

# DIVERSITÉ GÉNÉTIQUE TEMPORELLE (1984-2020) ET SPATIALE D'UNE POPULATION DE BLÉ SAUVAGE SOUS L'EFFET DU CHANGEMENT CLIMATIQUE



Avi Levy, March 2021

היכל וויצמן למדעים

WEIZMANN INSTITUTE OF SCIENCE

KIBBUTZ AMMIAD

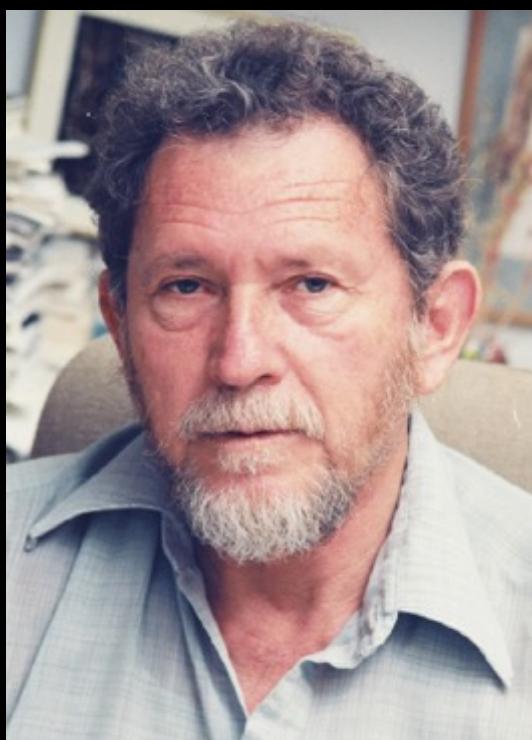


# Ammiad wild wheat population

- A project that started in 1984 under the umbrella of the Israeli Gene bank as a test case for *in situ* conservation
- Located at the center of distribution of the species as suggested by high genetic and phenotypic diversity
- Declared in 2007 by SPNI as a natural reserve for *in-situ* wheat biodiversity conservation
- Test case for spatial and temporal adaptation



Yehoshua  
Anikster (Head)  
TAU



Moshe  
Feldman  
WIS



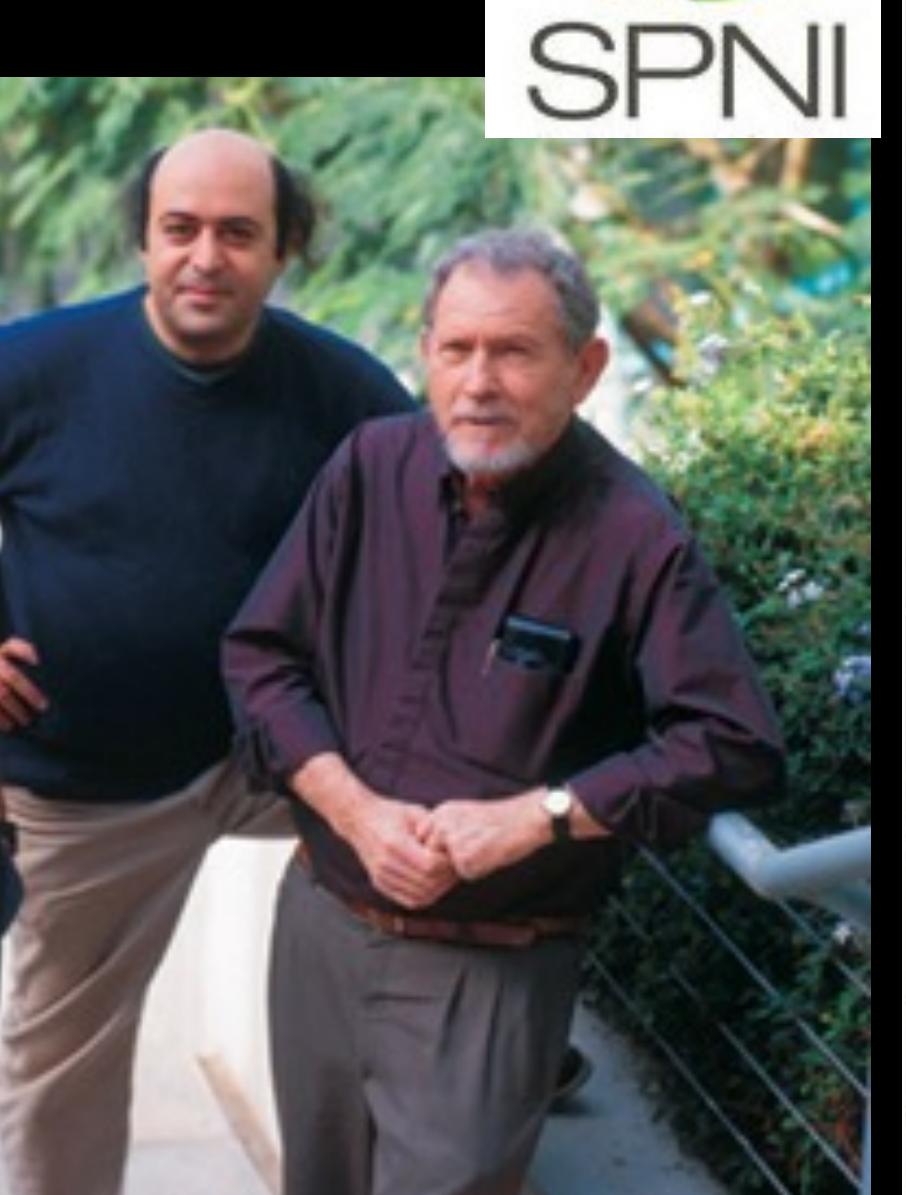
Theor Appl Genet (1988) 75:651–658



Magnus  
Nordborg  
GMI



Tal  
Dahan-Meir



Ecogeographical distribution of HMW-GS in populations of the wild tetraploid wheat *Triticum turgidum* var. *dicoccoides*

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# Wild emmer wheat distribution in the fertile crescent

*Triticum turgidum ssp. dicoccoides*



Self-pollinating ~ 99%

Annual plant (1 generation/year)

Tetraploid genome ( $2n=4x=28$ , BBAA)

Approximately 12 Gbp

Sequenced in 2017 (Avni et al. science)

