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Diversity of breeding technologies and impact for plant variety protection

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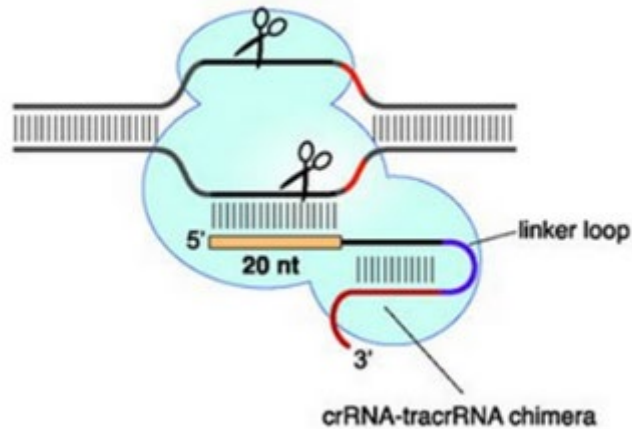
Key message

- The Essentially Derived Variety (EDV) issue is likely to be modified by the New Genomic Techniques (NGTs), their future use in breeding over a range of crops and the intellectual property regimes that are relevant in the domain

CRISPR/Cas method is today the dominant NGT technology

« Clustered Regularly Interspaced Palindromic Repeats » / « CRISPR Associated Protein 9 »

Cas9 programmed by single chimeric RNA



Major recent progresses

Technical and financial accessibility

Tool affordable to an increasing number of laboratories

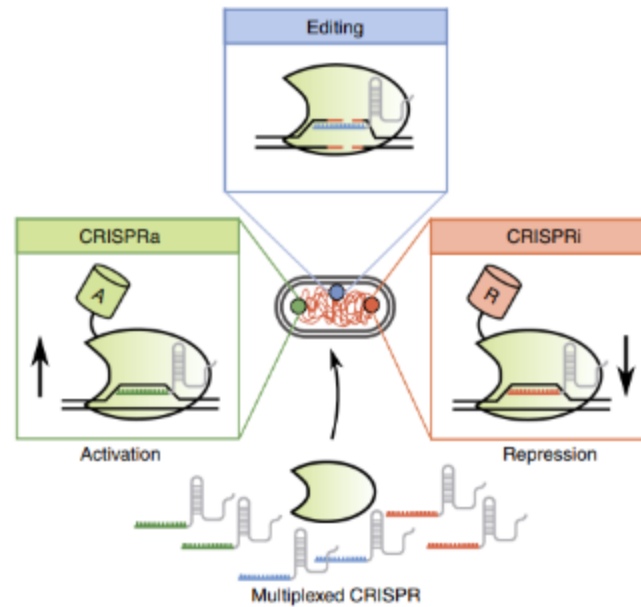
Possibilities of multiple targets editions

**Cas 9: nuclease with the highest number of studies and uses
(fundamental and applied research)**

Nombre de publications par année sur les méthodes CRIPR/cas9, TALEN et ZFN sur le site Pubmed



CRISPR/cas : recent technical progresses and use of this technology



(McCarty et al., 2020)

