



Accurate sweetpotato phenotyping of agronomic traits

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The 15th Sweetpotato Breeders Meeting,
June 6-7, 2016
BecA-ILRI Research Platform, Nairobi, Kenya

SWEETPOTATO ACTION FOR SECURITY AND HEALTH IN AFRICA

How do we measure genetic diversity



- **Morphological/ phenotypic markers**
- Biochemical markers
- Molecular markers

Variety/Clonal identification

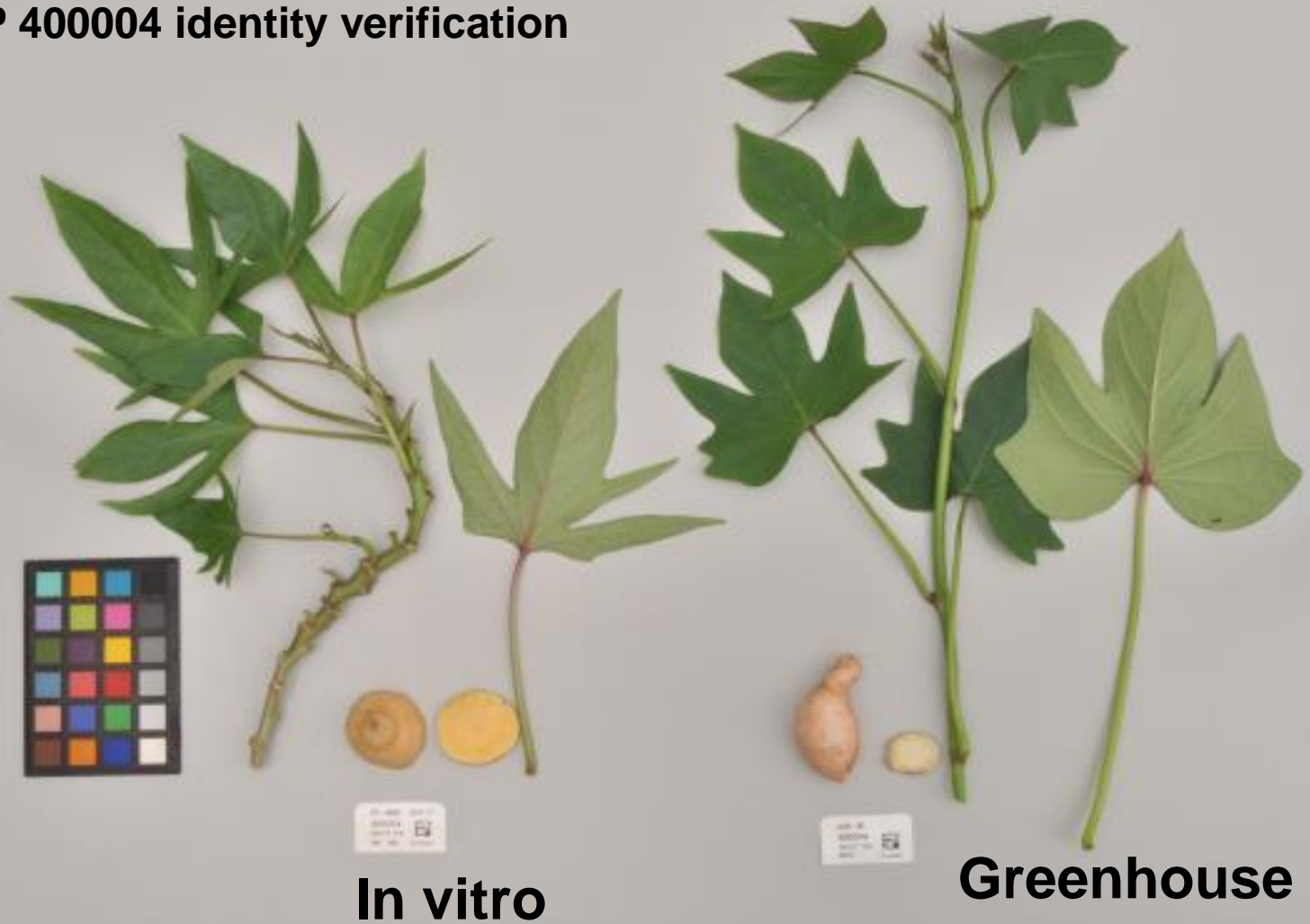


Why is it important?

