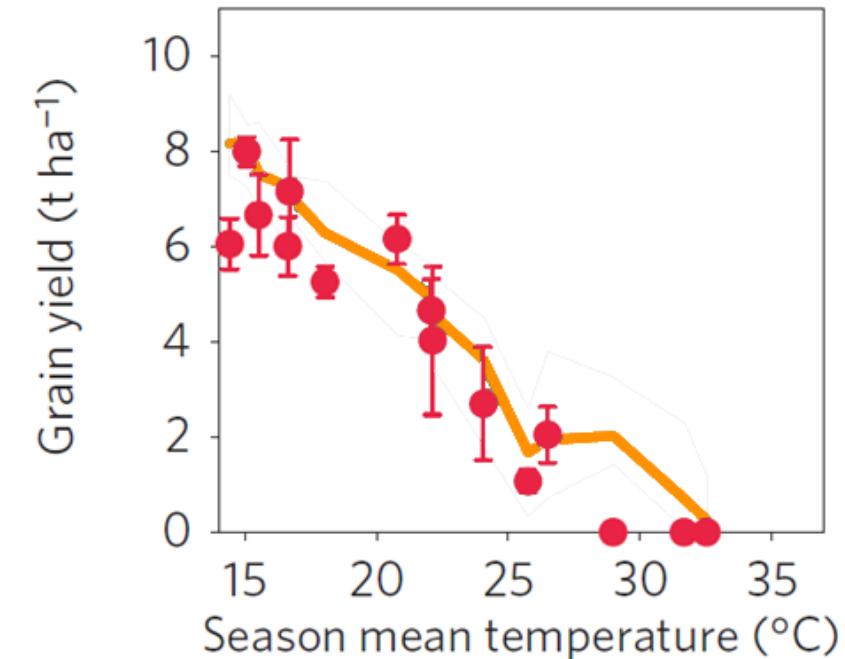
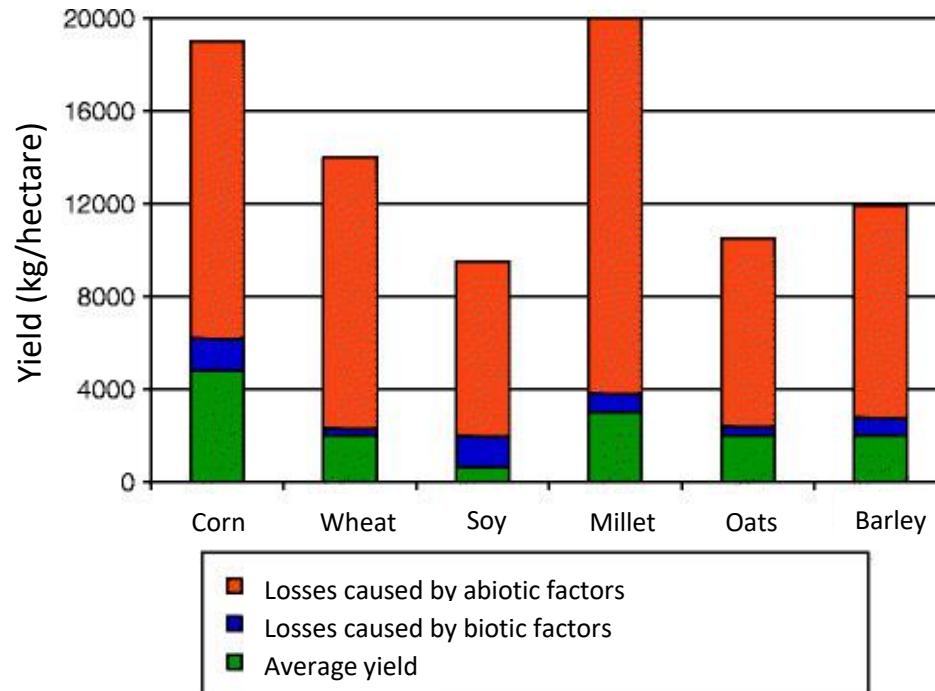
The background of the slide features a close-up photograph of a wheat field. The wheat ears are a vibrant golden color, indicating maturity. They are set against a clear, pale blue sky, which provides a sharp contrast to the yellow of the grain.

Comparative wheat phosphoproteome profiling pinpoints high temperature- associated breeding markers

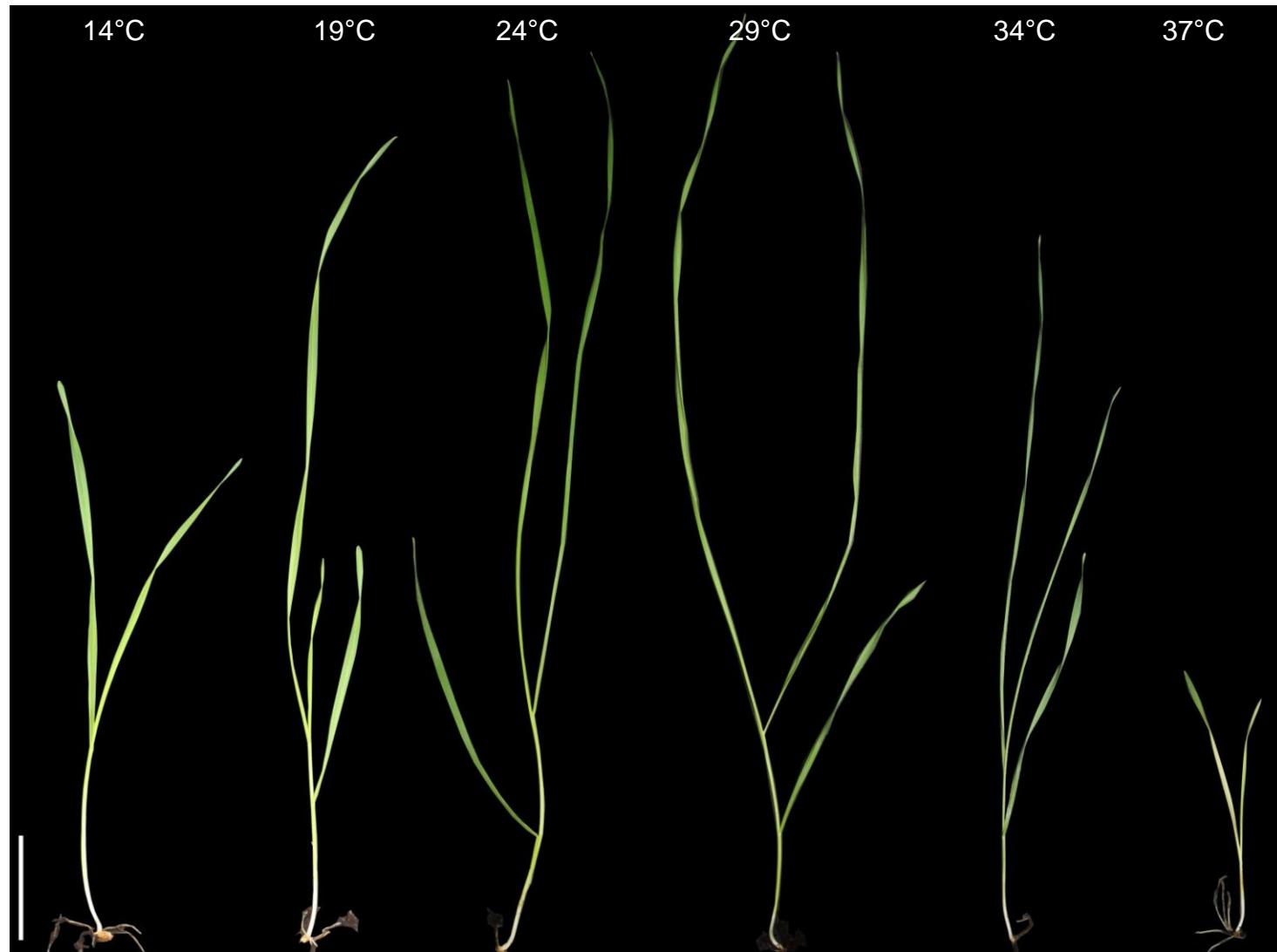
Ive De Smet

VIB-UGent Center for Plant Systems Biology

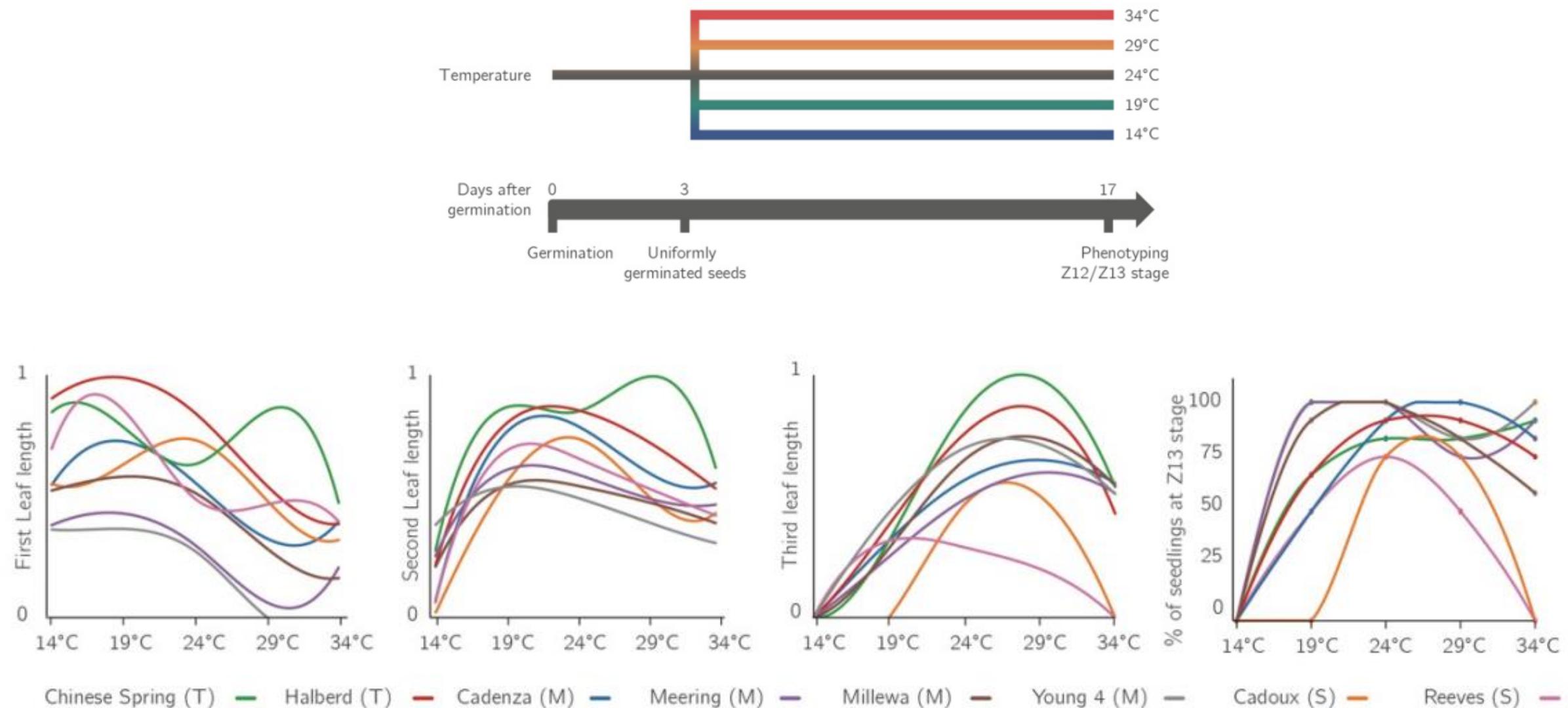
Abiotic stress impacts wheat yield



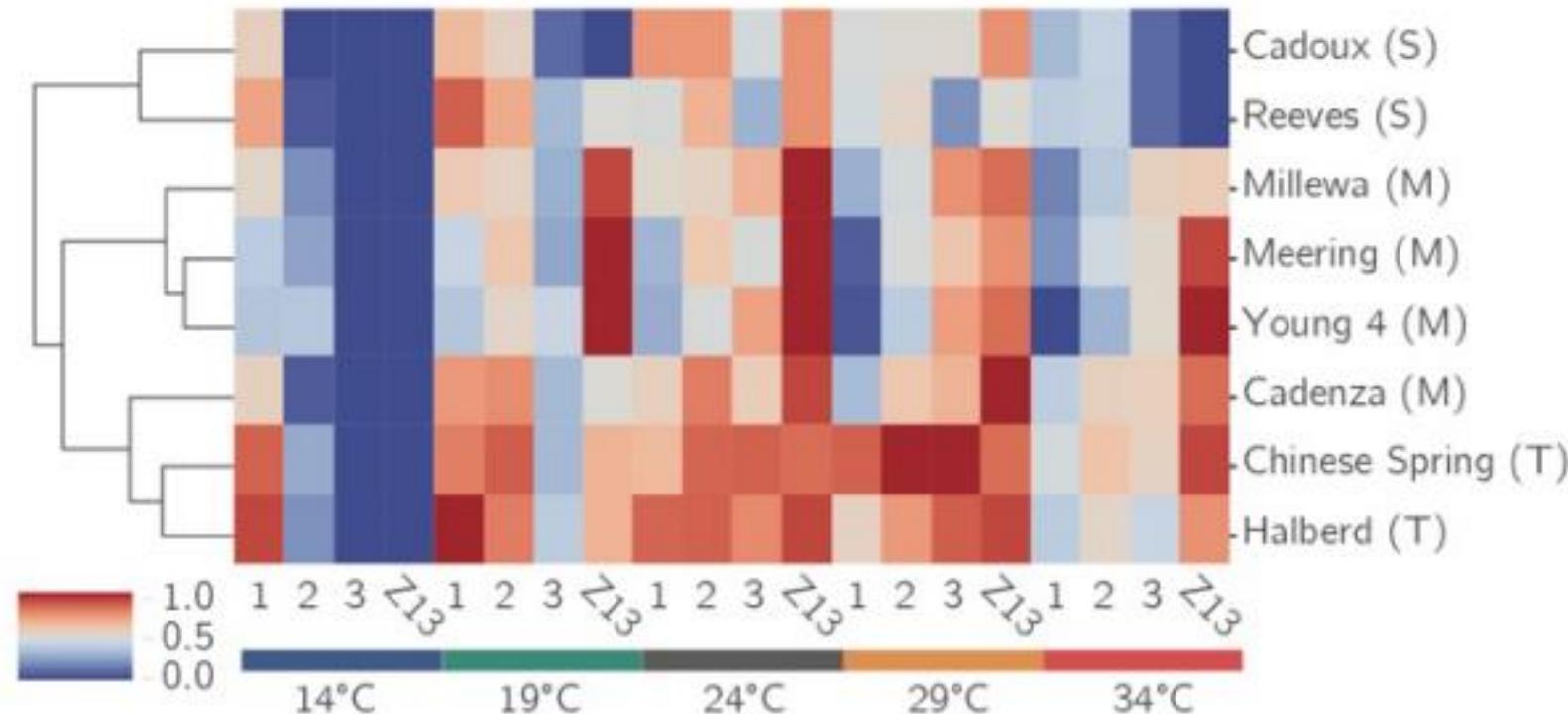
High temperature impacts wheat growth and development



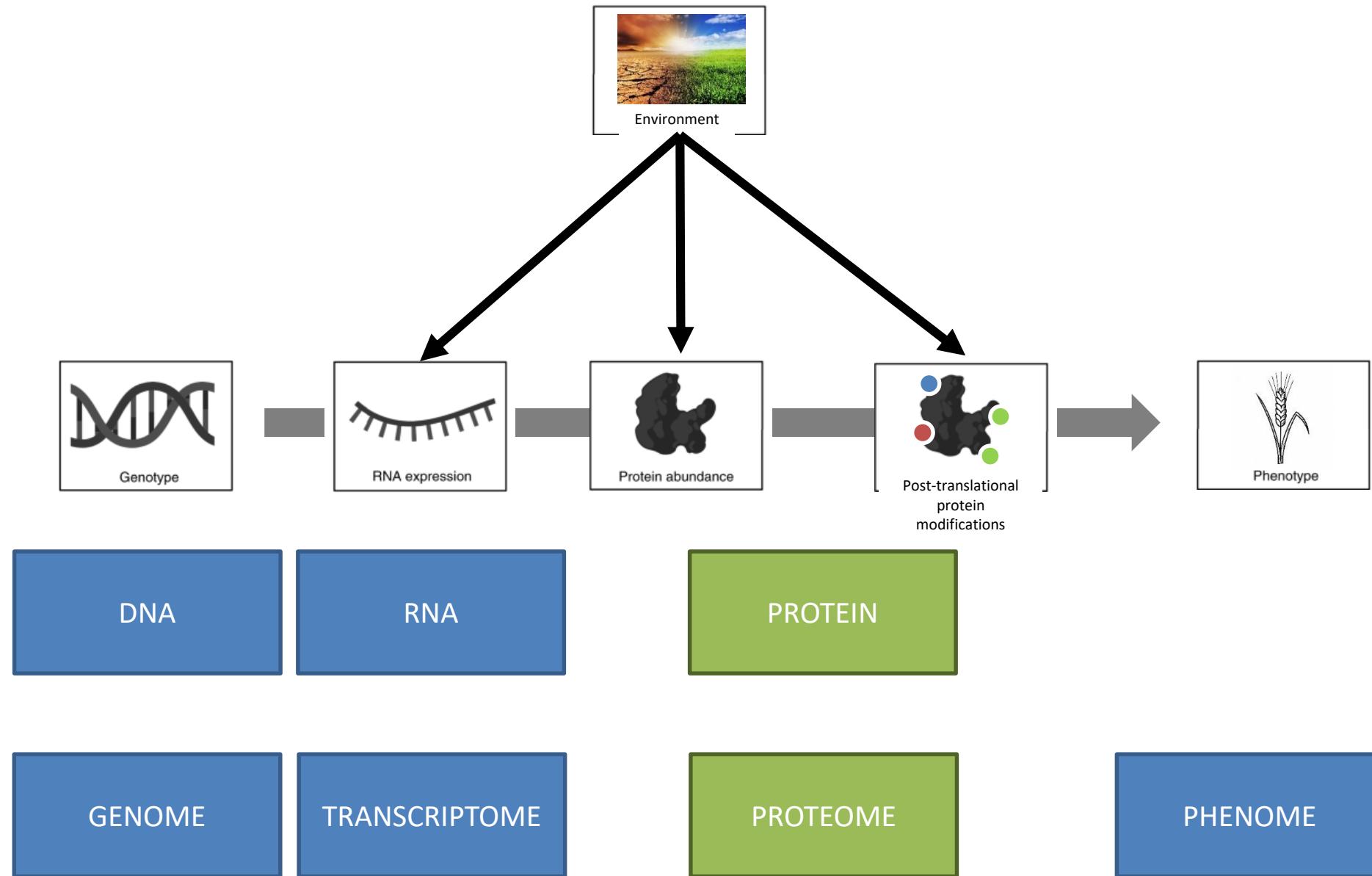
High temperature impacts wheat growth and development



High temperature impacts wheat growth and development



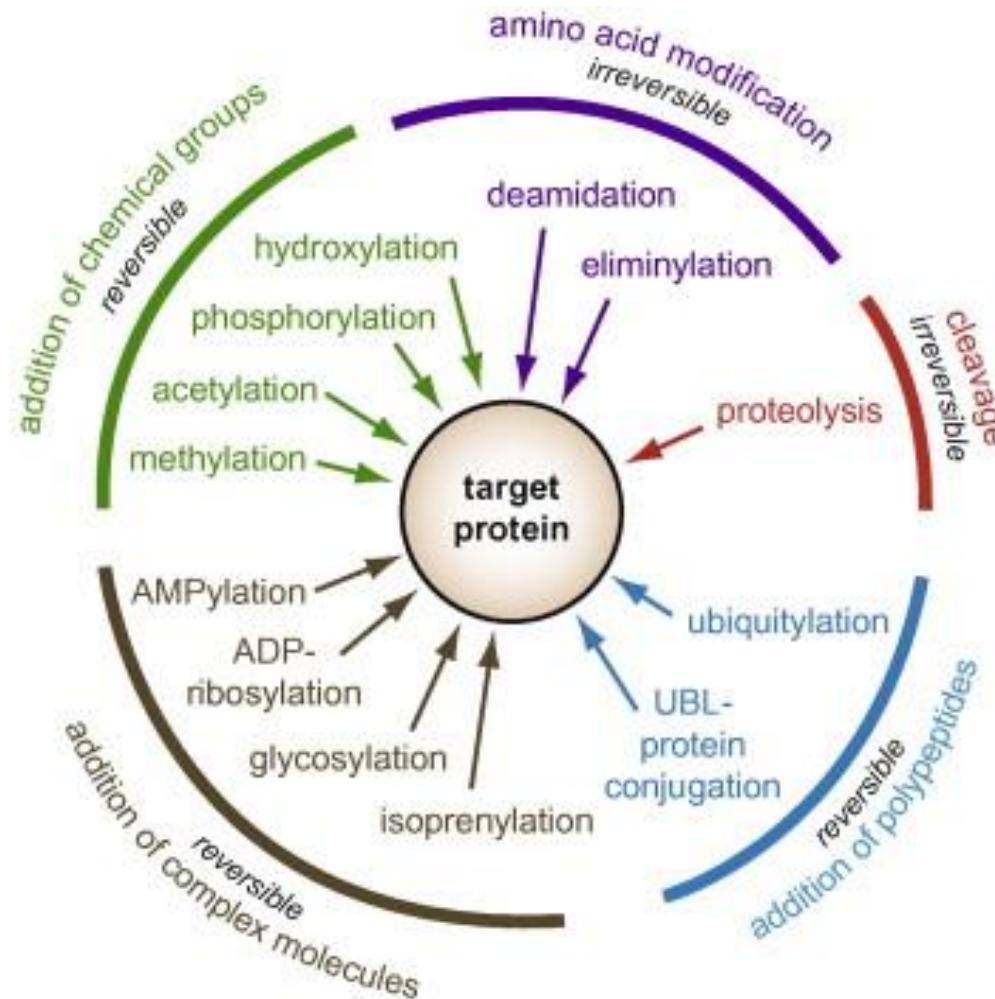
Focus on signalling in plants



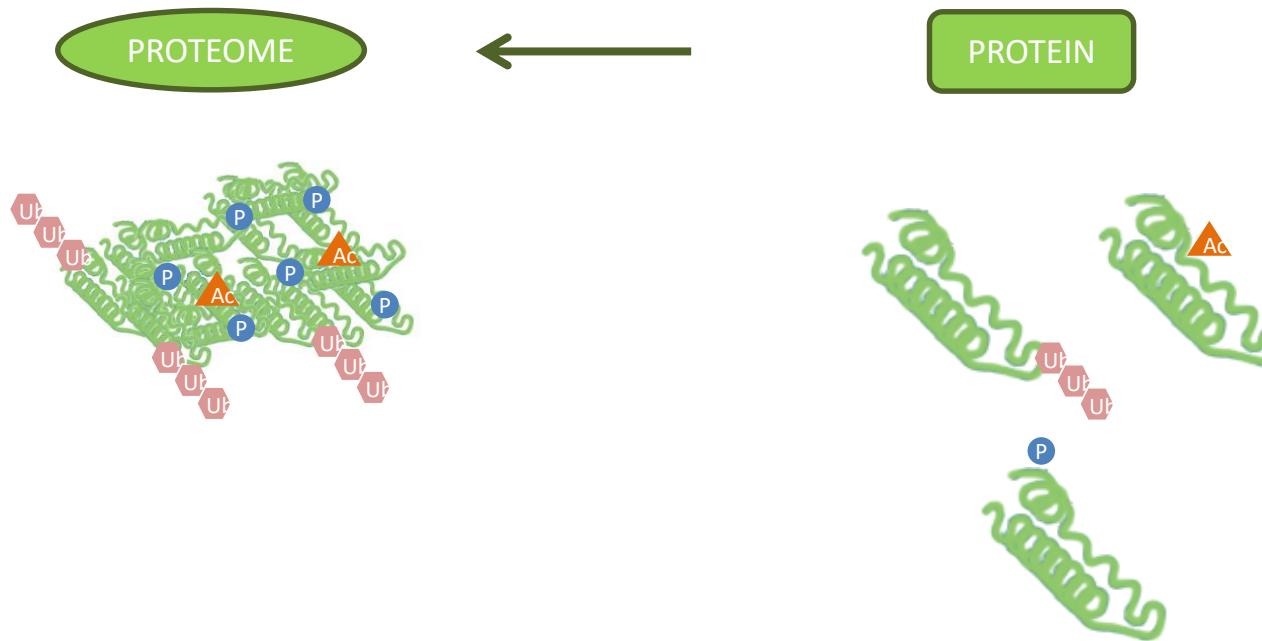
Protein pool expands through post-translational modifications



