

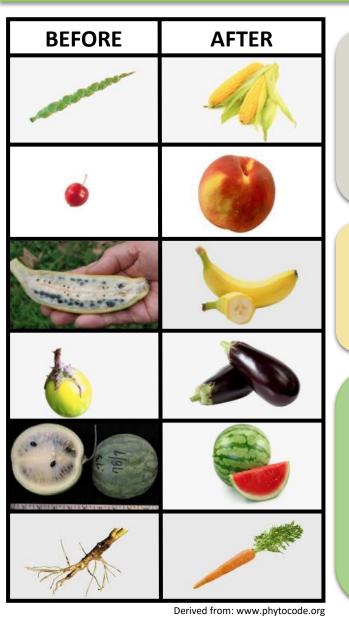
Canada



Forage Breeding for the Future

Stacy Singer, Ph.D. November 15, 2018

Plant breeding – a new thing?



- Around 13,000 years ago, humans began their own 'breeding program' that transformed wild plants into domesticated crops that better suited their needs
- The food that we grow and eat today did not always look and taste like it does now!
- Humans actively interfered with and manipulated crop evolution through selection
 - initially inadvertent
 - a very slow process
 - allowed improvement of numerous traits

New trait = genetic change

- Involved the 'improvement' of traits that were important at that time:
 - more robust plants
 - enlargement of fruits and/or grains
 - better flavour
 - synchronized flowering time
 - loss of seed dispersal



- Traits that need improving today are very different driven by changing human needs and agricultural conditions
- Each new trait derives from a genetic change

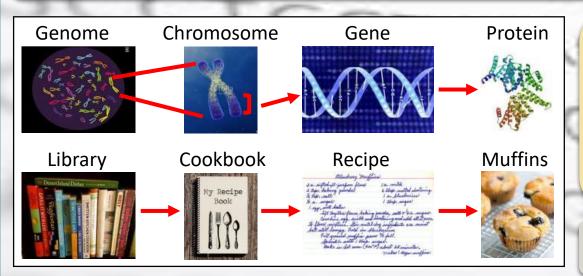


We've been tweaking the genetics of our favourite crops for millennia!

Source: www.gmoanswers.com

A little bit about genetics.....

- Every cell in a given individual contains the same DNA
 made up of 'alphabet' of four nucleotides A, T, C and G
- A sequence of these four 'letters' makes up a gene
 - only difference between different genes is the order of the four nucleotides
 - each gene has a different 'meaning' they translate into different proteins, which are responsible for different traits
- Many genes are placed alongside one another to form chromosomes
- Living creatures have thousands of genes that together make up their genome



- A genetic change (mutation) can mean the substitution of one nucleotide for another, or bigger deletions, insertions or rearrangements – occur naturally!
- Effect of mutation = new trait

History of crop breeding

'Non-GM':

Selective breeding:

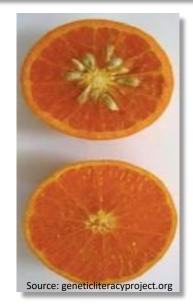
- ~ 13,000 years ago
- Select plants with naturally occurring mutations

Cross-breeding:

- ~ 300 years ago
- Cross plants to obtain desired genetic trait

Mutagenesis:

- ~ 80 years ago
- Use chemicals and radiation to induce mutations in plants



Almost everything we eat

e.g. pears, apples, grapefruit, rice, mint and some bananas

'GM':

Transgenics/cisgenics/RNAi:

- ~ 35 years ago
- Transfer gene from any organism into crop

e.g. corn, cotton, papaya, soybeans, squash, canola, alfalfa



'GM'/'non-GM':

Genome editing:

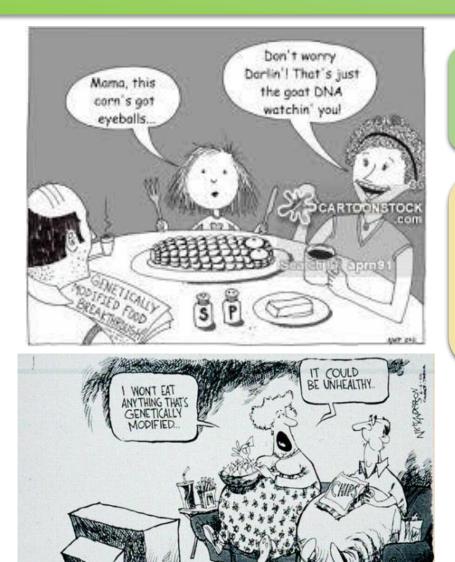
- ~ 10 years ago
- Precise genetic changes

e.g. canola, more to come soon!



