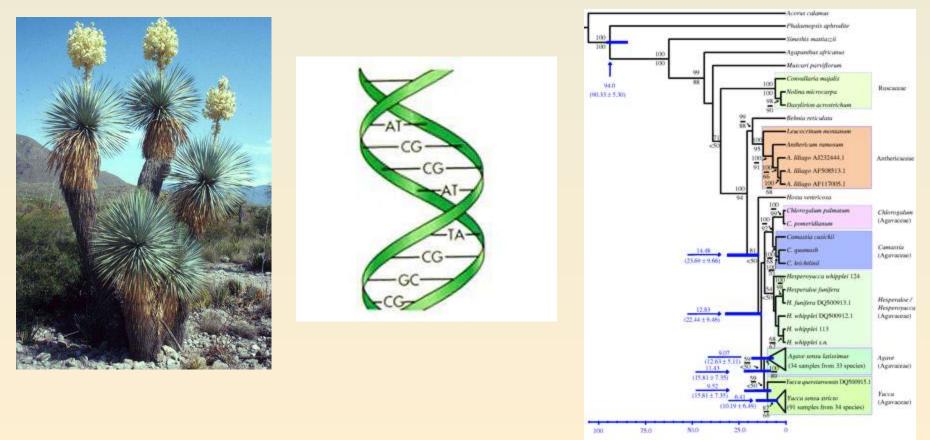
Classification and Phylogenetic Systematics:

A review of concepts with examples from the Agave Family

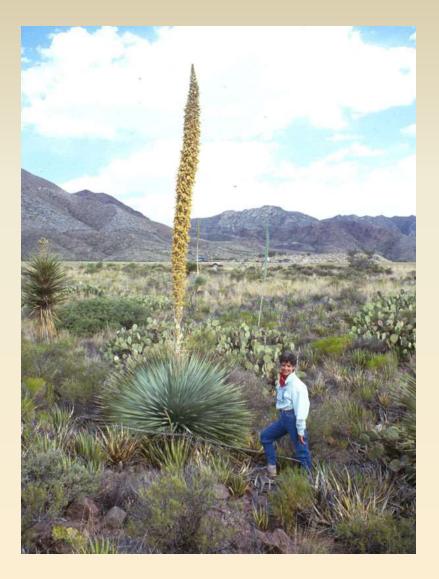
David Bogler Missouri Botanical Garden

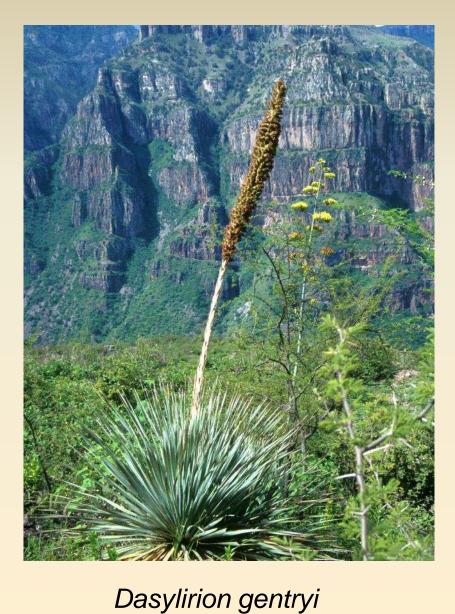


millions of yours before present

- **Taxonomy** the orderly classification of organisms and other objects
- Systematics scientific study of the diversity of organisms
 - Classification arrangement into groups
 - Nomenclature scientific names
 - Phylogenetics evolutionary history
 - Cladistics study of relationships of groups of organisms depicted by evolutionary trees, and the methods used to make those trees (parsimony, maximum likelihood, bayesian)

