

EXPLORING THE POTENTIAL  
OF *BRASSICA*  
CROP WILD RELATIVES:  
INSIGHTS FOR SUSTAINABLE  
AGRICULTURE AND BIODIVERSITY  
CONSERVATION

Makenzie E Mabry, PhD

USDA Postdoctoral Fellow

Florida Museum of Natural History, University of Florida

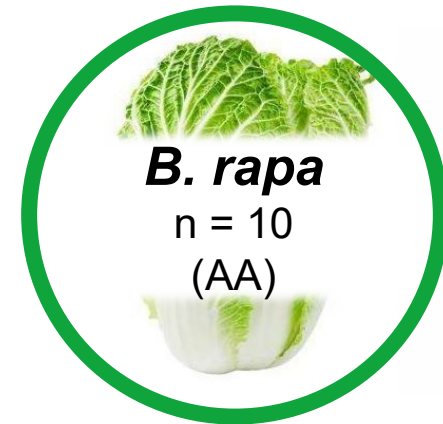
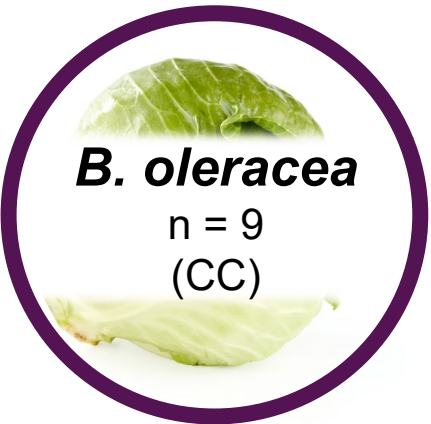
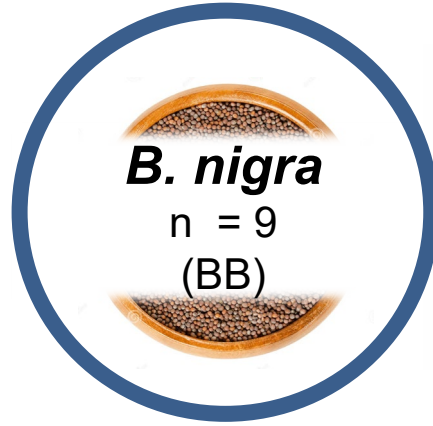
New York Botanical Garden



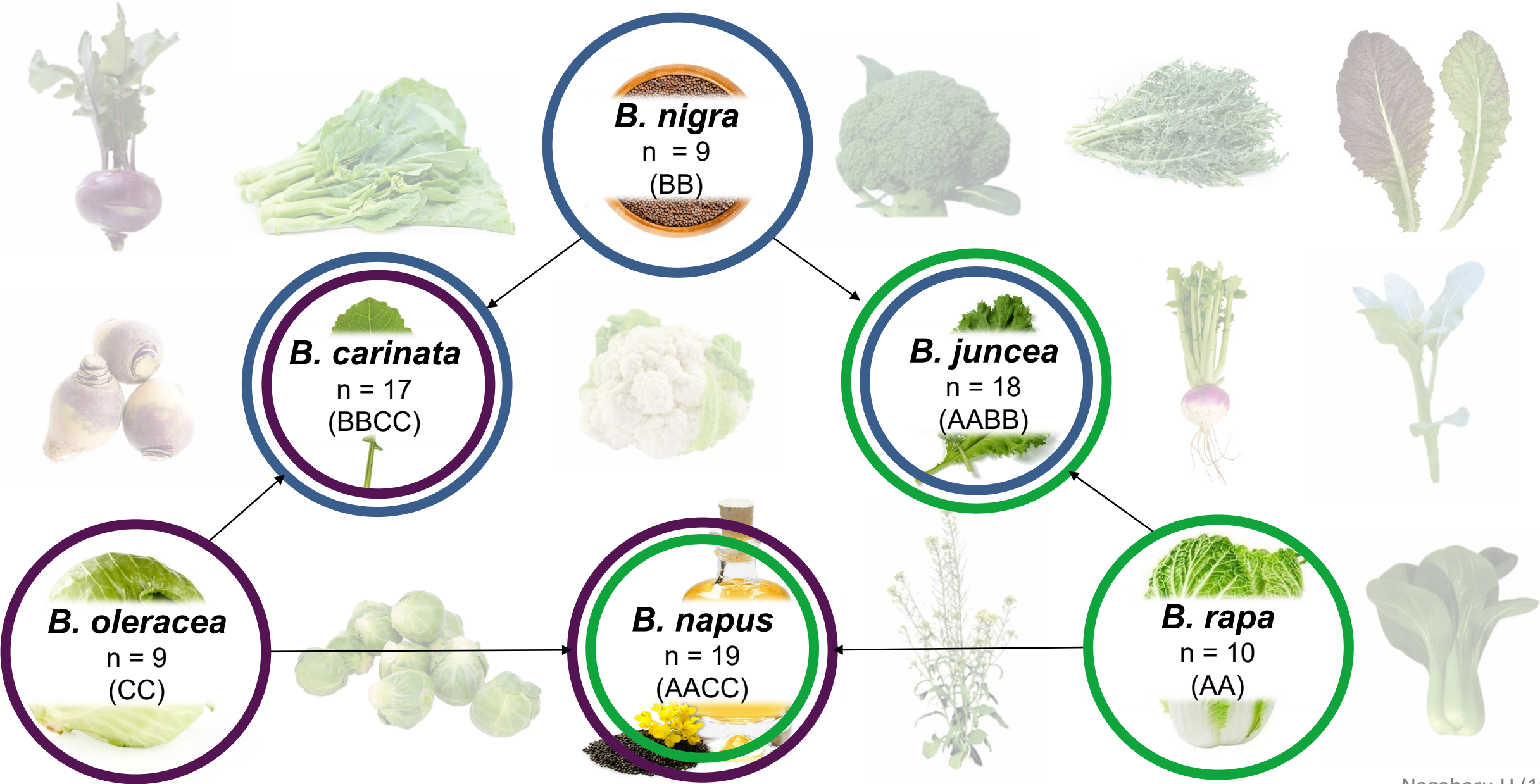
# BRASSICA CROP DIVERSITY IS UNMATCHED



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# TRIANGLE OF U



## WILD BRASSICAS – CROP WILD RELATIVES (CWRs)



- A **crop wild relative (CWR)** is a wild plant closely related to a domesticated plant. It may be a wild ancestor of the domesticated (cultivated) plant or another closely related taxon.

# WILD BRASSICAS – CROP WILD RELATIVES (CWRs)

*Includes 37 Accepted Species*



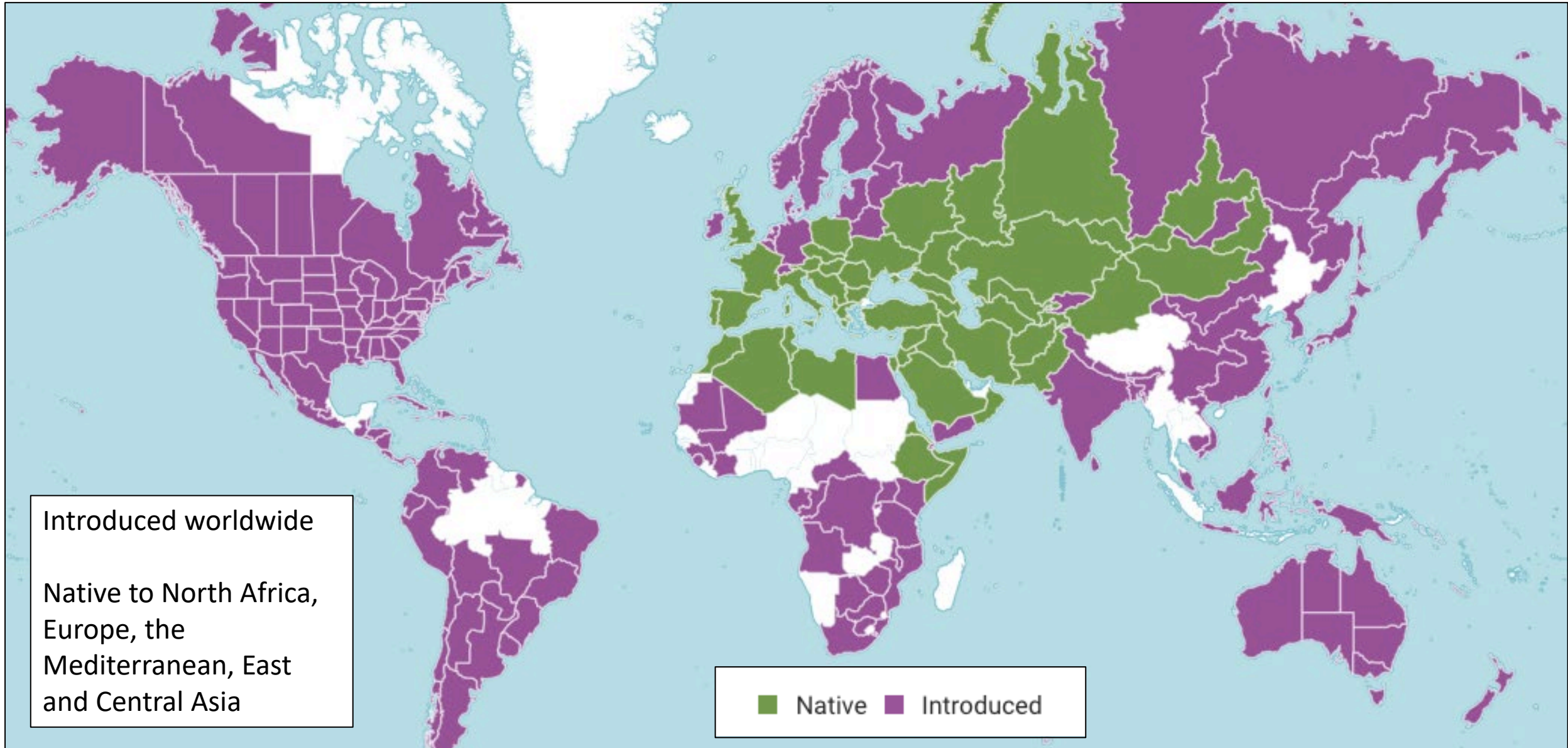
# WILD BRASSICAS – CROP WILD RELATIVES (CWRs)

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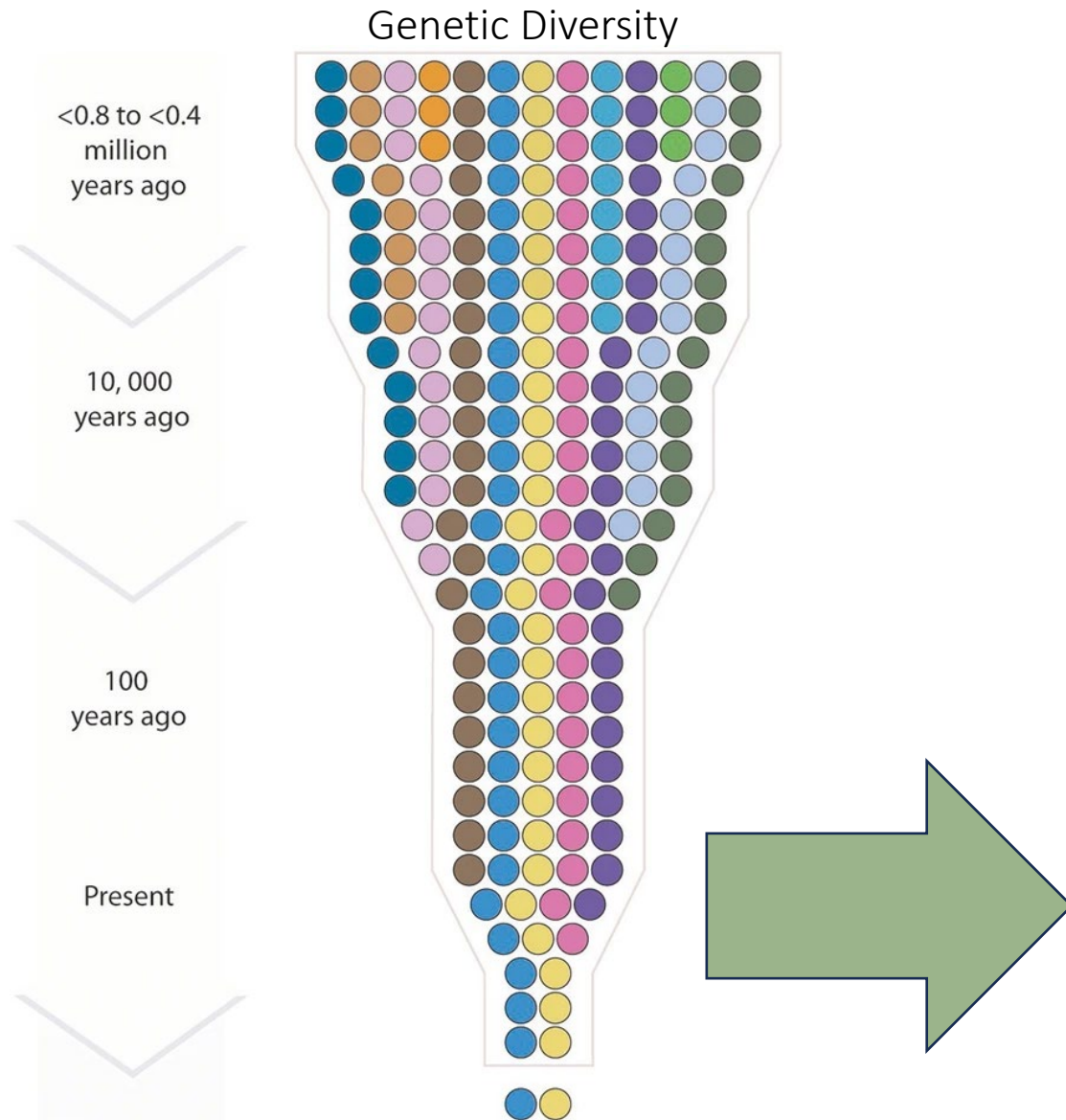


- *Brassica assyriaca*
- *Brassica aucheri*
- *Brassica baldensis*
- *Brassica balearica*
- *Brassica barrelieri*
- *Brassica beytepeensis*
- *Brassica bourgeauii*
- *Brassica cadmea*
- *Brassica cretica*
- *Brassica deflexa*
- *Brassica deserti*
- *Brassica desnottesii*
- *Brassica dimorpha*
- *Brassica elongata*
- *Brassica fruticulosa*
- *Brassica gravinae*
- *Brassica hilarionis*
- *Brassica incana*
- *Brassica insularis*
- *Brassica loncholoma*
- *Brassica macrocarpa*
- *Brassica maurorum*
- *Brassica montana*
- *Brassica nivalis*
- *Brassica oxyrrhina*
- *Brassica procumbens*
- *Brassica repanda*
- *Brassica rupestris*
- *Brassica setulose*
- *Brassica somalensis*
- *Brassica souliei*
- *Brassica spinescens*
- *Brassica taiwanensis*
- *Brassica taurica*
- *Brassica trichocarpa*
- *Brassica tyrrhena*
- *Brassica villosa*

# WILD BRASSICAS – WORLDWIDE DISTRIBUTION

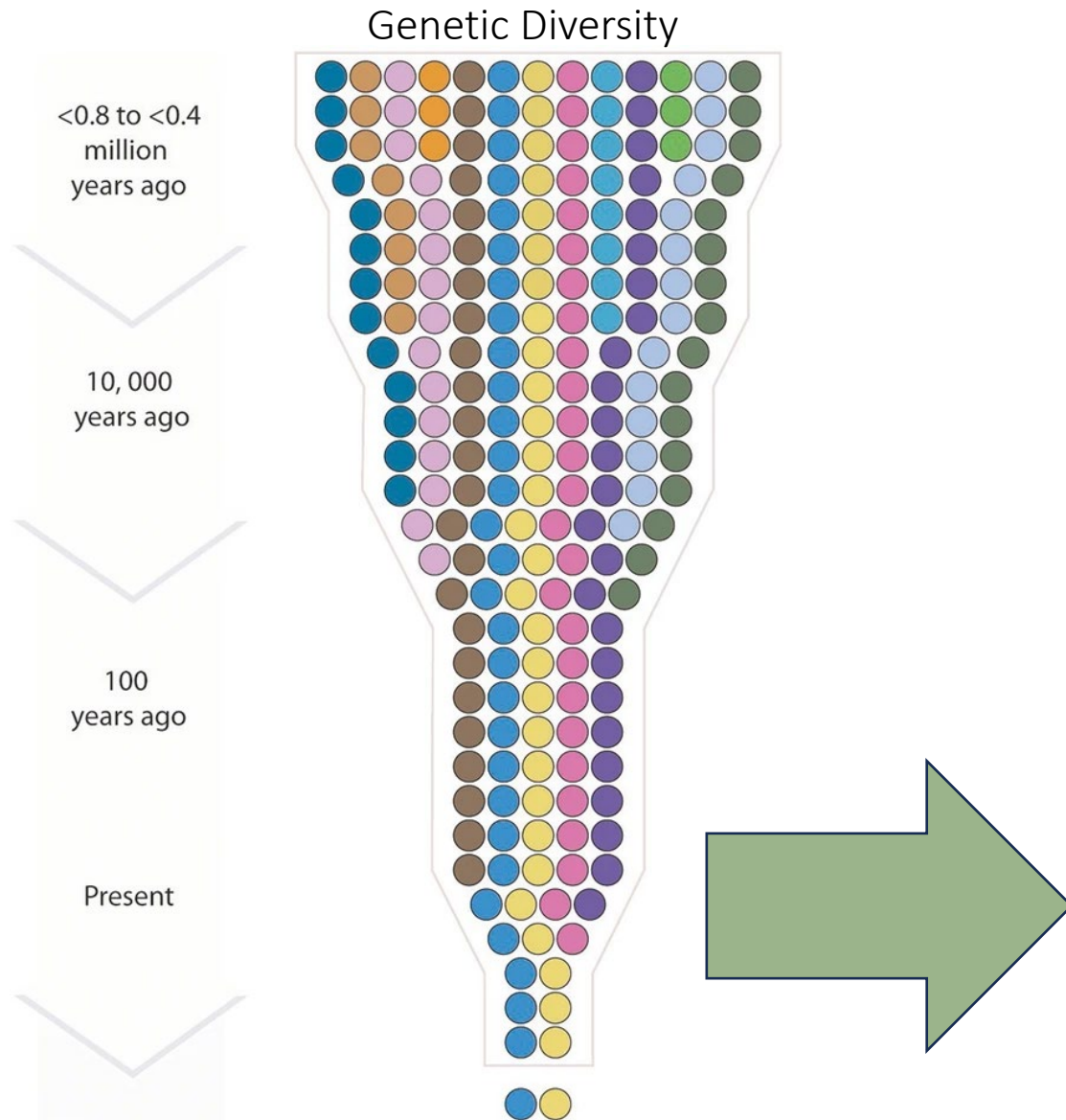


# WILD BRASSICAS HAVE HIGH GENETIC DIVERSITY



*Low genetic diversity*

# WILD BRASSICAS HAVE HIGH GENETIC DIVERSITY



*Wild Brassicas*



*High genetic diversity*



*Low genetic diversity*

# WILD BRASSICAS – CROP WILD RELATIVES (CWRs)



*Brassica cretica*



*Brassica montana*



*Brassica tournefortii*

Plants with high tolerance for drought, heat, salinity, etc.