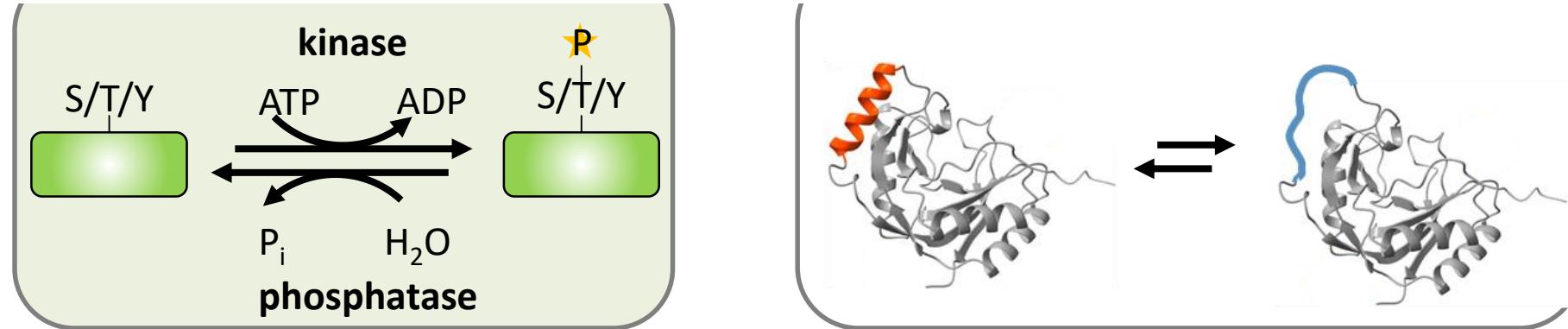
Protein pool expands through post-translational modifications



Protein pool expands through post-translational modifications



Focus on signalling in plants



POST TRANSLATIONAL PROTEIN
MODIFICATIONS
(MAINLY PHOSPHORYLATION)
PROTEIN STRUCTURE

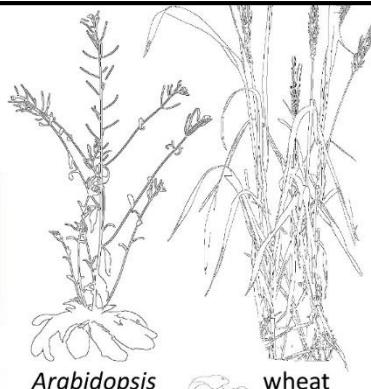
HIGH TEMPERATURE STRESS



DROUGHT STRESS



LOW TEMPERATURE STRESS



Arabidopsis

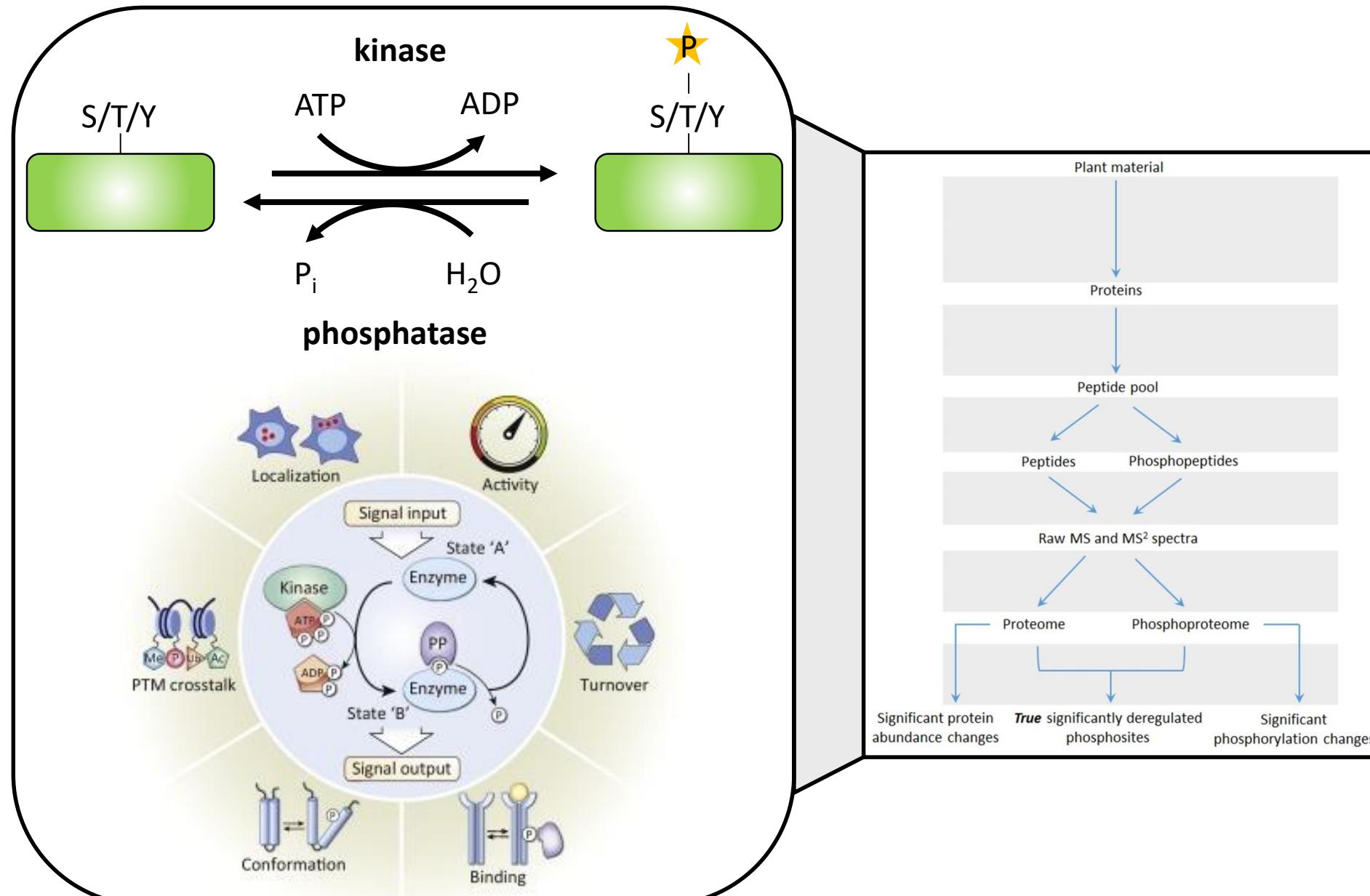
wheat

soybean

SHOOT GROWTH / FERTILITY



Phosphorylation is involved in everything



A resource for the community

The screenshot shows the Plant PTM Viewer 2.0 homepage. At the top, there's a banner with the VIB logo and a list of protein modifications: N terminus proteolysis, Methionine Oxidation, Lys S-methylation, Lys Methylation, N-Glycosylation, N-Acetylation, Myristylation, S-Glutathiolation, N-Ubiquitination, Phosphorylation, Lys Malonylation, Lys Acetylation, Lys Succinylation, Lys 2-Hydroxyisobutyrylation, Nitrosylation, O-GlcNAc, Lys Ubiquitination, Reversible Cys Oxidation, and Carbonylation. Below the banner, the title "Plant PTM Viewer 2.0" is displayed with the subtitle "From site-seeing to protein function". A navigation bar includes links for Home, Search, Analysis, Browse, Submit Data, and About. On the left, there's a sidebar with a "Peer reviewed MS/MS studies" section and a "From site-se" link. The main content area features a large green leaf with various protein modifications (phosphorylation, acetylation, methylation) shown as chemical structures. To the right, a text block describes the tool as a centralized resource for plant post-translational modifications, provides analysis tools, and accepts peer-reviewed data. It also links to features and statistics, and offers downloadable tutorials.

Plant PTM Viewer

Plant PTM Viewer is a centralized resource for plant post-translational modifications (PTMs) intuitive for wet- and dry-lab scientists.

Plant PTM Viewer provides innovative tools to analyze the potential role of PTMs for specific proteins or in a broader systems biology context.

Plant PTM Viewer is an open repository and accepts newly peer-reviewed plant PTM data - for more information click [here](#)

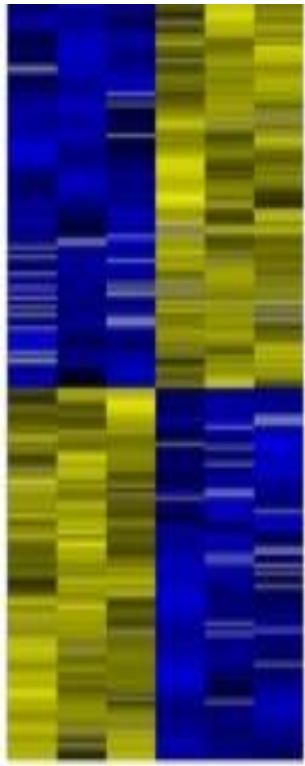
Below we provide a brief outline of Plant PTM Viewer [features](#) and current PTM data [statistics](#).

Downloadable tutorials can be found [here](#).

<https://www.psb.ugent.be/webtools/ptm-viewer/>

Discovery (and validation) in wheat / mode-of-action in *Arabidopsis*

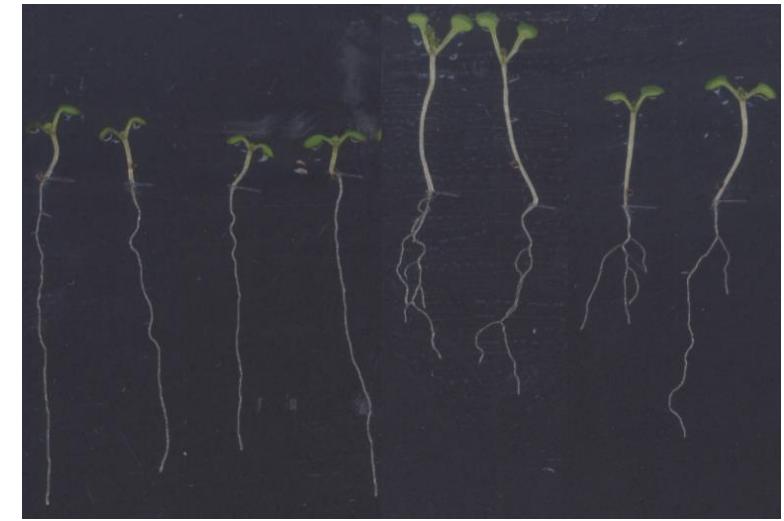
Relevant candidates in wheat



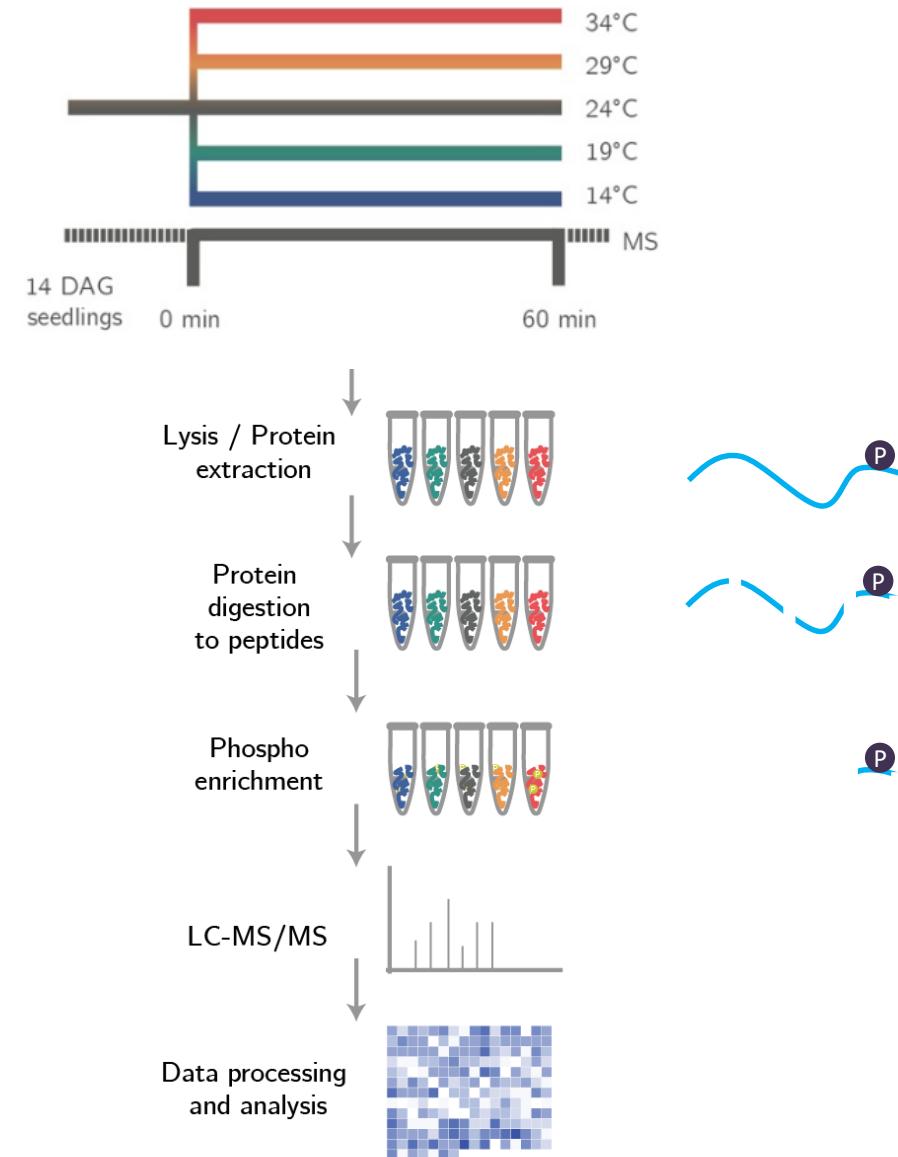
Validation in wheat



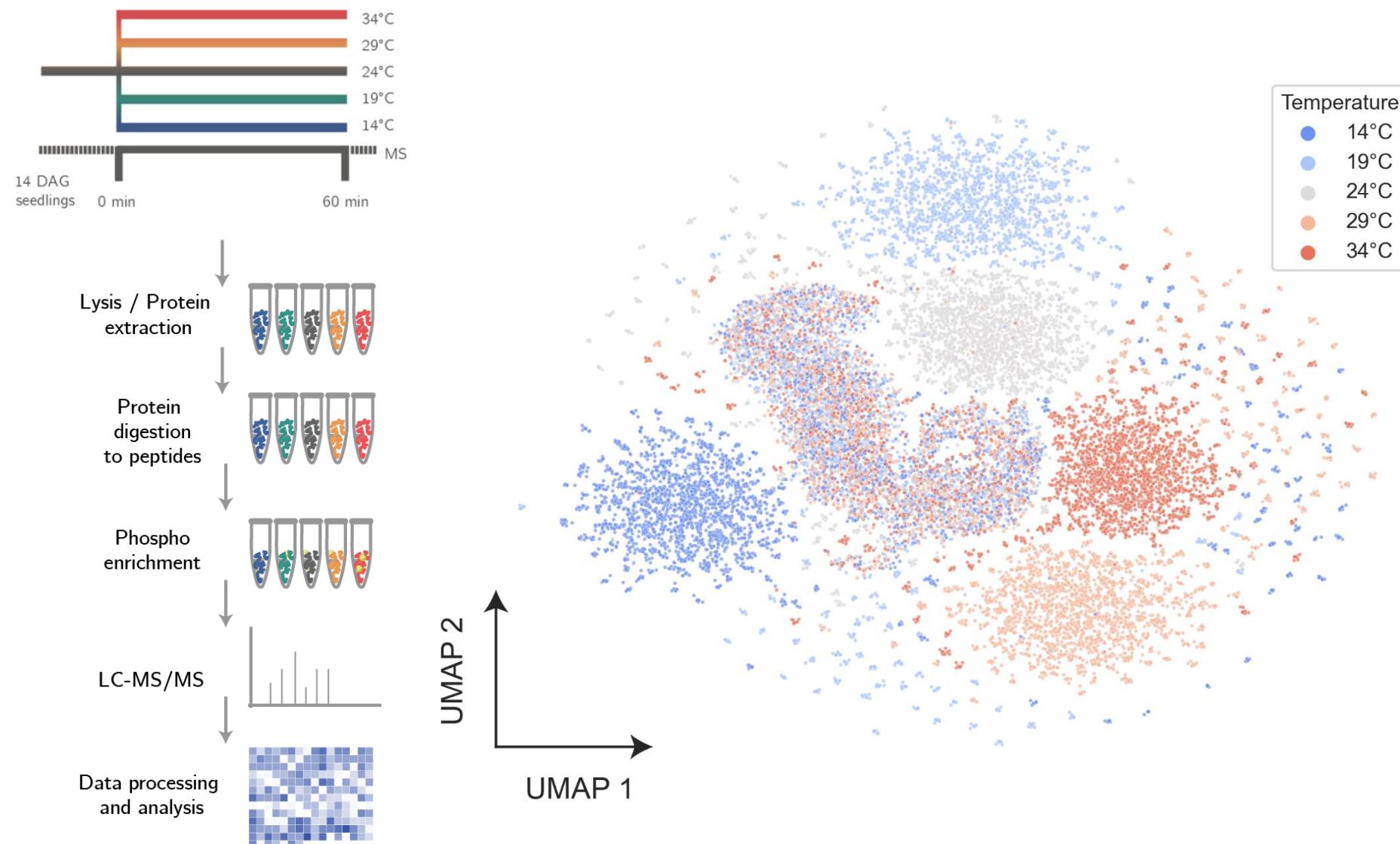
Mode-of-action in *Arabidopsis*



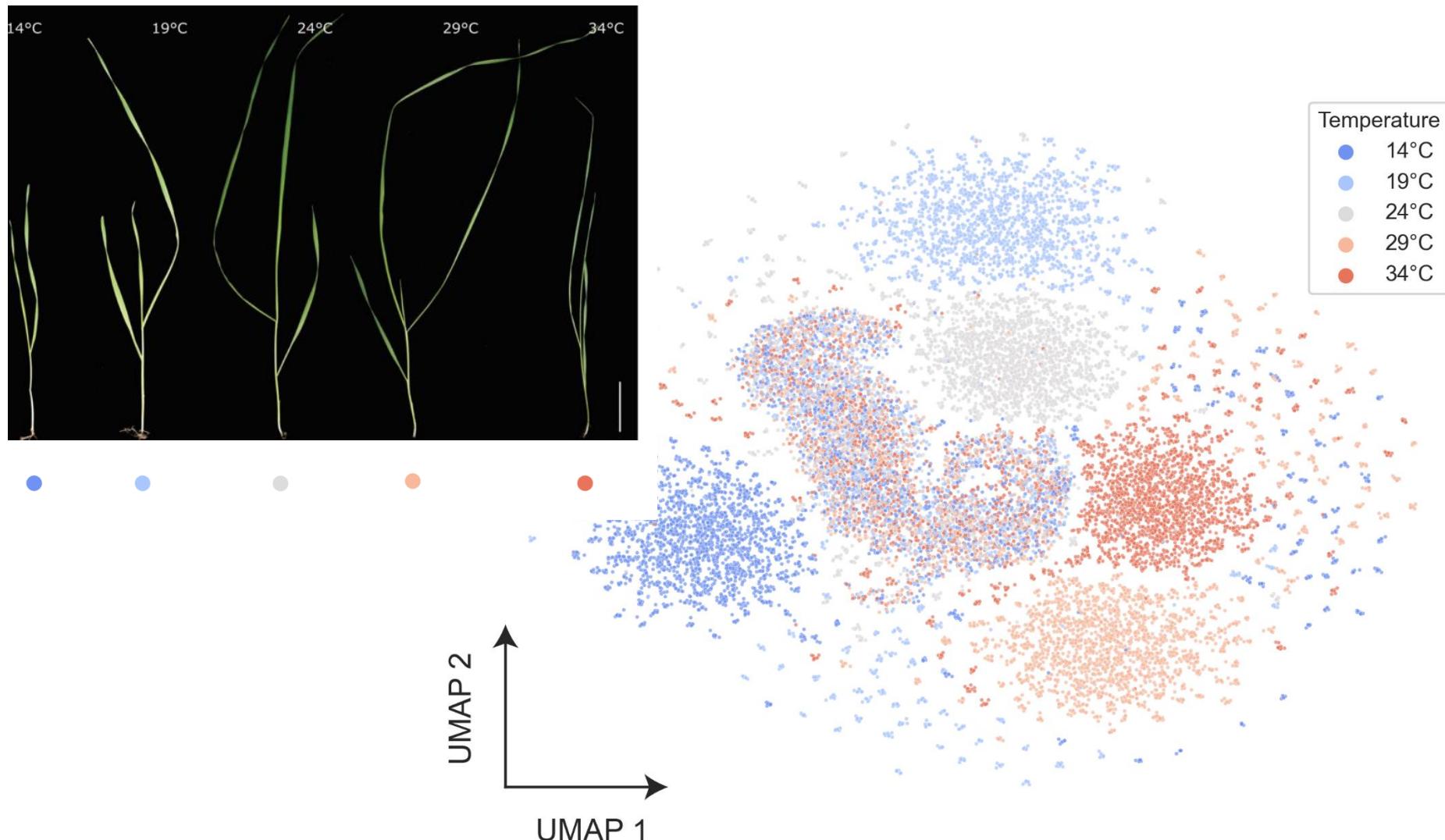
Temperature gradient phosphoprofiling in wheat