"El Sotol" - Dasylirion





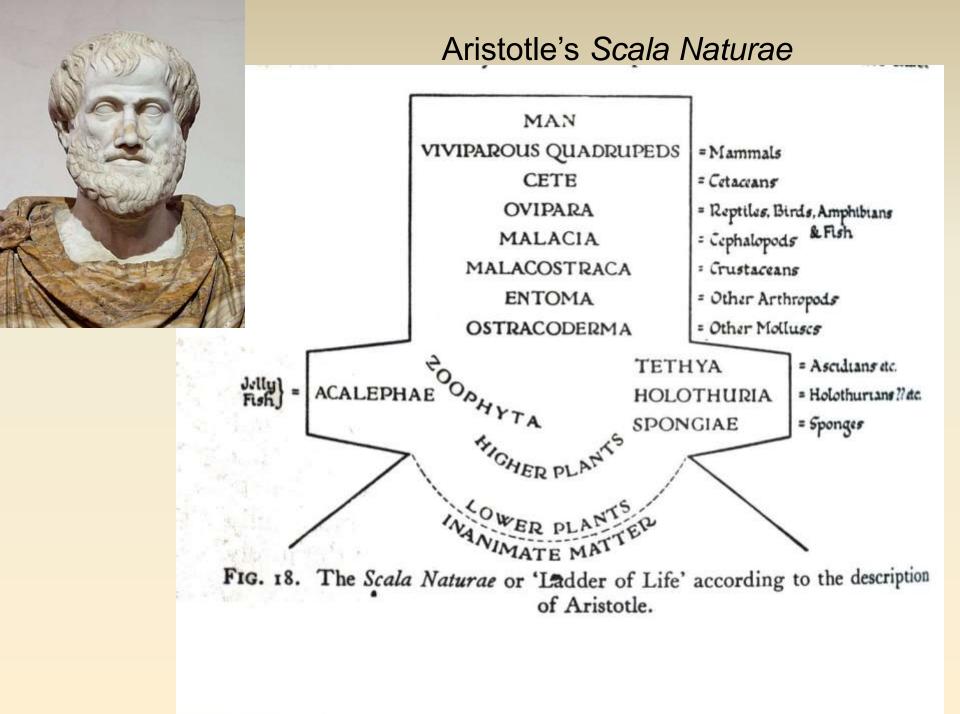
Dasylirion wheeleri



Agave havardii, Chisos Mountains

Agavaceae Distribution







Great Chain of Being

1579, Didacus Valades, *Rhetorica Christiana* hierarchical structure of all matter and life, believed to have been decreed by God

Middle Ages

Ruins of Rome Age of Herbalists Greek Authorities Aristotle Theophrastus Dioscorides Latin was the common language of scholars Plants and animals given Latinized names

Stairway to Heaven



From Llull (1304). Note that Homo is between the plant-animal steps and the sky-angel- god steps.

Systematics - Three Kinds of Classification Systems

- Artificial based on similarities that might put unrelated plants in the same category. - Linnaeus.
- Natural categories reflect relationships as they really are in nature. - de Jussieu.
- Phylogenetic categories based on evolutionary relationships. Current emphasis on monophyletic groups. - Angiosperm Phylogeny Group.



LINNAUS IN HIS LAVIAND DIMES.

Carolus Linnaeus

Tried to name and classify all organism

Binomial nomenclature Genus species

Species Plantarum - 1753

System of Classification

"Sexual System" Classes - number of stamens Orders - number of pistils Clarisf: LINNÆI.M.D. METHODUS plantarum SEXUALIS in SISTEMATE NATURÆ defcripta

N



A View of the Twenty-four Claffes of the SEXUAL SYSTEM of LINNEUS, with their Names and Characters; also the Number and Explanation of Orders, contained in each.

Number of the Claffes.	Their Names and Charafters.	Number of Orders in each.	Their Namet, expressive of the Number of Female Parts or Styles.	Nambo
и.	MONANDRIA. ne fertile flamen, i. having the <i>dathers</i> .	e.}2{1. 2.	Monogynia, Digynia,	. I Z
2. T	DIANDRIA. wo fruitful'Stamina male parts.	or 322.	Monogynia,	I 2 3
3. I	TRIANDRIA.	}3{1 3.	Monogynia, Digynia, Trigynia,	1 2 3
4 F	TETRANDRIA. our ditto, all of equ length, by which it diffinguished from t fourteenth clafs.	is > 3 2 2.	Monogynia, Digynia,	
5- _F	PENTANDRIA.	3334	Monogynia, Digynia, Triginia, Tetragynia, Pentagynia, Polygynia, m	H 1 1 3 4 5 1
6. s	HEXANDRIA. ix ditto, all of equilength, by which this diffinguithed from the fixteenth claft.	nal 2. his 5 3. om 4-	Monogynia, Digynia, Trigynia, Tetragynia, -	1 2 3 4 m
7. s	HEPTANDRIA.	3+22.	Monogynia, Digynia, Tetragynia, Heptagynia, -	1247
8. F	OCTANDRIA.	}+}2.	Monogynia, Digynia, Trigynia, Tetragynia,	
2· 2	ENNEANDRIA.	3.5%	Monogynia, Trigynia, Hexagynia,	0.0 1

10. DECAN-

HEXANDRIA MONOGYNIA: 313

e. Aloë foliis ovato-lanceolatis earnofis apice triquetris: angulis inerme dentatis. Hort. cliff. 131. Hort. apf. 86. Roy. lugdb. 24.

Aloë africana minima atroviridis, fpinis herbaceis numerofis ornata. Beerb. lugdb. 2. p. 131. t. 131. Habitat in Æthiopiæ campestribus. 2 Flores in bec generesspecier un certissimi indices conjun-

gunt Margaritiferam & Arachnoideam.