# WILD BRASSICAS – CROP WILD RELATIVES (CWRs)



*Brassica cretica*



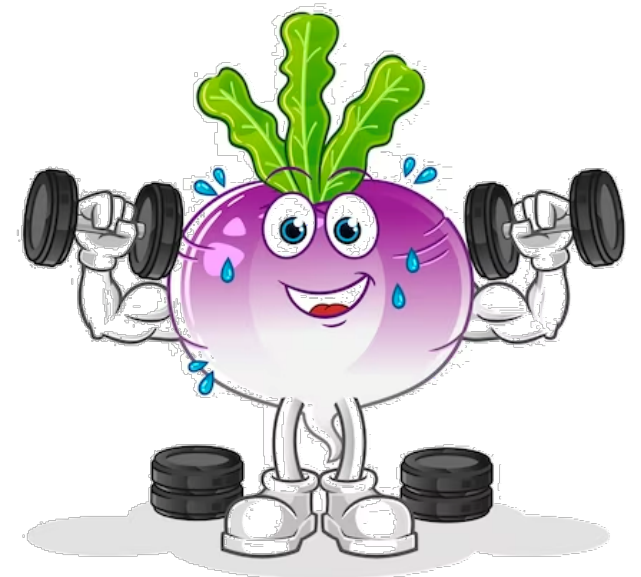
*Brassica montana*



*Brassica tournefortii*

Plants with high tolerance for drought, heat, salinity, etc.

*Breeding Resilient Crop  
Breeds or Climate Ready  
Crops*



In order to fully leverage CWRs,  
we must first work to understand their relationships  
to the crop of interest.



# THE ROLE OF CWRs IN DOMESTICATION

## 1 – *Brassica oleracea*



illustrations by Andi Kur

## 2 – *Brassica rapa*



## *Brassica napus*



# THE EVOLUTIONARY HISTORY OF WILD, DOMESTICATED, AND FERAL *BRASSICA OLERACEA* (BRASSICACEAE)

*Molecular Biology and Evolution*, Volume 38, Issue 10, October 2021, Pages 4419–443



Sarah Turner-Hissong



Evan Gallagher



Alex McAlvay



Hong An



Patrick Edger



Jonathan Moore



David Pink



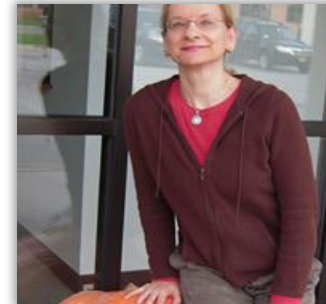
Graham Teakle



Chris J. Stevens



Guy Barker



Joanne Labate



Dorian Fuller



Robin Allaby



Timothy Beissinger



Jared Decker



Michael Gore



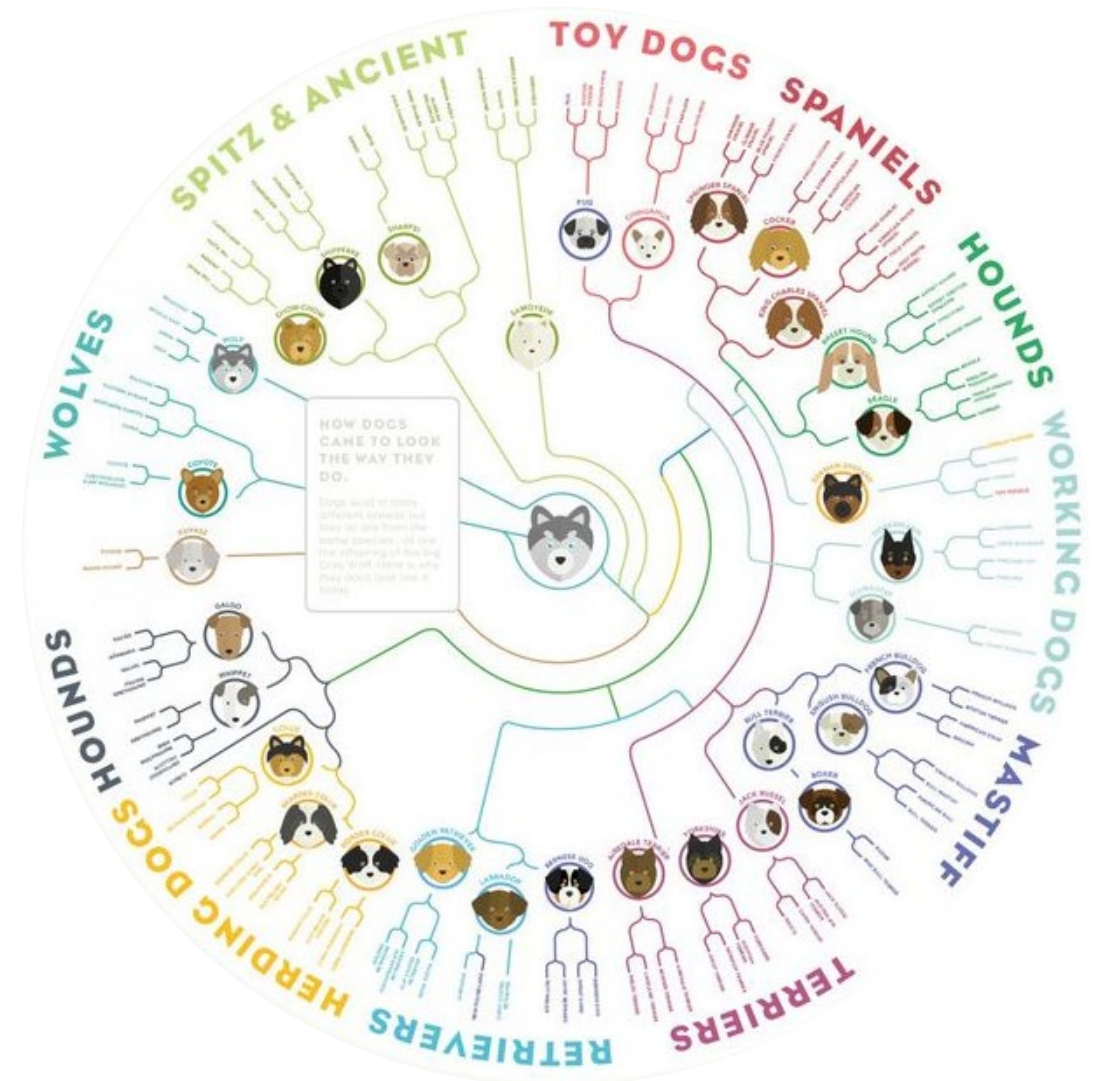
J. Chris Pires



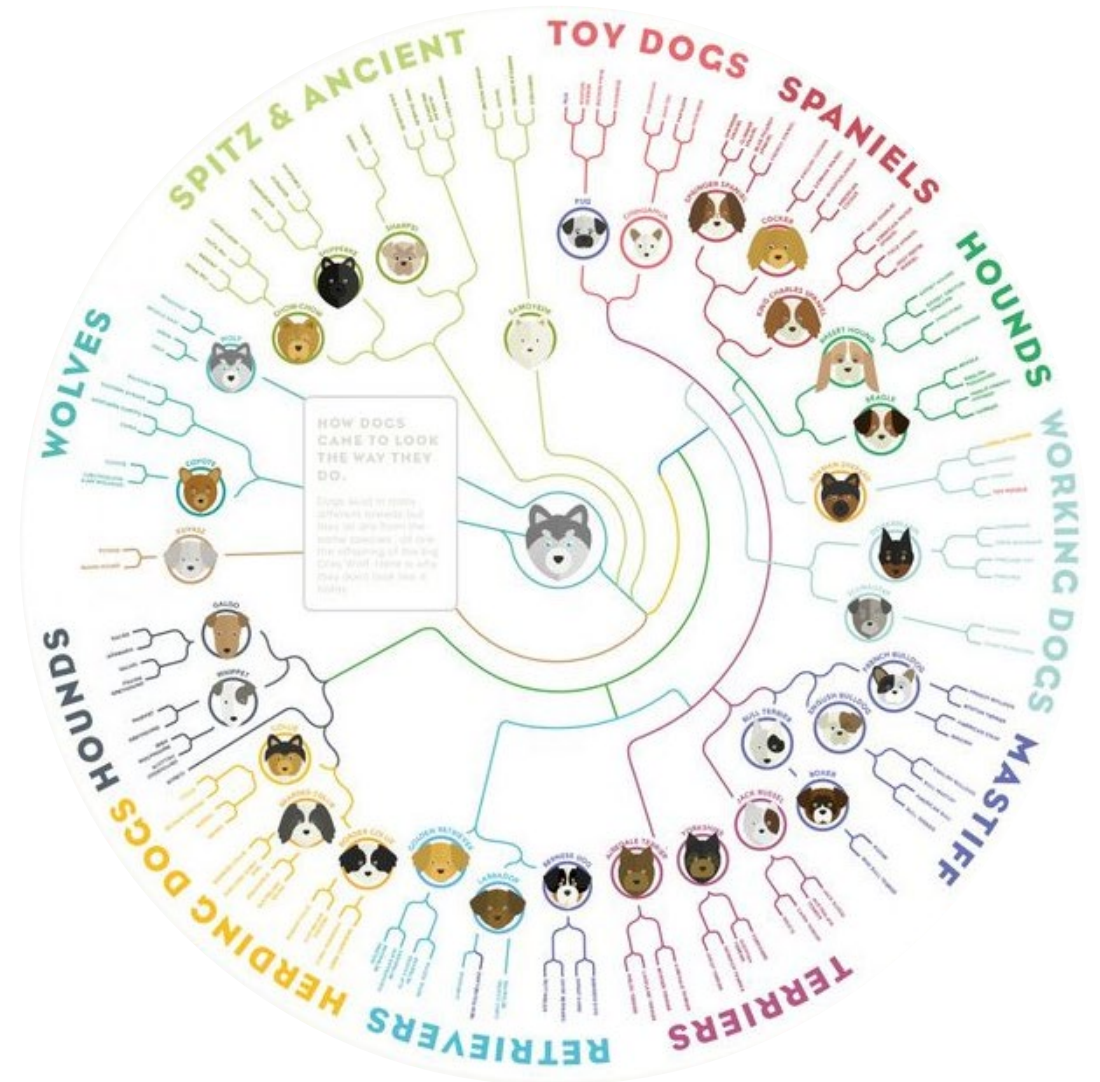
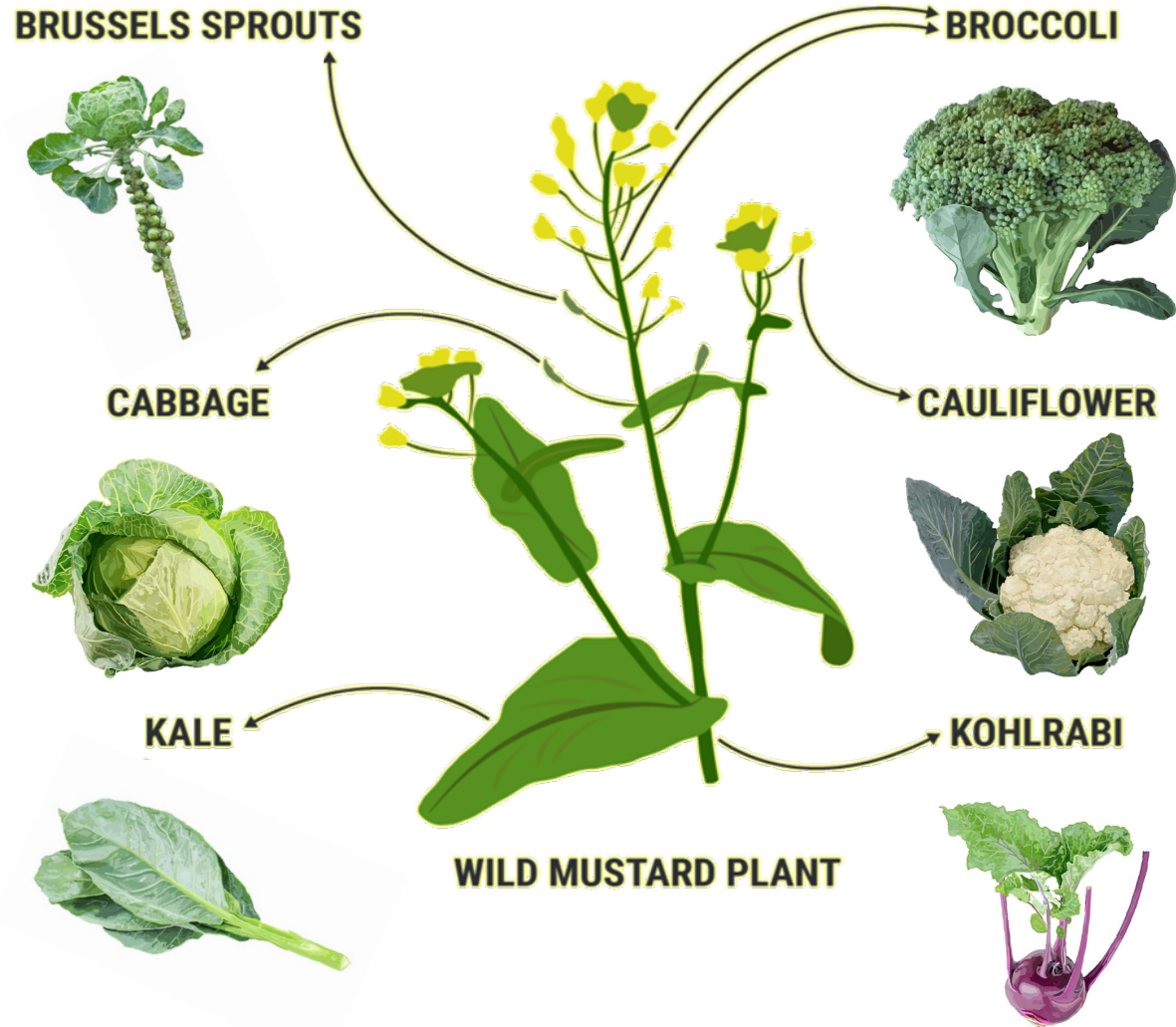
Scan to access the  
paper

