



VII International Congress on Cactus & Cochineal
Agadir, Morocco, October 22th 2010



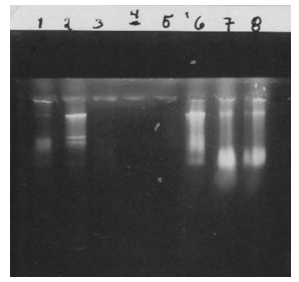
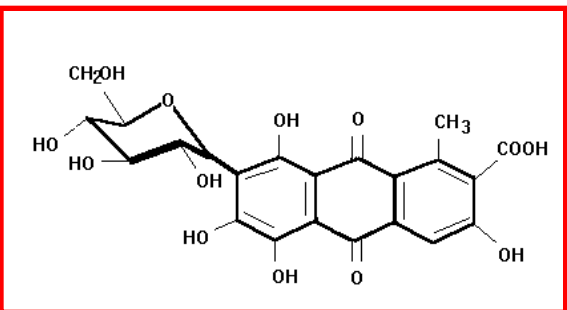
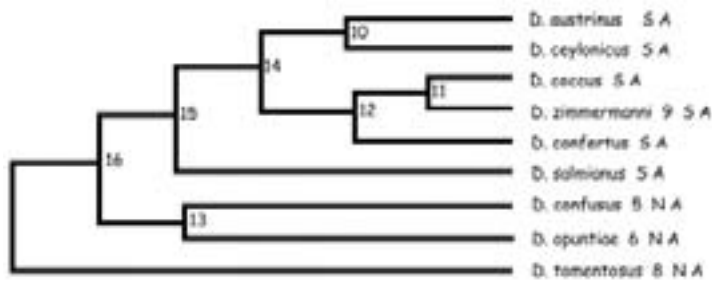
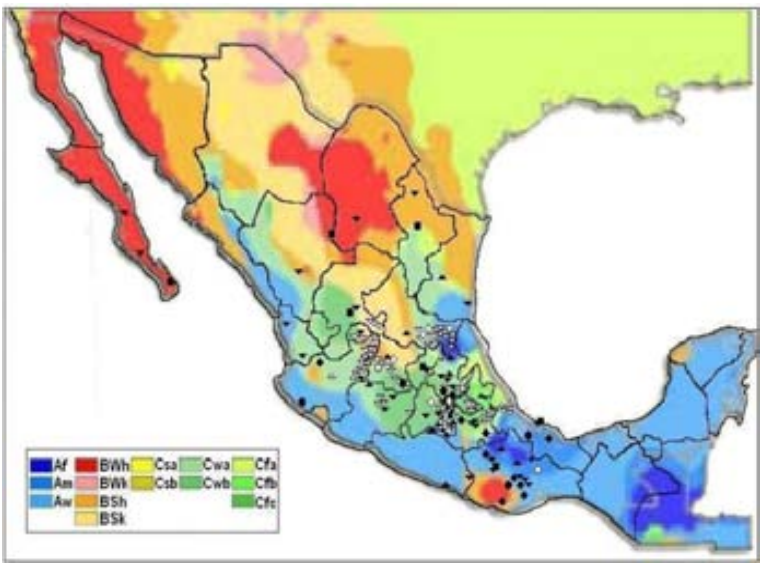
“The *Opuntia* (Cactaceae) and *Dactylopius* (Hemiptera: Dactylopiidae) in Mexico: a historical perspective of use, interaction and distribution with particular emphasis on chemical and phylogenetic aspects of the *Dactylopius* species”

Dr. Carla Karina Chávez Moreno
UNAM - UMSNH

Plant-insect: *Opuntia* – *Dactylopius* represents a **great challenge study because morphological complexity.**

The integration anthropological, geographical, ecological, chemical and molecular analysis, represent important information for understanding biological system.

Combination of different knowledge contribute to establish policies of conservation and strategies for a sustentable use of these natural genetic resources, great significance to Mexico.



INTERACTION

Opuntia spp.

Dactylopius spp.

Historical, cultural, geographical, ecological and Distribution: Expansion

Ecological characteristics of the distribution and habitat in Mexico

Metabolic profile of the colorant

Molecular analysis: Insects and endosymbionts

I

II

III

IV

Specific measures for protection of such biodiversity and generic resources in Mexico.

Strategies for *in situ* conservation combined with re-established use and cochineal production may enhance conservation policies.

The *Opuntia* (Cactaceae) and *Dactylopius* (Hemiptera: Dactylopiidae) in Mexico: a historical perspective of use, interaction and distribution

C. K. Chávez-Moreno · A. Tecante · A. Casas

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***Opuntia* (L.) Miller** (Opuntioideae: Cactaceae)

American continent 200 species

Mexico 83-104 species

- 50% endemic
- 20 domesticated species, management and artificial selection was focus on
 - 1) optimize their use for edible stems & fruits.
 - 2) cultivation of *Dactylopius*.
- More than 900 names.



O. joconoxtle



O. phaeacantha



O. robusta



O. streptacantha



O. hypthiacanta

Dactylopius Costa (Hemiptera: Dactylopiidae)

- D. confusus* Cockerell
- D. opuntiae* Cockerell
- D. tomentosus* Lamarck
- D. coccus* Costa
- D. ceylonicus* Green
- D. austrinus* De Lotto
- D. confertus* De Lotto
- D. salmianus* De Lotto
- D. zimmermanni* De Lotto
- D. bassi* Targioni Tozzetti

Mexico

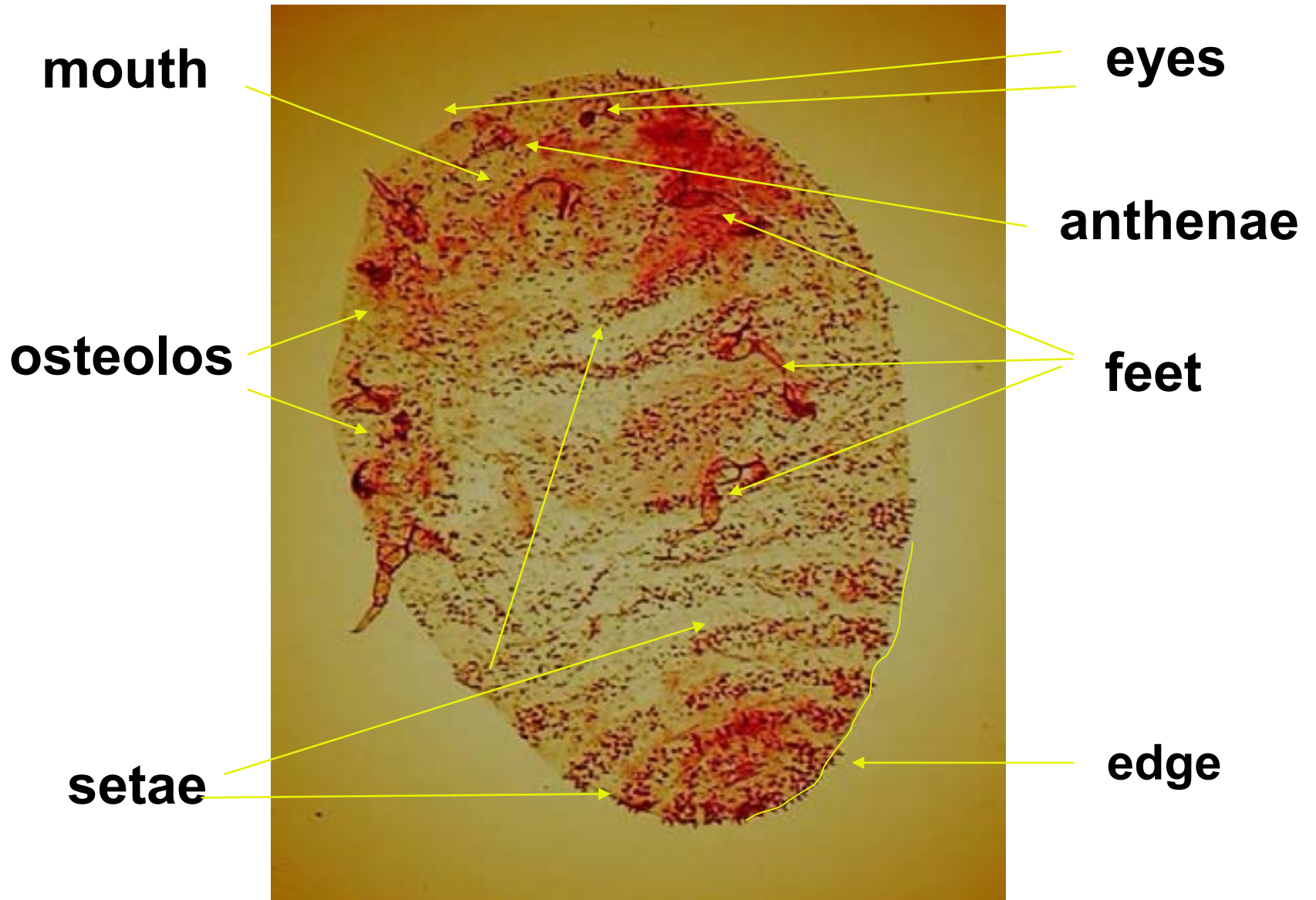
South
America



Feeding exclusively on cacti
~80 species hosts worldwide
22 in Mexico.



Possess very limited morphological characteristics



Opuntia – *Dactylopius*

Cactus ancestor South American center origin.

Molecular evidence 2 lineages:

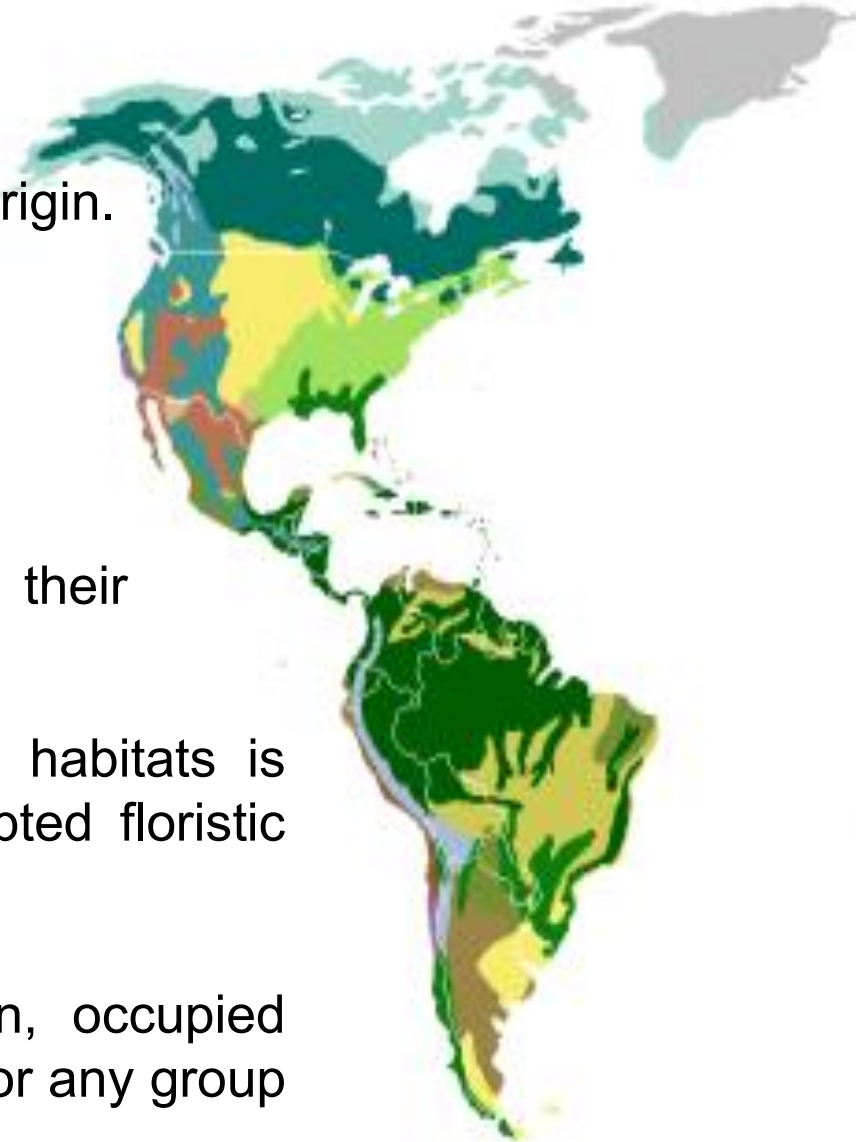
Cylindrical-stemmed

Flat-stemmed prickly pears,

Diverged in South America prior to their migration North, East & South America.

North-American desert and semi-desert habitats is result of parallel migration as xeric-adapted floristic cohort, later for *Dactylopius*.

Opuntia extensive evolutionary radiation, occupied the most widespread geographic region for any group within the Cactaceae (Anderson, 2001).