 ALOE floribus feffilibus reflexis imbricatis prifmaticis. Utaria: Aloè foliis linearibus radicalibus membranaceis. Hort. cliff. 133. Roy. Ingdu. 23.

Aloë africana folio triangulari longiffimo & angustiffimo, floribus luteis foetidis. Comm. bort. 2, p. 29. t. 15. Seb. thef. 1. p. 29. t. 19. f. 3. Habitar ad Cap. b. Spei. 2

AGAVE.

- t. AGAVE follis dentato fpinofis, fcapo ratiofo. Gen. americande.
 - Agave foliis spinoso-dentatis mucronatisque. Hort. apf. 81.
 - Aloë foliis lanceolatis dentatis fpina terminatis radicalibus. Hort. cliff. 130. Roy. lugdb. 22.
 - Aloë folio in oblongum mucronem abeunte. Banb. pin. 286.

Habitat in America ralidiore. 3

2. AGAVE foliis dentatis, staminibus corollam æquanti-vivipara: bus.

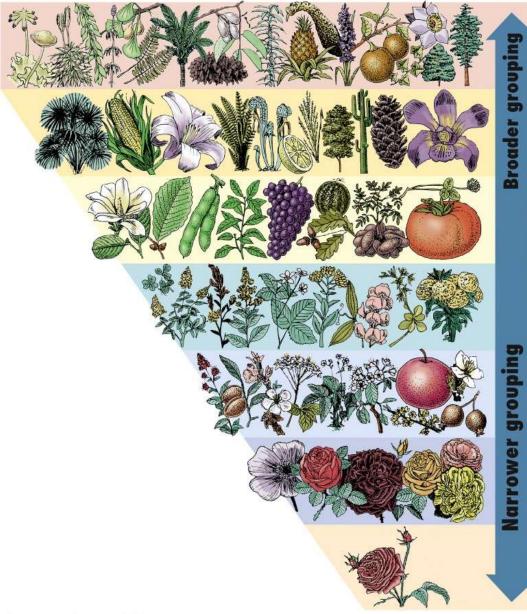
Aloë americana polygond. Comm. rar. 65. t. 65. Habitat in America. Confer. Aloe americana foboliferd. Herm. lugdb. 16. t. 17.

 3. AGAVE foliis dentato-fpinofis, feapo fimpliciffimo. virginica. Gen. nov. 1102: Aloe foliis lanceolatis fpina cartilaginea terminatis, floribus alternis feffilibus. Gron. virg: 152. Habitat in Virginia. 2

4. AGAVE foliis integerrimis. Gen. nov. 1102. fetida: Aloe foliis integerrimis patentiufeulis aculeo-terminatis, radice caulefcente. Hort: cliff. 132.

Aloe americana, viridi rigiditimo. & fortido folio, Piet dicta indigenis. Comm. bort. 2. p. 35. t. 15.

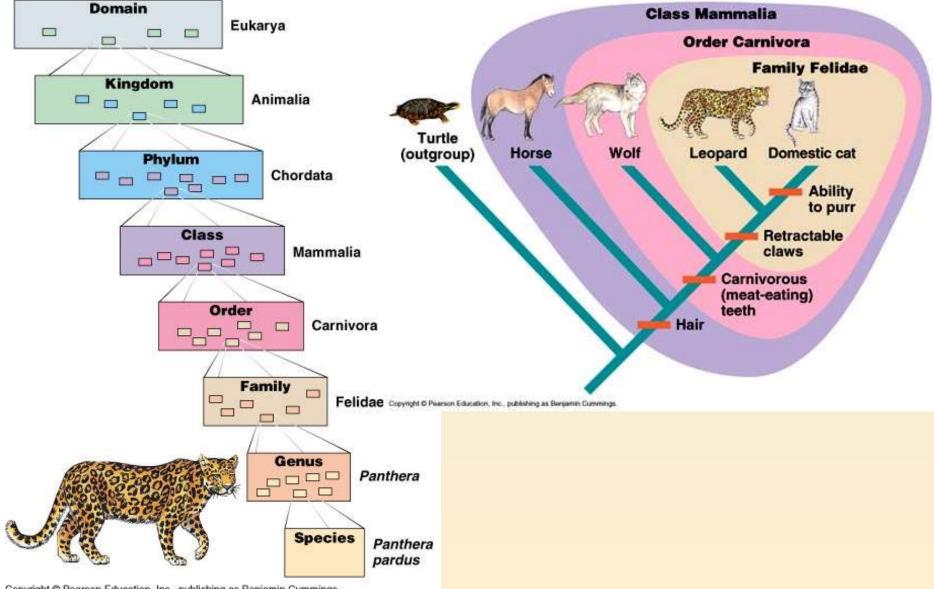
Linnaean Hierarchy



	Kingdom ± 280,000 species	Plantae (plants)
	Phylum ± 250,000 species	Angiospermae (flowering plants)
	Class ± 235,000 species	Dicotyledonae (dicots)
	Order ± 18,000 species	Rosales (roses and their allies)
	Family ± 3,500 species	Rosaceae (rose family)
	Genus ± 500 species	Rosa
-	Species Moss rose	Rosa gallica

Figure 2-6 Discover Biology 3/e © 2006 W. W. Norton & Company, Inc.

