

Genetic diversity of wild Coffee (*Coffea arabica*) and its implication for conservation

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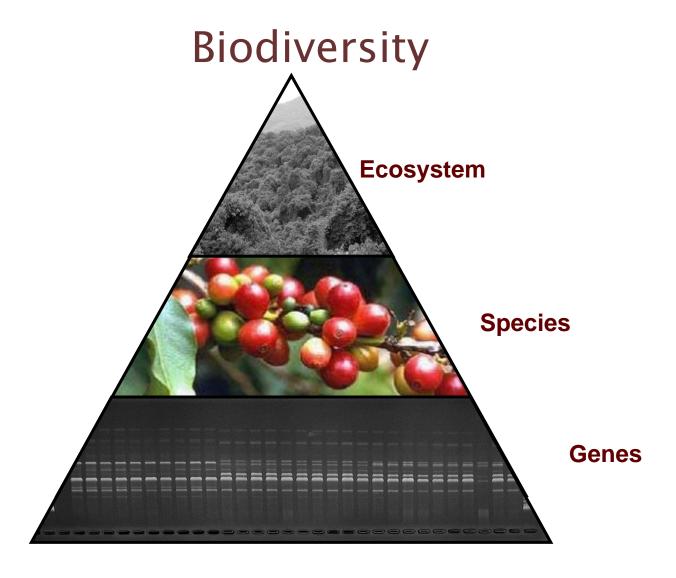














Objectives:-

- To clarify the maternal parentship among closely related *Coffea* species.
- To characterize cpDNA microsatellites & to detect SNPs within *C. arabica* to observe infraspecific variation among wild populations *C. arabica*.
- To evaluate if individuals from wild populations in forests are genetically different from landraces.
- To evaluate the relationships of wild C. arabica populations throughout Ethiopia & also to understand the patterns of distribution.



Chloroplast DNA
Microsatellites
-High level-

Interregional
Genetic Diversity
-Medium level-

Intraregional
Genetic Diversity
-Low level-







Genetic Diversity of *Coffea arabica* in Ethiopia

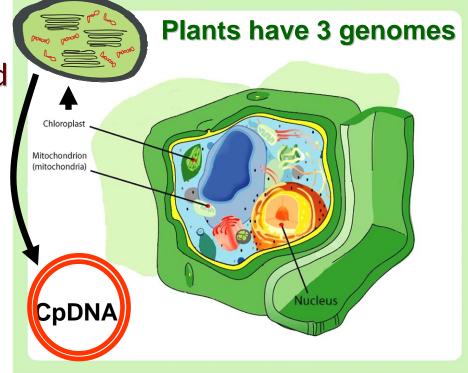


Conservation and Use Planning



Chloroplast DNA Microsatellites

- CpDNA is maternally inherited
- Intron & spacer usually variable within species
- Contain hyper-variable microsatelites

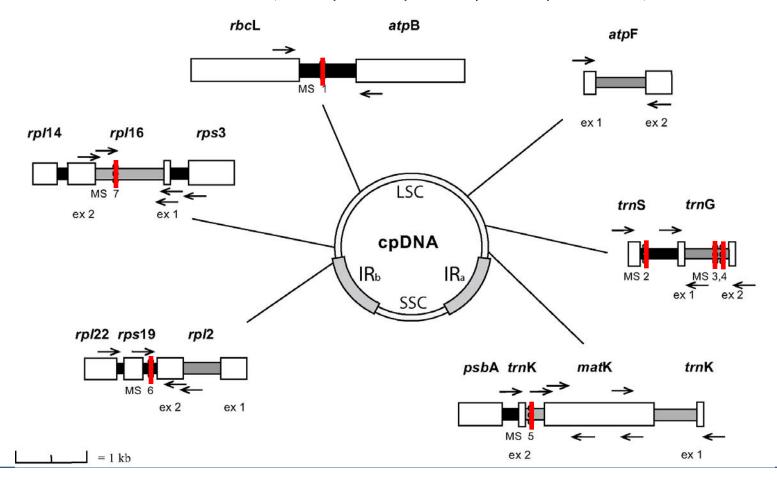


 To explain the genetic diversity in an evolutionary context



Chlorplast regions selected

- Screening cpDNA for its most variable regions for 17 taxa (5.5 %)
- Seven microsatellites (MS1, MS2, MS3, MS5, & MS7)