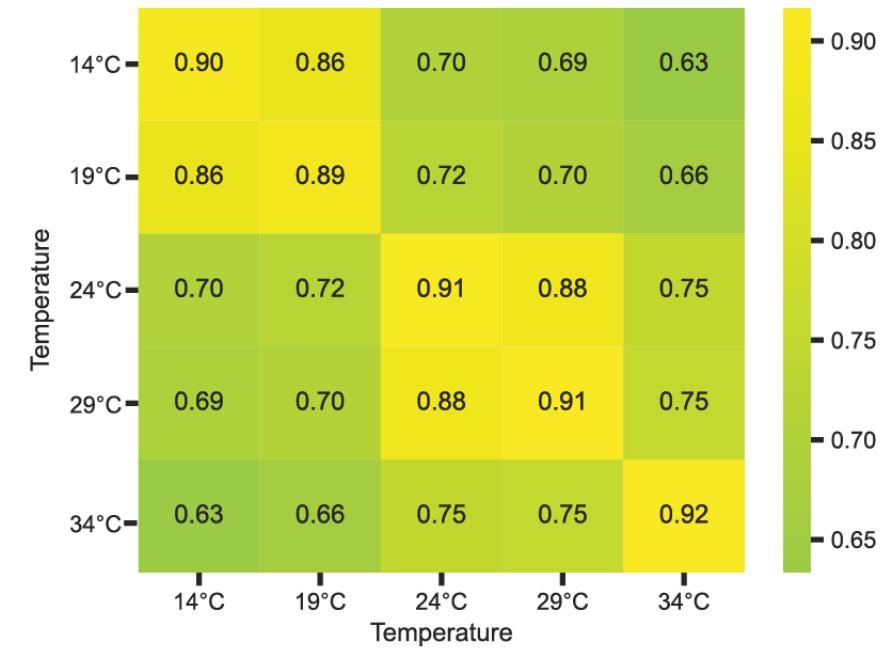
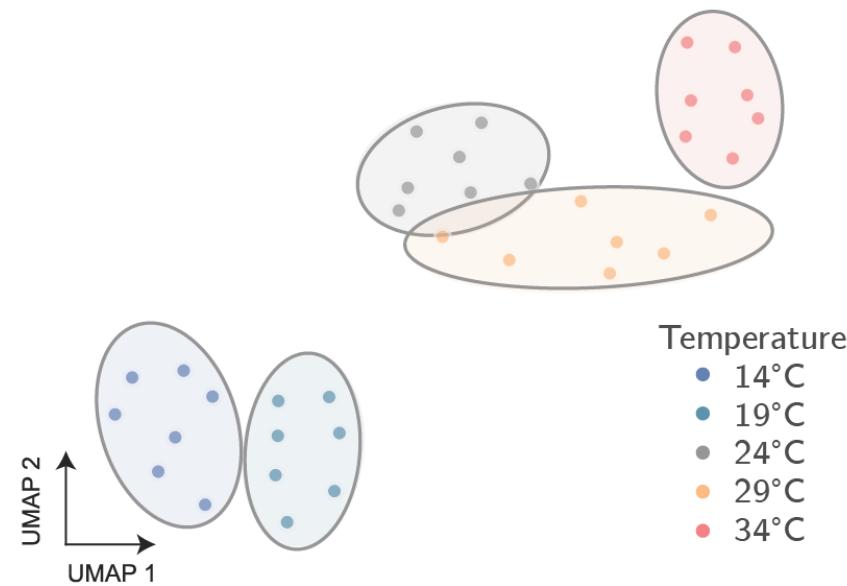
Temperature gradient phosphoprofiling in wheat



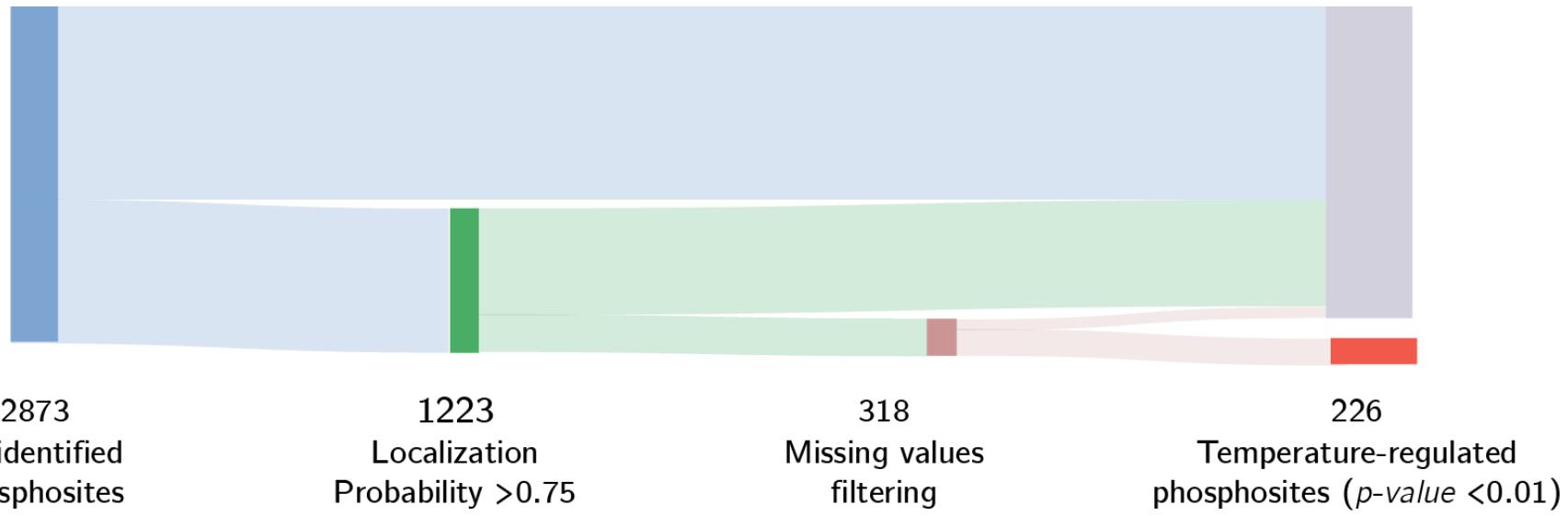
Temperature gradient phosphoprofiling in wheat



Temperature gradient phosphoprofiling in wheat



Temperature gradient phosphoprofiling in wheat



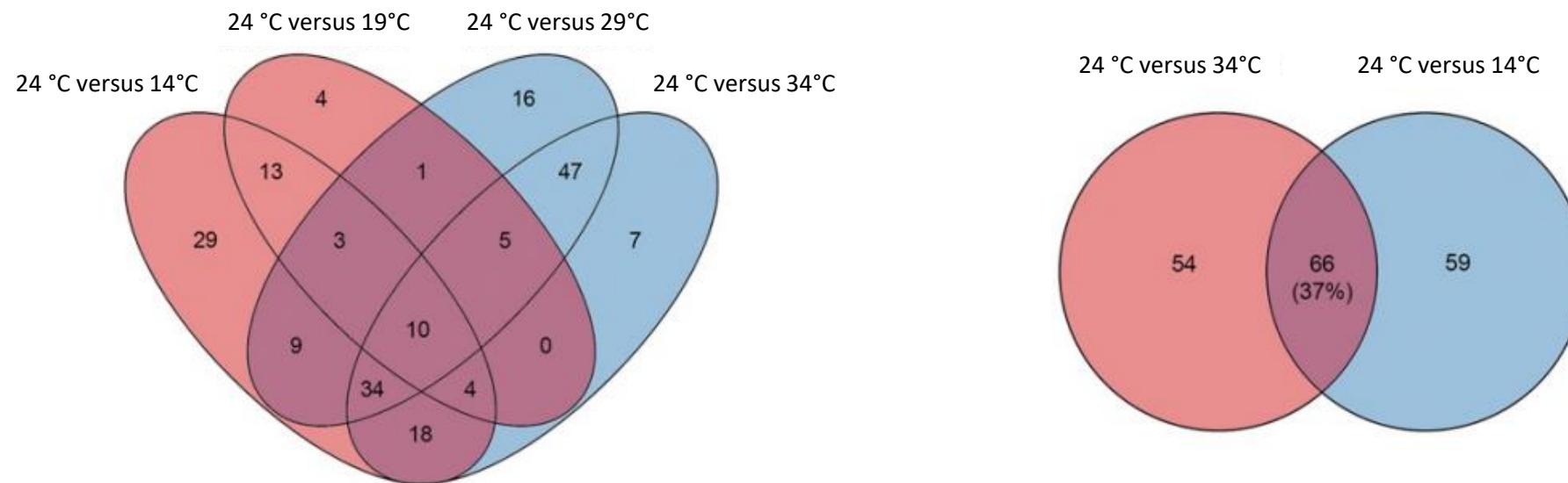
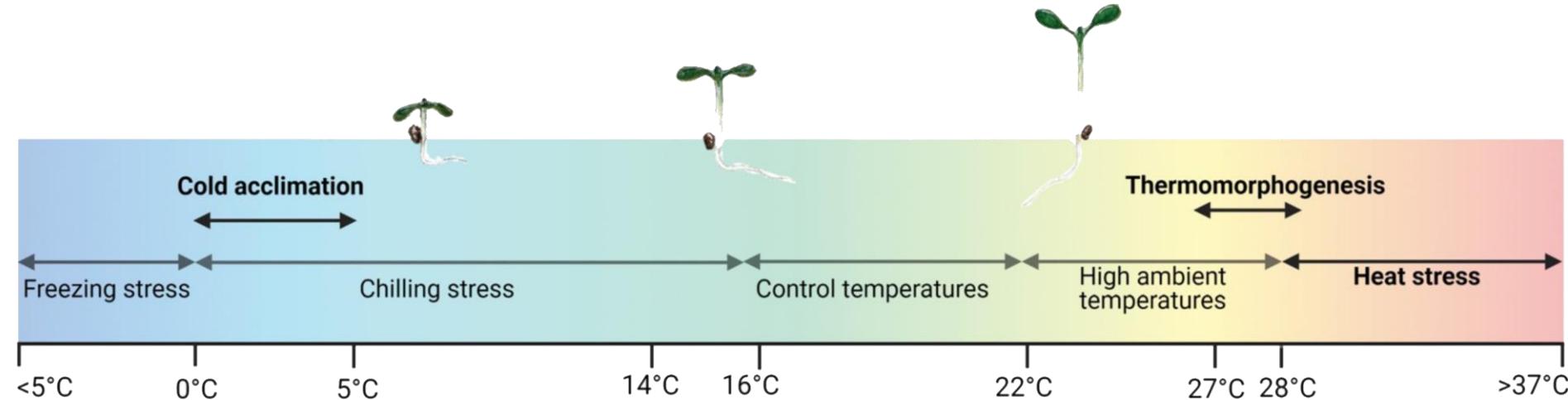
2873
All identified
phosphosites

1223
Localization
Probability >0.75

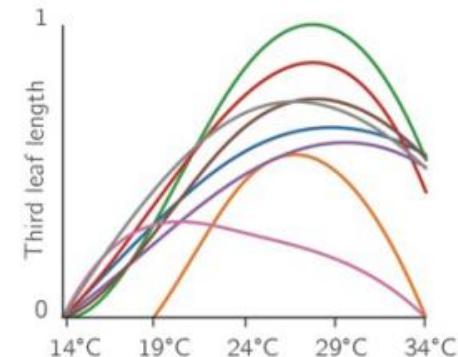
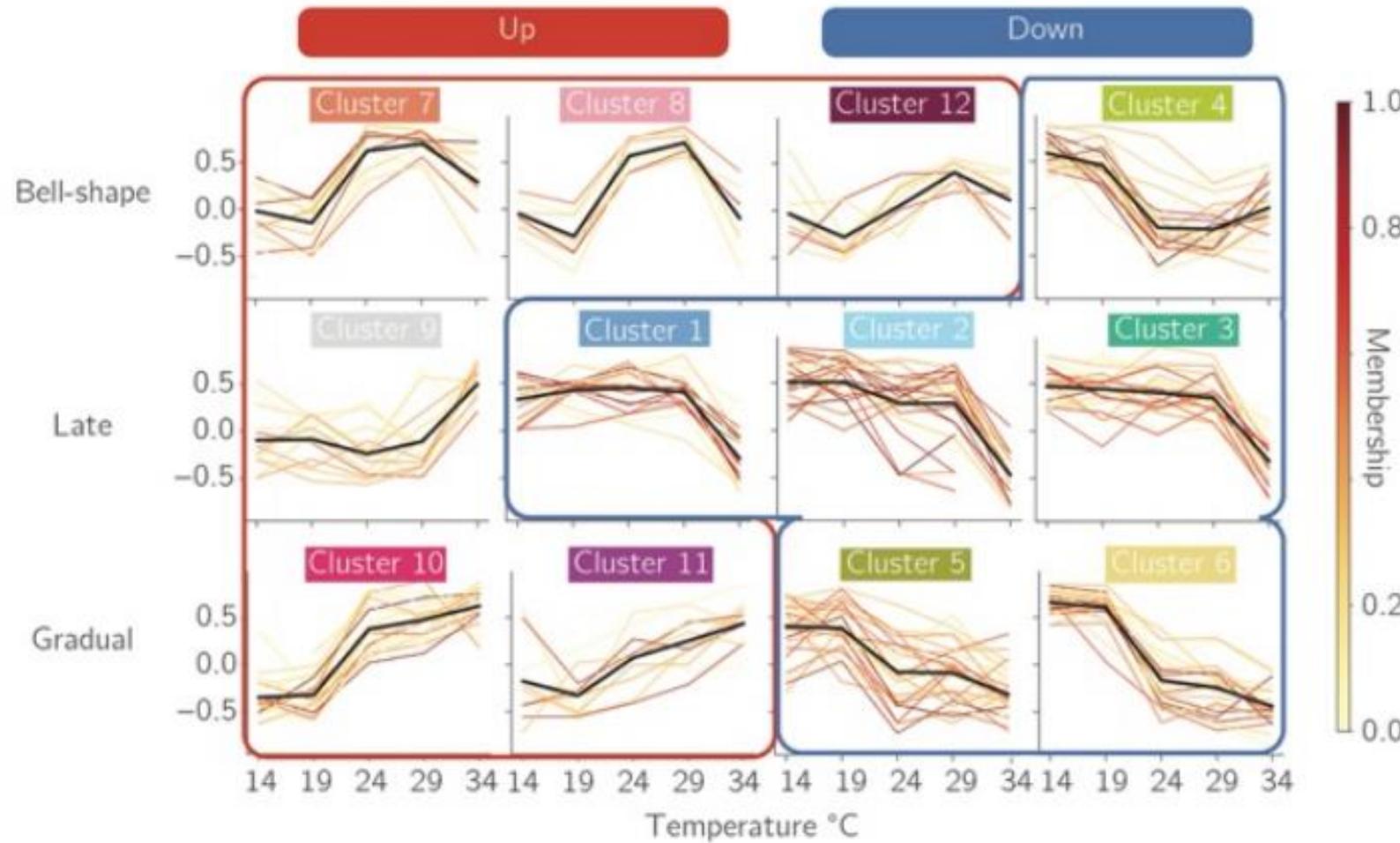
318
Missing values
filtering

226
Temperature-regulated
phosphosites ($p\text{-value} < 0.01$)

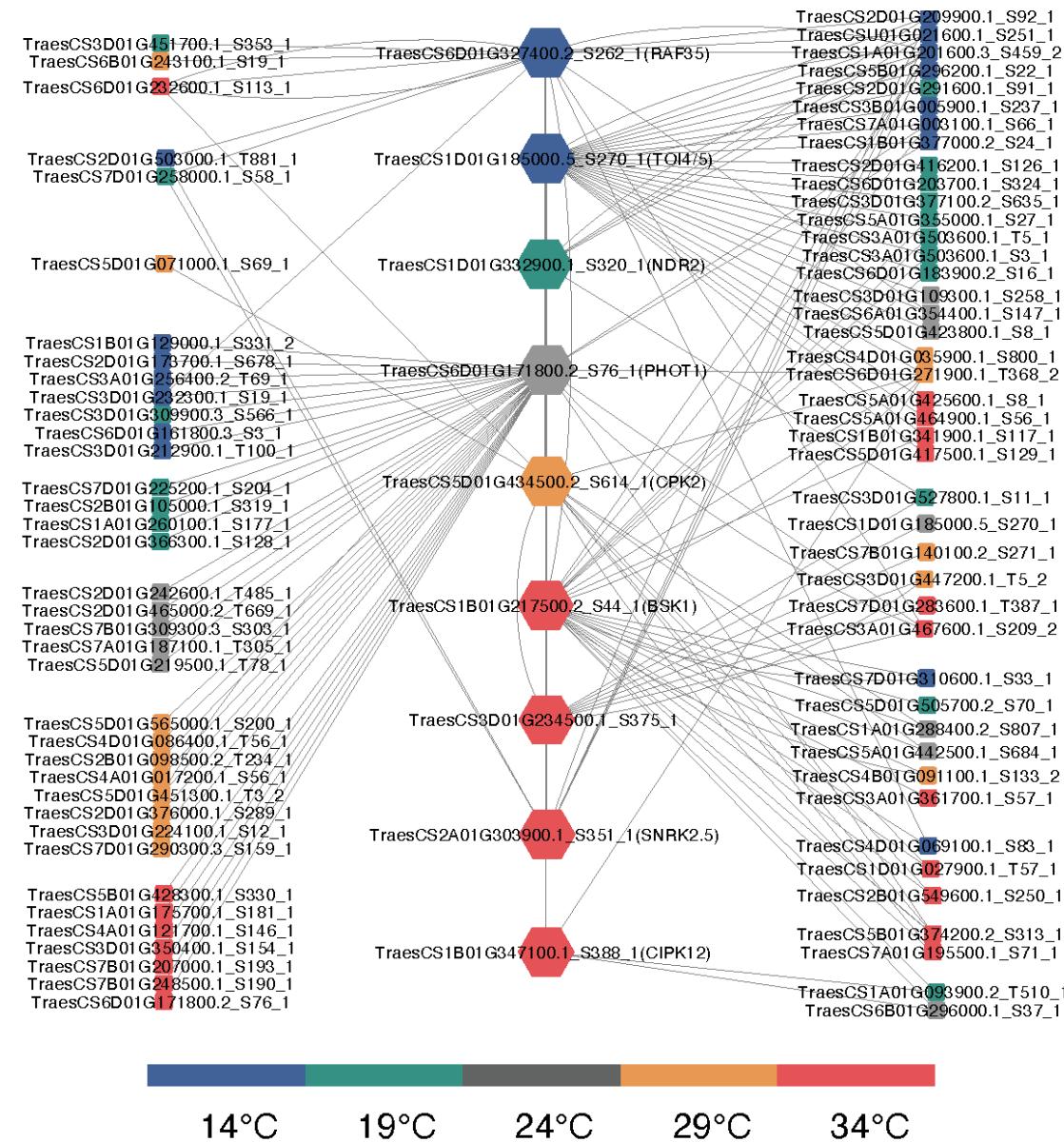
Temperature gradient phosphoprofiling in wheat



