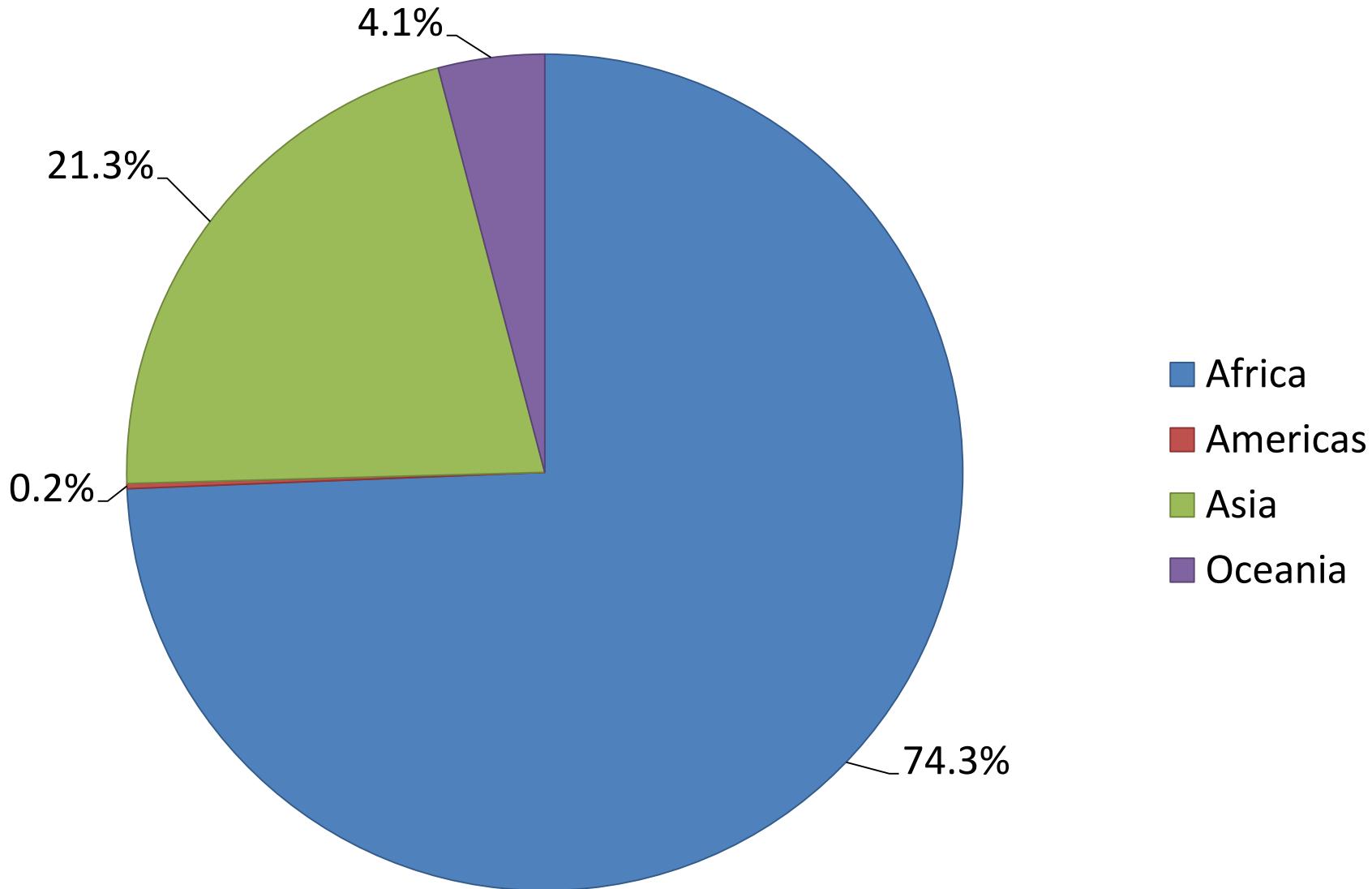
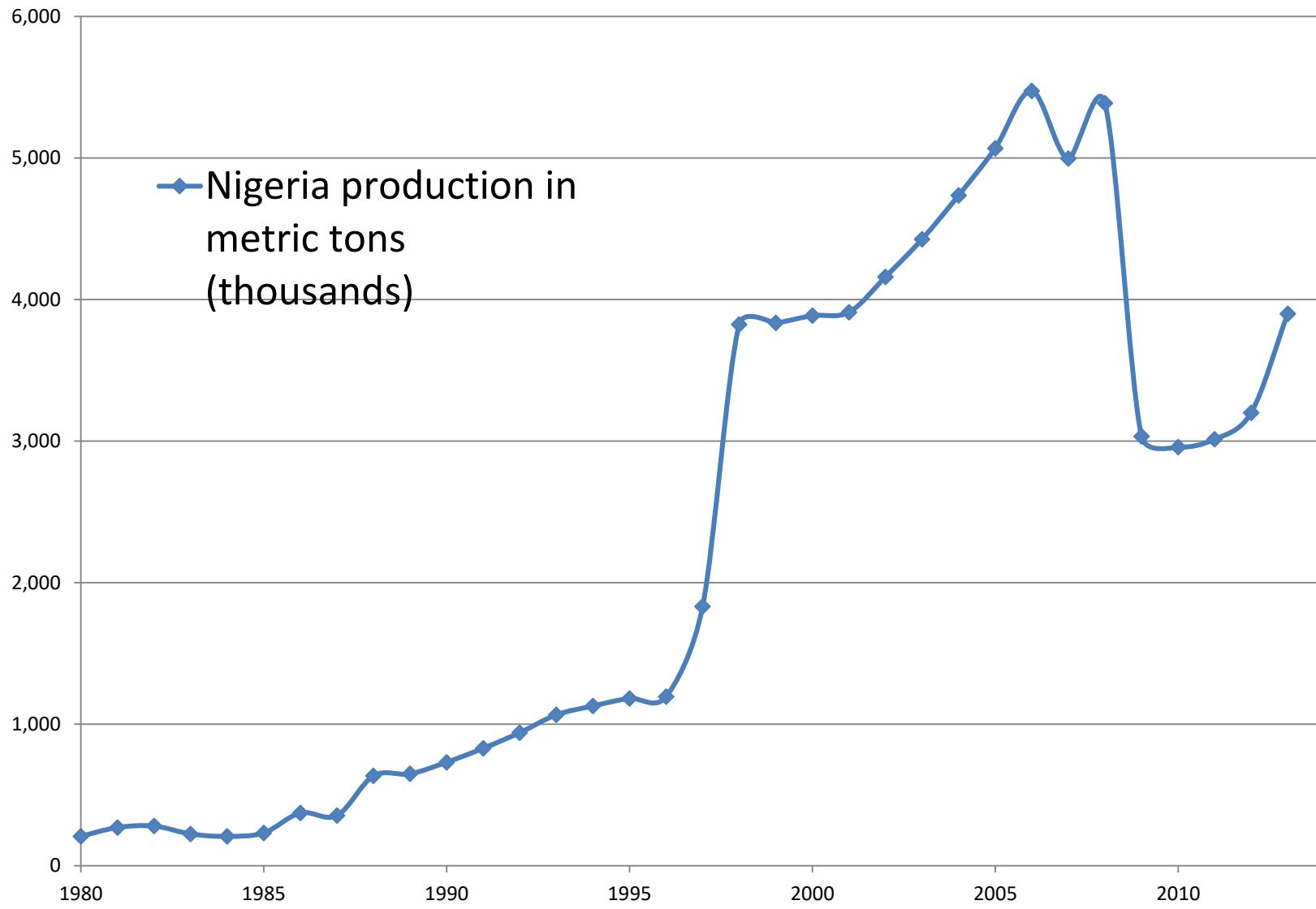


