

Breeding for freezing tolerance in plants

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Agriculture and Forestry



Outline

- Where I am from
- Freezing tolerance in plants
- Reverse genetics in winter wheat
- Association mapping in perennial ryegrass
- PPP pre-breeding project
- Future targets



LITHUANIAN RESEARCH CENTRE FOR AGRICULTURE AND FORESTRY (LRCAF)

was established in 2010 as a merger of three related research institutions:

- Institute of Agriculture
- Institute of Horticulture
- Institute of Forestry

Research staff: 183

PhD students: 58

Total staff: 636

Budget: 12 M€



HISTORY OF THE CENTRE

1911 Agricultural school was established in Dotnuva.

1919 Dotnuva Agricultural college was set up.

1922 Breeding Station was founded in Dotnuva.

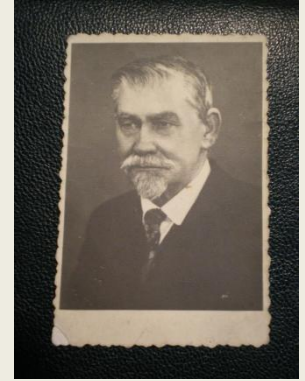
1923 Dotnuva experimental station started operating.

1924 Academy of Agriculture was established in Dotnuva,
transferred to Kaunas in 1947.

1927-1960 A network of experimental stations was established.

1956 Lithuanian Institute of Agriculture was established in Dotnuva.

2010 Lithuanian Research Centre for Agriculture and Forestry was
formed



D. Rudzinskas



LABORATORY OF GENETICS AND PHYSIOLOGY

Staff:

5 researchers
1 PhD and 2 Bsc students
3 technicians

Research:

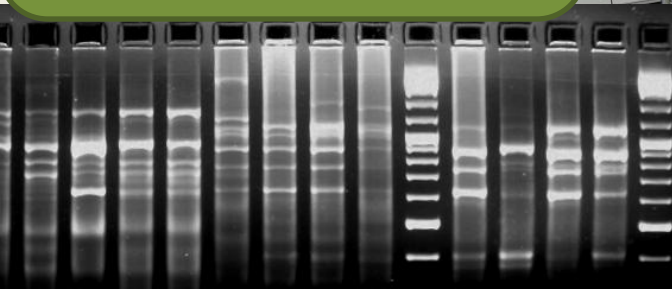
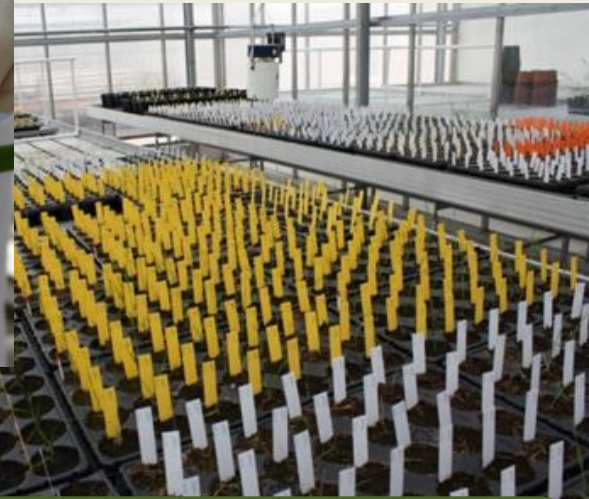
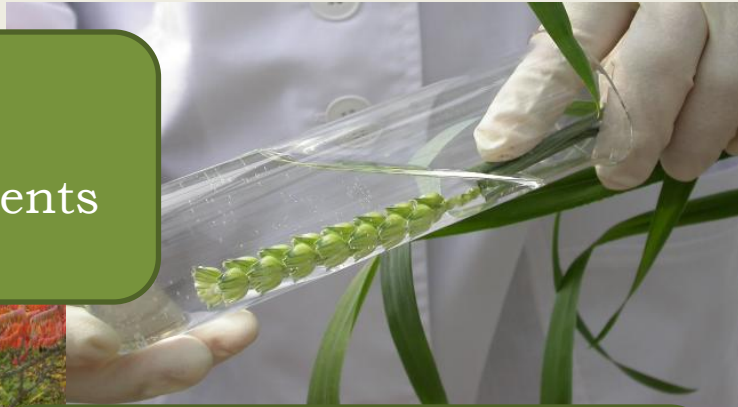
DNA marker development
DH production in wheat
Freezing tolerance testing

Facilities:

DNA laboratory
Experimental greenhouse
Growth and freezing chambers

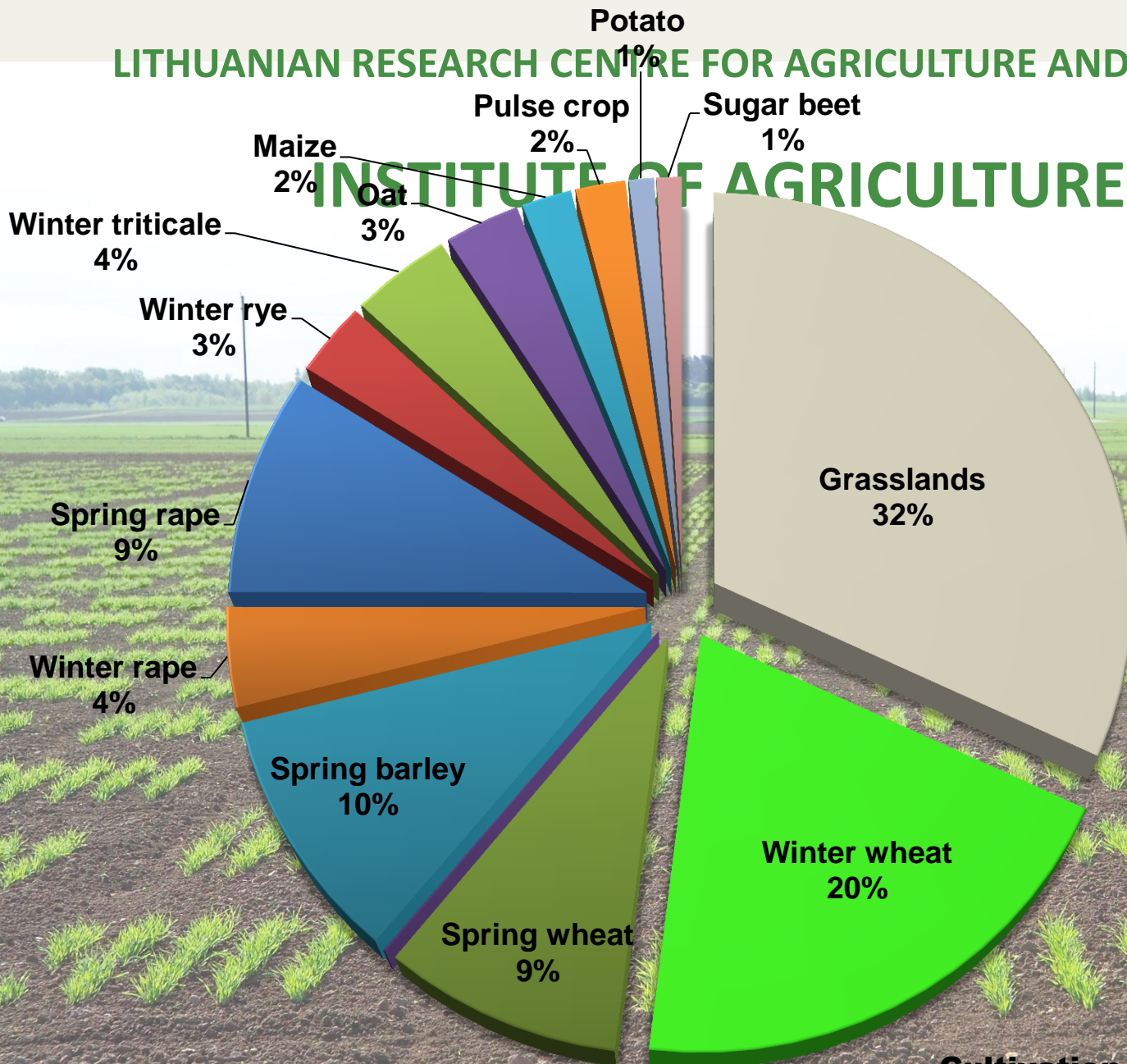
Collaborators:

Iowa State University
ETH Zurich
Aarhus University



LITHUANIAN RESEARCH CENTRE FOR AGRICULTURE AND FORESTRY

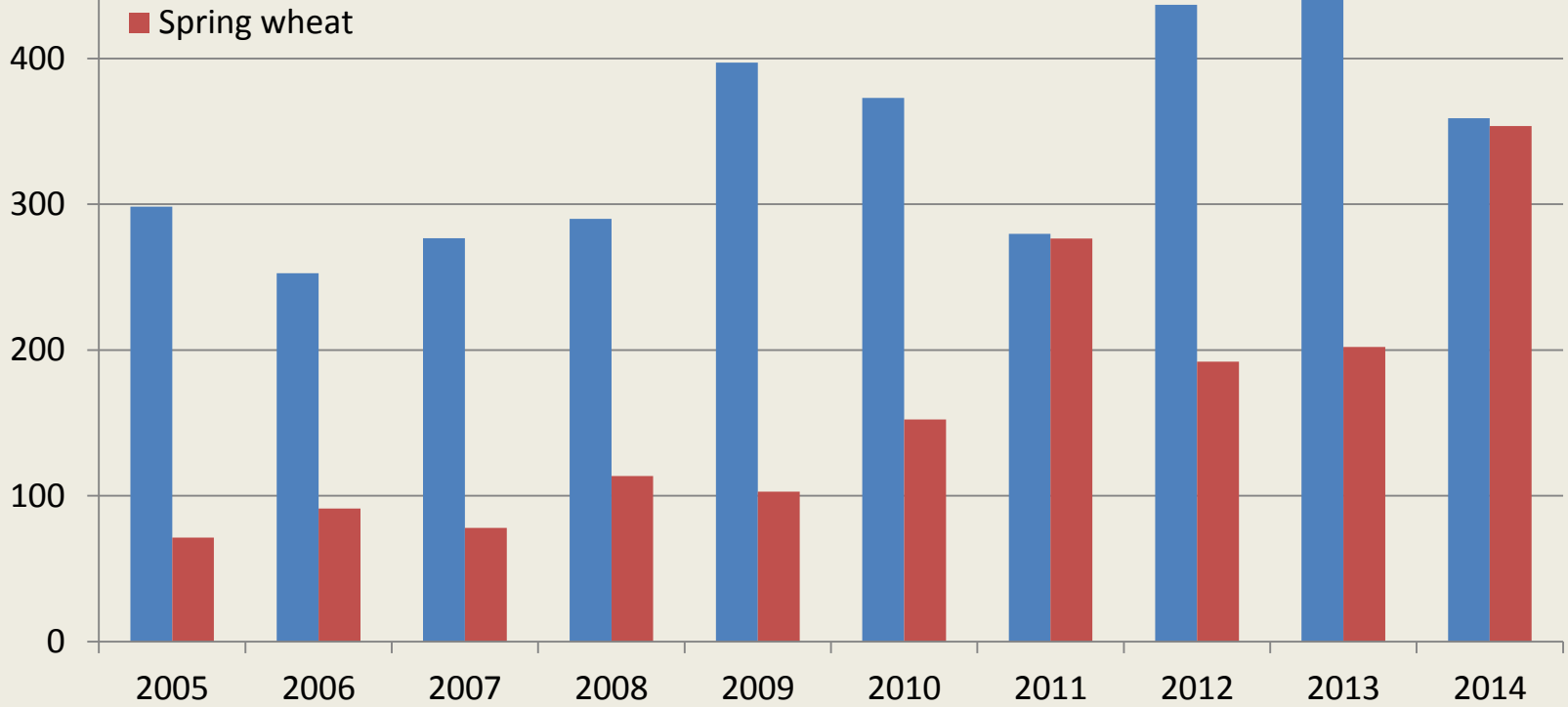
INSTITUTE OF AGRICULTURE



Cultivation area in Lithuania, %



kPa



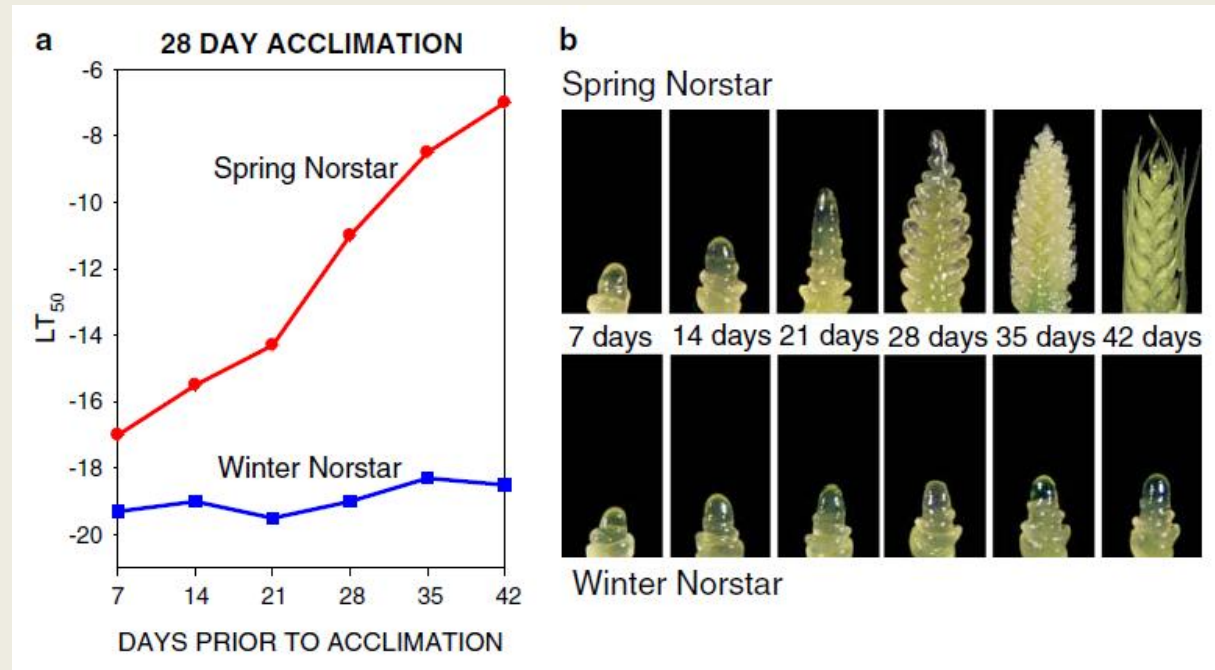
Plant responses to cold (and light)

- **Chilling** – low above zero temperatures
- **Freezing** - below zero temperatures
- **Cold acclimation** – acquisition of freezing tolerance (easy come easy go)
- **Vernalization** - acquisition of competence to flower (takes time and stays)
- **Winter hardiness** – ability to overwinter