All of these measures, whether 'GM' or 'non-GM', involve genetic changes!

Let's talk about GMOs.....



- Many misunderstandings and misconceptions about what GMOs actually are.
- Umbrella term that covers a range of very different molecular breeding technologies, which in some cases yield genetic changes that are no different than what you could achieve using conventional breeding ('non-GM').



Yet another unauthorized experiment in genetically-modified food

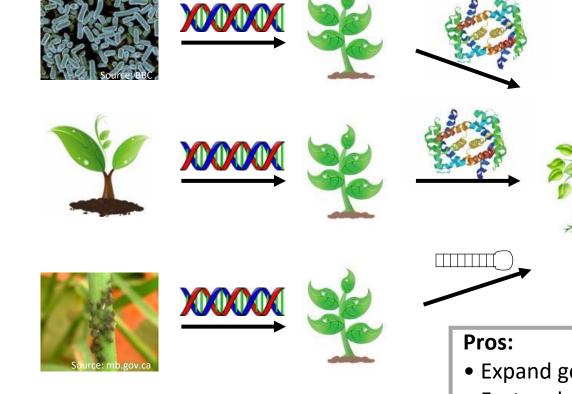
Anderson / Courier Journ

Transgenic, cisgenic, RNAi (GM)

Transgenic 'Foreign' proteincoding gene

Cisgenic Protein-coding gene from close relative

RNAi Non-protein-coding DNA



These methods of achieving genetic changes are very different!

Should be considered based on target gene, source, and whether it produces a new protein (toxicity, allergenicity)

- Expand gene pool
- Fast and easy
- Specific

Cons:

- Acceptance low (GM)
- Safety/environment

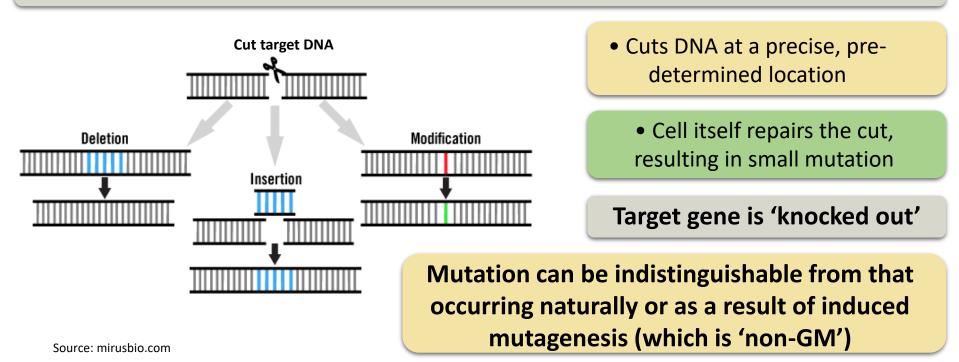
Genome editing (GM/non-GM)

A new breeding technology – a tool that has the potential to take breeding to a new level

Precision 'gene surgery'



• Introduce protein that acts as molecular scissors and is targeted to a specific genetic site



Genome editing: GM or not?