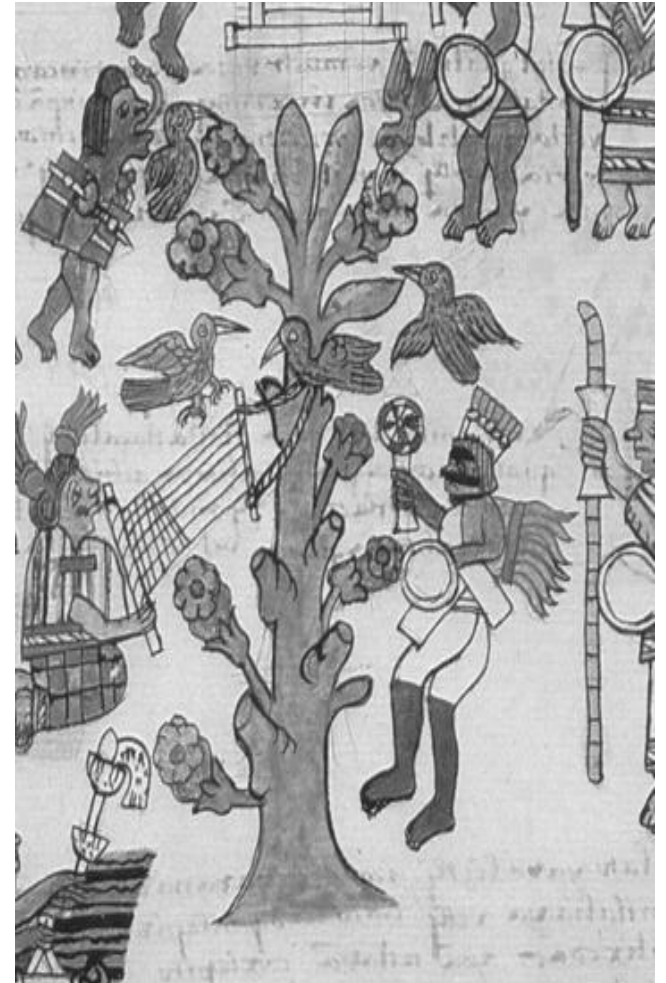
II. Historical use review

Opuntia main components of human diet during preagricultural times.
Gathering *Opuntia* data 12,000–14,000 years ago.

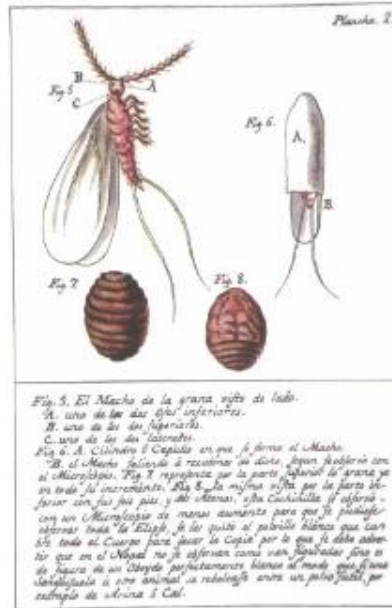


Cultivation more than 700 years, exhibited botanical gardens or iconographically represented on building walls.

Dactylopius. Rearing of cochineal, used and cultivated at least from X century, Toltec period, *amatl*.



Opuntia and *Dactylopius* production and marketing



Tribute to Aztecs, 394 communities Mixtec/4,400 kg/year

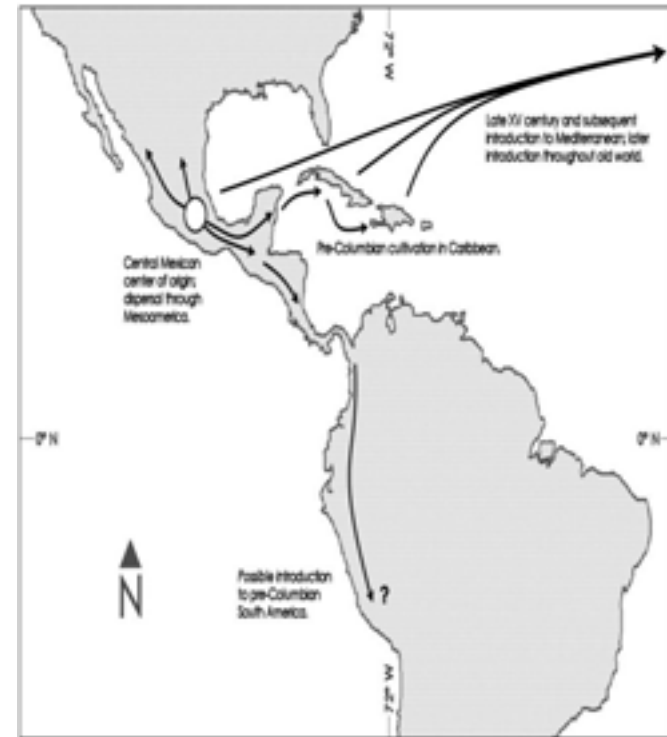
Prehispanic period

Mixtec production best quality dye, greatest commercial distribution.

Possible commercial and cultural links Mesoamerican and Andean exchange and propagation *Opuntia-D. coccus*.

Mexico and Peru shared use of dye from X-XII.

Chemical analyses Andean textiles (Peru, pre-Inca times) colored with *D. ceylonicus*, *D. confusus*, and *D. coccus* possible interchange materials & techniques.

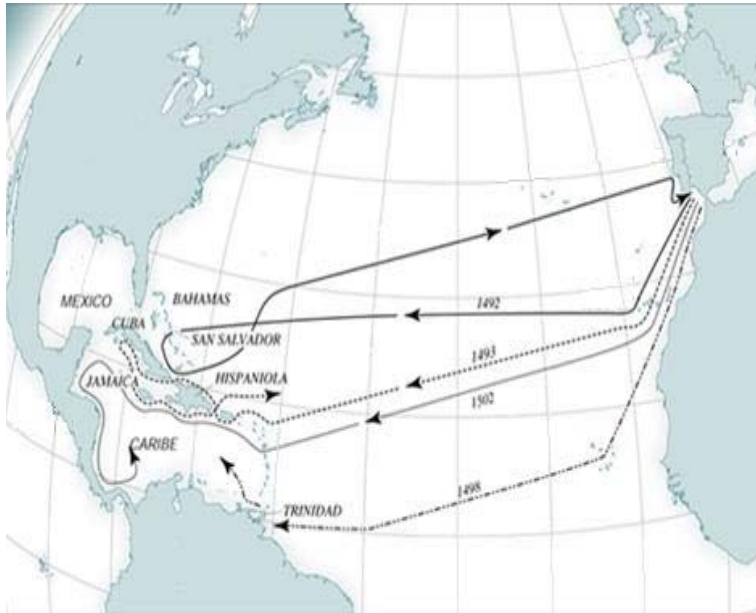


Conquest & Colony

1493 *Opuntia*,
Christopher Columbus
Lisbon



1523 *Dactylopius*,
Hernán Cortés
Spain

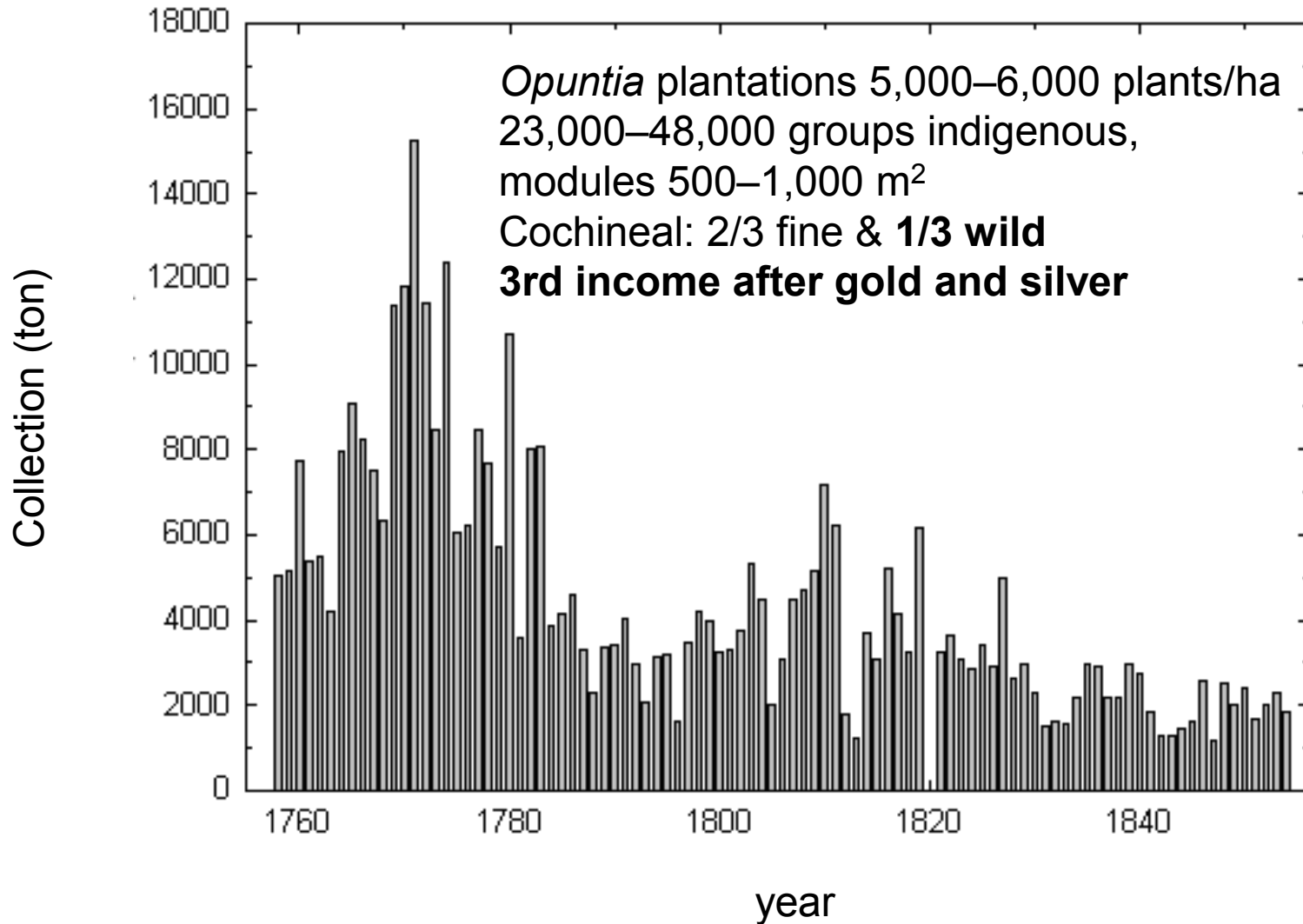


XVII-XVIII
America
Europe

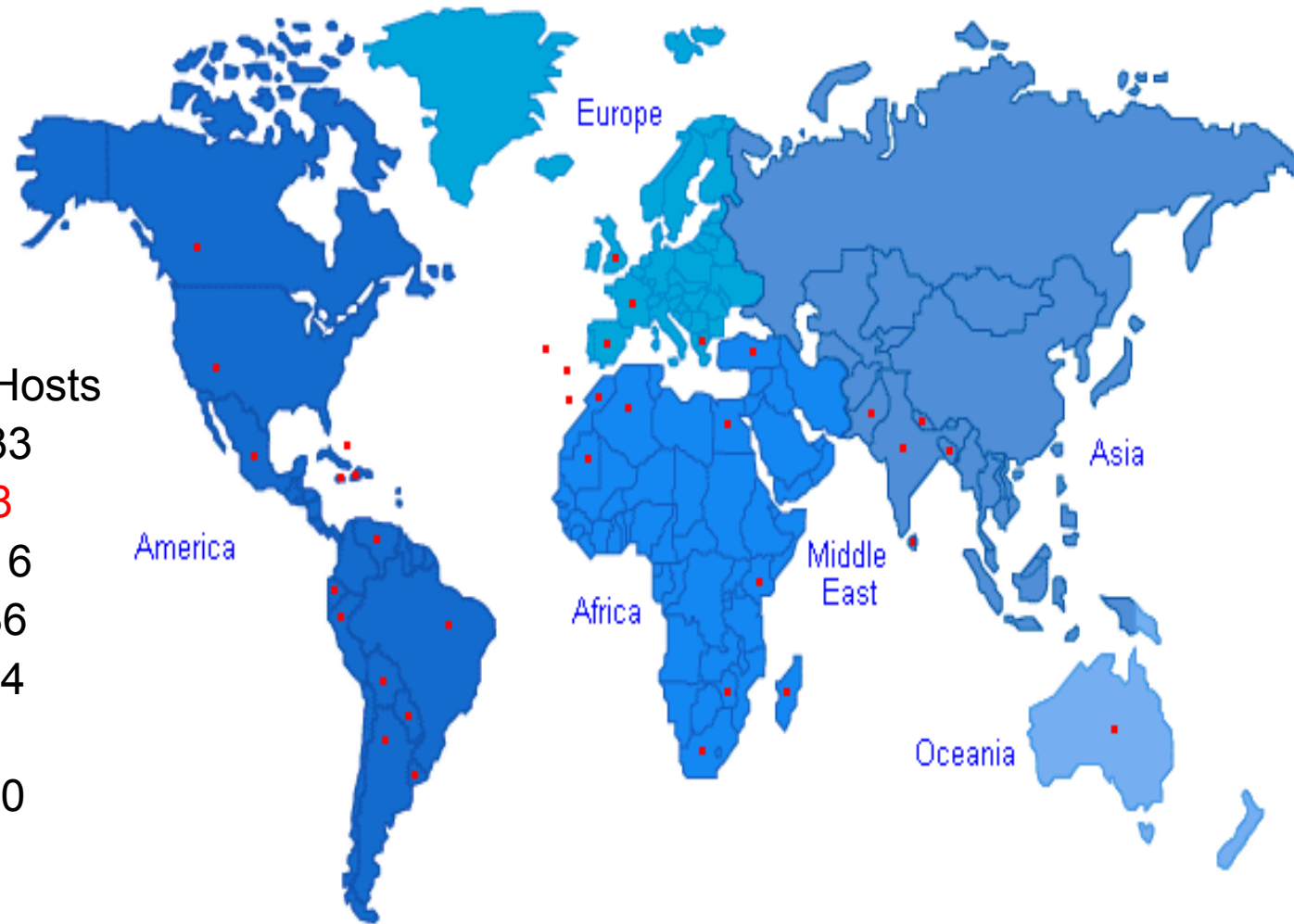


Collection grana cochineal Oaxaca (1760 to 1850)