Winter *versus* spring wheat

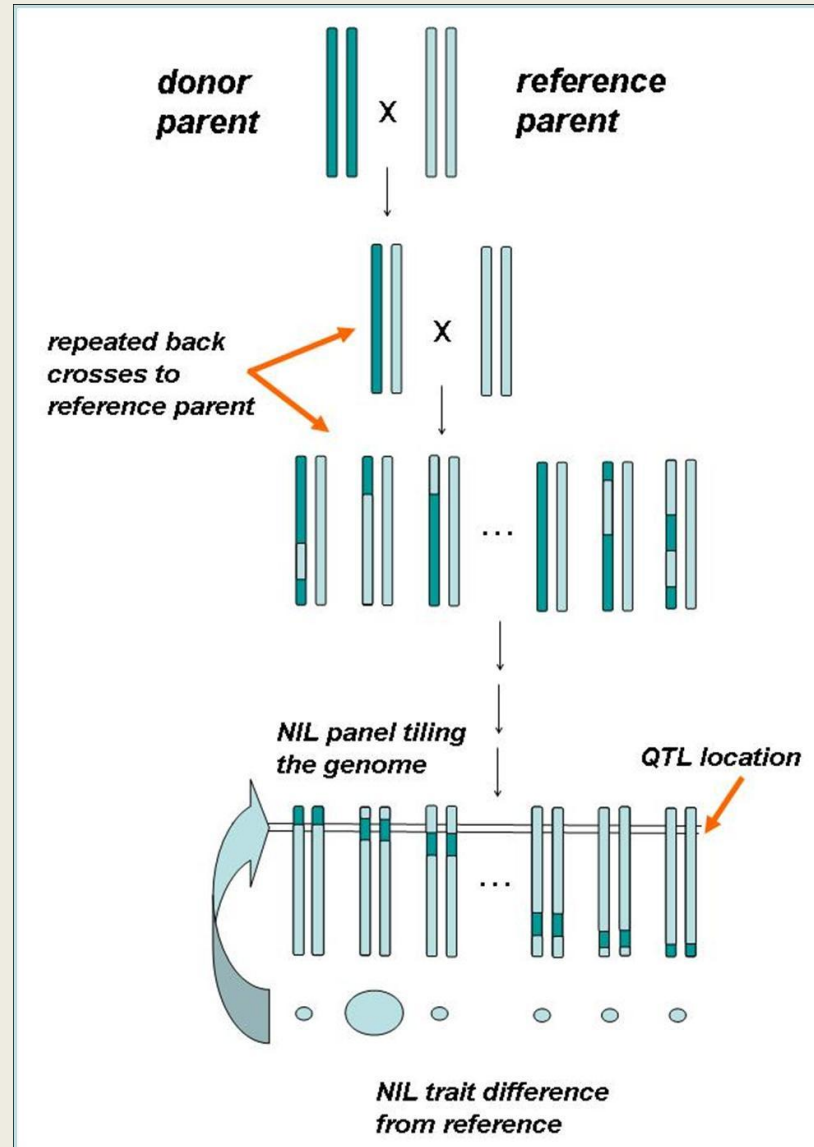


VRN1 controls vernalization

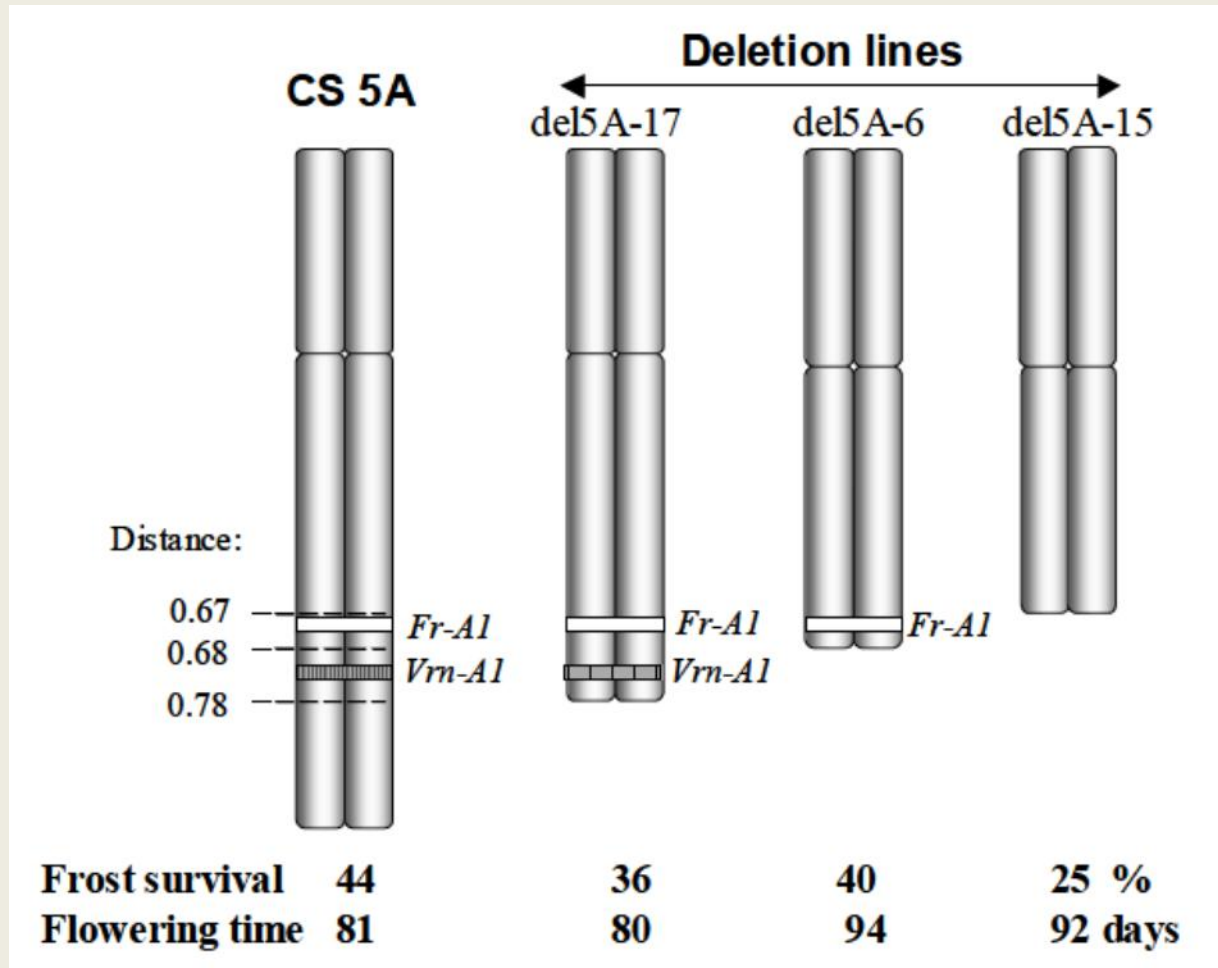


Norstar NILs were grown at 20°C and 16-h day from 7 to 42 days and then LT acclimated at 2°C for 28 days. a LT₅₀ versus plant age. b Dissected shoot apices indicating stage of development at the start of LT acclimation.

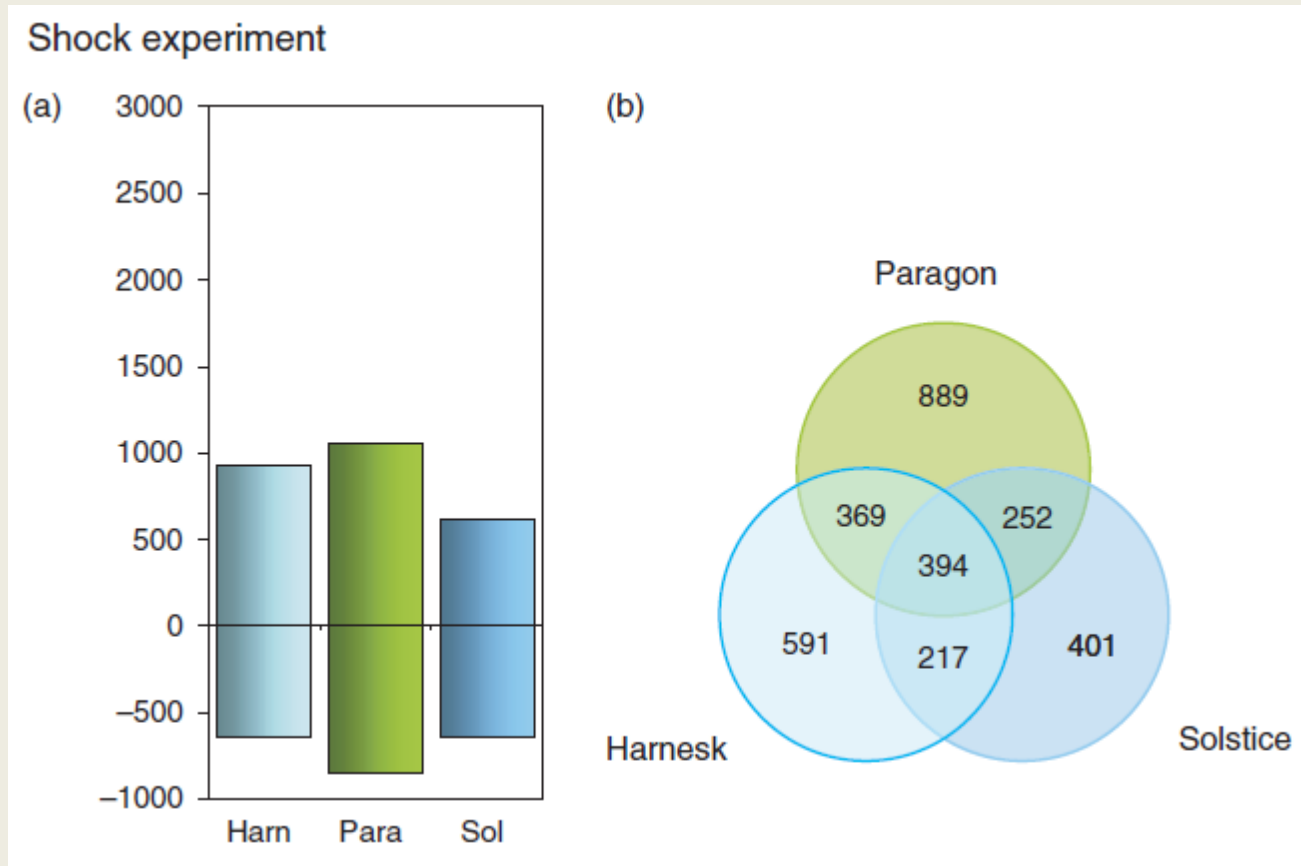
Near-isogenic lines (NILs)



Decoupling of *Vrn1* and *Fr1*

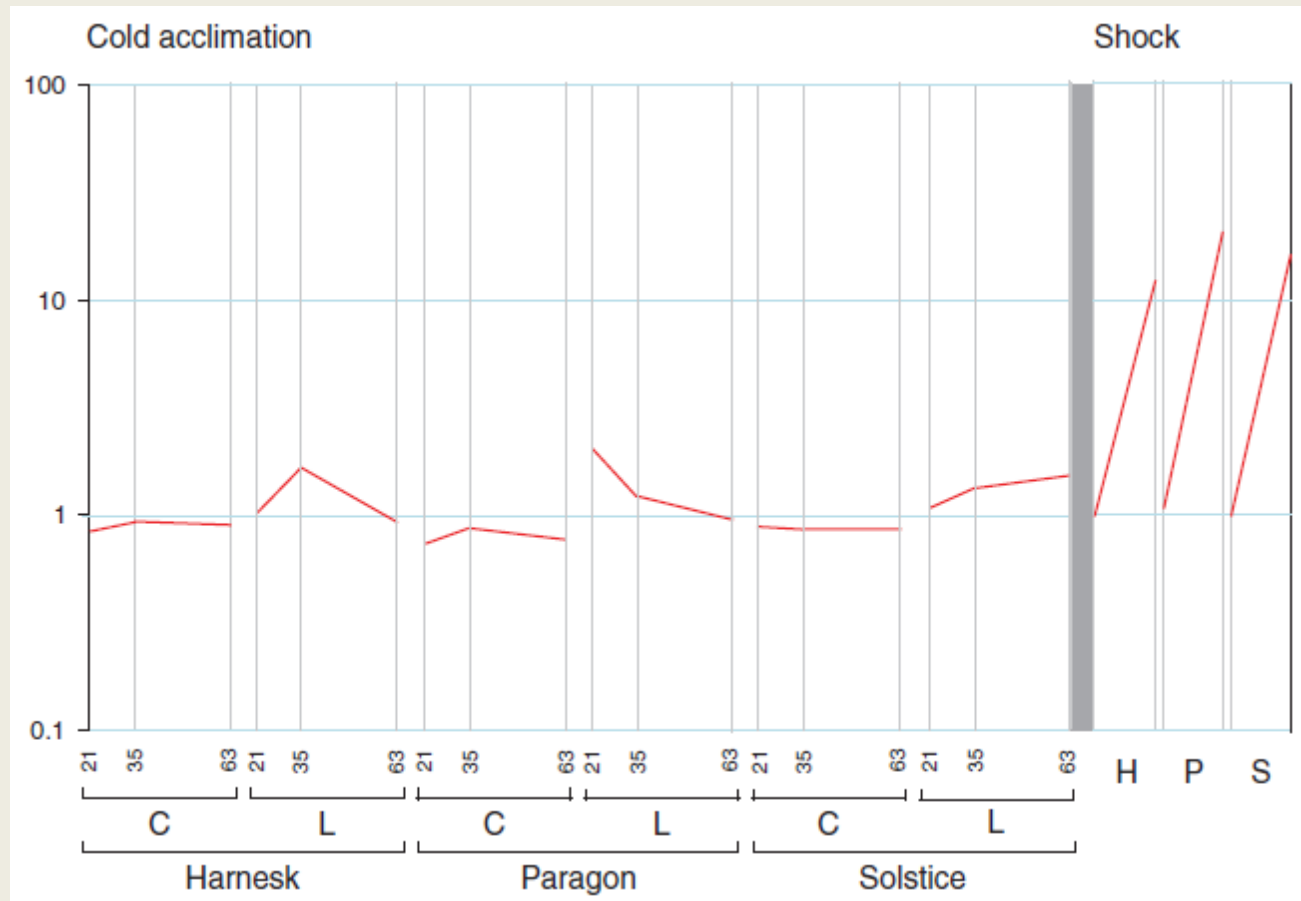


Wheat transcriptome response to cold



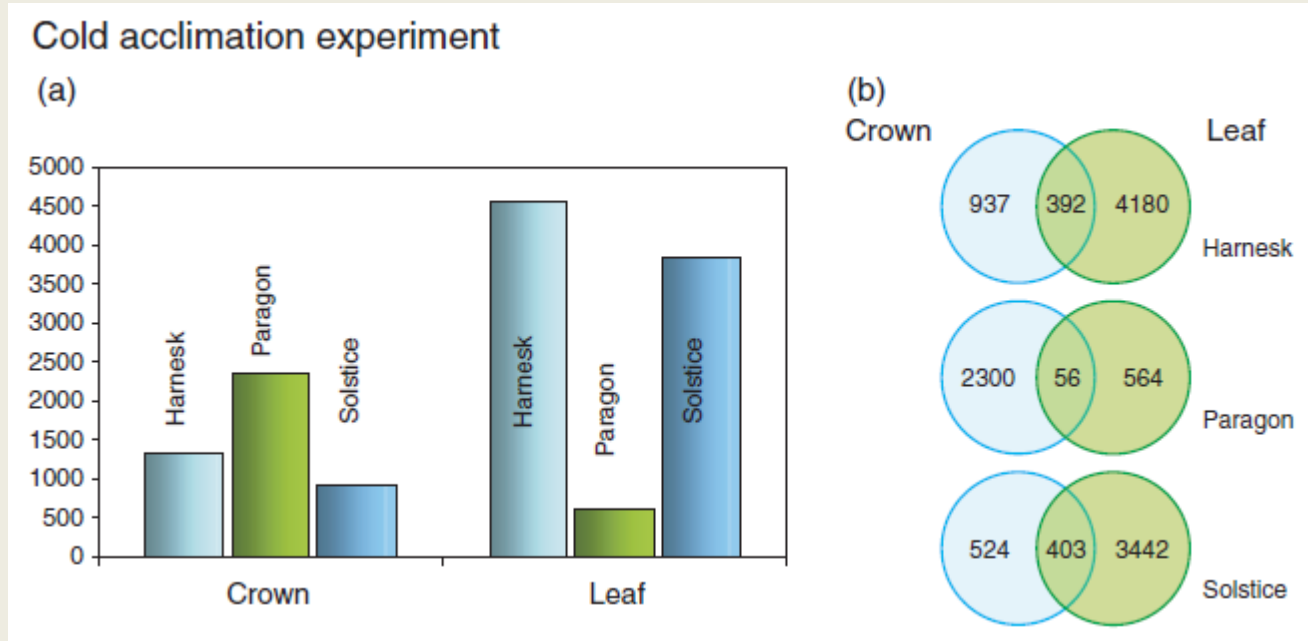
(a) Statistically significant two-fold or greater changes ($P = 0.05$) in transcript abundance in plants exposed to a 4 °C for 2 days. (b) Venn diagram showing the number of genes expressed in common between the three varieties after exposure to a 'cold shock'.

