

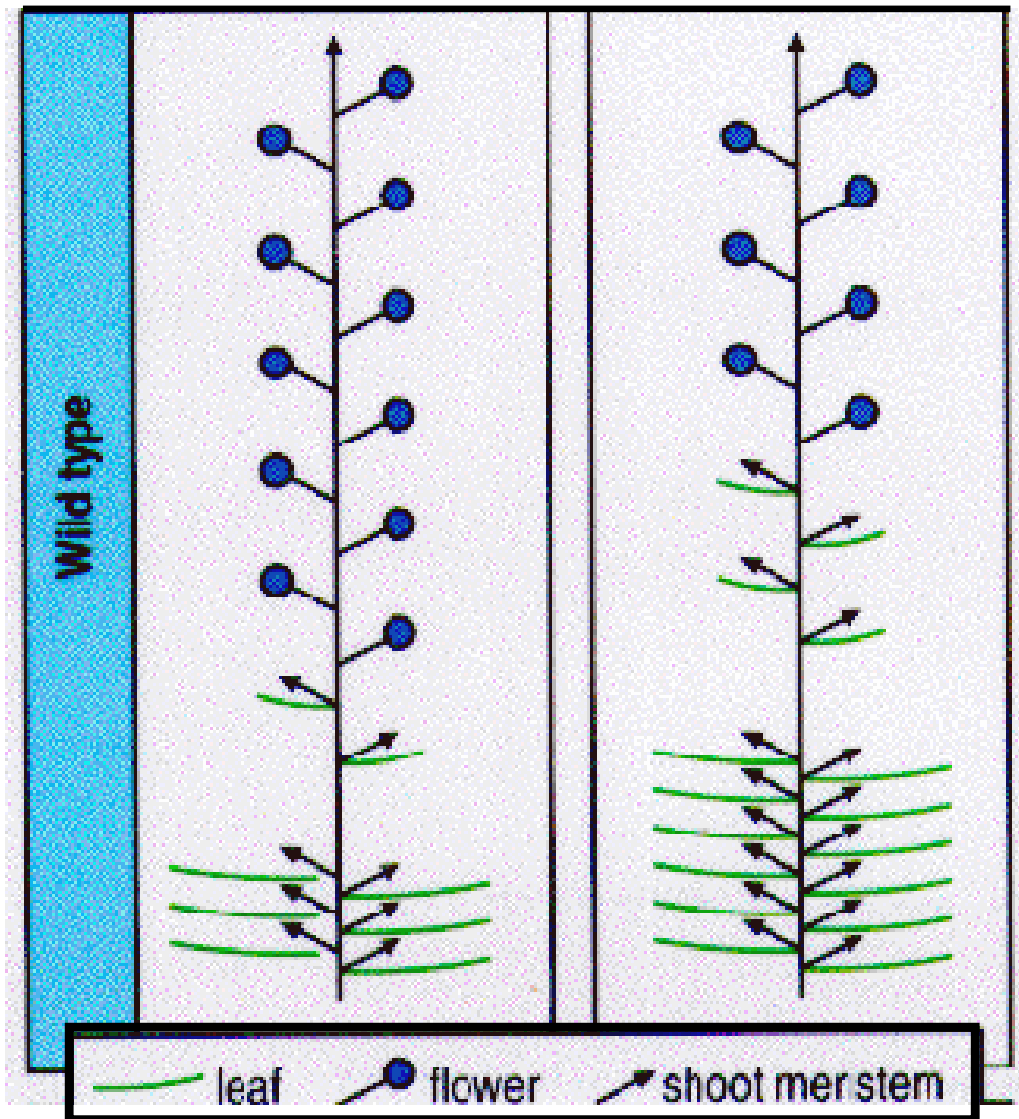
How is the transition from vegetative growth to flowering controlled ?

- How is it regulated by environmental conditions?

Environmental signals can influence the identity of the lateral organs formed at each node

Environment
promotes flowering

Environment
delays flowering

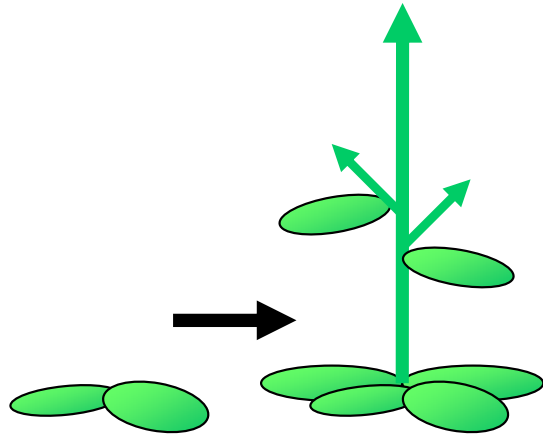


Environments that
Promote flowering
Of Arabidopsis are

-long daylengths,
photoperiod

-Long exposure to low
Temperature,
vernalization

Long-day plants flower when daylength exceeds a critical daylength



Examples:
Arabidopsis, wheat,
barley, sugar beet

Long days

Flowering

Short days

No Flowering

Two classes of mutation reduce the response to daylength

Long days



Wild-type



constans

Late-flowering in long days

No effect in short days

constans, gigantea, ft, cry2

Short days



Wild-type

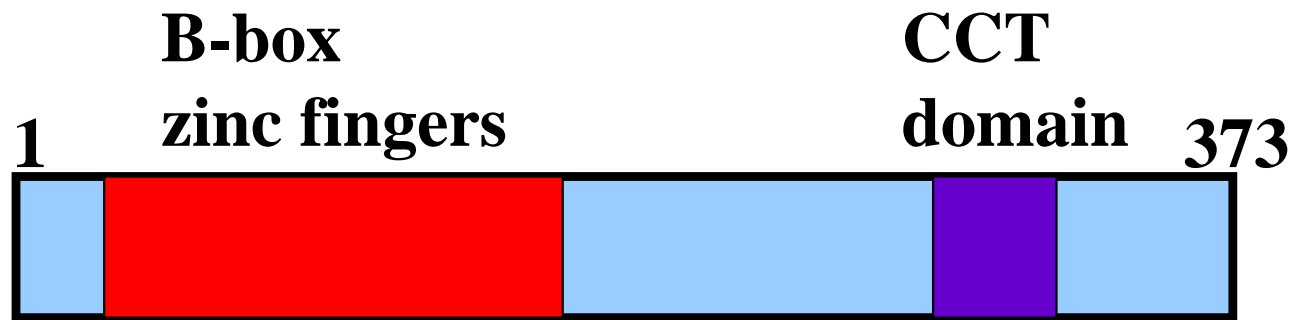
lhy-14

Early-flowering in short days

No effect in long days

lhy, toc1

CONSTANS has two motifs that are required for its function



**B-box proteins act in protein complexes
that regulate transcription in animal cells**

**CCT is plant specific named after CONSTANS,
CONSTANS-like and TOC1.**

CO Expression Under Long and Short Days

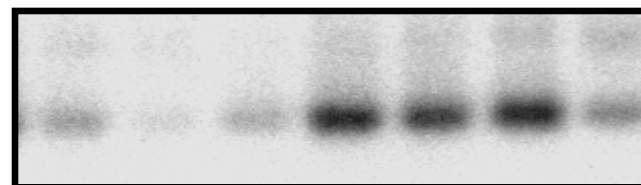
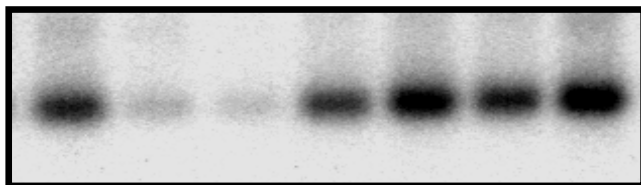
LONG DAYS