1700 T/year



Current distribution

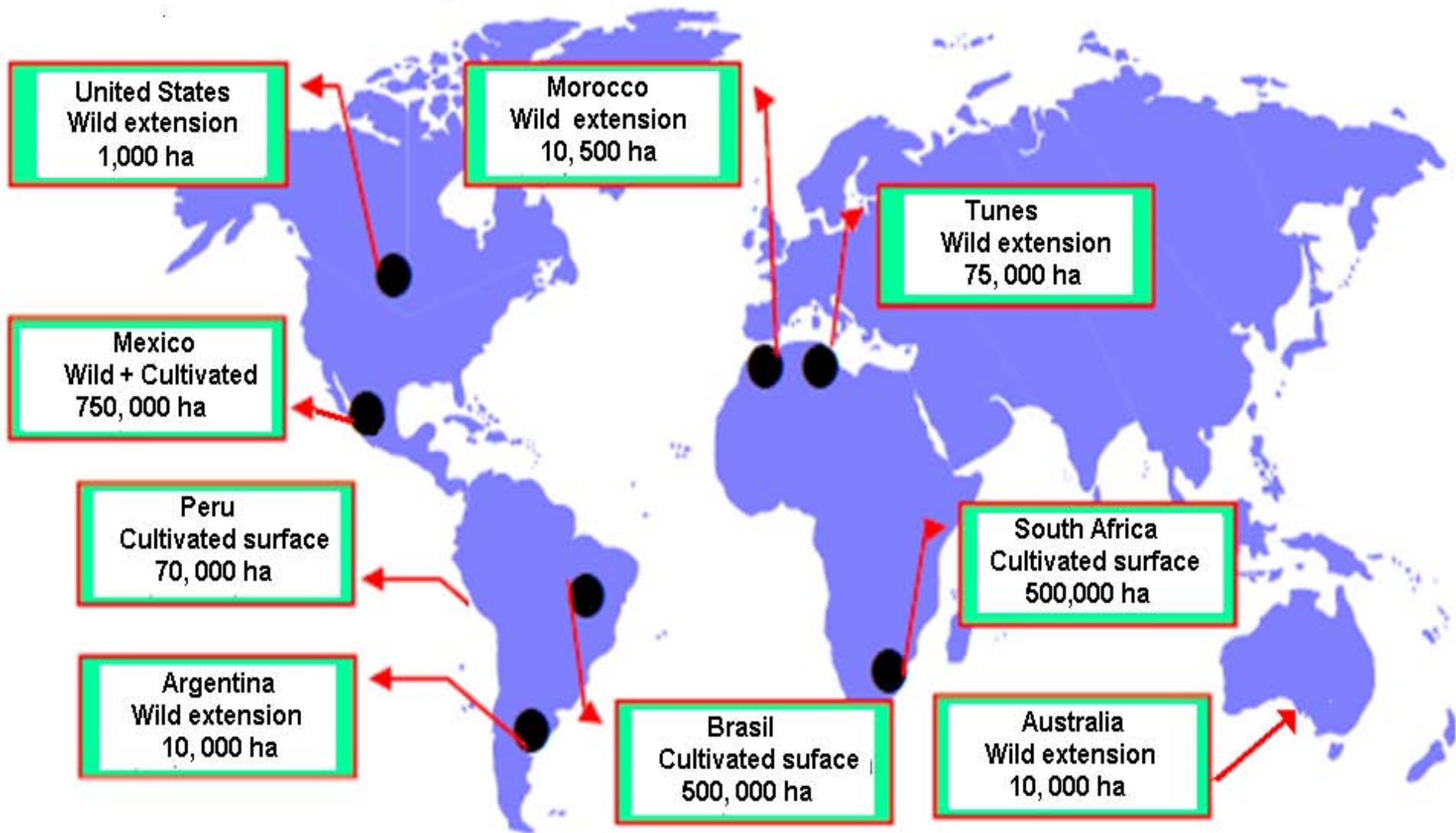


Countries / Hosts

<i>D. opuntiae</i>	16/33
<i>D. coccus</i>	15/8
<i>D. ceylonicus</i>	13/16
<i>D. confusus</i>	12/36
<i>D. tomentosus</i>	7/14
<i>D. austrinus</i>	3/8
<i>D. confertus</i>	3/10
<i>D. salmianus</i>	3/1
<i>D. zimmermanni</i>	3/7

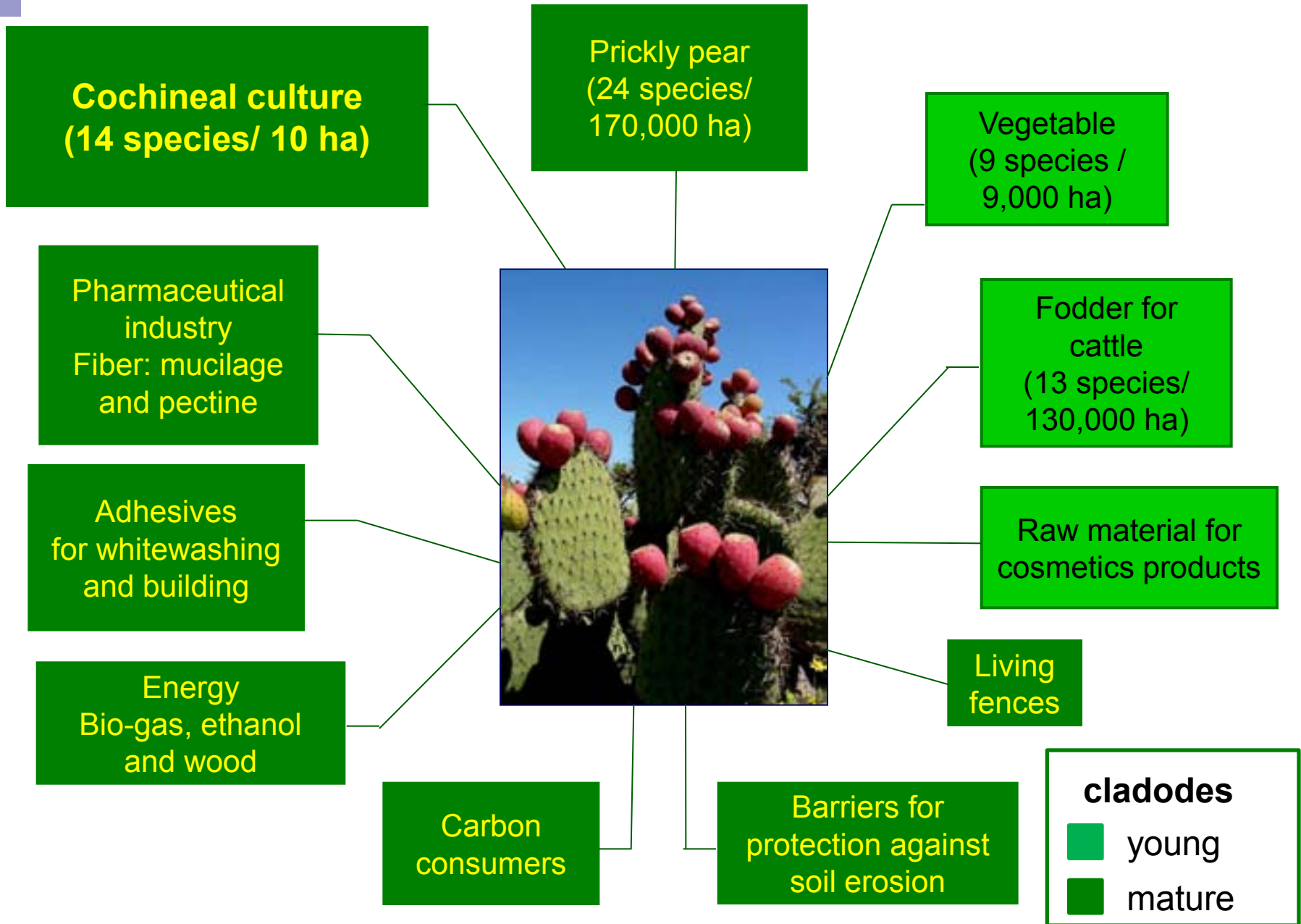
Opuntioideae *Cylindropuntia*, *Grusonia*, *Maihueniopsis*, ***Opuntia*** 60% *Tacinga*, *Tephrocactus*, *Tunilla*. **Other genus:** *Cereus*, *Cleistocactus*, *Denmoza*, *Echinopsis*, *Gymnocalcium*, *Harrisa*, *Maihuenia*, *Mammillaria*, *Pilosocereus*, *Selicereus*.

Opuntia



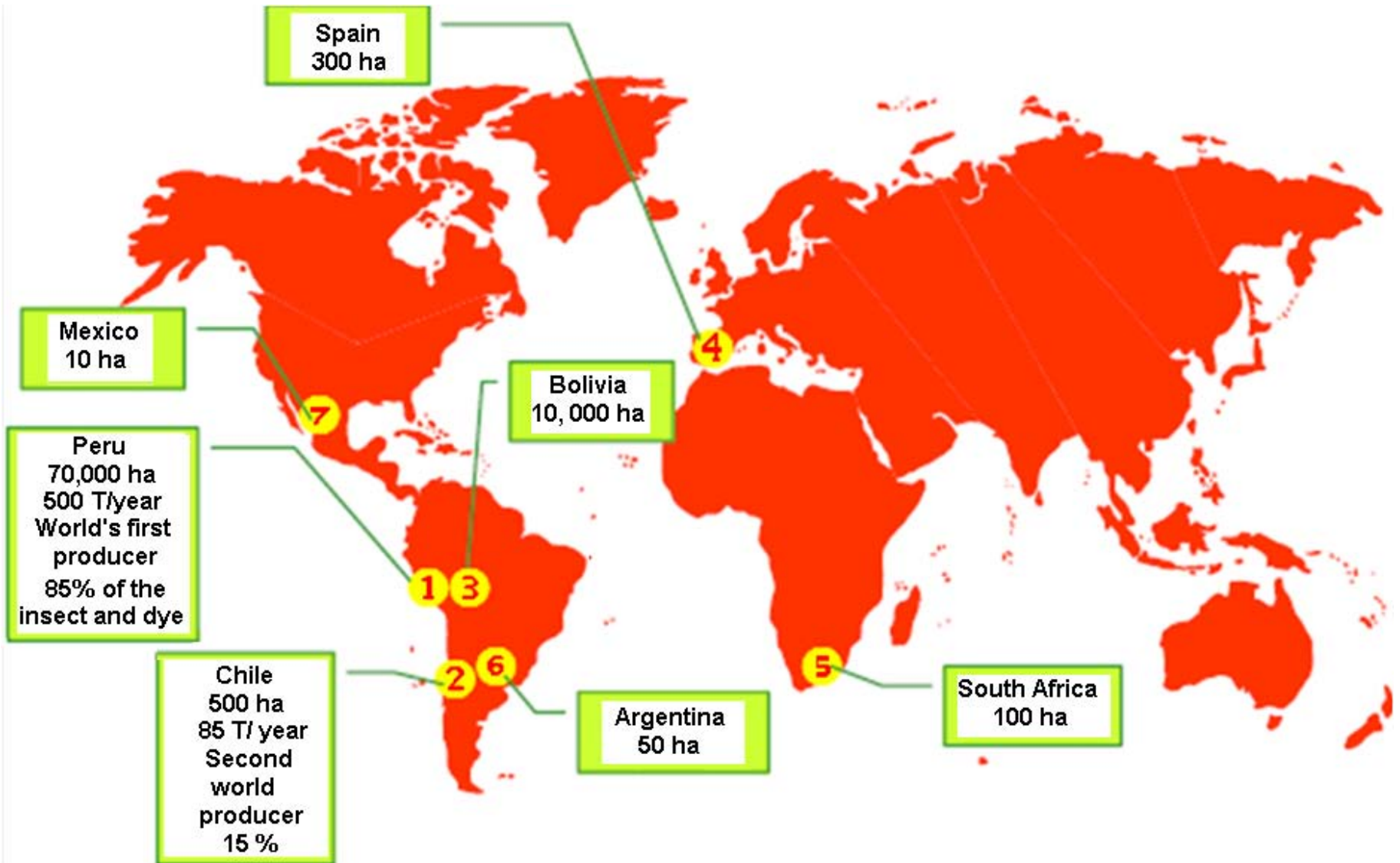
Market: Local markets, Mexico, United States, Japan