Cold acclimation *versus* cold shock



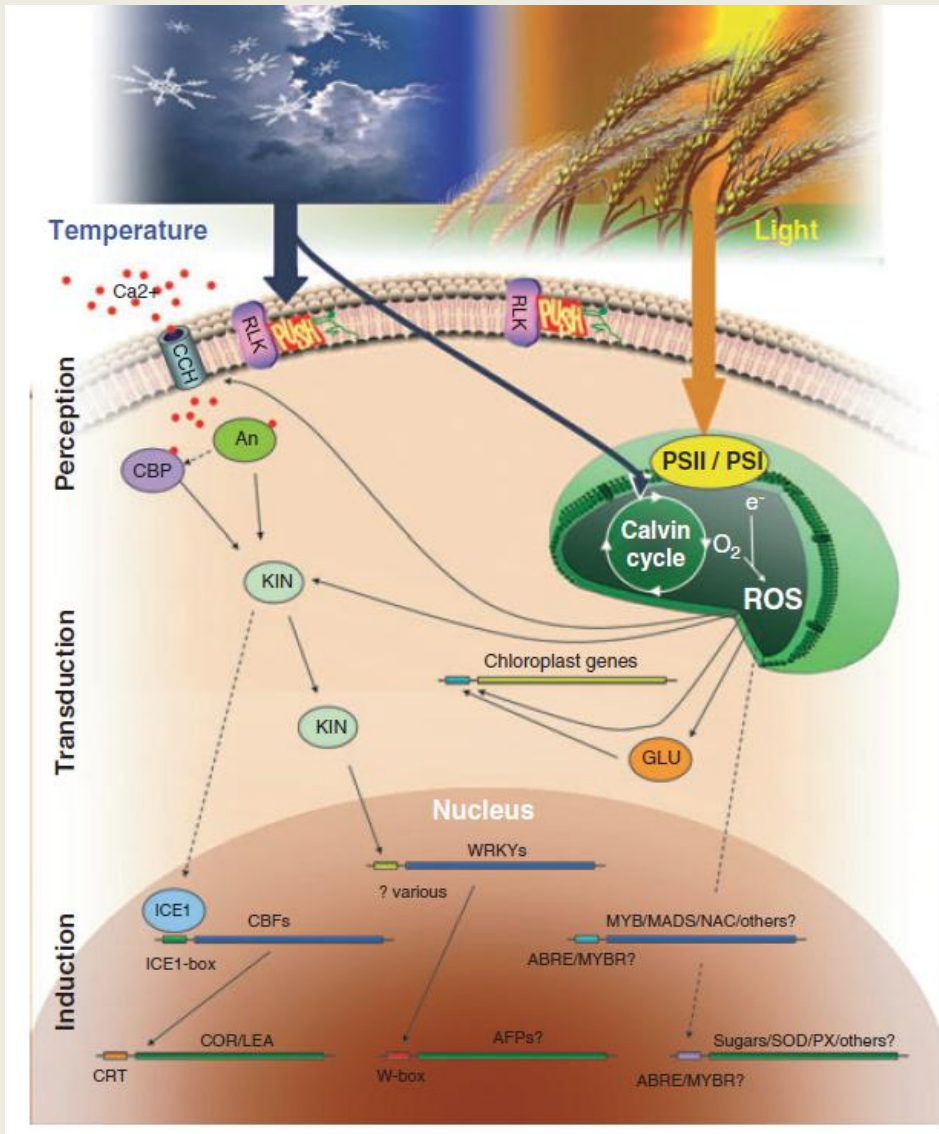
An early light-inducible protein (ELIP) showing a distinct response to a 'cold shock', but no response to a slow decline in temperature. C, crown; L, leaf; H, Harnesk; P, Paragon; S, Solstice.

Wheat transcriptome response to cold acclimation



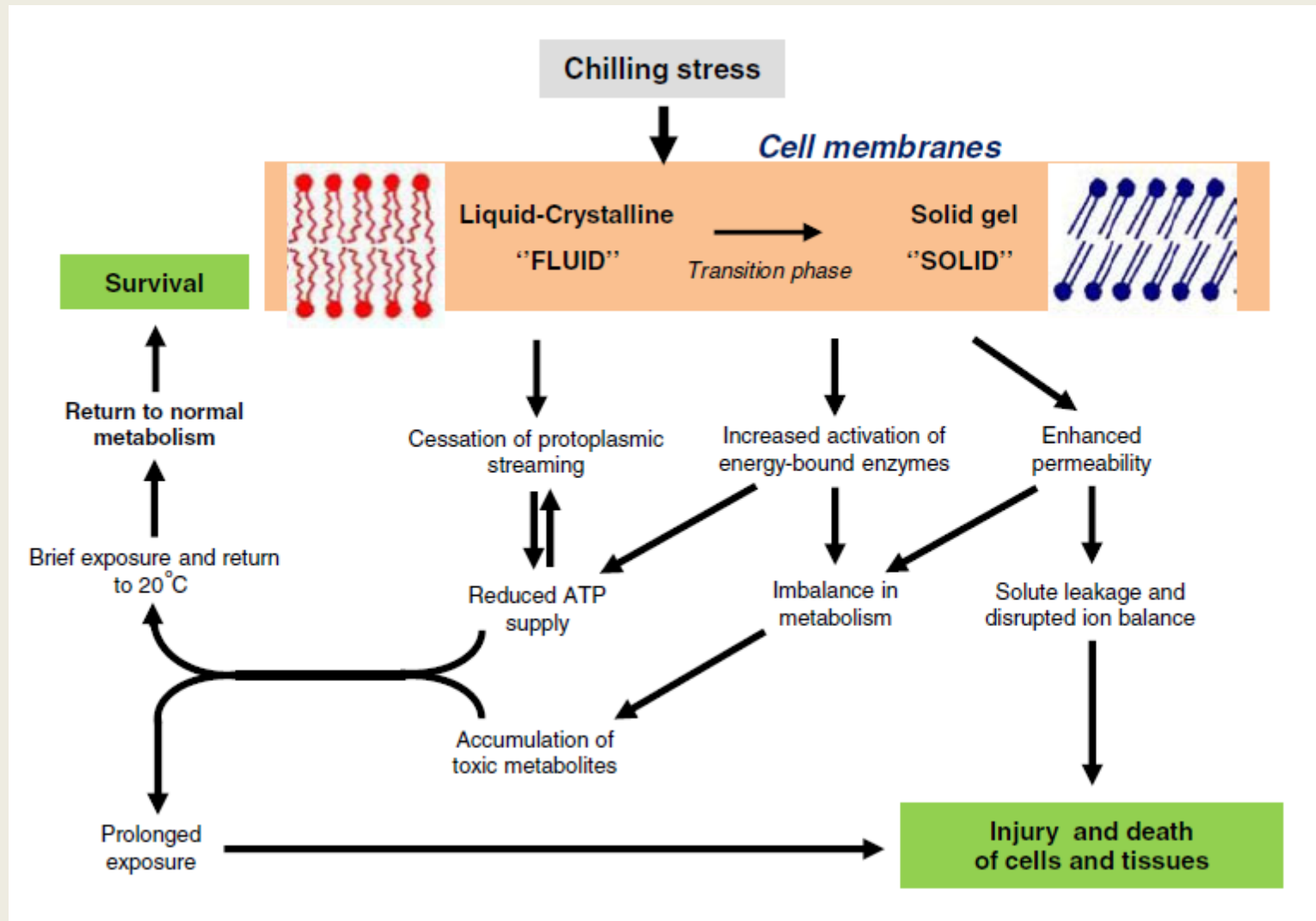
(a) transcript abundance in plants exposed to a gradual decline in temperature, light intensity and day length; (b) Venn diagrams showing the number of two-fold or greater changes ($P = 0.05$) in transcript abundance. The values refer to genes that changed in expression during the period 21–63 days.

Cold response in plant cells

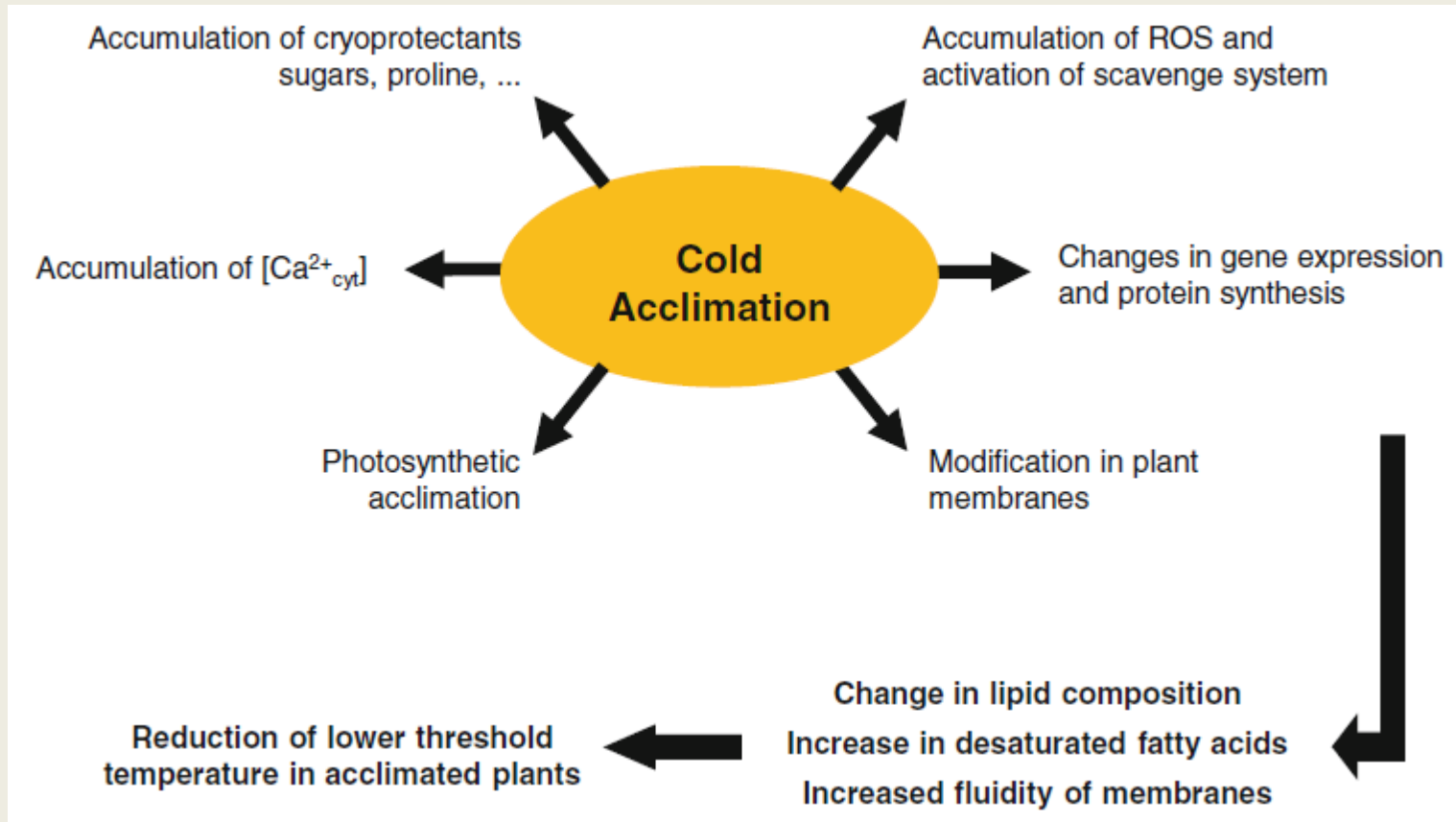


- ABRE, ABA response element*
- AFP, anti-freeze protein*
- An, annexins*
- CBF, C-repeat binding factor*
- CBP, calcium binding proteins*
- CCH, calcium channel*
- COR, cold-responsive genes*
- CRT, C-repeat elements*
- GLU, glutathione*
- ICE, inducer of CBF expression*
- KIN, kinases and phosphatases*
- LEA, late embryogenesis-abundant*
- PSI/PSII, photosystem I and II*
- RLK, receptor-like kinase*
- ROS, reactive oxygen species*