# *BRASSICA OLERACEA* – THE DOG OF THE PLANT WORLD

# BRASSICA OLERACEA – THE DOG OF THE PLANT WORLD



# BRASSICA OLERACEA – THE DOG OF THE PLANT WORLD



# UNKNOWN WILD RELATIVE

**BRUSSELS SPROUTS**



**BROCCOLI**



**CABBAGE**



**CAULIFLOWER**



**KALE**



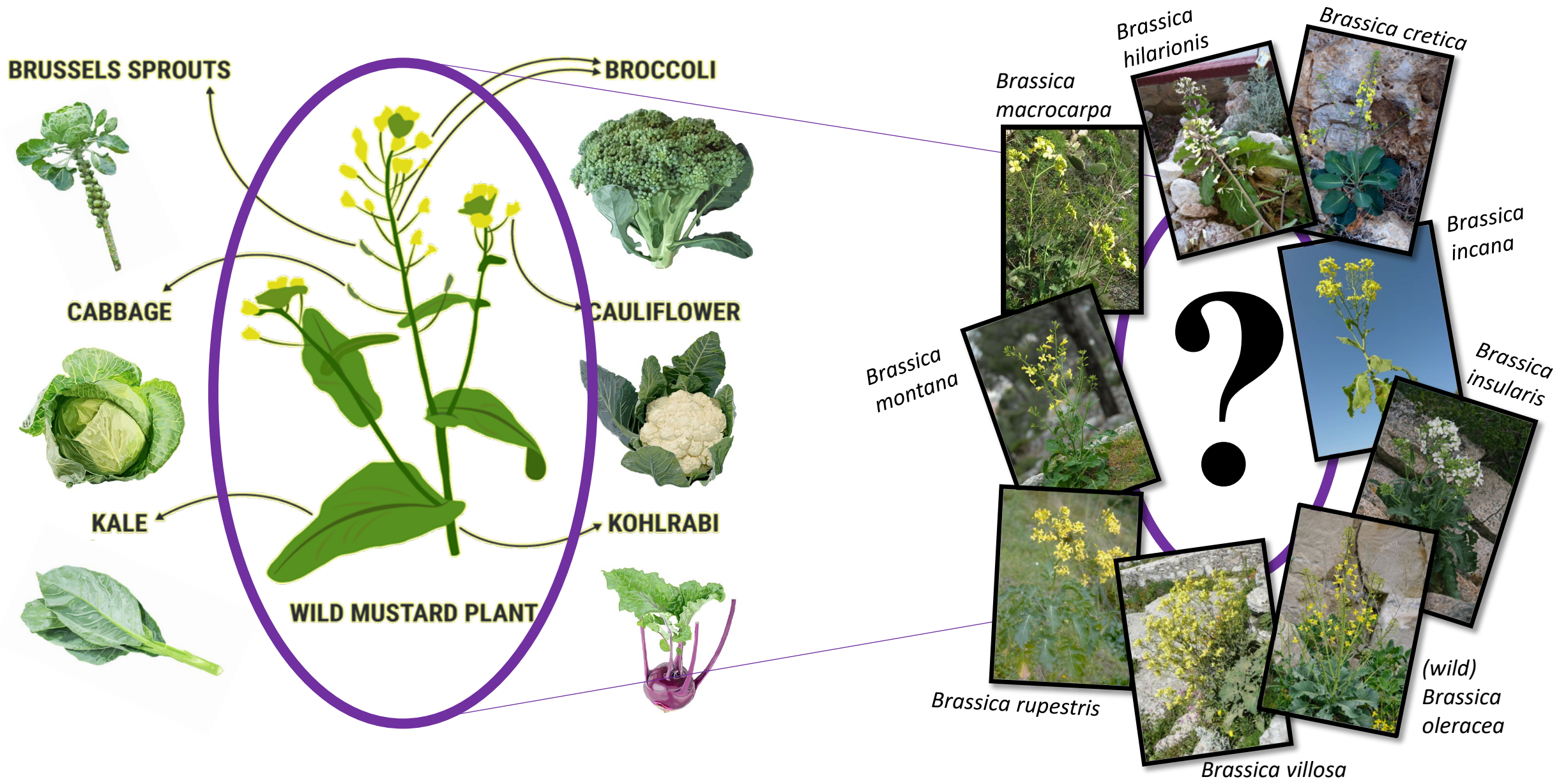
**WILD MUSTARD PLANT**



**KOHLRABI**



# UNKNOWN WILD RELATIVE



# WILD BRASSICAS – *BRASSICA OLERACEA* RELATIVES

8 species with the CC genome



- *Brassica assyriaca*
- *Brassica aucheri*
- *Brassica baldensis*
- *Brassica balearica*
- *Brassica barrelieri*
- *Brassica beytepeensis*
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- *Brassica cadmea*
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- *Brassica souliei*
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- *Brassica taiwanensis*
- *Brassica taurica*
- *Brassica trichocarpa*
- *Brassica tyrrhena*
- ***Brassica villosa***

# UNKNOWN DOMESTICATION ORIGIN

## 1. Italy/Sicily

-Center of diversity

(*B. macrocarpa*, *B. incana*, *B. rupestris*, *B. insularis*, *B. villosa*)



- Brassica cretica*
- Brassica hilarianis*
- Brassica incana*
- Brassica insularis*
- Brassica macrocarpa*
- Brassica montana*
- Brassica oleracea*
- Brassica rupestris*
- Brassica villosa*



# UNKNOWN DOMESTICATION ORIGIN

## 1. Italy/Sicily

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(*B. macrocarpa*, *B. incana*, *B. rupestris*, *B. insularis*, *B. villosa*)

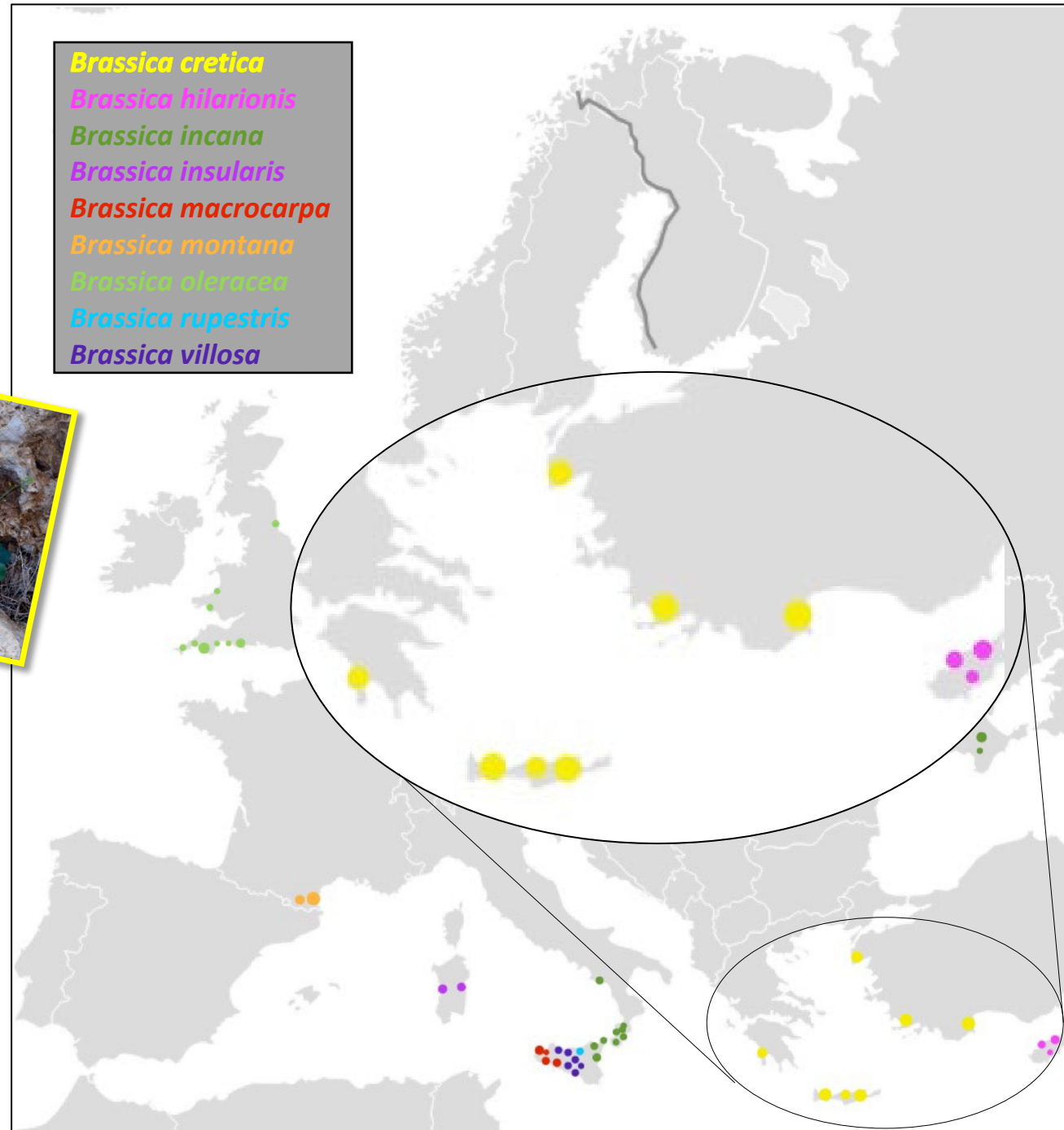


## 2. Eastern Mediterranean

-Linguistics and literary

(*B. cretica* and *B. hilarionis*)

*Brassica cretica*  
*Brassica hilarionis*  
*Brassica incana*  
*Brassica insularis*  
*Brassica macrocarpa*  
*Brassica montana*  
*Brassica oleracea*  
*Brassica rupestris*  
*Brassica villosa*



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## 2. Eastern Mediterranean

-Linguistics and literary

(*B. cretica* and *B. hilarionis*)

## 3. England

- Wild cabbage

(wild *B. oleracea*)



## MAIN QUESTION

- Which wild relative is most closely related to *Brassica oleracea* and where is the center of domestication?



- Samples from USDA, IPK, UPM, and John Innes Center
- 225 samples across wild relatives and diversity of *B. oleracea* cultivars

## DATA COLLECTION

- Wild *B. oleracea* - 3
- *B. cretica* - 3
- *B. incana* - 5
- *B. montana* - 2
- *B. hilarionis* - 1
- *B. insularis* - 5
- *B. macrocarpa* - 4
- *B. rupestris* - 7
- *B. villosa* - 5



- Genomic data
  - RNA-seq data

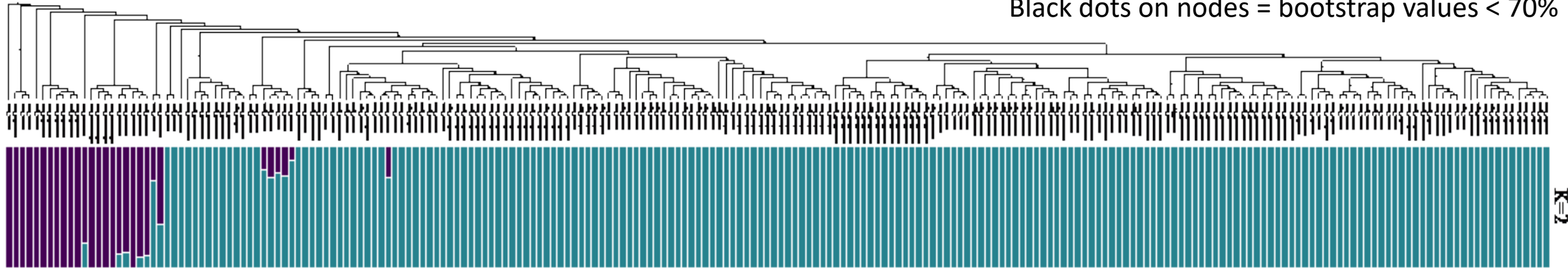
- Phenotypic data
  - Leaf scans
  - Glucosinolates
  - Ionomics

Black dots on nodes = bootstrap values < 70%



- Many cultivars are recovered as monophyletic.
  - Some, such as Kales, are not.

Black dots on nodes = bootstrap values < 70%



Wild C      wild C - clade 2      Chinese white kale      broccoli      cauliflower      kohlrabi      curly kale      Brussels sprouts      collards      cabbage