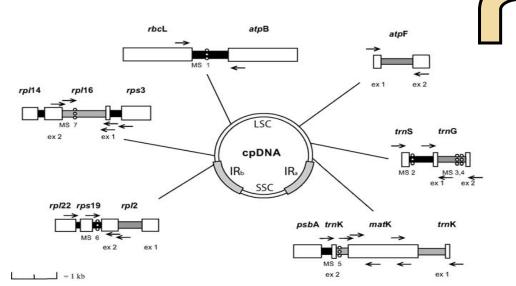


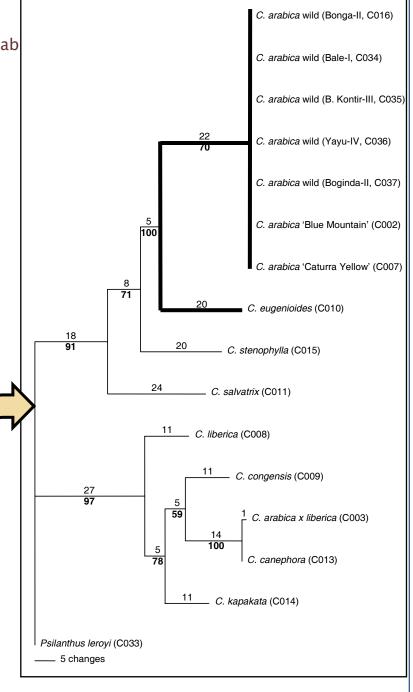


Conservation and Use of Wild Populations of Coffea arab

- 7.2 kb of cp-genome sequenced for 17 taxa
- C. arabica exhibits no cp-genome diversity (unlike most plants!)
- Very recent origin

• C. eugenioides sister to C. arabica







Summary of Chloroplast data

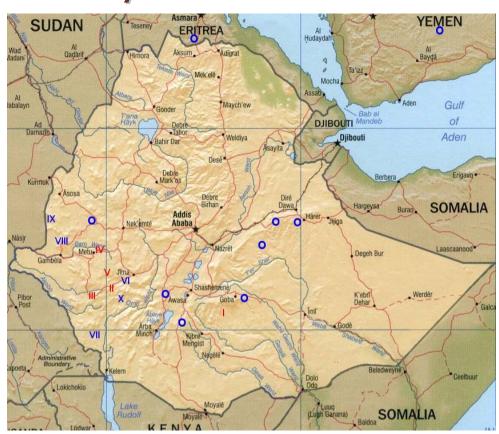
- An ancestor of C. eugenioides is evidenced to be the maternal parent for C. arabica.
- Both *C. arabica* & *C. eugenioides* are recently diverged within *Coffea*.

C. arabica has a single origin (one polyploidisation event).



Interregional diversity assessment

- 10 regions of wild coffee
- 1 sample per landrace
- 150 individuals



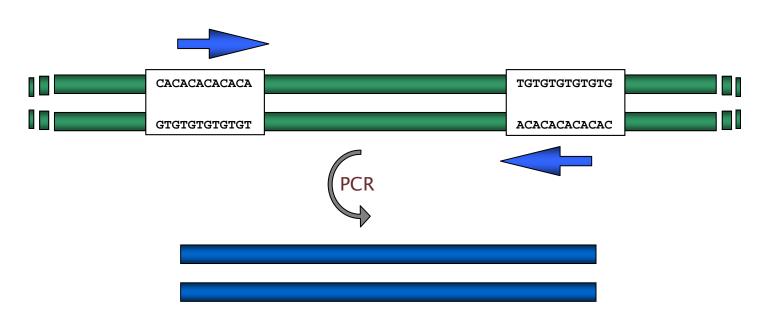
I to IV - Core CoCE sites (wild Coffee)

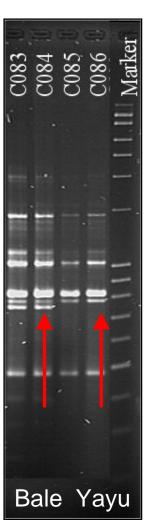
V to X - Additional wild Coffee sites

- Landraces (Farmer's varieties)