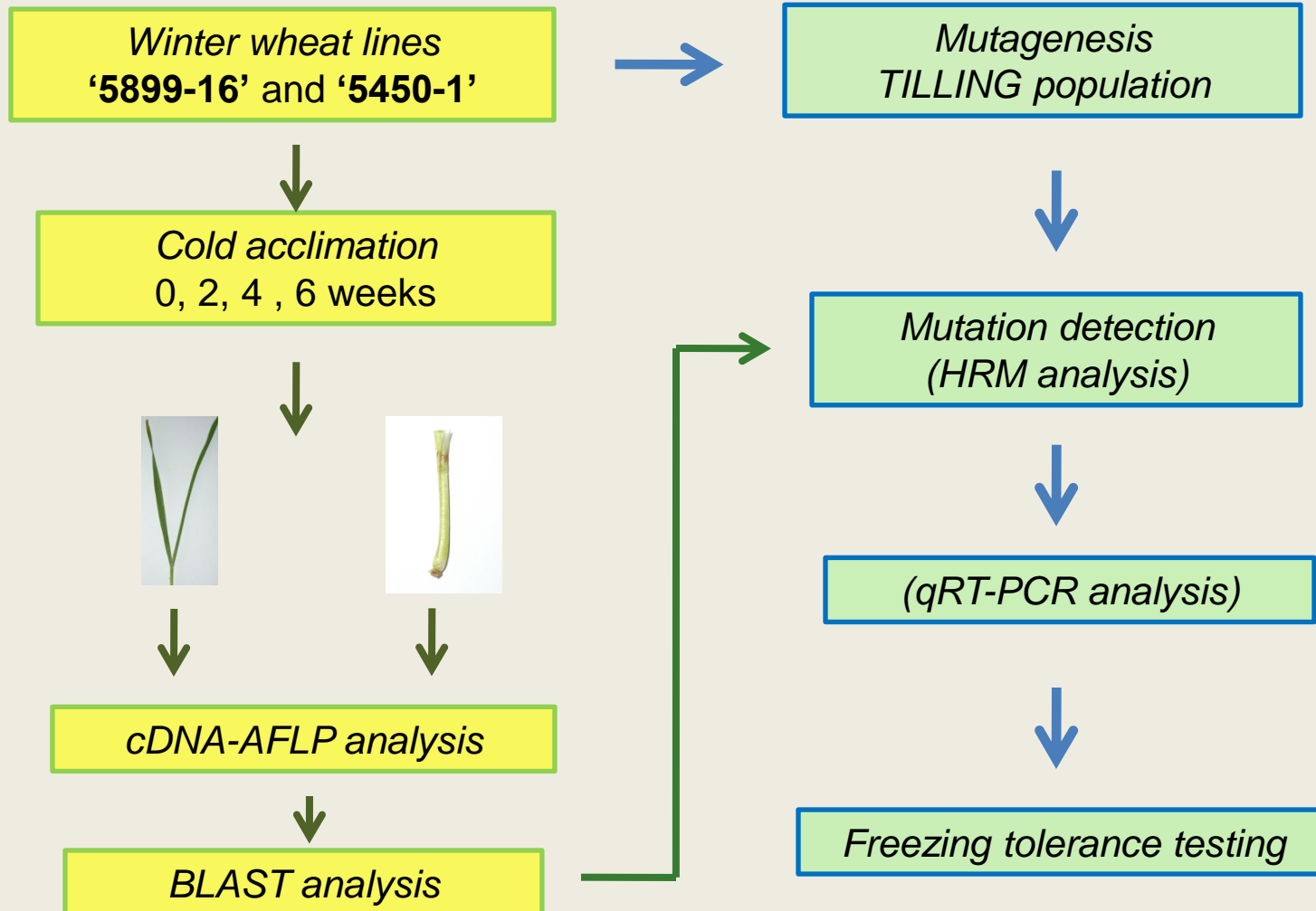
Chilling injury in chilling-sensitive plants



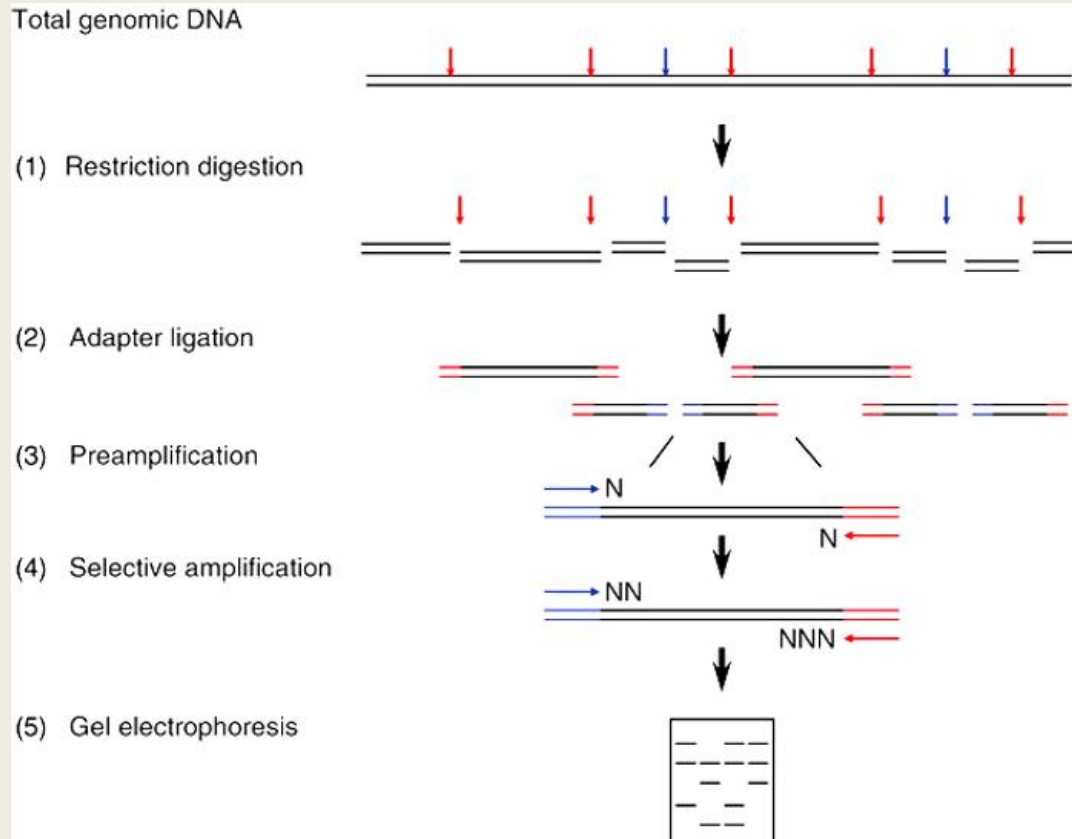
Cellular processes induced by cold acclimation



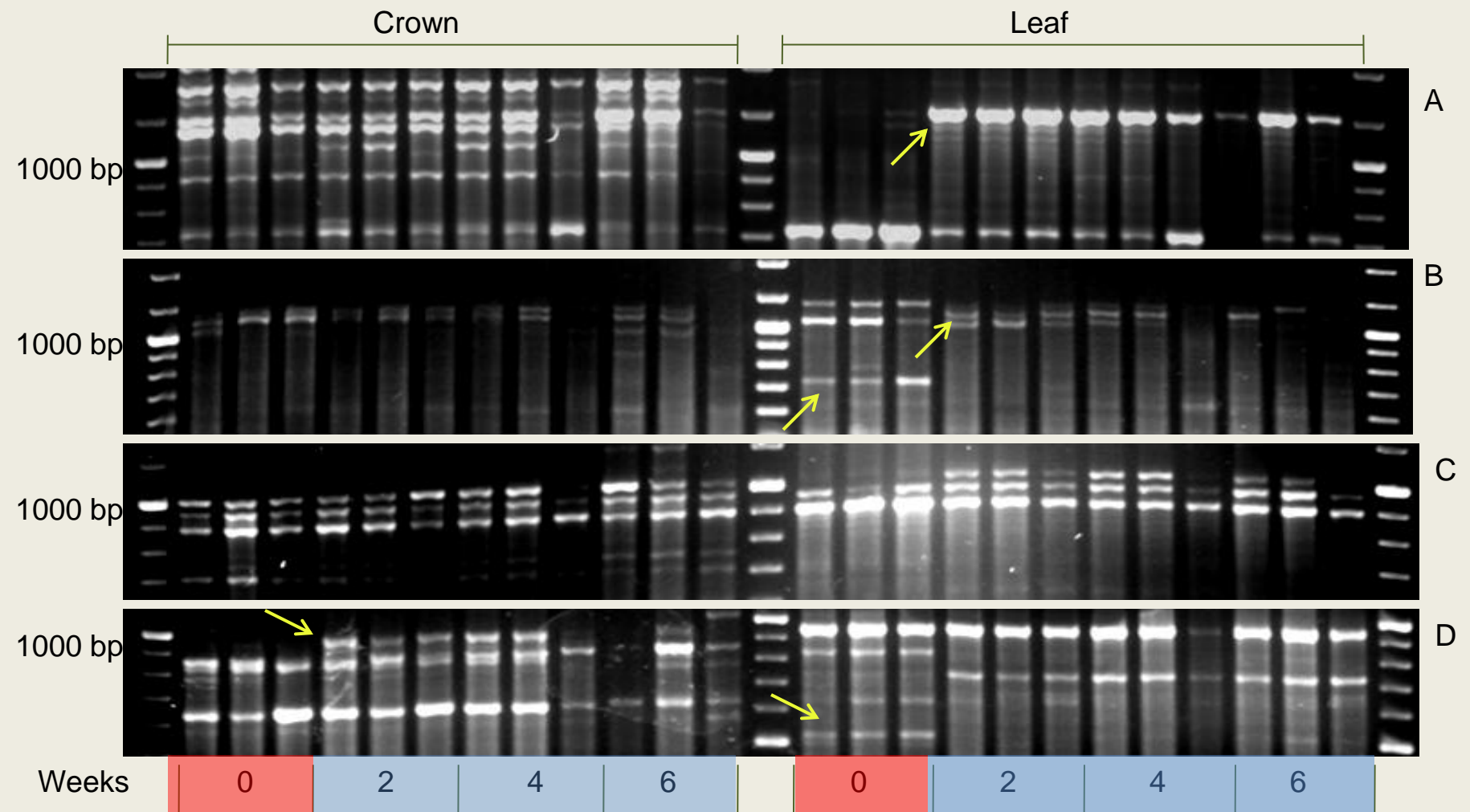
Identification and analysis of FT genes in winter wheat



AFLP and cDNA-AFLP



cDNA-AFLP results



Differentially expressed genes

2 weeks after the start of cold acclimation

Phosphoglycerate kinase, chloroplastic

Signal recognition particle 54 kDa protein, chloroplastic-like

ATP-dependent Clp protease ATP-binding subunit ClpC homolog 1

Clp-P protease subunit



Sucrose synthase 1

Heat-shock protein

Serine carboxypeptidase-like

Vacuolar-sorting receptor 1

Zinc finger AN1 domain



Crown



Beta-3-tubulin

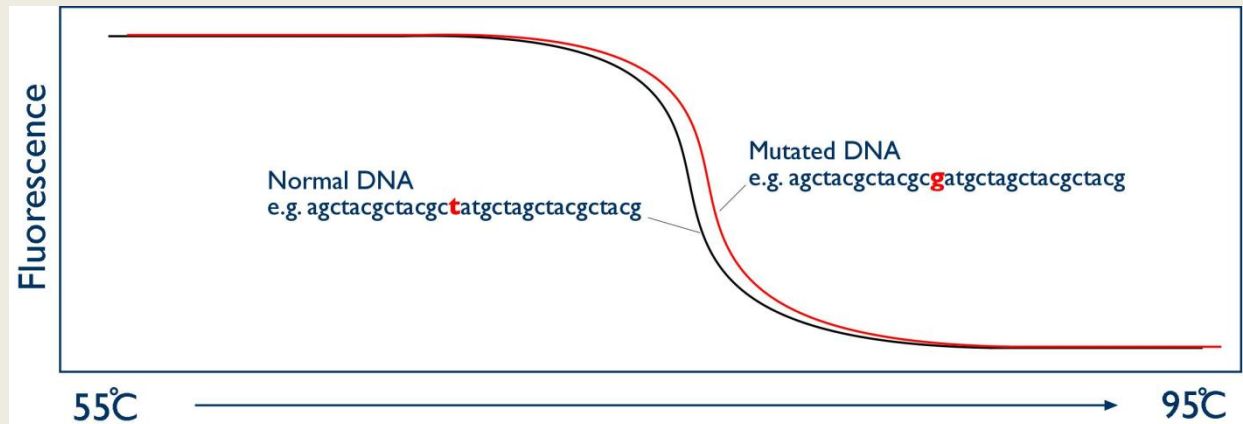
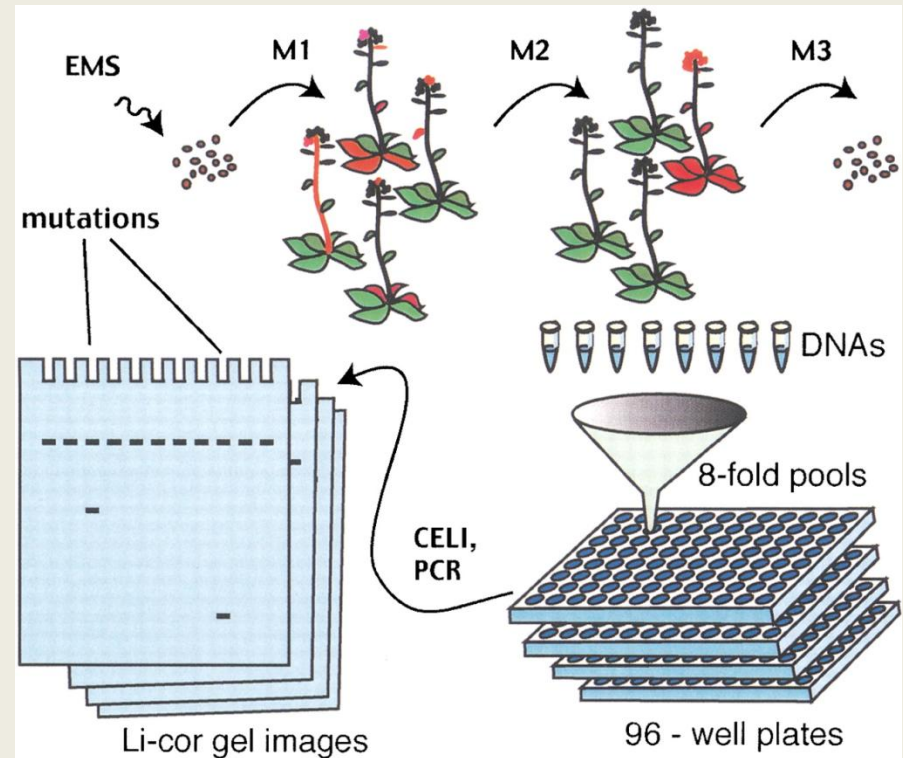
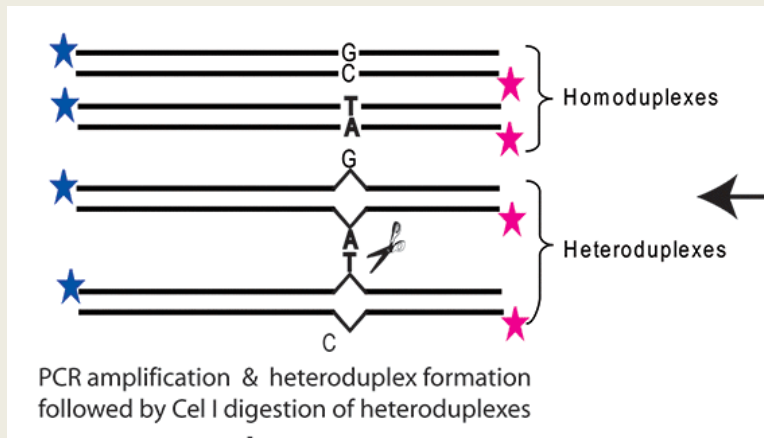
Phototropin-2-like