Example

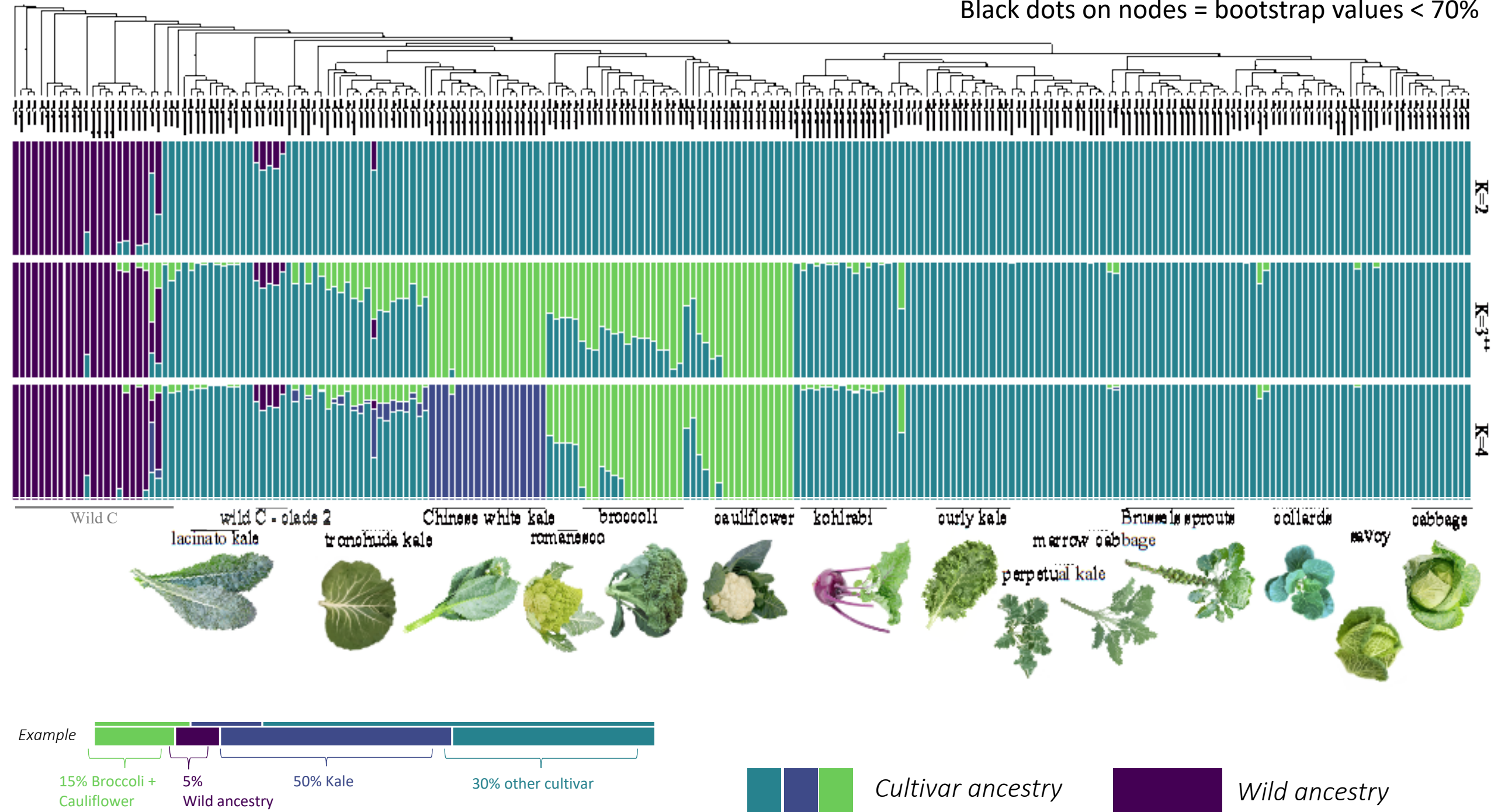


25% Wild ancestry

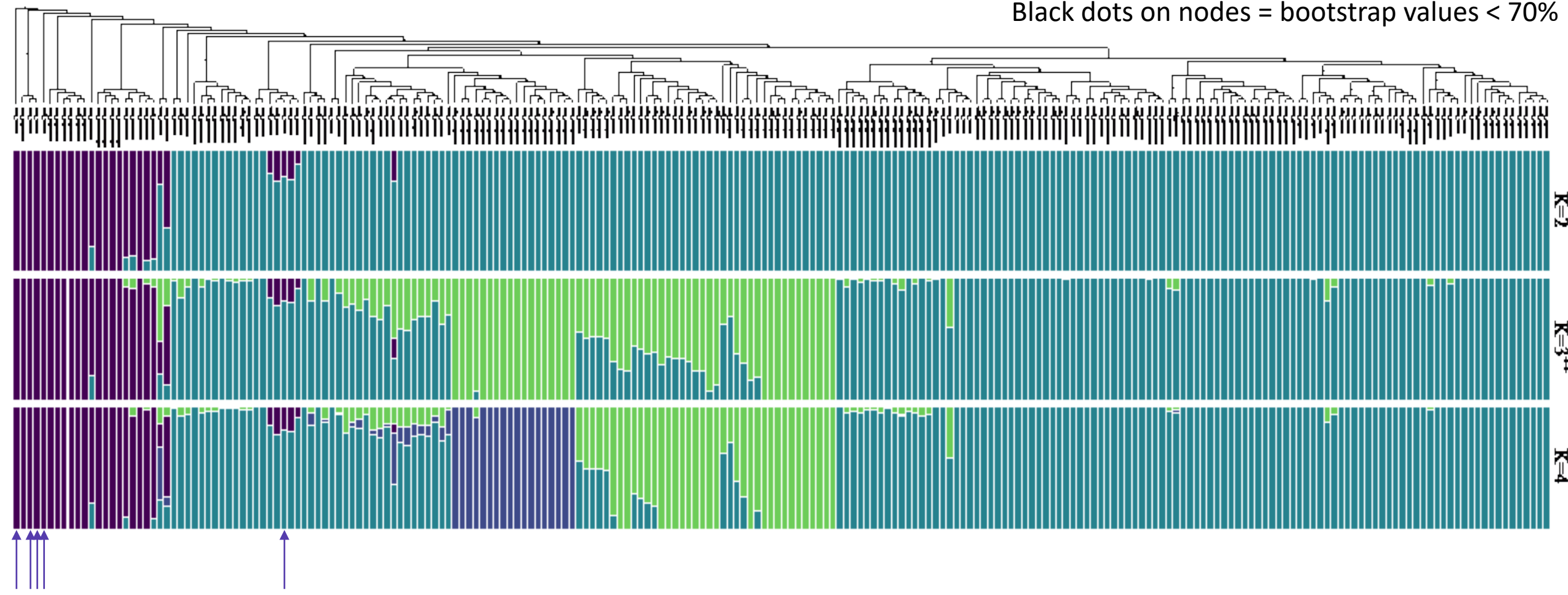
75% Cultivar ancestry

 Cultivar ancestry       Wild ancestry

Black dots on nodes = bootstrap values < 70%



Black dots on nodes = bootstrap values < 70%



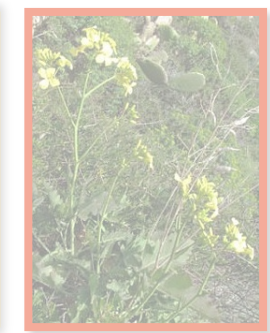
*B. villosa*



*B. rupestris*



*B. montana*



*B. macrocarpa*



*B. insularis*



*B. hilarionis*



*B. incana*

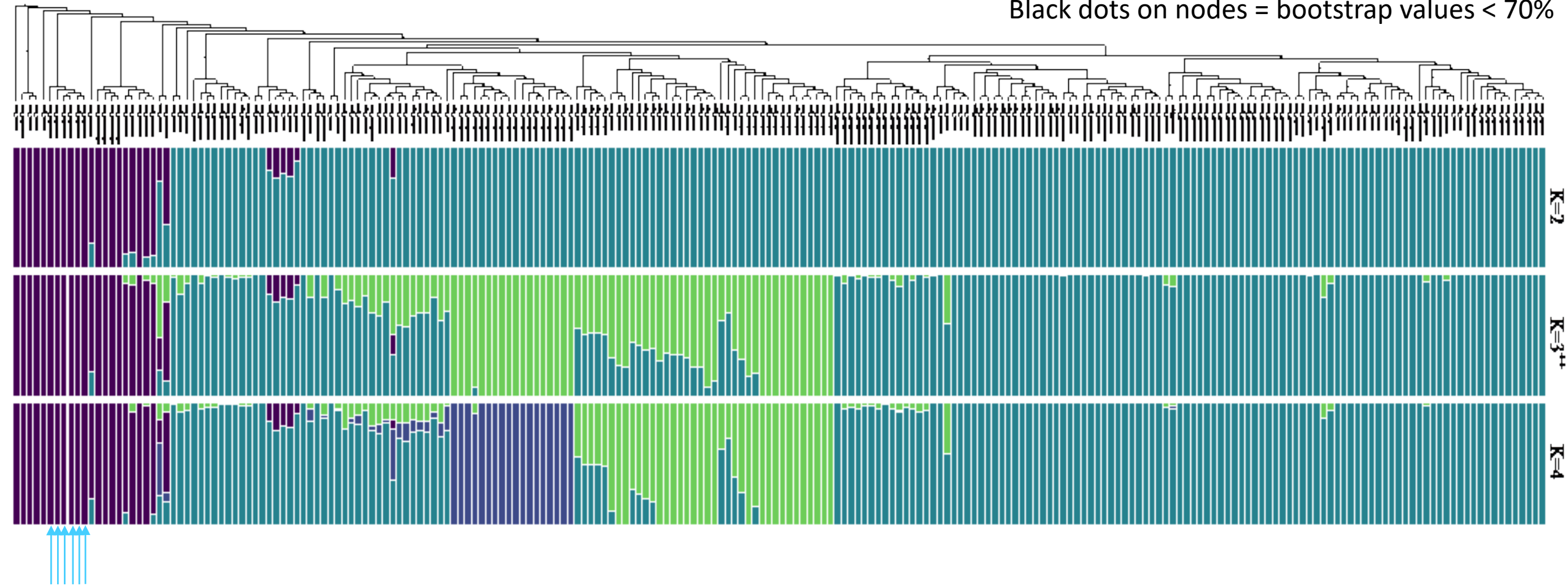


*B. cretica*



*B. oleracea*

Black dots on nodes = bootstrap values < 70%



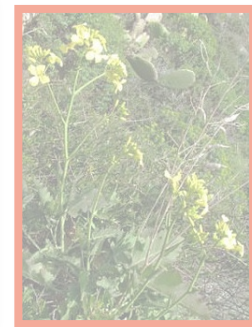
*B. villosa*



*B. rupestris*



*B. montana*



*B. macrocarpa*



*B. insularis*



*B. hilarionis*



*B. incana*

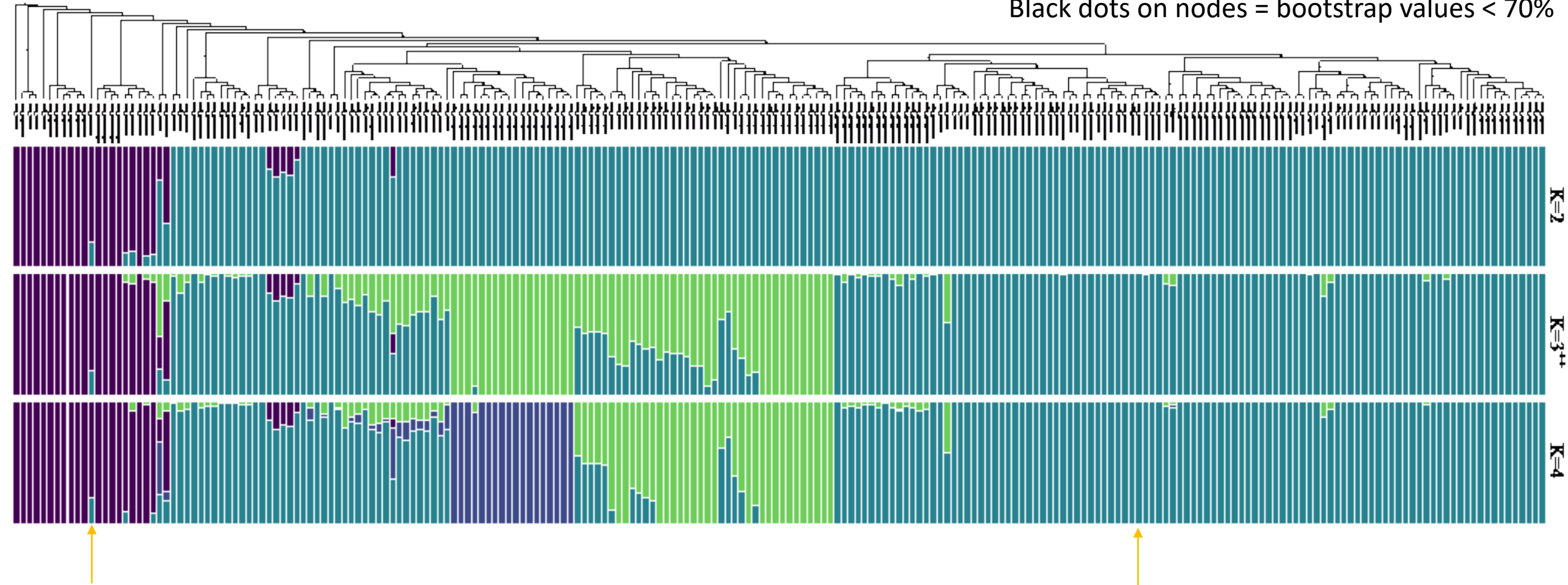


*B. cretica*



*B. oleracea*

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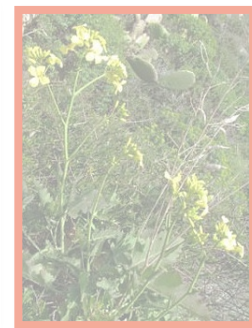
*B. villosa*



*B. rupestris*



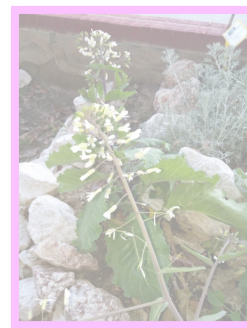
*B. montana*



*B. macrocarpa*



*B. insularis*



*B. hilarionis*



*B. incana*

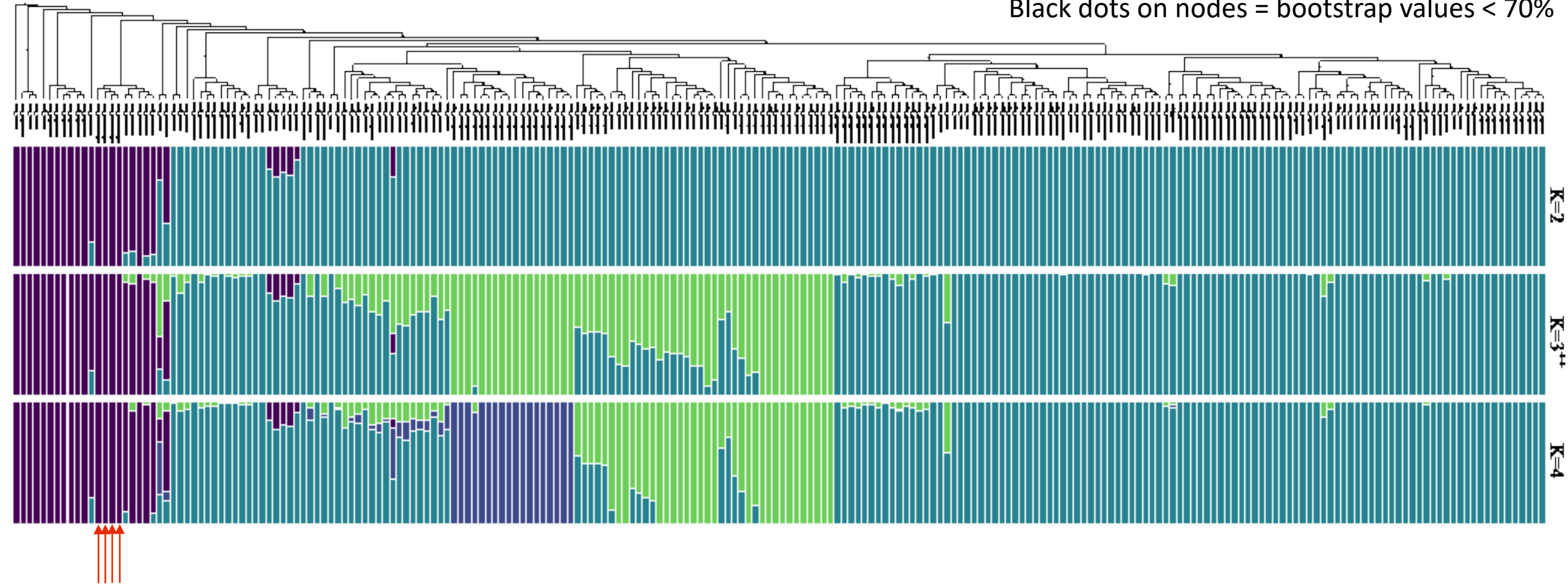


*B. cretica*



*B. oleracea*

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*B. villosa*



*B. rupestris*



*B. montana*



*B. macrocarpa*



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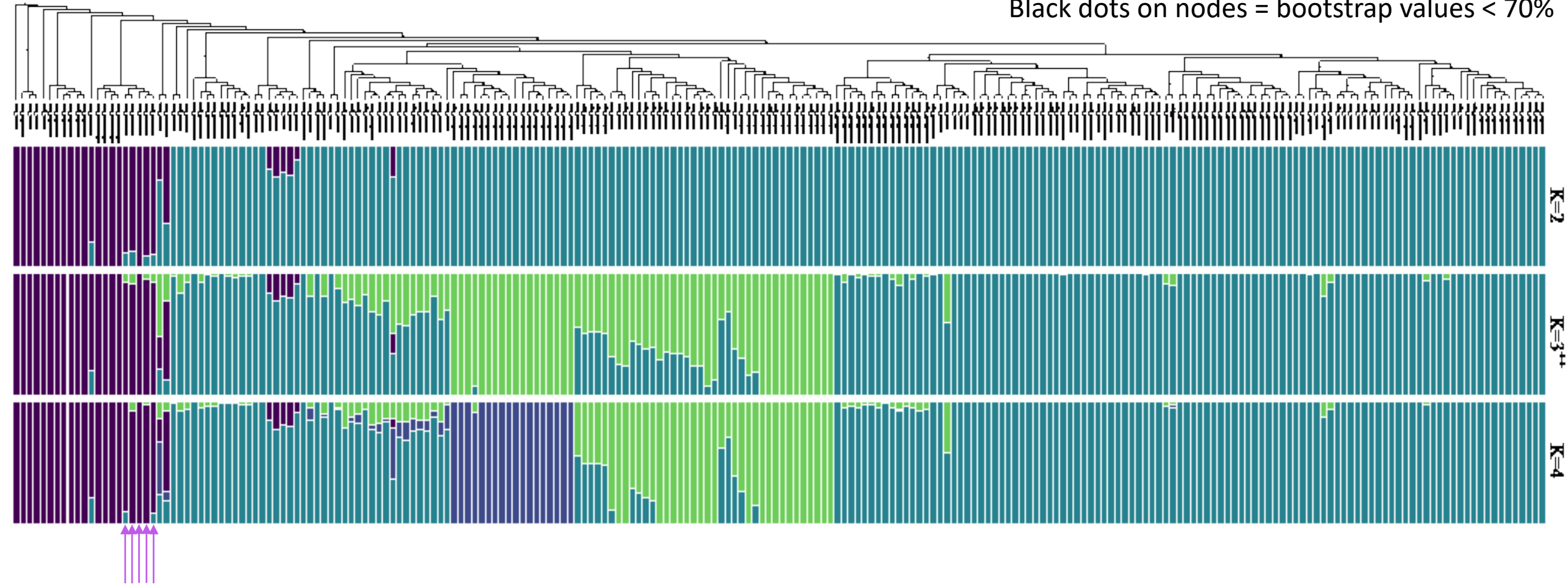


*B. cretica*



*B. oleracea*

Black dots on nodes = bootstrap values < 70%



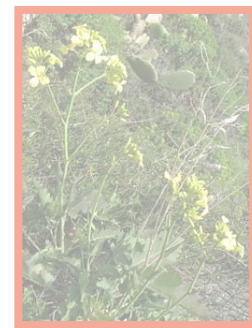
*B. villosa*



*B. rupestris*



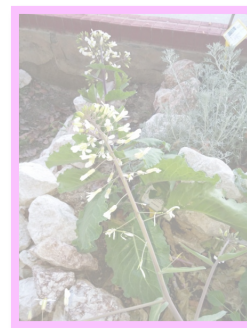
*B. montana*



*B. macrocarpa*



*B. insularis*



*B. hilarionis*



*B. incana*

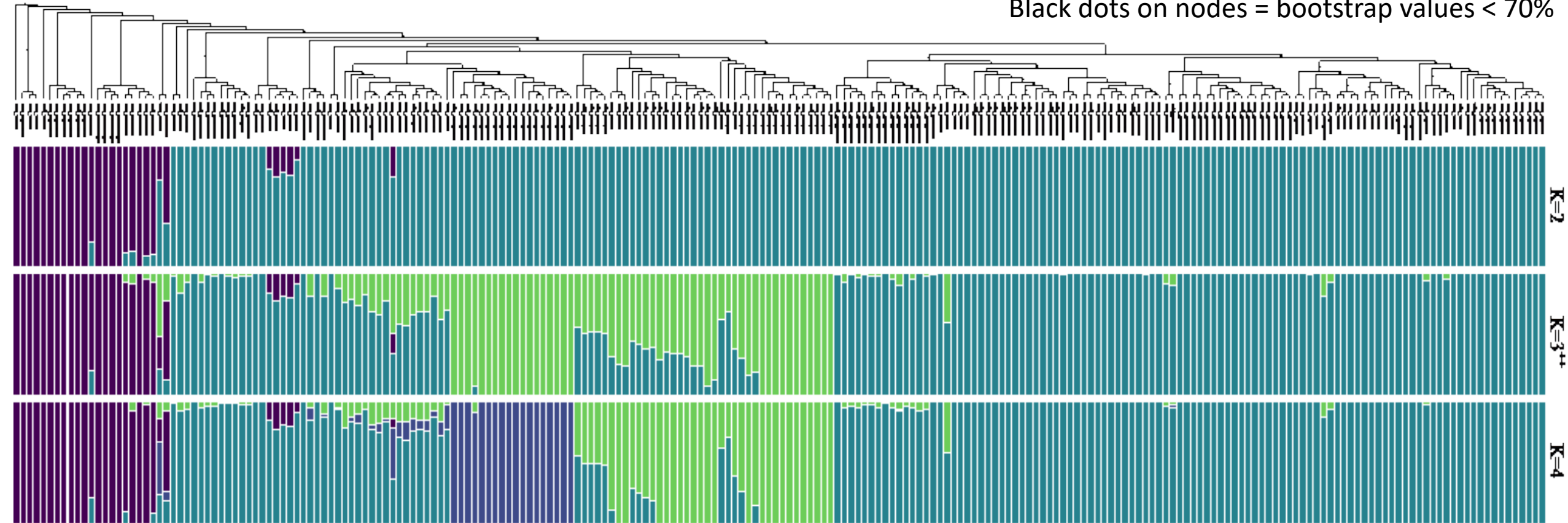


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*B. macrocarpa*



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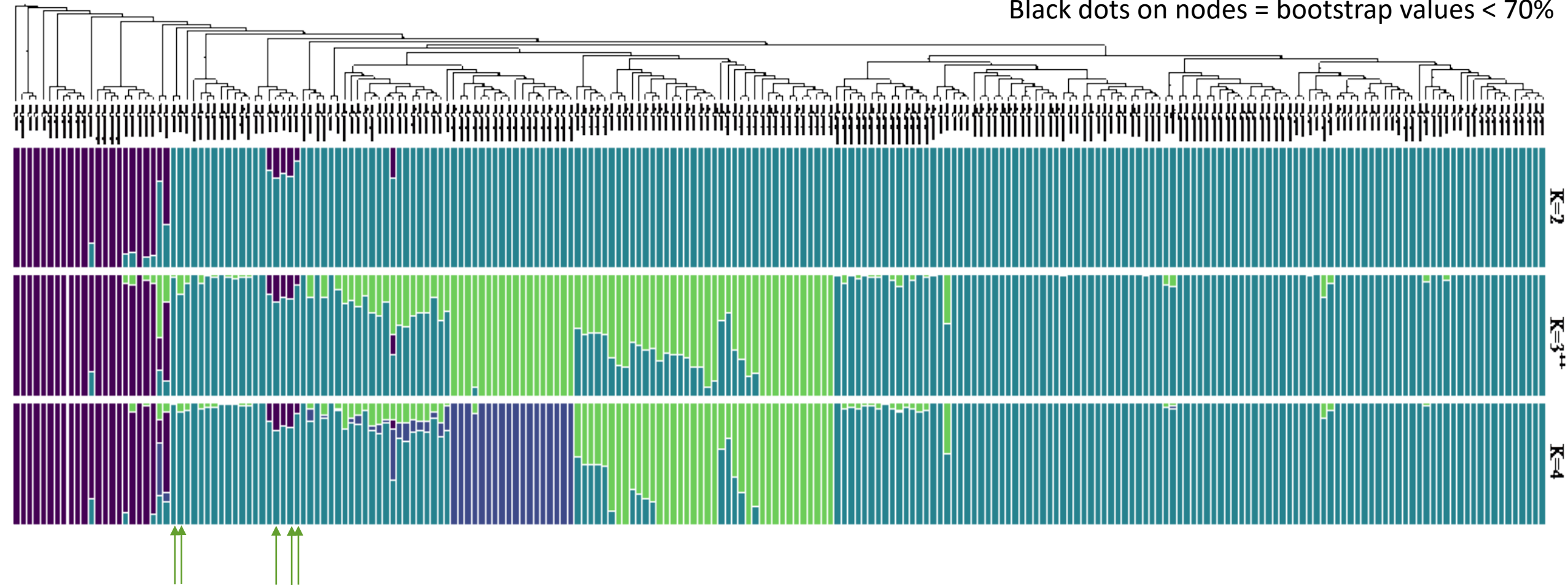