Uses in Mexico



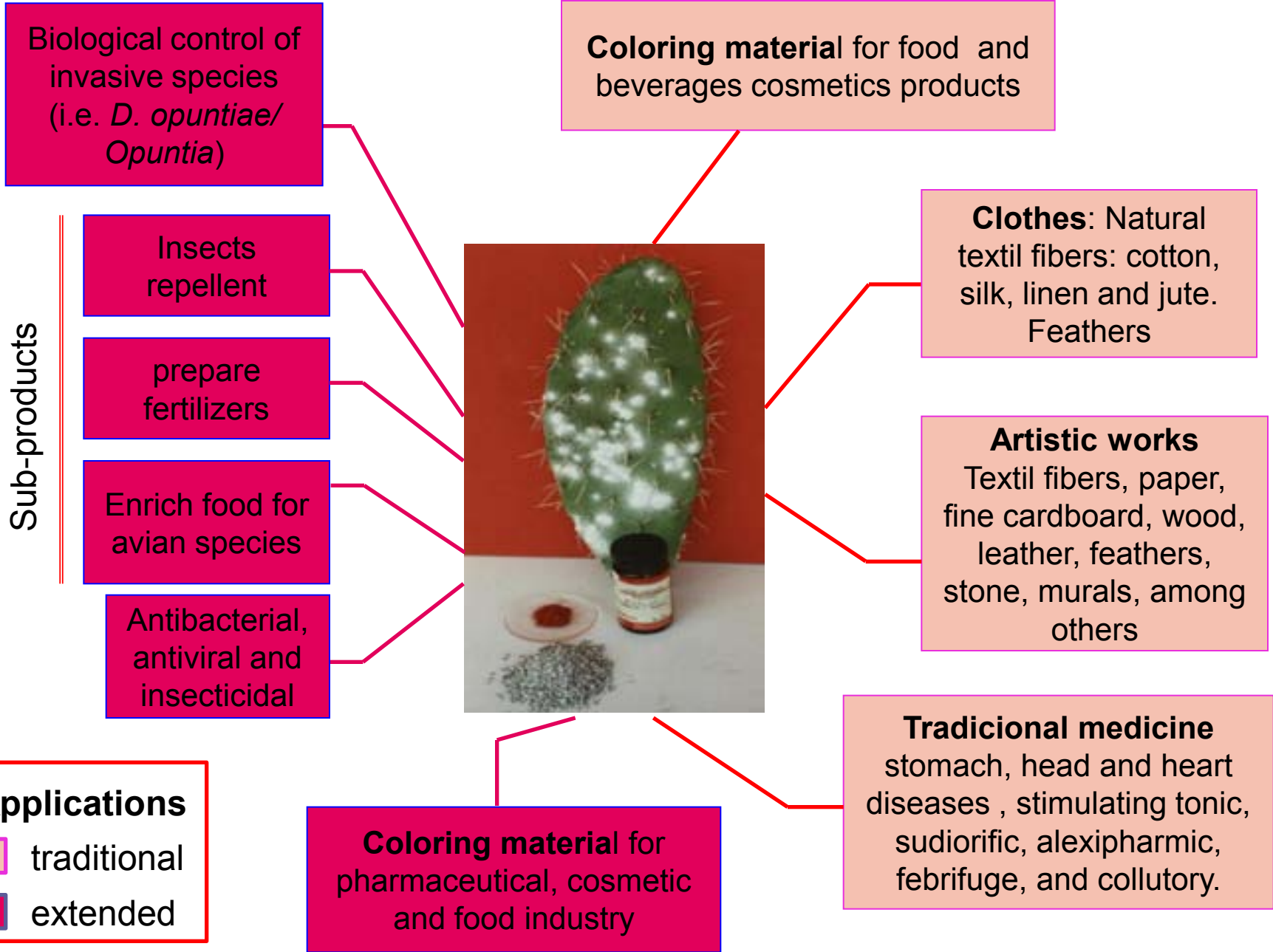


Dactylopius coccus



Market: Local, France, England, Italy, Japan, United States and Argentina

Uses in Mexico



- Food



- Cosmetics



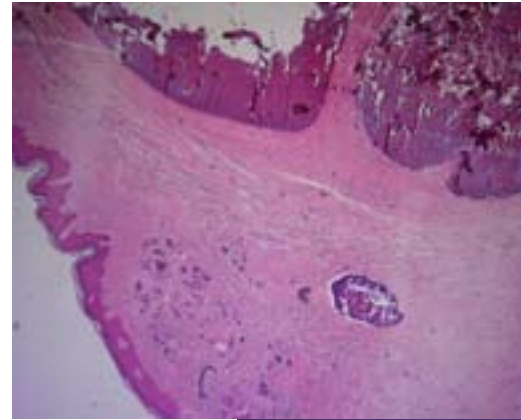
- Medicines



- Textiles



- Antibacterial, antiviral, and insecticidal



Conservation strategies

1. Constant fragmentation promoted by the extraction and exploitation of wild species, including *Opuntia*, without restrictions is reducing the possibility of studying the interaction between *Dactylopius* and *Opuntia*.



2. South American cactus-feeding moth, *Cactoblastis cactorum* a serious threat to the high diversity of native *Opuntia* species in Mexico.




51 endemic, 6 cultivated and 18 wild-growing species

Environmental Protection of native species in Mexico of wild flora and fauna, risk categories and specifications inclusion or change.

Opuntia 5/284. *Dactylopius* is not even considered.



Specific measures for protection of such biodiversity and generic resources, the patron of distribution and the habitat characteristics to maintain these resources.



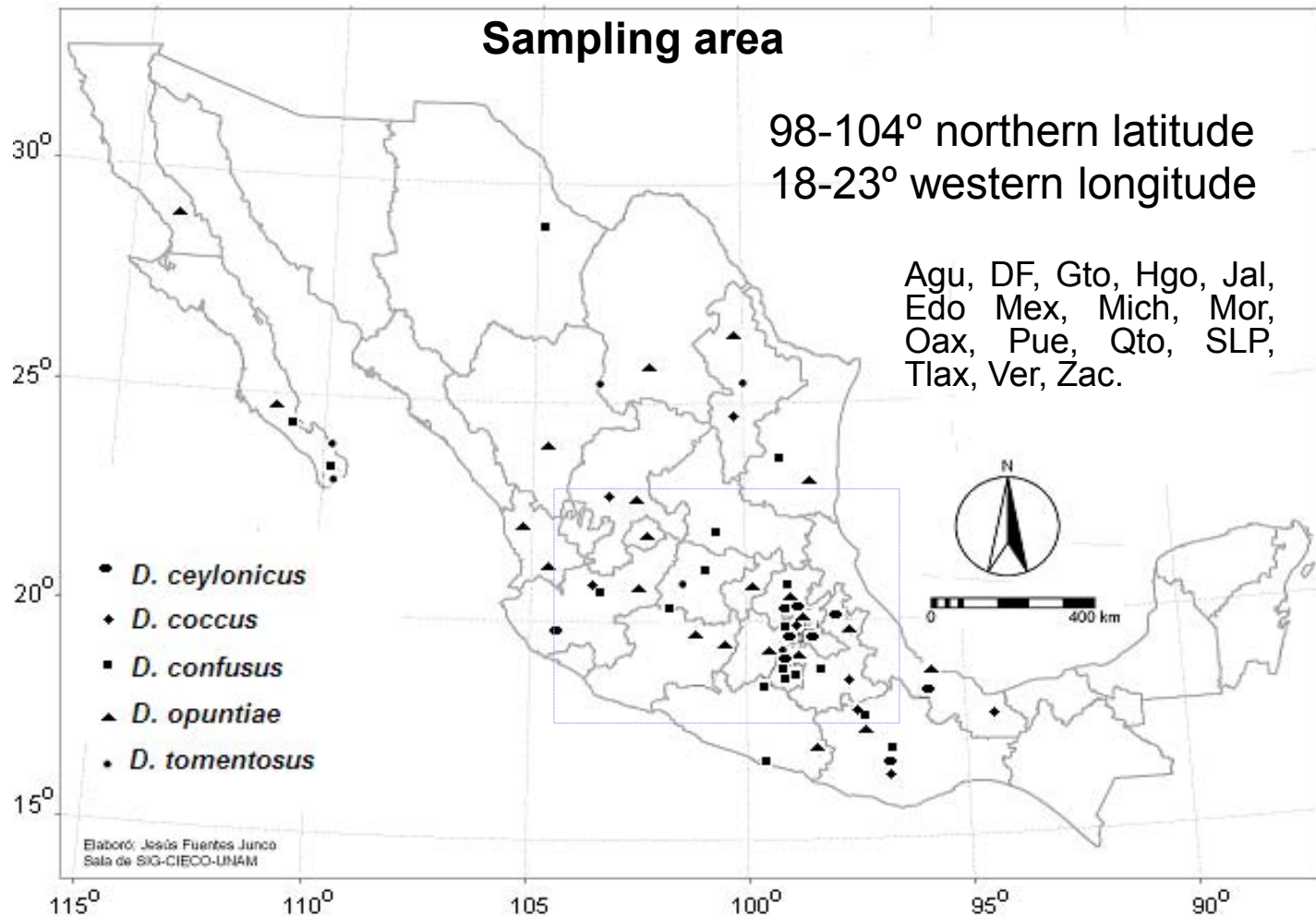
II. Chávez-Moreno, C.K.¹ Tecante A.¹, Casas A.², Claps L.E.³
Distribution and habitat in Mexico of *Dactylopius* (Hemiptera:
Dactylopiidae) and their hosts of the subfamily Opuntioideae
(Cactaceae)
Neotropical Entomology

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II. Distribution maps



NOTE: Records of distribution of *Dactylopius ceylonicus* and their hosts. Data compiled Britton & Rose 1963, Bravo-Hollis & Sánchez-Mejorada 1978, González *et al* 2001, Guzmán *et al* 2003, and herbaria MEXU and IBUG. Insect records Mann 1969, De Lotto 1974, Piña 1977, MacGregor & Sampedro 1983, Pérez-Guerra & Kosztarab 1992, Portillo & Viguera 2003a, b). New records (empty symbols) described in this study. CNI-IB-UNAM, GIS constructed ILWIS 3.3 mapping geographic location *Dactylopius* and hosts.

Field collection

25-100 specimens **208** populations **120** localities **14** states

2005 Feb, May –June, Sep; **2006** April-June; **2007** Feb, Nov.

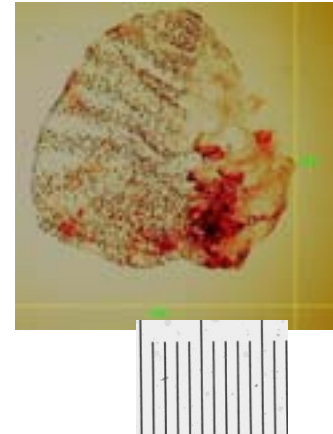
Dactylopius

Female

Male

stages of development
different portions host
collected separately
wild populations
production research centers
urban and rural zones

a live
preserved
70%, 96%
ethanol
CNI-IBUNAM



HOSTS

Opuntia

Nopalea

Cylindropuntia

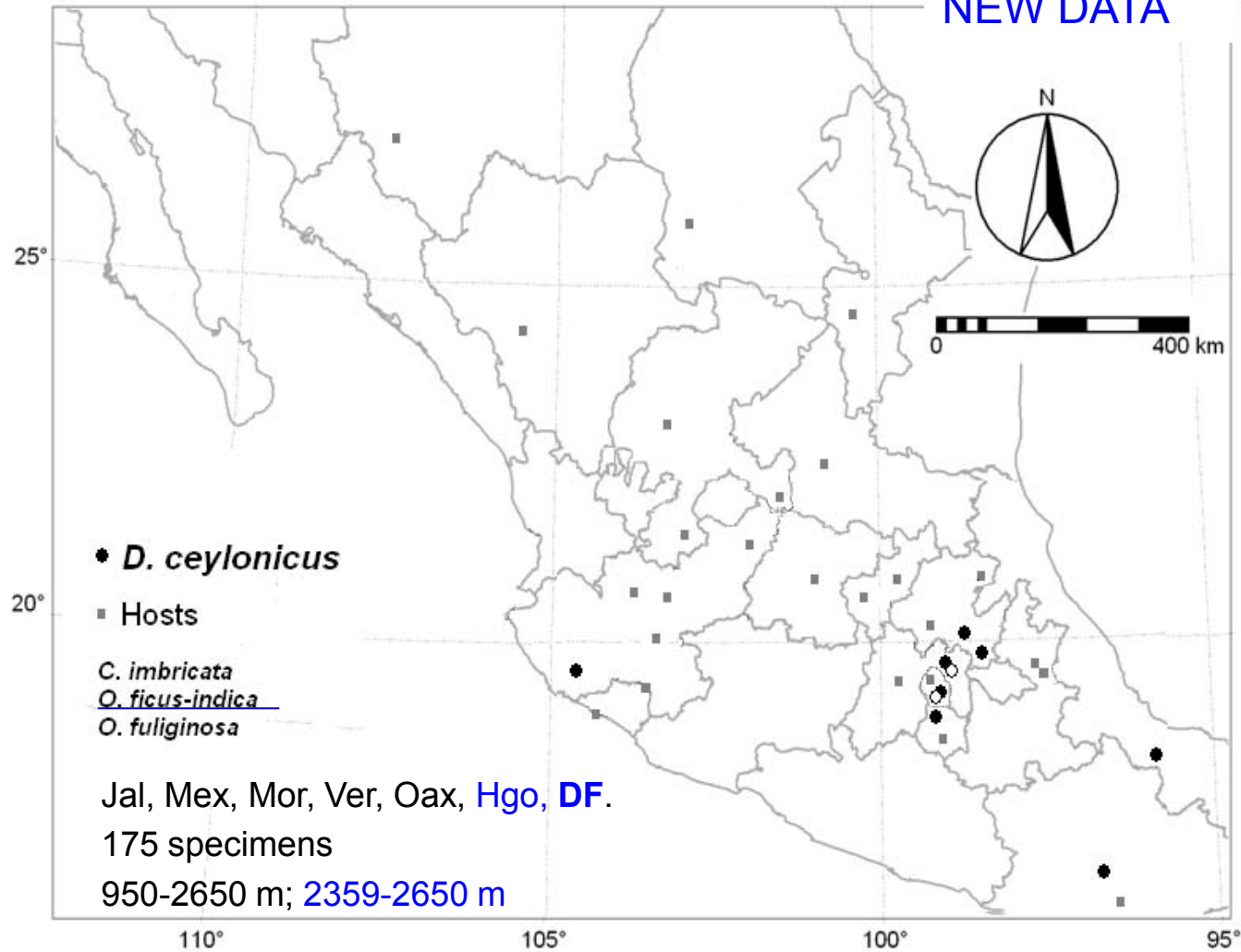
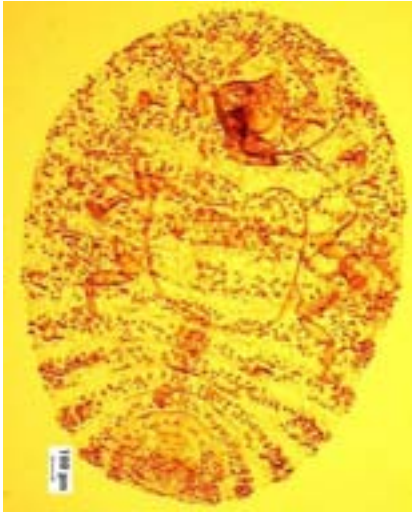
Collected 3X
Propagation