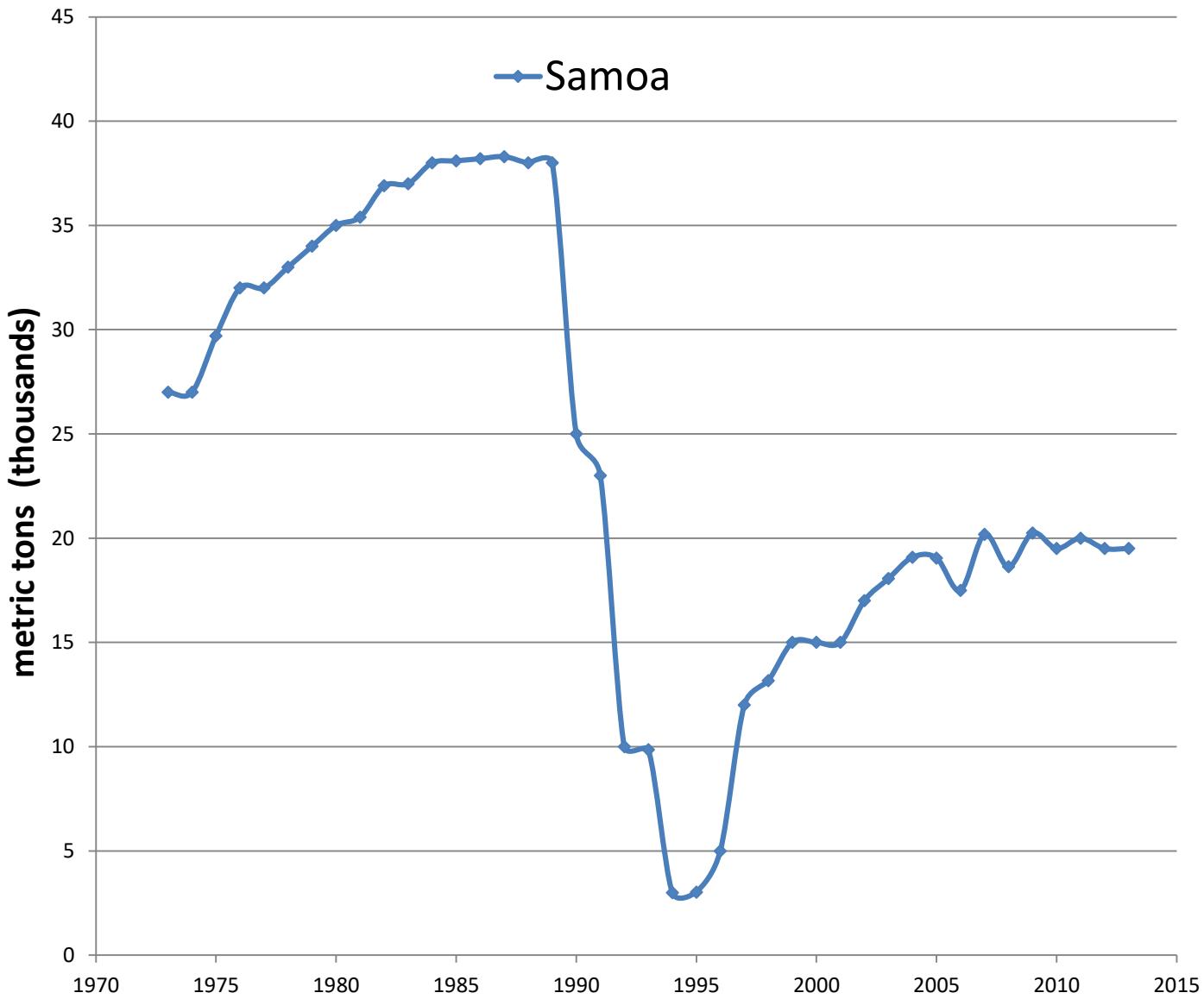
# Developing Genomic Tools for the Protection and Improvement of Taro