• Inter Simple Sequence Repeats-ISSR







Conservation and Use

Interregional NJ tree

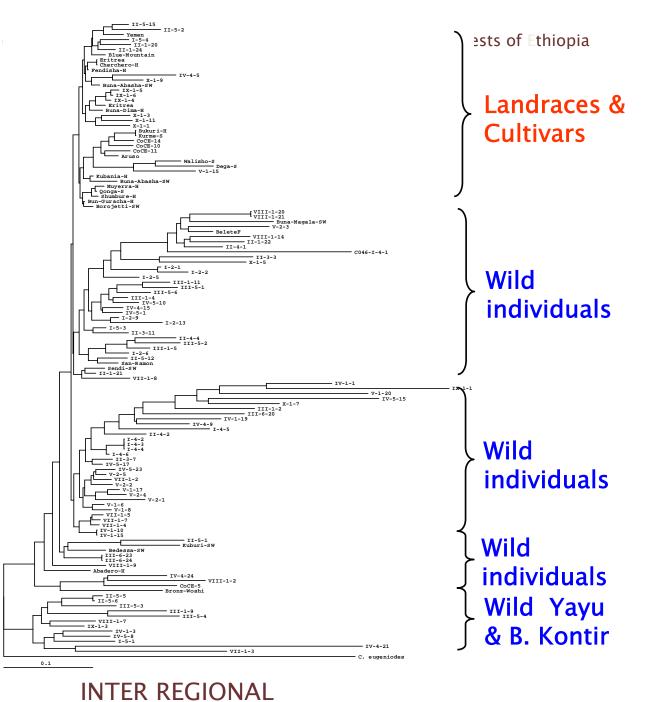
Landraces & Cultivars

different

from

Wild

individuals





Conservation and Use

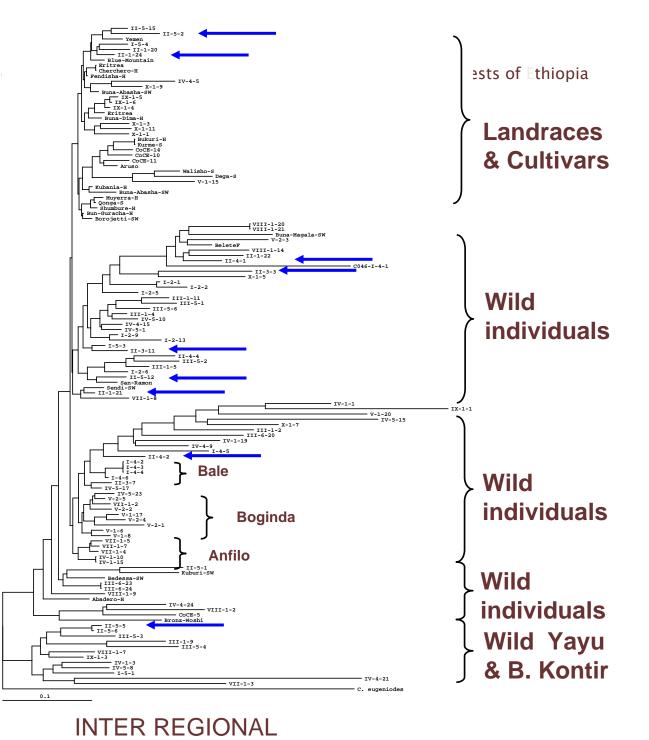
Interregional NJ tree

Bonga

Spread all over the tree

Boginda, Bale/Harrena & Anfilo;

Form its own groups





Shannon's diversity index (H) & percentage of ISSR band polymorphism (P)

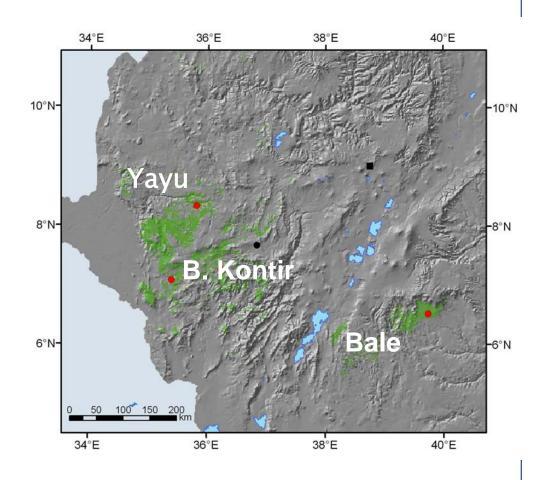
Region	Percent Polymorphism (%P)		Over all %P	Shannon's diversity Index (I)		Over all Index (I)
	Di/tri_ Primers	Tetra_ Primer	All primers	Di/tri_ Primers	Tetra_ Primer	All primers
Bale (Harrena)	13.7	80.7	27.7	0.18	0.53	0.25
Bonga	28.2	93.6	41.9	0.31	0.71	0.40
Berhane Kontir	20.5	87.1	34.5	0.28	0.56	0.34
Yayu/Geba Dogi	20.5	93.6	35.8	0.30	0.76	0.40
Boginda	16.3	83.9	30.4	0.18	0.59	0.27
Bench Maji	5.1	67.7	18.2	0.13	0.54	0.22
Anfilo	7.7	67.7	20.3	0.21	0.53	0.28
Daphe	2.6	71.0	16.9	0.17	0.60	0.25
Mankira	10.3	83.9	25.7	0.20	0.75	0.31
Sidamo	7.7	54.8	17.6	0.14	0.46	0.21
Hararegh	10.3	80.7	25.0	0.08	0.49	0.17
South West	15.4	58.1	24.3	0.17	0.40	0.22