- Outcrossing mating system
- Somatic mutation events
- Exchange system at local and regional levels

Characterization of morphological diversity is restricted to germplasm bank collections

CIP 400004 identity verification



In vitro

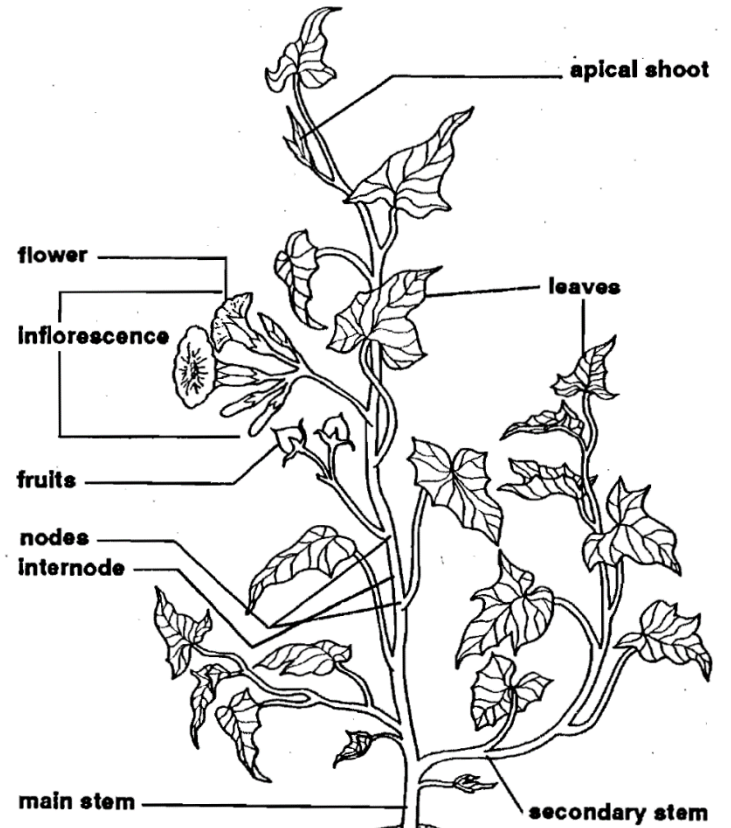
Greenhouse



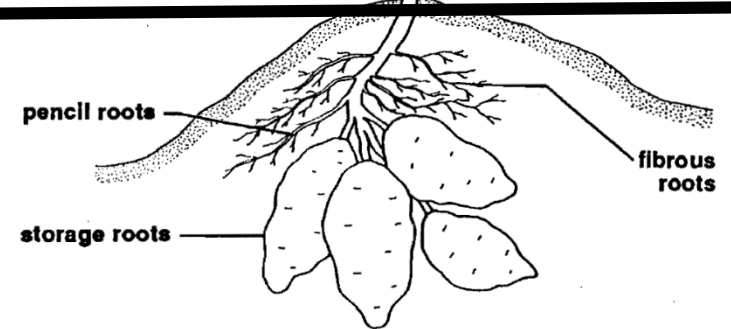
Figure: Leaf and root cross section: L “Orange-Chingova” and R Orange-Chingova

Morphological/Phenotypic

Vegetative aerial descriptors



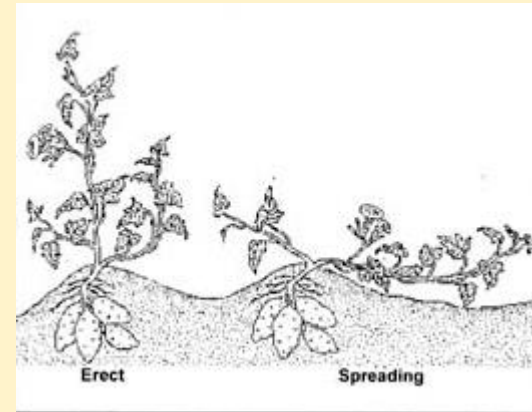
Storage root traits



Key sweetpotato descriptors (plant data)

Plant growth habit

- Erect (<75cm)
- Semi-erect (75-150 cm)
- Spreading (151-250 cm)
- Extremely spreading (>250 cm)



Twining (ability)