Nested box-within-box hierarchy is consistent with descent from a common ancestor, used as evidence by Darwin



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Nomenclature – system of naming species and higher taxa.

International Codes of Nomenclature - separate codes for plants, animals, fungi, bacteria

Binomial nomenclature – scientific name, usually Latinized

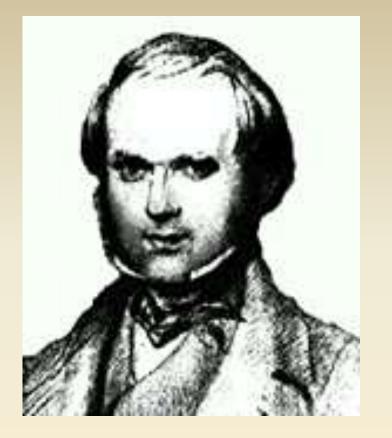
Each species has only **one scientific name**, eliminate confusion.

- **Priority** the correct name will be the **one that was published the earliest**, providing it is acceptable in terms of the rest of the code.
- Causes discontent when it turns out that the name with priority is not the one in common usage, often because it was published in some obscure place. Invalid names = synonyms.

Starts with Linnaeus publications:

Plants - Species Plantarum (1753) Animals - Systema Naturae (1758)

Type Concept - material on which an original description is based, fixes the meaning of a specific name. Type Specimen

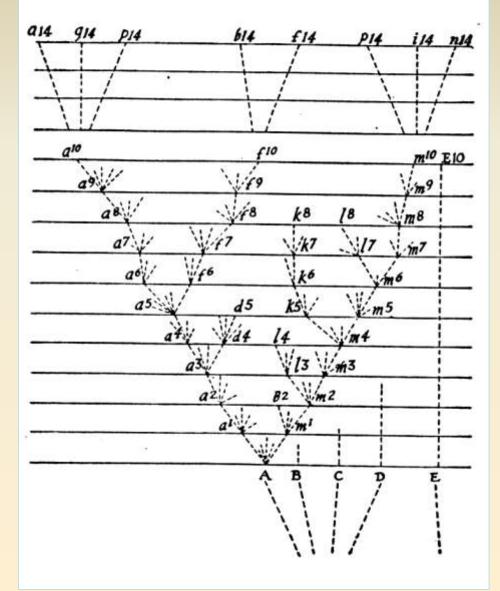


Charles Darwin 1859

•Origin of Species by Natural Selection, or the Preservation of Favored Races in the Struggle for Life

Descent from Common Ancestor

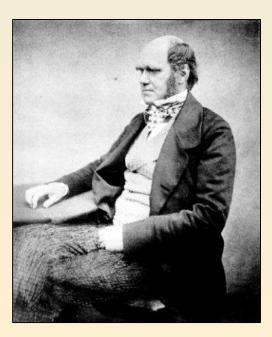
THE ORIGIN OF SPECIES



10

What is a Species?

"Certainly no clear line of demarcation has yet been drawn between species and sub-species – that is, the forms which...come very near to, but do not quite arrive at, the rank of species. ...A well-marked variety may therefore be called an *incipient species*. ...From these remarks it will be seen that I look at the term species as one arbitrarily given."



Darwin, The Origin of Species

Species

- The species is the basic biological unit around which classifications are based.
- However, what constitutes a species can be difficult to define and there are multiple definitions of species in use today.

Species Concepts

Table 15.1 The biological species concept and some recently proposed alternatives

(Futuyma 1997)

- **BIOLOGICAL SPECIES CONCEPT** A species is a group of individuals fully fertile inter se, but barred from interbreeding with other similar groups by its physiological properties (producing either incompatibility of parents, or sterility of the hybrids, or both). (Dobzhansky 1935)
- Species are groups of actually or potentially interbreeding natural populations that are reproductively isolated from other such groups. (Mayr 1942)
- **EVOLUTIONARY SPECIES CONCEPT** A species is a single lineage (an ancestral-descendant sequence) of populations or organisms that maintains its identity from other such lineages and which has its own evolutionary tendencies and historical fate. (Wiley 1978)
- **PHYLOGENETIC SPECIES CONCEPTS** A phylogenetic species is an irreducible (basal) cluster of organisms that is diagnosably distinct from other such clusters, and within which there is a parental pattern of ancestry and descent. (Cracraft 1989)
- A species is the smallest monophyletic group of common ancestry. (de Queiroz and Donoghue 1990)
- **RECOGNITION SPECIES CONCEPT** A species is the most inclusive population of individual biparental organisms that share a common fertilization system. (Paterson 1985)
- **COHESION SPECIES CONCEPT** A species is the most inclusive population of individuals having the potential for phenotypic cohesion through intrinsic cohesion mechanisms. (Templeton 1989)
- **ECOLOGICAL SPECIES CONCEPT** A species is a lineage (or a closely related set of lineages) that occupies an adaptive zone minimally different from that of any other lineage in its range and which evolves separately from all lineages outside its range. (Van Valen 1976)
- **INTERNODAL SPECIES CONCEPT** Individual organisms are conspecific by virtue of their common membership in a part of the genealogical network between two permanent splitting events or between a permanent split and an extinction event. (Kornet 1993)