Digalactosyldiacylglycerol synthase 2, chloroplastic

TILLING

Targeting Induced Local Lesions in Genomes

Colbert et al., 2001



Mutagenesis and mutation screening

(0.4% - 1.0%)

EMS



'5450-1' or '5899-16'
Winter wheat lines



M1 generation
(heterozygous and chimeric)



M2 seeds



M2 generation
(segregating)



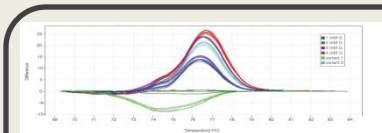
M3 seeds
(stored)

Creation of TILLING population

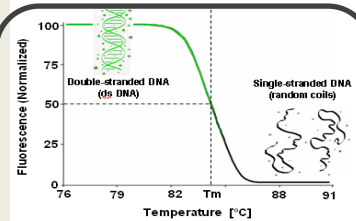


DNA isolation

HRM analysis



Data Analysis by
HRM Software v3.0



Sample Melt

7500 Fast Real-Time PCR System

(Applied Biosystems)

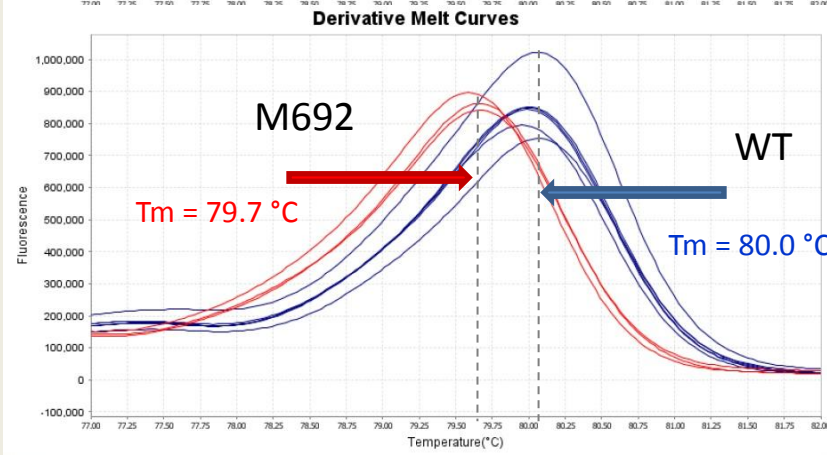
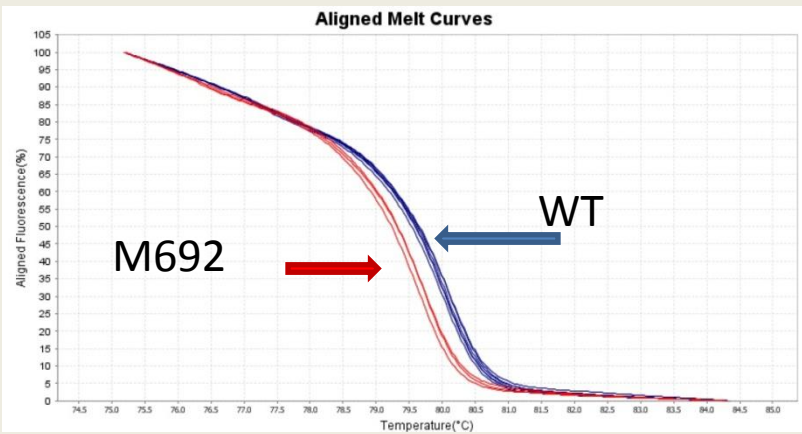
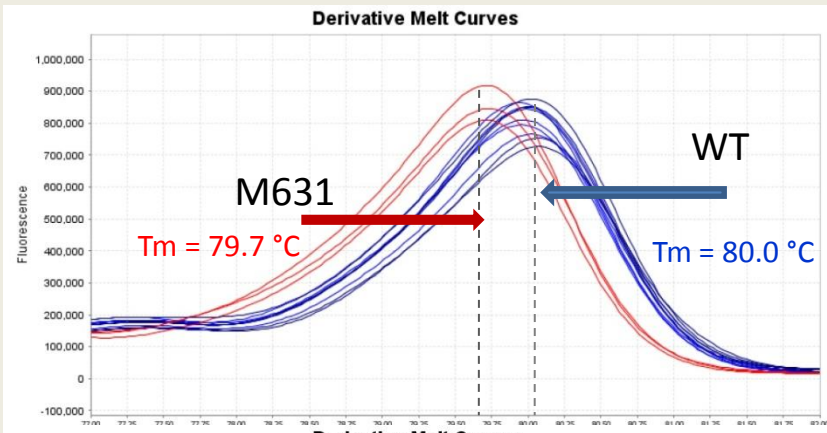
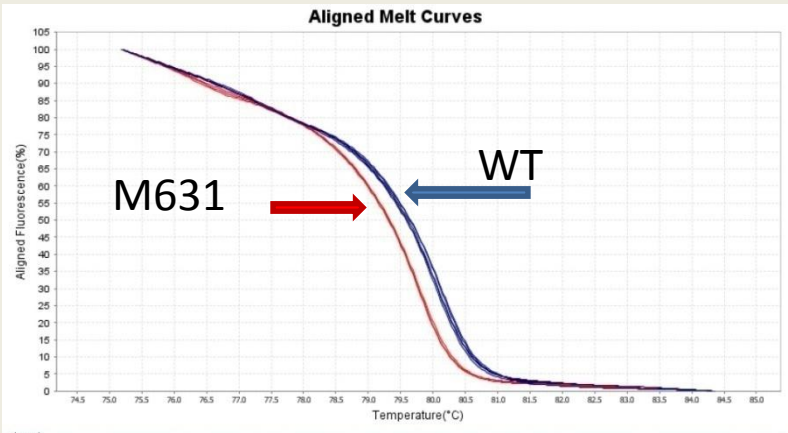
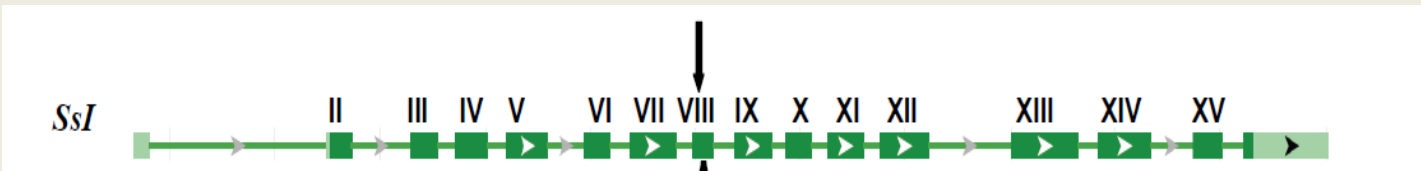
Exon X

2nd PCR
of amplicon with
lengths <250 bp

Gene X

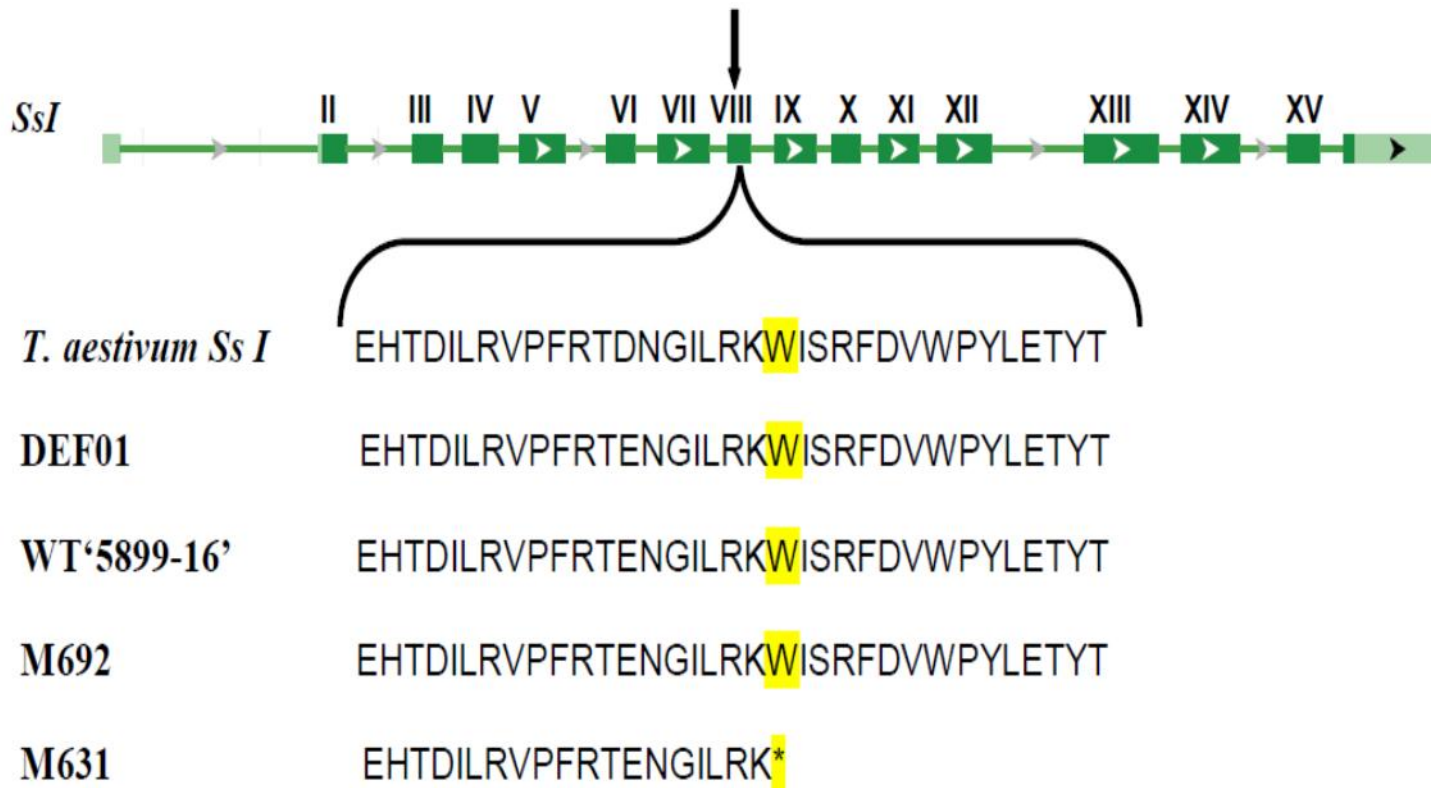
1st PCR
region of interest

Mutation discovery by HRM analysis

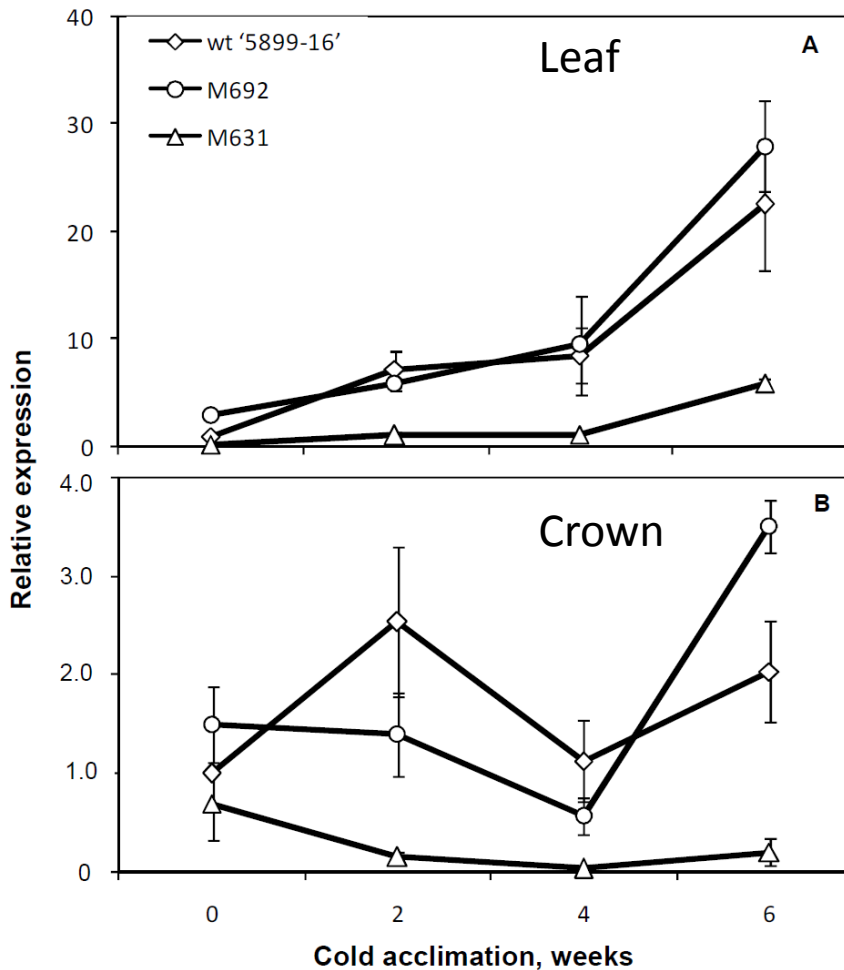


Aligned curves of fluorescence vs temperature showing wild type (blue) and mutant (red) samples.