- Non-twining
- Slightly twining
- Moderately twining
- Twining
- Very twining

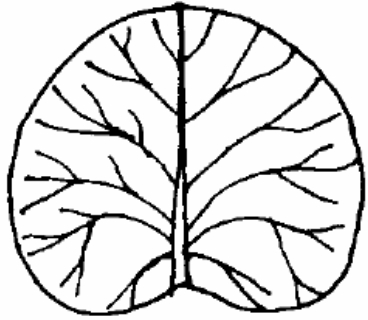
Ground cover

- Non-twining
- Slightly twining
- Moderately twining
- Twining
- Very twining

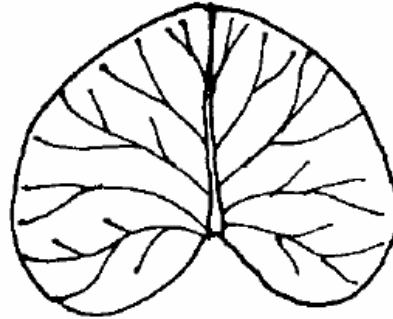
Vine

- Vine internode length
- Vine internode diameter
- Predominant vine colour
- Secondary vine colour

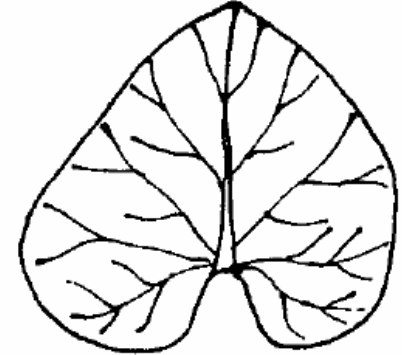
General outline of leaf



1 Rounded



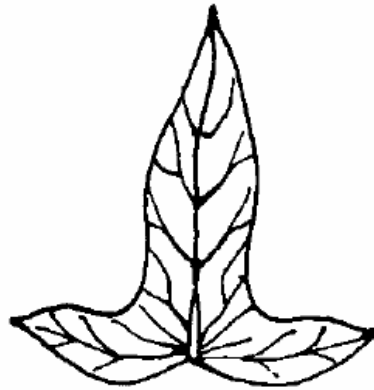
2 Reniform



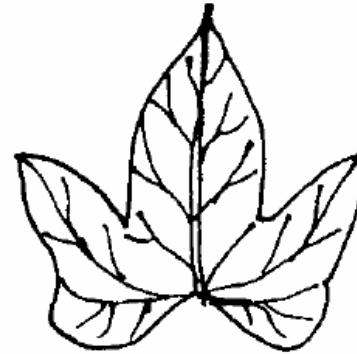
3 Cordate



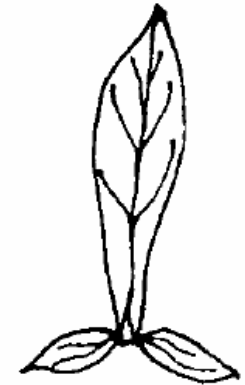
4 Triangular



5 Hastate

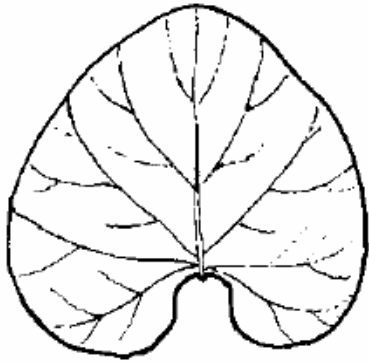


6 Lobed

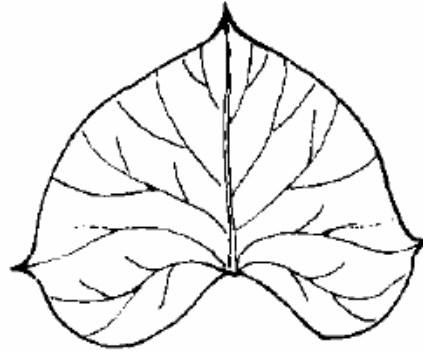


7 Almost divided

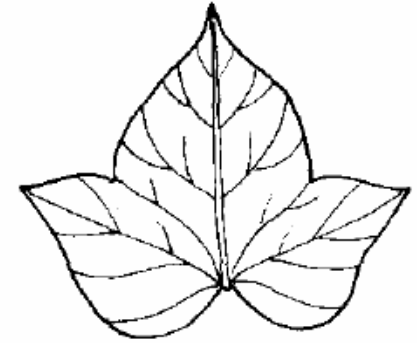
Leaf lobe type



0 No lateral lobes



1 Very slight (teeth)



3 Slight



5 Moderate

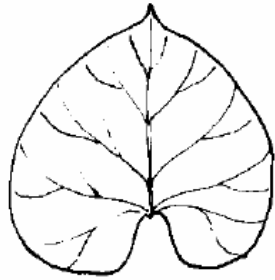


7 Deep