Temperature gradient phosphoprofiling in wheat

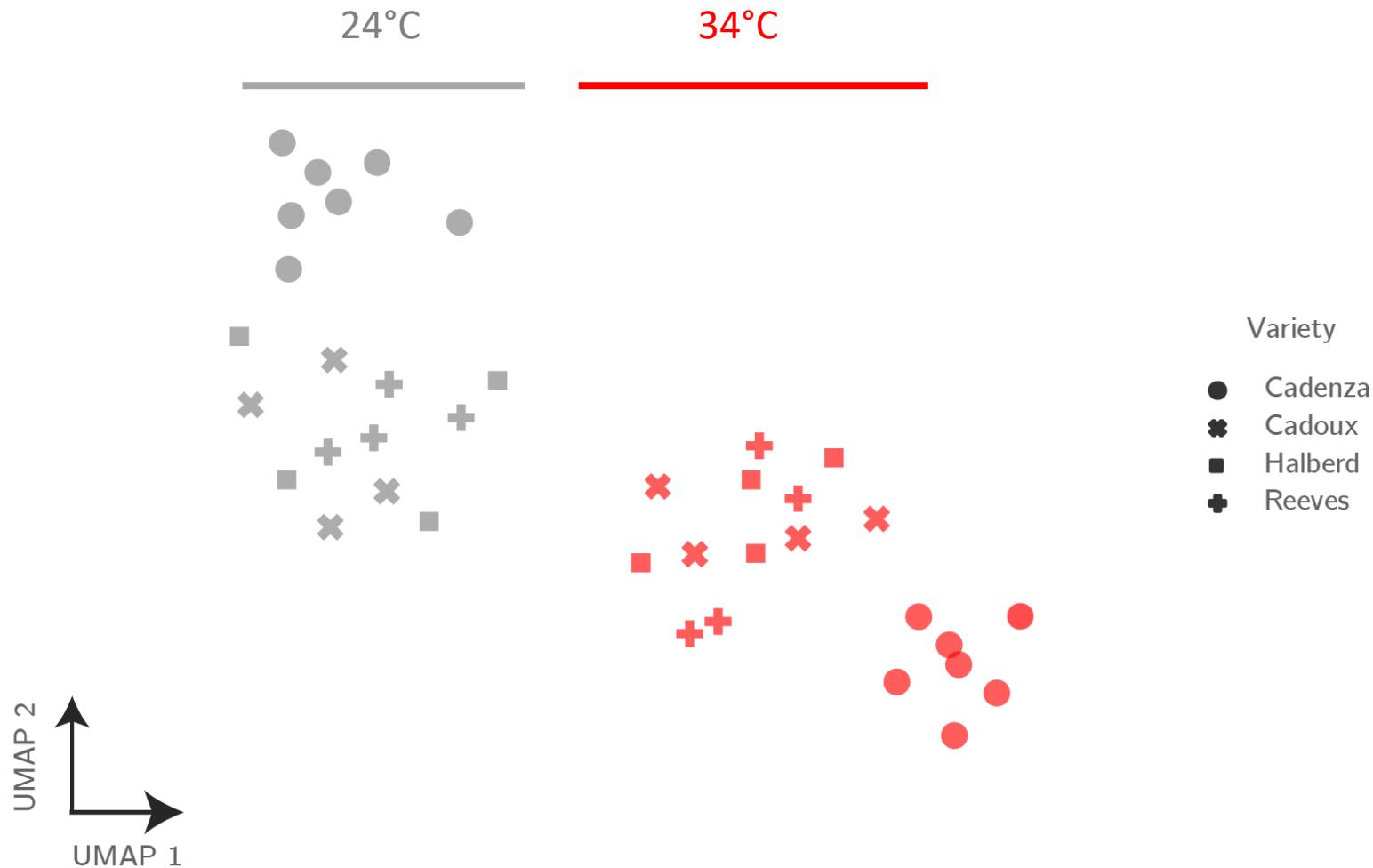


Signalling network from temperature gradient phosphoprofiling in wheat

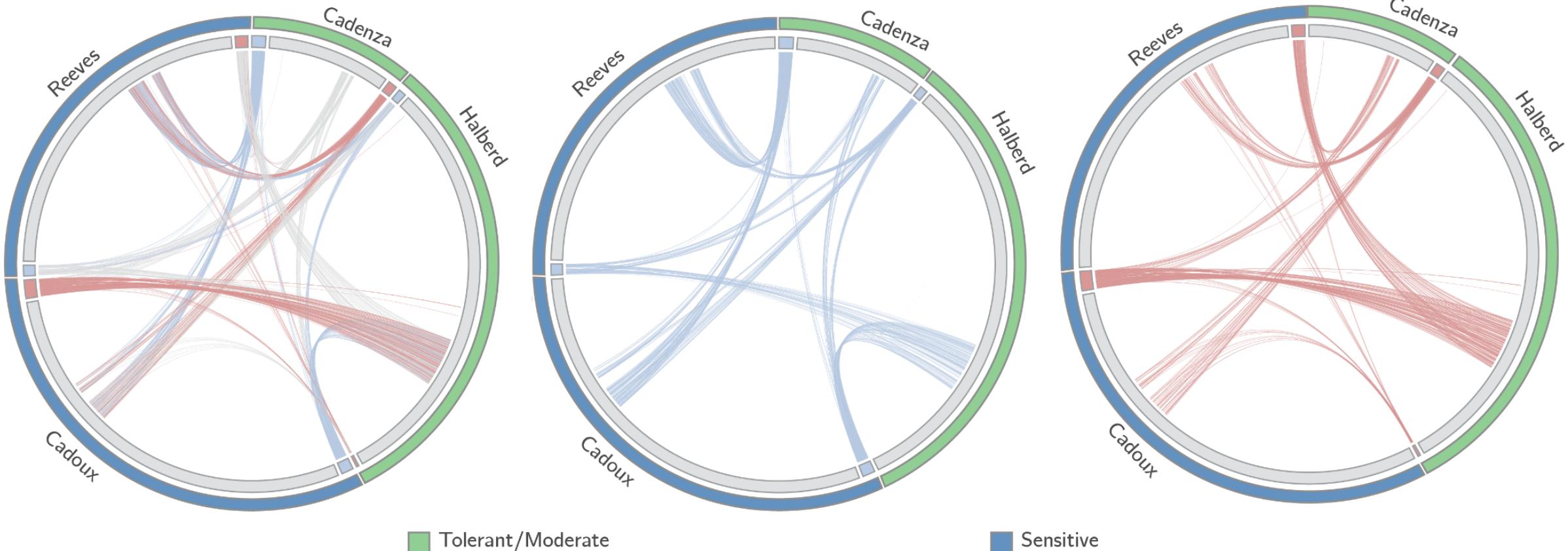


14°C 19°C 24°C 29°C 34°C

Temperature-regulated phosphoproteome: variety-dependent molecular responses



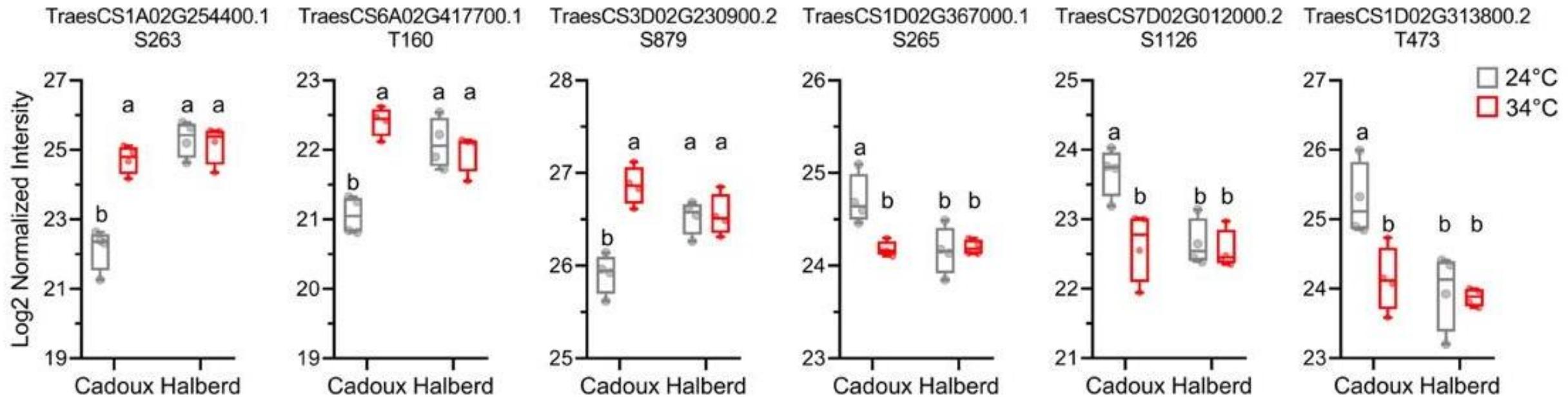
Temperature-regulated phosphoproteome: variety-dependent molecular responses



■ Tolerant/Moderate

■ Sensitive

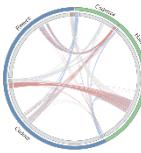
Temperature-regulated phosphoproteome: variety-dependent molecular responses



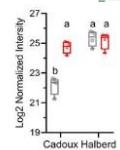
Predictive for breeding?



Capture environment-regulated phosphoproteome
in
sensitive and tolerant varieties
in
a growth chamber



Comparative analysis of phosphoproteome data



Identification
of
putative phosphoprotein-based biomarkers



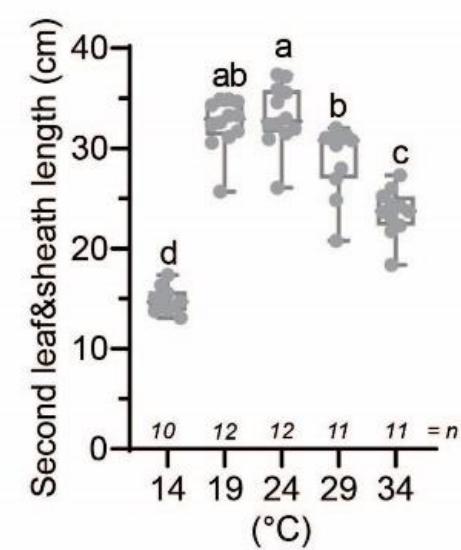
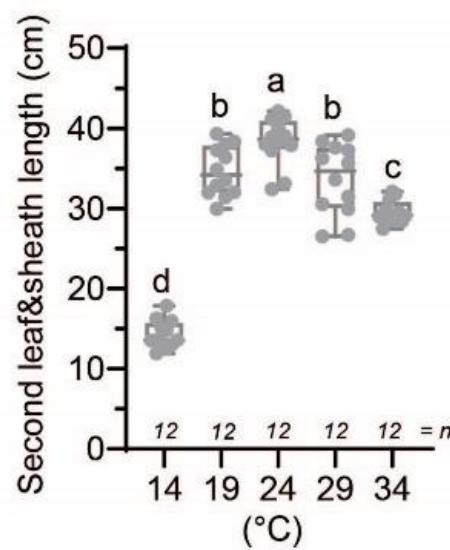
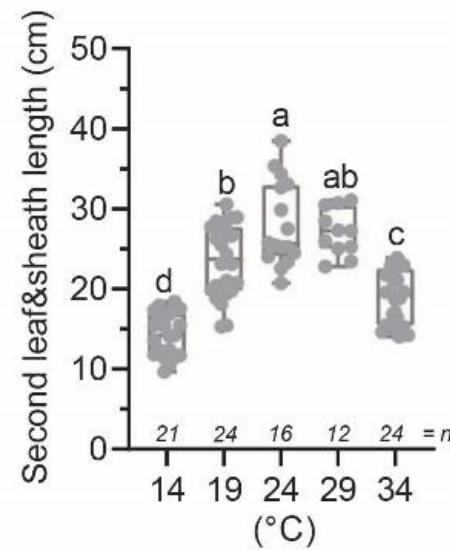
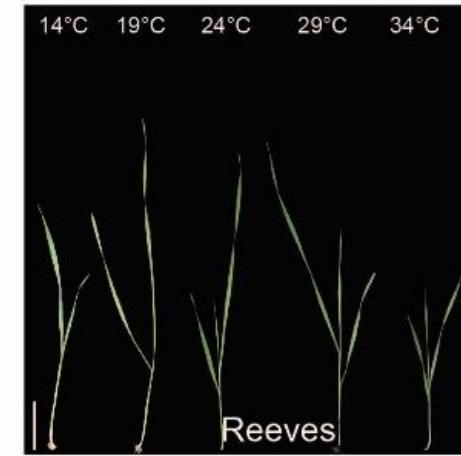
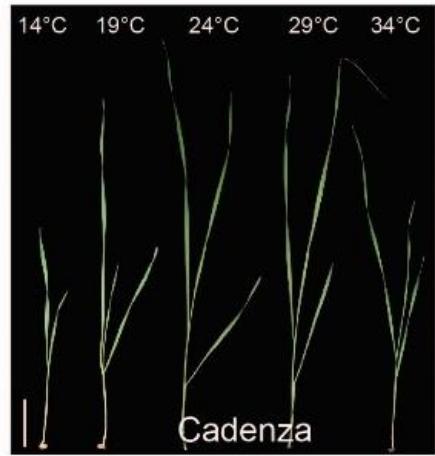
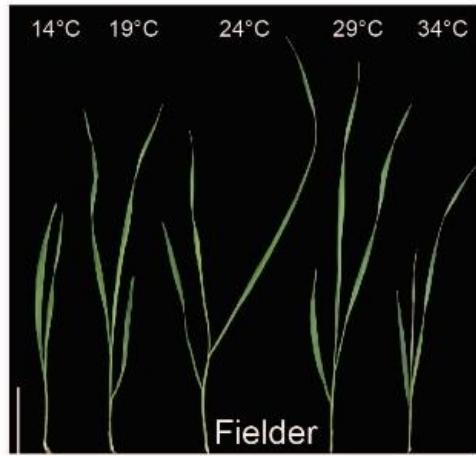
Breeding population

Physiological and genomic data
(sensitive versus tolerant variety)

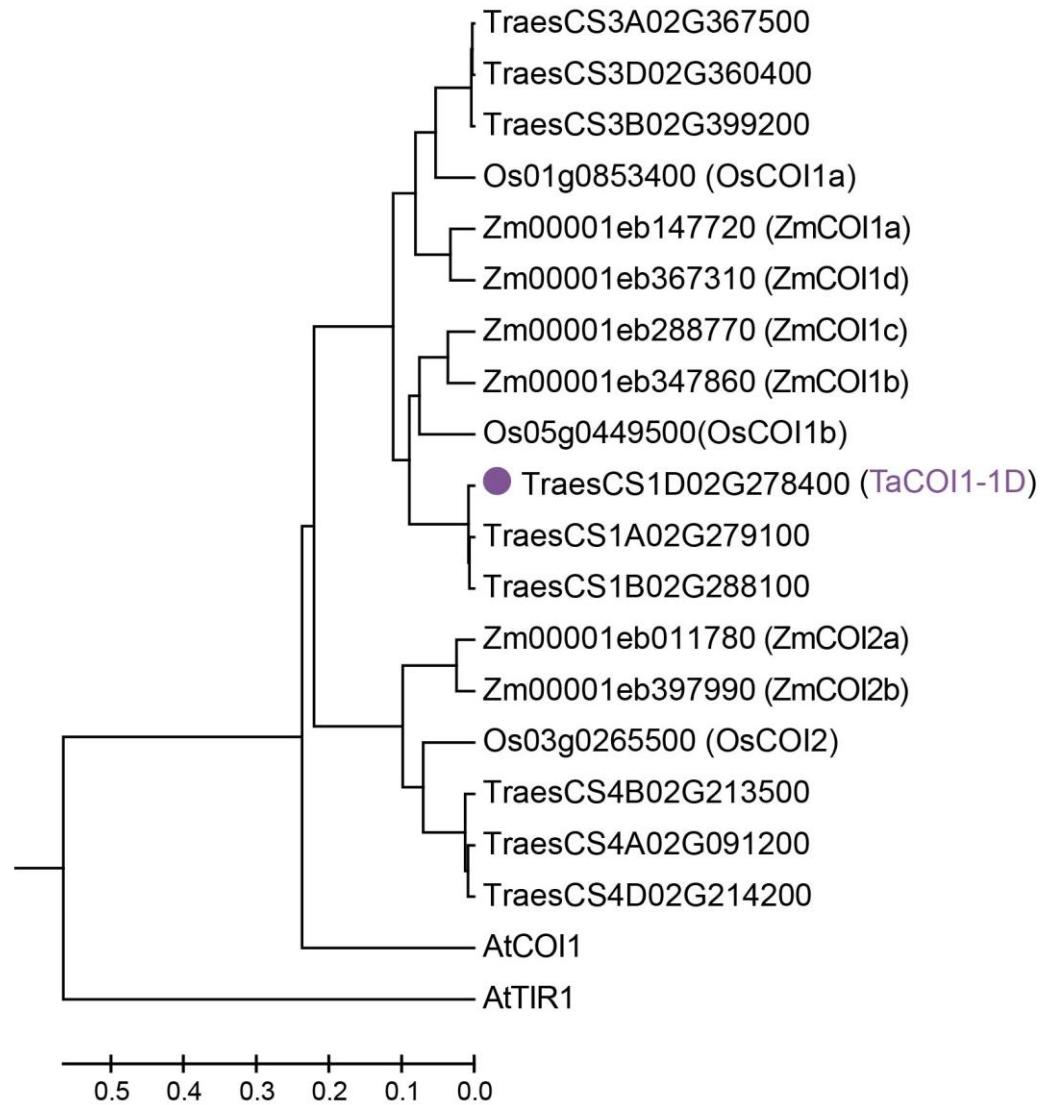
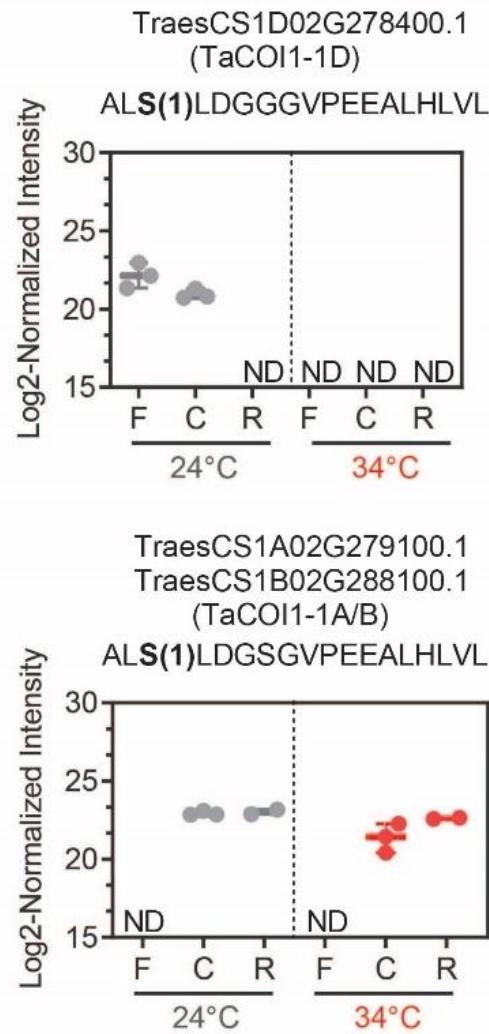
(Samples of) selected lines

Phosphoproteomics
(or phosphoantibody-based approach)

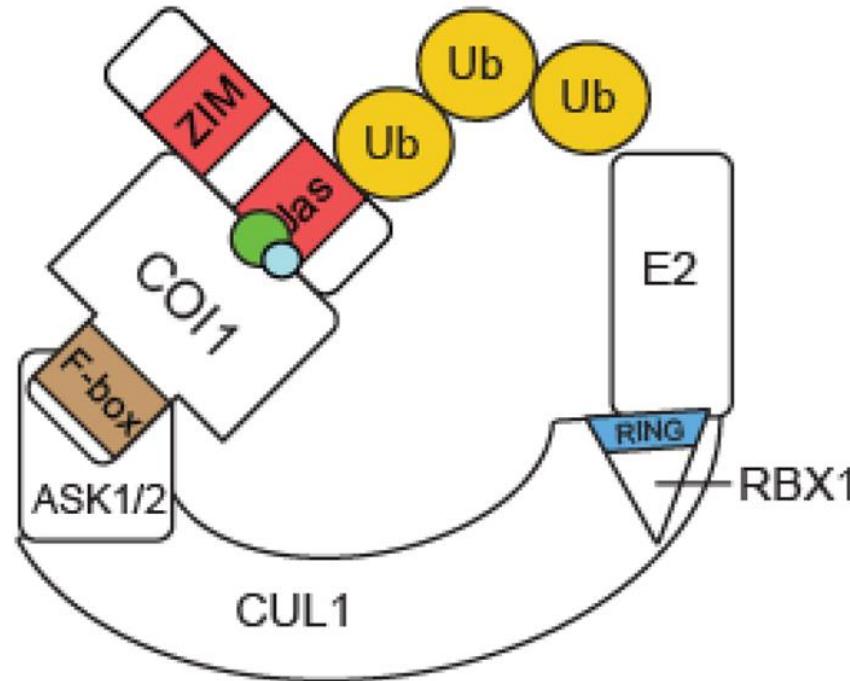
High temperature impacts wheat growth



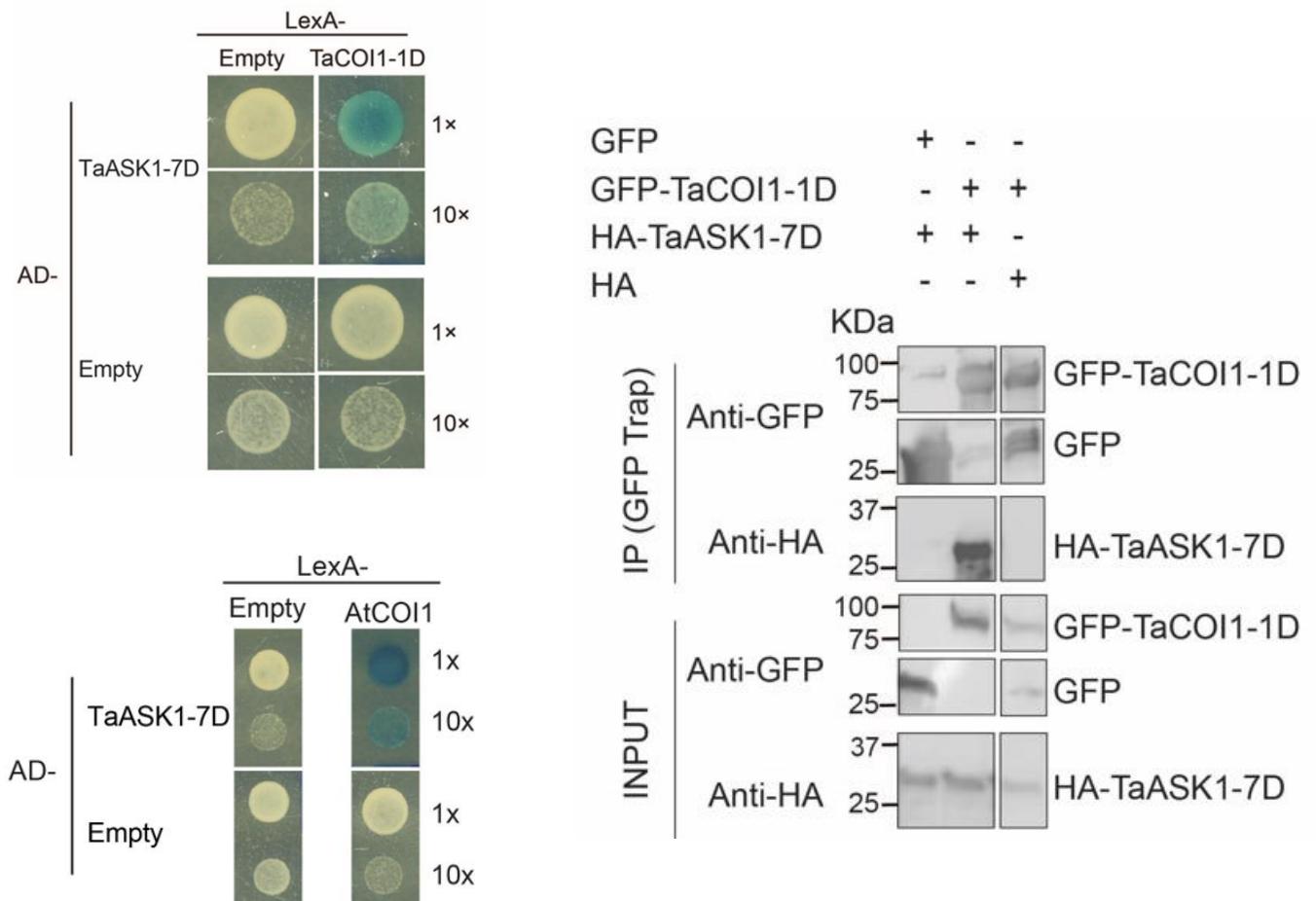
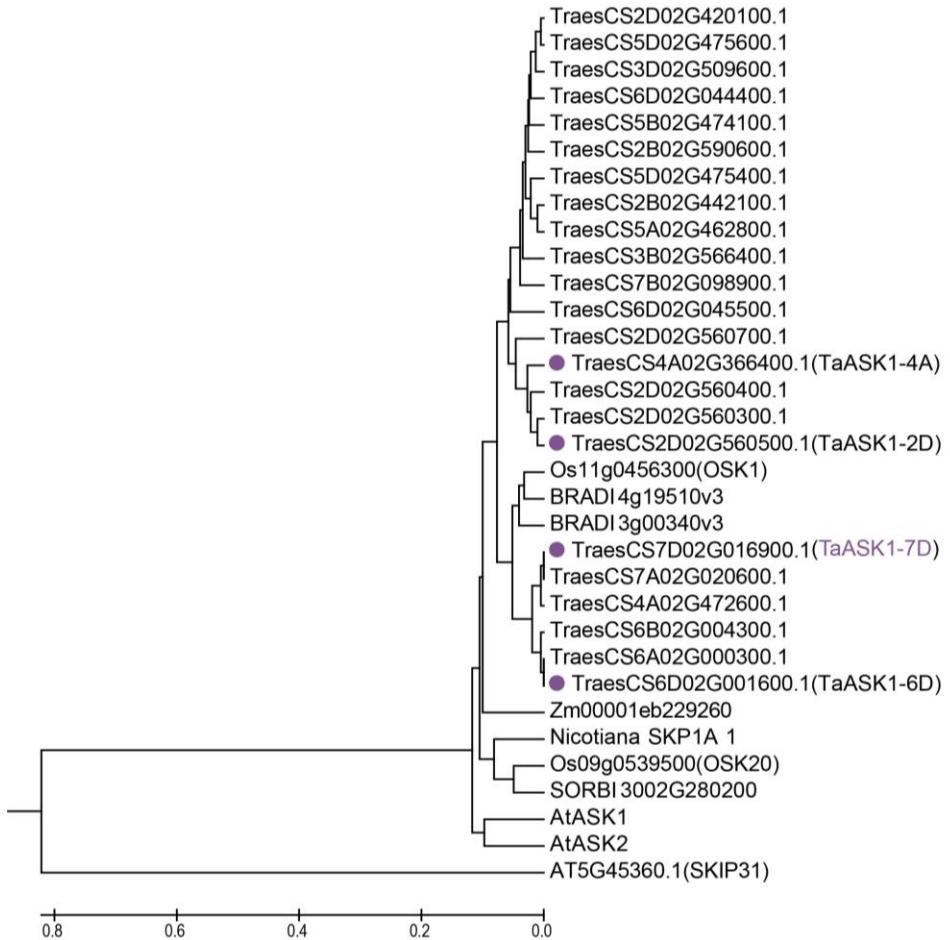
TaCOI11-D – A high temperature-regulated wheat (phospho)protein