### World Taro Production









# Taro Leaf Blight (TLB)

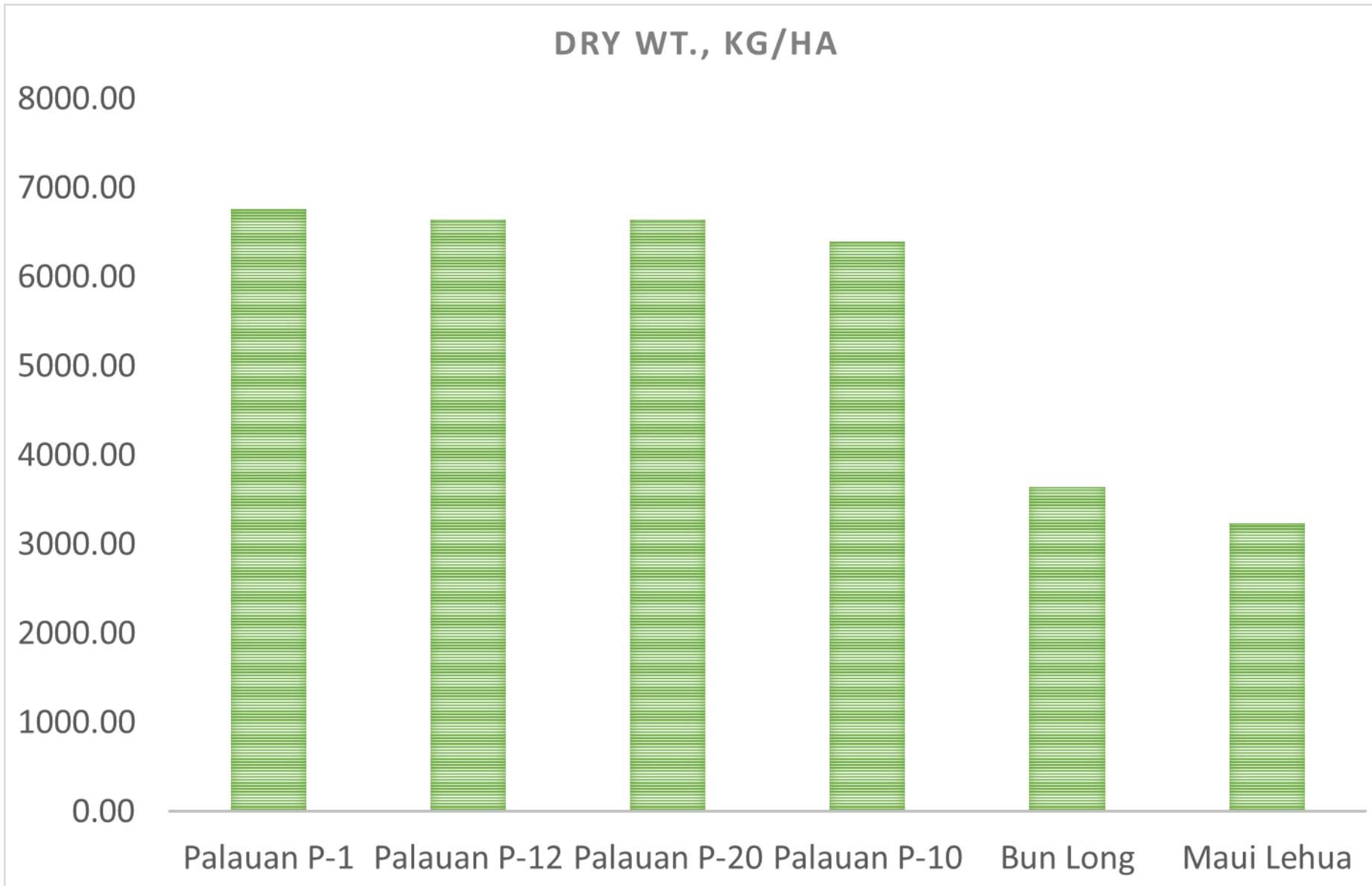
*Phytophthora  
colocasiae*



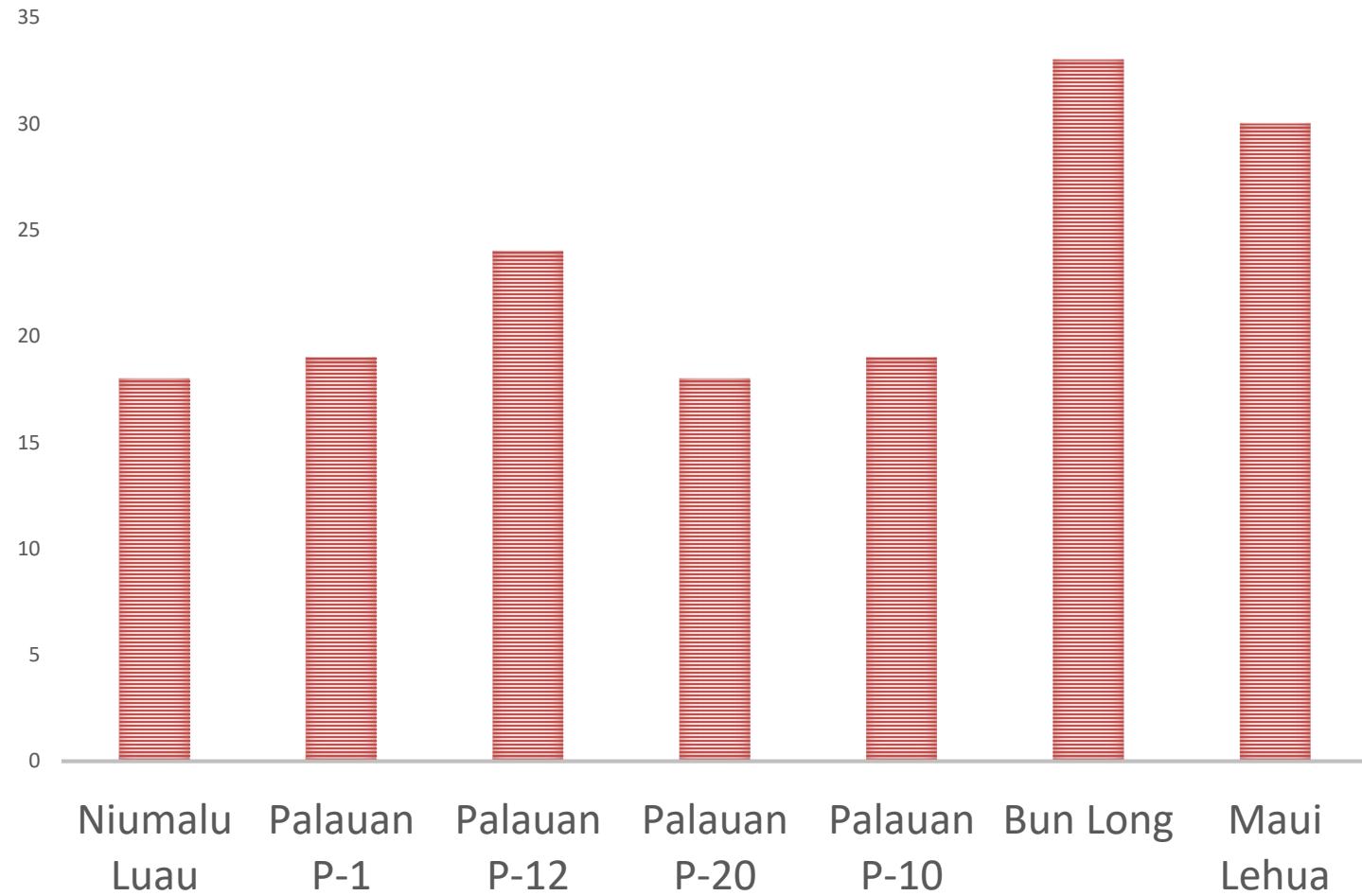


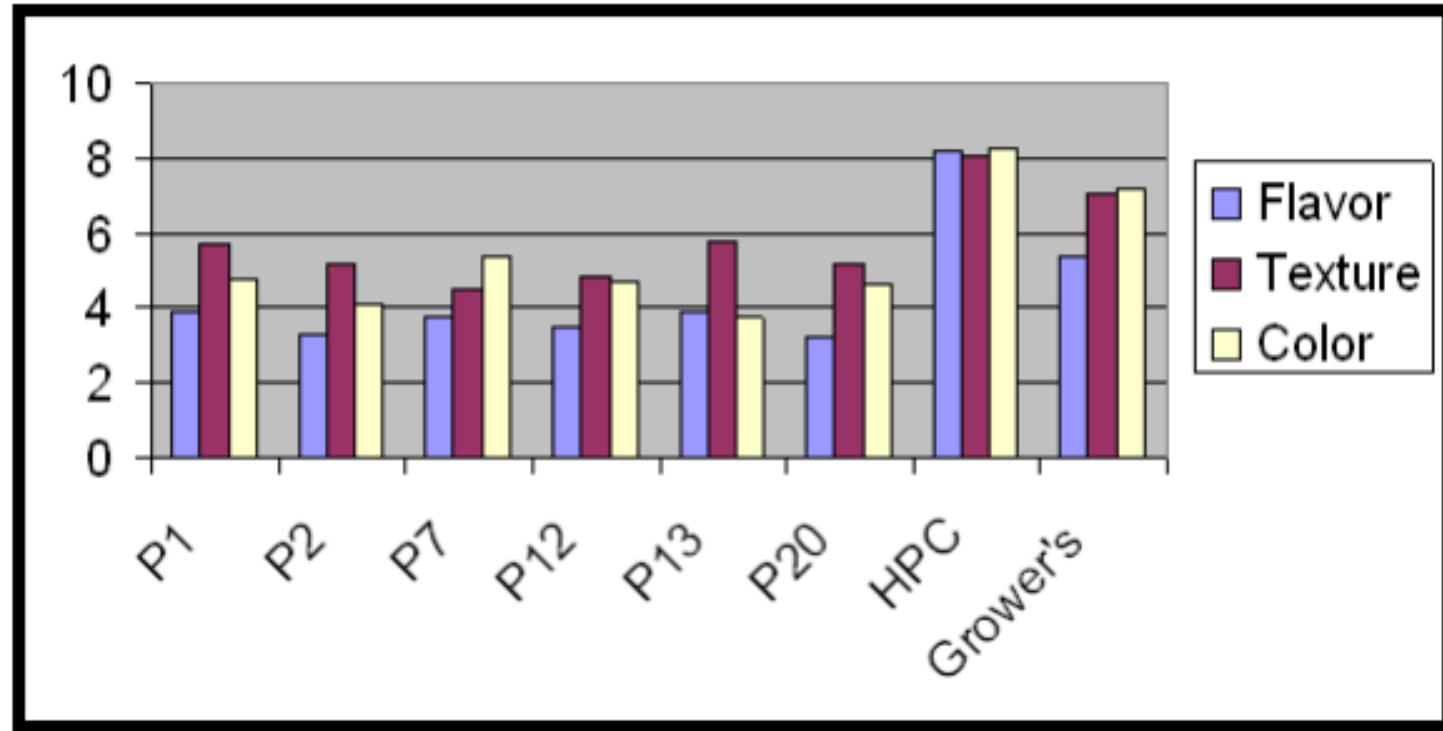


# Taro Cultivar Trials 1994-2006



## TLB RATINGS (HIGHER #'S= MORE BLIGHT)





#### PALAUAN VARIETIES:

P1 = Ngesuas

P2 = Terrekakl

P7 = Ochelochel

P12 = Merii

P13 = Dirraiuosch

P20 = Dirratengadik

The following were included in the taste test for comparison:

HPC = Taro Brand Poi

Grower's = Poi from Kauai Wetland

"Poi products made from the Palauan taros all failed at the grower and processor level in relation to color and taste." D. Sato 1999

Arnold Melim of Palolo was crossing Hawaiian taro varieties in the 1970's.

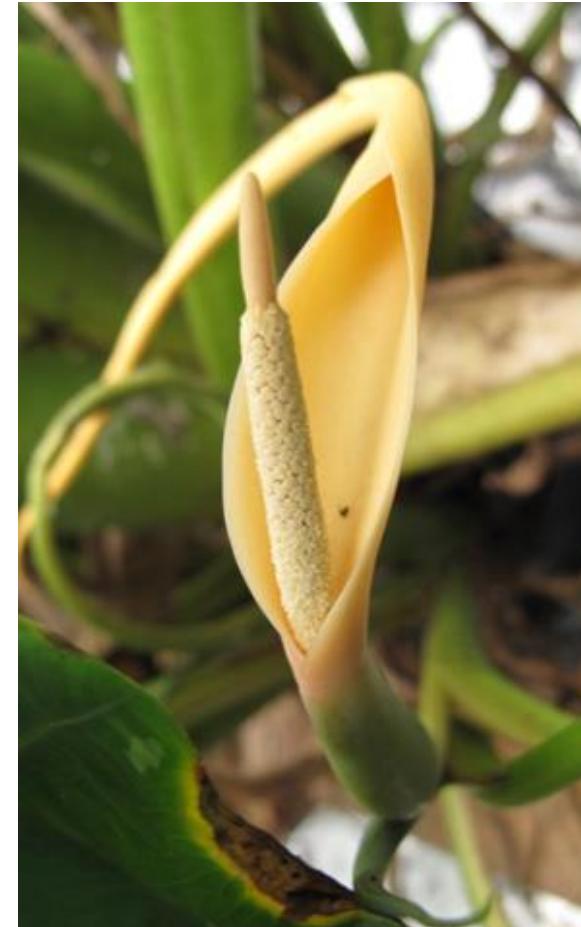
In the 1980's, Dr. Ramon Dela Pena on Kauai crossed Hawaiian and South Pacific varieties.

In the early 1990's, Dr. Eduardo Trujillo crossed Hawaiian and Palauan taro varieties.