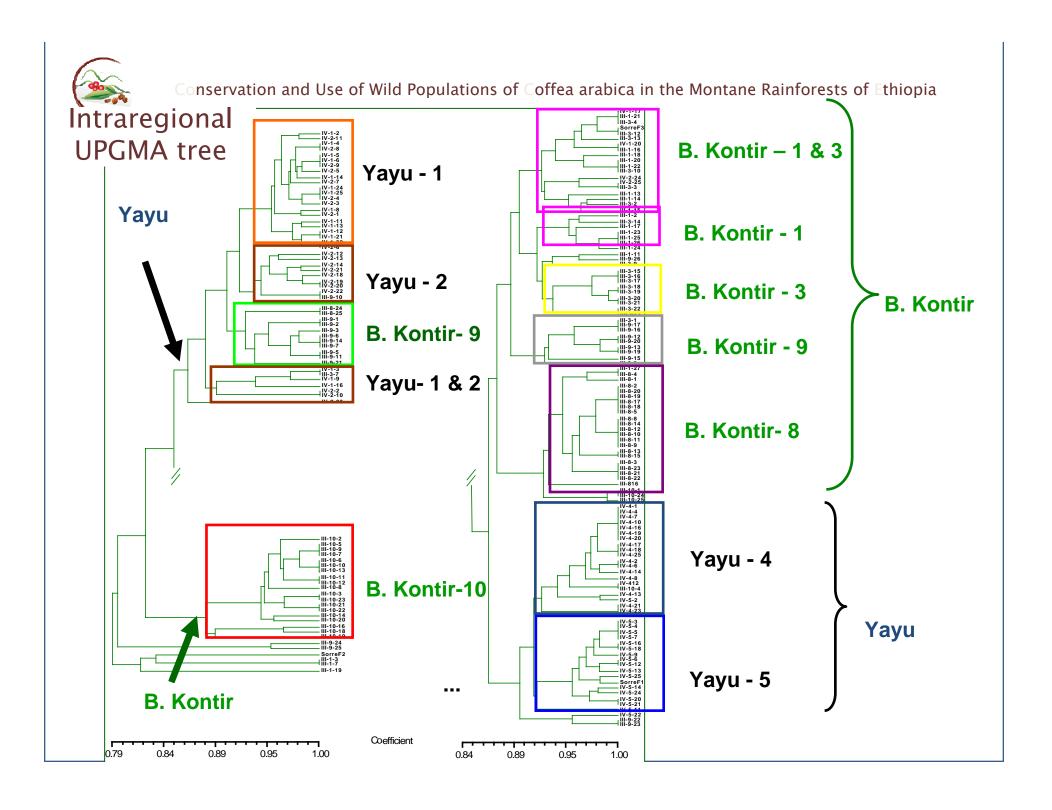
INTER REGIONAL



Intraregional diversity assessment

- Three regions
 (Yayu, B.Kontir & Bale)
- 250 individuals (Yayu & B.Kontir)
- ISSR marker system







ISSRs show that: -

- Wild coffee populations can be clearly distinguished from landraces "true wild coffee".
- High levels of genetic diversity within geographical regions and vary from region to region.
- Even though Bonga, Yayu and Berhane Kontir showed higher diversity indices, it is observed that most regions have their unique genotypes.
- Individuals belonging to landraces often have lower genetic distances to each other than individual of wild populations.
- There is fine scale spatial genetic structuring among populations of Berhane Kontir and Yayu.
- Semidisturbed plots tends to show higher diversity than undisturbed.



Recommendations for conservation & use planning

- Genetic evidence for true wild coffee Justify conservation
- Unique genotypes in different regions Multi-site in situ conservation approach
- The existing genetic divergence among regions –
 Prospects for use in national breeding (heterosis)



Implementation-oriented research activities

- Establishing coffee in situ gene bank:-
- The aim is to have a 'copy' of the genetic diversity in the natural populations at the forest margin.
- To re-introduced elite breeding materials originally came from the same forest into the forest margin to increase the yield and income of the local farmer.

 In situ gene bank have both conservation and use value for the local farmers who are fully dependent on wild coffee.



In situ gene bank-Yayu

Naturally regenerated seedlings were collected from different forest population









In situ gene bank

 Totally 5,000 coffee seedlings were disseminated for the 3 farming community-Year 2007









Seeds of forest tree species like *Cordia africana, Albizia* grandibracteata, Albizia gummifera, Acacia abyssinica, & Acacia mellifera were collected and raised in Nursery

-2,000 seedlings were disseminated









*Local Youth Association

Agreement were made with Youth Association organized to manage of the forest

- 10,000 seedlings of coffee along with 6,200 tree species were given for the association in Yayu.





Leaders of the Association





Selection of elite individual coffee for re-introduction to the forest margin -Jima Agricultural Research Center















Conservation and Use of Wild Populations of Coffea arabica in the Montane Rainforests of Ethiopia