2 mutations per 75.68 kb

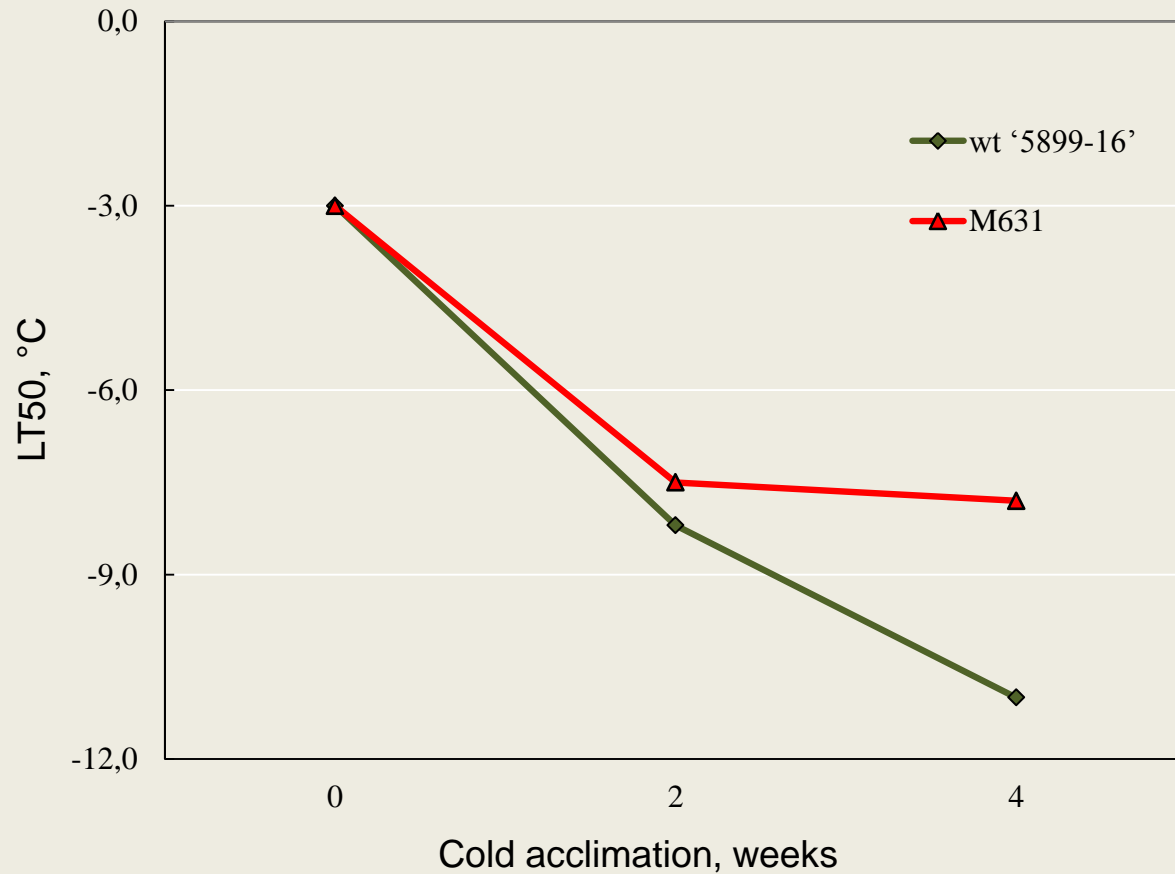


Nonsense-mediated *Sucrose synthase 1* mRNA decay during cold acclimation in winter wheat



wt – wild type
M692 – missens mutation
M631 – nonsense mutation
A – leaf tissue
B – crown tissue

Reduced freezing tolerance of M631



Freezing tolerance of wild type '5899-16' and M631 winter wheat acclimated at 5 °C for 0–4 weeks.

Perennial ryegrass (*Lolium perenne* L.)



- The main forage grass species in Denmark and further south and west in Europe
- Finland: 2%, Estonia: 13%, Norway: 13%, Lithuania: 30% of certified seed sale
- Superior feed quality and productivity under frequent cutting regimes
- Is expected to expand further north due to milder winters with shorter periods of snow cover

Main problems with perennial ryegrass

- Susceptibility to low-temperature fungi
- Inadequate growth cessation in autumn to allow for sufficient cold hardening
- Can de-harden too early with increased risk of frost injury in spring



Festuca arundinacea and *L. perenne*



F. pratensis and *L. perenne*

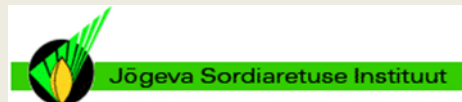


L. perenne at Vågønes, Bodø

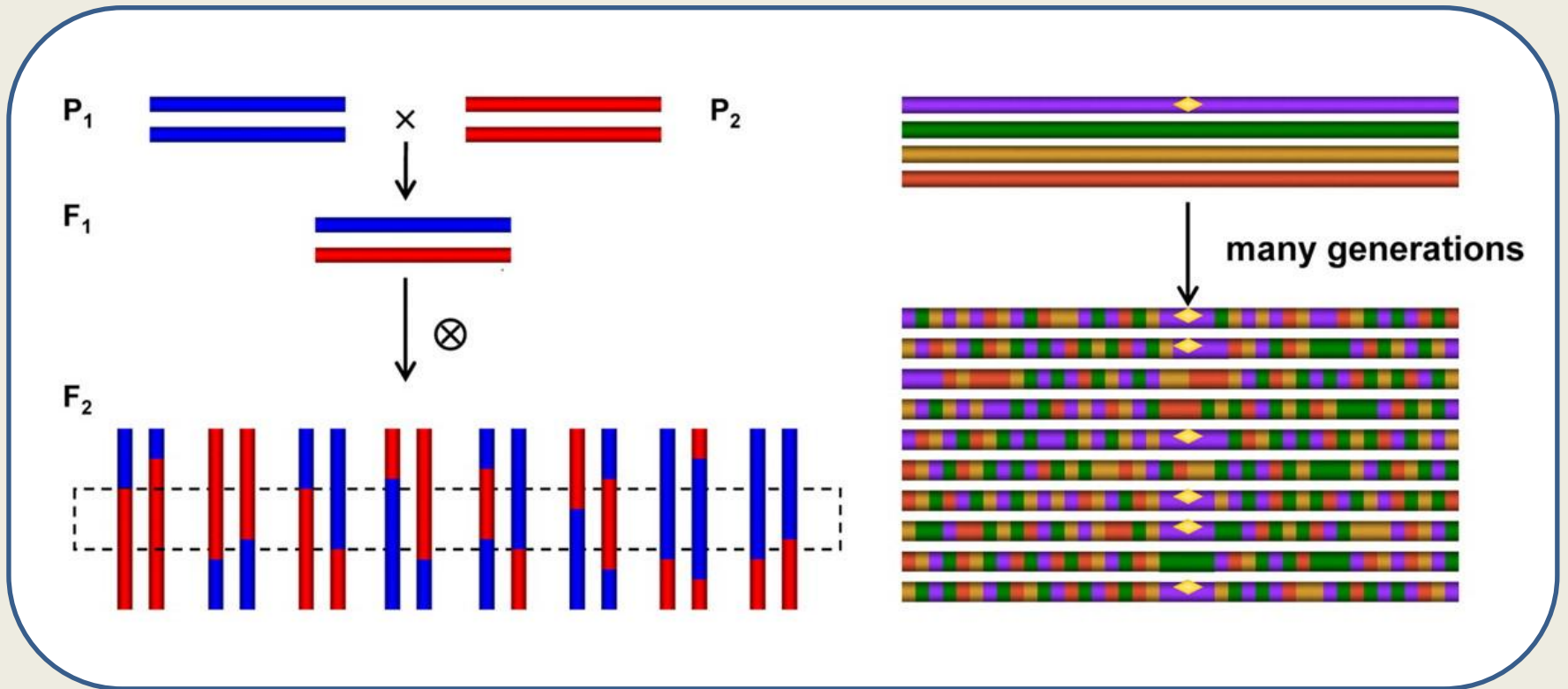
PPP perennial ryegrass pre-breeding project

- Aim:

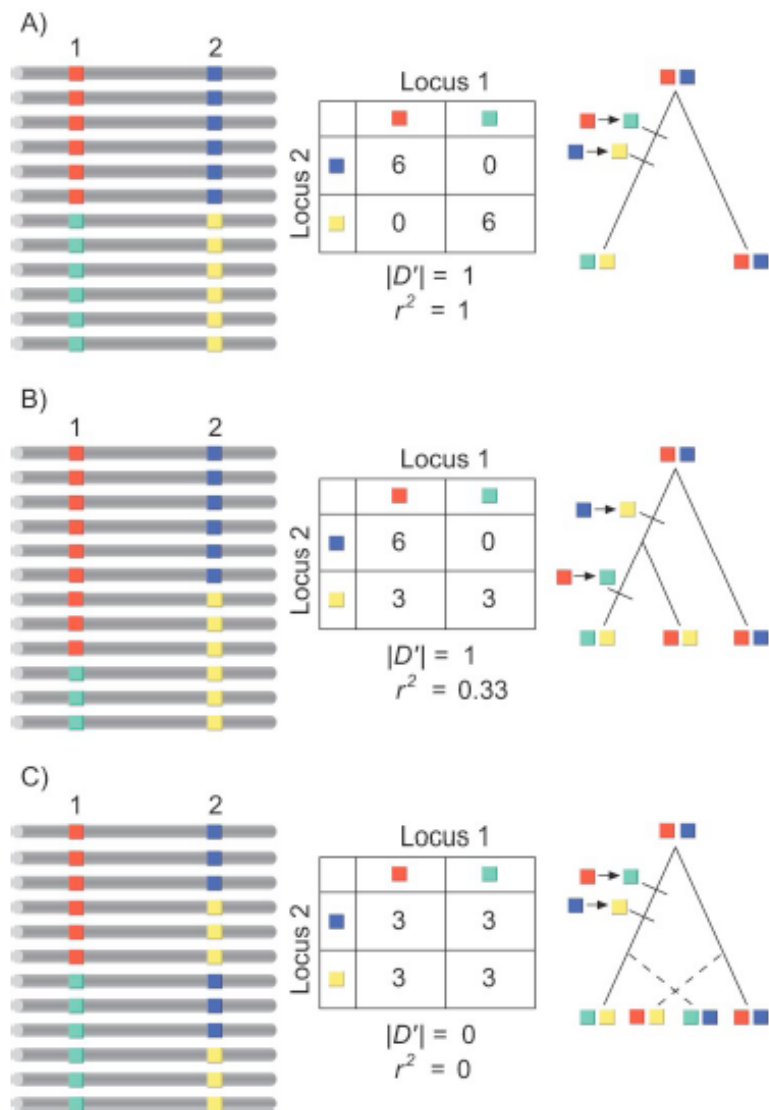
- Identify and select new plant materials for development of cultivars with a suitable adaptation to future climates
- Recombine exotic materials with existing germplasm to create new genetic resources



Biparental versus Association mapping



Linkage disequilibrium (LD)

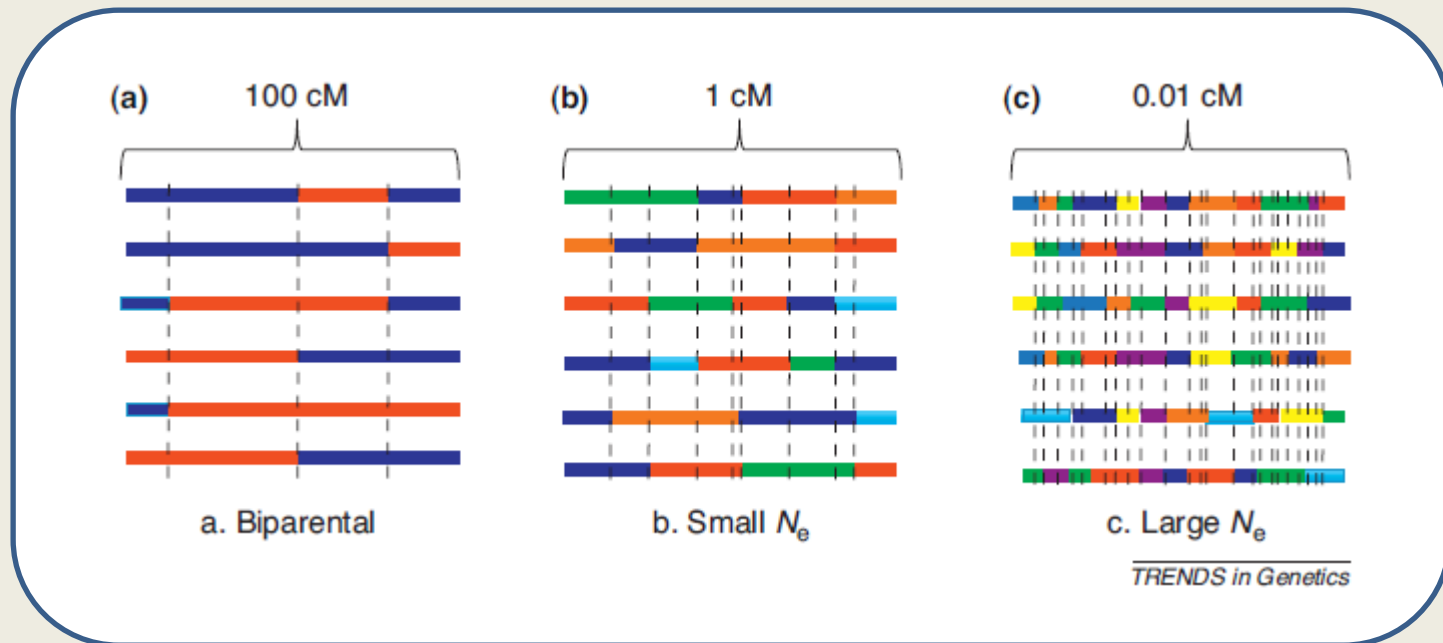


$$D_{ab} = (\pi_{AB} - \pi_A\pi_B).$$

$$r^2 = \frac{(D_{ab})^2}{\pi_A\pi_a\pi_B\pi_b}.$$

LD in plants

- Mating type
- Population structure
- Genetic drift
- Selection intensity



Sample preparation for freezing tolerance test



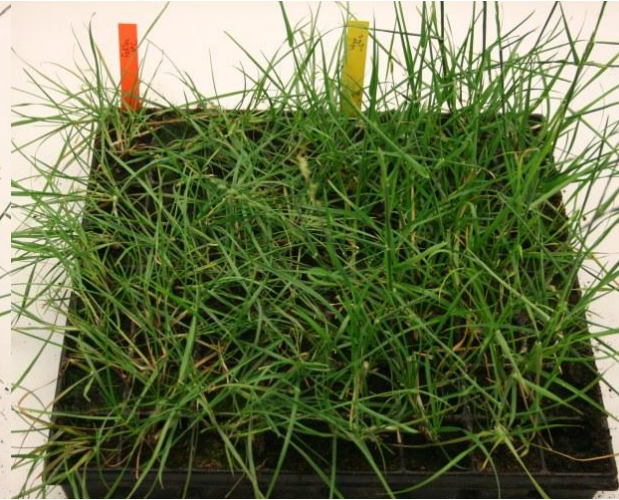
**Freezing
test**

Regrowth at 3 weeks after freezing

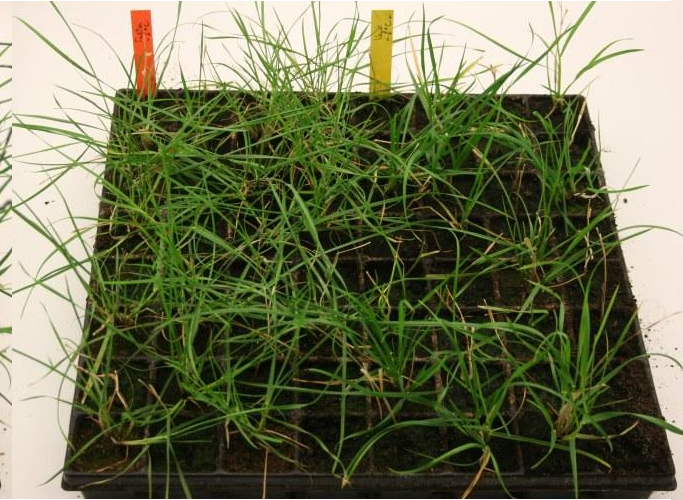
Control



-4°C



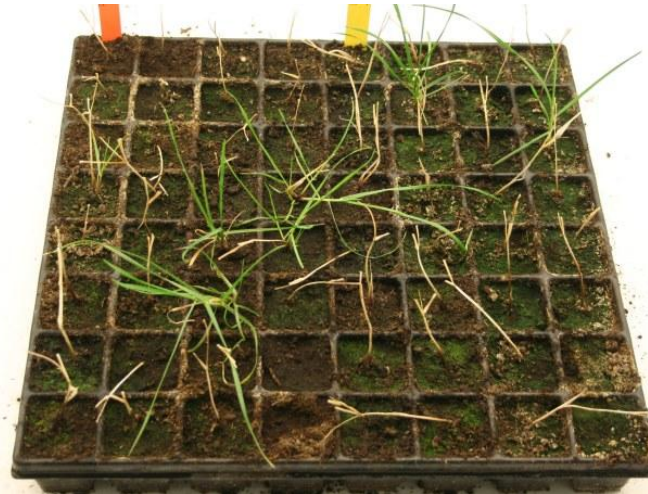
-6°C



-8°C



-10°C



-12°C



Phenotypic variation for freezing tolerance traits

Trait	Ecotypes				Cultivars			
	Mean	Minimum	Maximum	Sd ^a	Mean	Minimum	Maximum	Sd ^a
EL at -8°C [%]	14.27	6.09	25.80	5.23	15.83	7.82	25.06	4.82
EL at -12°C [%]	38.91	22.27	57.06	9.14	43.09	23.85	63.51	11.48
PTS at -8°C [%]	51.12	0.00	100.00	35.24	56.28	0.00	100.00	30.33
PTS at -12°C [%]	11.02	0.00	91.67	22.10	9.92	0.00	53.33	16.54
PC [$\mu\text{g g}^{-1}$ DW]	2558.86	473.34	6157.35	1366.34	2521.75	519.68	5840.46	1342.00

