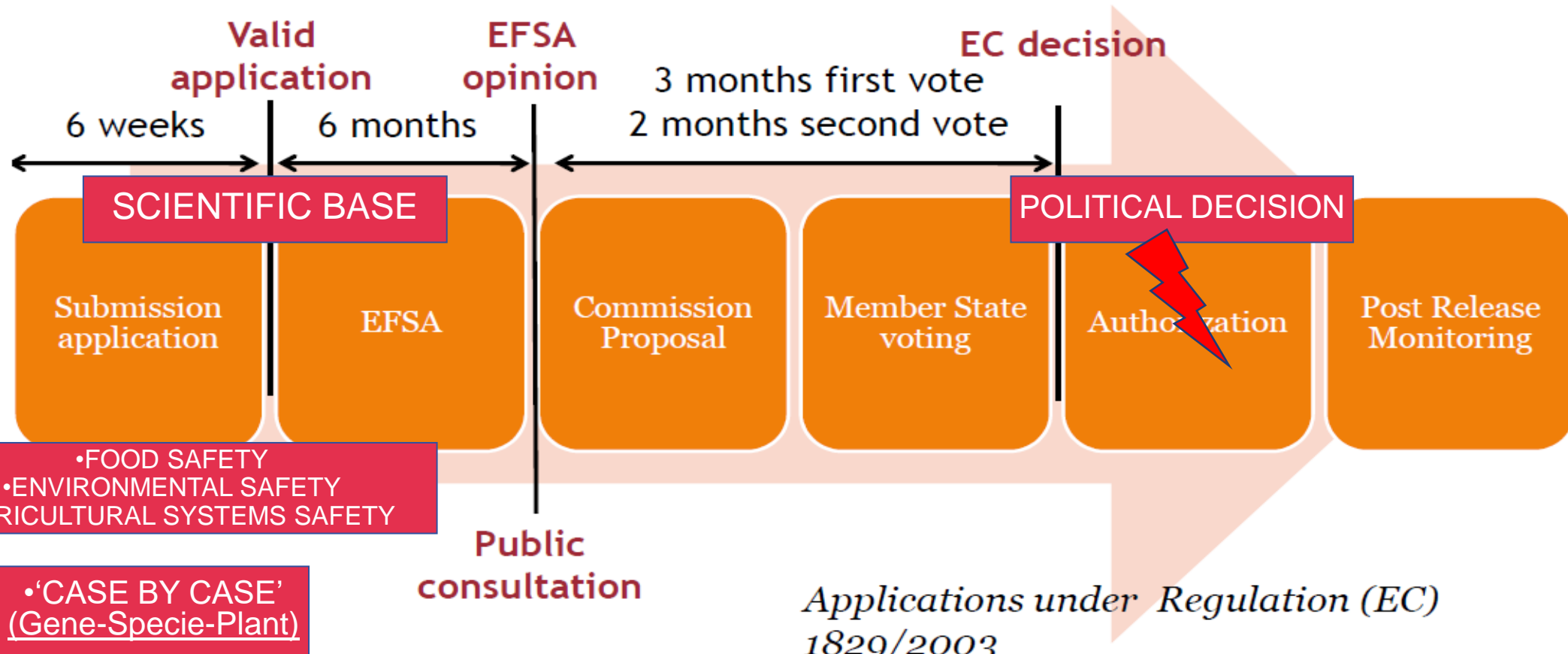
More at https://www.europarl.europa.eu/news/en/press-room/20240202IPR17320/new_genomic-techniques-meps-back-rules-to-support-green-transition-of-farmers.



EU regulatory system

1829/2003 - IMPORT More than 100 EVENTS -----2001/18 - CULTIVATION = MON810 ONLY

Approval process in EU for food, feed, import and processing



Proposal for actions:

- Union between **SCIENCE, PRODUCTION and POLICY** to deal appropriately and efficiently with climatic and health emergencies. Don't be afraid to say that they are GMOs and that they are ALL GMOs.
- Communication on **UNATTACKABLE SCIENTIFIC BASIS** to gain consensus. It is impossible to reach unanimity - the majority is enough - working on biotech is a life that I am in the minority.
- It is true that biotechnologies are in the hands of multinationals but for conveniences (corn, soy, ...), **there is still room for public research and for small businesses interested in applying biotechnology** to solve emergencies in fruit and vegetable plants. The blockade from the 'organic' world favors multinationals and blocks research at the local level.
- Only by overcoming these barriers can we think of using biotechnologies to find useful solutions to reduce the impact of the emergencies that destroy our production.
- RNAi technology offers the double possibility of obtaining new products / formulations for defense and resistant plants.
- To extend the discussion to different institutional levels, on the procedures to be followed to ensure the **immediate experimentation in a controlled environment and field of new resistant RNAi/Gene Editing/cis/intragenic plants**.
- The application of **2001/18** can only be of help in demonstrating the benefits, even for organic crops, and the absence of risk.
- IT IS FUNDAMENTAL THAT AGRICULTURAL ENTREPRENEURS AND ALL CITIZENS BE ABLE TO SEE **AND "TOUCH WITH HAND" THE RESULTS OF THE RESEARCH**



INTERNATIONAL AND NATIONAL PROJECTS



RESO: REsilience and SUSTAINABILITY of the fruit and vegetable and cereal supply chains to enhance the territories



BREEDINGVALUE: Pre-breeding strategies for obtaining new resilient and added value berries



Med-Berry PRIMA Project: Developing new strategies to protect strawberry crop in Mediterranean countries.



GOODBERRY: Improving the stability of high-quality traits of berry in different environments and cultivation systems for the benefit of European farmers and consumers



iPLANTA: Modifying plants to produce interfering RNA. OC-2015-2-20281



MIUR-PRIN2017: Small RNAs and peptides for controlling diseases and development in horticultural plants 20173LBZM2,





ALMA MATER STUDIORUM
UNIVERSITÀ DI BOLOGNA



CENTRO DE EDAFOLOGÍA Y BIOLOGÍA
APLICADA DEL SEGURA



C.I.V.V. Ampelos
Consorzio Italiano
Vivaisti Viticoli



D3A – UPM RESEARCH GROUP



**Prof. Mezzetti e
Prof. Capocasa
supervisors**



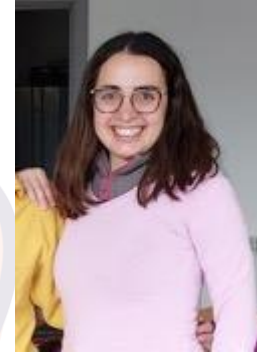
**Angela Ricci
Research Grant**



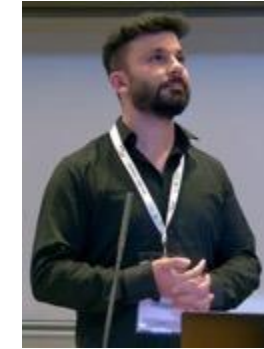
**Luca Capriotti
Researcher**



**Valeria Pergolotti
Research Grant**



**Rohullah Quaderi
Research Grant**



**Victoria Sunico
Research Grant**



**Giammarco
Giovannetti
PhD student**



**Silvia Sabbadini
Researcher**



**Luca Mazzoni
Researcher**



**Davide Raffaelli
PhD student**



**Federica Mecozzi
PhD student**



**Irene Piunti
PhD student**