living
collection
Botanical
garden
CIEco-UNAM



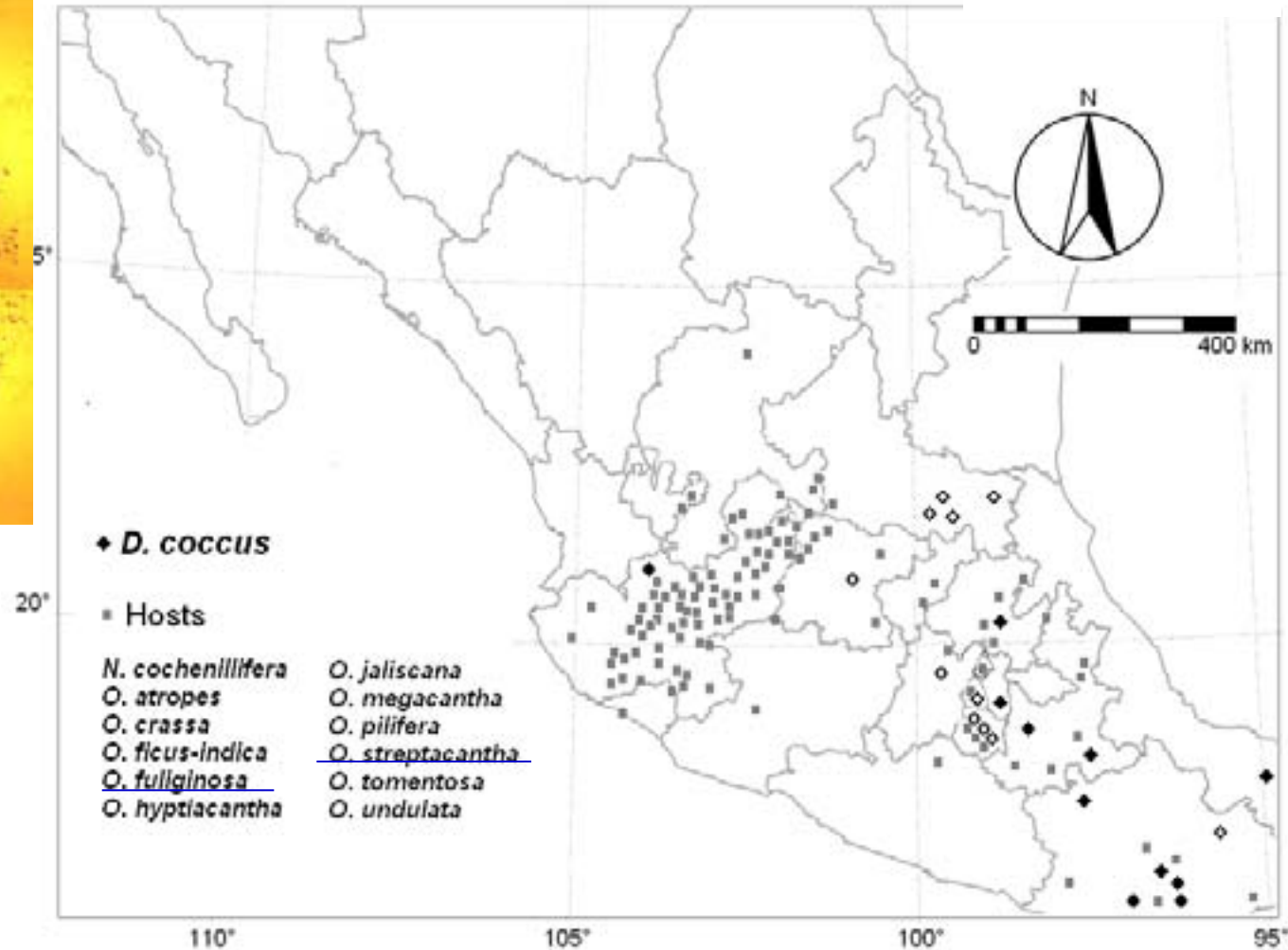
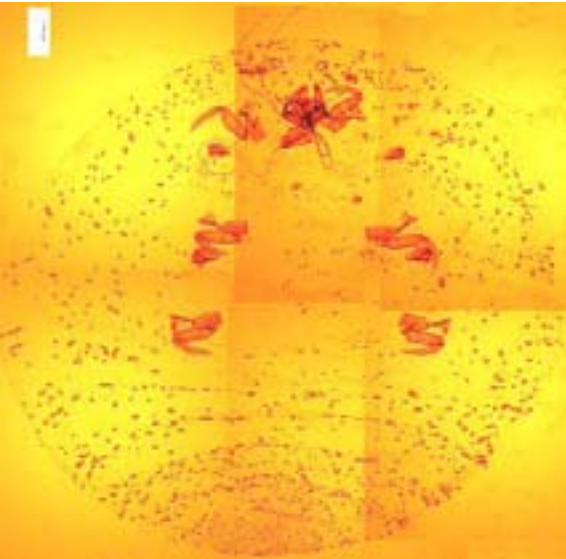
Identification of species Taxonomic keys Support
(De Lotto 1974, Pérez-Guerra y Kosztarab 1992).

Environmental database: insect, host, place of collection, vegetation and soil types, weather Peel *et al* (2007).



NOTE

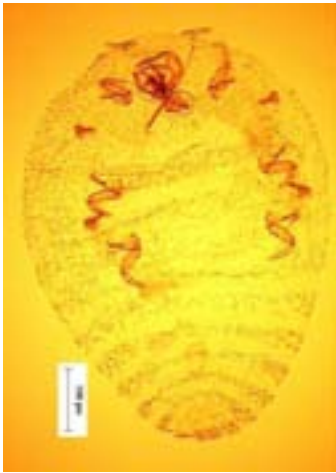
Root nodules in winter / summer xerophilous thickets & arenosol



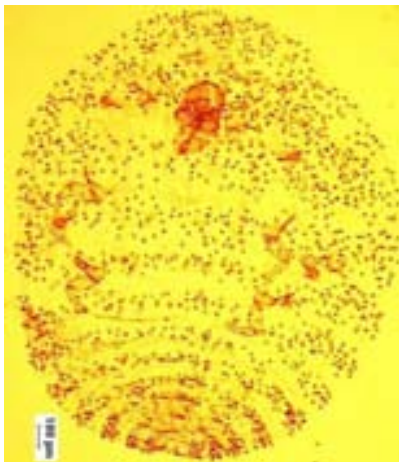
Jal, Hgo, Ver, Pue, Oax ©, Gto, Mex ©, Mor © y SLP© 575 specimens
1250-2200; 1654-2845 m

Research & production centers, living fences /xerophilous thickets, agricultural fields/calcisol, xerosol, vertisol, regosol, leptosol, fozem

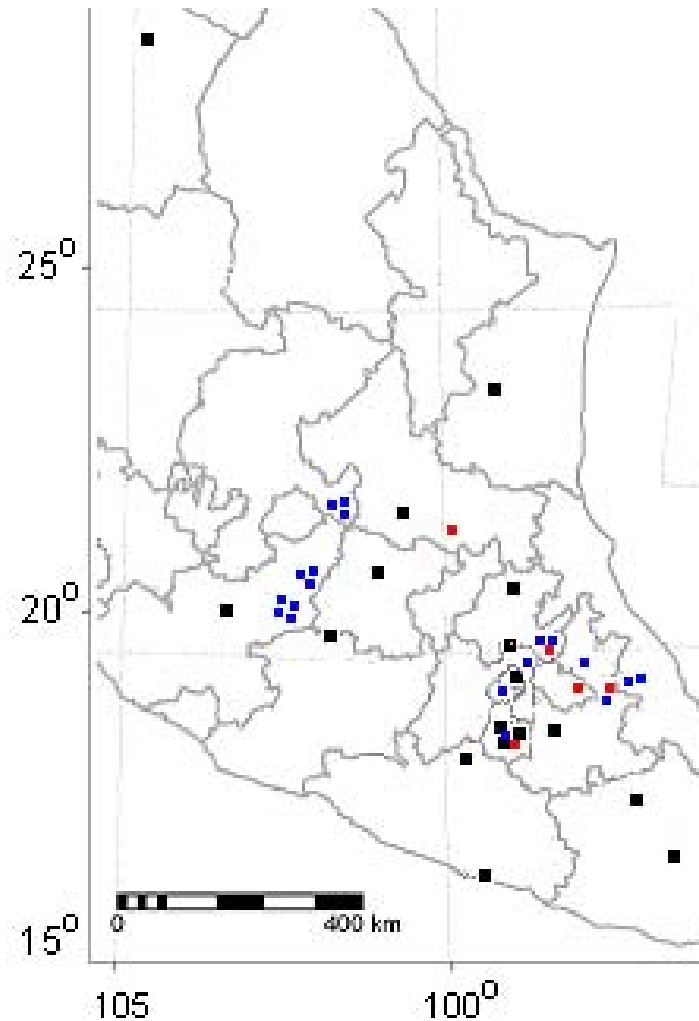
D. confusus



D. confusus



D. salmianus



- Review
- *D. confusus*
- *D. confusus* biotype 1

Son	Coah	NL
Dgo?	Chih	Tam
Gto	Gro	Oax
Jal	DF	Pue
Hgo*	Mor*	Ver
Zac	SLP	Tlx

1100-2200 m
 1200-547 m
 1654-2773 m

Opuntia y *Nopalea* *O.grahamii*, *C.imbricata*, *C.leptocaulis*, *C.tunicata***,
O.fuliginosa, *O.jaliscana*, *O.joconostle*, *O.spinulifera*, *O.streptacantha*,
O.phaeacantha, *O.hyptiacantha*, *O.ficus-indica**, nopal tuna roja