80% repetitive elements  
~65,000 genes

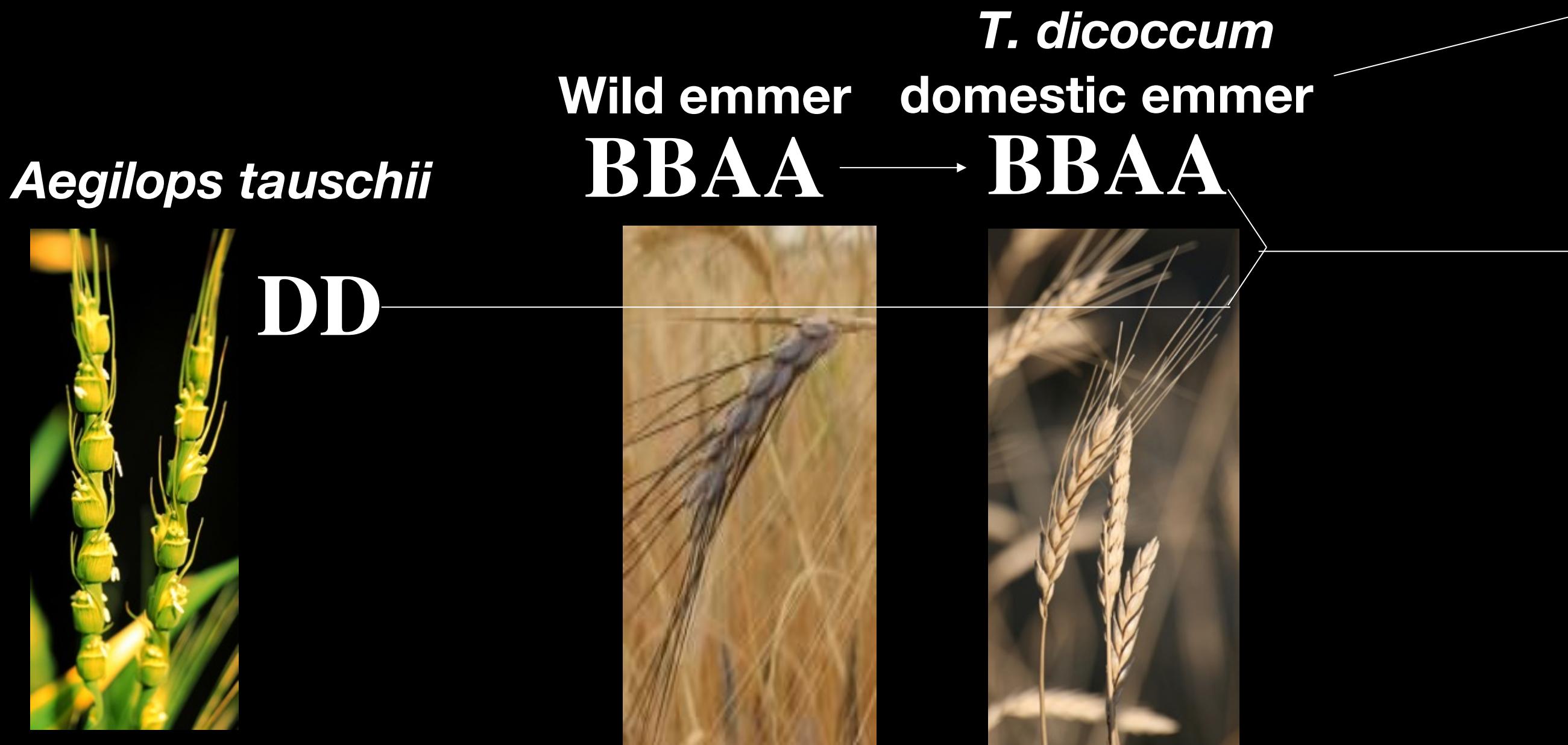


South Levant: Center of variation (= Center of origin?)  
Center of domestication of wild emmer wheat

From Salamini et al.  
2001, Nat Rev

Wild emmer: direct progenitor of modern wheat;  
rich source for genes conferring resistance to  
diseases, abiotic stress, yield and quality

*durum*  
wheat  
BBAA



Important for food security:

- Conservation
- Characterization
- Exploitation

Bread wheat  
BBAADD



20%  
calories  
consumed  
by Human

20%  
proteins

750 Mt/Yr

AABBDD

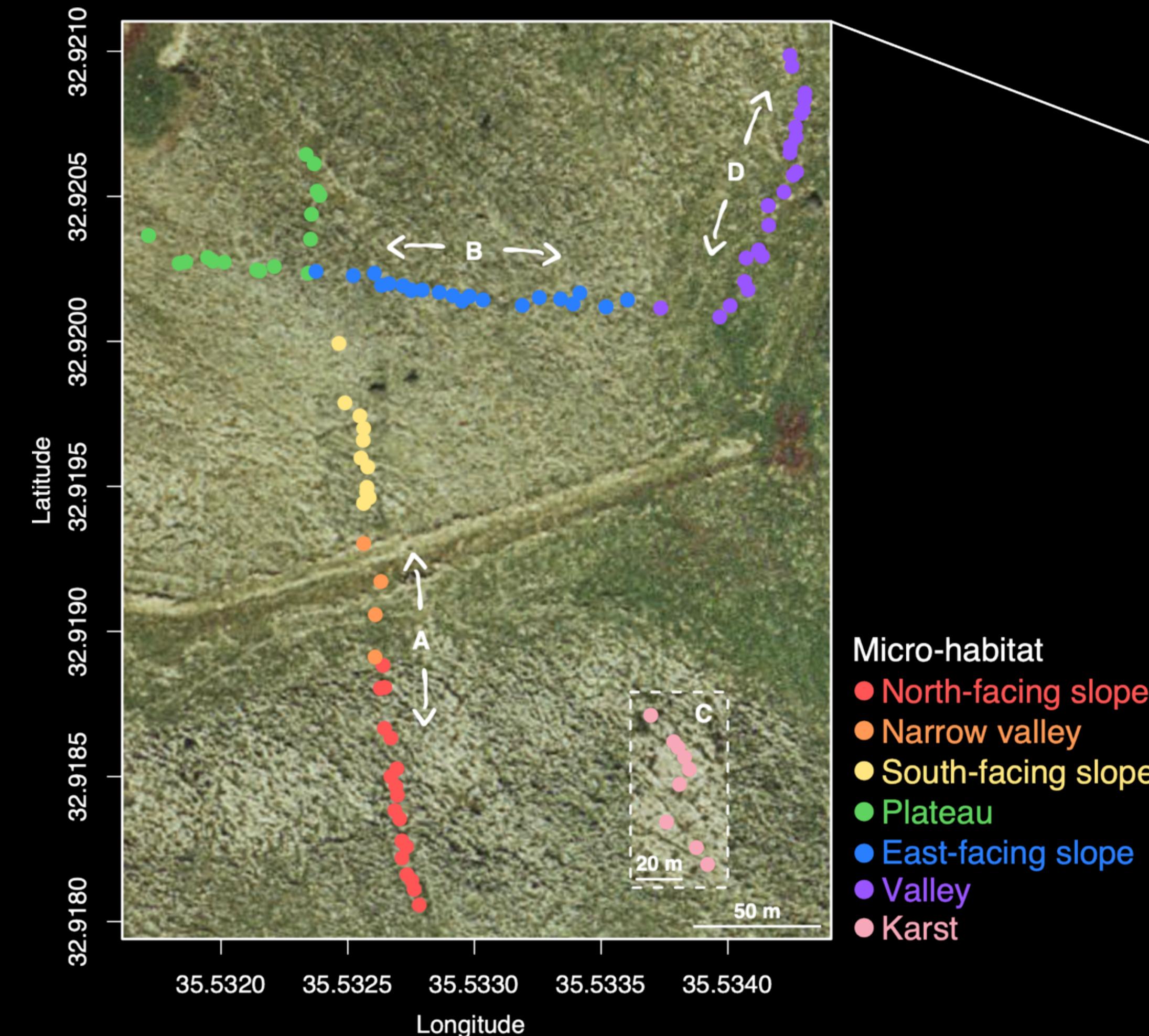
# AMMIAID POPULATION

- Four transects - 100 marked (GPS+Peg) points, sampled between 1984 to 2020
- Transects diverse in topography, soil/rock composition and flora

Collection in:

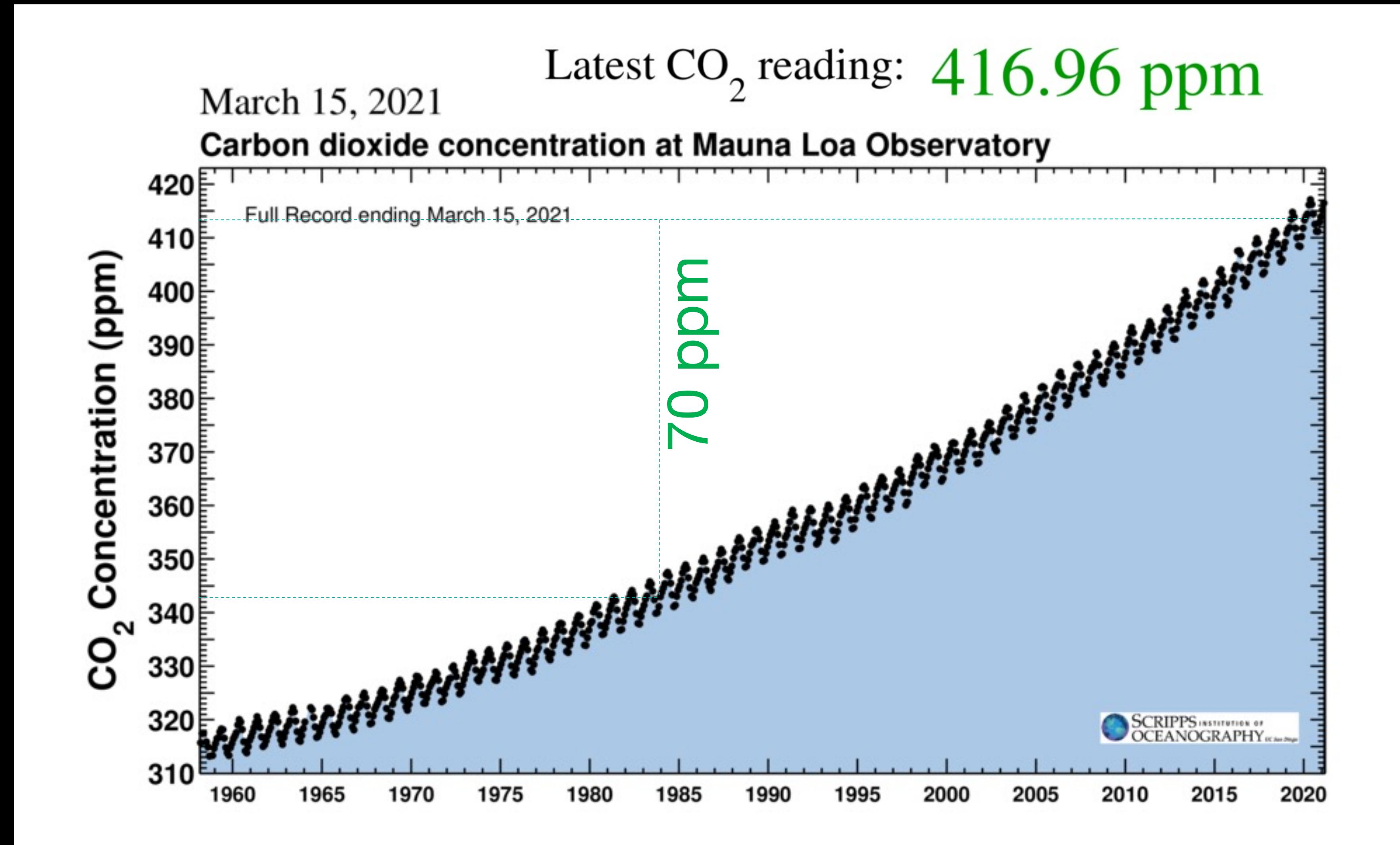
1984  
1988  
1992  
1996  
2002  
2014  
2016  
2018  
2020