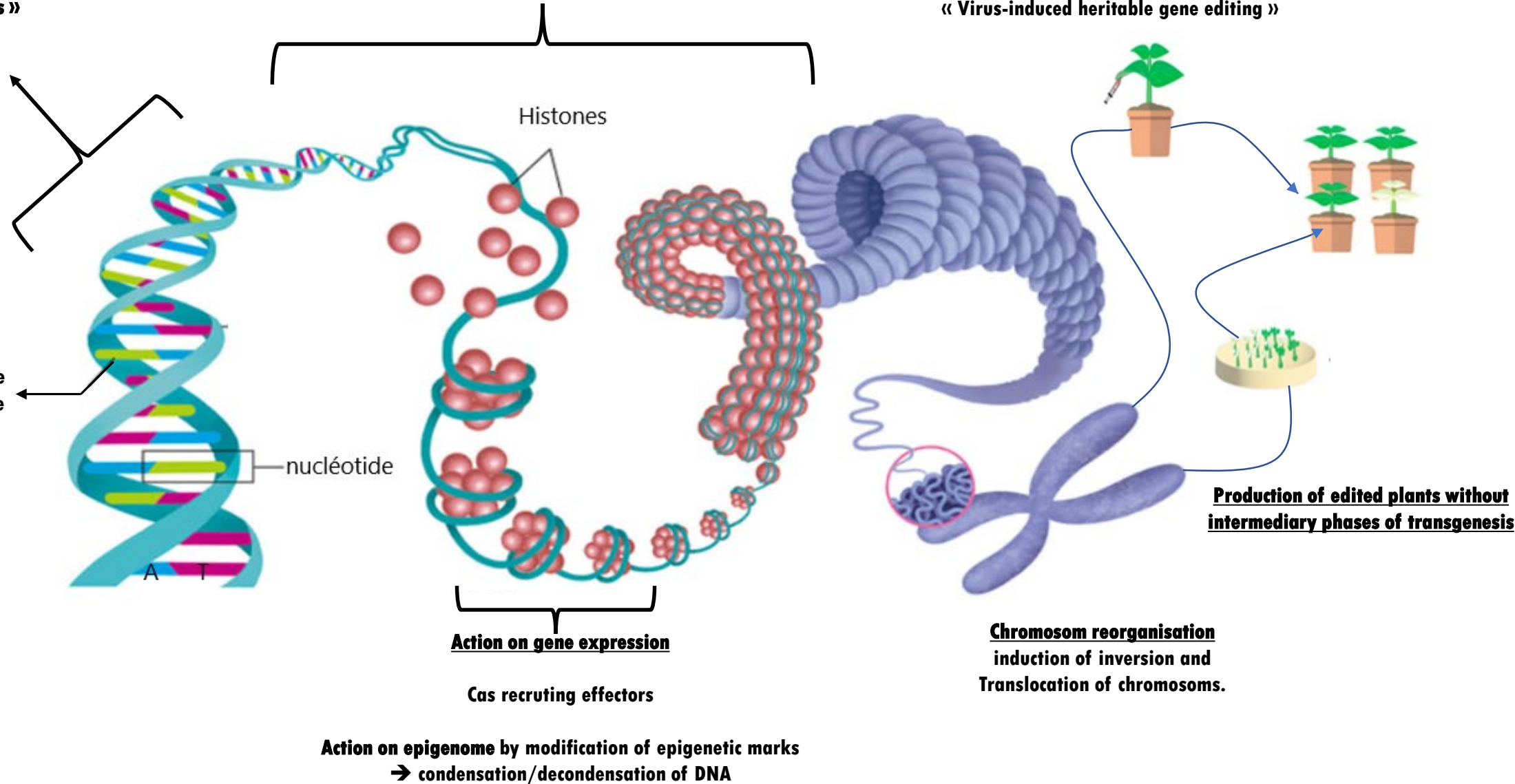
Technical advances

Multiplex modifications

To simultaneously target several DNA sequences. To modify expression of several genes of multigenic families.

More variability in PAMs motives +
« PAMless »