SHORT DAYS

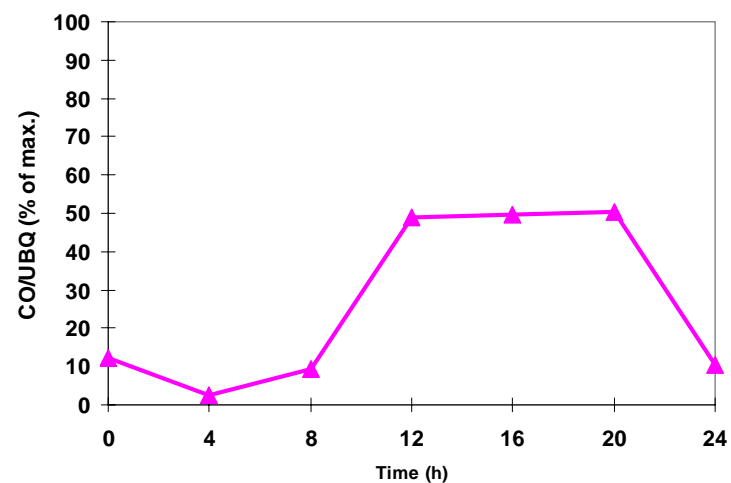
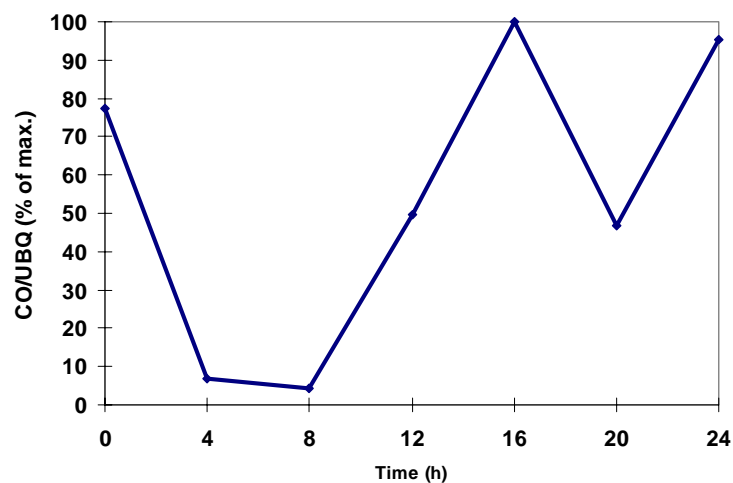
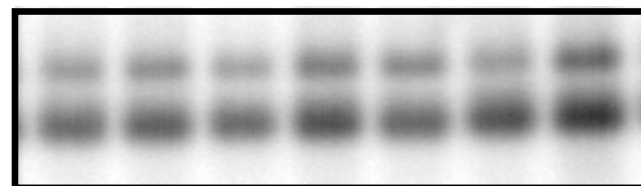
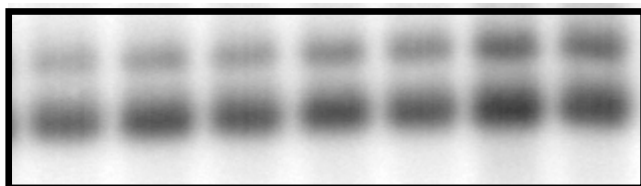
Time 0 4 8 12 16 20 24

0 4 8 12 16 20 24

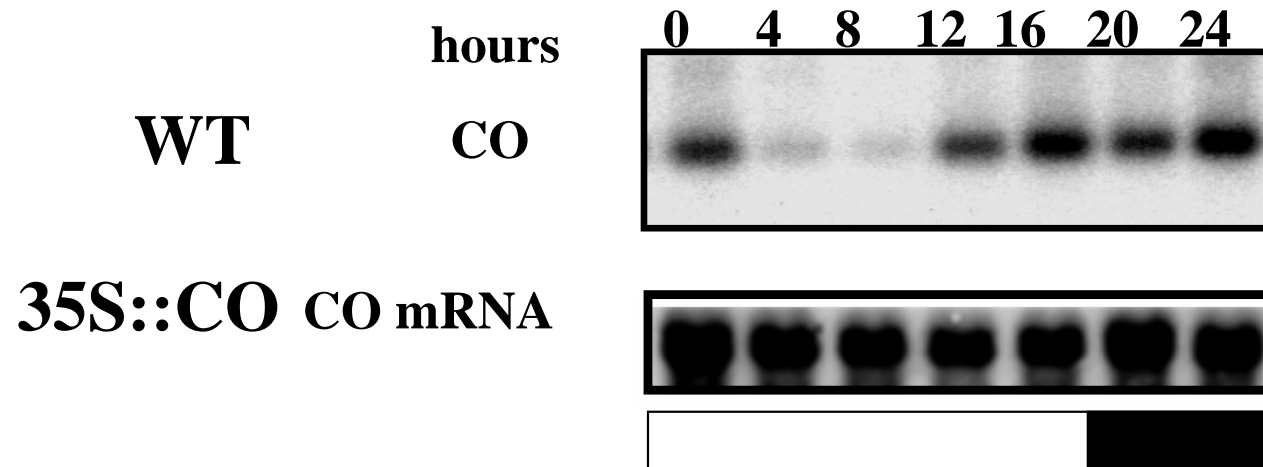
CO



UBQ

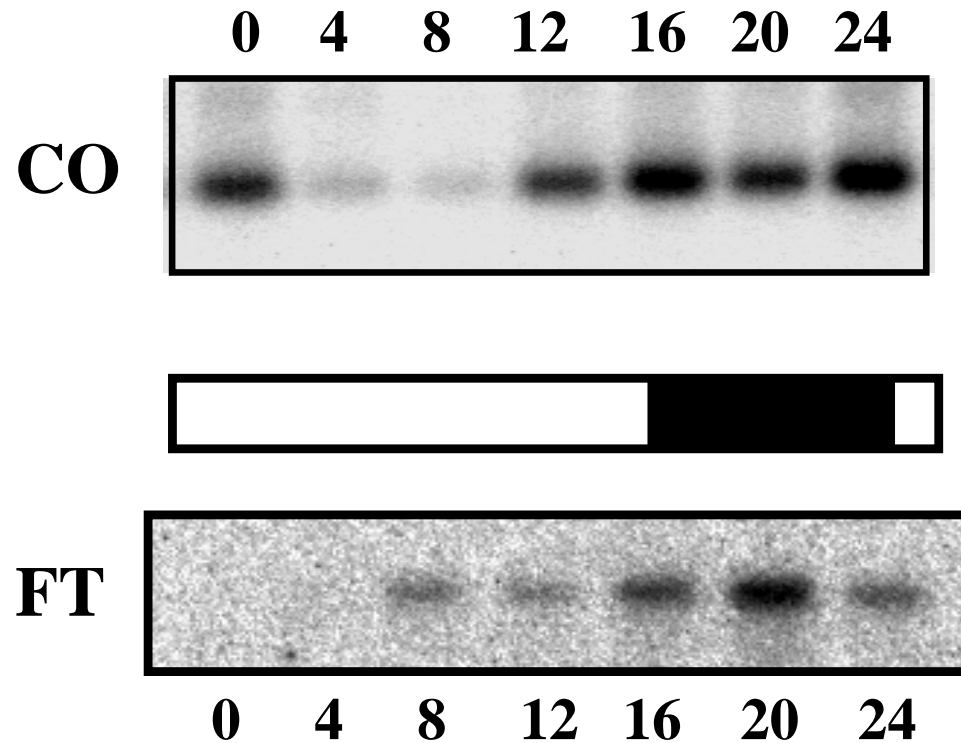


**CO mRNA is expressed throughout the day
and at higher levels in 35S::CO plants,
and this causes early flowering in all daylengths**



		Leaf Number
35S::CO		
Long Days		5.0
Short Days		4.8
Wild-type		
Long Days		8.6
Short Days		27.0