*B. cretica*



*B. oleracea*

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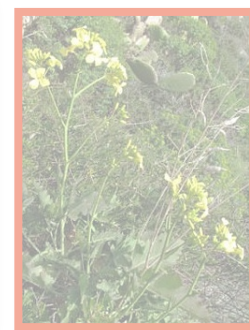
*B. villosa*



*B. rupestris*



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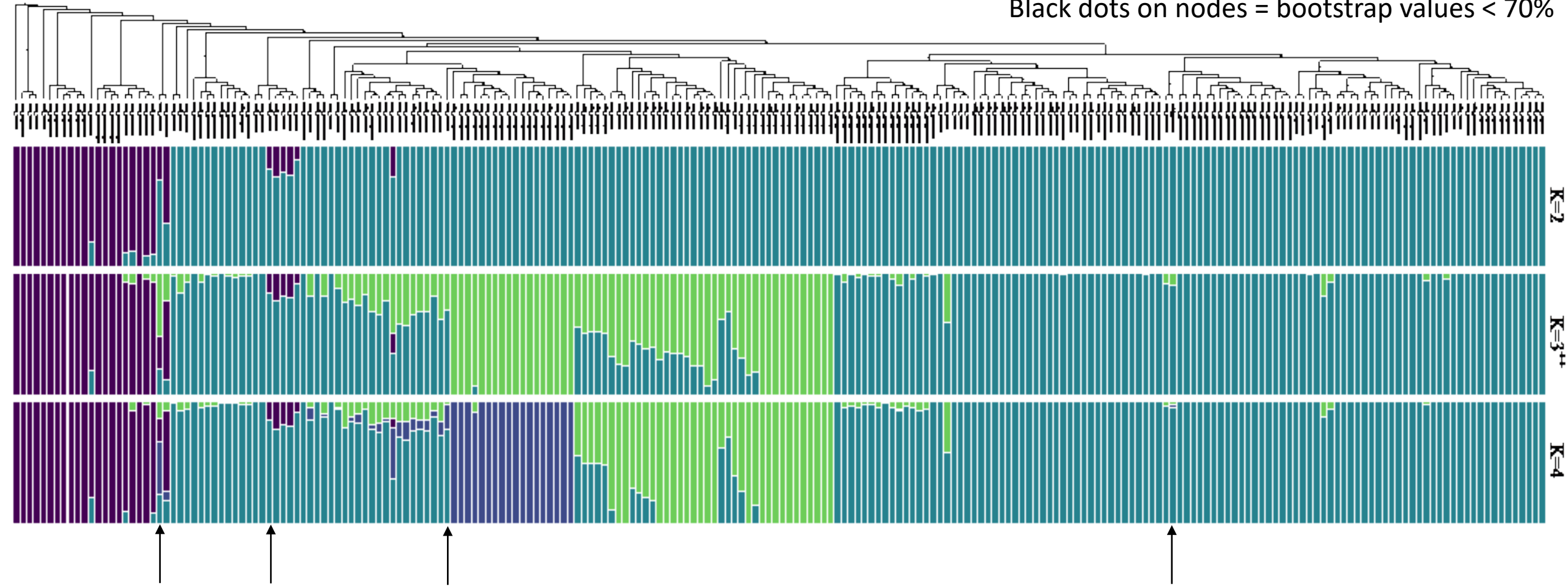


*B. cretica*

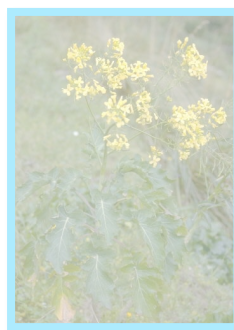


*B. oleracea*

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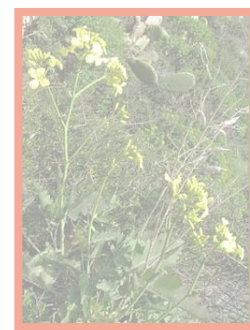
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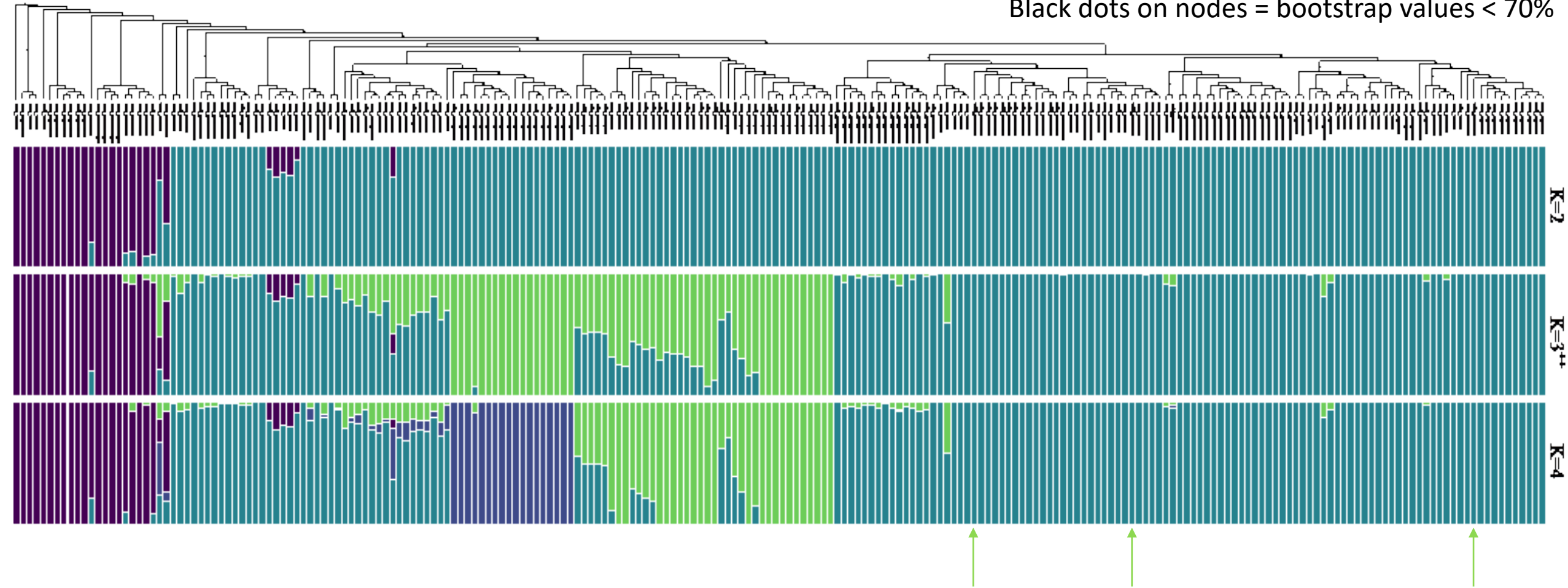


*B. cretica*

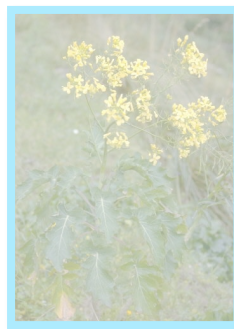


*B. oleracea*

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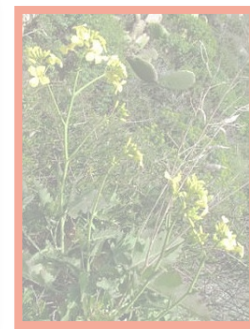
*B. villosa*



*B. rupestris*



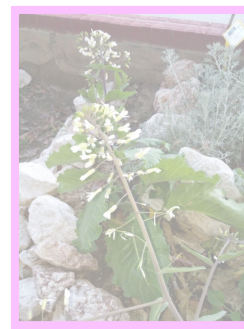
*B. montana*



*B. macrocarpa*



*B. insularis*



*B. hilarionis*



*B. incana*

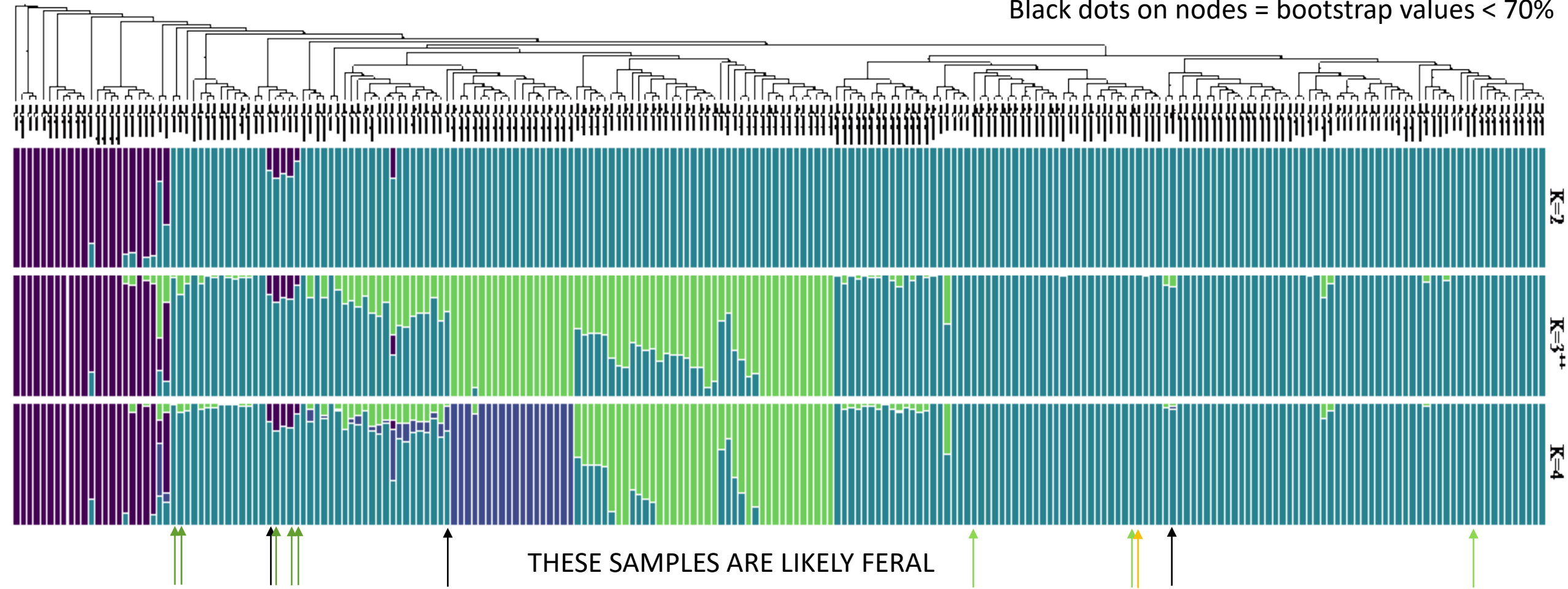


*B. cretica*



*B. oleracea*

Black dots on nodes = bootstrap values < 70%



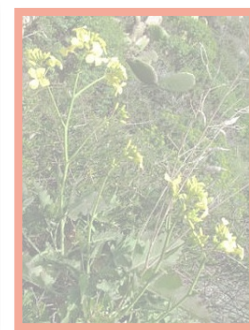
*B. villosa*



*B. rupestris*



*B. montana*



*B. macrocarpa*



*B. insularis*



*B. hilarionis*



*B. incana*

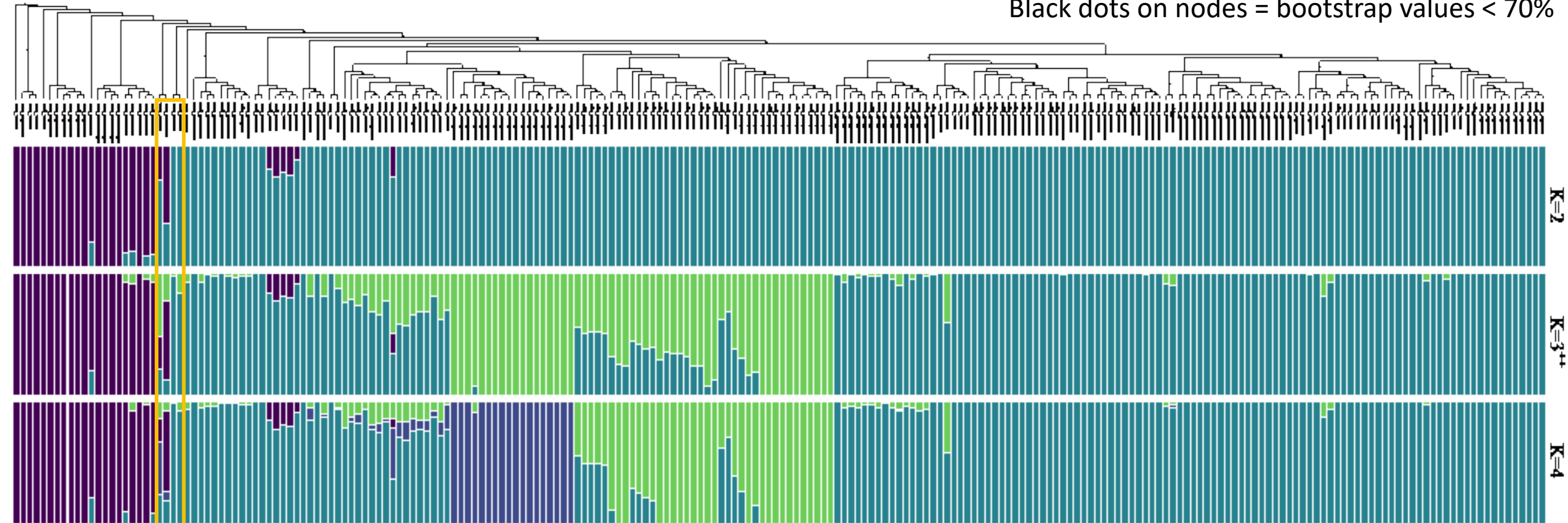


*B. cretica*

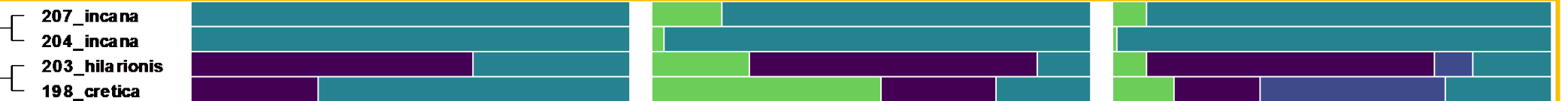


*B. oleracea*

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Closest living wild relatives... but....