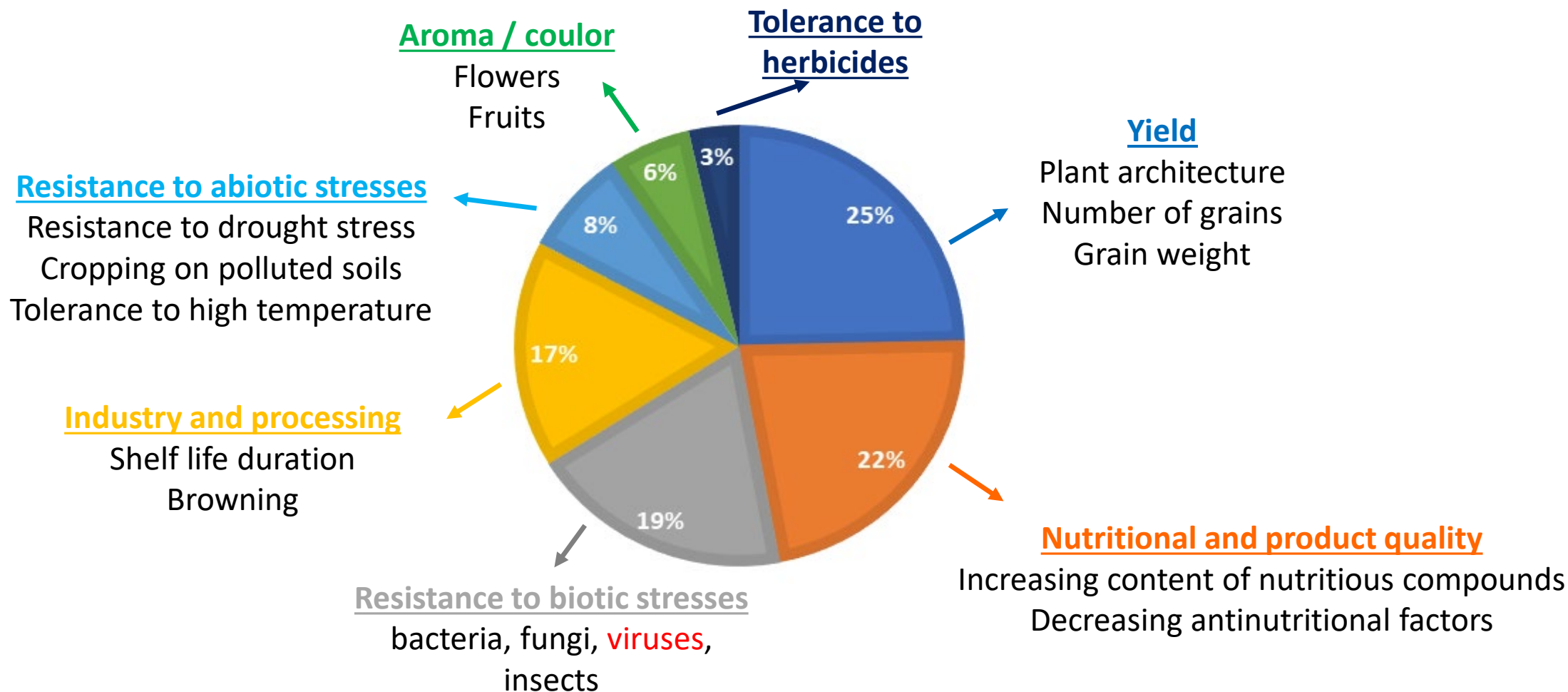
Base Editing and Prime
Editing methods to induce
modification of 1 or more
bases.



- **Promising tool whose all possibilities for plant breeding have not yet been documented**
 - **Accelerating innovation to foster agriculture sustainability**
 - **More accessible traits**
 - **New variation available**

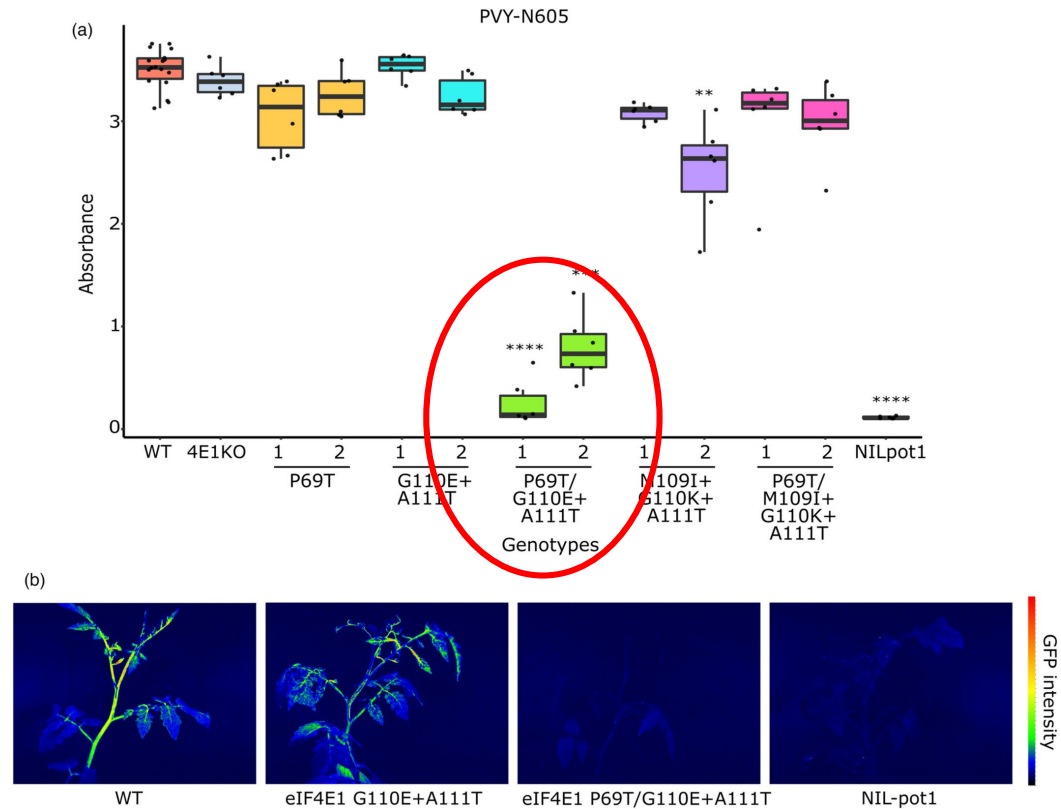
- Many proofs of concept of the ability to modify agroecological traits