Total ~900 samples

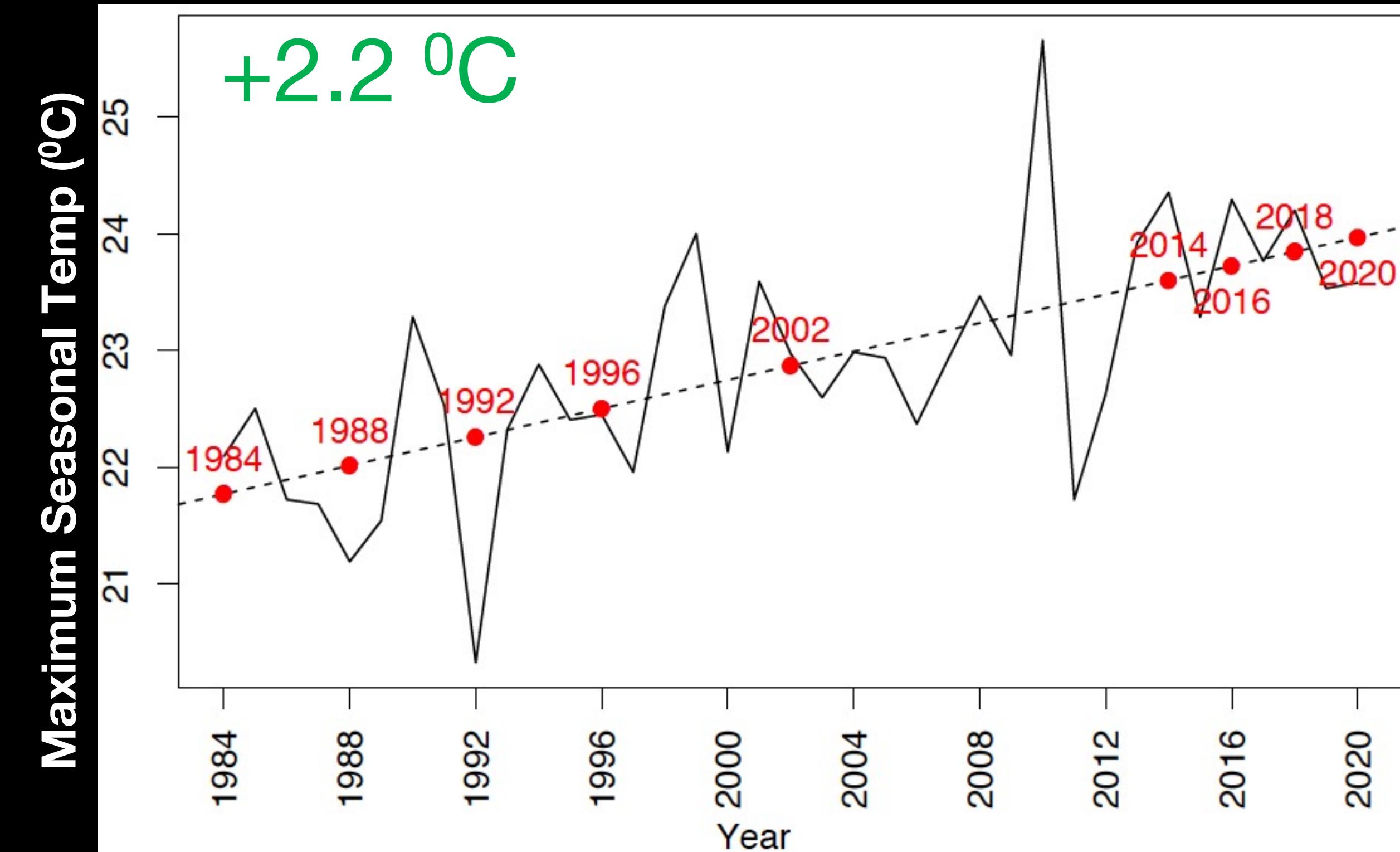
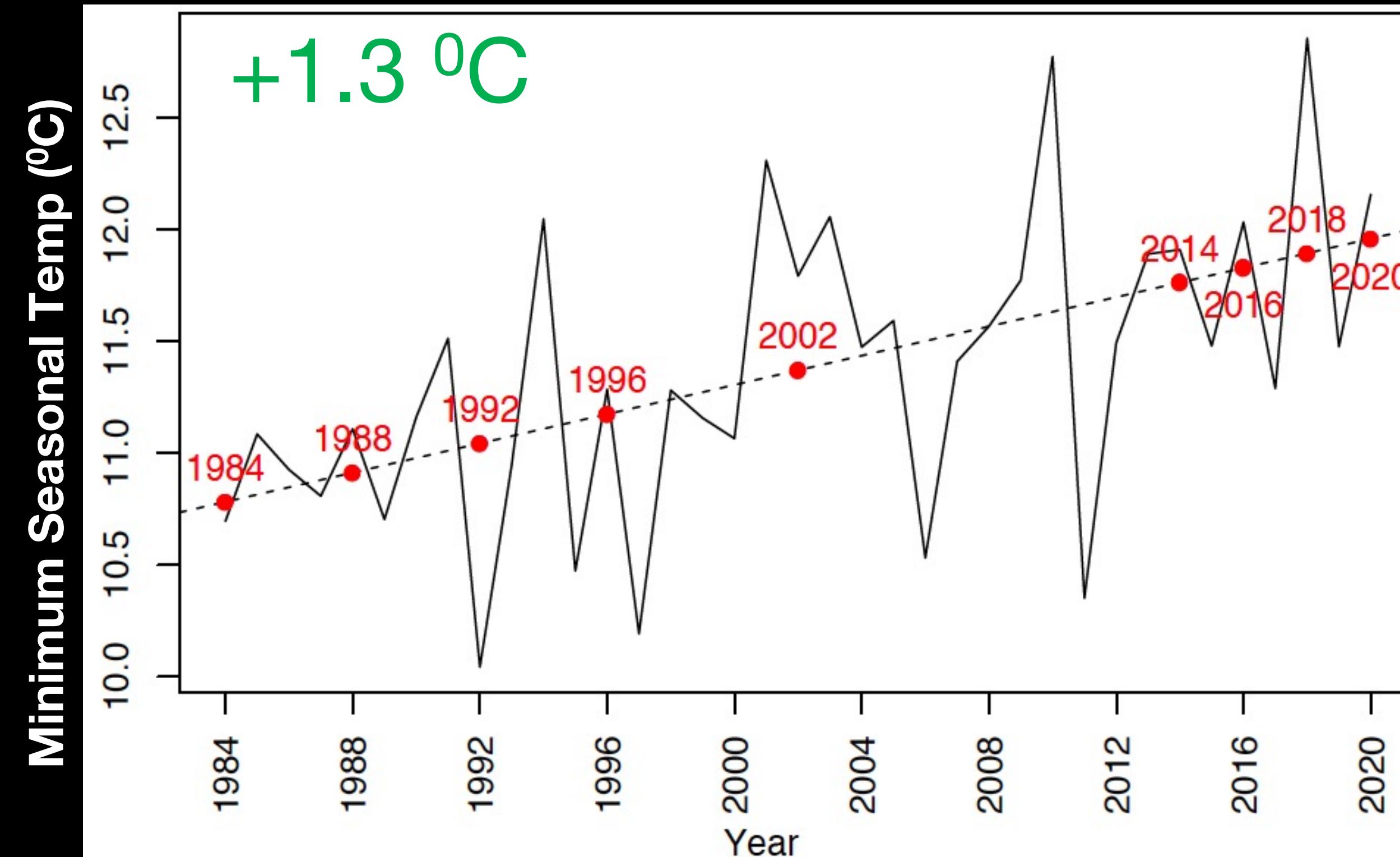