- *B. incana* 100% shared ancestry with cultivars
- *B. hilarionis* ~60% shared ancestry with cultivars + ancestry from all groups
- *B. cretica* ~30% shared ancestry with cultivars + ancestry from all groups

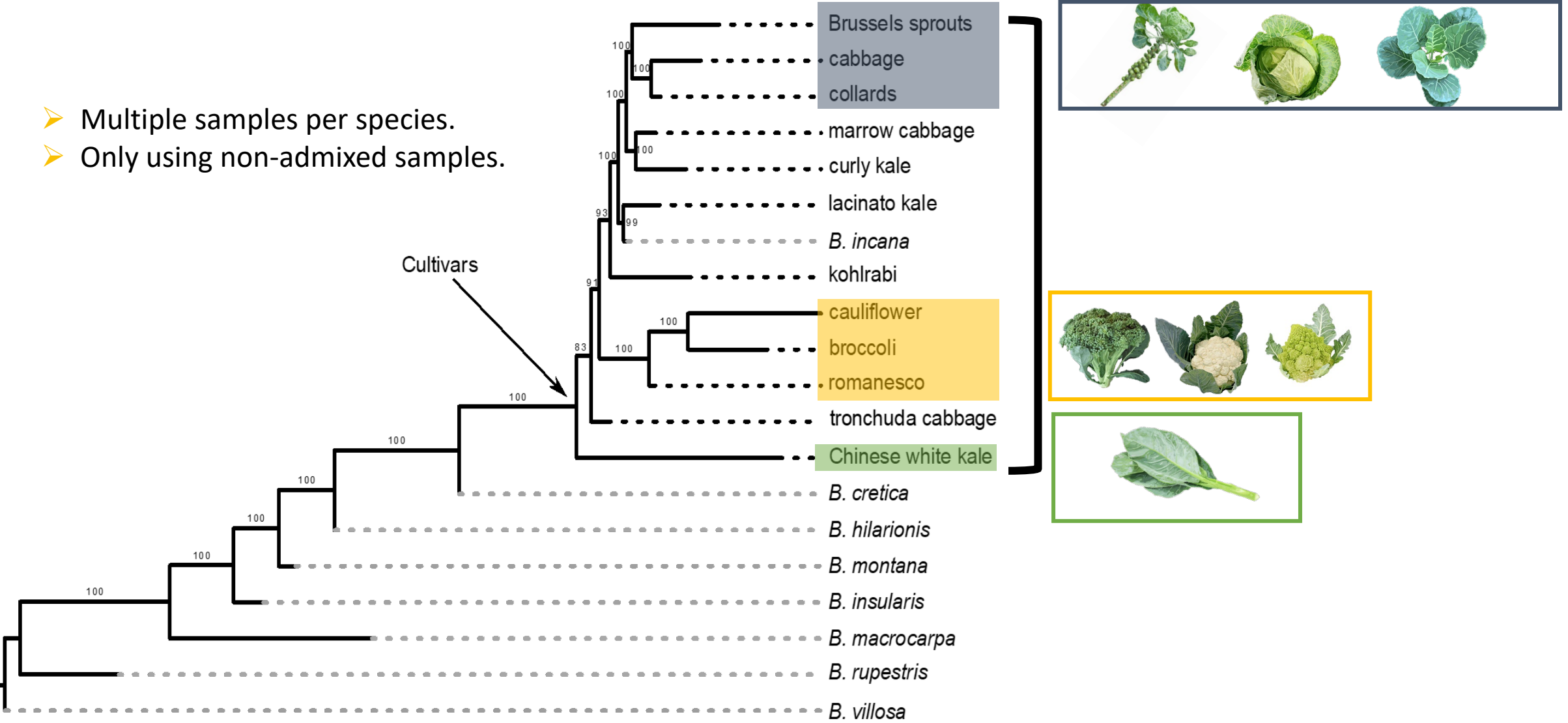


➤ Wild C – clade 2 shares ancestry with cultivars and wild taxa



# SPECIES PHYLOGENY INDICATES *BRASSICA CRETICA* AS CLOSEST LIVING WILD RELATIVE

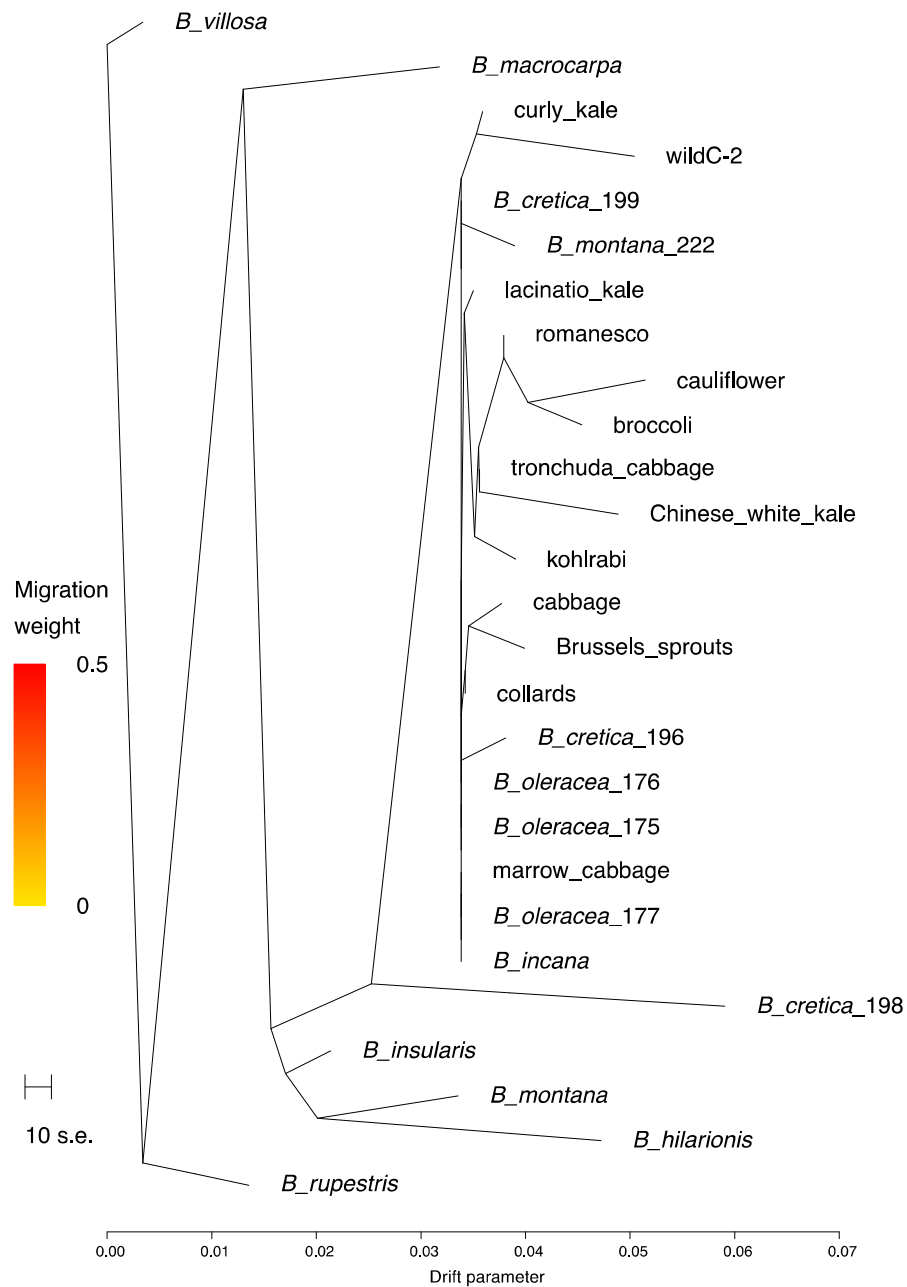
- Multiple samples per species.
- Only using non-admixed samples.



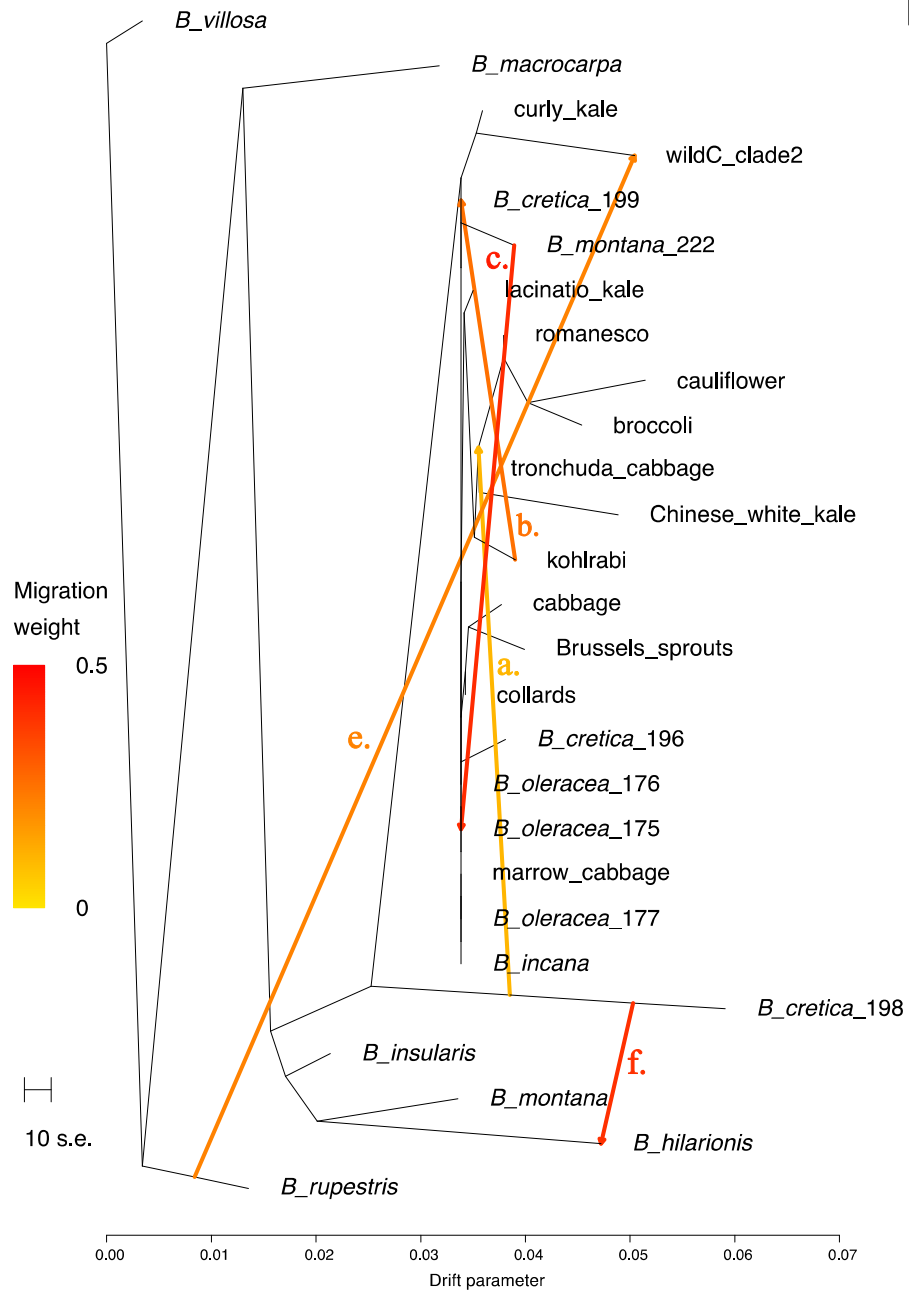


# FERAL ORIGINS

➤ Without migration = 87.3% of the variance in the dataset



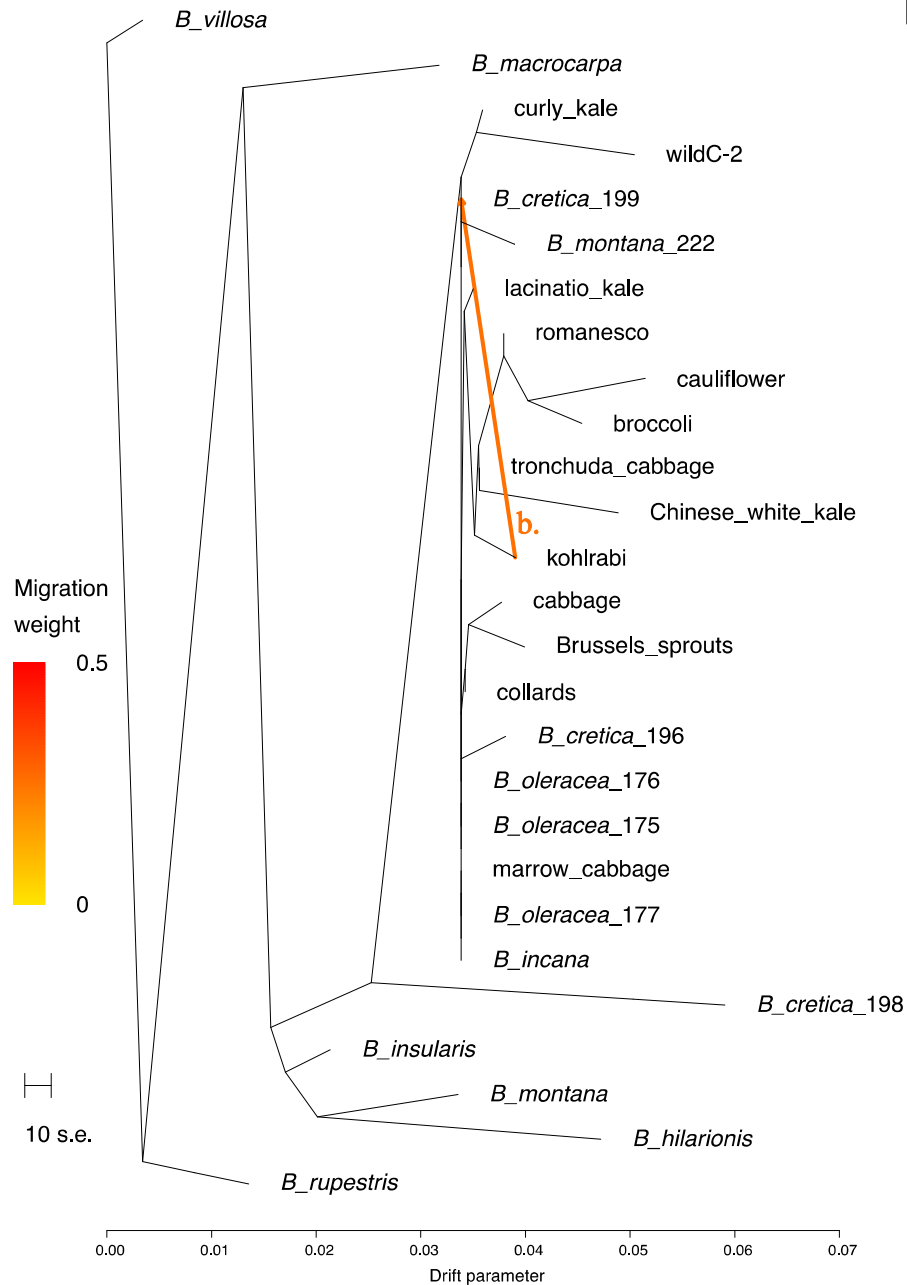
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➤ Without migration = 87.3% of the variance in the dataset

➤ Five migrations = 92% of the variance in the dataset

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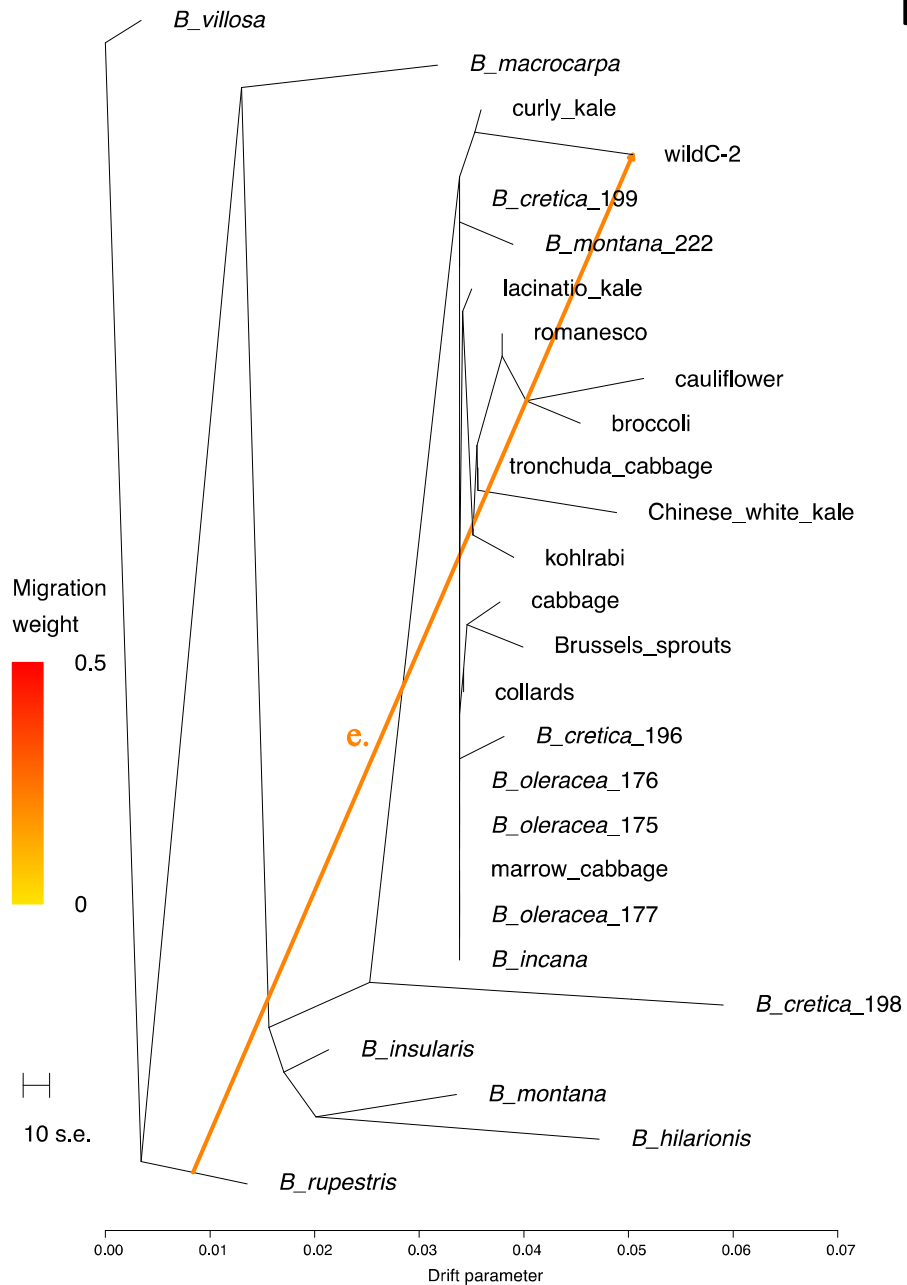


➤ Without migration = 87.3% of the variance in the dataset

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➤ b. = migration from kohlrabi to *B. cretica* 199,  
➤ seen in swollen stem phenotype when growing the plants out.