SLP

Morelos

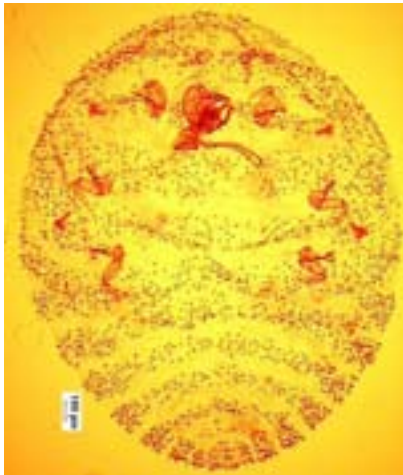


**Living fences / urban zones
Production centers foezem y
regosol**

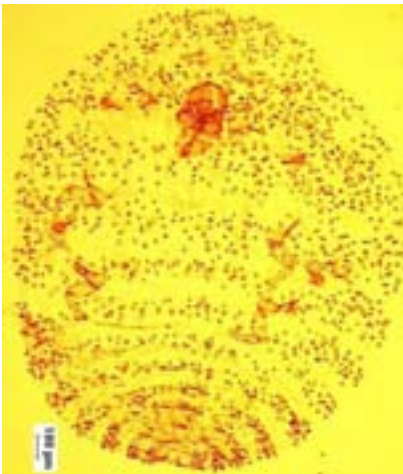


Reesearch centers / xerophilous thickets*/arenosol*, xerosol*

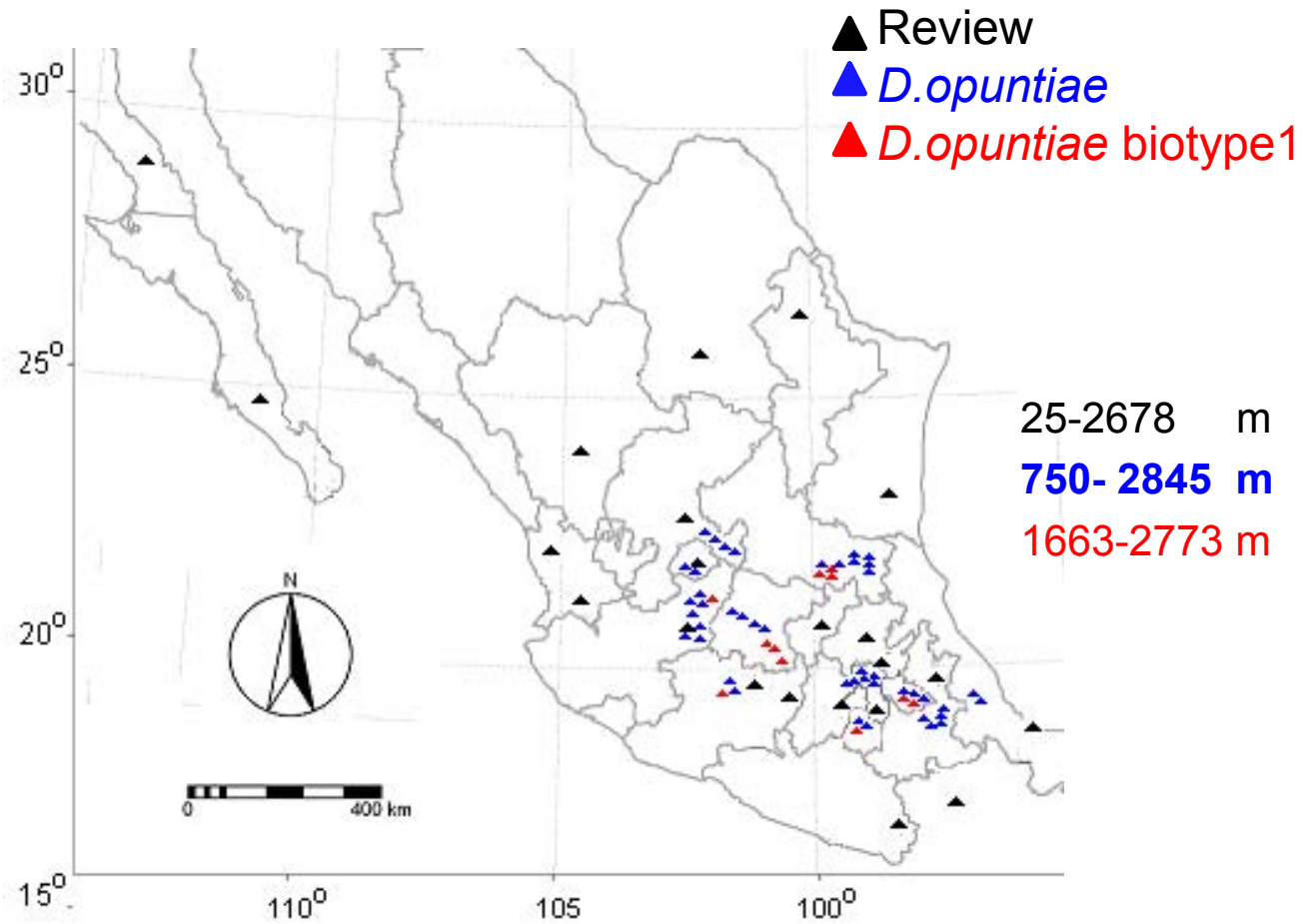
D. opuntiae



D. opuntiae



D. salmianus



Oax, Gro, BC, Coah, NL, Tam, Dgo, Nay, Qro, Mex, Hgo, Chia Zac, Ags, Jal*, Mich* DF, Pue, Ver, Gto*, Mor*, SLP*, Tlx*

*O. ficus-indica*** , *O. tomentosa** , *O. robusta* , *Opuntia* , *Nopalea* y *Cactus* , *O. hyptiacantha* , *O. megacantha* , *O. joconostle* , *O. phaeacantha* , *O. atropes* , *O. albicarpa* , *C. tunicata* , *O. streptacantha** , variantes * y cultivares *

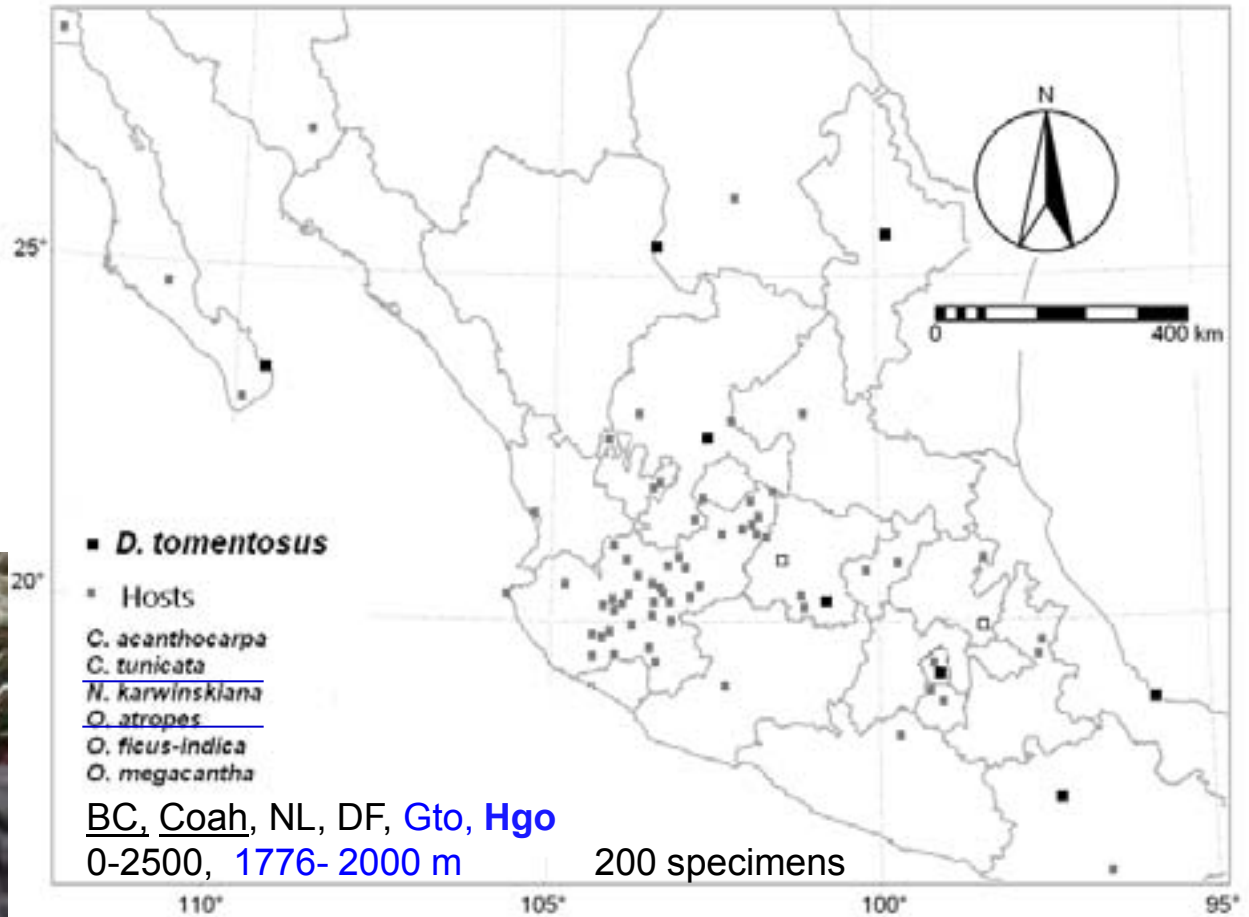
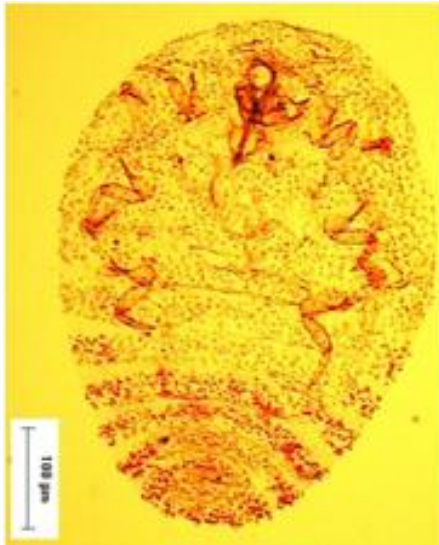
Estado de Mexico

1200 specimens



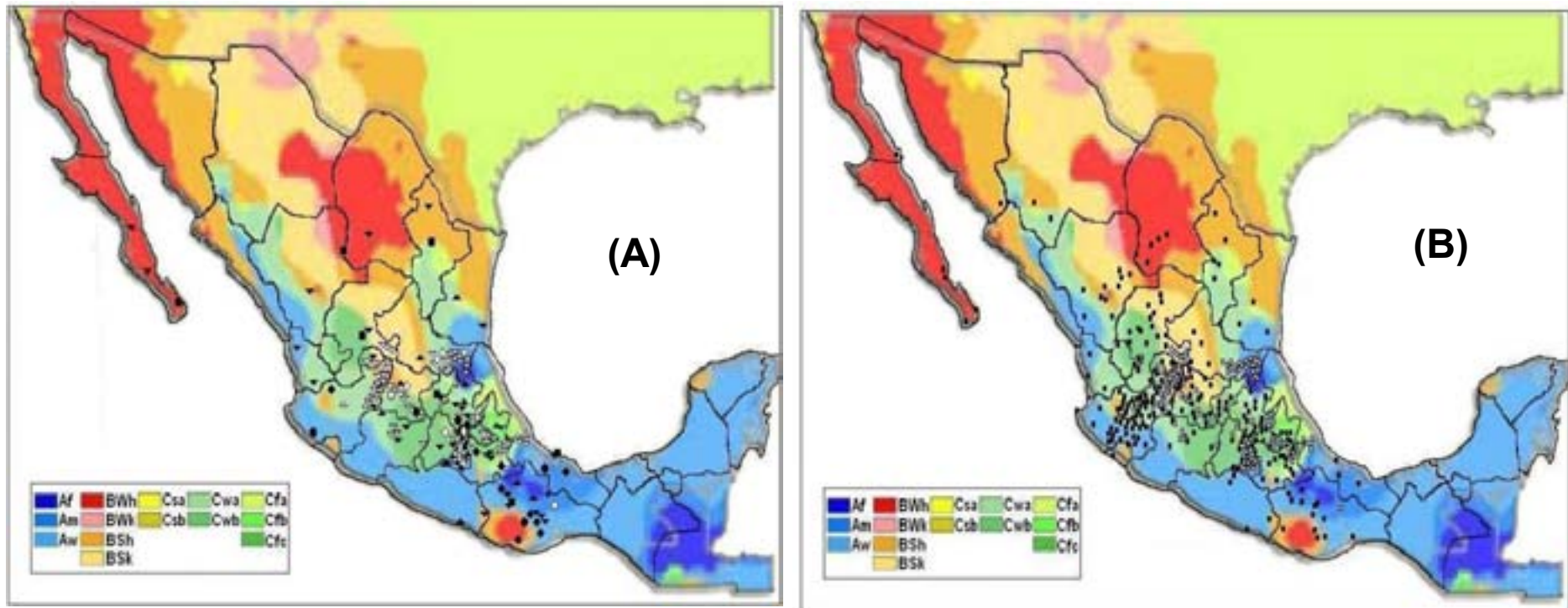
Living fences, urban zones Production centers /
Cultivated Forest oak-pine / calcisol, regosol, fozem,
regosol, vertisol

D. tomentosus



Areolas of cladodes / living fences xerophilus thickets vertisol, arenosol

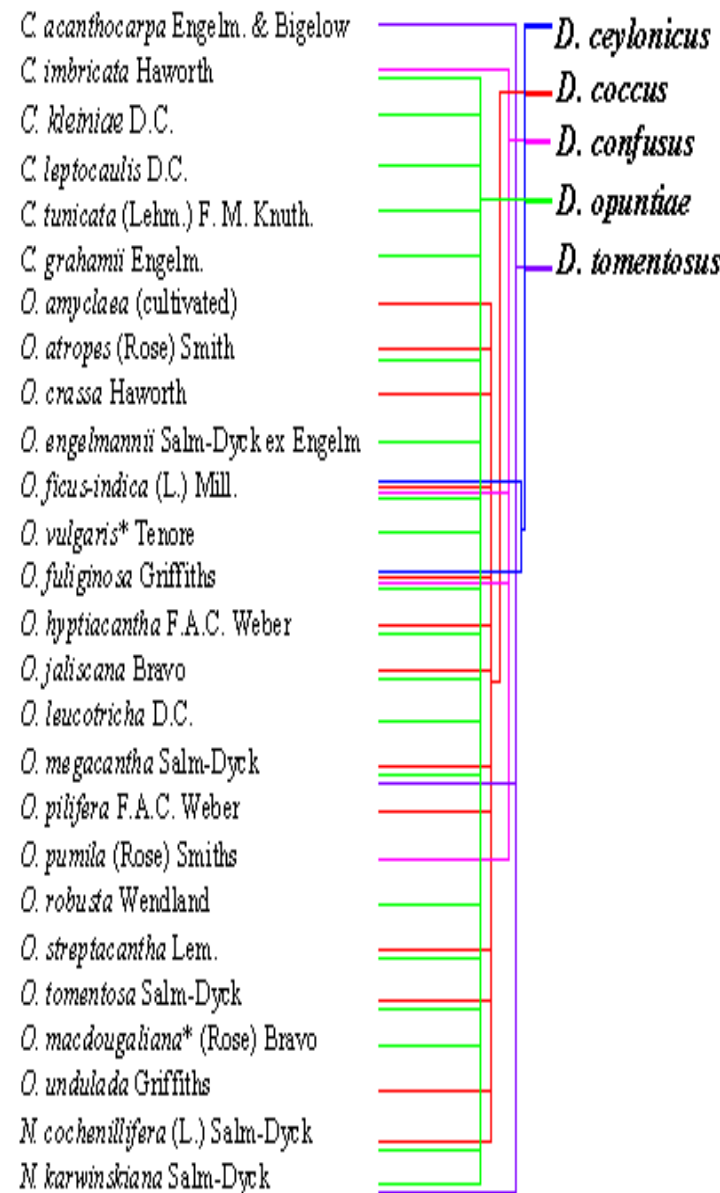
Biogeographical region



Distribution of **A) *Dactylopius*** *D. ceylonicus* (circles), *D. coccus* (diamonds), *D. confusus* (triangles), *D. opuntiae* (inverted triangles), *D. tomentosus* (rectangles) and **B) *Opuntia*** over the Köppen-Geiger climate type map of Mexico (North America) extracted from Peel *et al* (2007).