# 70 ppm increase (~20%) in atmospheric CO<sub>2</sub> concentration between 1984-2020

THE KEELING CURVE

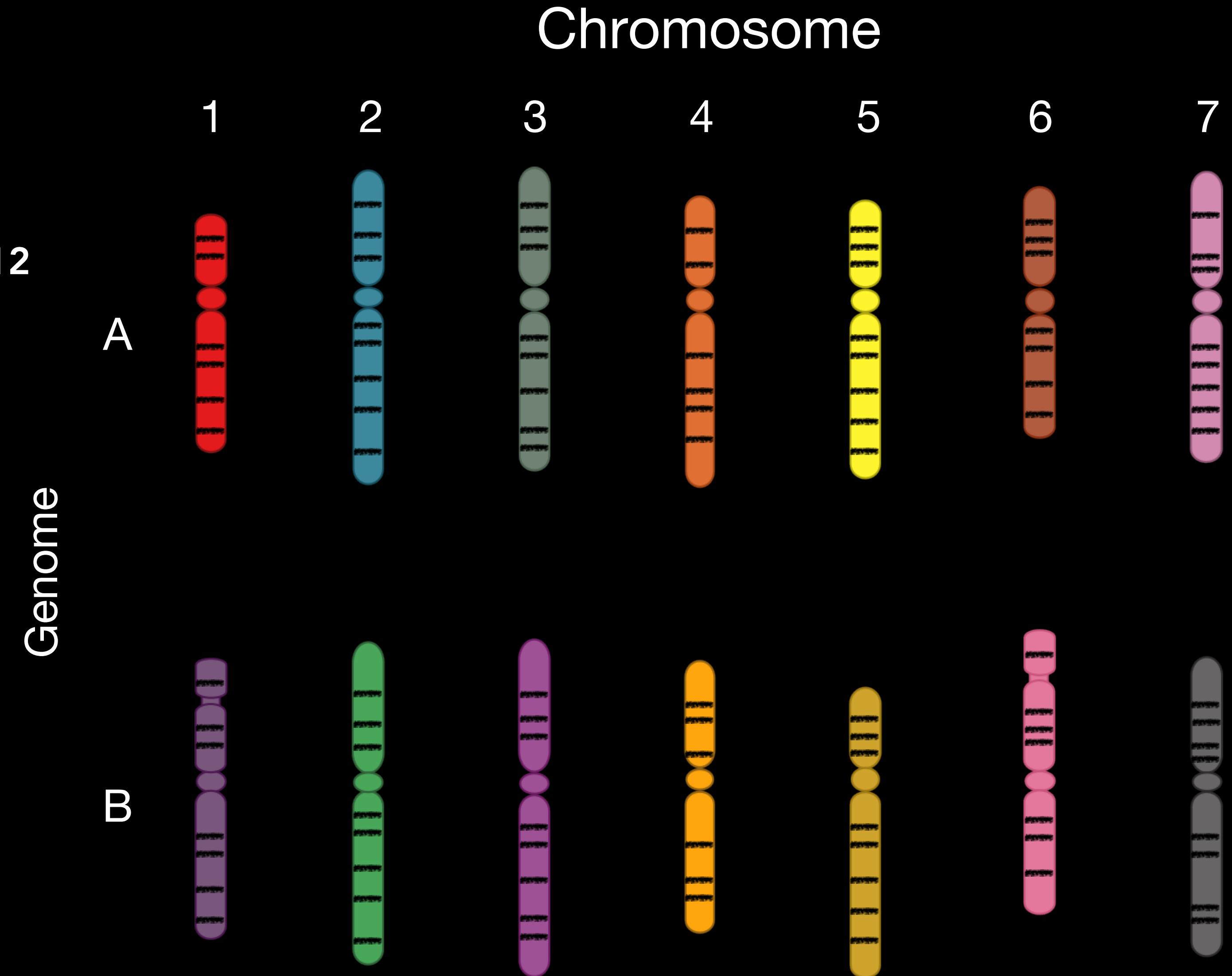


# Temperature increase between 1984-2020 during growth season (Nov-May) near Kibbutz Ammiad



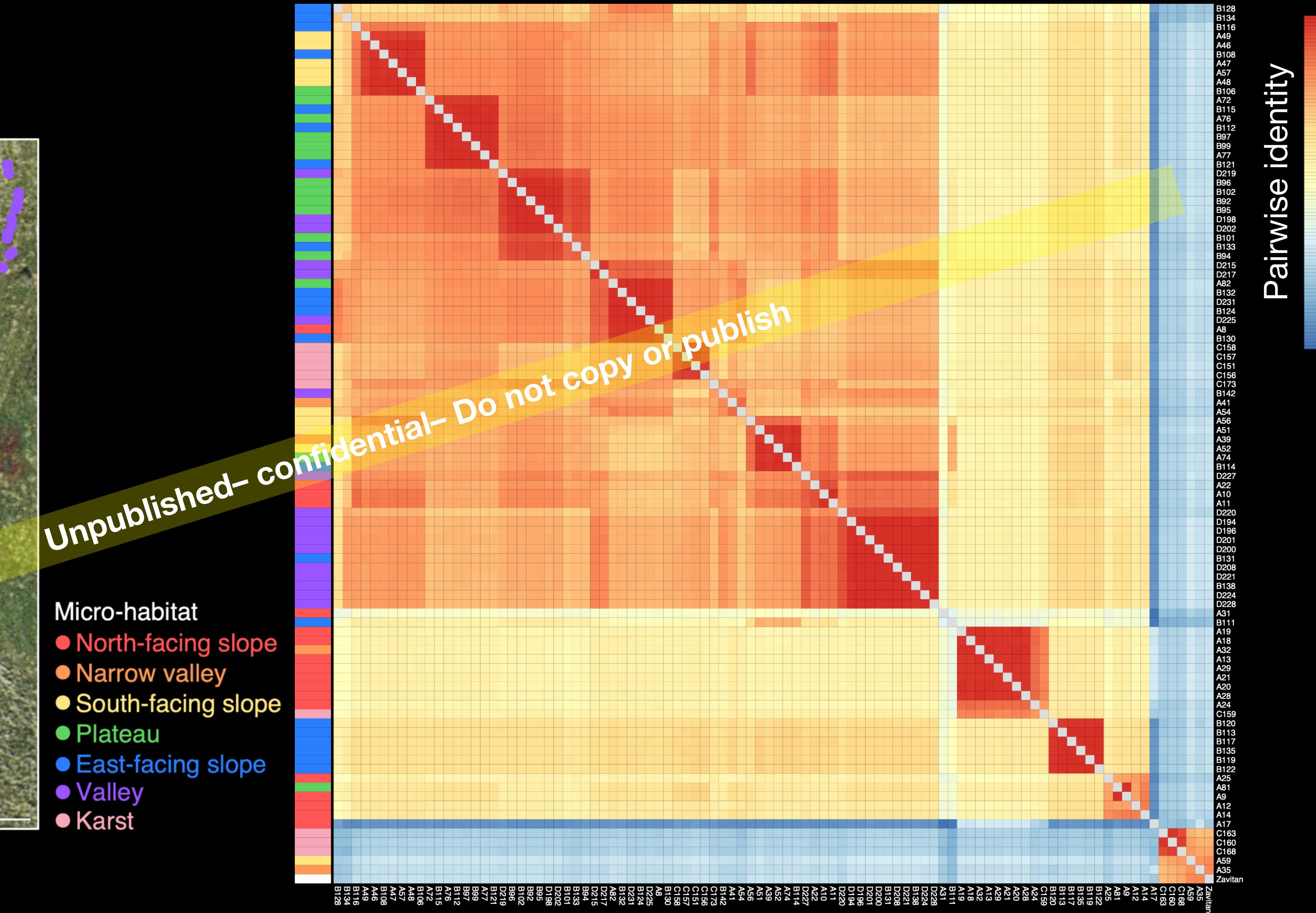
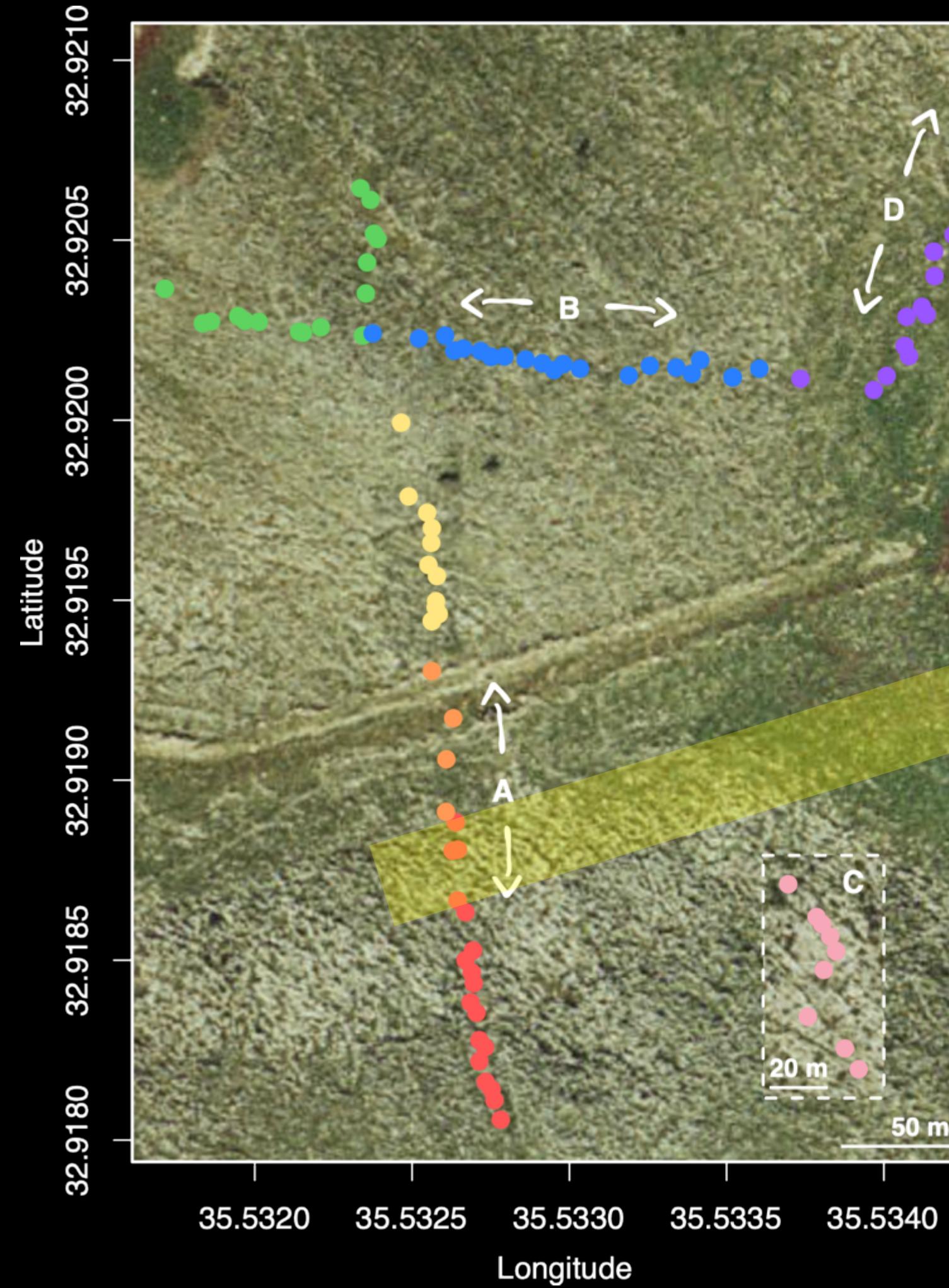