Comparison of diurnal rhythms in CO mRNA and mRNA of FT, a target gene of CO



FT expression –

Reduced in co mutants

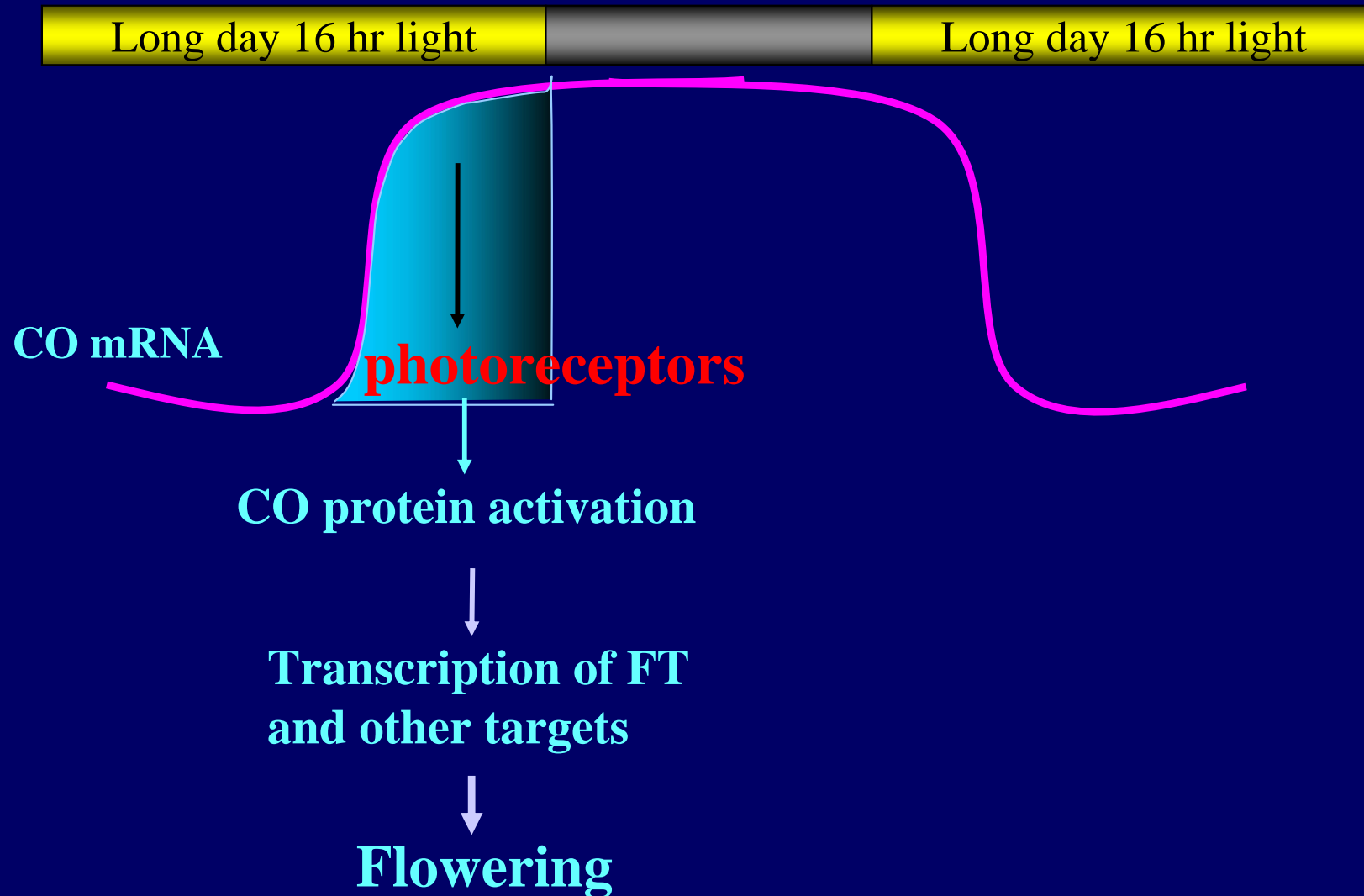
Reduced in short days

Overexpressed in 35S::CO

ft mutants similar

phenotype to co mutants

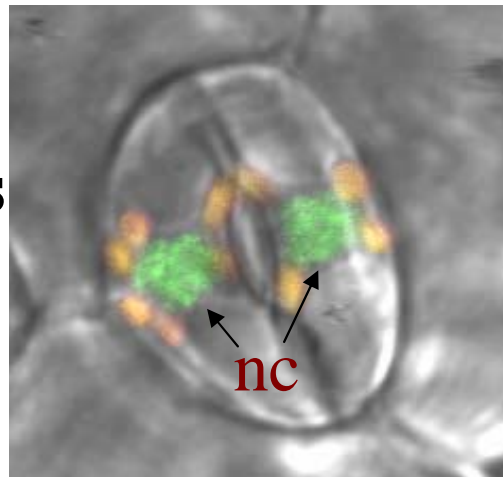
A model for the response of Arabidopsis to daylength



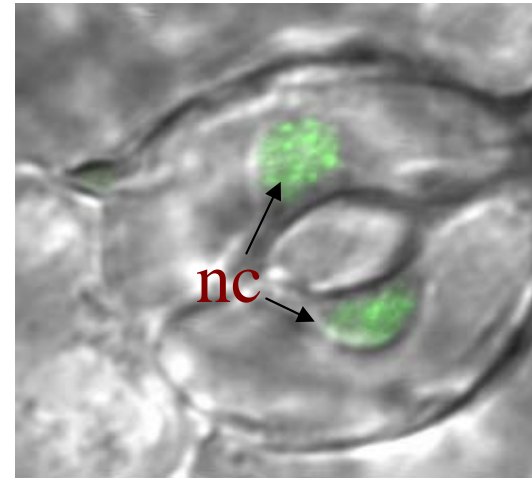
Exposure to light stabilises CO protein in the nucleus