In the early 2000's, Dr. John Cho on Maui crossed Hawaiian varieties with varieties from throughout the Pacific and Southeast Asia

# Hand pollination required

- Male flowers produce pollen approx. 1-2 days **after** female flowers ready
- Cross-pollination is required for fertilization
- Insect that naturally pollinates taro in Australia, Papua New Guinea & Indonesia isn't found in Hawaii



# Seed pods and Taro seedlings



One seed pod of taro could produce up to 4000 seedlings



Slide courtesy Dr. Susan Miyasaka



June 9, 2016

July 20, 2016  
(6 weeks later)



# Taro seedlings



Agricultural Technician C. Bernabe with 4-5 month-old taro seedlings in greenhouse

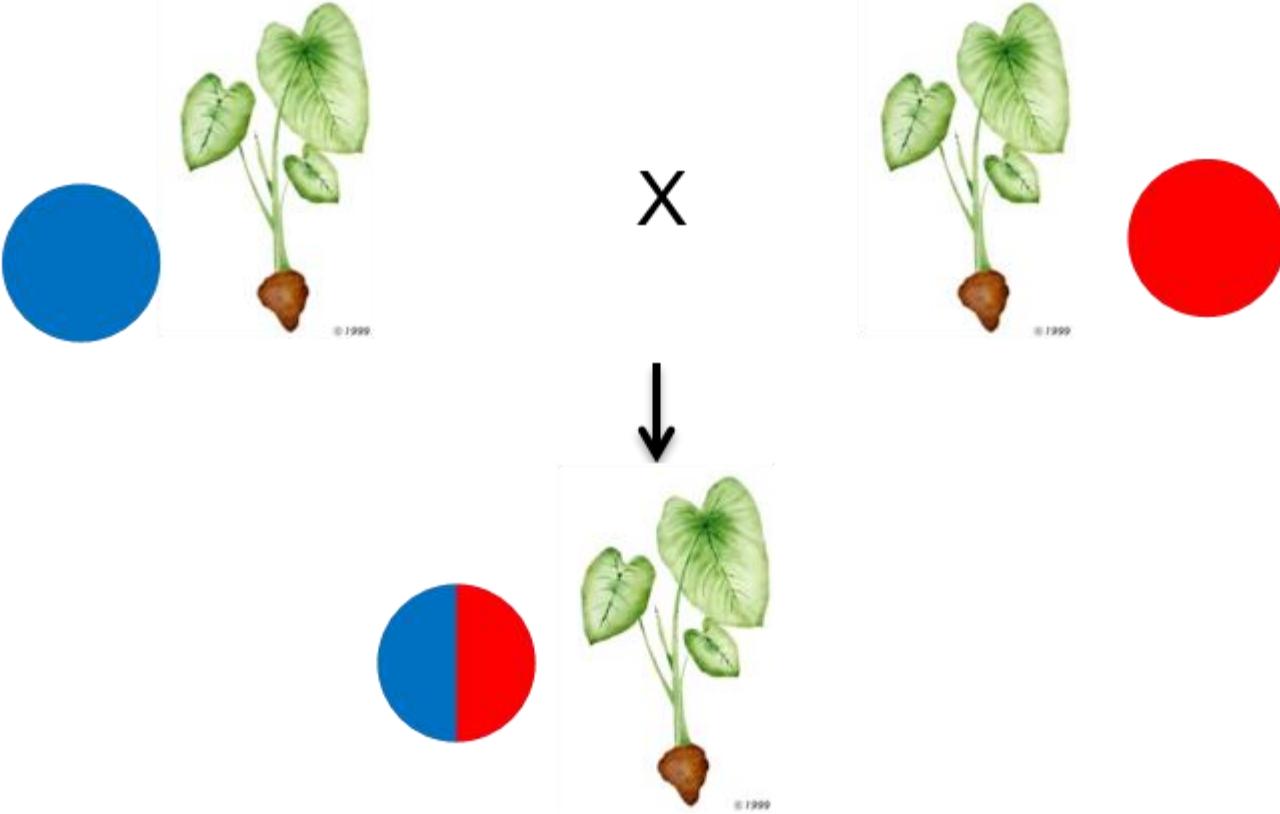


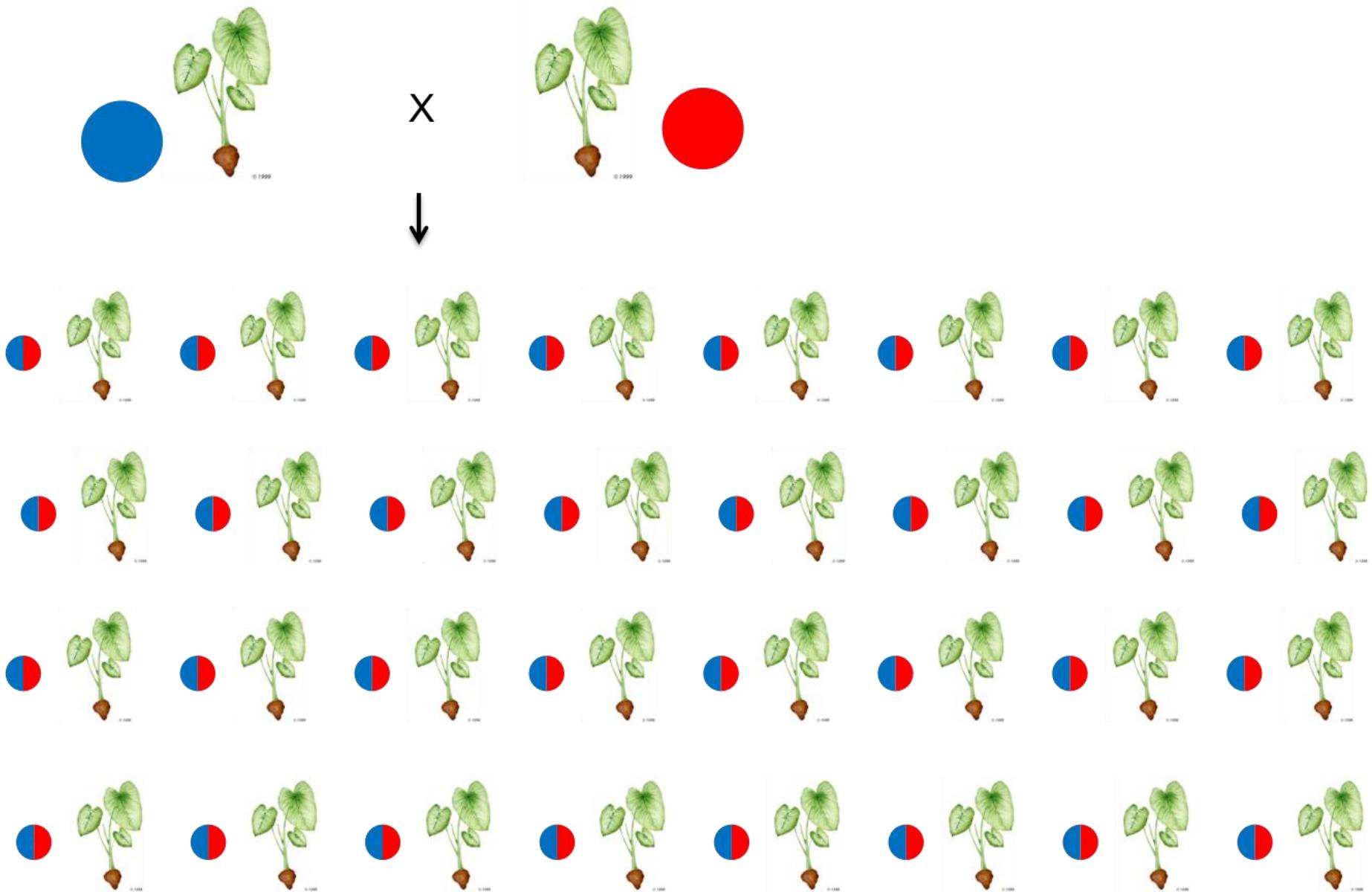
Field of taro seedlings 2 months after planting