According to scientific literature (WOS): use of genome editing techniques for plant breeding objectives



Example of gene-editing of tomato, copying eiF4E1 pepper allele. Inrae Avignon, France

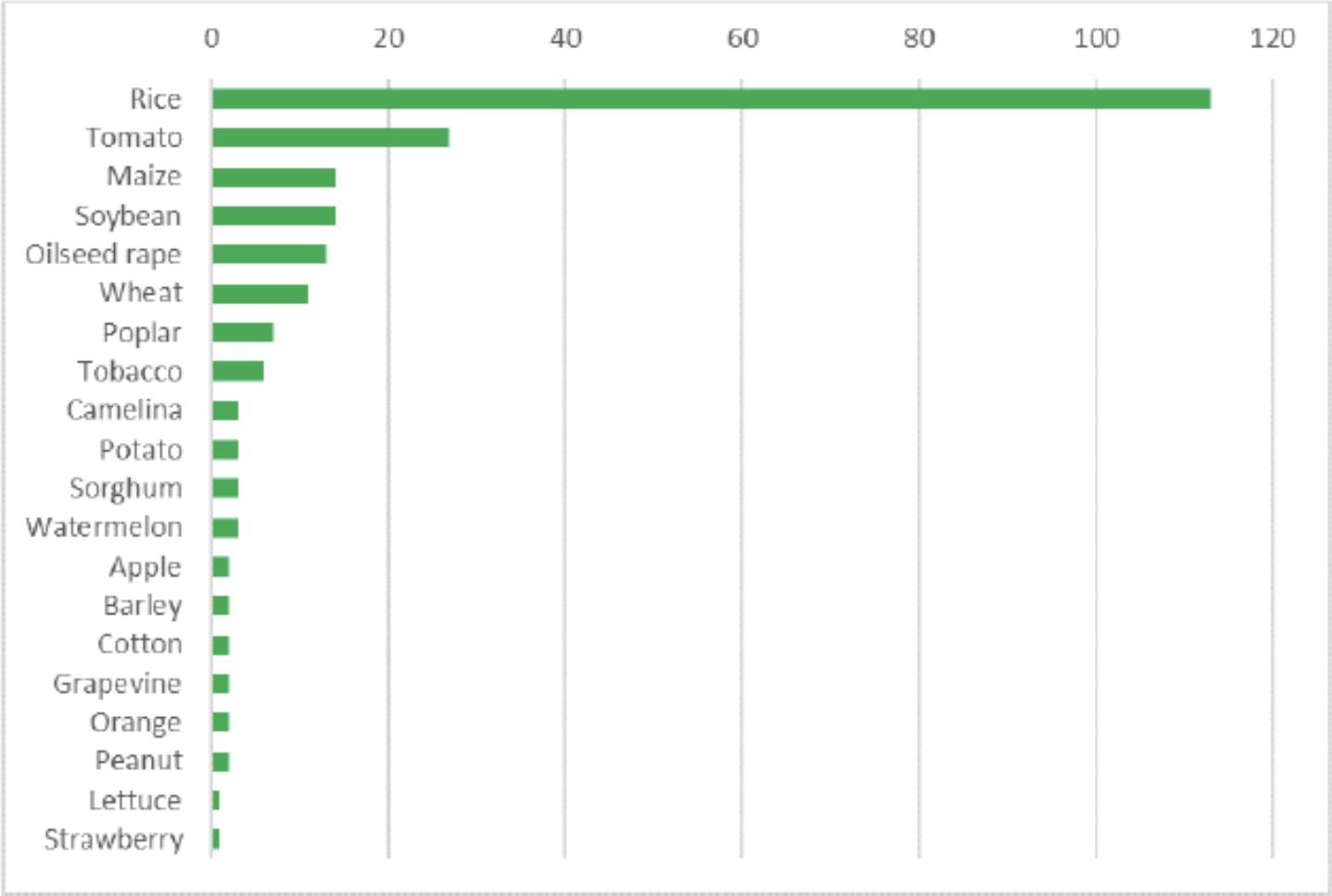
An iterative gene-editing strategy broadens eiF4E1 genetic diversity in *Solanum lycopersicum* and generates resistance to multiple potyvirus isolates