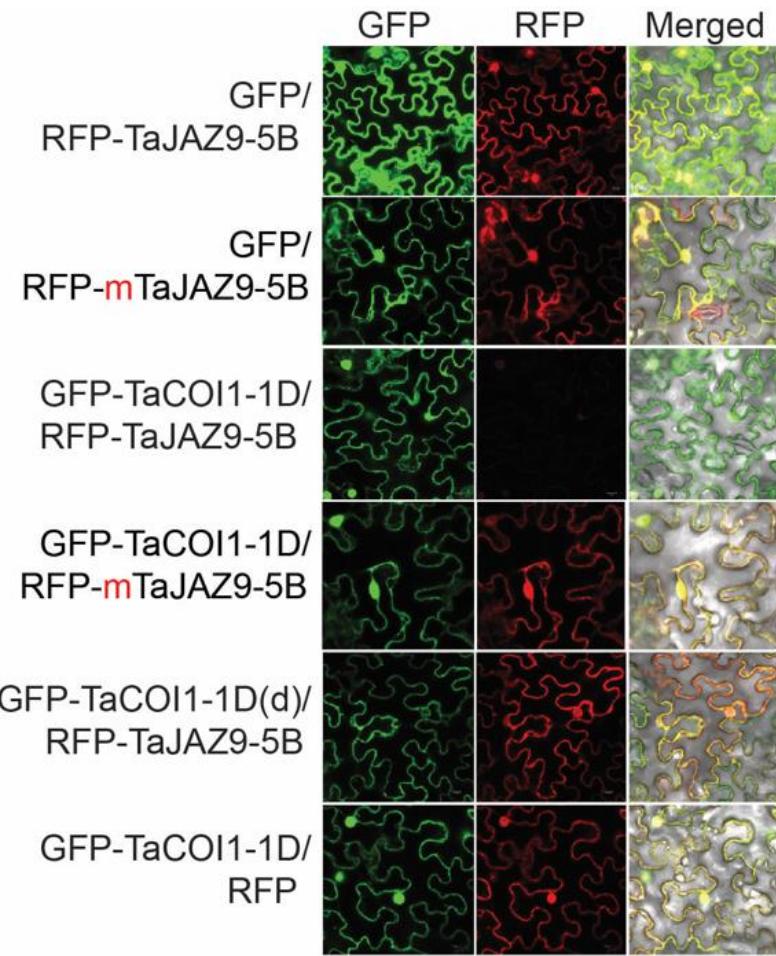
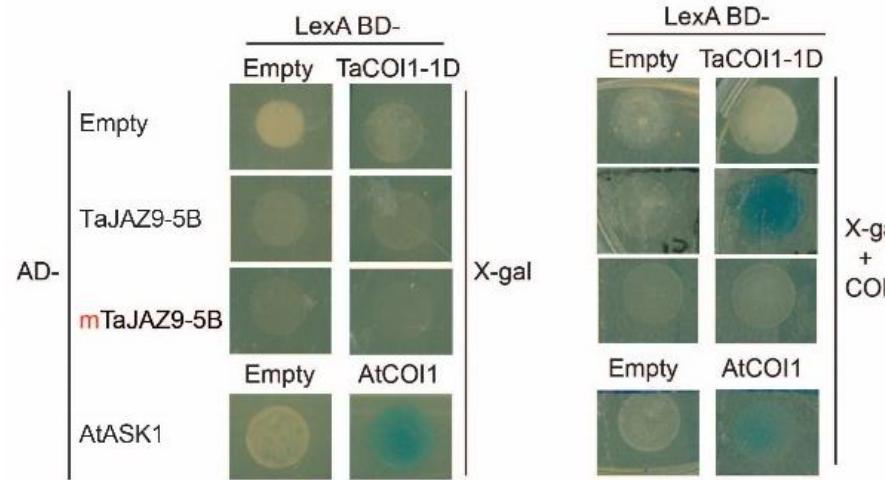
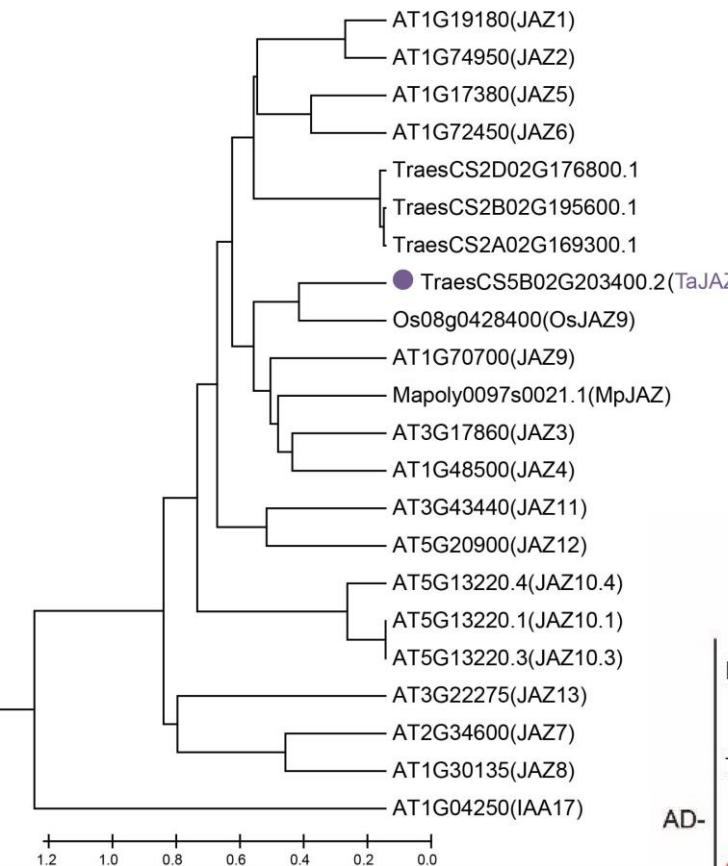
TaCOI11-D – A high temperature-regulated wheat JA receptor?



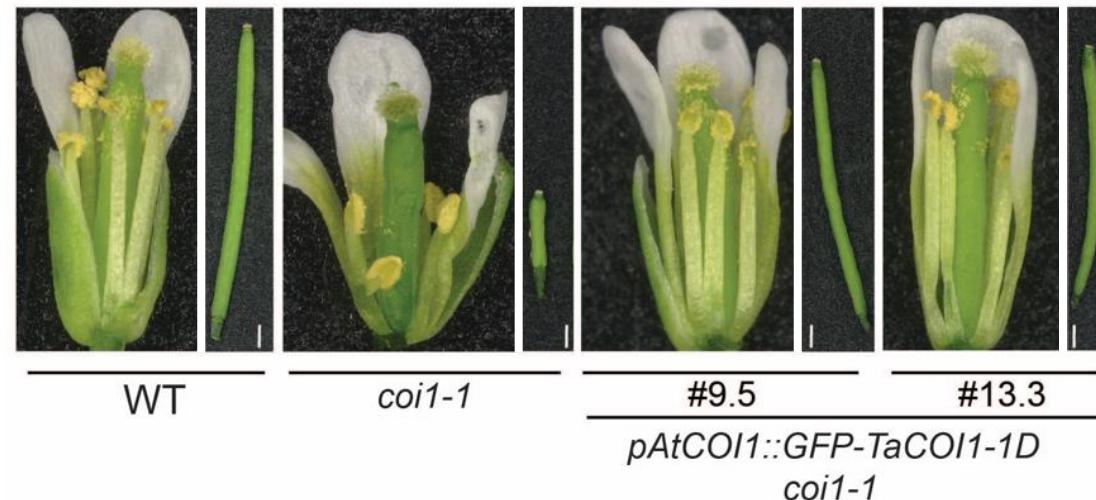
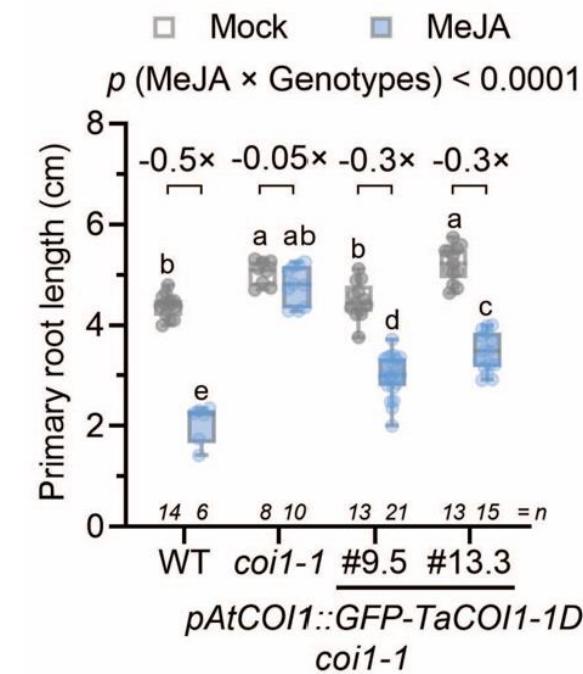
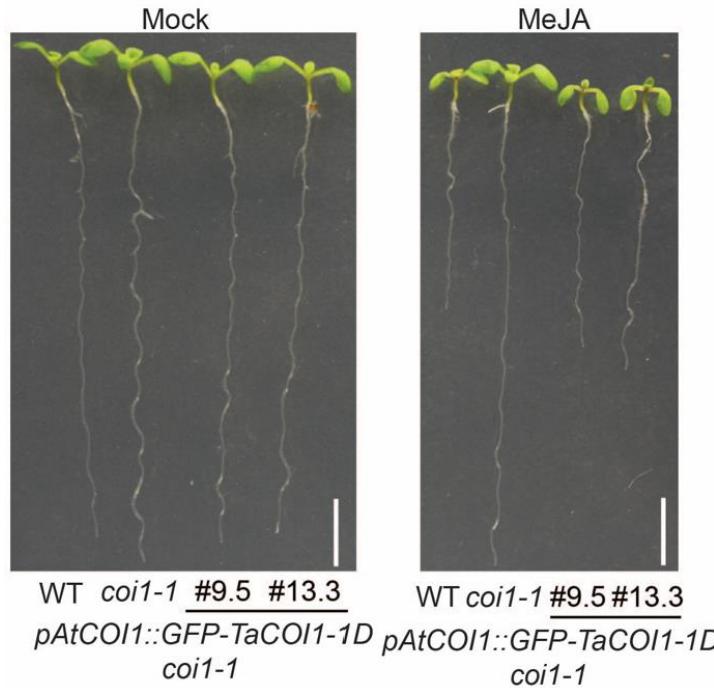
Wheat TaCOI1-1D interacts with TaASK orthologues



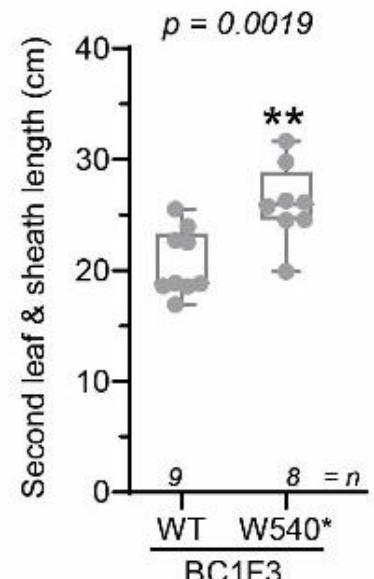
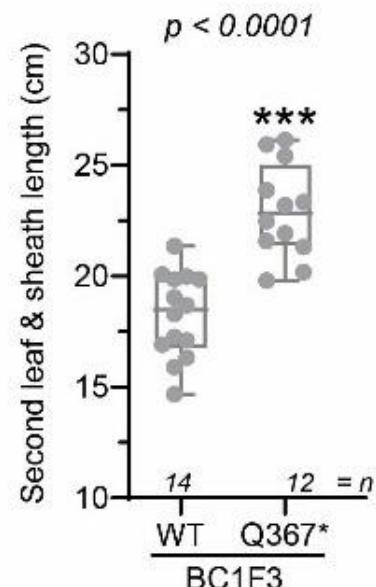
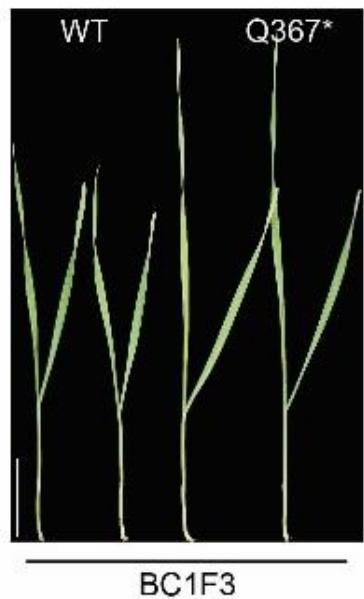
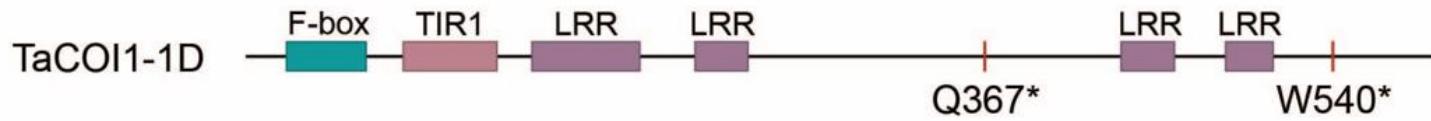
TaCOI1-1D interacts with TaJAZ9-5B in CORONATINE-dependent way



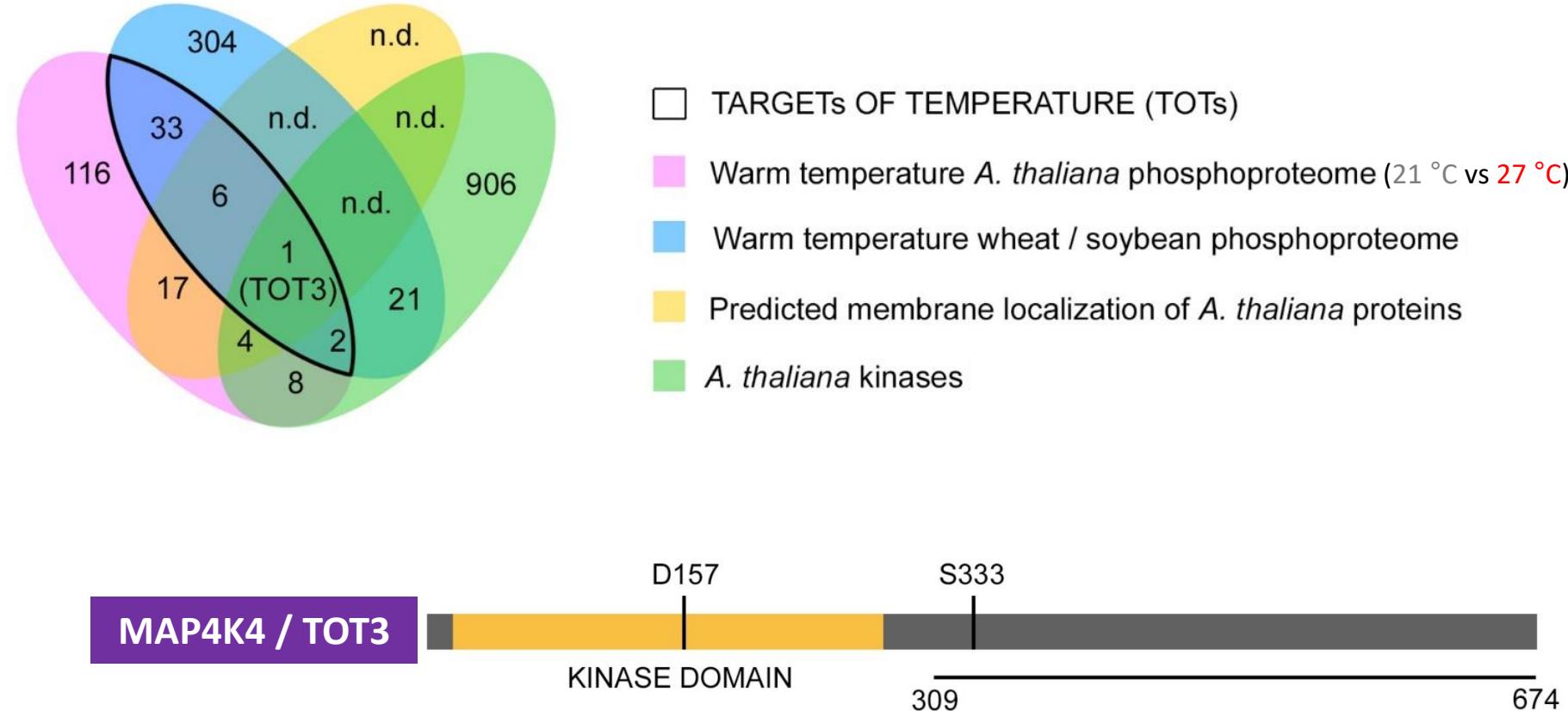
Wheat TaCOI1-1D can rescue Arabidopsis *coi1-1* mutant



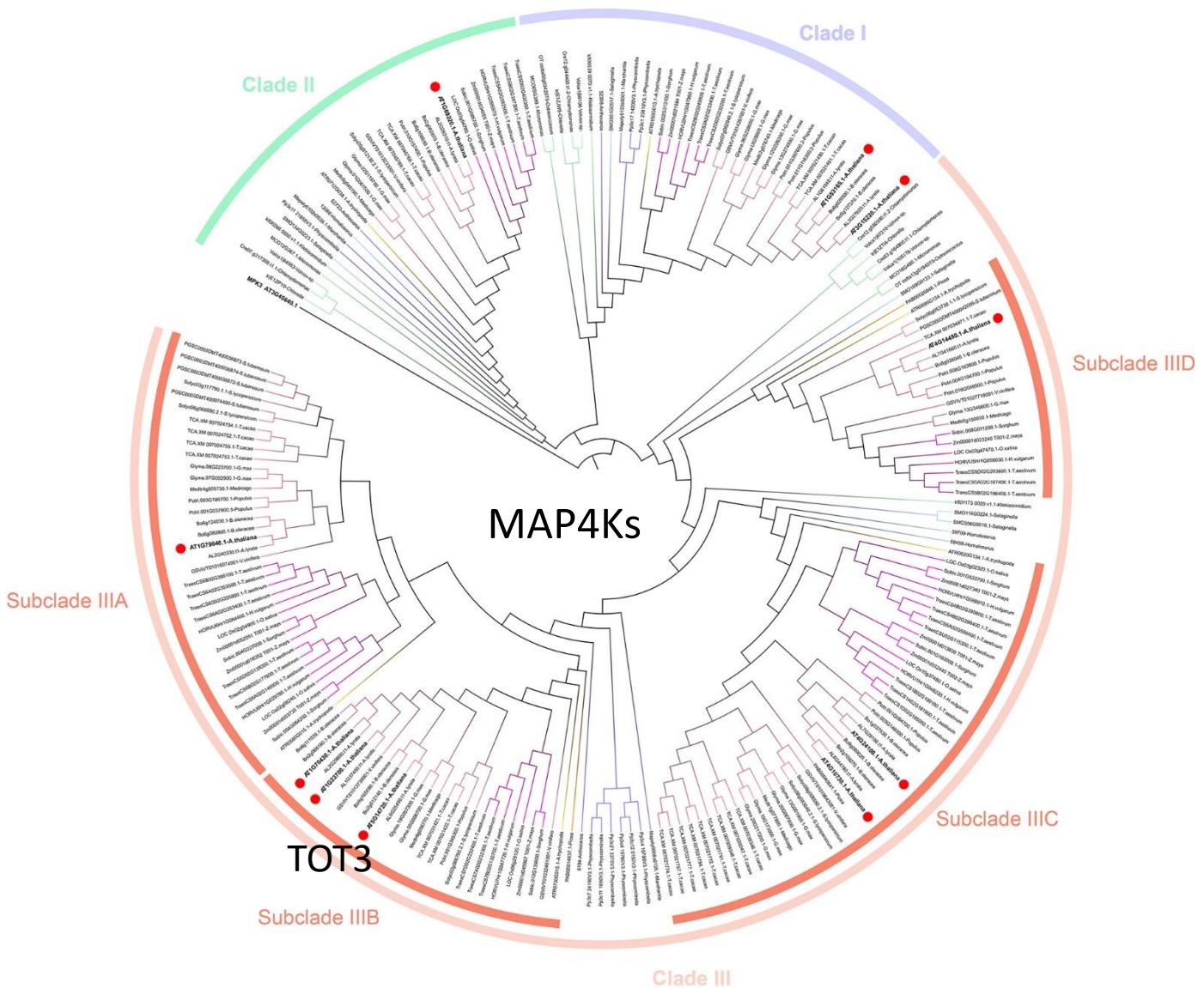
TaCOI1-1D mediates TaJAZ9-5B degradation