# GENOTYPING OF EACH PLANT

- Genotyping-by-Sequencing (GBS - ddRADSeq) Poland et al. 2012
- Reduced representation of the wheat genome (<1%)
- Filtering for coverage >6x
- Filtering for unique sequences
- Identification of heterozygotes



4286 POLYMORPHIC MARKERS ALONG THE GENOME

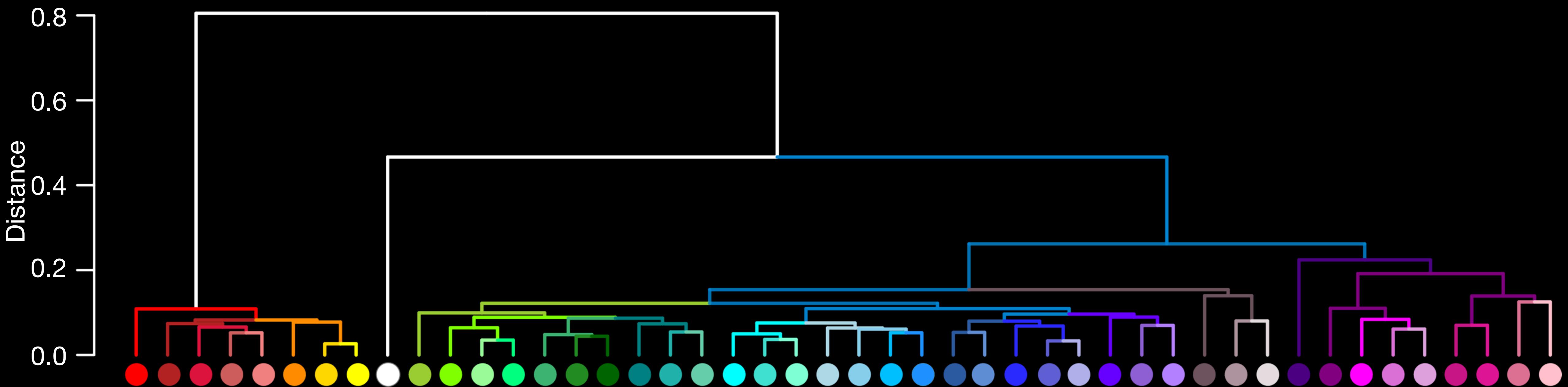
# WHAT IS THE POPULATION STRUCTURE? 1984