Increasing number of crop species altered using genome editing that the USDA has stated will not be regulated

Source: Pacher and Puchta (2017) Plant J 90:819-833



Canola:

 herbicide resistant (Cibus)
 already on market in Canada and USA



Soybean:

- high oleic acid (Calyxt)
- low linoleic acid (Calyxt)
- stress tolerant (USDA)



Mushroom:

 non-browning (Penn State)

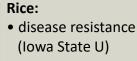


Corn:

- waxy (Dupont)
- high yield (Benson Hill)
- low phytate (Dow)
- disease resistant (Dupont)



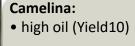








- non-browning (Calyxt)
- low acrylamide (Calyxt)





- Alfalfa: • low-lignin (Calyxt)





• low-nicotine (NCSU)

Tobacco:

Green foxtail: • late flowering (Donald Danforth Plant Science Center)

Flax:

- herbicide resistant (Cibus)
 expected to be on
 - market by 2019

Genome editing: GM or not?

Globally, things are less clear, but discussions are ongoing

Lack of cohesion in terms of regulatory policies surrounding gene edited crops:

- US, countries in South America, Japan will not regulate, not 'GM'
- EU will regulate, are 'GM'
- Canada regulates depending on the trait, could be 'PNT'
- Many countries are still undecided

What constitutes a 'GM' crop????

Cost and time for de-regulation of a 'GM' crop is prohibitive, and therefore often only large corporations are capable of commercialization



It is essential that we develop coordinated, efficient, fact-based and product-focused regulatory systems to avoid loss of innovation in agriculture and delays in the diffusion of these new technologies

How do we apply this to forages?

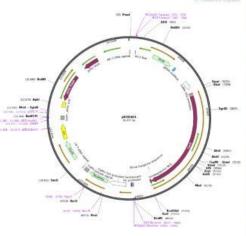


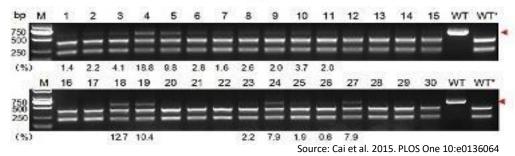
Cource: passel.unl.edu

We apply molecular genetic techniques and biotechnology to advance the pace and ease of forage improvement

Perennial legumes

- Alfalfa
- Sainfoin









Forage traits for today

1) Population growth



2) Climate change

Estimated that we need to increase food production by 70% in the next 40 years to keep up with population growth

Source: fao.org

In warm regions, crop yields drop ~3-5% with every 1°C increase in temperature

Source: Burney and Ramanathan 2014

Improve biomass yields
Improve stress tolerance
Increase leaf lipid content
Improve digestibility
Improve pest resistance
Improve disease resistance
Improve nutrition
Reduce health issues
Reduce fertilizer use

ource: ag.ndsu.edu

Source: noble.org

We need to come up with ways to grow more forages that are more efficient with better nutrition on less land in a harsher environment – in a sustainable and safe manner

Finding new sources of genetic variation

- 1) Identify candidate genes that when modulated might elicit the desired trait
 - from previous studies in other plant species
 - from alfalfa's wild relatives, which although not agronomically useful, often possess certain beneficial traits



Drought treatment:





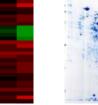
Wild relatives

Cultivated alfalfa

Assess plants for traits of interest

- drought tolerance
- salt tolerance
- pest resistance





Proteomics

Protein



Metabolomics Metabolites

Use comparative 'omics' approaches to identify candidate genes

 find out what the genetic basis of these beneficial traits are

Genomics DNA

Transcriptomics **RNA**

Validating candidate gene function

- 2) Validate the function of identified candidate genes (or those identified previously in other plant species) using transgenic and RNAi approaches
 - proof-of-concept work





Introduce genetic cassette into alfalfa

• use Agrobacterium-mediated transformation

Confirm presence of genetic cassette



Assess plants for trait of interest



Developing germplasm – genome editing

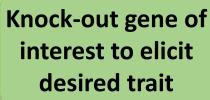
3) Use genetic knowledge to generate improved alfalfa germplasm

Use CRISPR-Cas9-mediated genome editing

- 1) Can we increase efficiency?
- 2) Can it be used directly in adapted cultivars?
- 3) Can we achieve it without the initial use of a transgene?