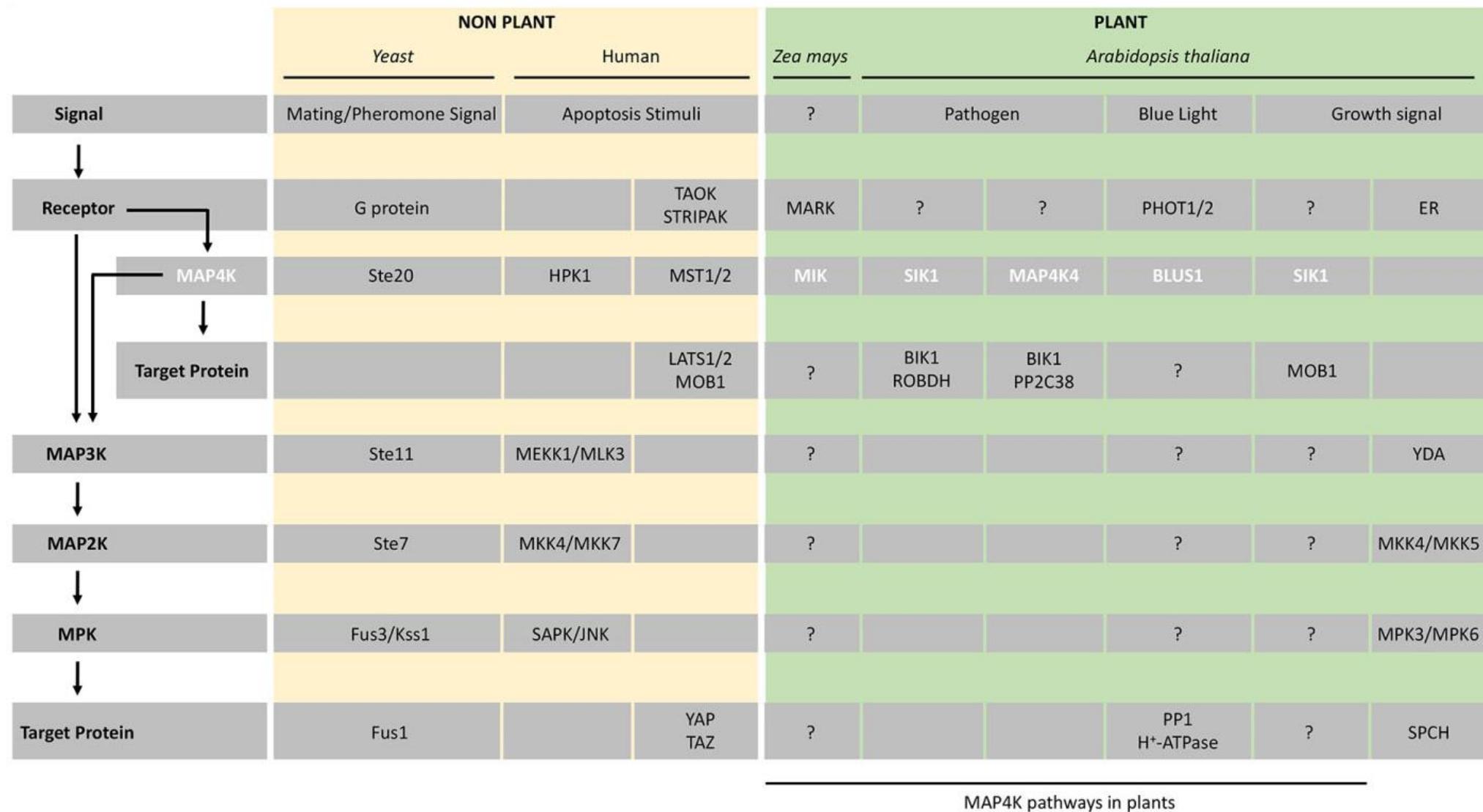
Are there conserved temperature-regulated proteins?



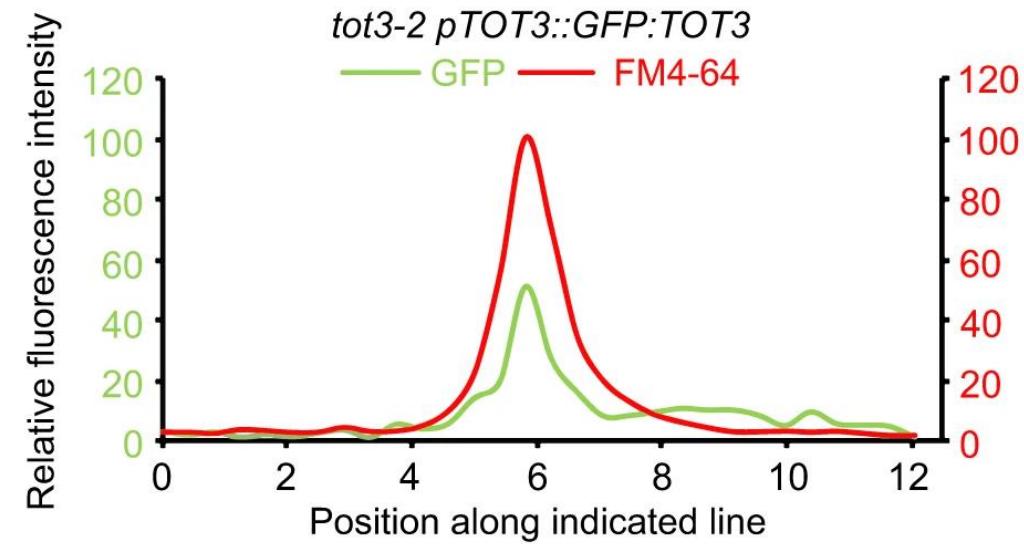
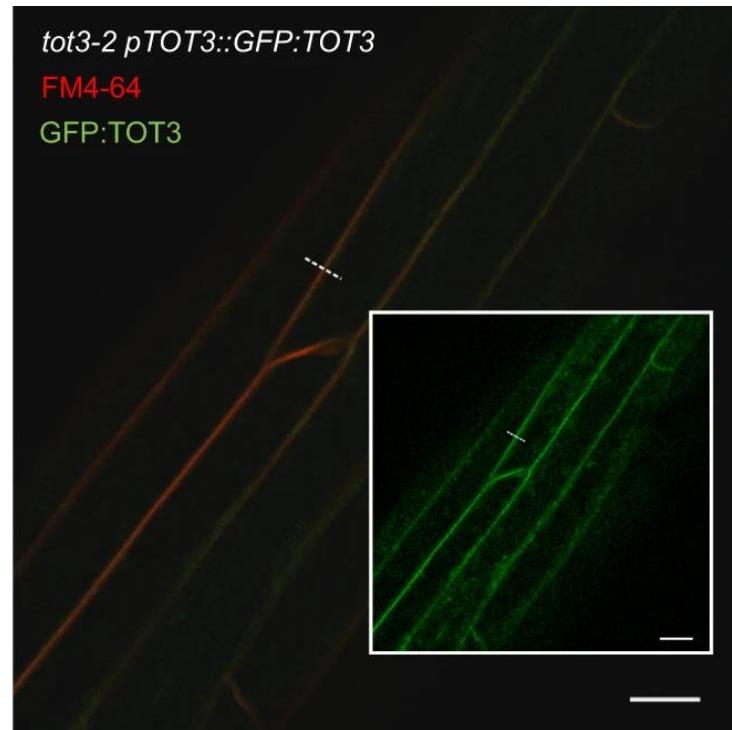
TOT3 is a MAP4K



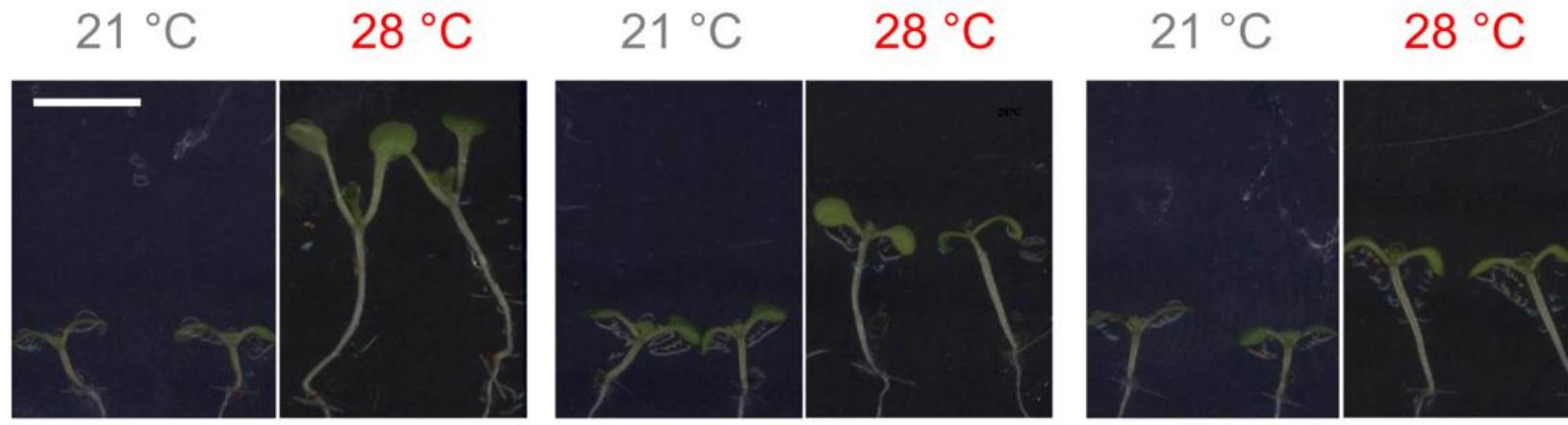
MAP4Ks are hardly explored in plants



TOT3 localizes to the plasma membrane



TOT3 is required for warm temperature-mediated growth in *Arabidopsis*

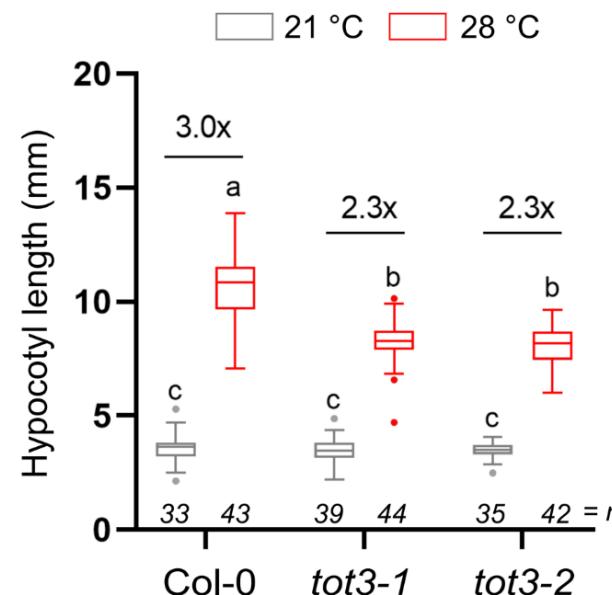


Col-0

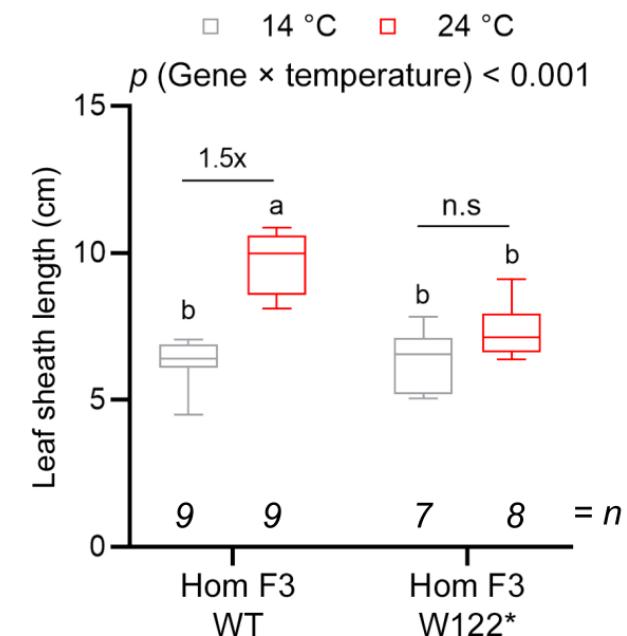
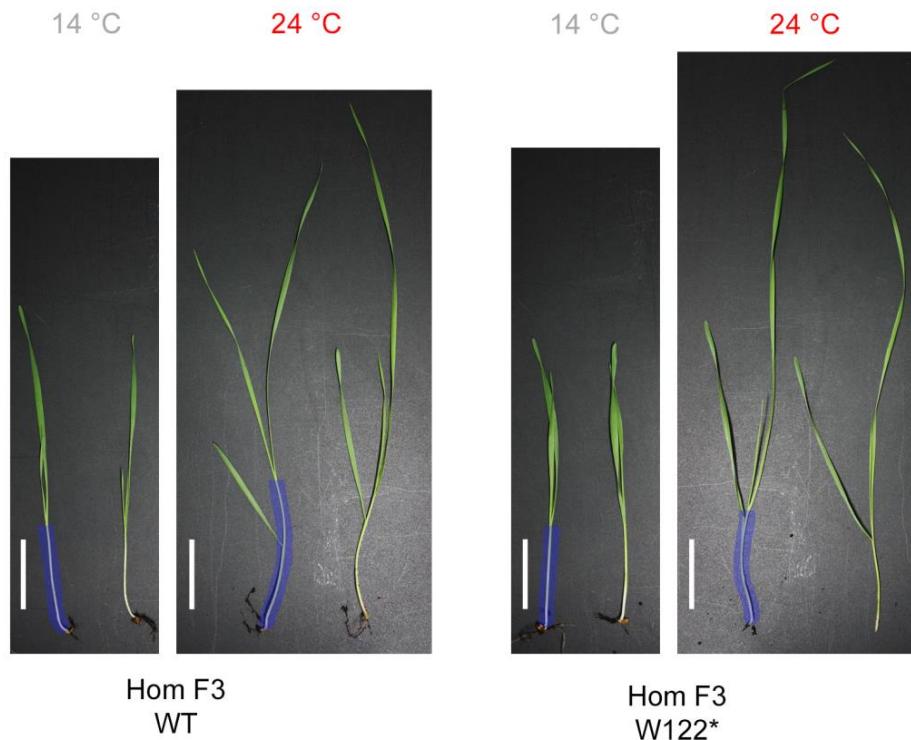
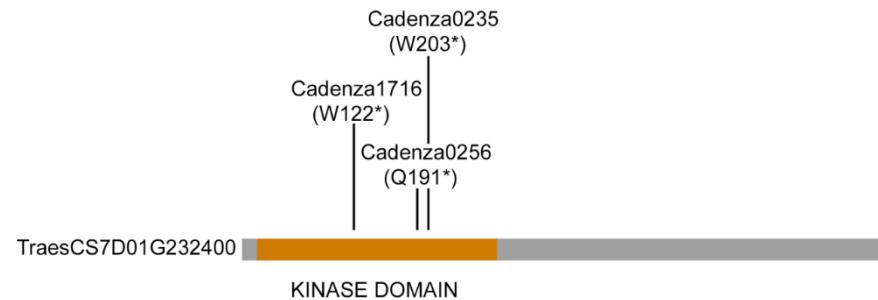
tot3-1

tot3-2

p (Gene \times temperature) < 0.0001



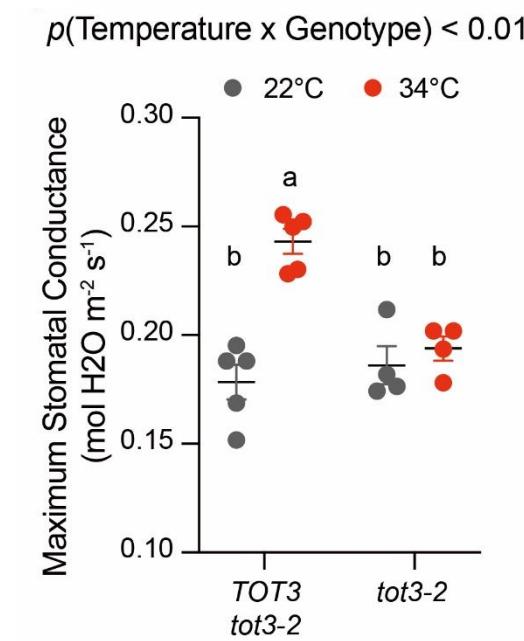
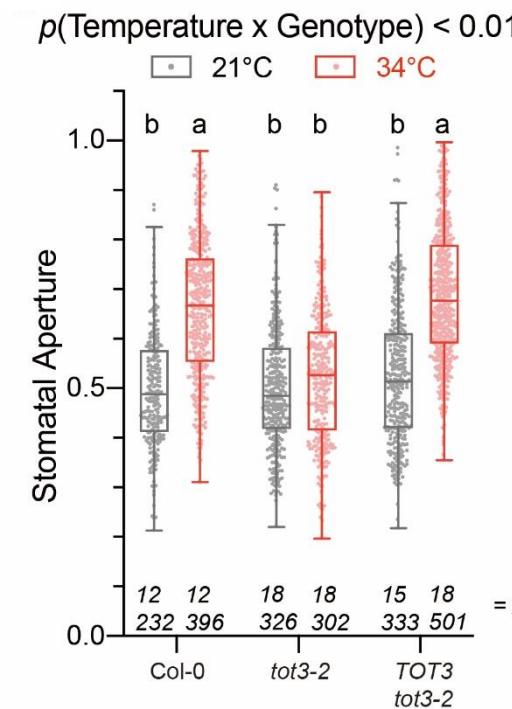
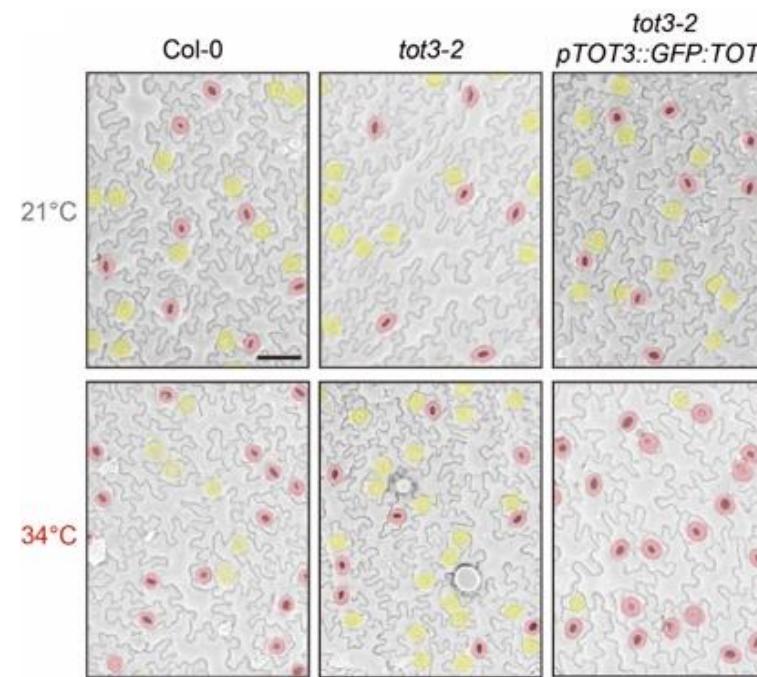
TOT3 is required for warm temperature-mediated growth in wheat