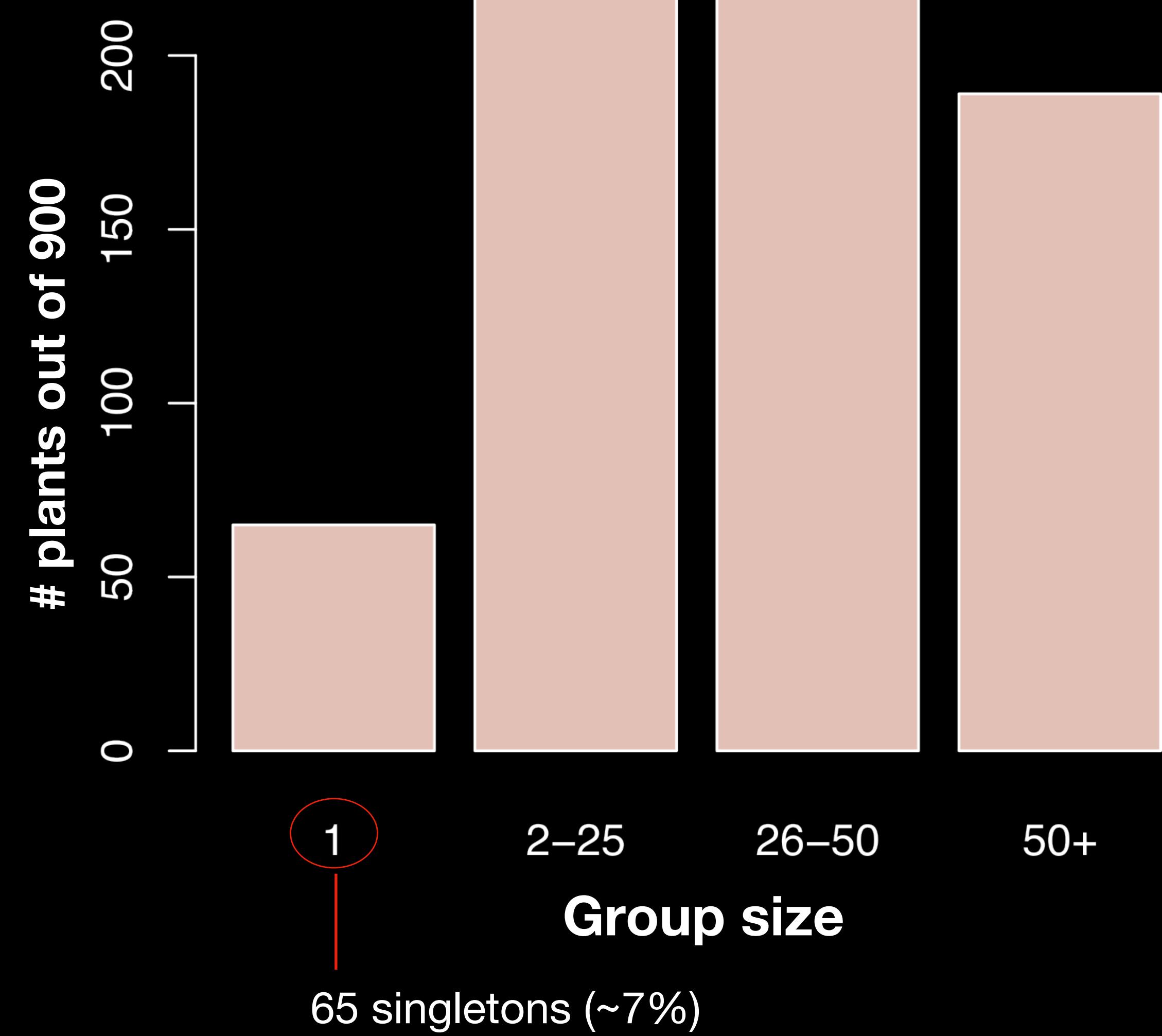
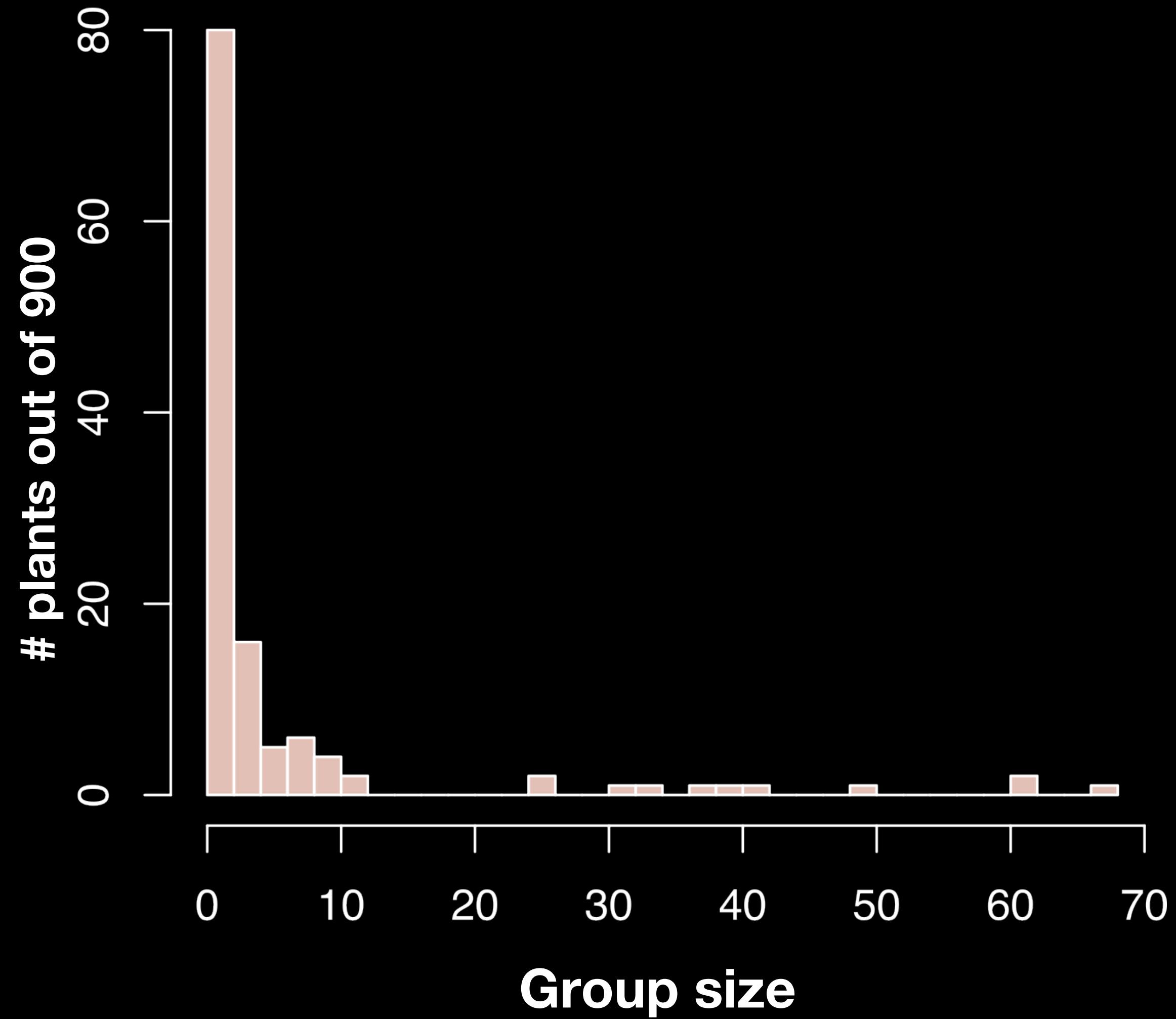
# IDENTICAL GENOTYPE GROUPS (IGGs)

