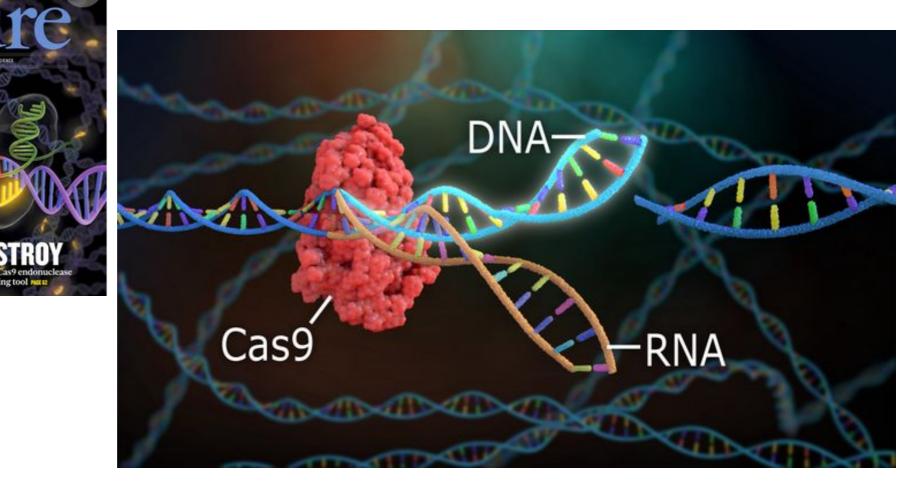
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New breeding techniques

Luigi Cattivelli CREA research centre for Genomics and Bioinformatics



Genome editing based on CRISPR-Cas9



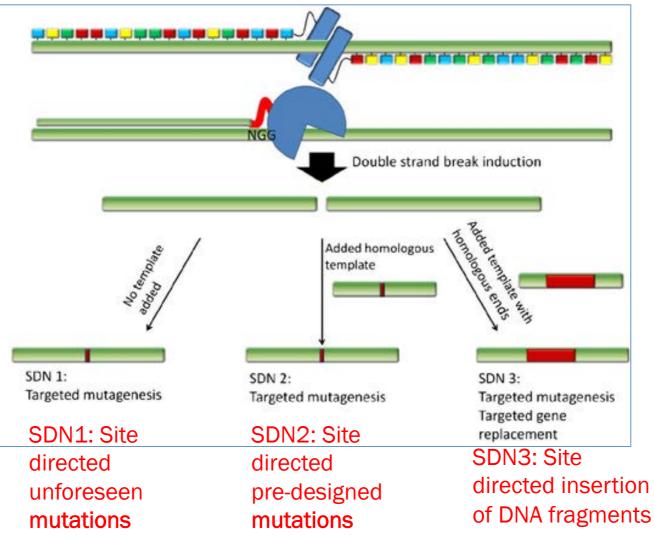
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in CRISPR/Cas9 genome-editing tool no



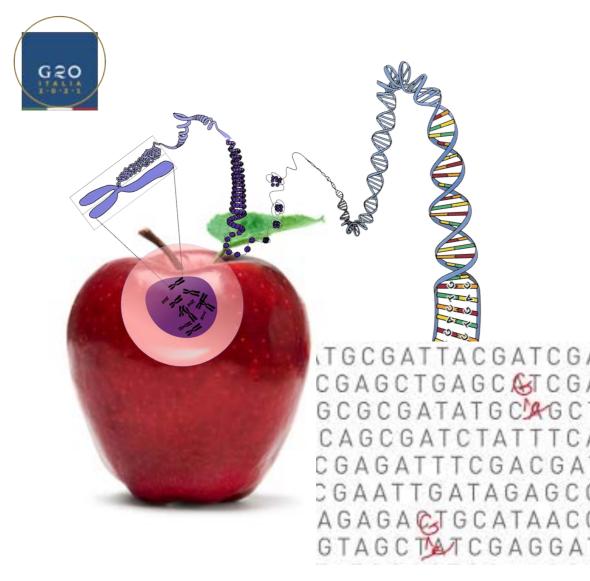
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CRISPR-Cas9: a simple technology for Site Directed