35S::GFP:CO plants exposed to different light qualities

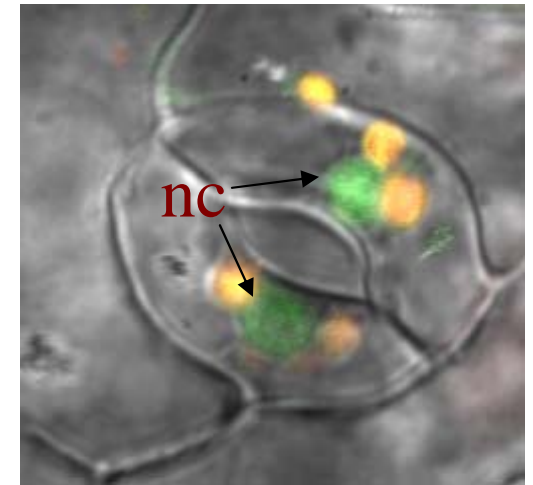
Light conditions
in which CO is
active



White

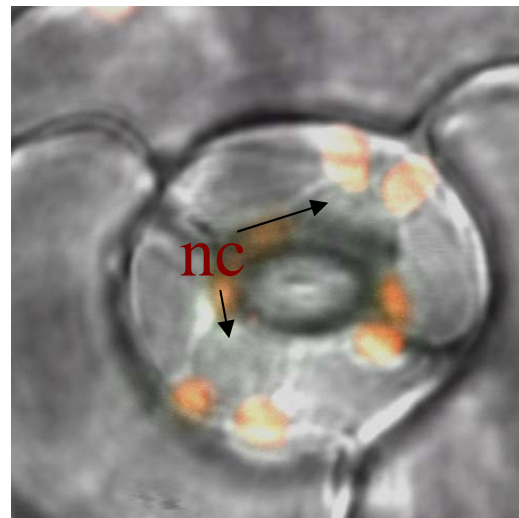


Blue

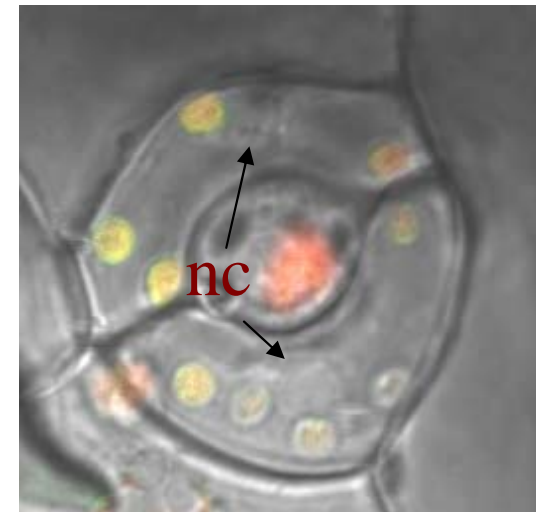


Far-red

Light conditions
in which CO is
inactive

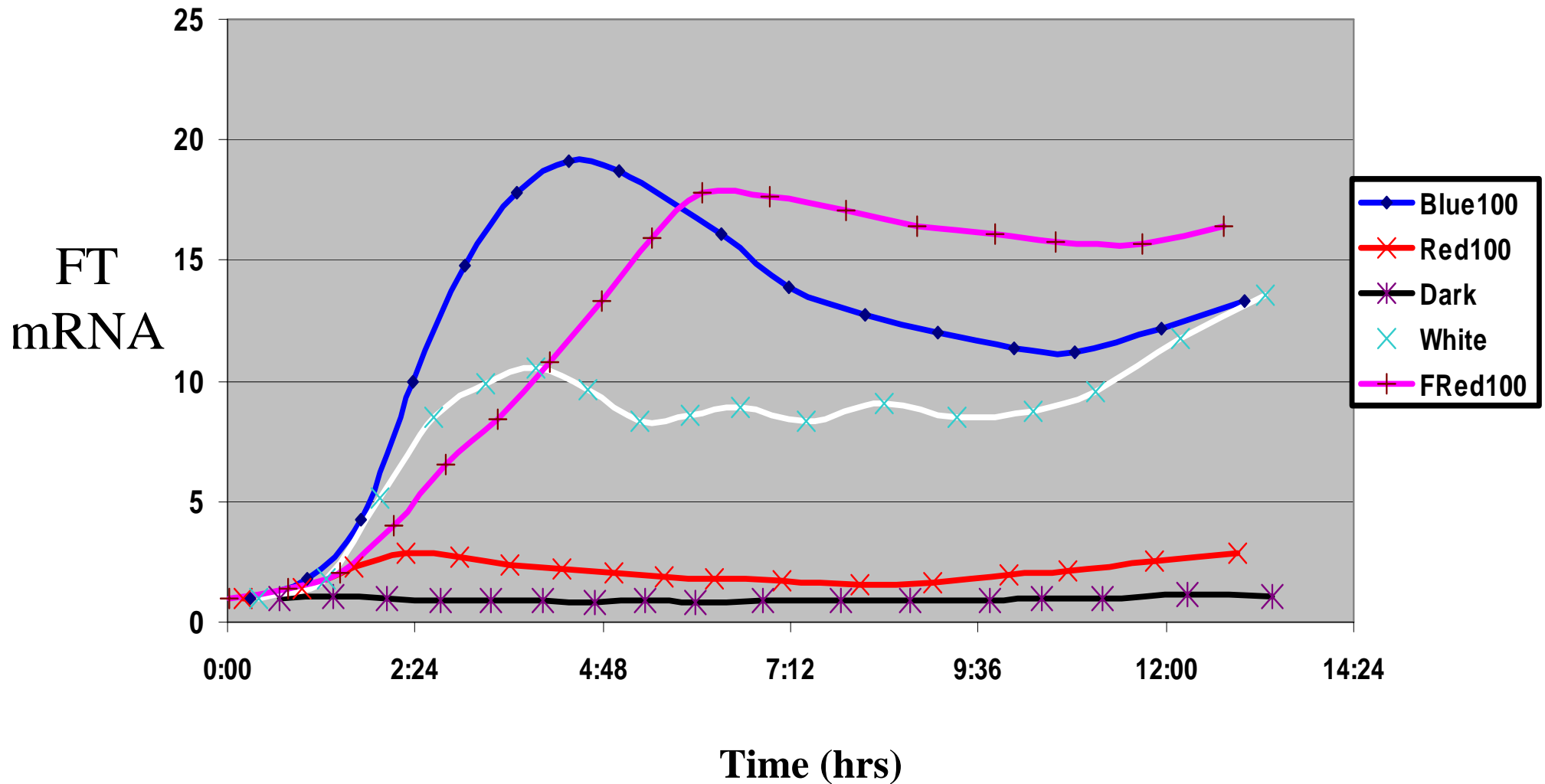


Dark

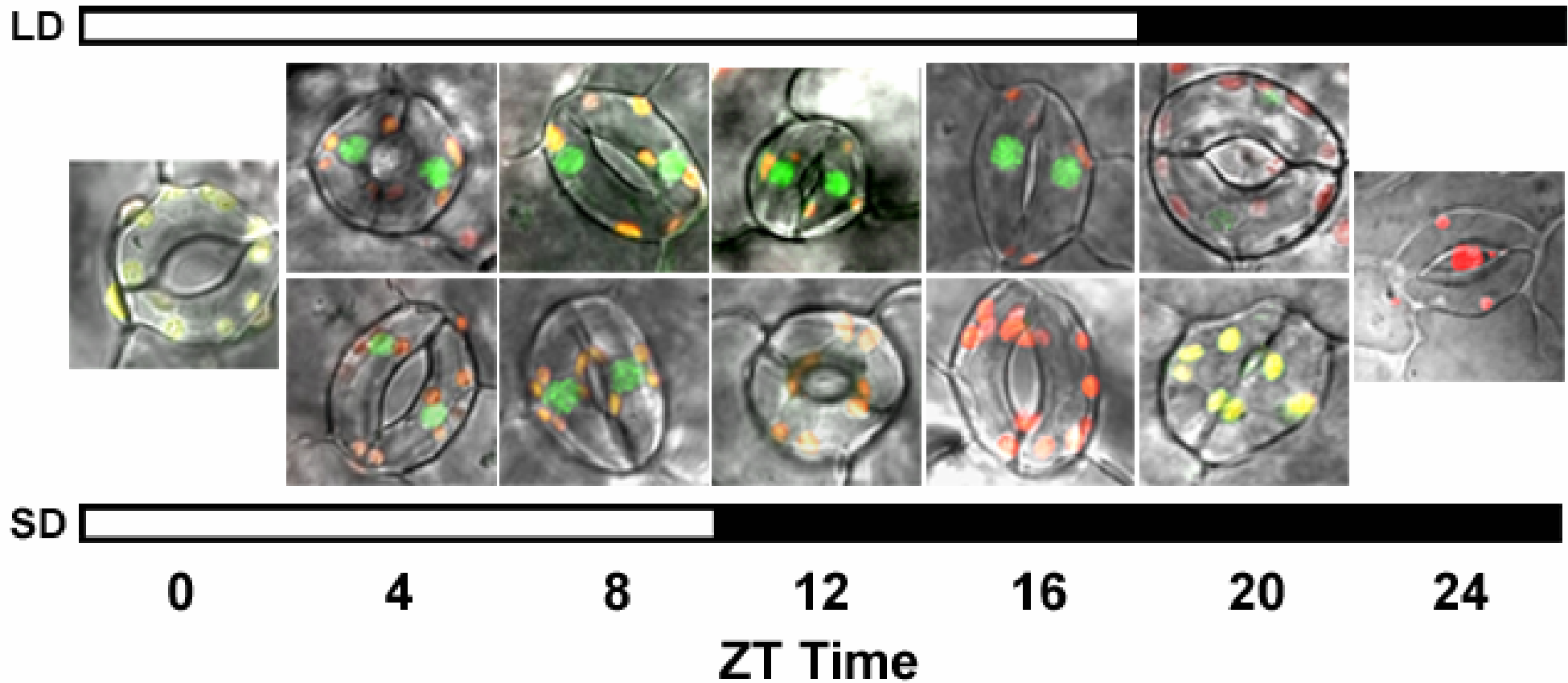


Red

Activation of FT by 35S::CO occurs in blue and far-red light, but not red light

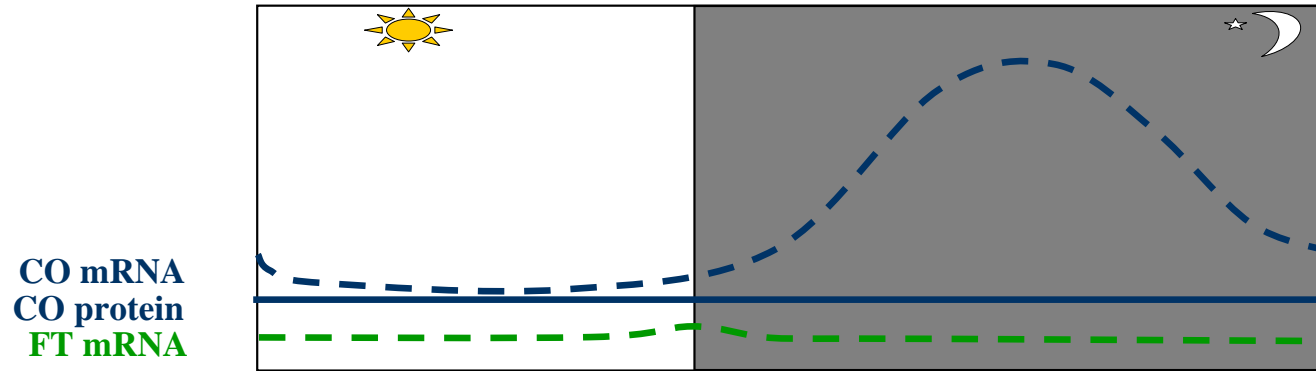


CO:GFP accumulates during the photoperiod in *35S::CO:GFP* plants

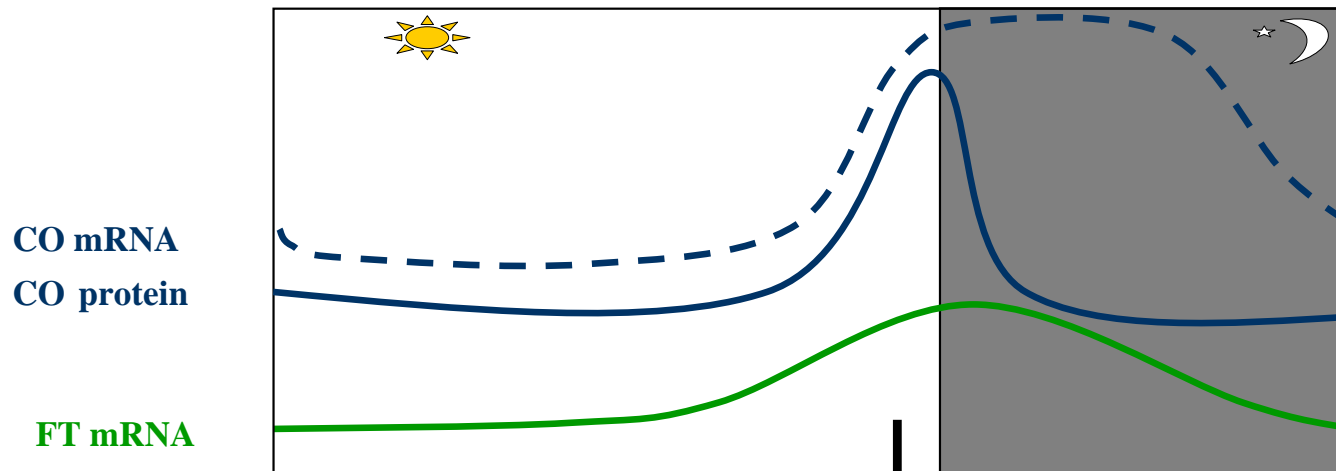


Coincidence of light and CO mRNA leads to stabilisation of CO protein and activation of FT under long days