EL – electrolyte leakage

PTS – plant tiller survival

PC – proline content

Correlation between phenotypic traits

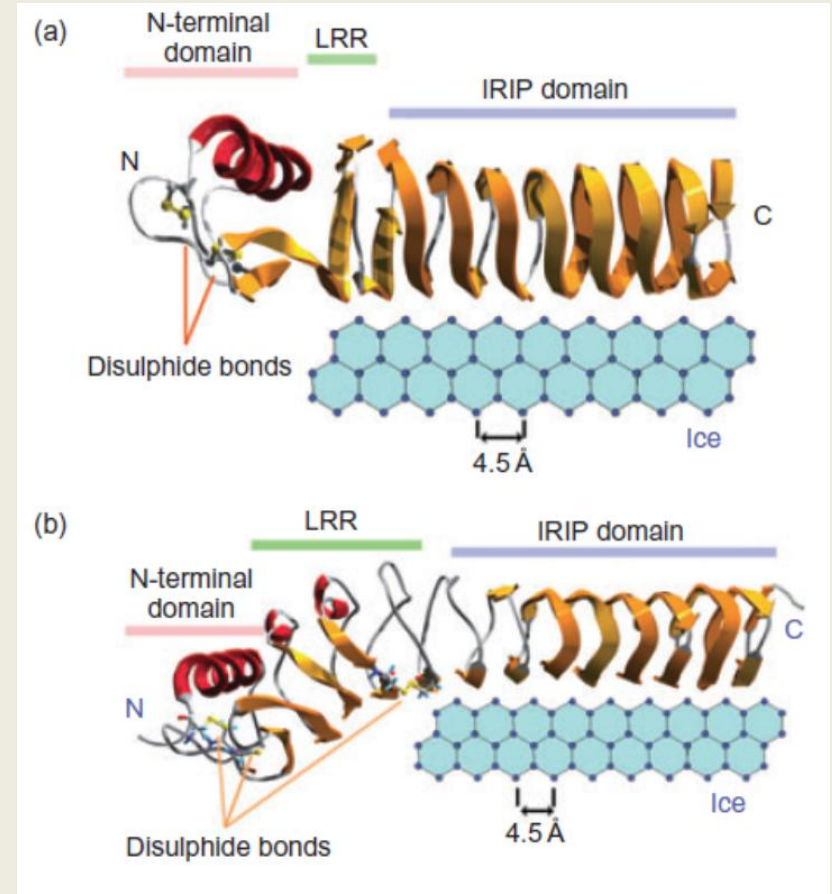
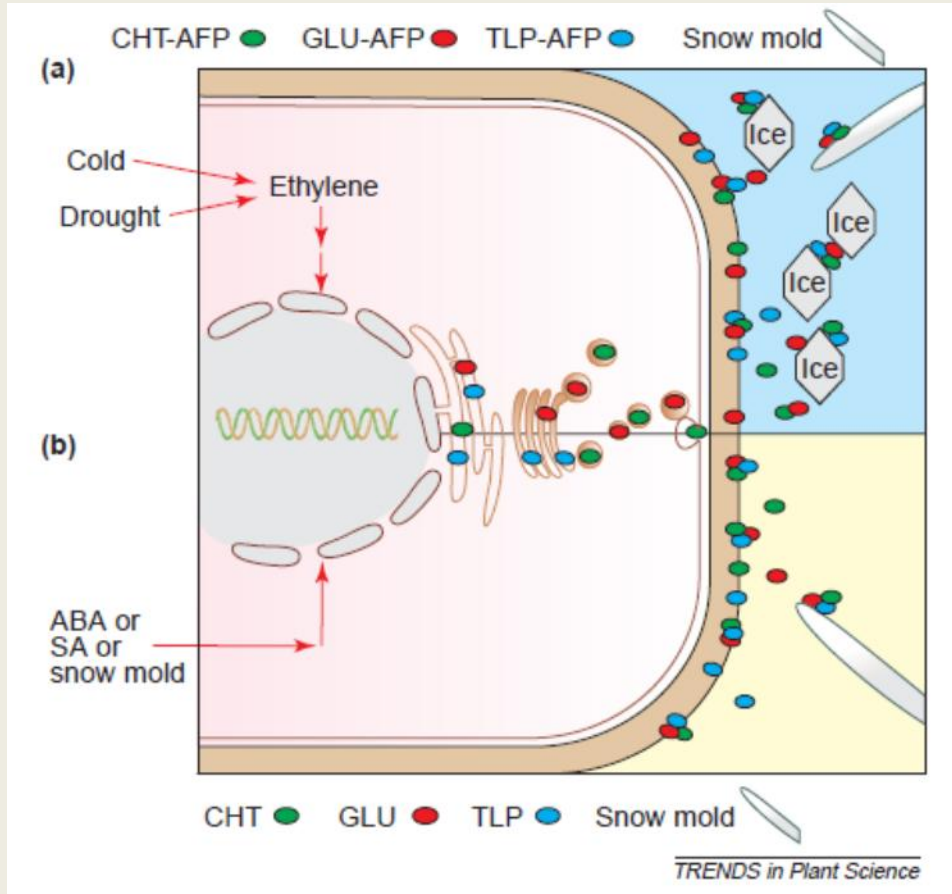
Trait	EL at -8 °C	EL at -12 °C	PTS at-8 °C	PTS at -12 °C
EL at -8 °C	-			
EL at -12 °C	0.43***	-		
PTS at-8 °C	-0.40***	-0.21 ns	-	
PTS at -12 °C	-0.14 ns	-0.49***	0.46***	-
PC	-0.06 ns	0.02 ns	-0.20 ns	0.03 ns

EL – electrolyte leakage

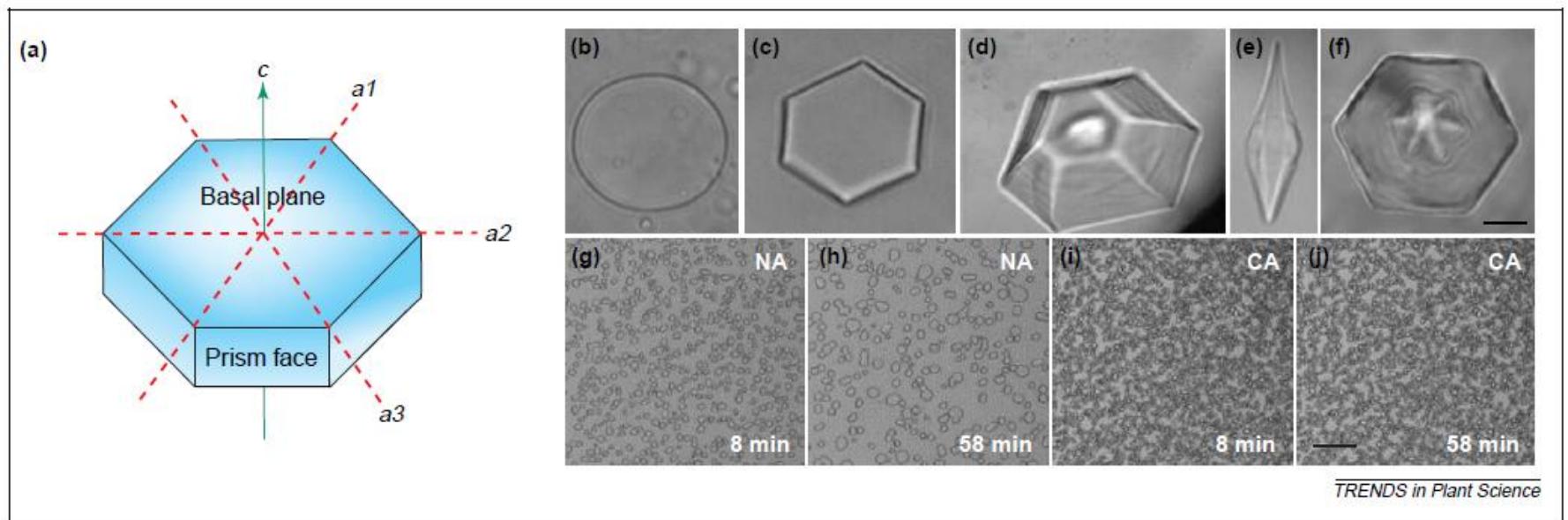
PTS – plant tiller survival

PC – proline content

LpIRI1 – ice recrystallization inhibition

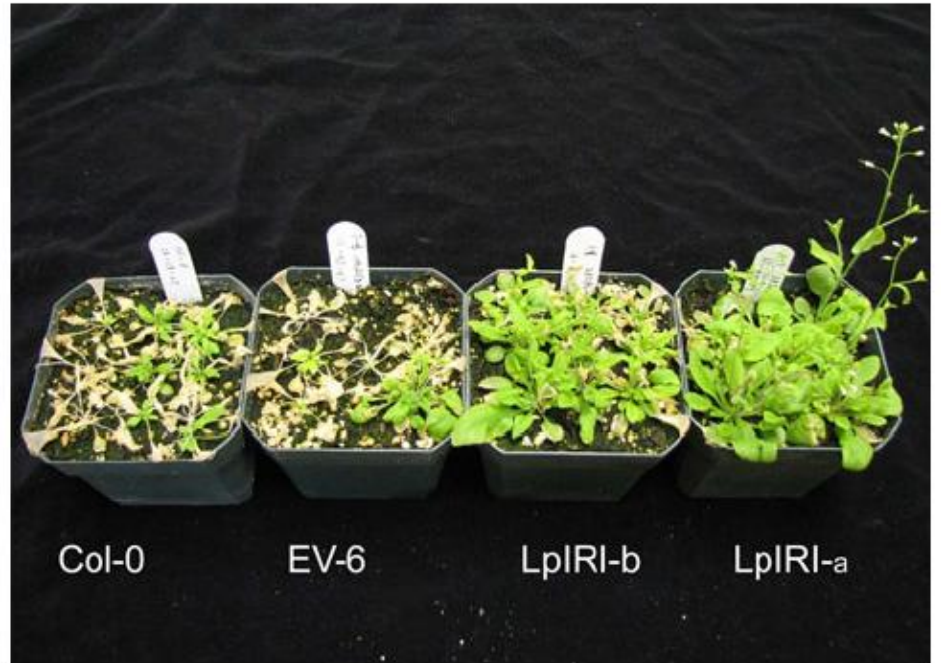
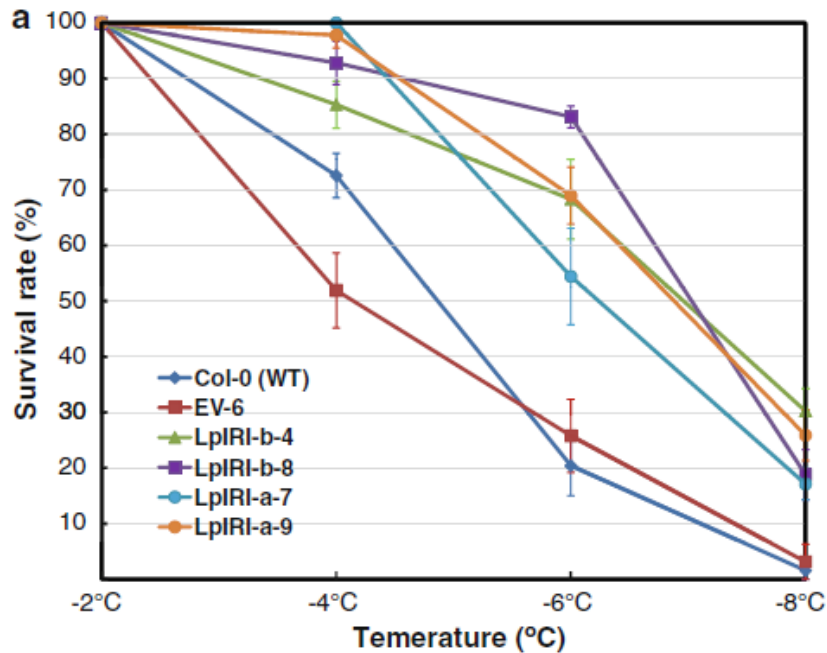


Controlled ice crystal growth



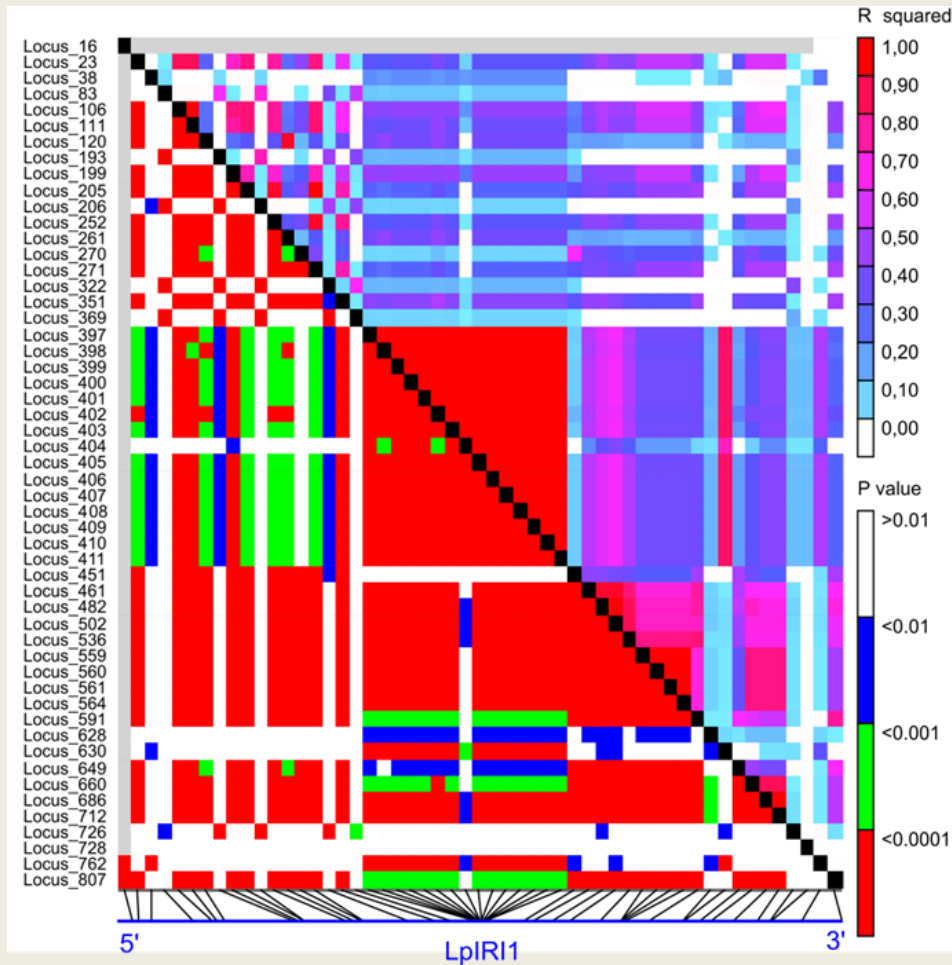
(a) ice crystal morphology in (b) water (c) dilute AFP (d-f) concentrate AFP solution. Scale bar = 10 μm . Recrystallization of ice (g-j) in nonacclimated (NA) and cold-acclimated (CA) winter rye extracts. Scale bar = 1.75 cm.