Source: Coyne (1994).

Species Concepts

- There are many difficulties associated with the definition of "species."
- Definitions that work well for some groups of organisms do not necessarily work for other organisms (extant versus fossil species).
- Some species concepts take evolution into account and attempt to address problems that are associated with a species being an evolving rather than an immutable biological entity.

Morphological Species Concept (MSC)

- A species is a group of organisms that resemble one another and are distinct from other such sets.
- More or less the same as Phenetic Species Concept. What makes two organisms members of the same species is some measure of their <u>similarity</u> across morphological, genetic or behavioral dimensions.

Biological Species Concept (BSC)

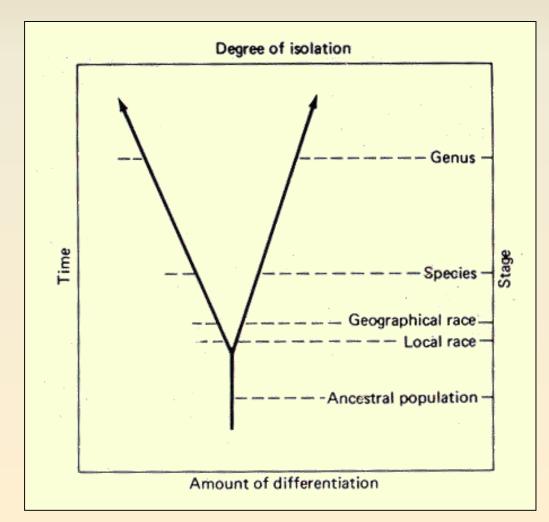
"Species are groups of actually or potentially interbreeding populations that are reproductively isolated from other such groups."

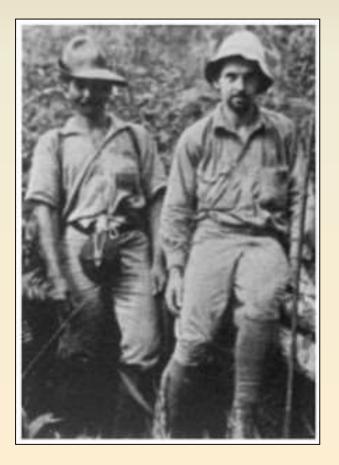
E. Mayr (1942)

Main criterion is reproductive isolation.



Biological Species Concept (BSC) Ernst Mayr - Investigated the question of how species originate, and the importance of geographical isolation





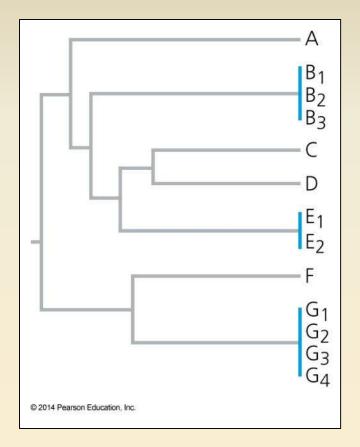
Ernst Mayr

Biological Species Concept (BSC)

- A biological concept of a species is a population or group of populations that are able to *interbreed*, under *natural conditions* to produce *fertile offspring*.
- According to the BSC, speciation occurs when populations evolve reproductive isolating mechanisms.

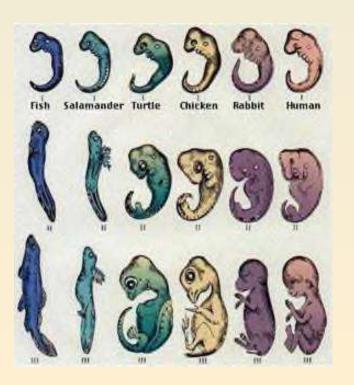
Phylogenetic Species Concept (PSC)

- The phylogenetic species concept <u>emphasizes common descent</u> and covers both sexually and asexually reproducing organisms.
- Under the PSC any population that has become separated and has <u>undergone character evolution</u> will be recognized as a species.