**IGGs = Groups with >99% pairwise identity**

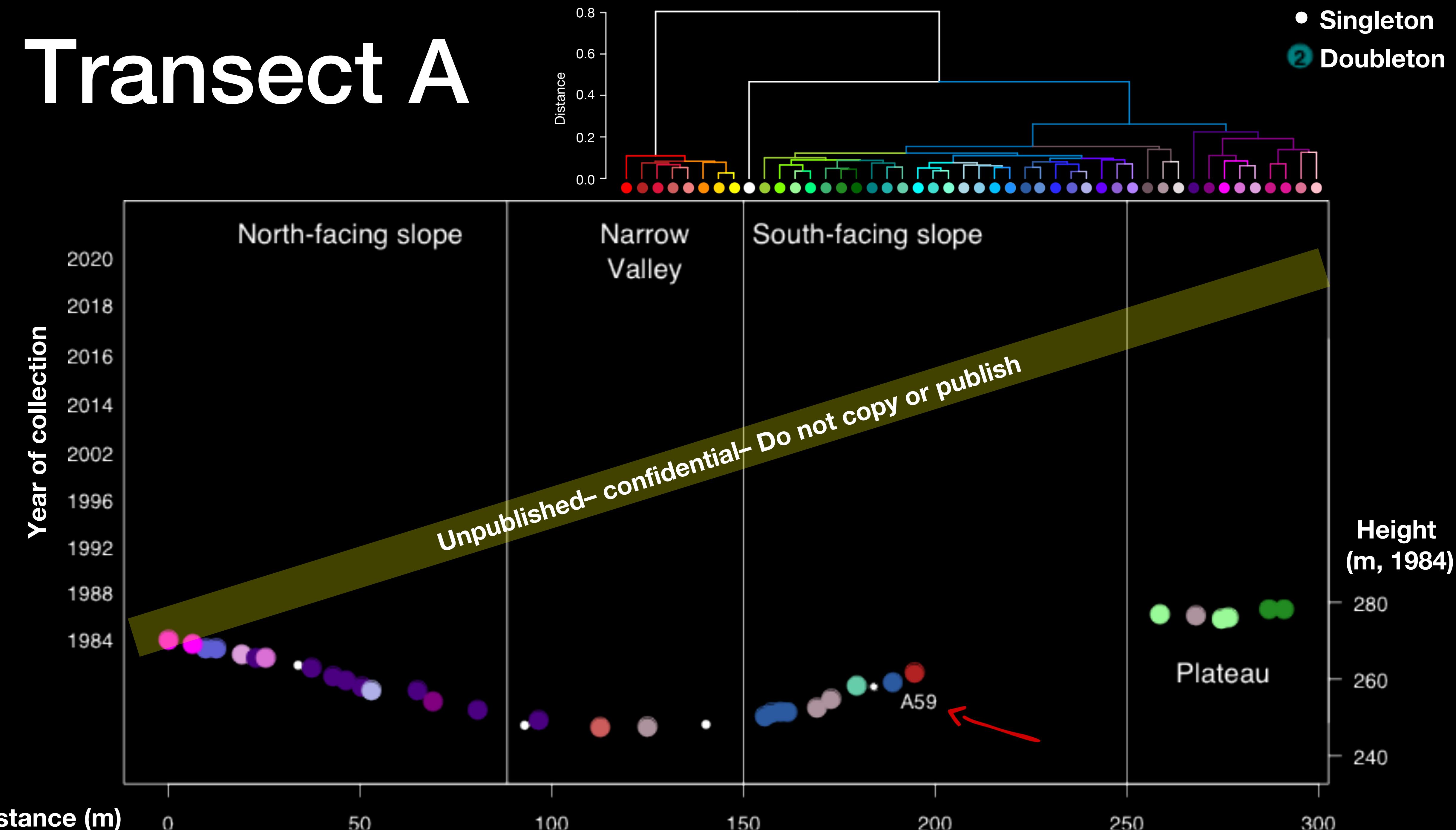
- 127 IGGs out of 900 plants



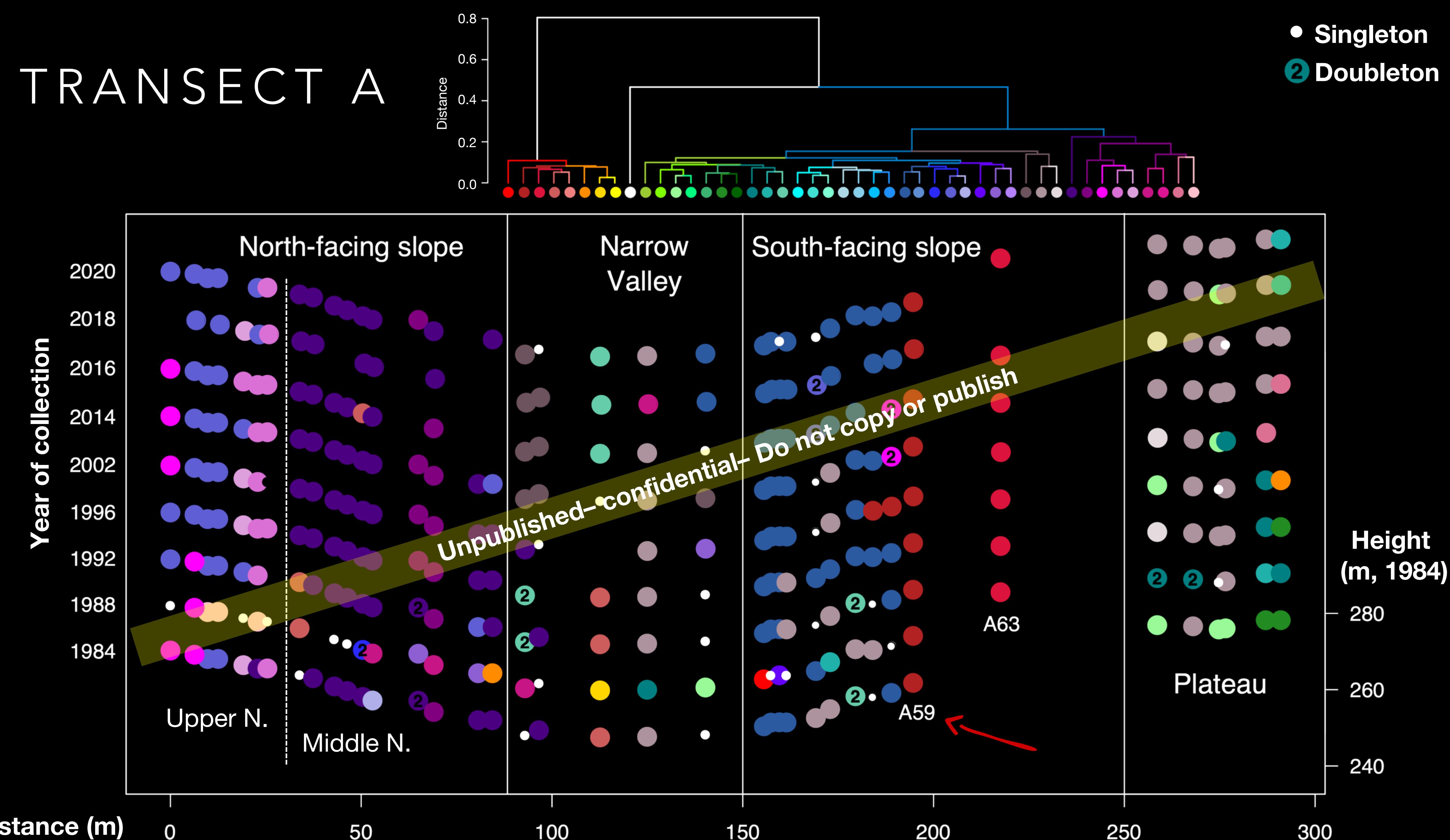
# Sizes of Identical Genotype Group (IGGs)



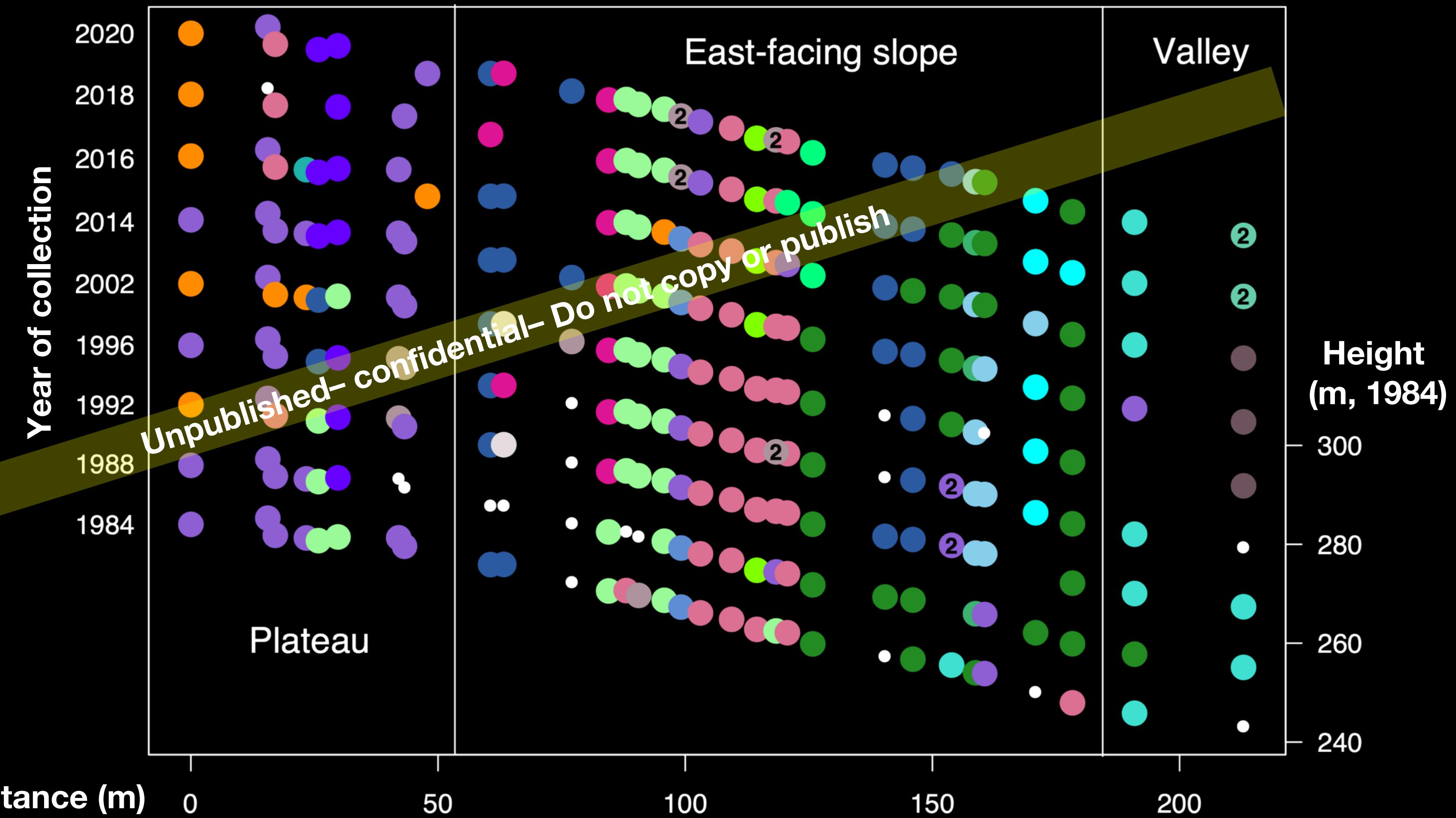
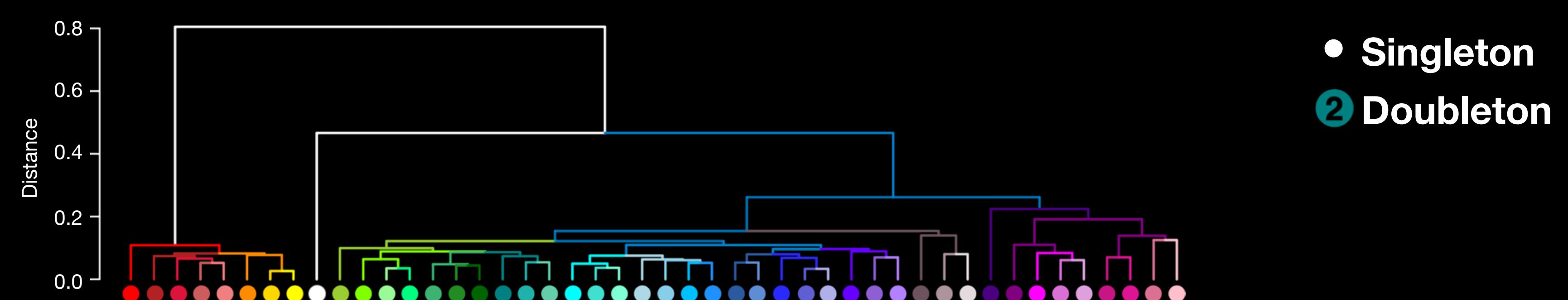
# Transect A