Conclusion

DATA	BEFORE	AFTER
Region	14	23
Species	5	5 or more
Biotypes	--	2
Hybrids	--	¿?
Cohabit	--	2, 3, 4 species
Hosts	26	41+26 = 67
Vegetation	Hosts`s	xerophilous thickets, cultivars, collections, urban zones & forest (pine, oak)
Soil	Hosts`s	foezem, vertisol, xerosol, arenosol, calcisol, regosol, leptosol
Weather	Hosts`s	Template, dry arid and semiarid, Forest pine-oak
Elevation	---	0-2845



Chemical and genetic analysis contribute to establish interaction specificity

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Metabolic profiling of *Dactylopius* (Hemiptera: Dactylopiidae) species pigments by geographical origin and hosts using multivariate data analysis

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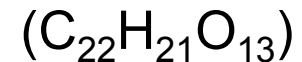
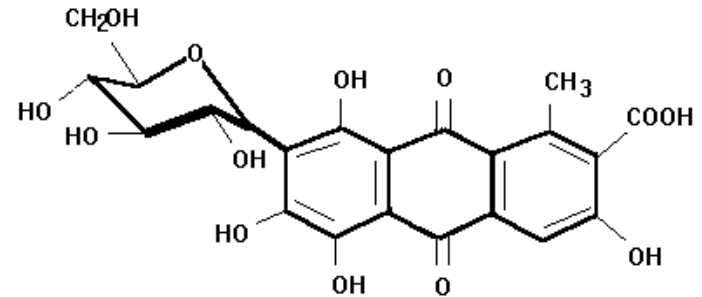
III. Chemical analysis

Dactylopius coccus

- Source of natural colorant (Food & Drug Administration. 2009).

- Historically wild species could serve as alternative.

or additional sources of the colorant and its derivatives.



- Studies HPLC–MS–NMR

- Origin, constituents hydroxyanthraquinone (González *et al.*, 2002),
- Chemical structure characteristics (Méndez *et al.*, 2004; Maier, 2004),
- Identify CA: foods, antique textiles, (Yamada *et al.*, 1993; González *et al.*, 2002; Szosteketal, 2003; Maier *et al.*, 2004; Surowiec *et al.*, 2007; Peggie *et al.*, 2008; Karapanagiotis *et al.*, 2008, 2009).
- Optimize extraction conditions, quality (Wouters, 1985; González *et al.*, 2002; Méndez *et al.*, 2004; Szosteketal, 2003; Peggie *et al.*, 2008; Karapanagiotis *et al.*, 2009).

OBJETIVE: compare the metabolic profile 5 mexican species and the argentinan.

- Material populations
 - 35 Mexico *Dactylopius*
 - 2 Argentina *D. ceylonicus*
 - Outgroup *K. Vermilio*

Reheodyne 117 volt AC, 50-60htz

Binary LC: pump Water 600

Column C18 waters

Dual λ absorbance detector

0.5 mL/min / 20 μ L

Quintuplicate

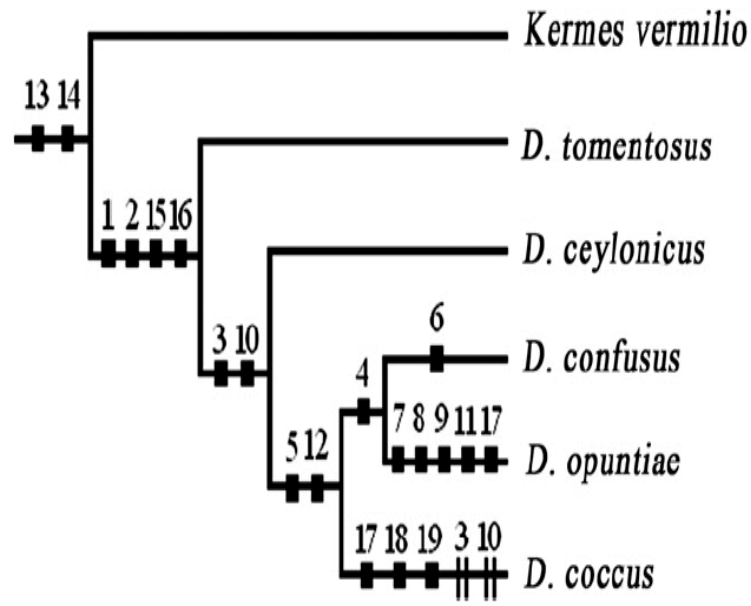


(Méndez *et al.*, 2004; González *et al.*, 2002)

Peak tR(min) *D. ceylonicus* *D.coccus* *D.confusus* *D.opuntiae* *D.tomentosus*

Peak	tR(min)	<i>D. ceylonicus</i>	<i>D.coccus</i>	<i>D.confusus</i>	<i>D.opuntiae</i>	<i>D.tomentosus</i>
1	0.678	1	1	1	1	1
CAc	1.84	1	1	1	1	1
3	3.29	1	0	1	1	0
4	4.58	0	0	1	1	0
5	5.36	0	1	1	1	0
6	6.05	0	0	1	0	0
7	8.31	0	0	0	1	0
8	9.73	0	0	0	1	0
9	10.6	0	0	0	1	0
10	12.5	1	0	1	1	0
11	13.4	0	0	0	1	0
12	14.3	0	1	1	1	0
FkAc	15.4	1	1	1	1	1
KAc	16.6	1	1	1	1	1
15	17.5	1	1	1	1	1
16	18.8	1	1	1	1	1
17	19.3	0	1	0	1	0
18	21.9	0	1	0	0	0
19	22.1	0	1	0	0	0

1 Presence: **unique** or **shared**
 0 Absence

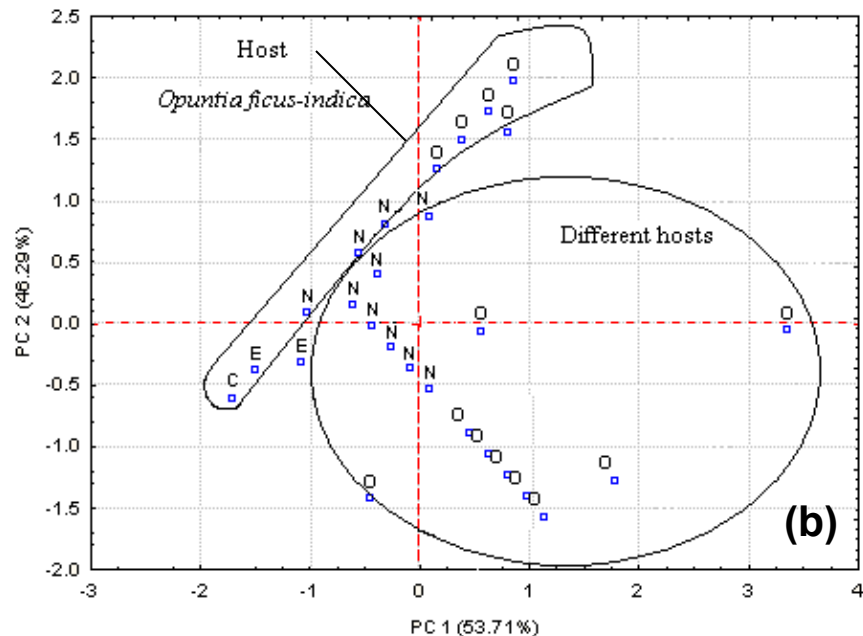
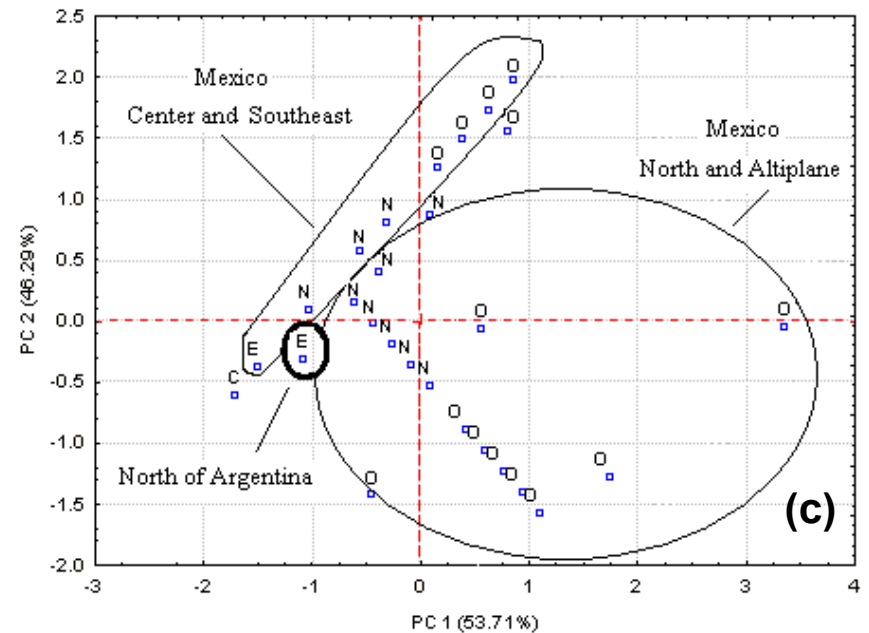
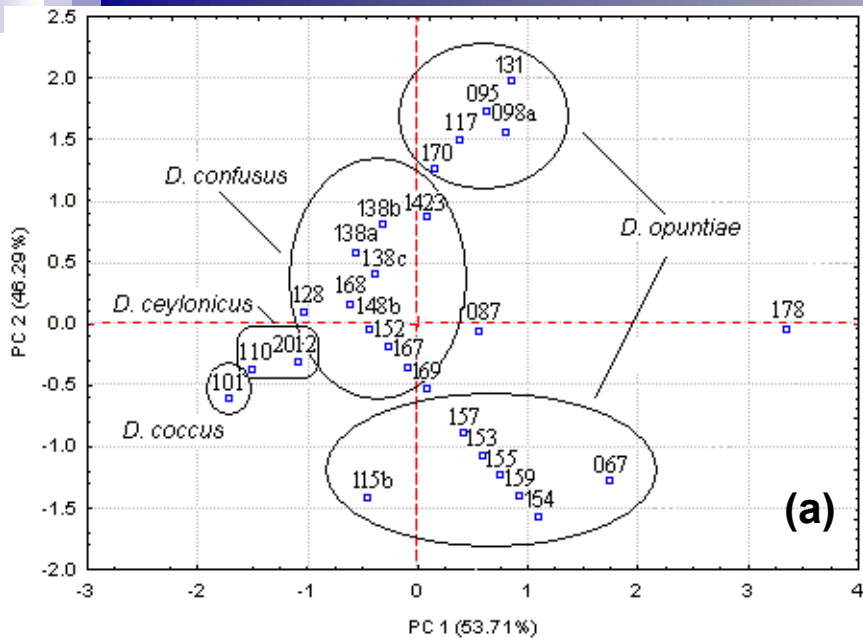


Dendrogram (1-0). Resulting from a cluster analysis using principle of parsimony for *Dactylopius* & *K.vermilio*. The numbers above the lines indicate the peaks of the chromatographic profile.

Total peak 8 11 12 16 6

Quantitative analyses. Absolute area (*10⁻⁴) of the HPLC profile of commercial carminic acid and five *Dactylopius* species (0 = absence, CAc = carminic acid, KAc = kermesic acid).

Pico	Acido carminico	<i>D. ceylonicus</i>	<i>D. coccus</i>	<i>D. confusus</i>	<i>D. opuntiae</i>	<i>D. tomentosus</i>
Cac	9.83 ± 3.13	3.70 ± 0.241	8.60 ± 3.12	3.45 ± 0.91	4.70 ± 0.41	6.37 ± 0.280
3	1.83 ± 3.17	> 0 ± 0.002	0	0.50 ± 0.13	0.24 ± 0.028	0
4	0	0	0	0.30 ± 0.12	0.18 ± 0.31	0
5	0	0	1.08 ± 0.38	0.30 ± 0.12	0.12 ± 0.13	0
6	0	0	0	0.020 ± 0.023	0	0
9	0	0	0	0	> 0 ± 0.001	0
10	0.010 ± 0.016	0.13 ± 0.23	0	0.06 ± 0.05	> 0 ± 0.0002	0
12	0	0	2.39 ± 1.21	0.62 ± 0.34	> 0 ± 0.0003	0
Kac	0.83 ± 0.05	0.75 ± 0.17	0.03 ± 0.05	0.07 ± 0.13	0.59 ± 0.09	1.92 ± 0.286



Component analysis the plane formed by the first two principal components.