Kuroiwa K, et al, *Plant Biotechnology Journal*, First published: 30 January 2023, DOI: (10.1111/pbi.14003)

A limited number of species concentrates most efforts:

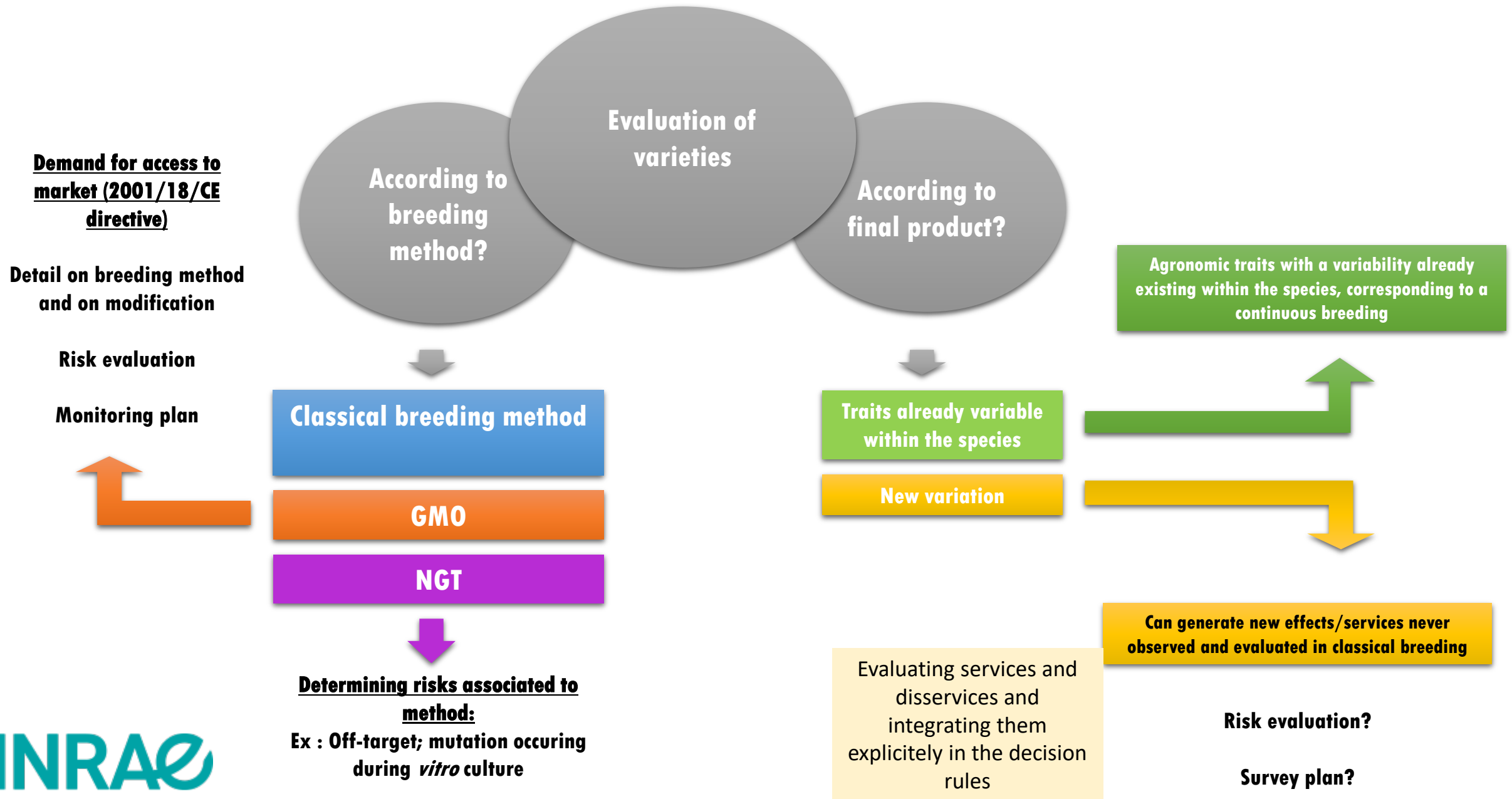
- Market size
- High-quality genome sequencing required
- Efficient *in vitro* regeneration