LpIRI improve FT in *Arabidopsis*



Survival rates and phenotypic appearance of control and transgenic *Arabidopsis* plants after freezing at -4°C and recovery. From left to right wild-type (Col-0); plants carrying an empty pMDC32 vector (EV-6); transgenic plants overexpressing LpIRI-b; transgenic plants overexpressing LpIRI-a.

Linkage disequilibrium in *LpIRI1*



- ORF 855 bp
- 52 SNPs (MAF >5)
- SNP density 1/16 bp
- 1 INDEL of 15 bp

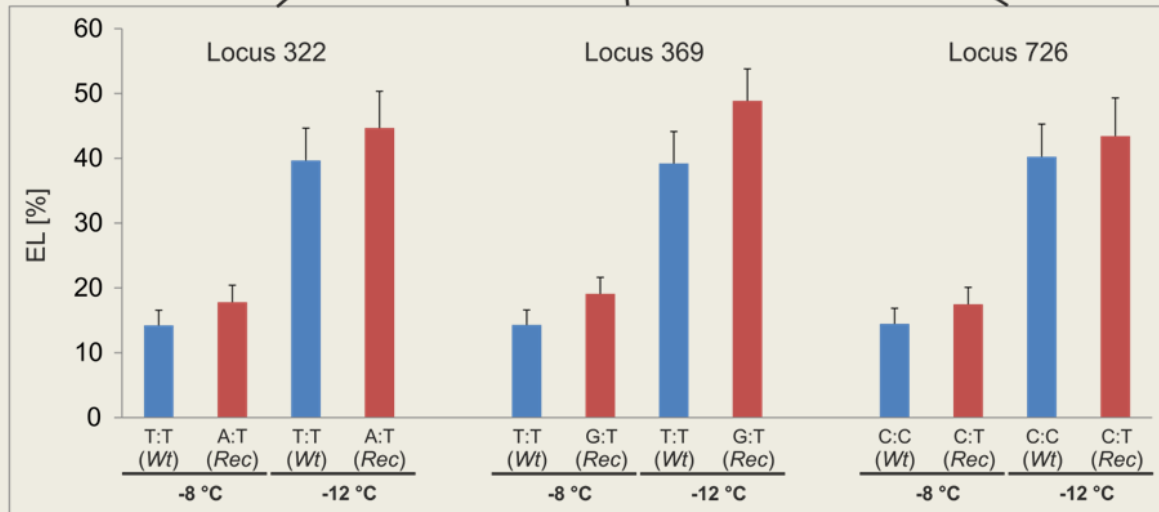
Marker-trait associations in *LpIRI1*

(A)



Genotype	Number of accessions	Locus 322	Locus 369	Locus 726	EL at -8 °C [%]	EL at -12 °C [%]
<i>Wt</i>	52	T:T	T:T	C:C	14.02	39.45
<i>Rec-1</i>	5	A:T	G:T	C:C	18.16	46.94
<i>Rec-2</i>	4	T:T	T:T	C:T	16.82	42.41
<i>Rec-3</i>	6	A:T	T:T	C:T	14.83	34.86
<i>Rec-4</i>	8	A:T	G:T	C:T	19.78	50.32

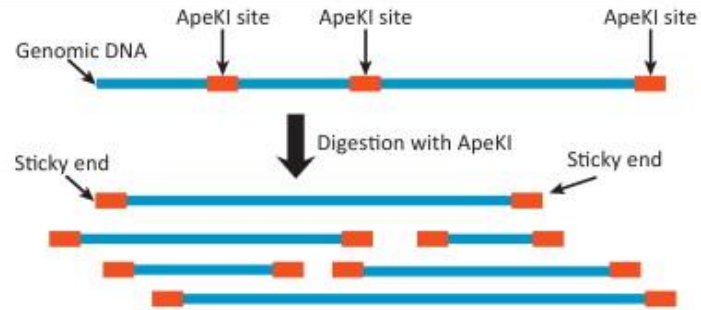
(B)



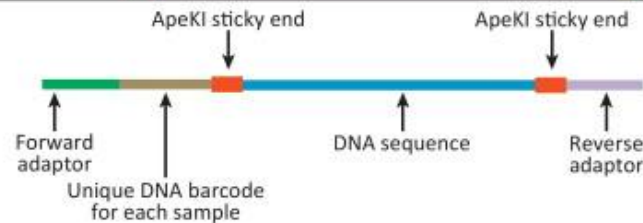
Wt – wild type
Rec – recombinant
 EL – electrolyte leakage

Genotyping by Sequencing (GBS)

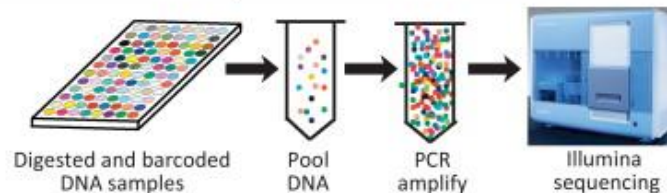
Step 1
Construct reduced representation libraries (RRLs) by digesting each DNA sample with a restriction enzyme (ApeKI)



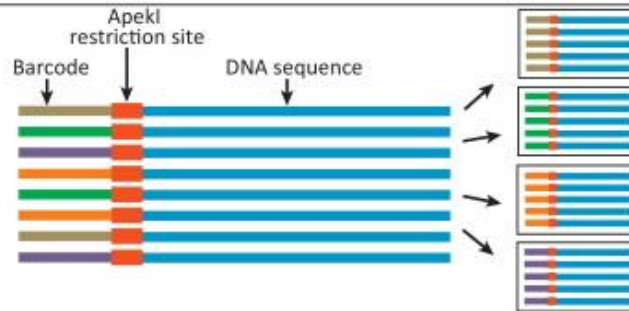
Step 2
Ligate custom 'barcoded' adaptors to sticky ends of restriction site. Each sample has its own unique barcode sequence



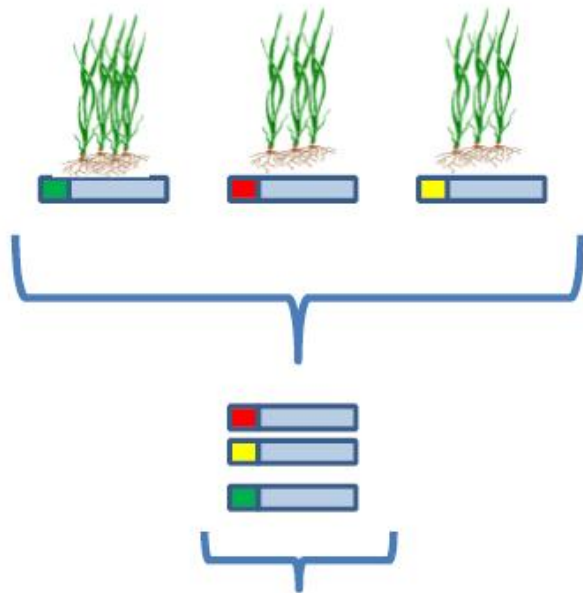
Step 3
Pool digested and barcoded DNA into a single tube. Perform PCR amplification, library preparation, and sequencing on Illumina platform



Step 4
Use barcodes to assign sequences to samples. Produce a file of DNA sequence data for each sample



Genotyping of 380 PPP populations

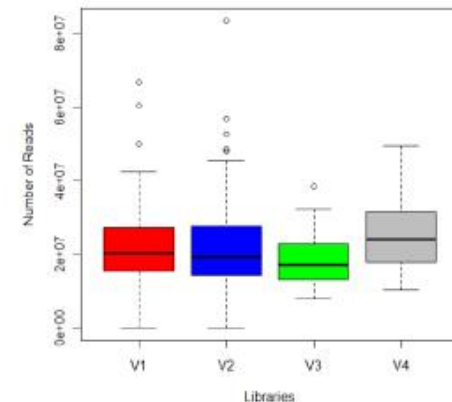


4 libraries, each with 88 populations, 1 library with 28 populations

Each Library
sequenced on 12 (4) lanes

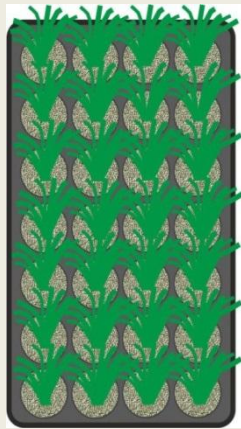


Boxplot of Reads per population



Evaluation of LT_{50} value in perennial ryegrass for PPP (workflow)

1 . The collection



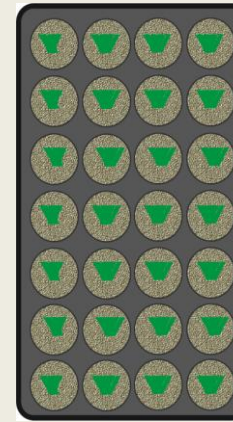
20 seed per population;
3 replicates;
soilless substrate;
30.000 plants in total.

2 . Cold-acclimation



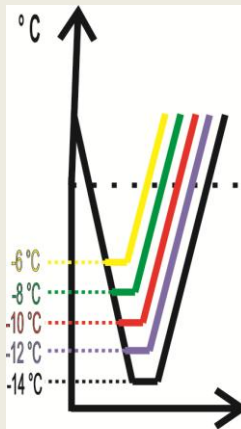
At +2 °C for
15 days,
12/12 h
photoperiod.

3 . Sample preparation



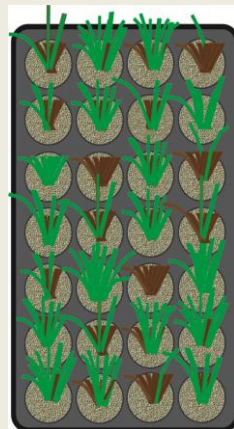
Leaves will be
trimmed and
the tillers will
be counted
prior the
freezing test.

4 . Freezing test



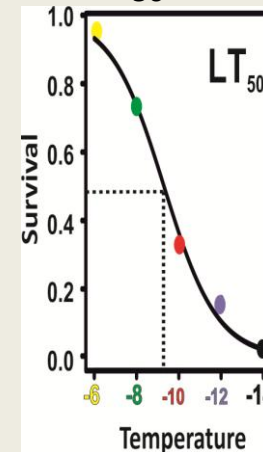
6 freezing
temperatures;
temperature
probes in
each cell pack.

5 . Regrowth



After freezing
plants will be
moved to
phytotron;
regrowth will
be evaluated
after 21 days.

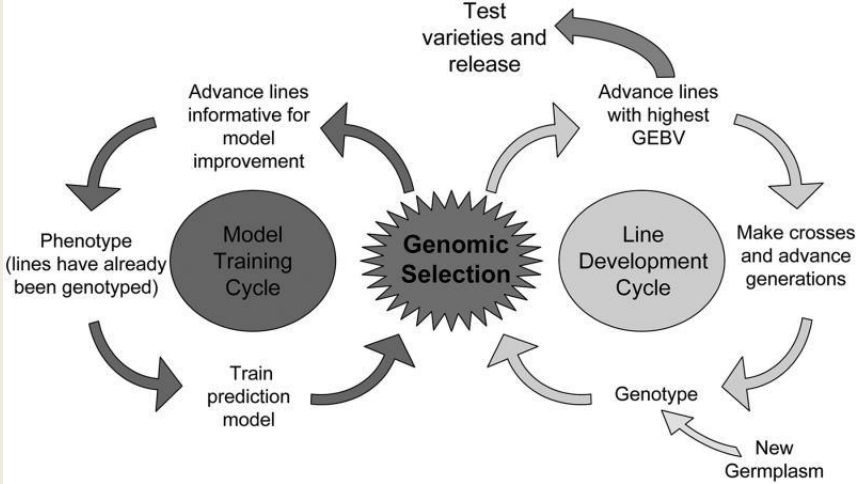
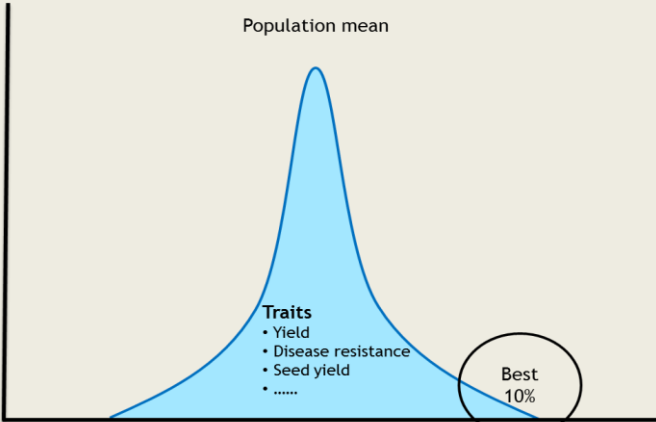
6 . LT_{50} calculation



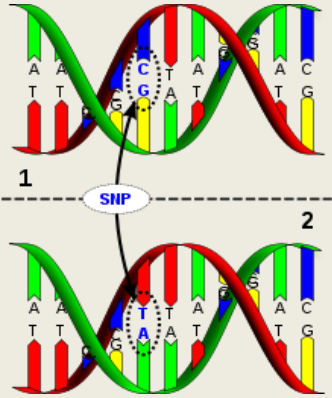
LT_{50} values will be
calculated for
each population.



GWAFF and Genomic Selection



Heffner et al., 2009



Acknowledgements