No flowering
In short days



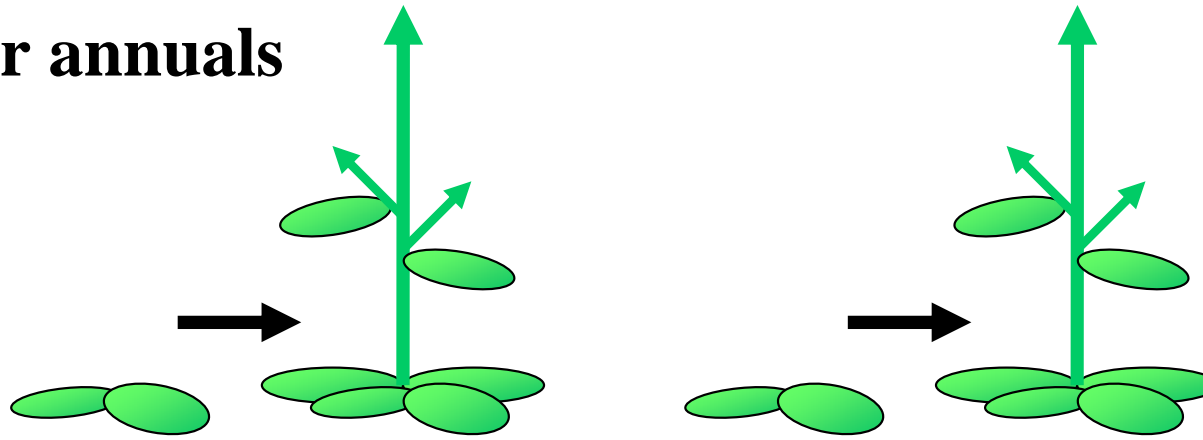
Flowering in
Long days



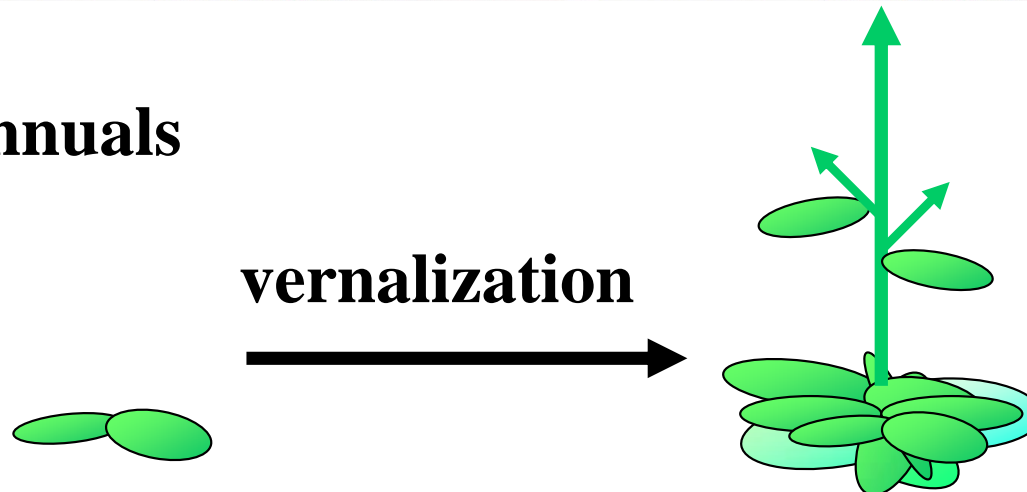
Flowering

Arabidopsis varieties differ in their reproductive strategies

Summer annuals



Winter annuals



Features of vernalization

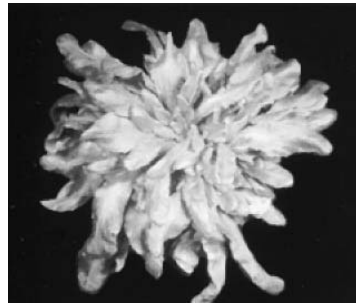
1. Occurs at shoot meristem
2. Requires several weeks of exposure to low temperatures.
This treatment called **vernalization**.
3. Effects of vernalization maintained through many mitotic divisions.
If a plant is vernalized as a young seedling this effect can be „remembered“ for several months.
4. Effects of vernalization reset at meiosis.
The progeny of a vernalized plant behave do not inherit the effect of vernalization, and behave as if they have not been vernalized.

Analysis of the genetic basis of vernalization response

No vernalization

8 weeks vernalization

**Santa Fe
(winter annual)**



**Columbia
(Summer annual)**



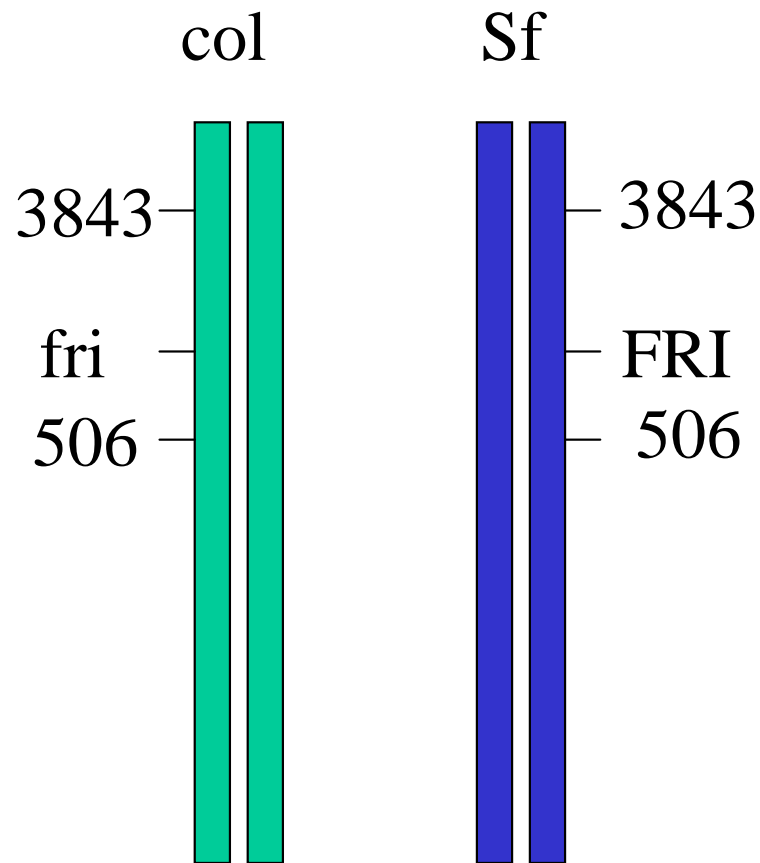
**To identify genes that confer a vernalization response
- intercross Santa Fe and Columbia**

Genetics of crossing Stockholm with Li-5

	Late flowering	Early flowering
Columbia	0	55
Santa Fe	40	0
Sf x Col	135	46

Indicates 3:1 segregation with late flowering dominant.

Gene that confers late flowering, FRIGIDA, also Occurs in other vernalization responsive varieties

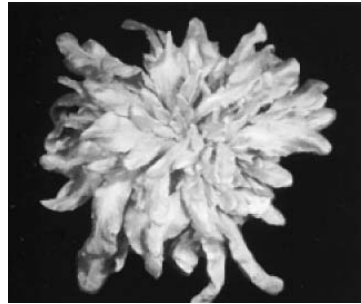


In crosses between other
Examples of vernalization
responsive and non-responsive
strains alleles of FRIGIDA
shown to underlie the difference.
Including European strains
Stockholm and Limburg-5.

Chromosome 4

**FRI encodes a 609 amino acid protein of unknown function
containing two coil-coil domains implicated
in protein protein interaction.**

How does FRIGIDA delay flowering?



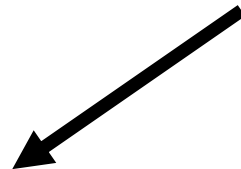
FRI plant – very late flowering



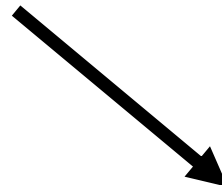
Mutagenise with radiation



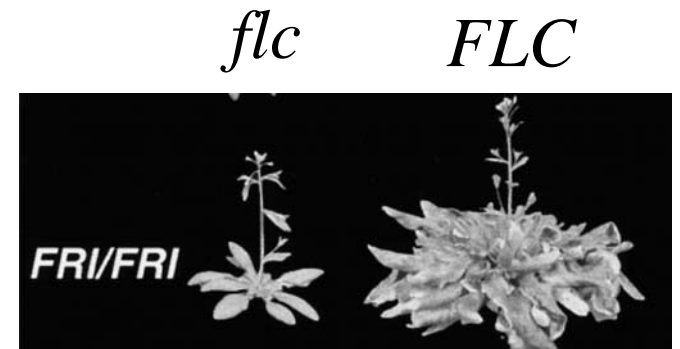
Screen for early flowering plants



**3 loss of function
Alleles of FRI**



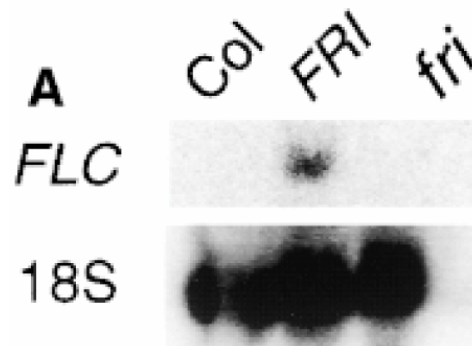
**4 loss of function
Alleles of FLC**



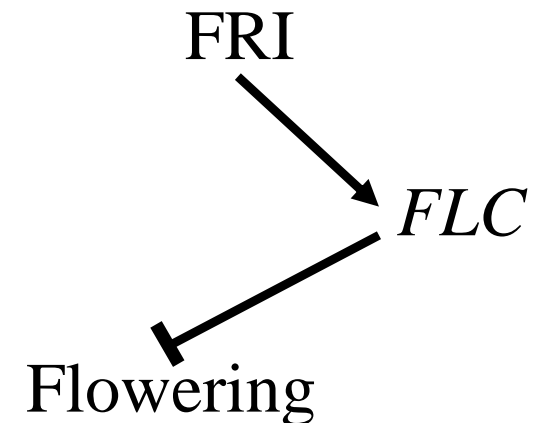
FLC encodes a MADS box transcription factor that represses flowering



Overexpression of FLC delays flowering



FRIGIDA promotes expression of the floral repressor FLC



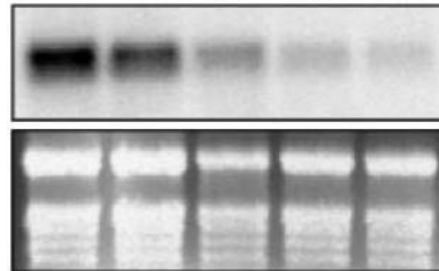
In winter annuals vernalization reduces abundance of the floral inhibitor FLC and accelerates flowering