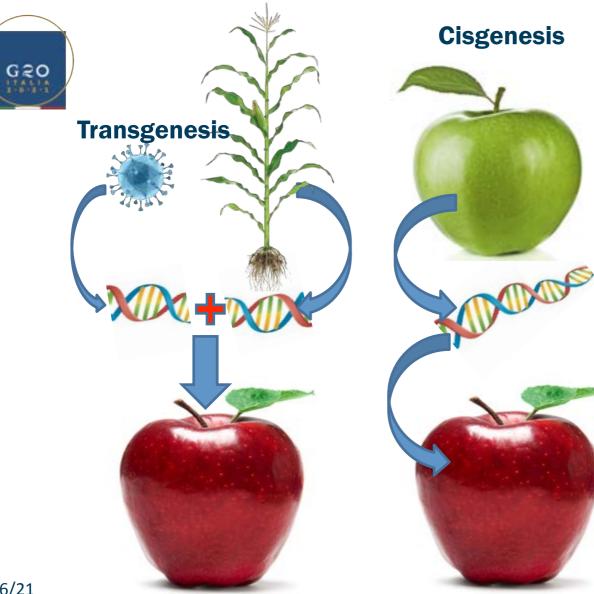
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Nuclease



Genome Editing

- Genome editing (SDN1 and SDN2): modify a gene within the target species (without inserting any external gene).
- The result is equivalent to what is obtained with mutagenesis
- The mutation concerns a single gene defined a priori
- After the mutation the editing mechanism (Cas19) is removed



Cisgenesis

- Cisgenesis: when transformation is limited to the transfer of whole genes between plants of the same species (or interfertile).
- The inter-species barrier is fully respected and in principle, the gene transfer could be obtained with a backcross

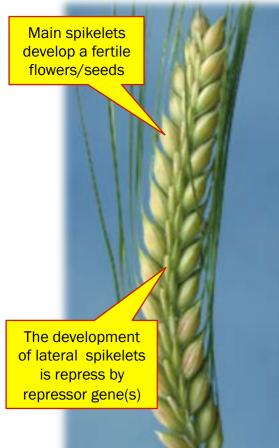


Is it possible to demonstrate a genome edited events?

Although existing detection methods are able to detect even small mutations, this does not necessarily confirm the presence of a Genome Edited product; the same alteration could have been obtained by conventional breeding (natural mutation, mutagenesis)



Natural mutants that could have been generated by genome editing: barley



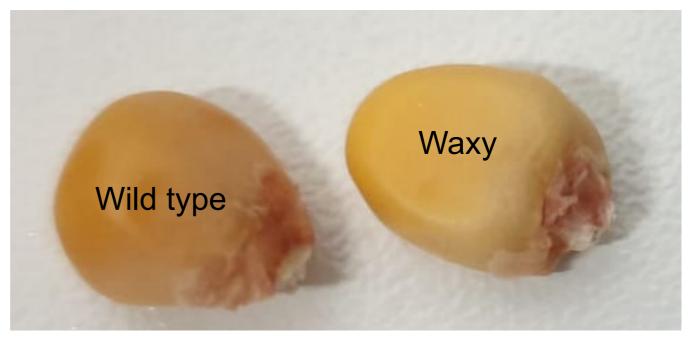
A single loss-offunction mutation transform two row barley (wild type allele) into six row barley (most of the barley grown today).





Natural mutants that could have been generated by genome editing: waxy corn

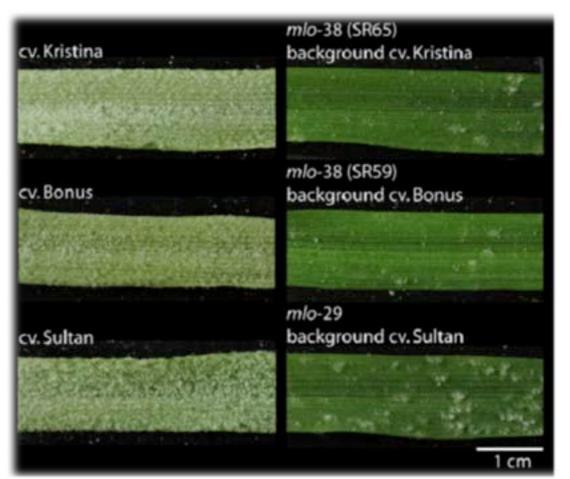
A knock out of the Granule Bound Starch Synthase gene leads to a maize waxy (with starch made of amylopectin only). Waxy mutants are known since 1908, many waxy-hybrids have been released. Waxy starch is easy to gelatinize and it is used in adhesives industry. **Corteva has made a waxy corn with genome editing.**





Natural mutants that could have been generated by genome editing: powdery mildew resistance in barley







Natural mutants that could have been generated by genome editing: powdery mildew resistance





- A knock-out mutation in the barley **mlo** gene confers resistance to powdery mildew
- Many mlo mutants have been selected after artificial mutagenesis, but are also known natural *mlo* mutants
- Powdery mildew exists in many plants and in all species there is an *mlo* gene homologs
- For example, a mutation induced by Genome Editing in the *mlo* gene of grapevine, bread wheat, strawberry, etc. leads to plants resistant to powdery mildew