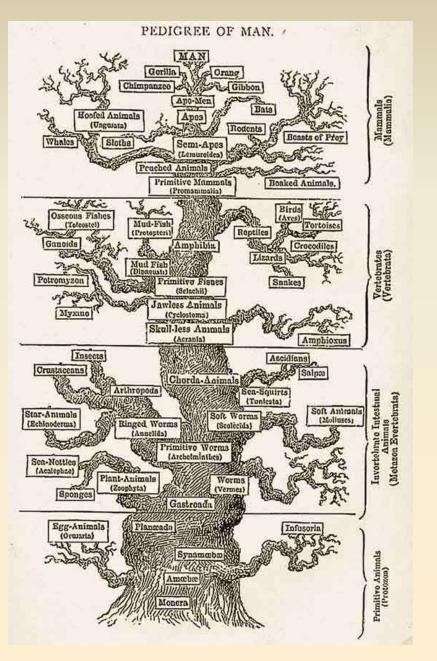
Ernst Haeckel - 1860s

- Coined the terms Phylum, Phylogeny, Ecology
- Drew complete Tree of Life
- Proposed Biogenetic Law -"ontogeny recapitulates phylogeny"

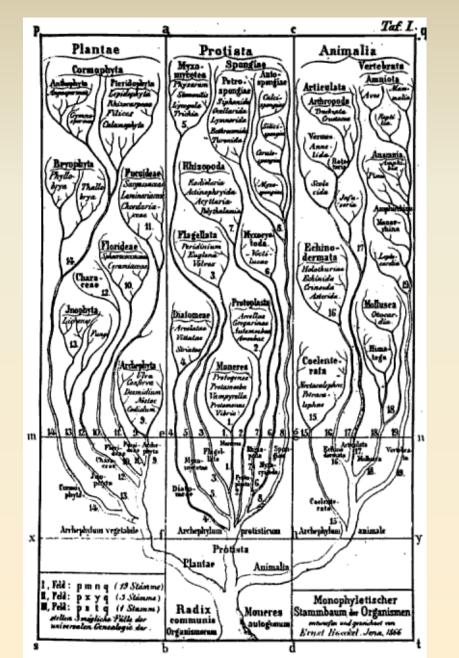




Ernst Haeckel 1860s



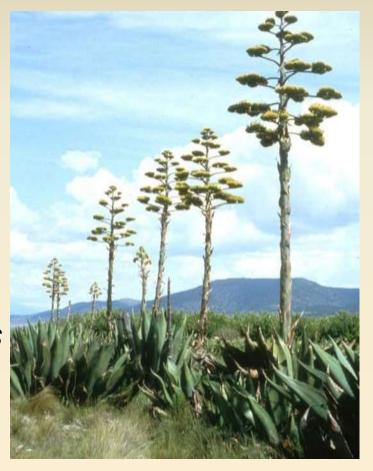
Three Kingdom System



"Agavaceae" sensu stricto - The Agave Family APG3 Asparagaceae – Agavoideae



Yucca Hesperaloe Beschorneria Furcraea Agave Manfreda Polianthes Prochnyanthes



Yucca faxoniana Big Bend N.P., *Texas* *Agave salmiana* Nuevo Leon, Mexico



Beschorneria





Manfreda



Polianthes

Prochnyanthes

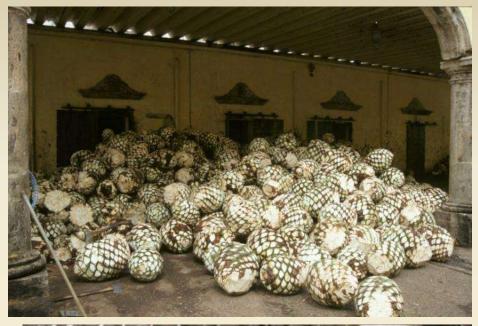






Blue Agave Tequila Plantation, Jalisco, Mexico

El Cuervo Tequila Distillery Tequila, Jalisco













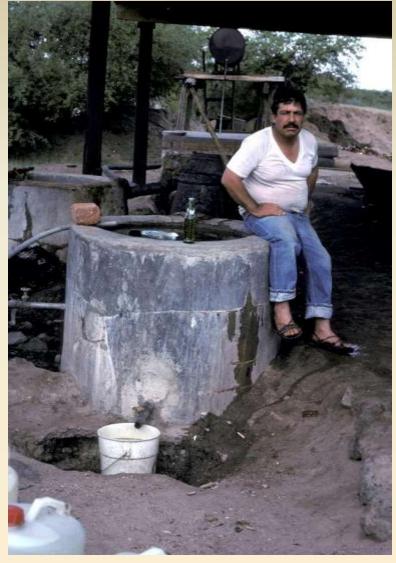
Sampling mescal, Oaxaca

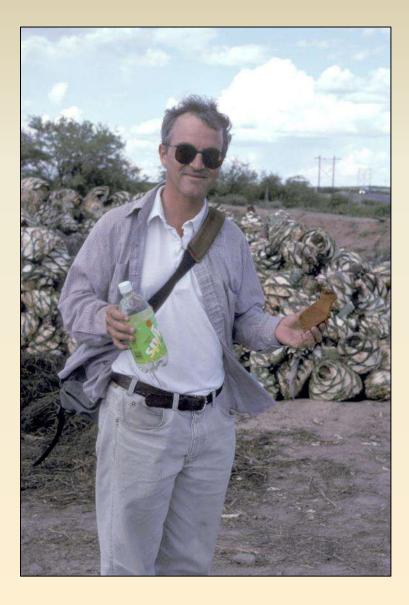


Mescal Distillery, Durango











Dasylirion - Sotol

Dasylirion leiophyllum, Sotol Vista, Big Bend N.P.

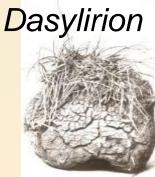
D. wheeleri, Organ Mtns, New Mexico

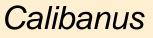
"Nolinaceae" APG3 Asparagaceae – Nolinoideae

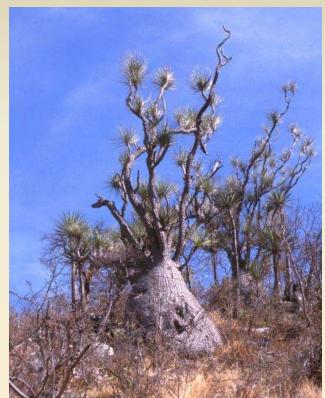






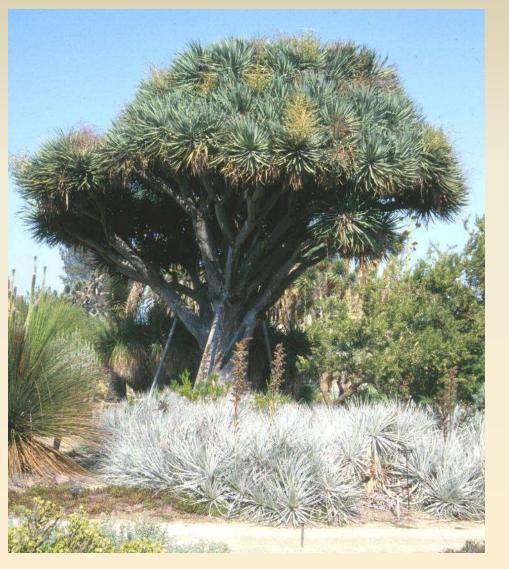






Beaucarnea

"Dracaenaceae" APG3 Asparagaceae – Dracaenoideae





Dracaena draco

Sansevieria trifasciata