***FRI FLC* plants**

Time at 4C

(days) 1 7 14 21 28

FLC



No vernalization

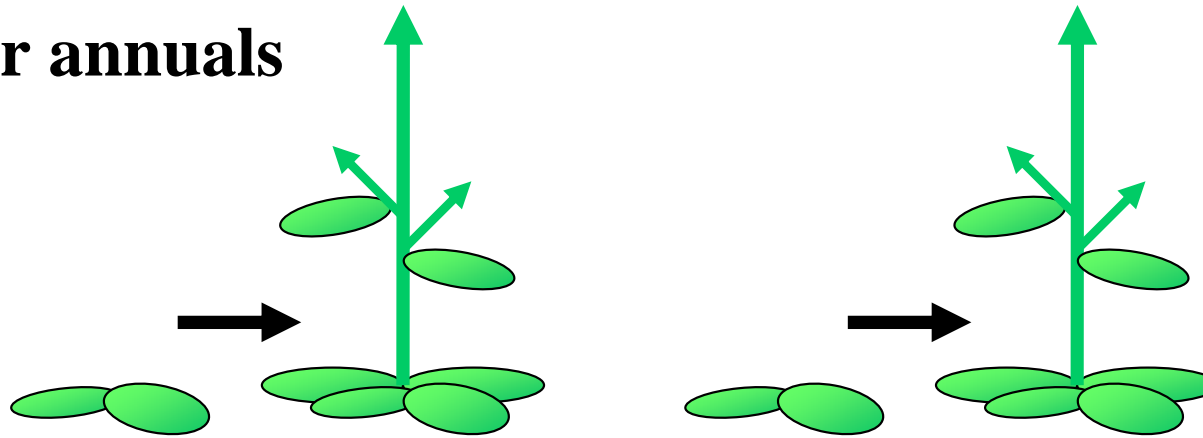
8 weeks vernalization

Santa Fe

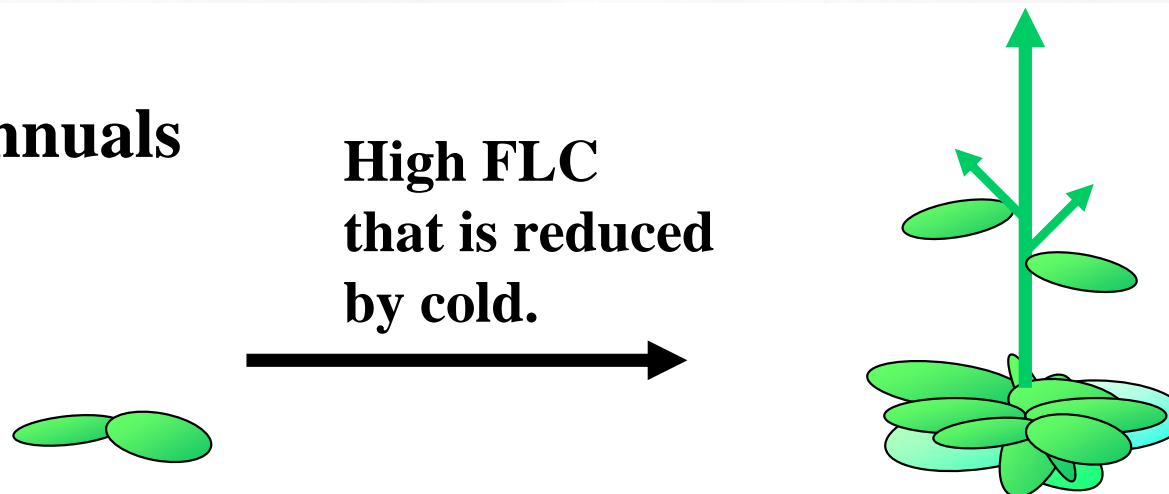


Arabidopsis varieties differ in their reproductive strategies

Summer annuals



Winter annuals



Identification of genes required for vernalization response

- how is FLC expression controlled by low temperatures?

Late flowering, high FLC



Treat with mutagen - EMS



**Give vernalization treatment to
all plants**

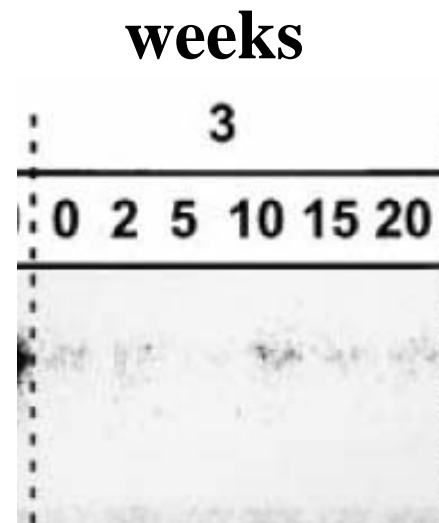
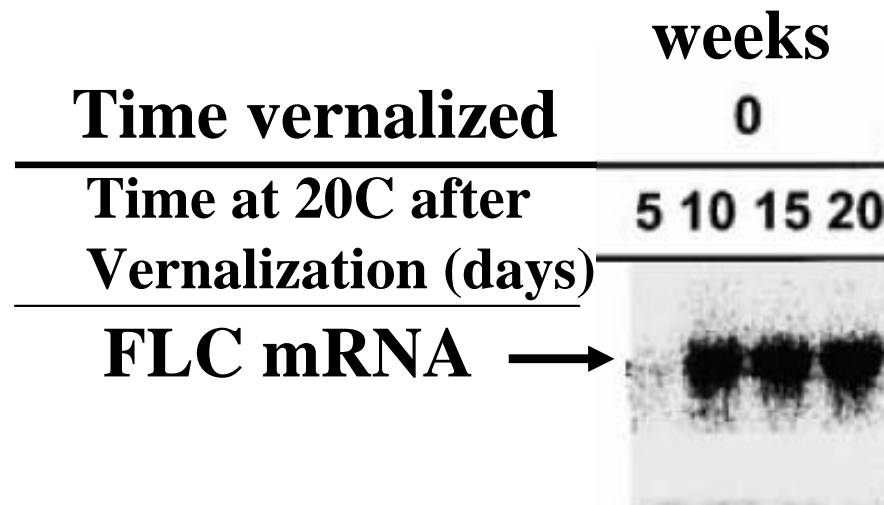


**Most plants flower
early.**

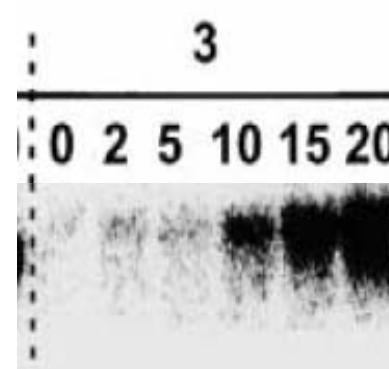
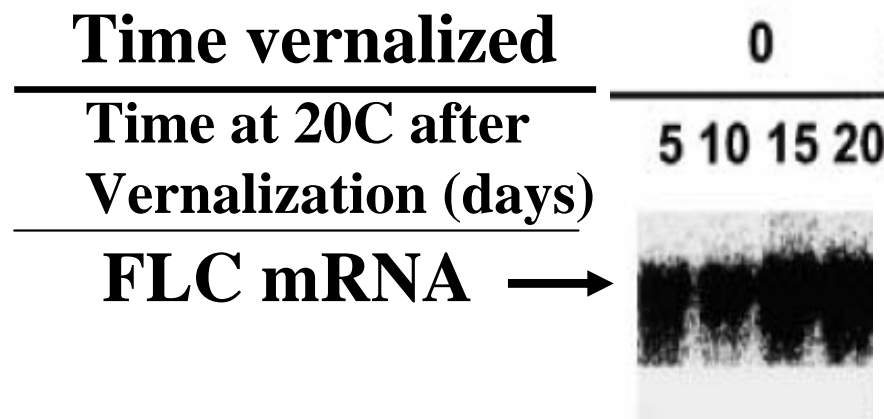