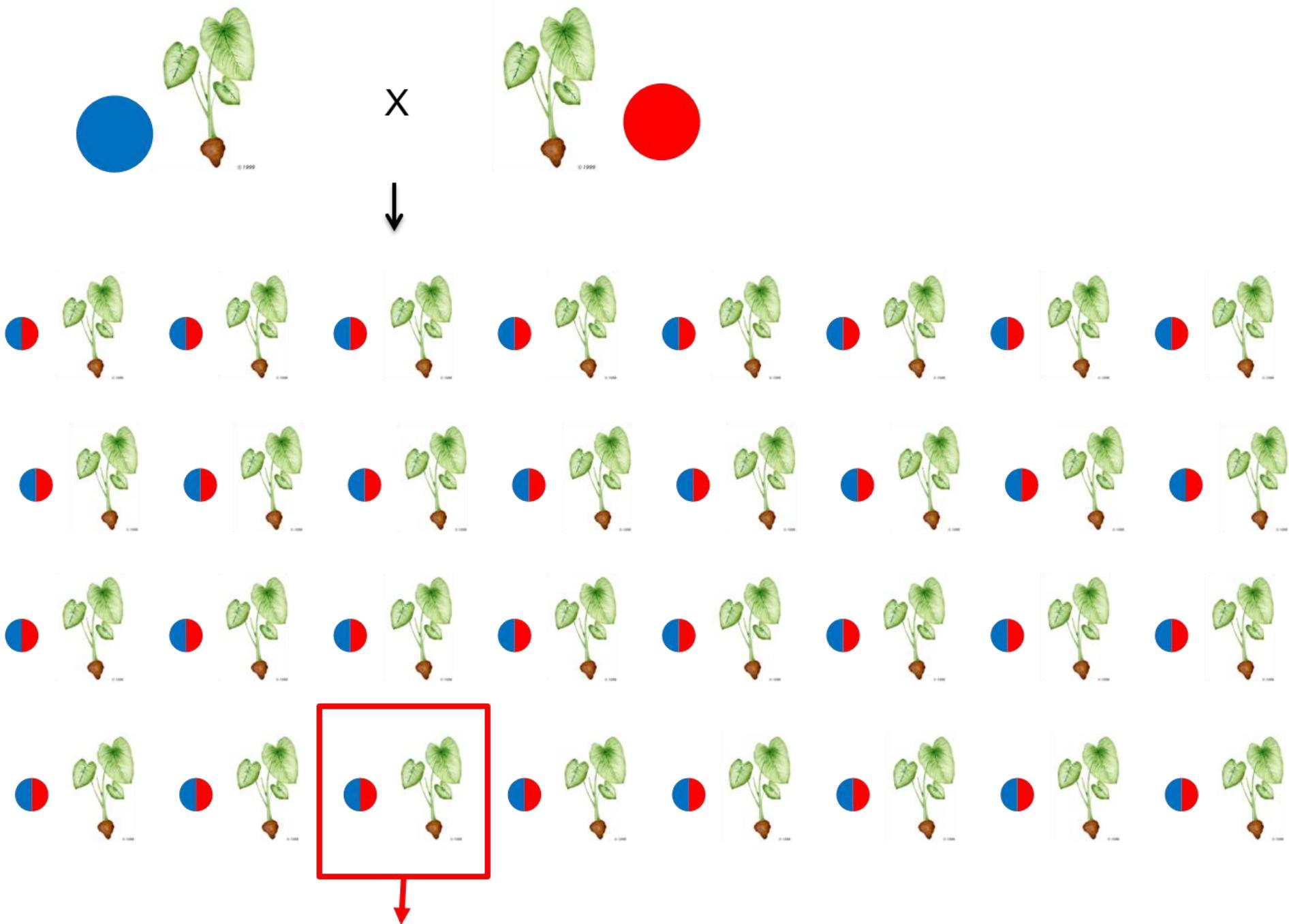
Slide courtesy Dr. Susan Miyasaka

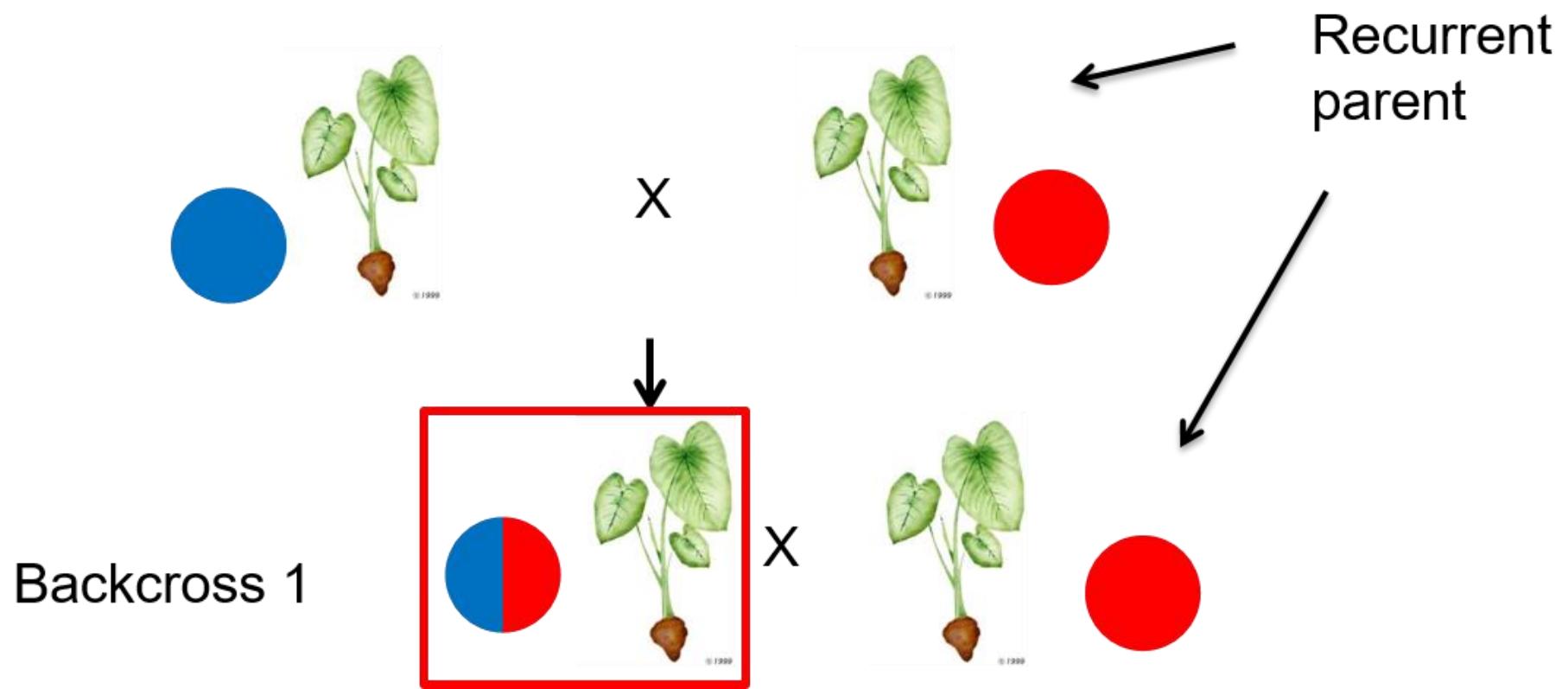
- Taro seedlings are evaluated in field over 1-2 years
- Most promising genotypes are multiplied and evaluated in replicated plots



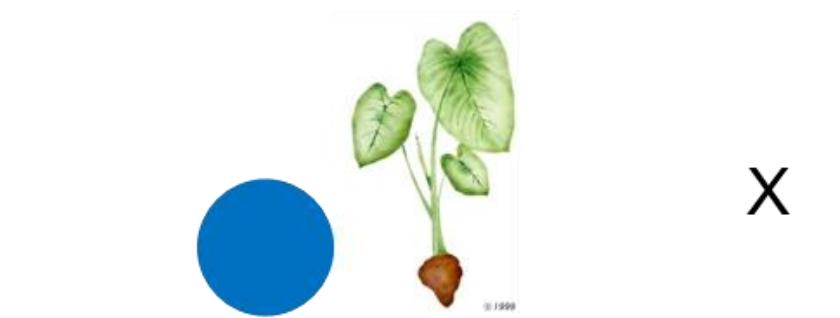




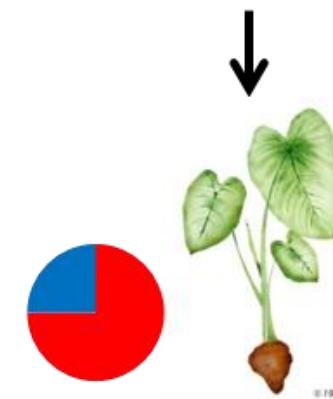
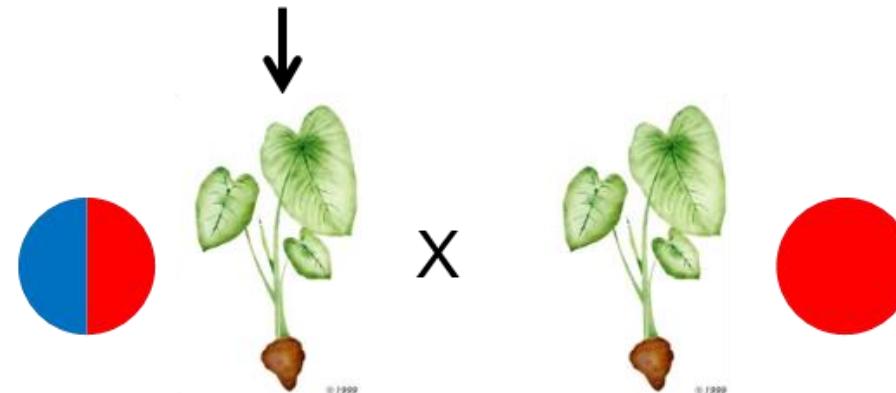