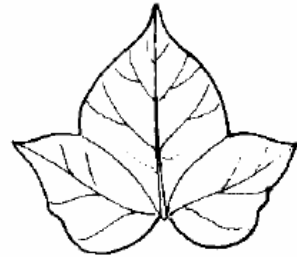


9 Very deep

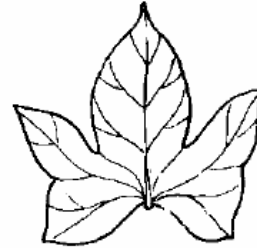
Leaf lobes number



1



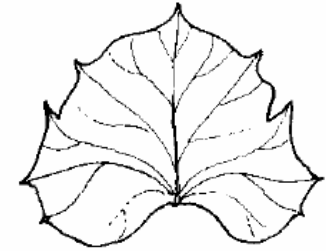
3



5

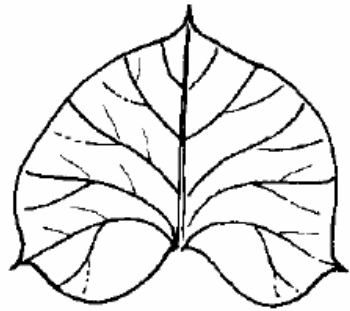


7

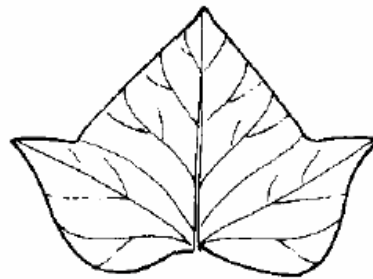


9

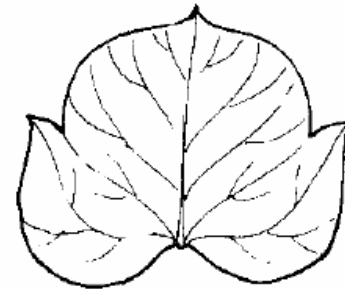
Shape of central leaf lobe



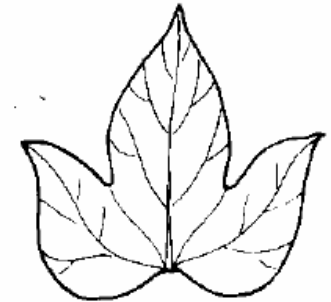
1 Toothed



2 Triangular



3 Semi-circular



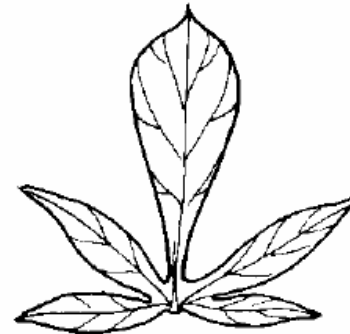
4 Semi-elliptic



5 Elliptic



6 Lanceolate



7 Oblanceolate



9 Linear (narrow)

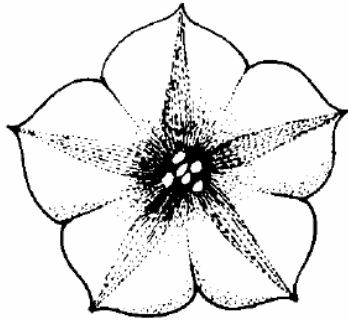
Other leaf attributes



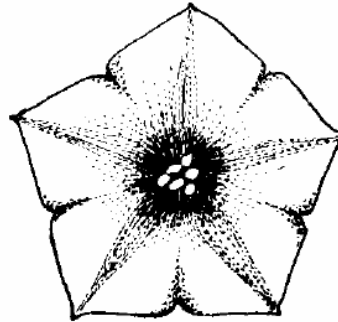
- Mature leaf size
- Abaxial leaf vein pigmentation
- Mature leaf colour
- Immature leaf colour
- Petiole length
- Petiole pigmentation

Inflorescence

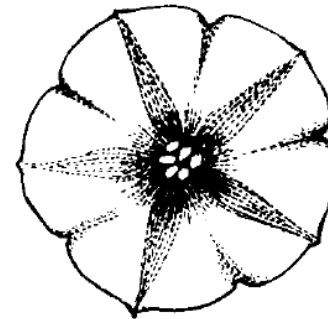
- Flower habit
- Flower color
- Flower size-length and width
- Shape of limb