**Some plants flower late.
These are vernalization
and vernalization insensitive
mutants.**

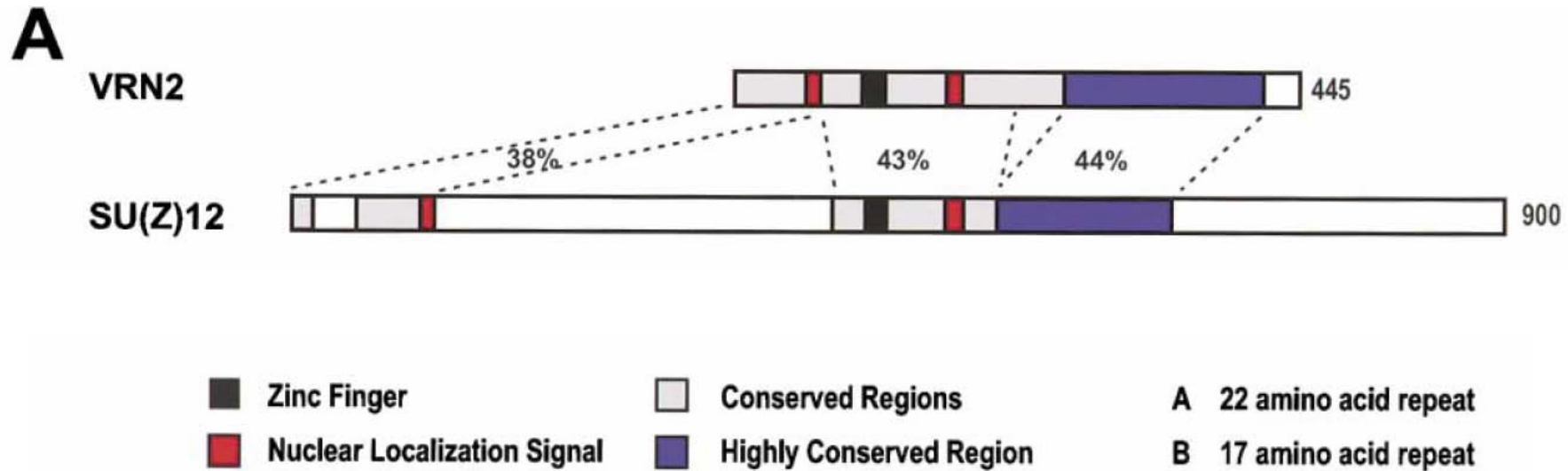
The role of VERNALIZATION2 is to maintain the repression of FLC expression



vrn2 mutant



VERNALIZATION2 encodes a gene related to *Drosophila* Polycomb-group (PcG) genes



In *Drosophila*, PcG proteins act in large protein complexes. They maintain the repression of transcription of homeotic genes, once the pattern of expression of these genes has been established during early embryo development.

Polycomb-group complexes in *Drosophila* repress gene expression by modifying histones.



A nucleosome contains two copies of H2A, H2B, H3 and H4 wrapped around 147 bp of DNA

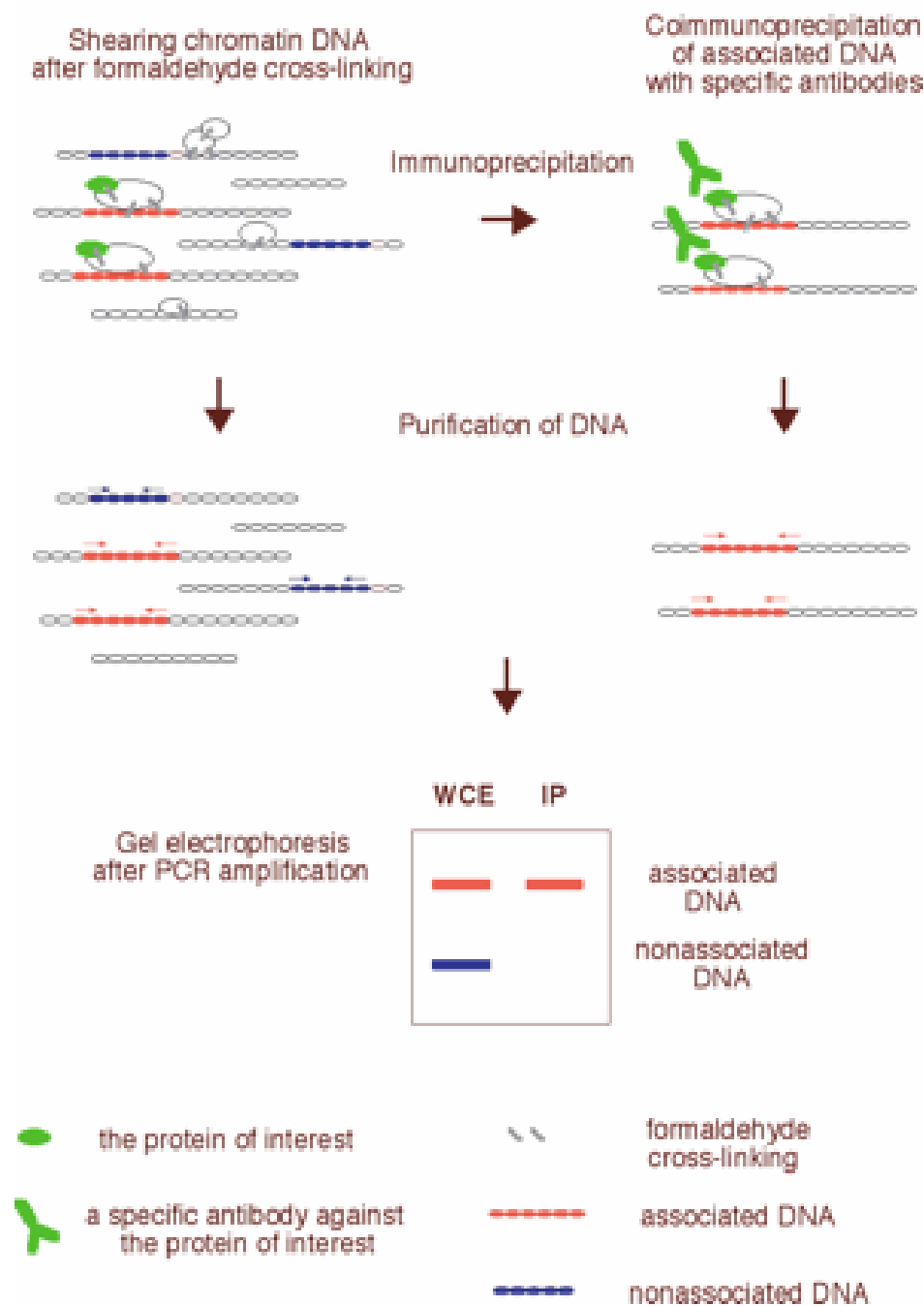


Modification of histones can alter gene expression.

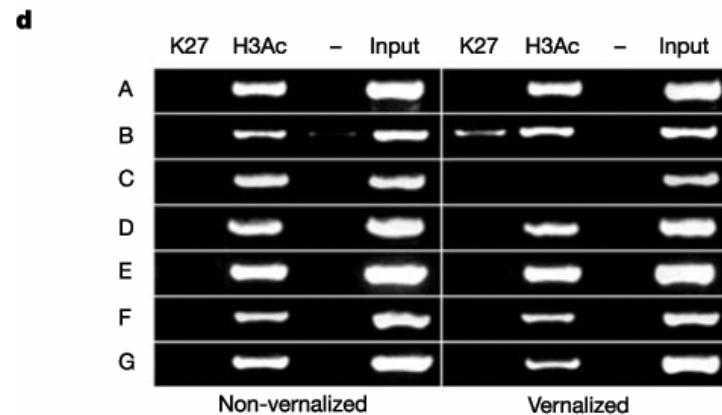
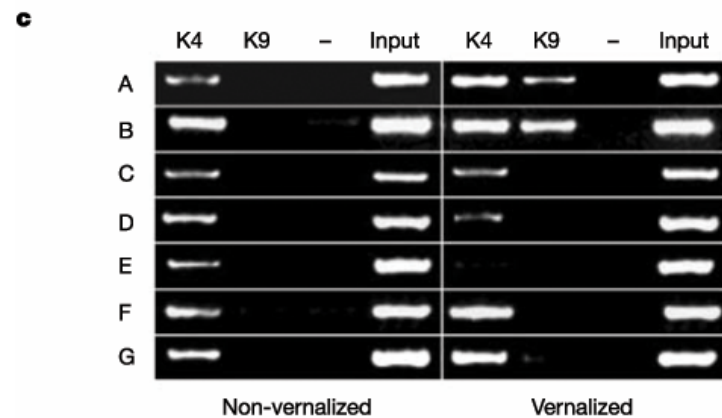
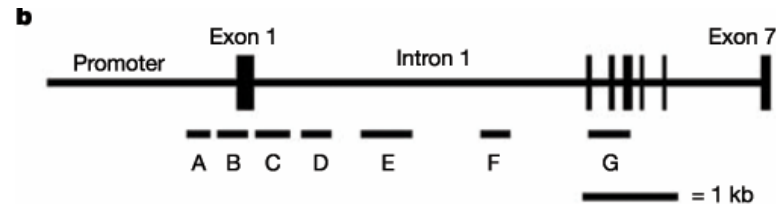
Histone 3 is a target for modifications – those above activate gene expression, those below repress it.