"Convallariaceae" Asparagaceae – Convallarioideae



Genera Plantarum - A. L. de Jussieu, 1789 A natural system

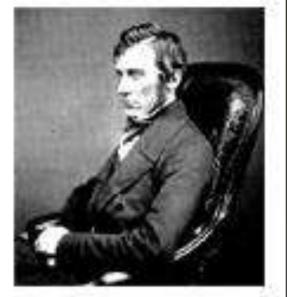
Class III - Plantae Monocotyledones, Stamina Perigyna

Orders:

- Palmae Calamus, Phoenix etc.
- Asparagi Dracaena, Asparagus, Trillium, Convallaria, Dioscorea etc.
- **Junci -** Juncus, Xyris, Commelina, Melanthium, Colchicum, etc.
- **Lilia -** *Lilium, Uvularia, Yucca* etc.
- Bromeliae Puya, Bromelia, Agave etc.
- Asphodeli Aletris, Aloe, Asphodelus, Allium etc.
- **Narcissi -** *Crinum, Hemerocallis, Narcissus, Polianthes, Tacca,* etc.
- Irides Tigridia, Iris, Crocus, Gladiolus etc.



George Bentham



Jospeh Dalton Hooker

Genera Plantarum - 1862-1883

- All genera described anew, in Latin
- Recognized 202 Orders (=Families)
- Dicots come before monocots
- Ranales placed first, Apetalae last
- Importance of epigyny exaggerated

Monocotyledons - 7 Series

- I. Microspermae Orchidaceae, Burmanniaceae
- II. Epigynae Bromeliaceae, Iridaceae
- III. Coronariae Liliaceae, Pontederiaceae
- IV. Calycinae Juncaceae, Palmae
- V. Nudiflorae Pandanaceae, Typhaceae
- VI. Apocarpae Alismaceae
- VII. Glumaceae Gramineae, Cyperaceae

Adolf Engler (1844-1930)



Die naturlichen Pflanzenfamilien Das Pflanzenreich

Treatment of all known plants Families with simple or reduced flowers are placed first (primitive?) Widely used system