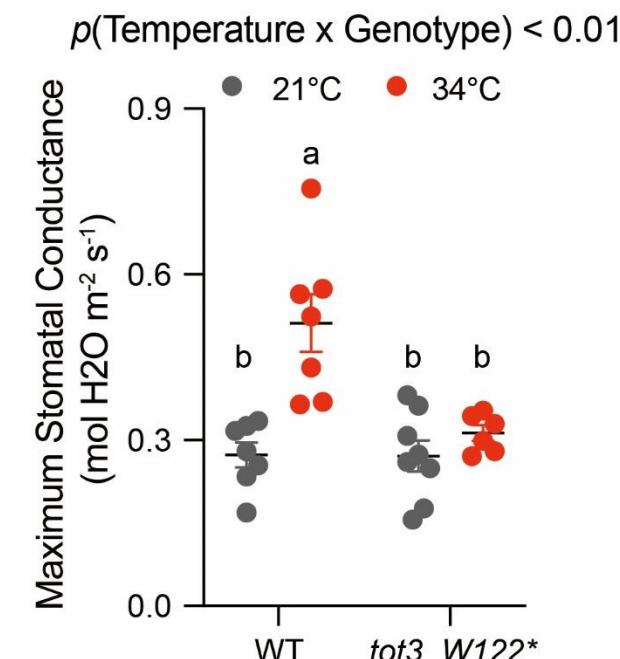
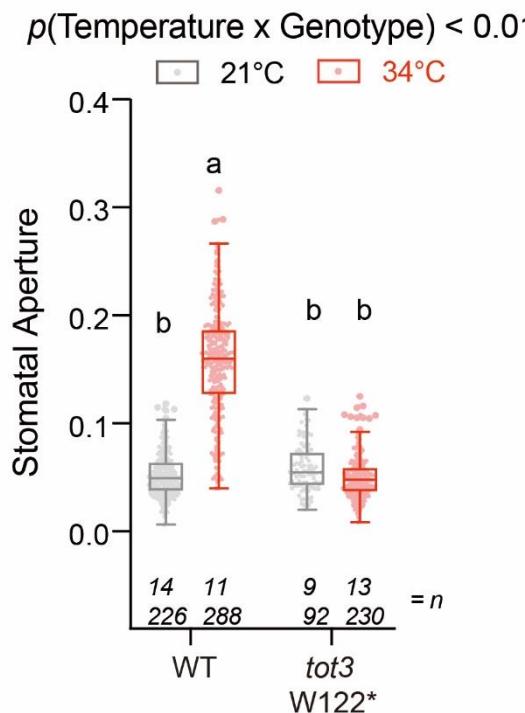
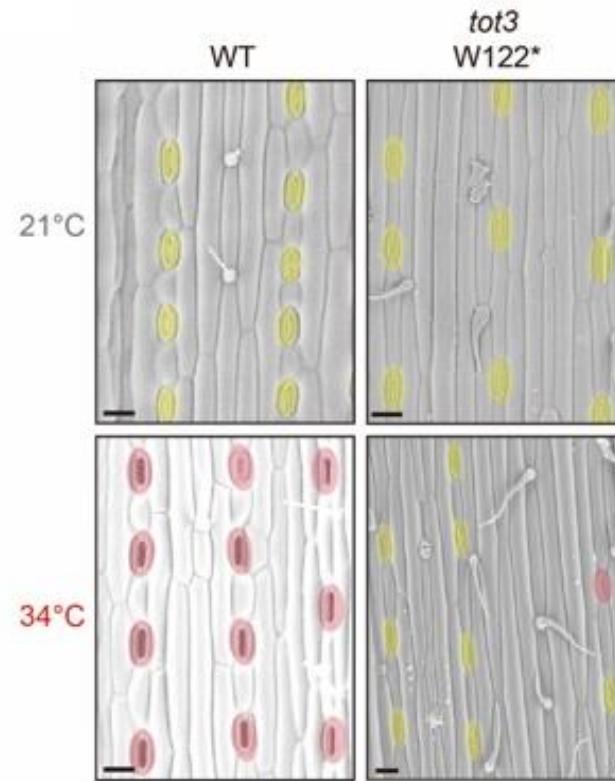
TOT3* is expressed in stomata in *Arabidopsis



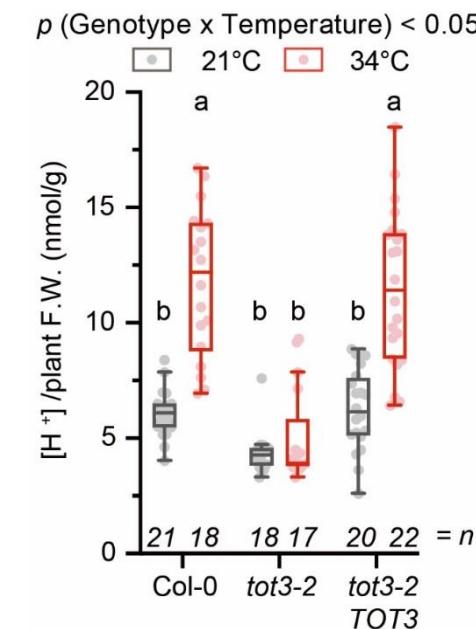
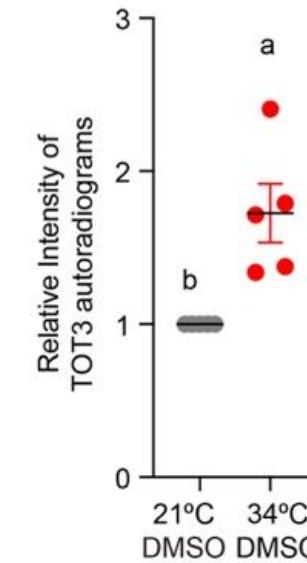
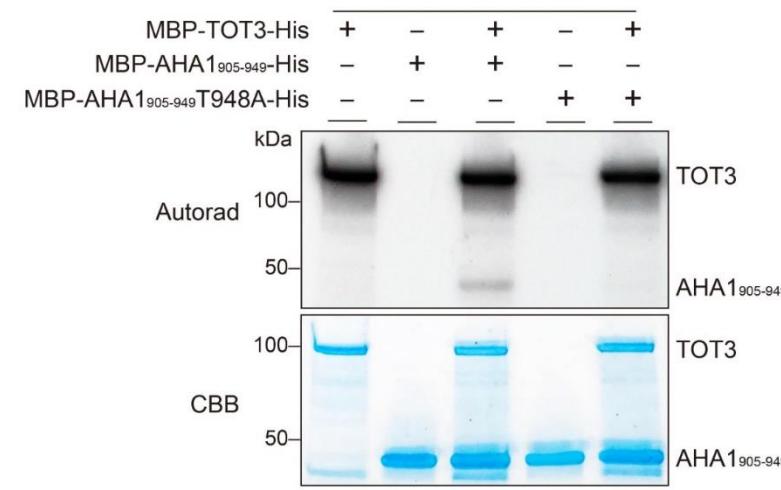
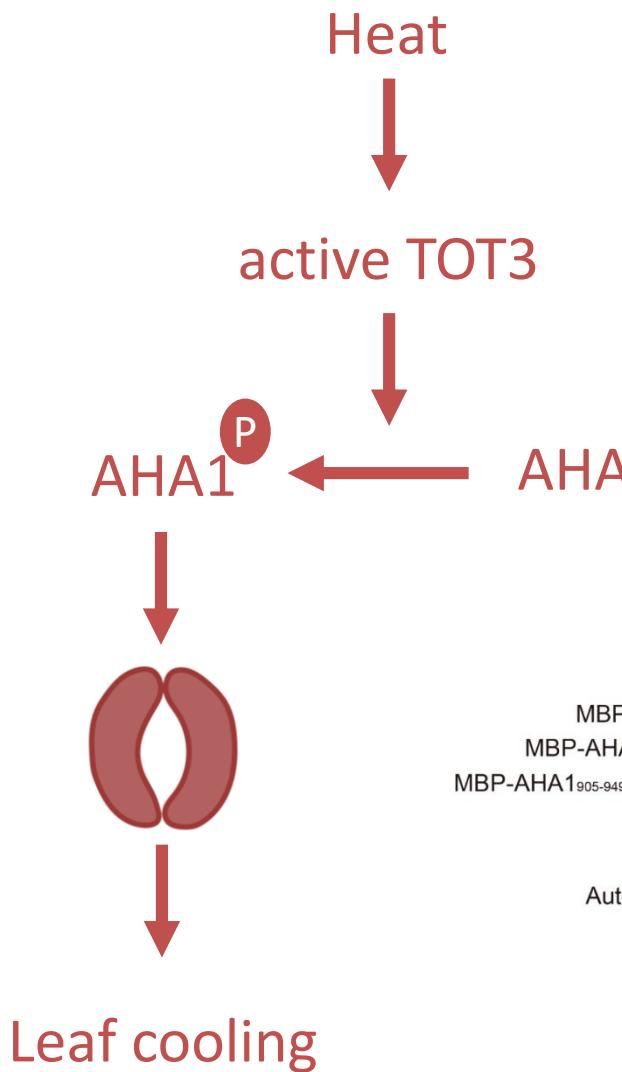
TOT3 is required for stomatal opening at high temperature in *Arabidopsis*



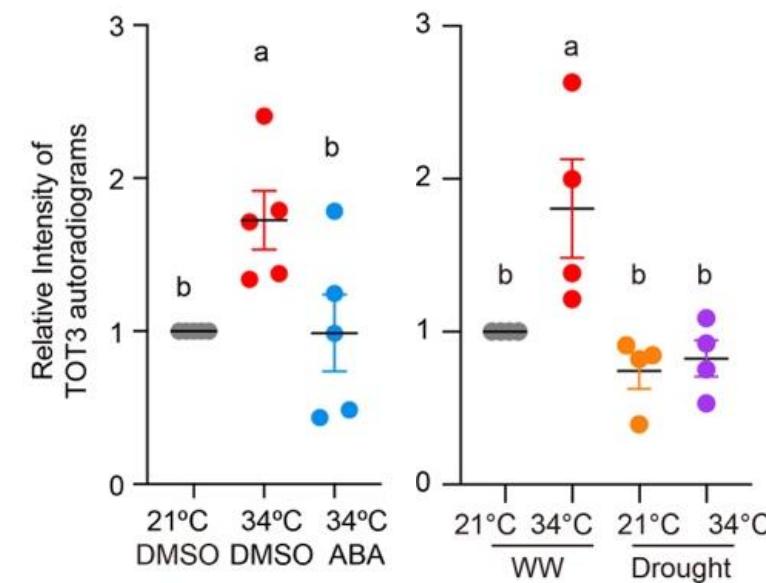
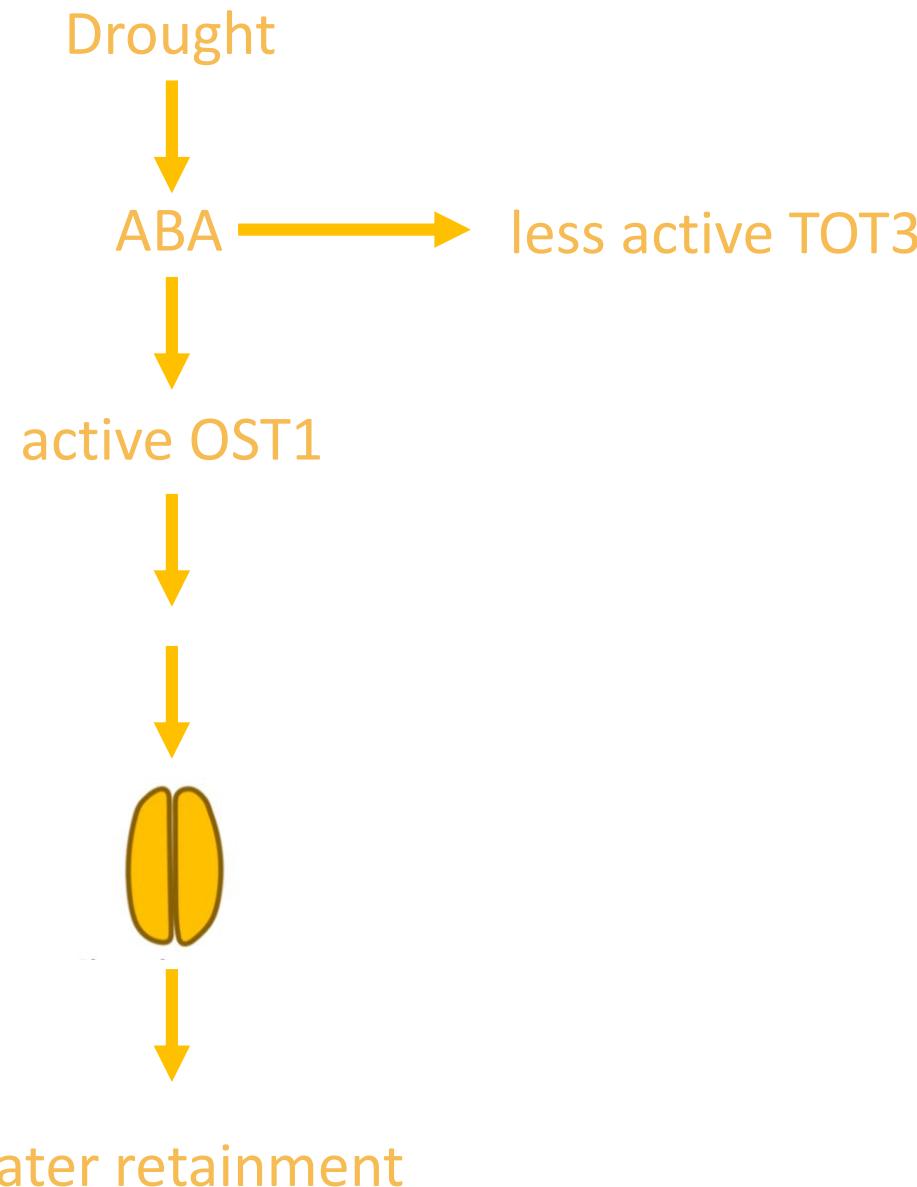
TOT3 is required for stomatal opening at high temperature in wheat



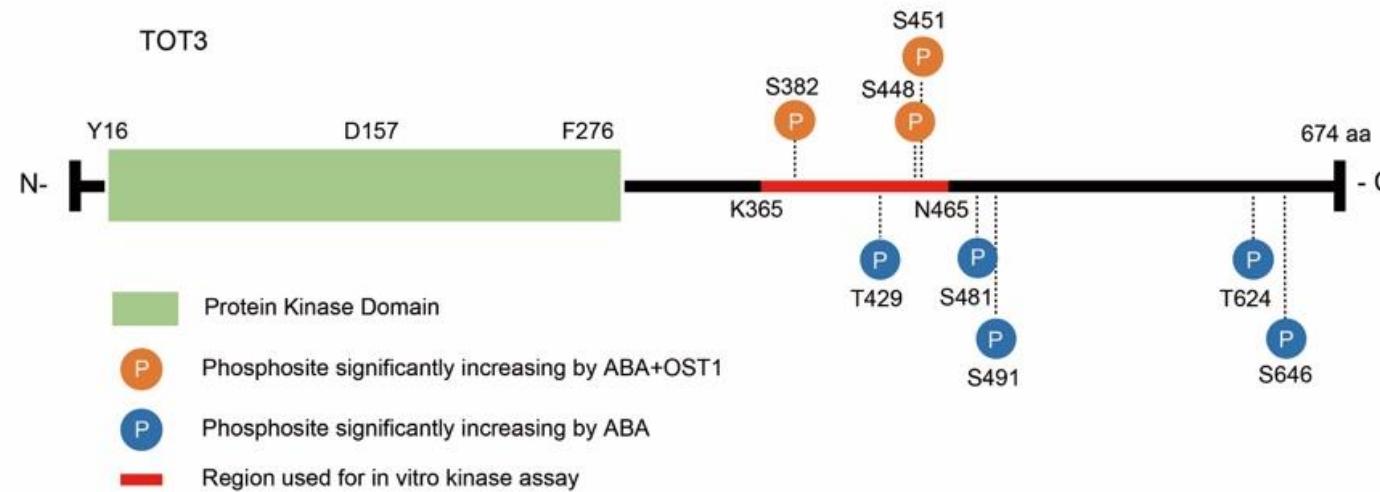
TOT3 controls AHA1 activity to open stomata by phosphorylating T948 upon heat



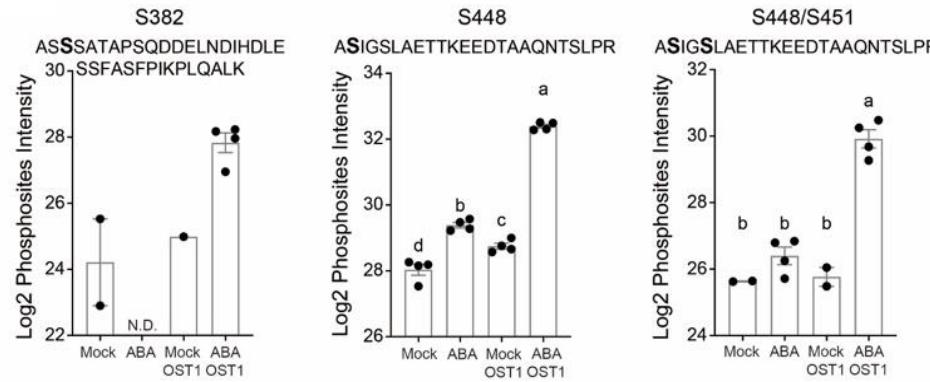
What happens under drought?



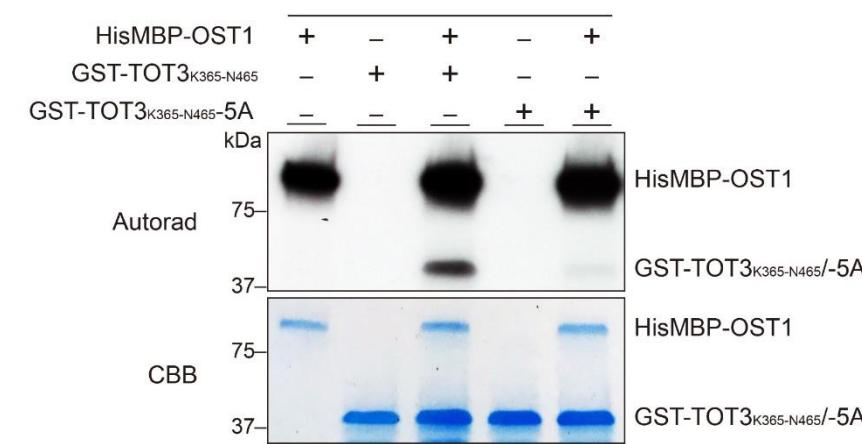
OST1 directly phosphorylates TOT3



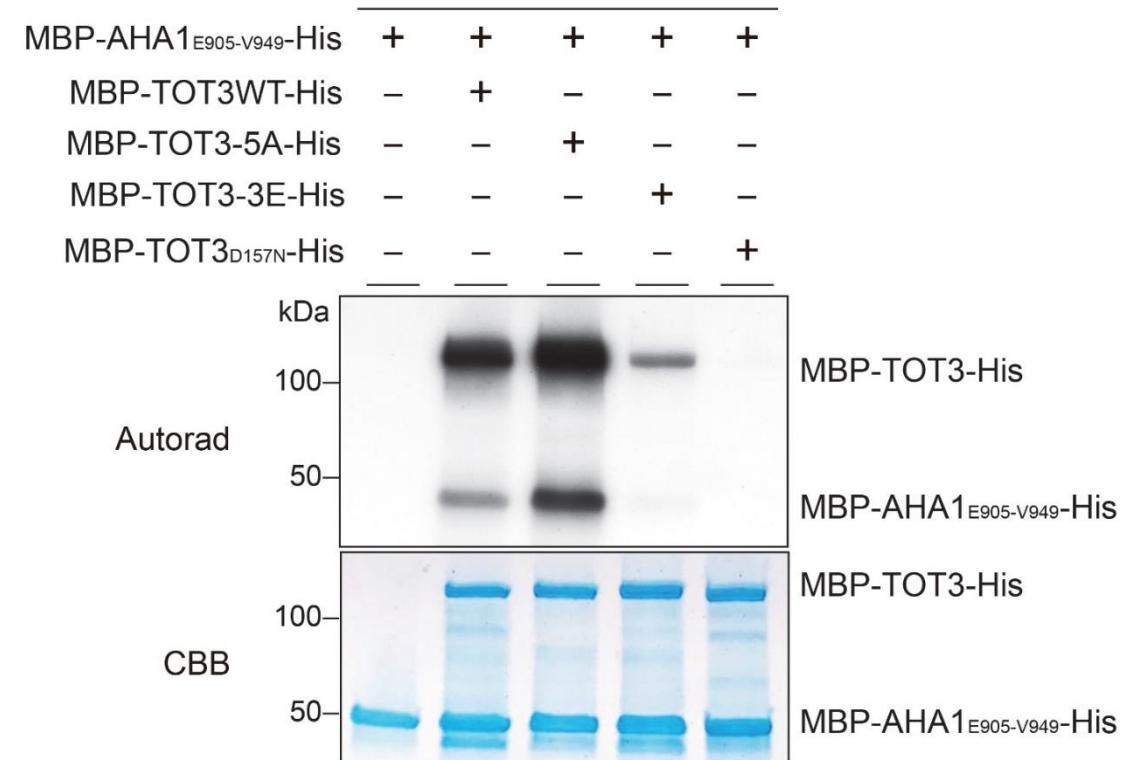
Transient tobacco assay



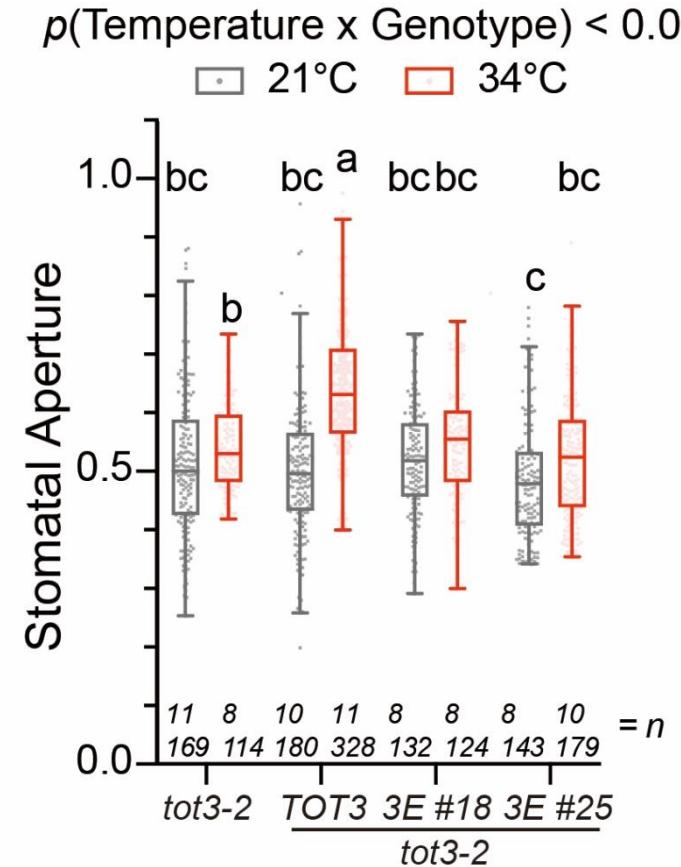
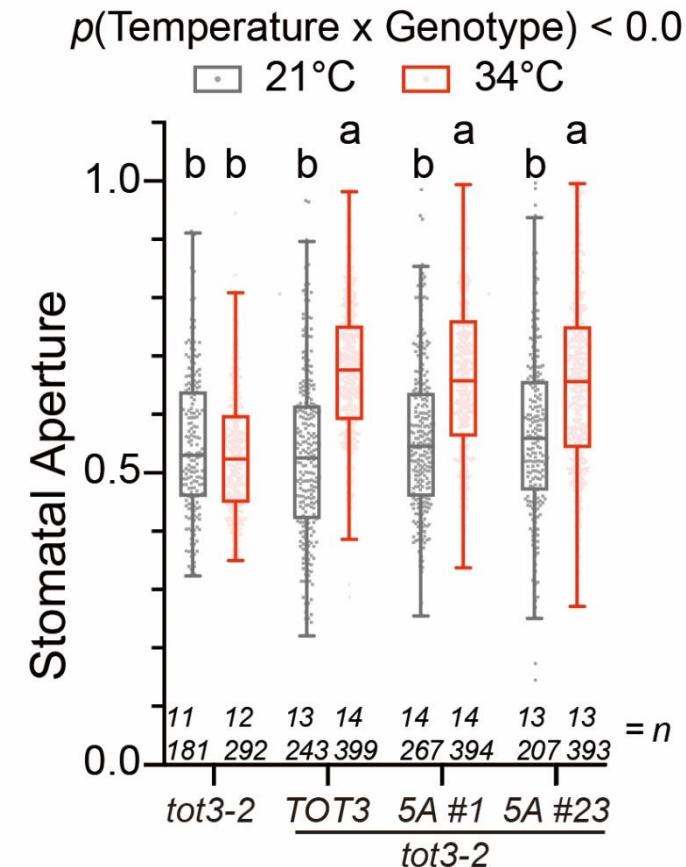
In vitro kinase assay