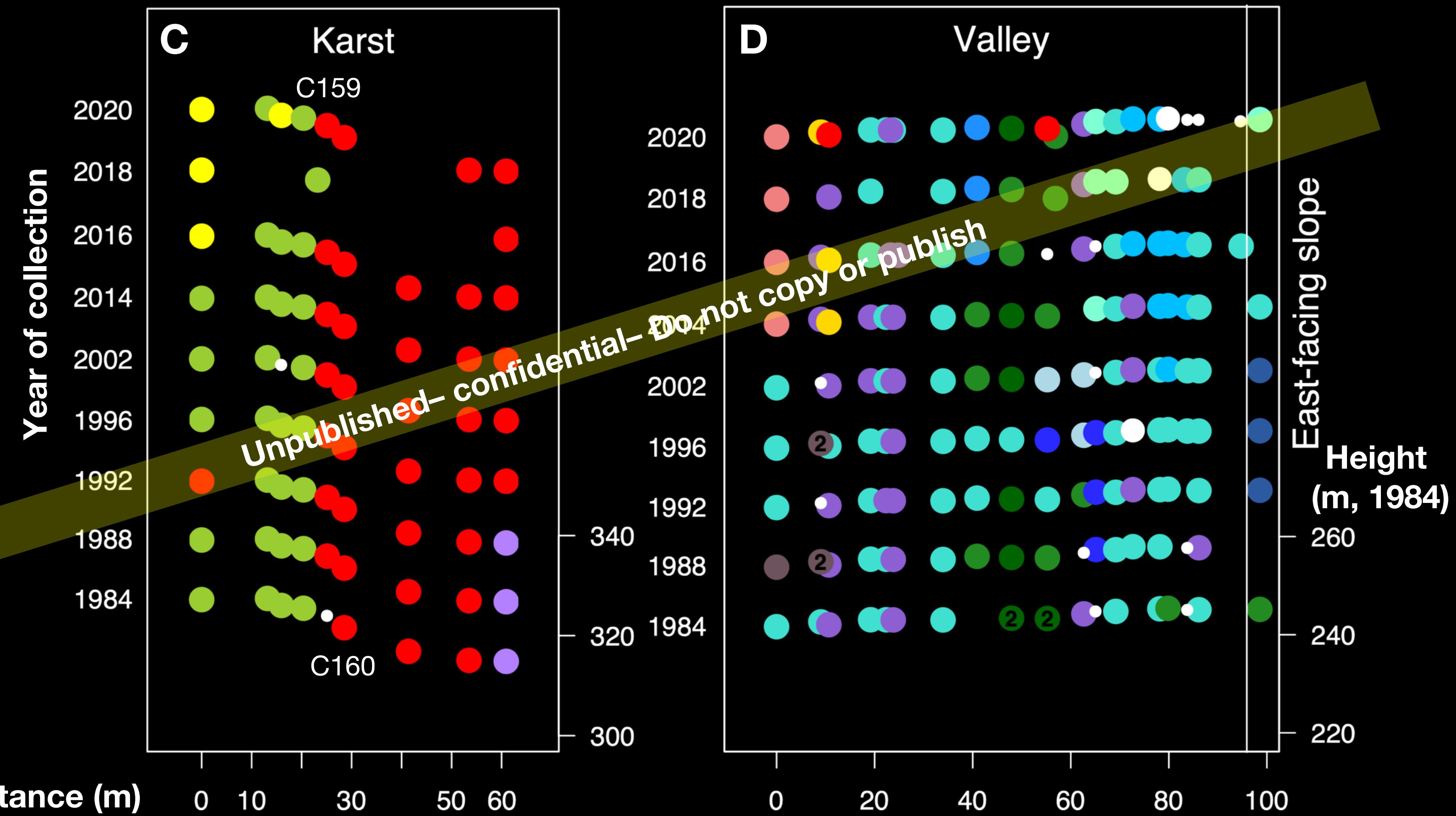
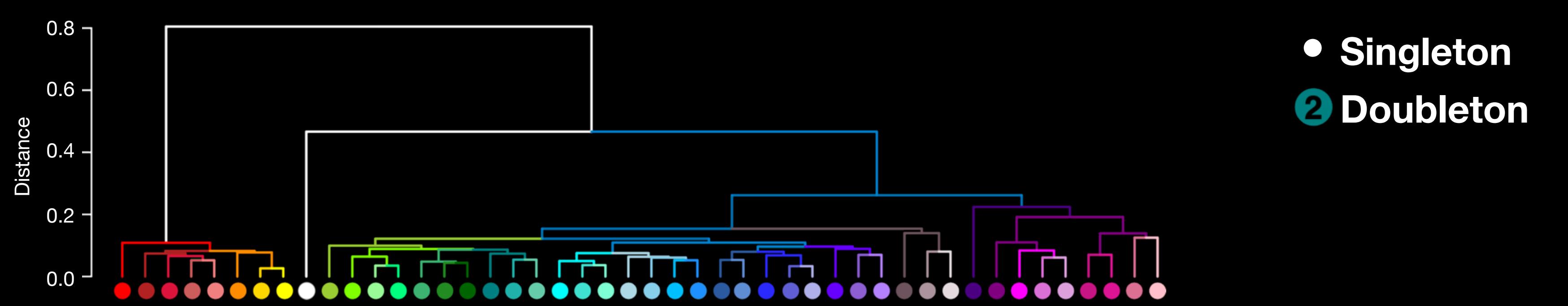
# TRANSECT A



# TRANSECT B



# TRANSECTS C, D



# SIMULATIONS, MODELLING & STATISTICS- TOM (NORDBORG GROUP)



Thomas Ellis

## Could the spatial and temporal structuring of IGGs be explained by neutral processes?

On-Going analysis:

- What is the probability for IGGs to be found in clusters assuming random order?
- What is the probability that a given sampling point will be occupied by the same genotype over years of collection?
- Indications for adaptation?

# CONCLUSIONS

- Highly structured population
- Genetic structure and diversity are robust through time despite climatic changes during the past 36 years
- Highly unlikely that the population stability can be explained by neutral processes
- *In situ* conservation of wild wheat biodiversity:
  - Sample once in a few decades
  - Collect taking microhabitats into consideration to cover most genetic diversity
  - Importance of keeping undisturbed and protected habitats, even if of limited size. (several wild wheat populations are disappearing due to urbanization or over grazing)



MERCI



THANK YOU!



מכון ויצמן למדע  
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