# FERAL ORIGINS



- Without migration = 87.3% of the variance in the dataset
- Five migrations = 92% of the variance in the dataset
- b. = migration from kohlrabi to *B. cretica* 199,
  - seen in swollen stem phenotype when growing the plants out.
- e. = migration from *B. rupestris* to wild C - clade 2
  - Explains admixture result
  - Perpetual kale (*B. oleracea* var. *ramosa*)



# ARCHAEOLOGICAL EVIDENCE FOR *B. OLERACEA* DOMESTICATION



- Oldest
  - ca. 3550-3350 BP Middle Bronze Age
  - 3 charred seeds on a hilltop settlement in southern alps (Schmidl and Oeggel 2005)

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- Difficulty in telling *Brassica* seeds apart from one another.
  - Many suggest these ancient finds to be that of *B. cretica* (van der Veen 2001; Kaniewski et al. 2011)

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    - 3 charred seeds on a hilltop settlement in southern alps (Schmidl and Oeggel 2005)
- Difficulty in telling *Brassica* seeds apart from one another.
  - Many suggest these ancient finds to be that of *B. cretica* (van der Veen 2001; Kaniewski et al. 2011)
- Other than seeds
  - Residues of Brassica leaves identified within pottery shards (Evershed et al. 1992)
    - Early Medieval (12th century)
    - England



# CLASSICAL AND MEDIEVAL WRITTEN SOURCES FOR *B. OLERACEA* DOMESTICATION



- Earliest textual reference
  - 2500-2000 BP
  - Hipponax
  - Greece
- Refers to a "seven-leaved" cabbage in iambic verse
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- 3 varieties
  - smooth seedless
  - larger leafed
  - sweeter parsley/curly-leaved
- Refers to a wild type with bitter taste, many branches, and smaller leaves may be *B. cretica*

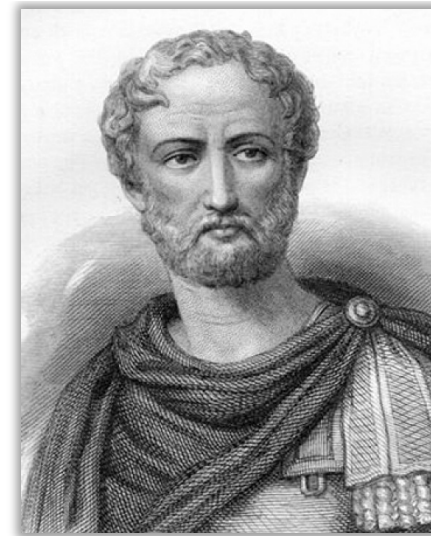
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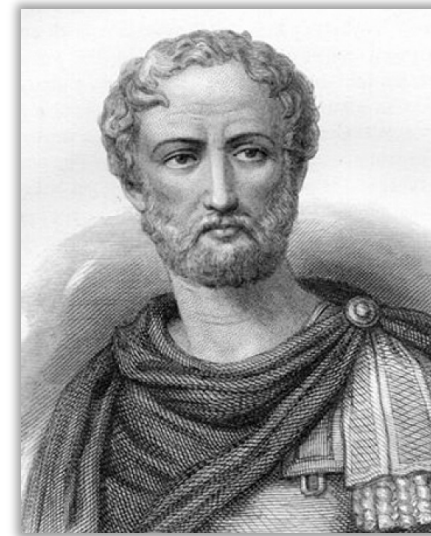


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- Modern-day distribution of *B. cretica* includes south-east Greece and the Aegean Islands
- *B. incana* and *B. oleracea* are not known from Greece
- Therefore, these earliest types may belong to *B. cretica* which is incredibly diverse itself



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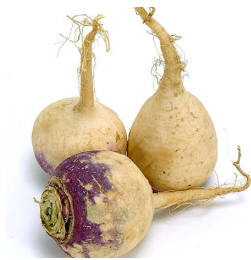


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# THE ROLE OF CWRs IN DOMESTICATION

## 1 – *Brassica oleracea*

## 2 – *Brassica rapa*

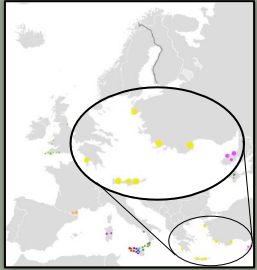


## *Brassica napus*



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- Taking all data together, genetic, archeology, and literature, *B. cretica*, is best supported as the closet living wild relative
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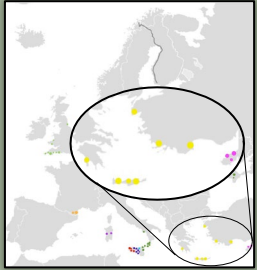


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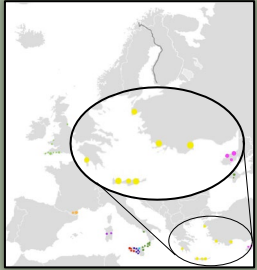


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## *Brassica napus*



# BRASSICA RAPA DOMESTICATION: UNTANGLING WILD AND FERAL FORMS AND CONVERGENCE OF CROP MORPHOTYPES

*Molecular Biology and Evolution*, Volume 38, Issue 8, August 2021, Pages 3358–3372



Alex McAlvay



Aaron Ragsdale



Makenzie Mabry



Xinshuai Qi



Kevin Bird



Pablo Velasco



Hong An



J. Chris Pires



Eve Emshwiller



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paper

## MAIN QUESTIONS



- Is weedy *B. rapa* wild or feral?
- What is the center of domestication of *B. rapa*?