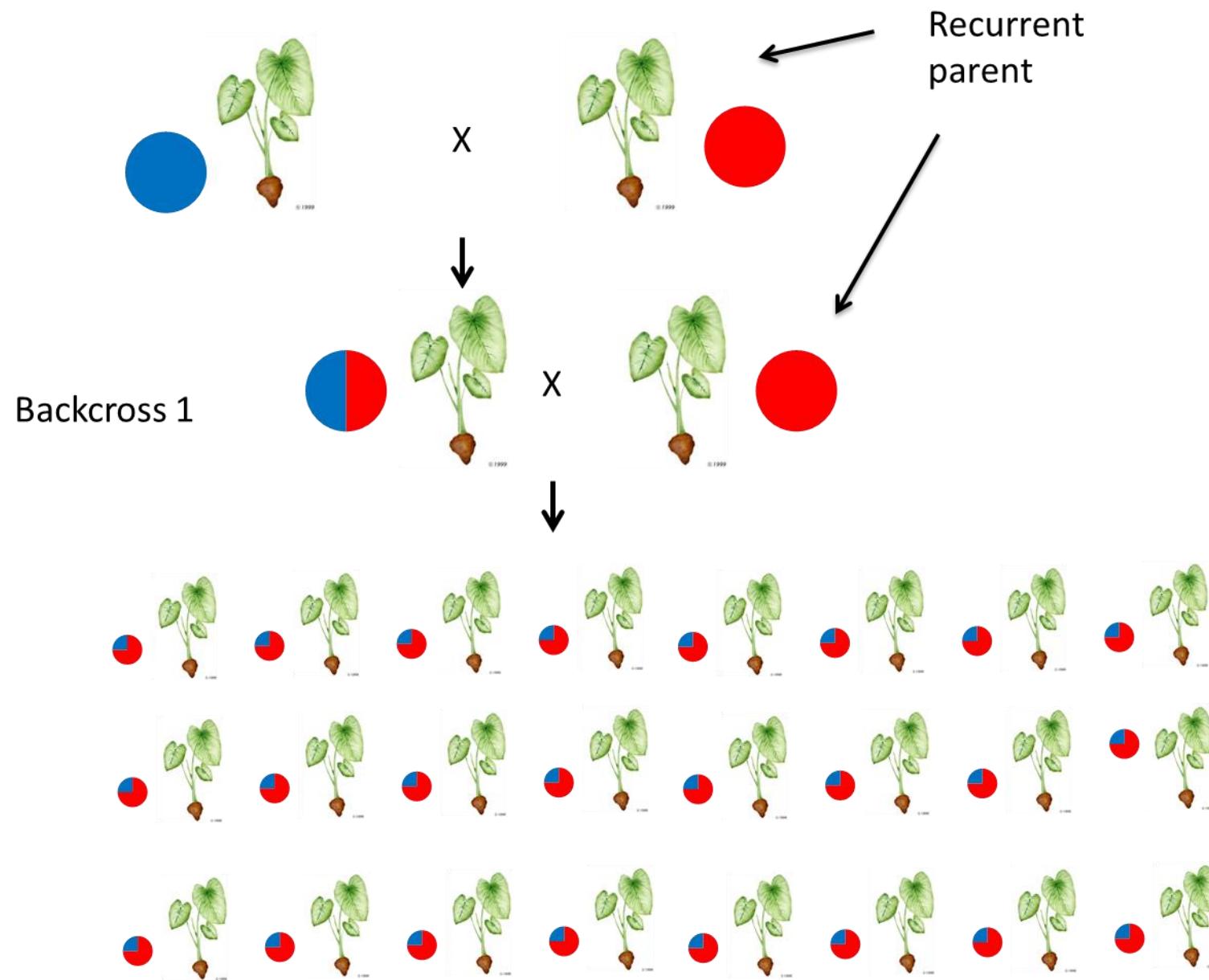


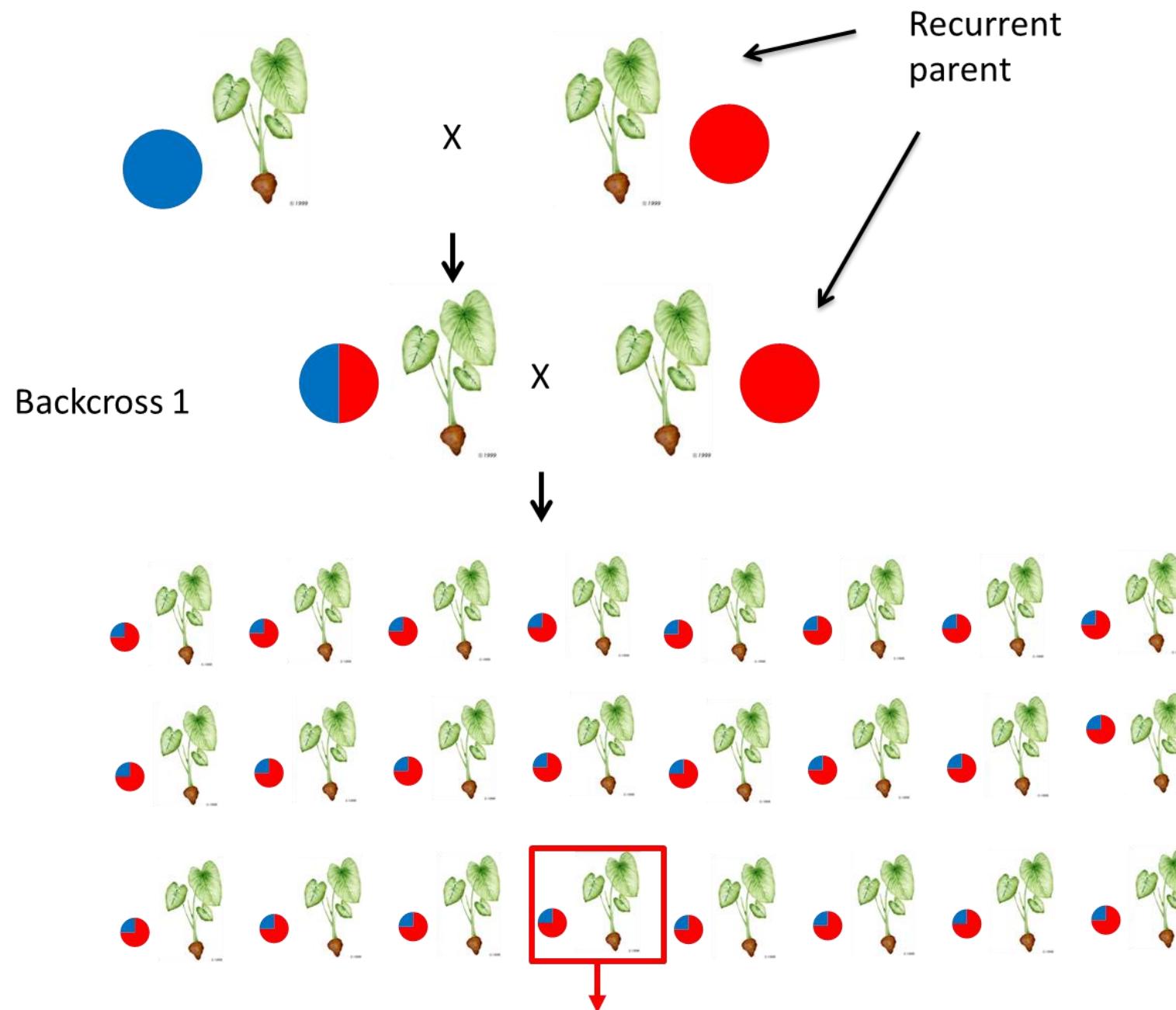
Recurrent  
parent

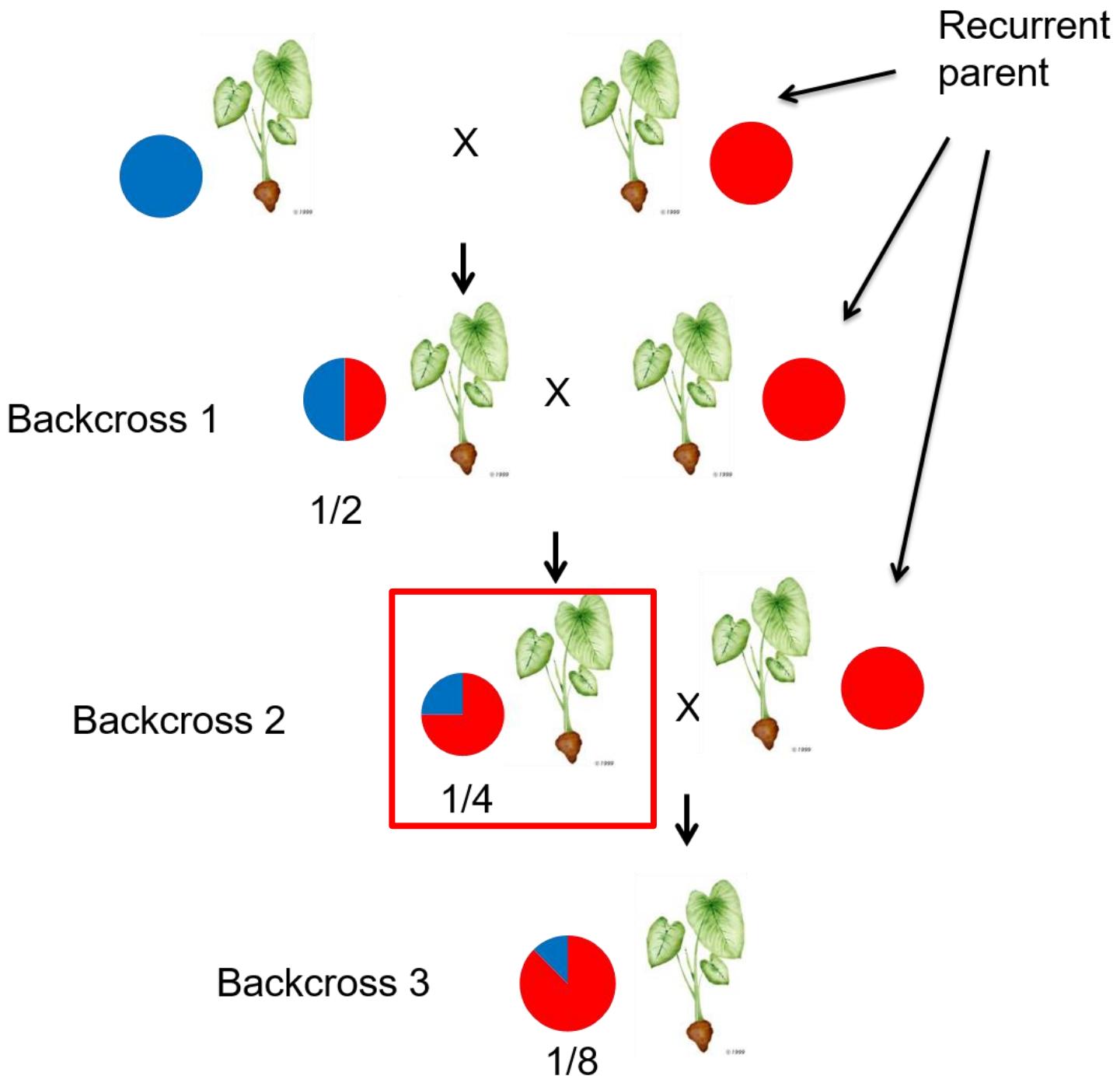


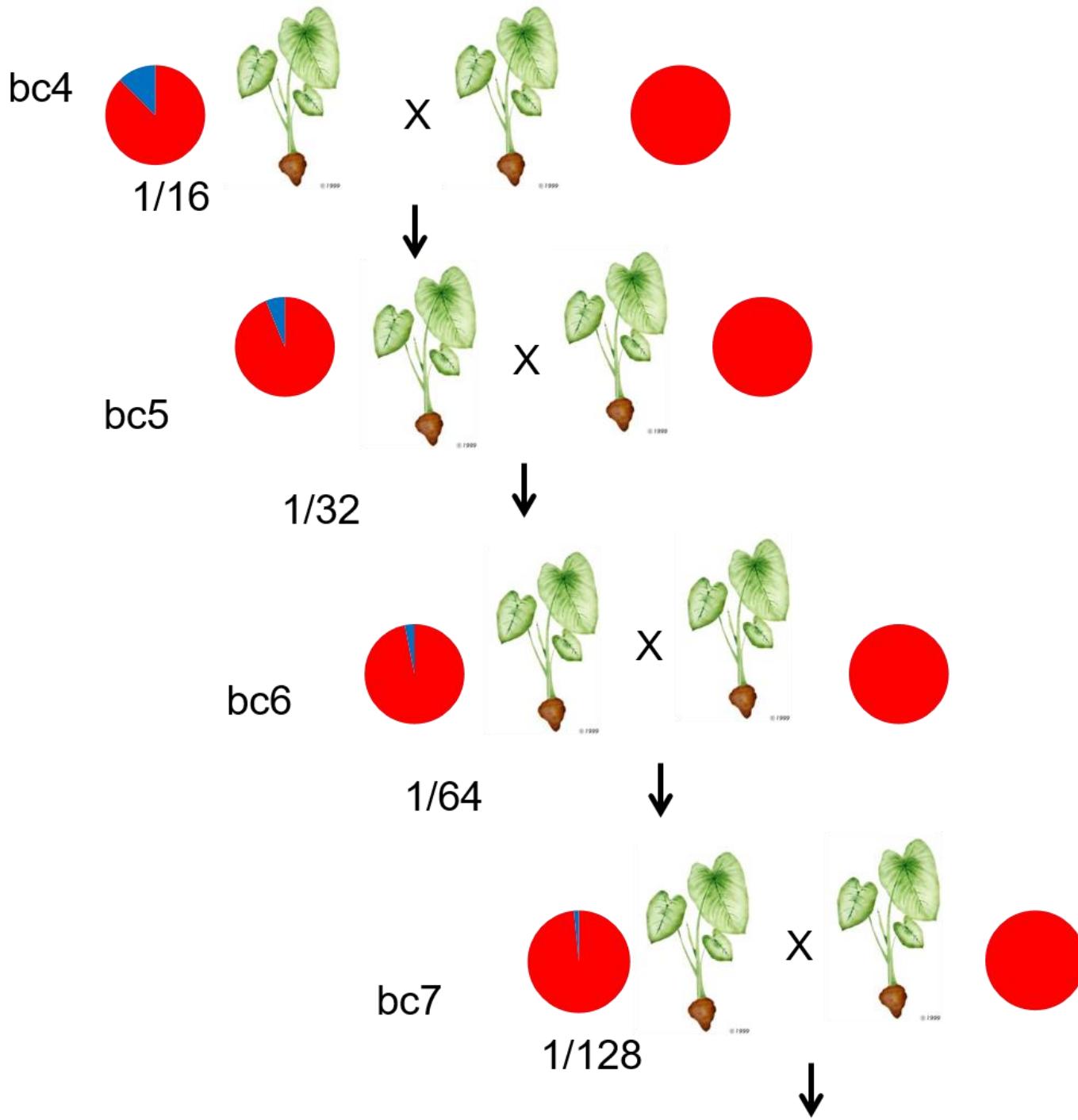
Backcross 1











# DNA Sequencing

- Extracted DNA from
  - 73 Hawaiian varieties
  - 12 hybrids
  - 6 Palauan
- Over 220 million DNA reads,  
about 100 bases each

- Over 220 million DNA reads, about 100 bases each

A diagram illustrating a single DNA sequence read. At the top left, the sequence ACGTACCC is shown in large black letters. A blue line extends from the right side of this sequence down to a much longer sequence below it. The long sequence is composed of many smaller black letters representing individual bases. A horizontal blue bracket is positioned under the long sequence, spanning its entire length. Centered below this bracket is the text "100 bases".

ACGTACCC

ACGTACCGTTACCAGAGAGGCCATGCCACGGTACTACGGGCCAAACCGCCACGTACGTTACCAGAGAGGCCATGCCACGGTACTACGGGCCAAACCGCC

100 bases



ACGTACCGTTTACCAAGAGAGGCCATGCCACGGTACTACGGCCAAACCGCACGTACGTTA...



ACGTACCGTTACCAAGAGCCCATGCCACGGTACTACGGCCAAACGCCACGTACGTTA...  
ACGTACCGTTACCAAGAGCCCATGCCACGGTACTACGGCCAAACGCCACGTACGTTA...