3 Semi-stellate



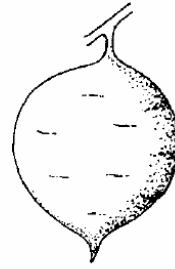
5 Pentagonal



7 Rounded

Storage root

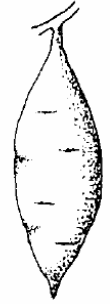
- Storage root shape
- Skin color
- Storage root flesh color
- Type root formation



1 Round



2 Round elliptic



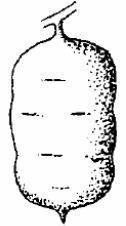
3 Elliptic



4 Ovate



5 Obovate



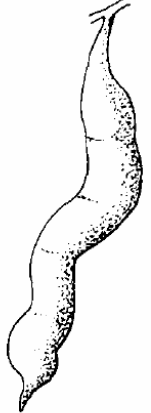
6 Oblong



7 Long oblong

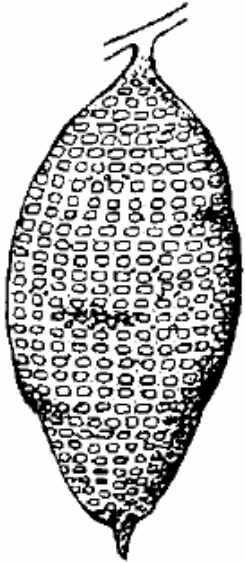


8 Long elliptic

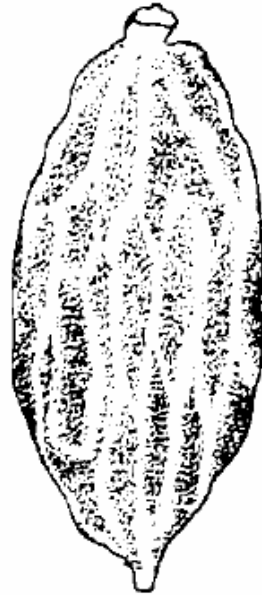


9 Long irregular or curved

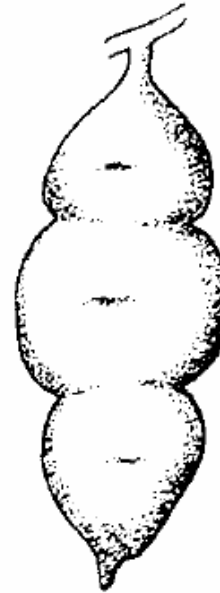
Storage root surface defects



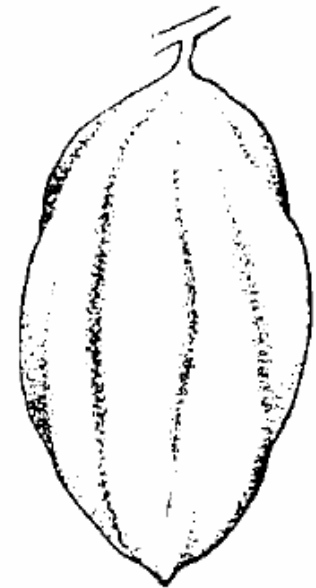
1 Alligator-like skin



2 Veins



3 Horizontal constrictions



5 Longitudinal grooves

Other storage root attributes



- Storage root dry matter content
- Storage root nitrogen content
- Storage root starch content
- Storage root total alcohol soluble sugar content
- Storage root carotene content
- Consistency of boiled storage root
- Texture of boiled storage root flesh

Abiotic Stress

- Reaction to drought
- Reaction to flooding
- Reaction to heat
- Reaction to salinity

Biotic Stresses

- Sweet potato weevil (*Cylas* spp.)
- Root-knot nematode (*Meloidogyne* spp.)
- Fusarium wilt or stem rot (*Fusarium oxysporum* f. sp. *batatas*)
- Black rot (*Ceratocystis fimbriata*)
- Java black rot (*Diplodia gossypina*)
- Scab or spot anthracnose (*Elsinoe batatas*)
- Charcoal rot (*Macrophomina phaseoli*)
- Bacterial stem and root rot (*Erwinia chrysanthemi*)
- Sweet potato virus disease (SPVD complex)
- Sweet potato chlorotic stunt virus (SPCSV)

Overcoming Bottle-Necks



Information management

- Standardized data formats
- Tools for accurate, field-based phenotyping
- Data repositories (distributed centralized) - CIPTCL
- Data-analysis & visualization tools
- Support & assistance

Conclusion



- Genetic diversity is basic ingredient of all breeding & conservation of plant genetic resources
- Bridging two worldviews
 - Genomes & genes = Germplasm & traits

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