# DATA COLLECTION – GENOTYPING BY SEQUENCING (68,468 SNPs)



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*384 crop samples*

Grelos



Rapini



Pak choi



Komatsuna



Choy sum



Turnips



Mizuna



Napa cabbage



Wutacai



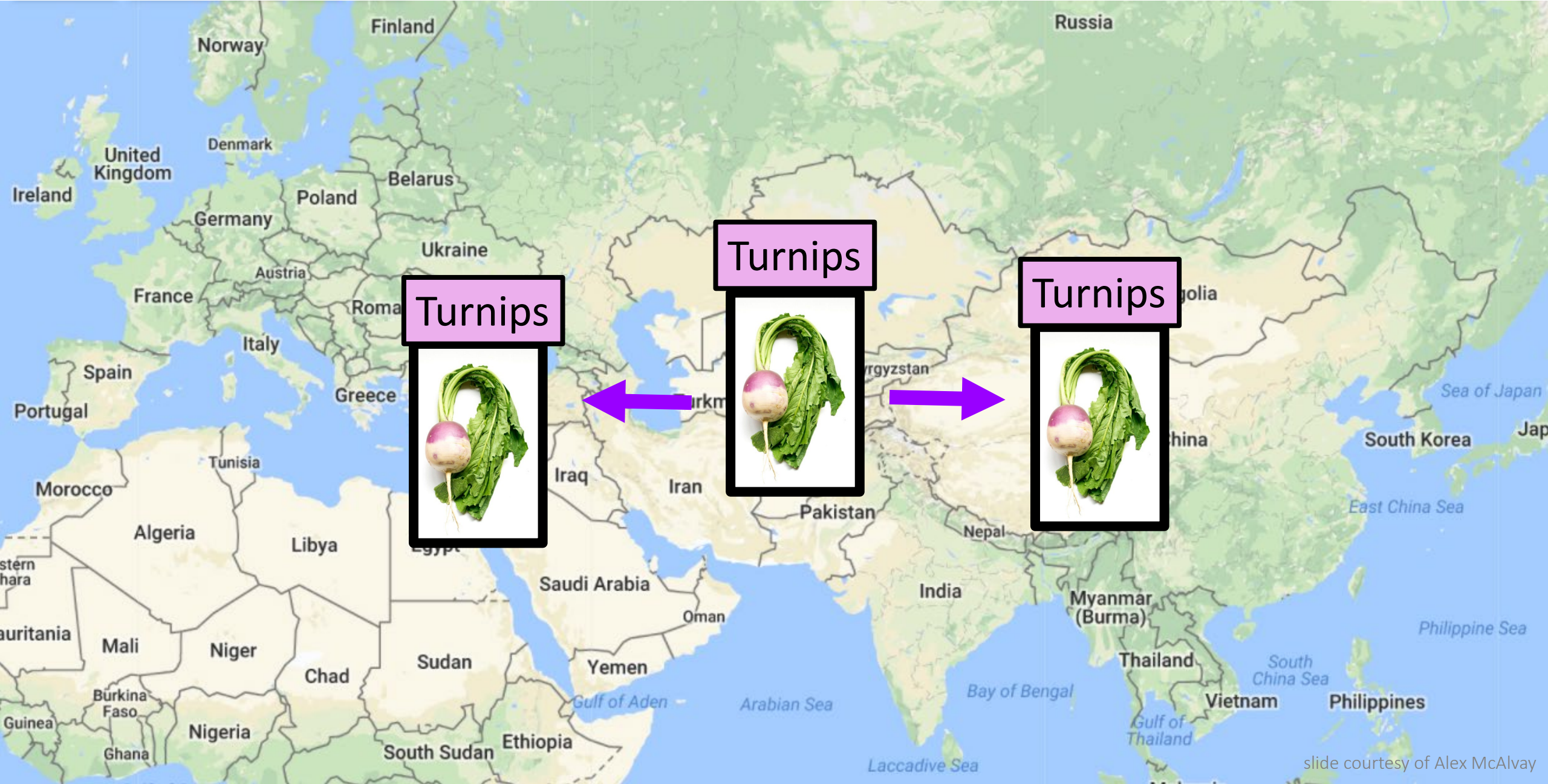
Brown sarson



Yellow sarson

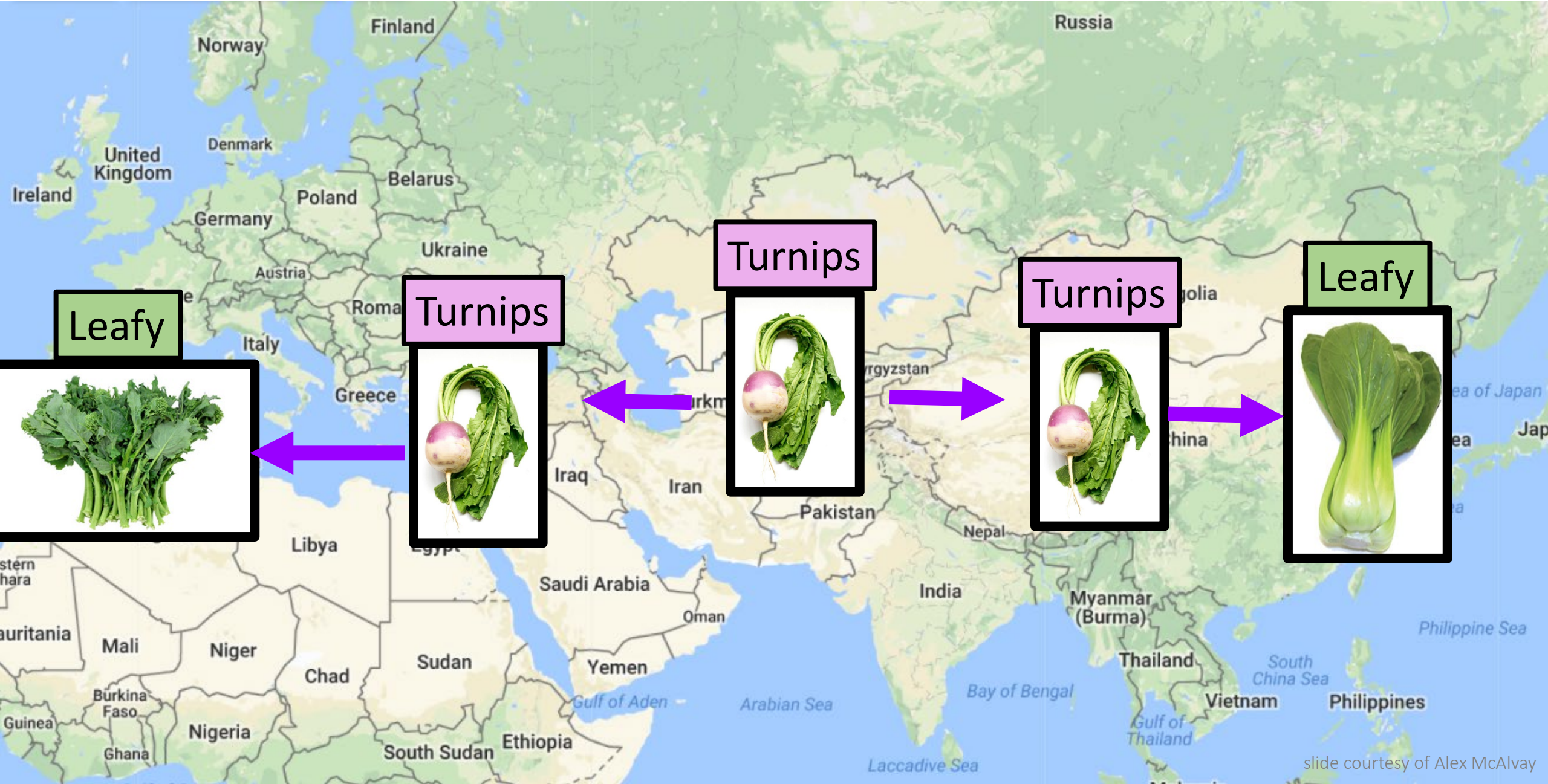


# INDEPENDENT SELECTION FOR LEAFY TYPES FROM TURNIPS



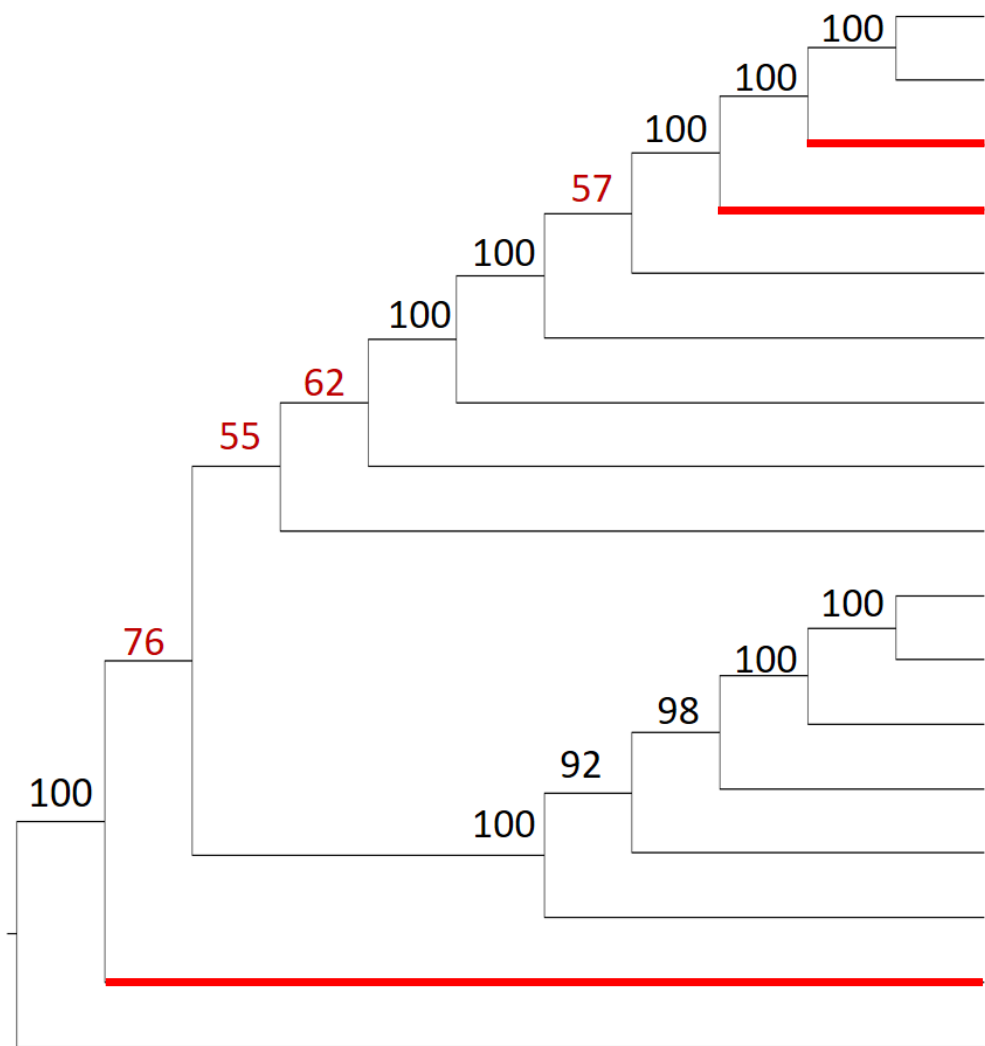
slide courtesy of Alex McAlvay

# INDEPENDENT SELECTION FOR LEAFY TYPES FROM TURNIPS



# MULTIPLE ORIGINS OF WEEDINESS /FERALS

*Originated from different crops*



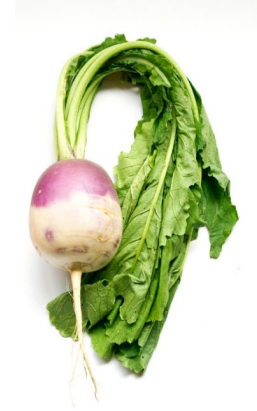
**European crops**

**Weedy Americas**  
**Weedy Europe**

**European crops**

**East and South Asian crops**

**Weedy Caucasus**  
**Outgroup (*B. oleracea*)**



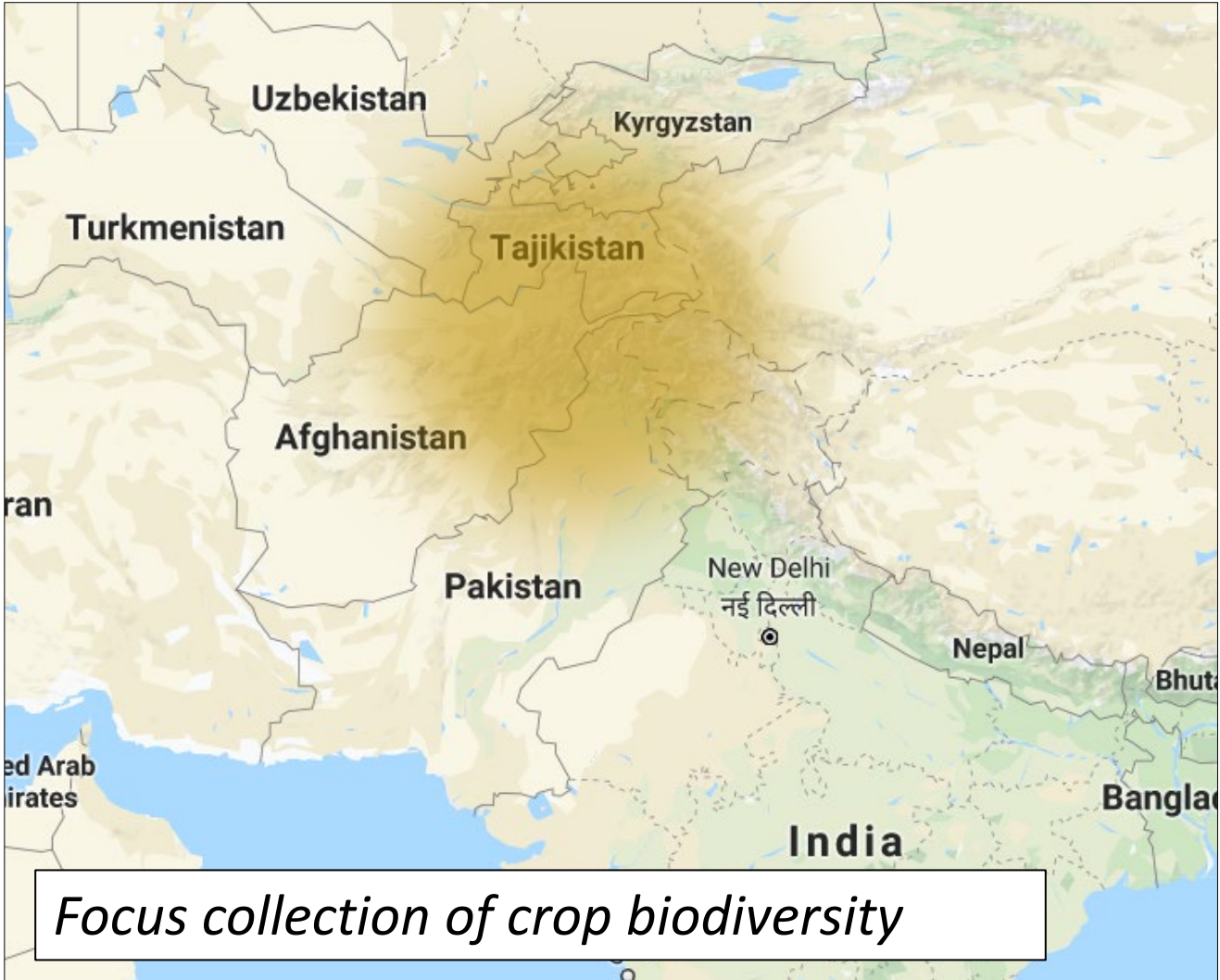
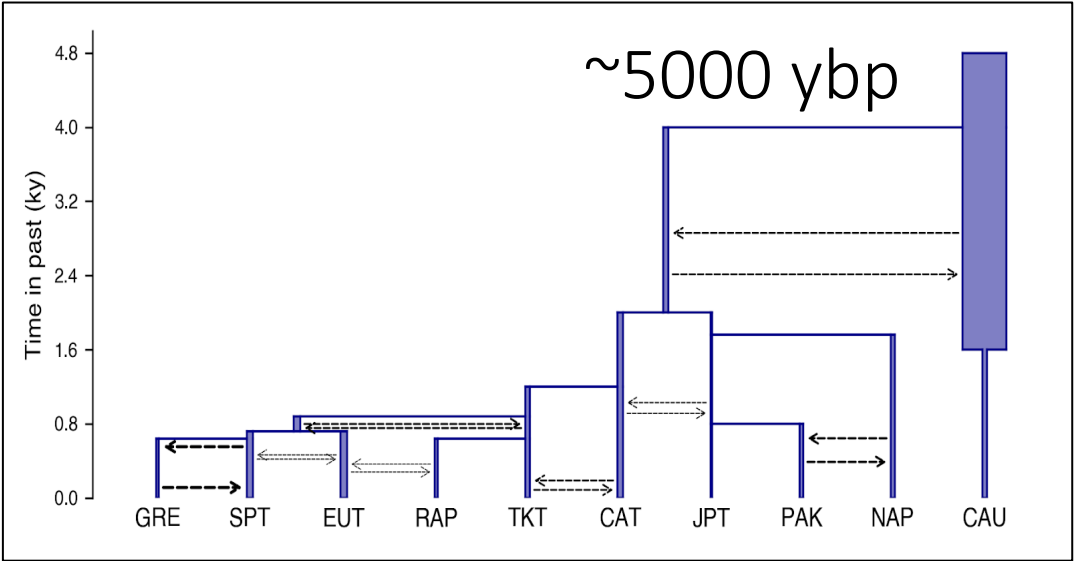
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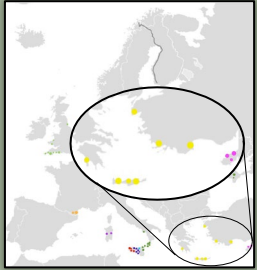


# DEMOGRAPHIC MODEL SUPPORTS CENTRAL ASIAN ORIGIN



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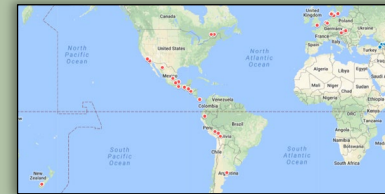
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## 2 – *Brassica rapa*



- Samples from the Caucasus may be truly wild.
- Weedy *B. rapa* occurring in the Americas and much of Europe are feral.



- Demographic model supports central Asian origin.

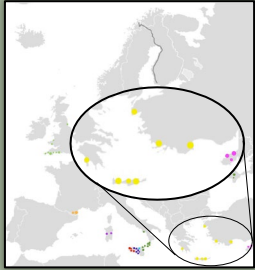


## *Brassica napus*



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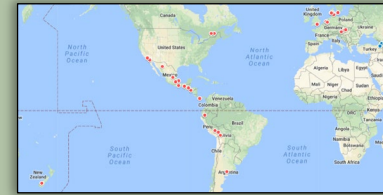
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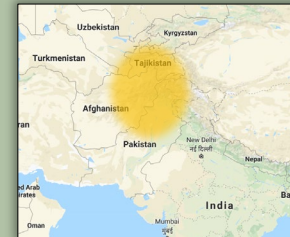
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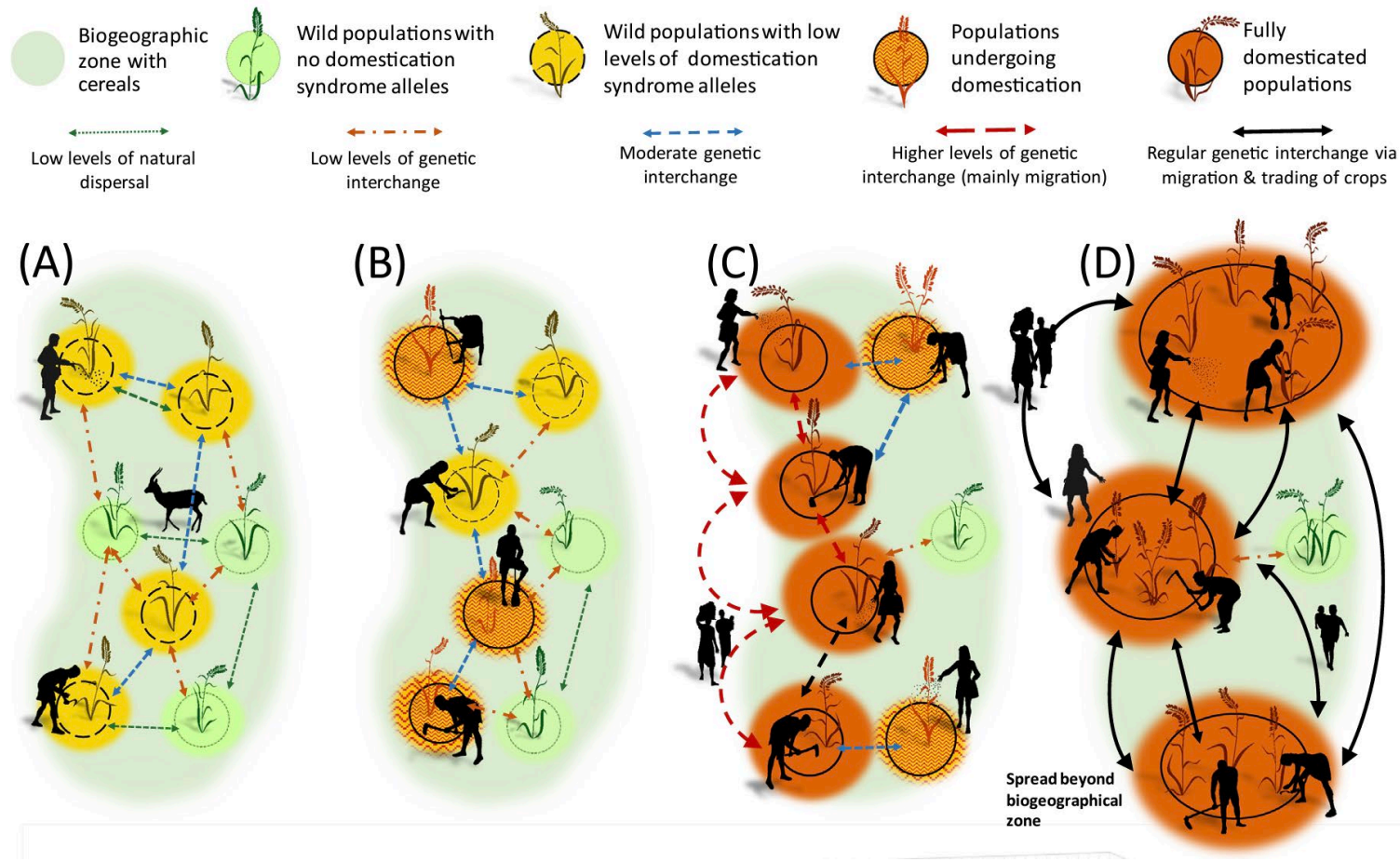
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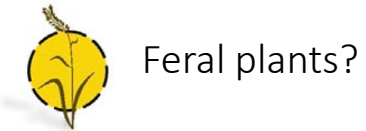
- What could the role of these CWRs be for *B. napus*?
  - Recent work highlights that there is a single origin, but lots of admixture (An et al. 2019, Wang et al. 2023).
- Knowledge of *B. oleracea* and *B. rapa* domestication histories (including the role of feral populations) will help inform our basic and applied understanding of *B. napus*.

# A FERAL FUTURE

*Allaby et al. 2022 – Emerging evidence of plant domestication as a landscape-level process*

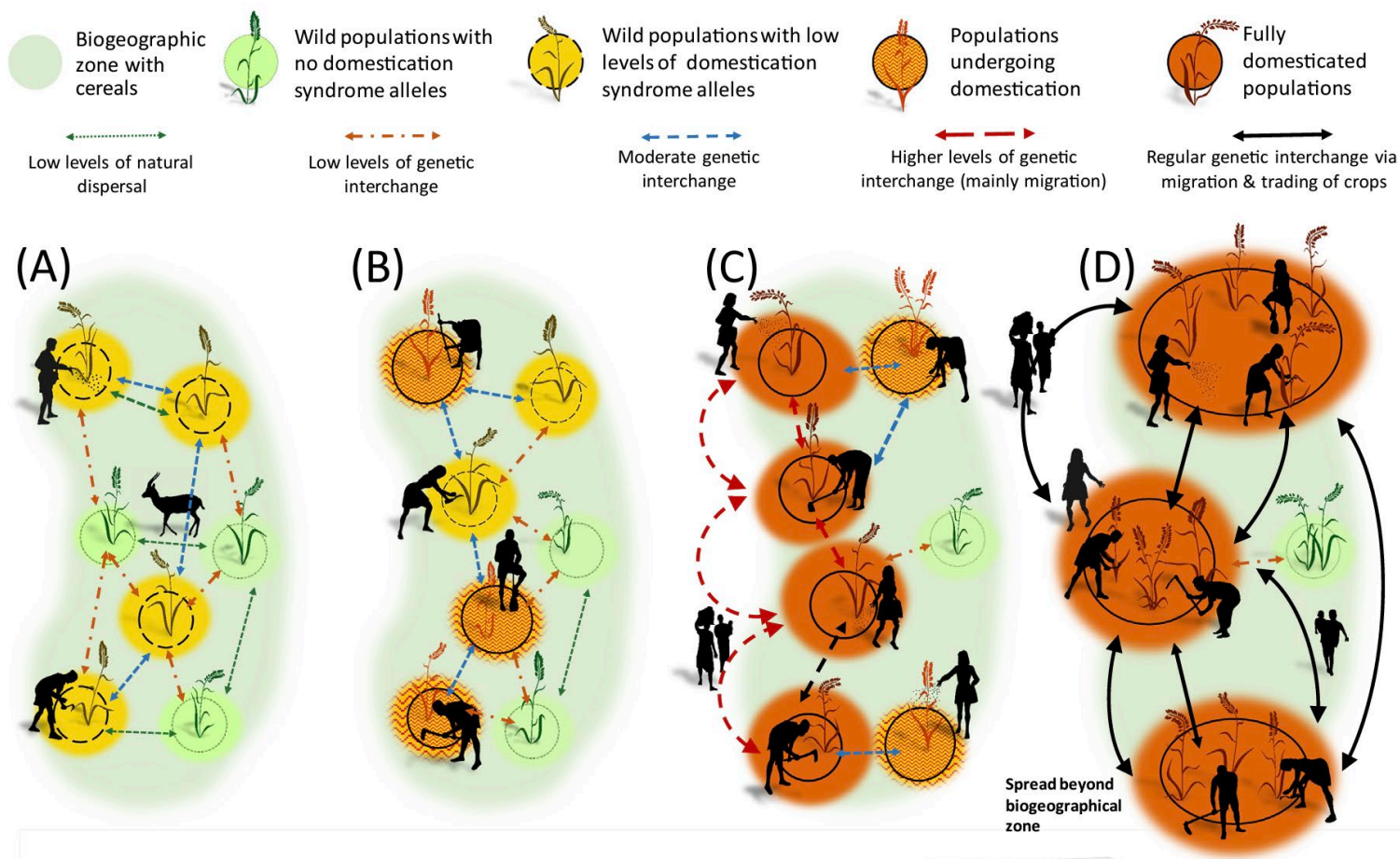


➤ Feral plants always a part of the story in introducing variability during domestication!

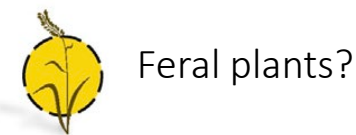


# A FERAL FUTURE

Allaby et al. 2022 – Emerging evidence of plant domestication as a landscape-level process



➤ Feral plants always a part of the story in introducing variability during domestication!



➤ But we know so little about them!

➤ Mabry et al. (2023). Building a feral future: Open questions in crop ferality. *Plants, People, Planet*.



Scan to access the paper

# A FERAL FUTURE

- *Can we harness the genetic diversity found in feral plants to develop crops that are better adapted to withstand harsher environments?*



*Feral Brassica oleracea*

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- Can we harness the genetic diversity found in feral plants to develop crops that are better adapted to withstand harsher environments?



Feral *Brassica oleracea*



Alex McAlvay



Pam and Doug Soltis

- Current USDA postdoc fellowship to investigating feral *B. rapa* and *B. oleracea*.
- Let us know if you know of any populations we should include!



# ACKNOWLEDGMENTS + FUNDING

Genetics, Genomics and Breeding section organization  
committee:

Matt Nelson and Wallace Cowling

Troy Rowan, Harly Durbin, Sarah Unruh, Marianne  
Slaten, Bob Schnabel,  
Jeff Ross-Ibarra, and Paul Blischak



University of Missouri



**WISCONSIN**  
UNIVERSITY OF WISCONSIN-MADISON

- Staff at the MIZZOU DNA Core
- Research Computing Support Services (RCSS) and Informatics Research Core Facility (IRCF)

- Staff of UW-Madison Biotechnology Center and Bioinformatics Resource
- Staff at the Walnut Street Greenhouse at UW-Madison



Project No. 8060-21000-024-00D



Project No. DOE HDTRA 1-16-1-0048



Award No. DEB-1601430  
Award No. NSF IOS 1339156  
Award No. PRFB DEB-1711347

QUESTIONS?