Polycomb-group proteins promote the methylation of K9 and K27

Chromatin immunoprecipitation (ChIP) to identify DNA bound by specific proteins

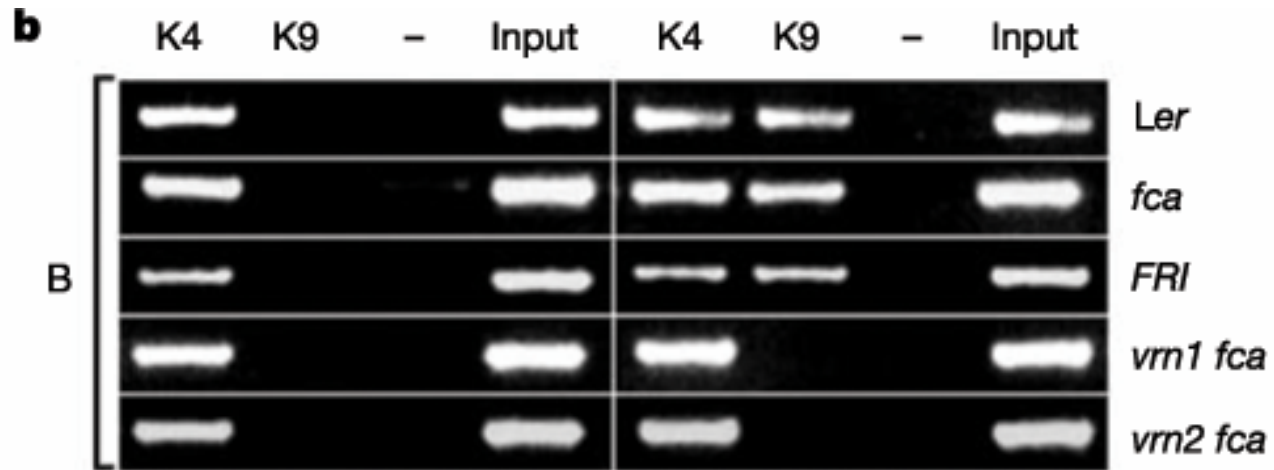


Methylated histones appear on the FLC gene after vernalization

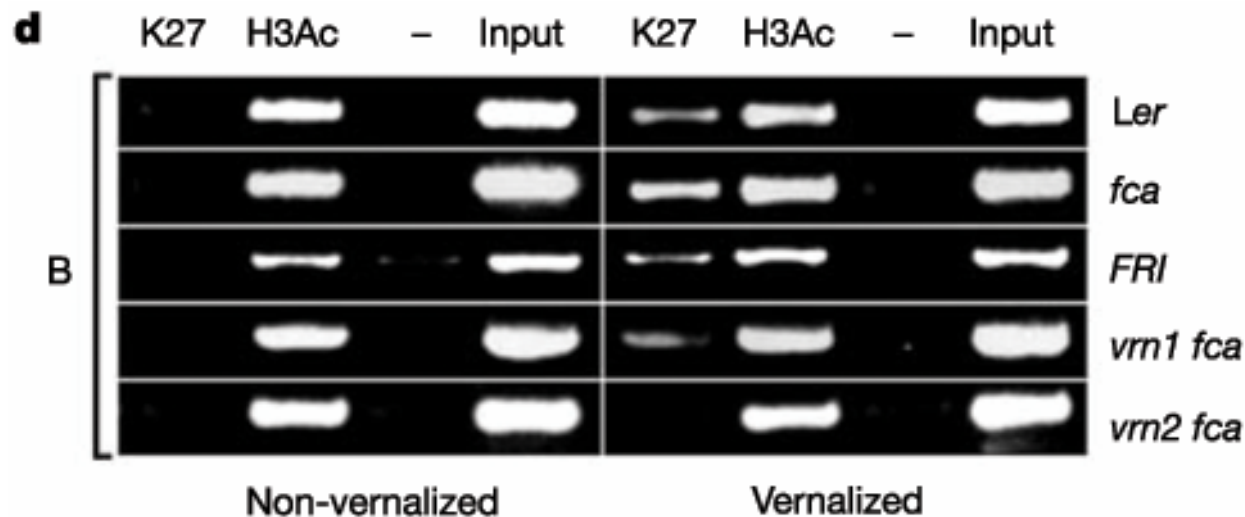


Methylation of H3
K9 and K27 appears only
After vernalization when
The FLC gene is
repressed

ChIP of B fragment at 5'end of FLC gene using Specific antibodies against modified H3 histones



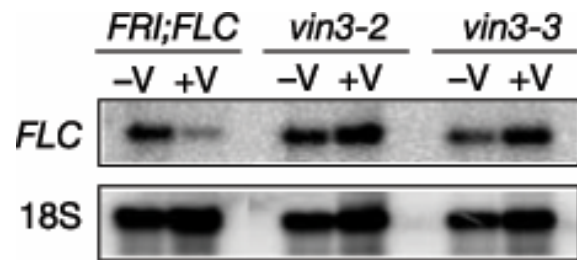
Methylation of K9
And K27 after
Vernalization
Requires VRN2



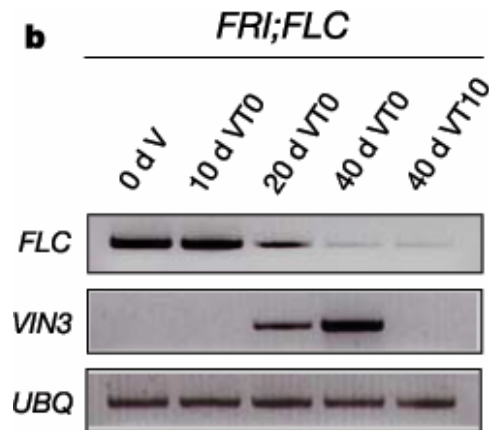
FLC expressed_F FLC repressed

L
C

VERNALIZATION INSENSITIVE 3 gene is required to reduce FLC



In the *vin3* mutant FLC expression
Is not reduced by cold



VIN3 expression rises during
Vernalization, suggesting has an
I early role in vernalization response

N
3

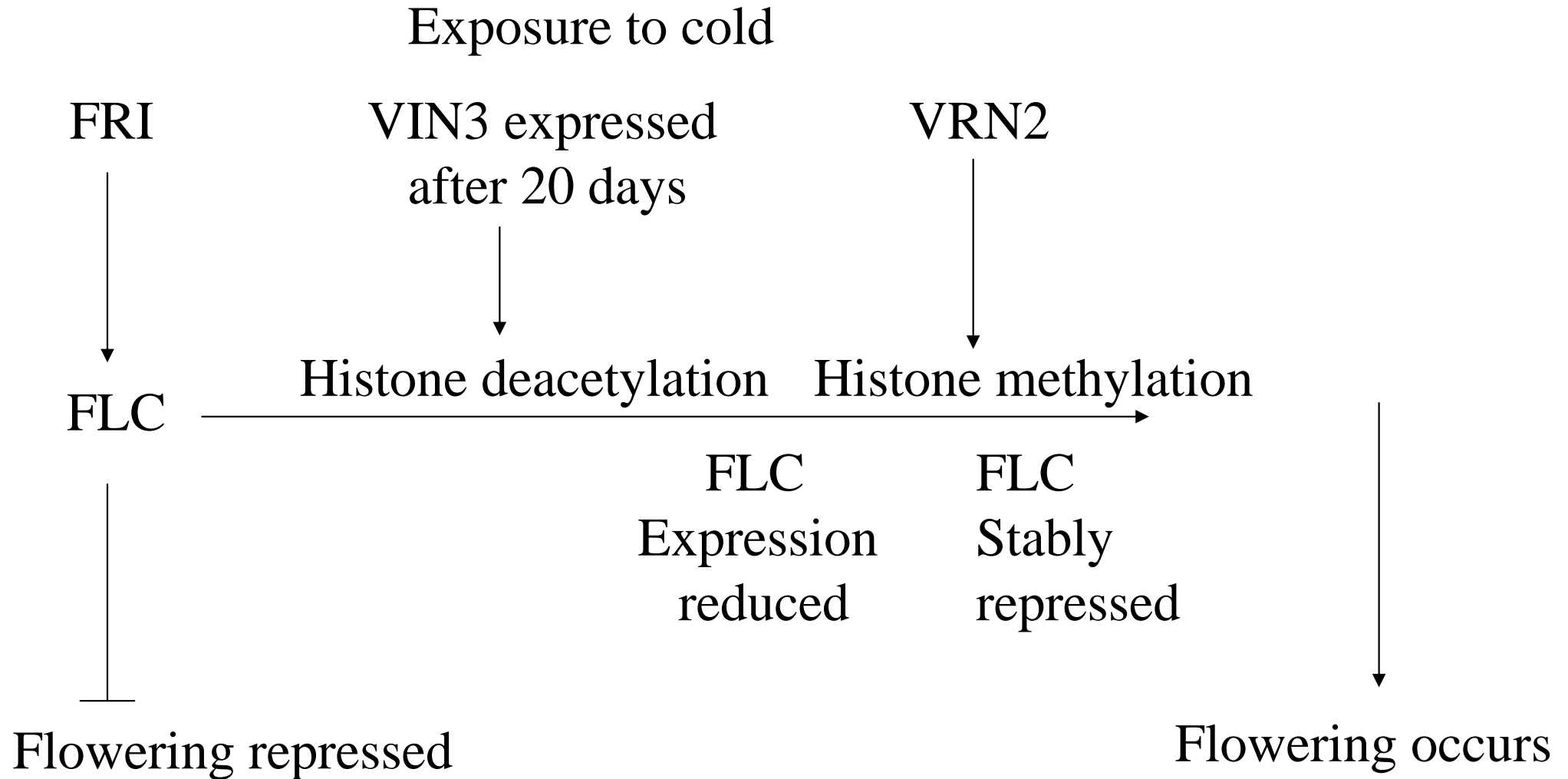
In the *vin3* mutant deacetylation and methylation of histones
on the FLC gene are blocked

x

p

e

Effect of vernalization on FLC expression and flowering



The photoperiod and vernalization responses converge on the same target genes: SOC1 and FT