D. coccus (C), *D. ceylonicus* (E),
D. confusus (N) and *D. opuntiae* (O).

a) species

b) geographic distribution 3, 9, 15

c) host *O. ficus-indica* 3, 7, 11 CAc

Molecular Phylogeny of the Genus *Dactylopius* (Hemiptera: Dactylopiidae) and Identification of the Symbiotic Bacteria

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Systematic of *Dactylopius*

■ Molecular

- Phylogeny of Coccoidea Dactylopiidae-Eriococcidae. Mitochondrial gene secuenciation 12S rARN and nuclear 18S rARN (Cook et al., 2002)

■ Morphological analysis

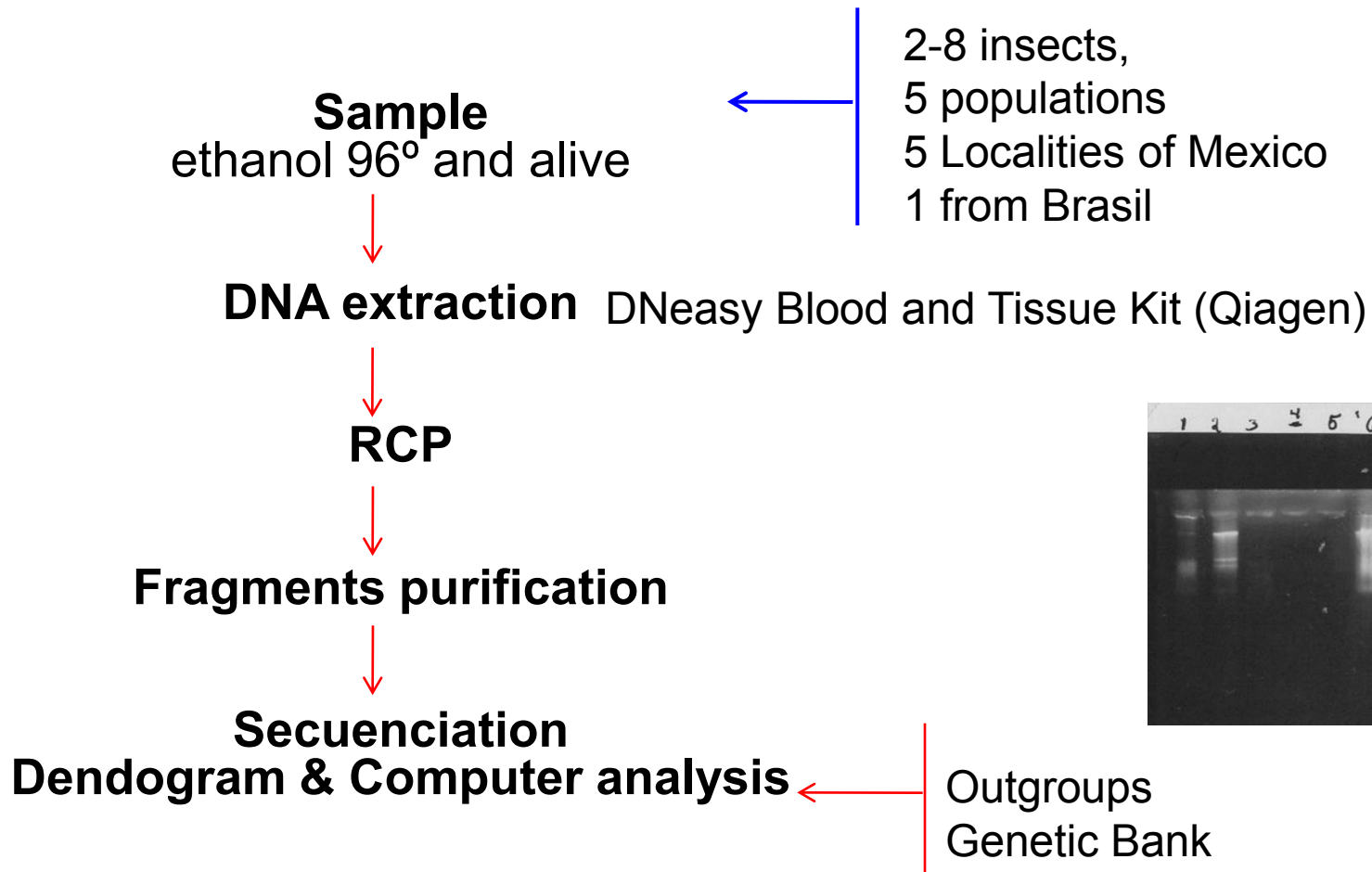
- Taxonomical keys (De Lotto, 1974; Pérez-Guerra & Kosztarab, 1992), and Rodríguez *et al.* (2001)

■ Endosymbionts

- *Wolbachia* present in *Dactylopius* sp. eggs secuenciated by nuclear 16S rARN gene. (Thao et al., 2002, Baumann, 2005; Moran, 2006, (Gruwell et al., 2007, Pankewitz et al., 2007).

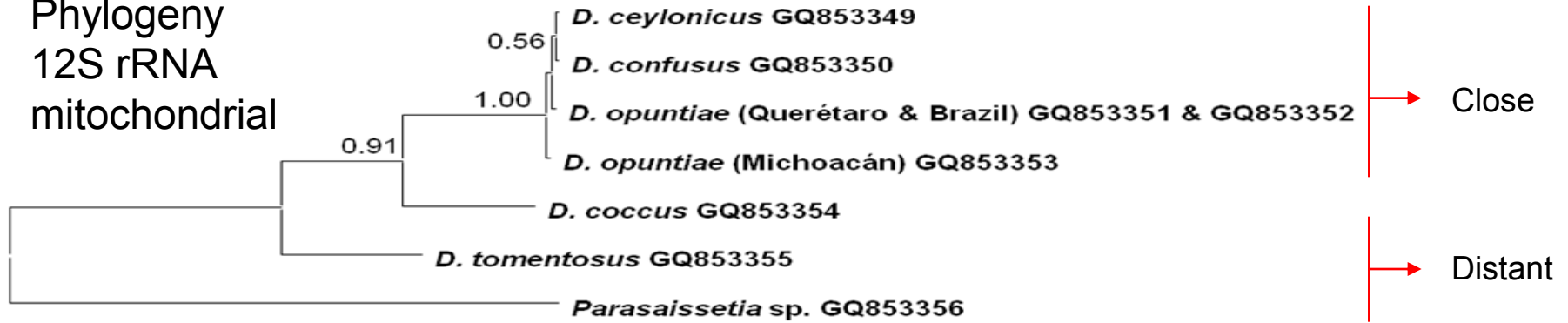
OBJETIVE. *Dactylopius* phylogeny of 5 mexican species with mitochondrial 12S rARN and nuclear 18S rARN genes for insectos.

Endosymbionts phylogeny of insects molecular gene 16S rARN.

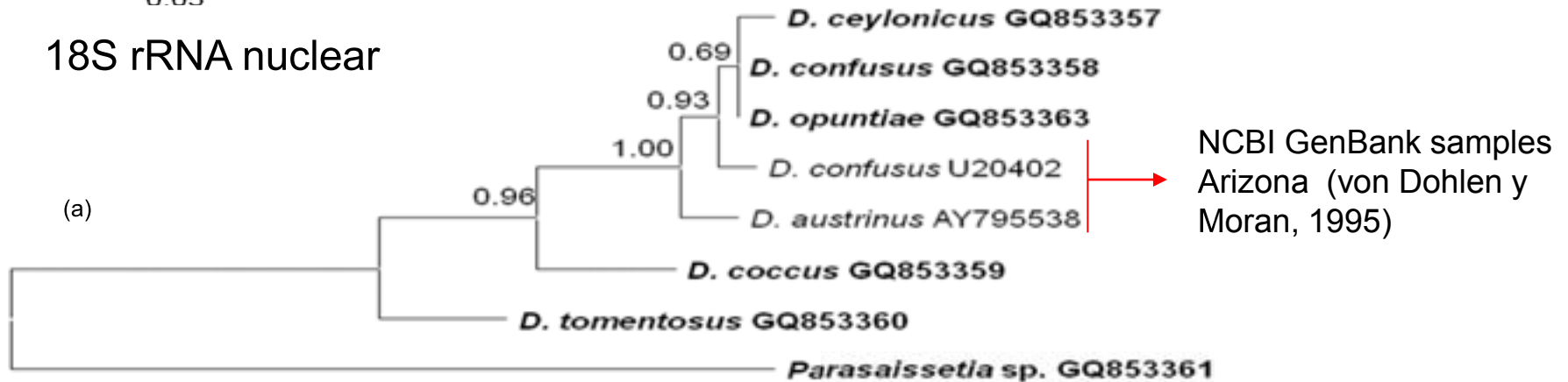


III. Molecular analysis

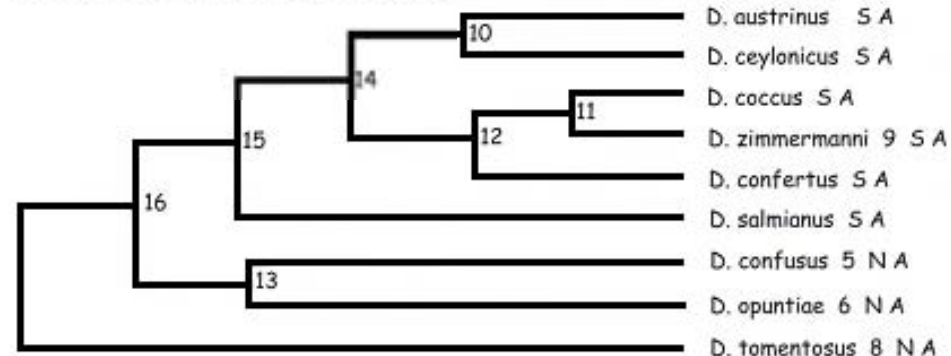
Phylogeny
12S rRNA
mitochondrial



18S rRNA nuclear

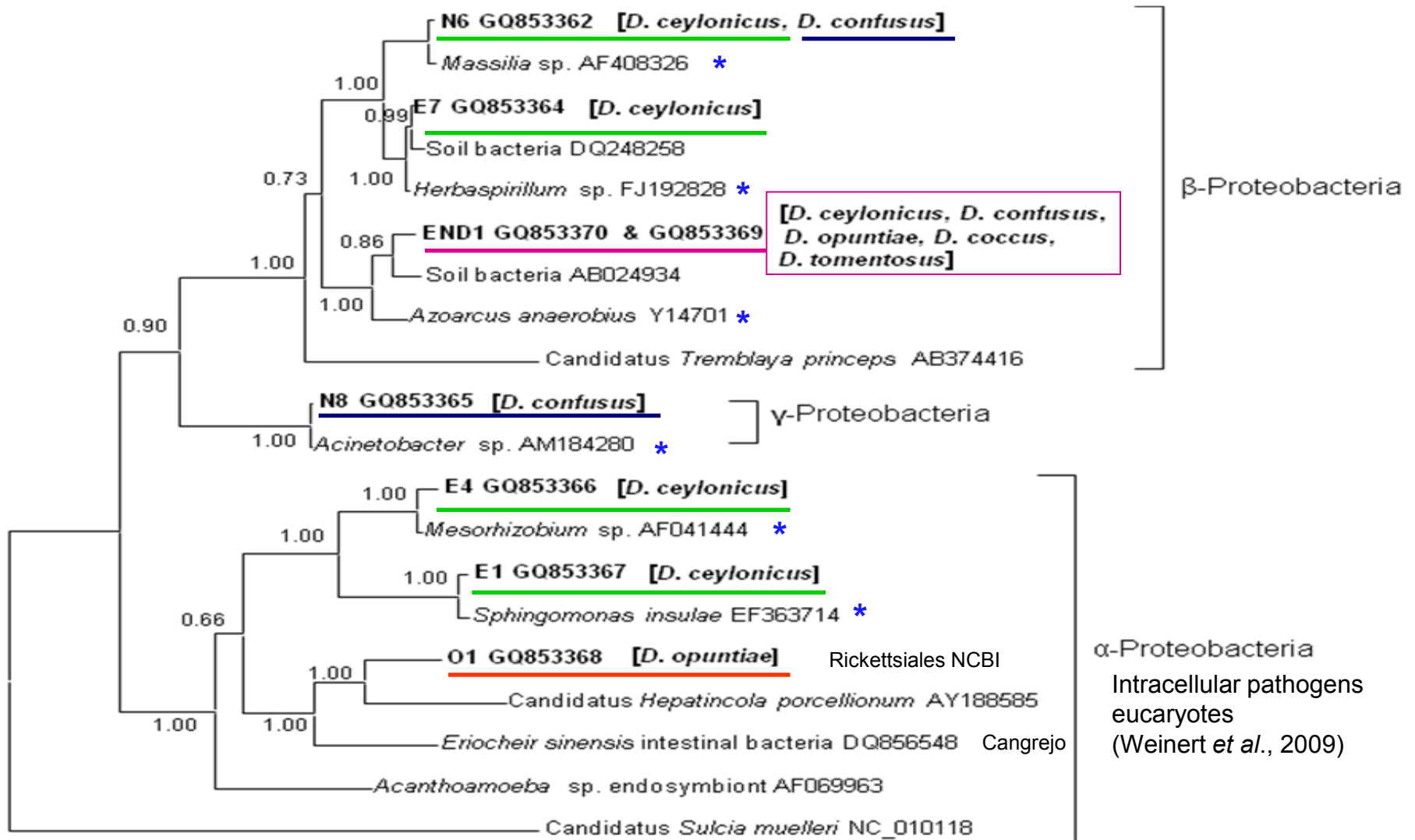


Comparison of molecular phylogeny (18S rRNA nuclear) Morphological phylogeny (Rodríguez *et al.*, 2001), resulted in different topographies.



Molecular analysis

Dactylopius endosymbionts phylogeny 16SrRNA



0.05

* Free living bacteria in the intestine similar to those present in plants/soil

Conclusions

- Mexico Center origin genetic diversity of *Dactylopius* and their hosts
 - Real and potencial uses of these genetic resources in Mexico.
 - Geographic patrons and habit characteristics, main protected areas.

 - Chemical analysis:
 - Metabolic profile Colorant of 5*Dactylopius*, diferentiate according species, geographical origin and host and
 - recognise *Dactylopius* source of colorant.

 - Molecular analysis:
 - Molecular phylogeny 5 Mexican *Dactylopius* sequenced mitochondrial genes 12S rRNA and nuclear sequenced 18S rRNA for insects.
 - Comparison between molecular phylogeny and morphology dendogram characters resulted in different topographies.
 - Insect endosymbiont phylogeny 16S rRNA demonstrated presence:
 - α -protobacteria (E1, E4, O1)
 - β -protobacteria (N6, N7, END1)
 - γ -protobacteria (N8)
- } → Bacteria intestine/insects, and free living/plants and soil.
-
- The Conservation for the interaction plant-insect resources must be the guide to use in a sustentable maner these genetic resources.

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