Number of scientific publications, with proofs of concept (in 2022)

Evaluating edited varieties

Evaluating edited varieties for their registration



Intellectual property issues and EDV

Access to genetic diversity

- Concerns regarding the accessibility to the genetic diversity, existing either in the current varieties under Plant Breeders Rights or in the genetic resource
 - Patents on edited traits are a key issue
 - The coexistence of the various intellectual property regimes is really at stake

Intellectual property

	Patent/technology	Patent/trait	P.B.R.
Technology	✓	✗	✗
Gene	✗	✓	✗
Variety	✗	✗	✓
Breeder exemption	✗	✓	✓
Farmer privilege	✗	✓	✓
Crossed licence	✗	✓ ↔	✓


NGT variety could be under P.B.R. and patent

For breeders


For the use of a variety carrying a **patented gene/allele**, the breeder must:

- Pay a licence right if the gene is preserved in the final variety
- Withdraw the gene and not to pay a licence right

Multiplicity of patented genes would create a minefield for breeders




Full transparency required on PI on varieties



More challenging work for breeders

Survey of patents databases (OEB, PINTO, ACLP, ILP, ...)



Patents on traits could be a source of rejection by the society

Essentially Derived Varieties

- EDV key issues (according to B. Kiewiet, 2006)
 - Retaining the expression of the essential characteristics of the initial variety
 - It conforms (essentially in the Basic Regulation) to the initial variety
 - EDV must be related to phenotypical characteristics and must be genetically heritable
 - EDV are determined as
 - Originating from an act of derivation
 - Phenotypically similar to the initial varieties except for the difference due to the derivation
- How does this fit with NGT?
 - The genome editing may go far beyond a simple/single modification
 - Regulation genes
 - Deep physiology
 - Multiplex possibility
 - **With NGTs progresses → high-throughput editing by a specialized operator on a variety protected by a PBR (*delivered to another owner*).**

- Integration of NGTs in breeding programs
 - Creating new genetic diversity to be used in breeding programs (*the induced genetic diversity may require the adaptation of many other physiological traits*). This is likely to be the situation for most edited traits (phenology, resistance to abiotic stresses, architecture).
 - In that case, the technology will be used early in the breeding programs and no EDV will be relevant
 - Providing extra traits in optimum genetic backgrounds.
 - In that case, EDV is relevant. This is likely to be the case for pest and disease resistance traits/genes

➤ **NGT: large possibilities**

- Precision, multiple targets
- Today, mainly proofs of concept
- Importance of technological sovereignty

➤ **NGT: some hurdles as not all species can be edited**

- Need of high-density genome sequencing
- In vitro regeneration

➤ **Variety evaluation**

- Variable traits : no need of modification of the evaluation pathway
- Disruptive traits : characterization of services and disservices at the scale of the crops and cropping systems

➤ **Traceability and transparency required for coexistence**

➤ **Intellectual property**

- Different possible regimes
- An important issue on patenting edited traits

➤ **Concerns**

- Acceptability: sharing benefits, access to technology, focussing on traits of societal relevance (climate change, agroecological transition)
- Genetic resources and diversity of cultivated species
- Boosting effort of public research

Conclusions

➤ **Technologies (cost, accessibility) that could speed up concentration of breeding companies**

- Today 6 companies= 50% of variety markets

➤ **Possible weakening of the PBR system** (which showed a tremendous efficiency since 1961 to foster genetic gains in all cultivated species)

➤ **Essentially Derived Varieties**

- Varieties carrying edited traits could be EDV, but...
- Edited traits or multiplex edition could modify expression of essential characteristics
- High-throughput edition could become a reality for some 'important' species

Thank you for your attention ...