Source: Li et al. 2016. Front Plant Sci 7:12217







Developing germplasm - TILLING

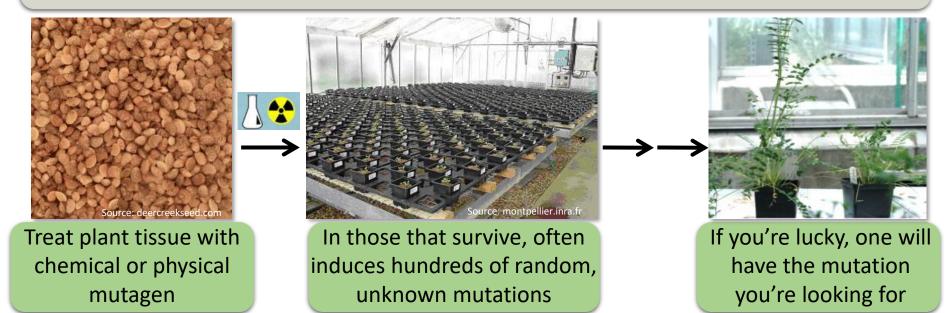
Screen candidate genes for mutations in mutagenized alfalfa population (non-GM)

Targeting Induced Local Lesions IN Genomes (TILLING)

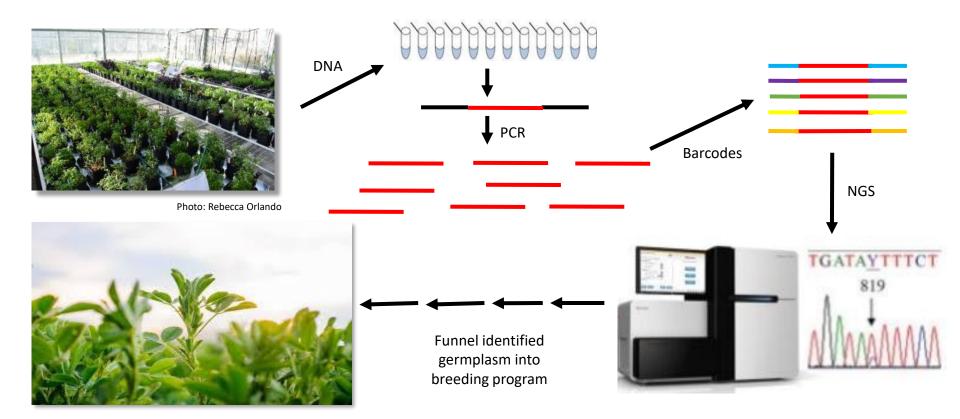


First step is to generate a large mutagenized population (3000-5000 plants)

• we use ethyl methanesulfonate (EMS) as the mutagen



Developing germplasm - TILLING



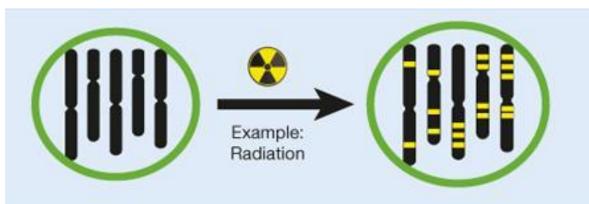
• Self-incompatibility of alfalfa makes this a bit more challenging

- however, it's being done in other self-incompatible plants, including radish and forage grasses

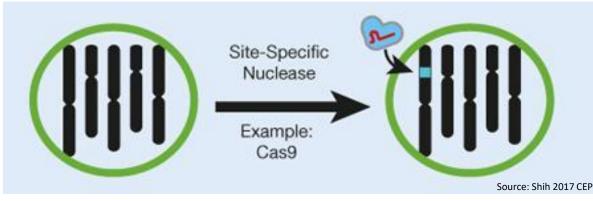
Kohzuma et al. 2017. Breed Sci 67:268 Manzanares et al. 2016. N Biotechnol 25:594

Mutagenesis vs. genome editing

Induced mutagenesis (non-GM):



Genome editing (GM or non-GM?):





Together with conventional breeding, molecular breeding technologies have the potential to be instrumental in a sustainable future of forage and livestock production

Acknowledgments

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Thank you!