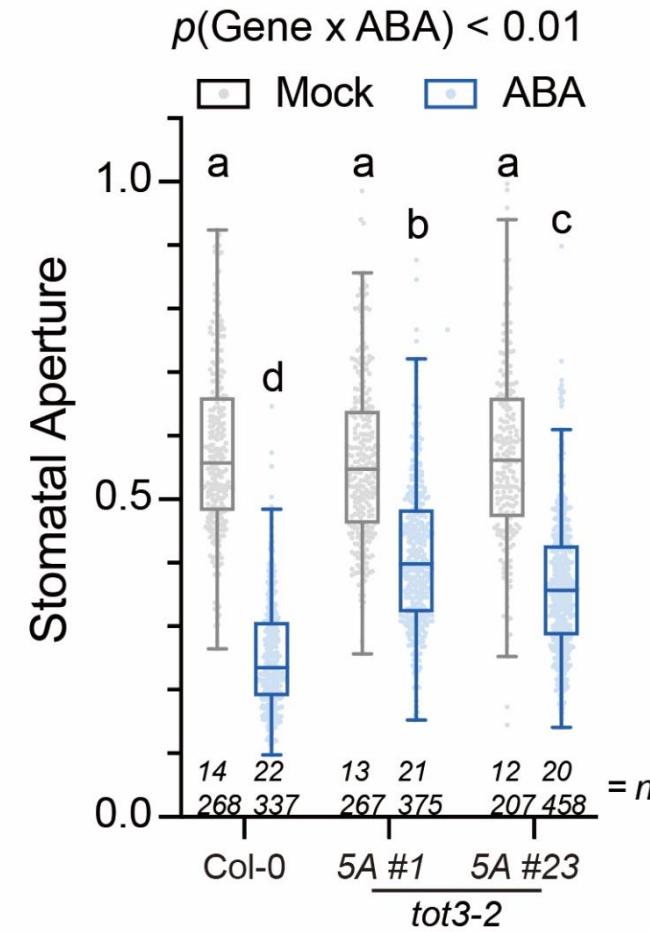
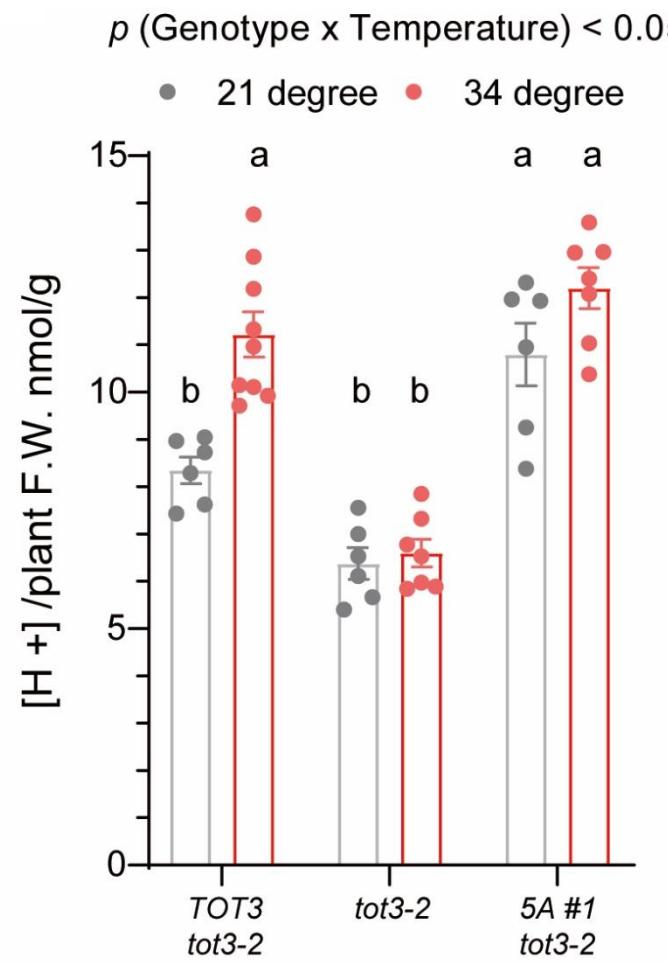
OST1-mediated TOT3 phosphorylation leads to loss of kinase activity



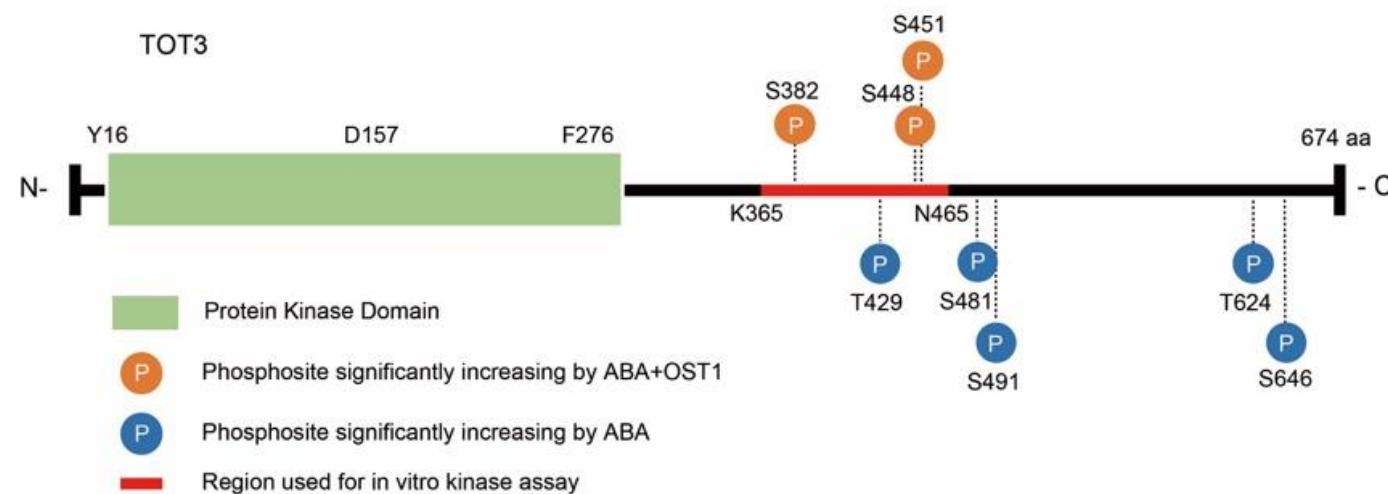
TOT3-5A = active / TOT3-3E = inactive



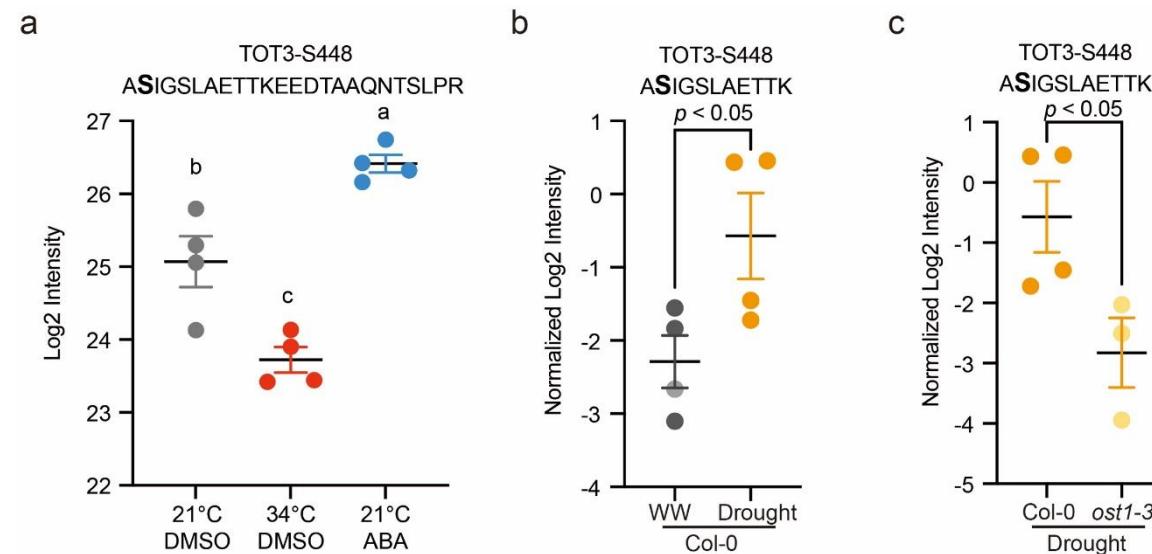
Impaired OST1-mediated TOT3 phosphorylation leads to more proton pump activity and reduced ABA-activated stomatal closure



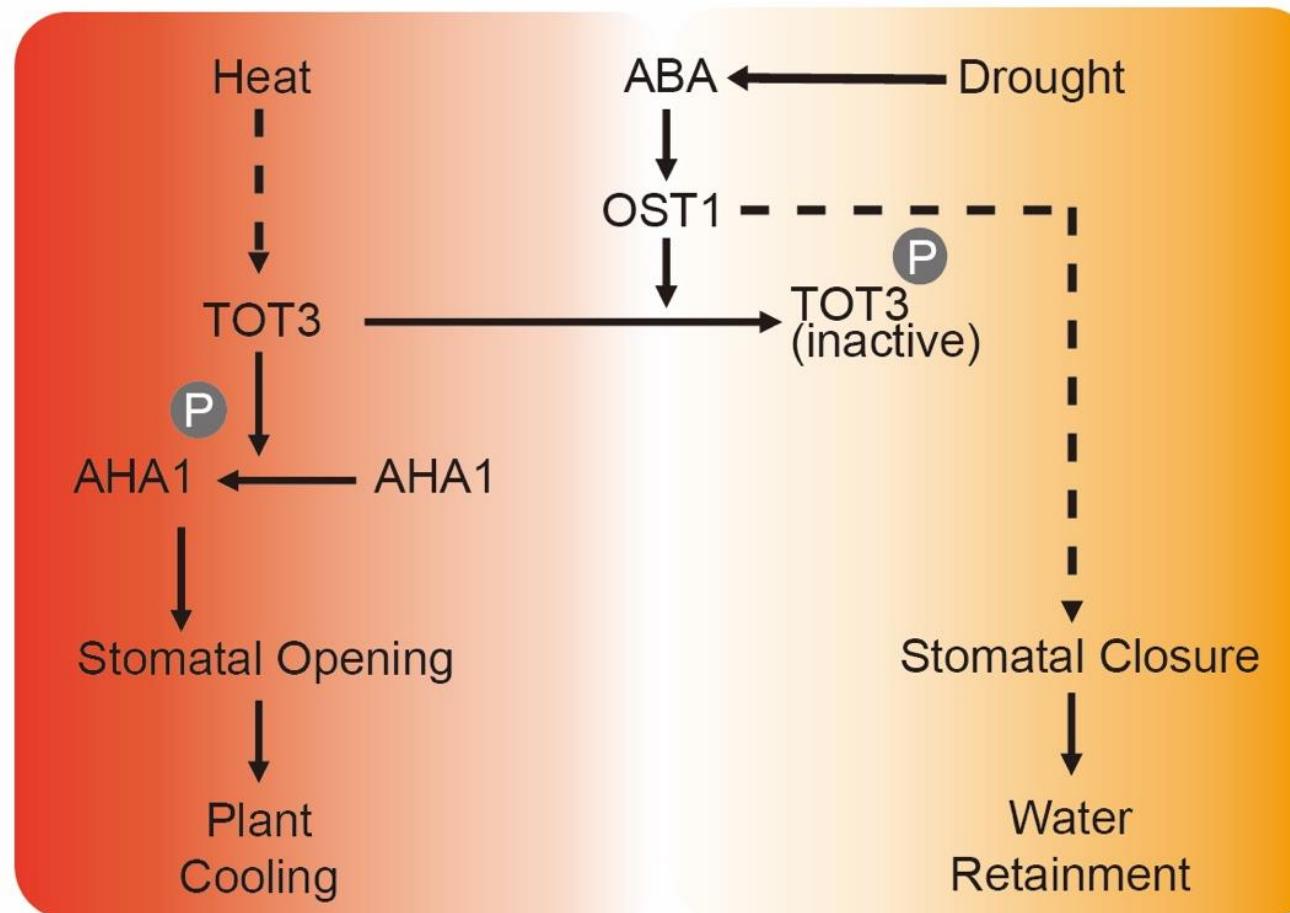
OST1 phosphorylates TOT3 upon ABA or drought



Arabidopsis



OST1 and TOT3 interplay balances water loss / cooling under drought / heat



Take Home Messages

- Capturing protein phosphorylation allows the identification of novel signaling pathways, including components that control temperature signaling and plasticity
- Phosphorylated proteins can be valuable breeding markers
- TOT3 function is conserved in wheat and *Arabidopsis*
- The TOT3-AHA1-OST1 signaling module integrates environmental signaling
- TaCOI1-1D encodes a JA receptor that is involved in regulating wheat growth

WHO DID THE WORK?



Group alumni

L.D. Vu, E. Stes, L. Joos, I. Verstraeten, S. Zhu, N. Nikonorova, P. Neyt, X. Xu

Collaborators

K. Gevaert, C. Uauy, IWGSC, M. van Zanten, A.M. Locke, E. Russinova, G. De Jaeger, D. Van Damme, M., G. A. Pizzio, Z. Jiang, B. Van De Cotte, N. Leonhardt, P. L. Rodriguez, S. Driever, R. Sozzani, L. Van den Broeck, A. Goossens, ...



desmetlab.be and ive.desmet@psb.vib-ugent.be

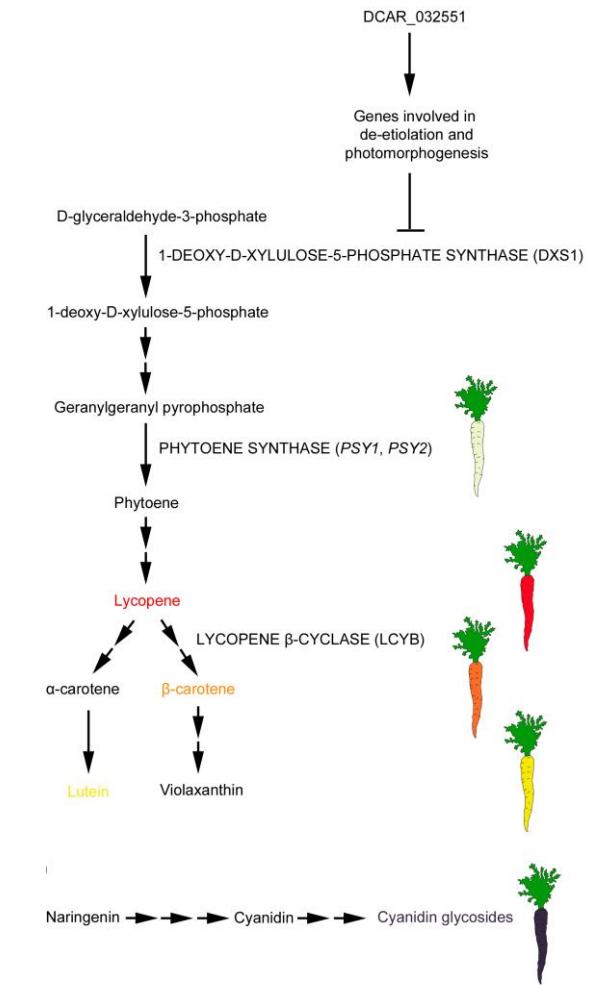
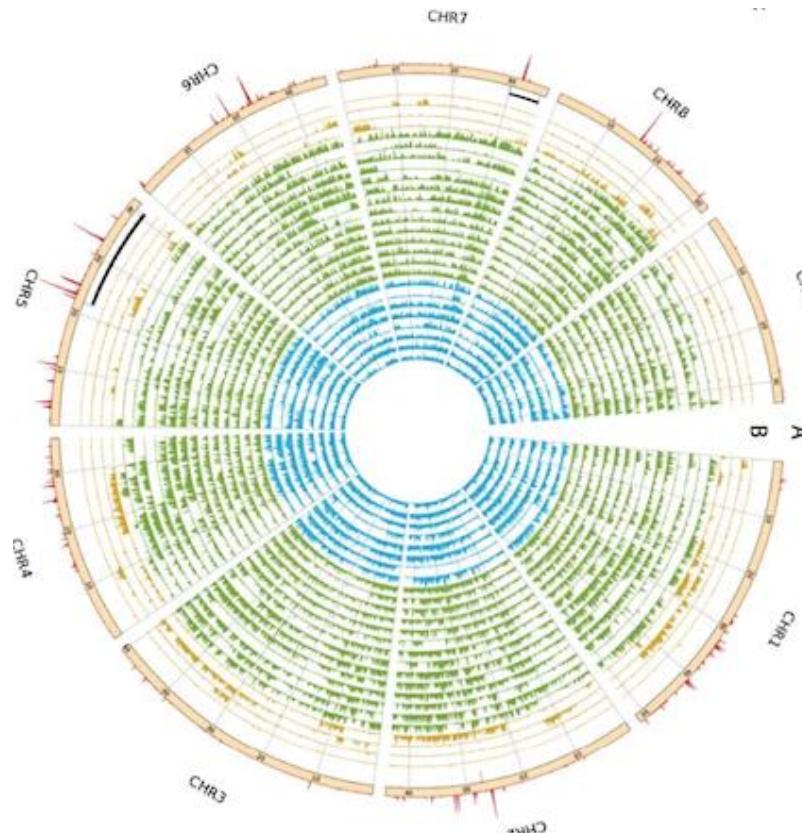


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#artgenetics - Belgian researchers

@david.vergauwen (art historian) and @desmetive (VIB-UGent) reveal plant diversity over time 🍉🍍



... meer

Trends in Plant Science

Science & Society

Down the Rabbit Hole—Carrots, Genetics and Art

David Vergauwen¹ and Ivo De Smet^{2,3,*}

The recent carrot genome assembly provides insight into carotenoid accumulation in carrots, and allows—together with other genetic information—to provide a molecular explanation for color differences observed in carrots painted throughout the centuries.

Carrots and Art

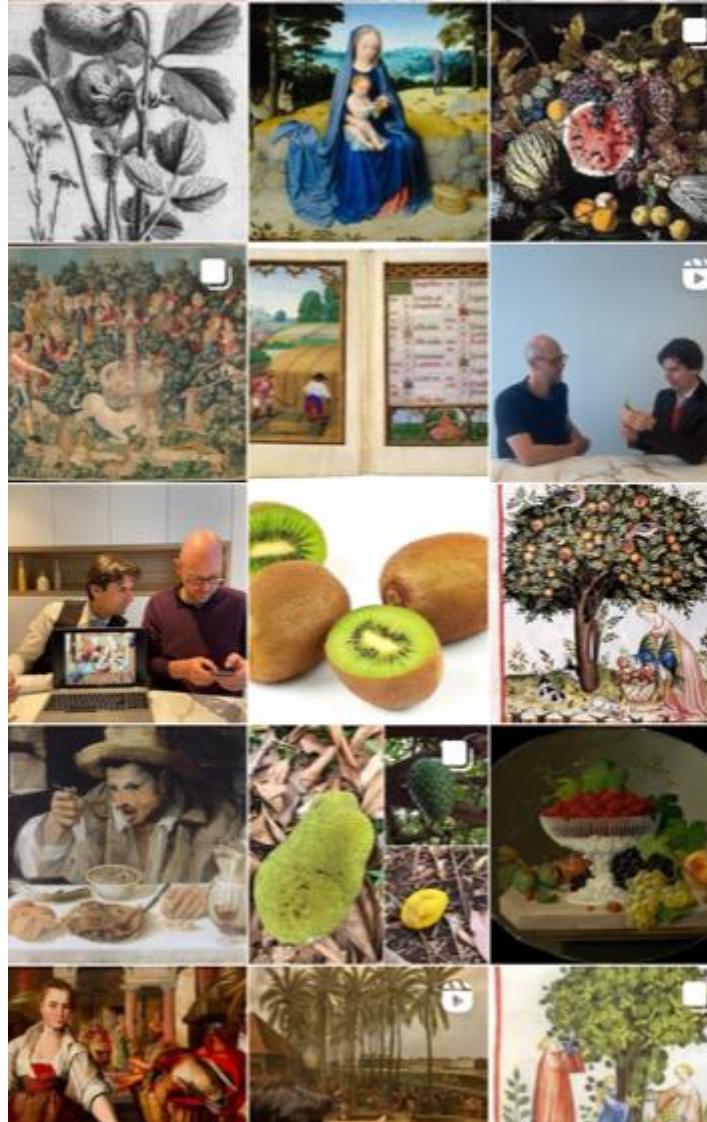
The large, unbranched cultivated carrot (‘*Kitāb al-Tabi’*), a book with about 600

some of the molecular mechanisms and underlying genetic network responsible for the observed colour differences.

On the Origin of the Carrot

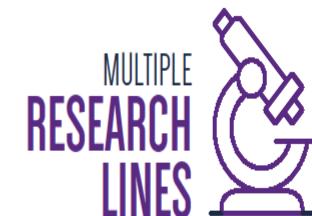
The origin of carrot is complex, but recent studies shed some light on the geographical distribution and domestication origin of wild and cultivated carrot [3]. Eastern and Western type carrots can be recognized, with the origin of the former in western to central Asia (purple, red or orange-yellow roots) and the latter derived from the Eastern type (yellow, orange, or sometimes-white roots) [3,4]. The carrot was probably first domesticated somewhere in the area between the north east of Kabul (Afghanistan) and north Kashmir (India) around AD 900 [5]. Later, in the 10th century, a Bagdad-based cook, Ibn Sayyar al-Warraq, mentioned in his *Juliana Anicia Codex*. From the 11th century drawings of (yellow) carrots started to

CellPress



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