ACGTACCGTTACCAAGAGCCCCATGCCACGGTACTACGGGCCAACCGCCACGTACGTTA...  
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AAACCGCCAAACCGCCAAACCGCCAAACCGCCAAACCGCCAA



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# SNPs = Single nucleotide polymorphisms



5' -GGCAACACATTGCC- 3'

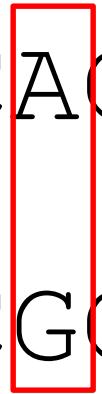


5' -GGCAACGCATTGCC- 3'

# SNPs = Single nucleotide polymorphism



5' -GGCAACACATTGCC-3'



5' -GGCAACGCGATTGCC-3'



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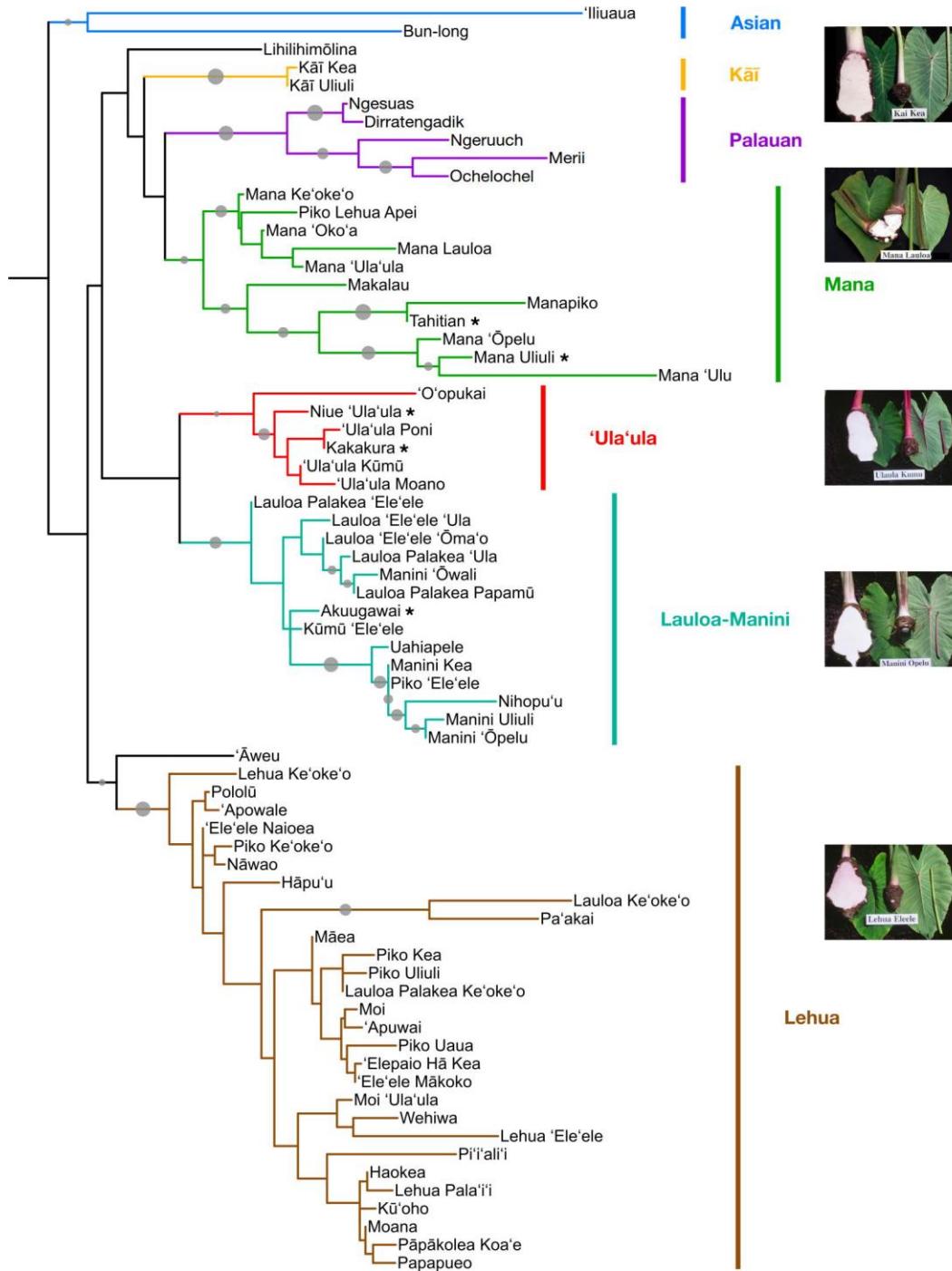


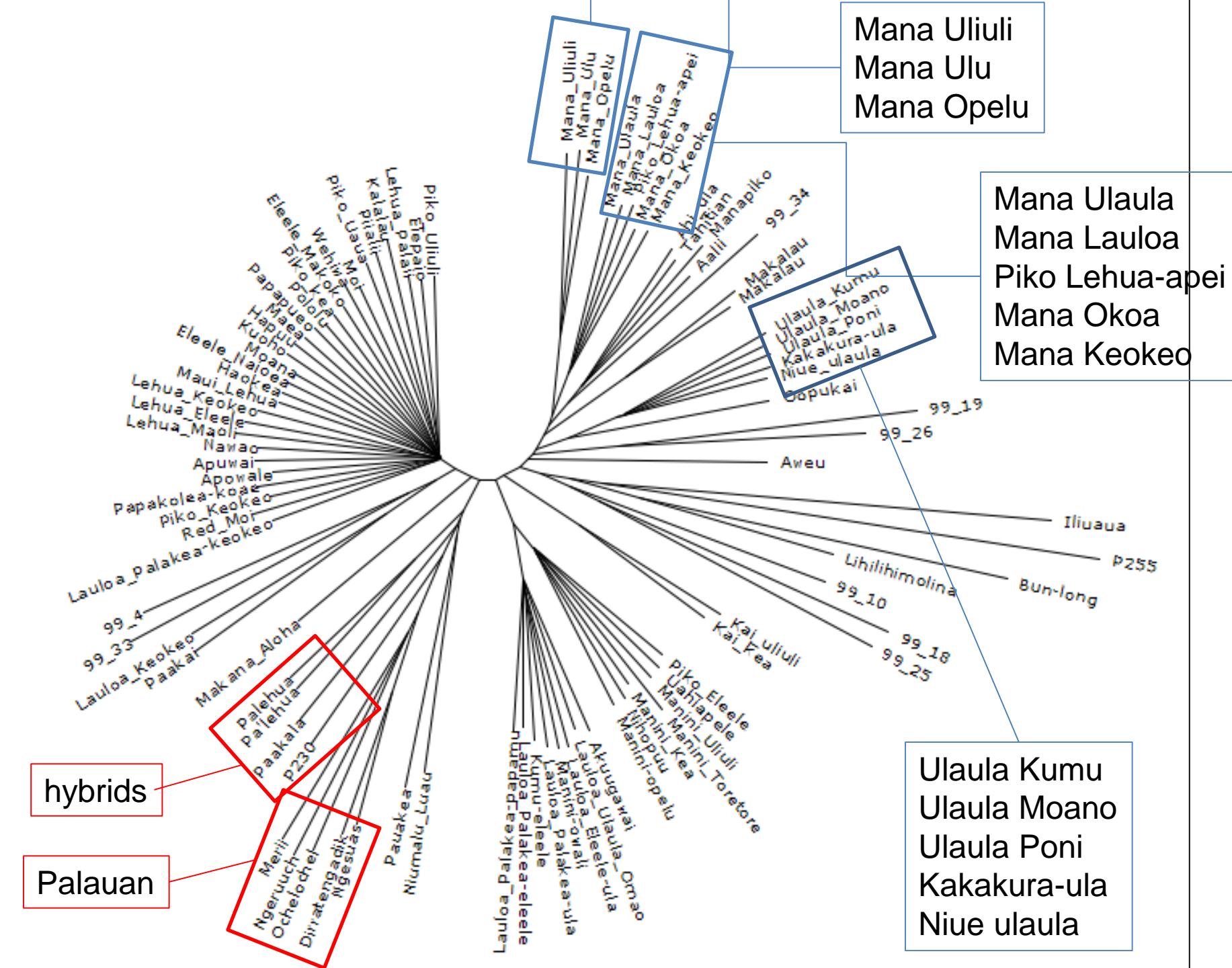
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CGGTAA



CGTACCGTTACCAAGAGGCCATGCCACGGTACTACGGCCAAACGCCACGTA  
..CGGGCCAAACCGCCACGTACGTTA...

Tree scale: 0.01





## **Acknowledgements:**

Susan C. Miyasaka UHM-CTAHR

Christopher (Popo) Bernabe UHM-CTAHR

Michael B. Kantar UHM-CTAHR

Martin Helmkampf UHH

Thomas K. Wolfgruber UHM-CTAHR

M. Renee Bellinger UHH

Roshan Paudel UHM-CTAHR

Heather Kimball UHH

Ashley Brown UHH (currently Quidel Corp., Ohio)

Anne Veillet UHH

Andrew Read UHH (currently Cornell University)

Jerry Konanui