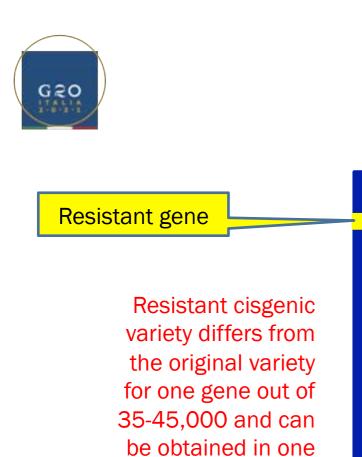




Cisgenesis transfers the resistant gene only

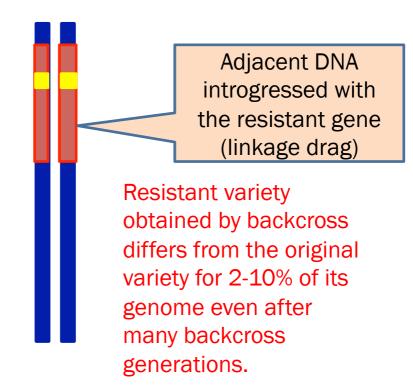
Backcrossing wild resistant genes on cultivated vines is possible, but a large genome region is transferred

GRO



single generation

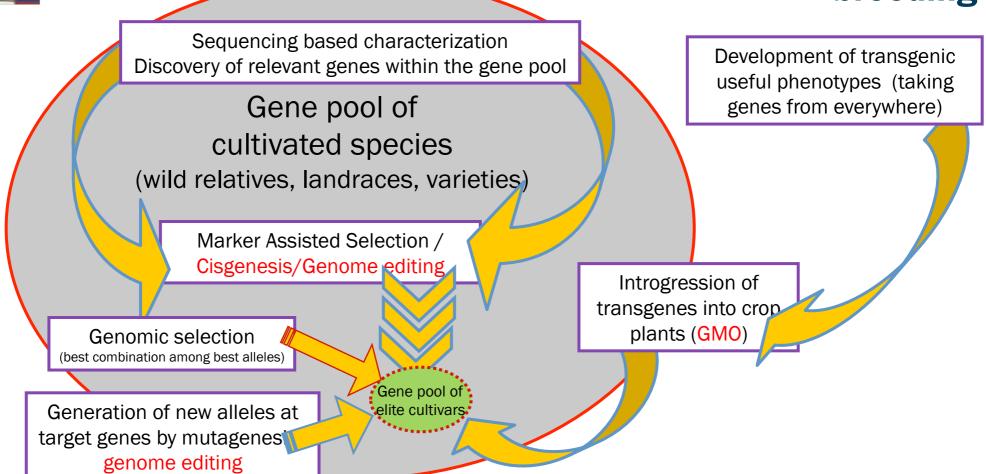
Backcross or cisgenesis?



Cisgenesis varieties are identical to the original variety (except the introgressed gene) e.g. cisgenic grapevine for a disease resistant gene have the same quality traits of the original one

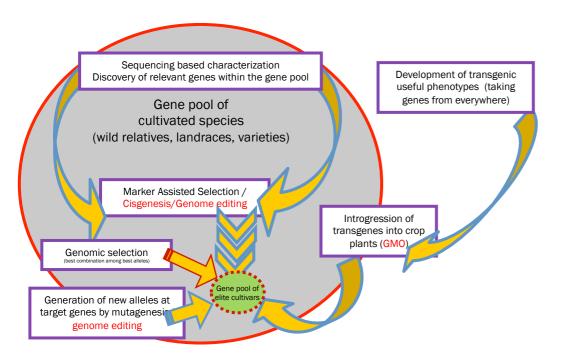


The new breeding technique in the contest of plant breeding





The new breeding technique in the contest of plant breeding



The impact of the New breeding techniques should be seen in the global contest of all breeding tools and strategies (diversity, sequencing, markers and gene mapping, mutants, NBT, crop modelling, ...)

A global coordination of the research aiming to enhance synergies could be useful.

Wheat Initiative is an example.





New breeding techniques have been proven to be very useful for selecting the plants for the future
Mutations in specific genes may confer very useful phenotypes and they are fully equivalent to natural or chemically induced mutations

 It is a fact that regulatory frameworks of genome editing differ from one country to another and a debate is ongoing in several countries on the role and regulation of genome edited plants

Concluding remarks and points for the discussion