Monocots - 10 Reihen (Orders)

Pandanales - Typhaceae, Pandanaceae Helobiae - Alismaceae etc. Glumiflorae - grasses and sedges Principes - palms Syanthae - Cyclanthaceae Spathiflorae - Araceae Farinosae - Bromeliaceae, Commlinaceae etc. Liliiflorae - Liliaceae, Amaryllidaceae etc Scitamineae - Musaceae, Zingiberaceae Microspermae - orchids

Die Naturlichen Pflanzenfamilien (K. Krause, 1930)

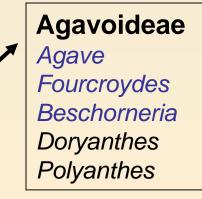
Liliaceae 12 Subfamilies Melanthioideae Herrerioideae Asphodeloideae Allioideae Lilioideae Scilloideae Dracaenoideae Asparagoideae Mondoideae Aletroideae Luzuriagoideae Smilacoideae

Haemodoraceae Amaryllidaceae Dioscoreaceae Iridaceae

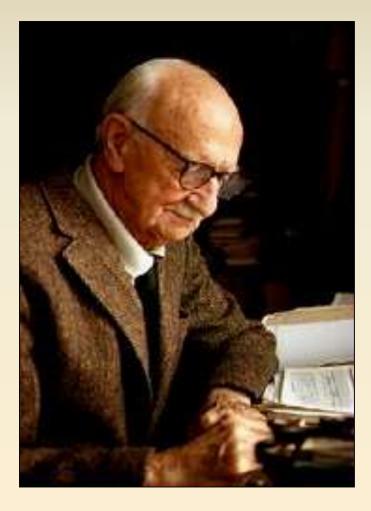
Dracaenoideae Yucceae Nolineae Dracaeneae

Asparagoideae

Asparageae Polygonatae Convallarieae Parideae



John Hutchinson (1884-1972)



Families of Flowering Plants

Dicots fundamentally divided Lignosae Herbaceae

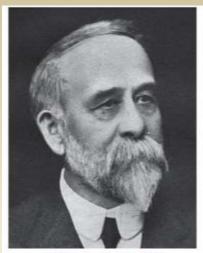


Monocots

Derived from Ranales 3 Divisions, 29 Orders Calyciferae Corolliferae Glumiflorae

Agavaceae - added Yucca

Yucca, Hesperaloe, Agave, Manfreda Furcraea, Beshchorneria, Dasylirion, Nolina, Dracaena, Sansevieria, Cordyline, Phormium, Doryanthes



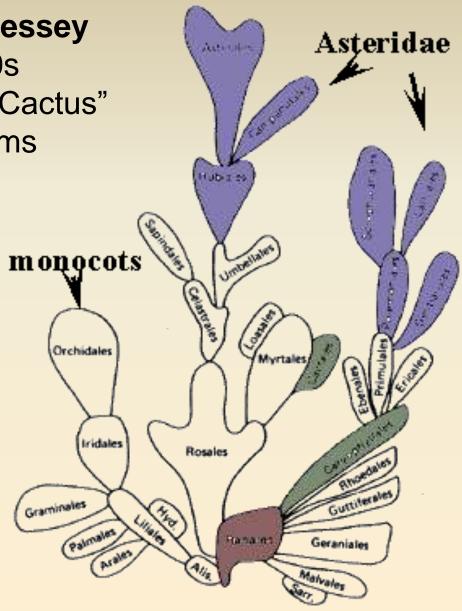
(HARLES EDWIN BESSEY (1845–1915, American)

Charles Bessey

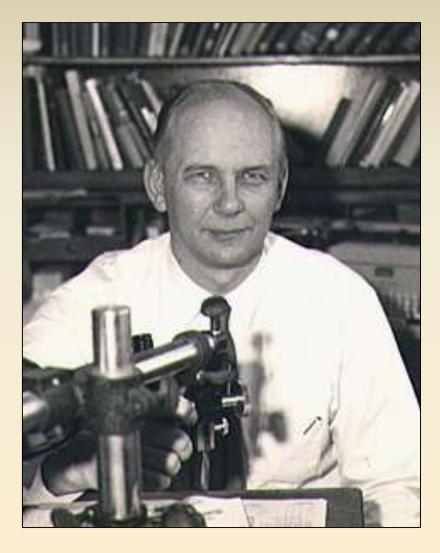
Early 1900s "Bessey's Cactus" Angiosperms

He developed a set of "dicta" (rules) stating which characters were primitive and which were advanced in flowering plants. Not all considered correct today but many are (as Cronquist said, "we are all Besseyans"). Magnolias primitive





Arthur Cronquist (1919-1992)



An Integrated System of Classification of Flowering Plants, 1981

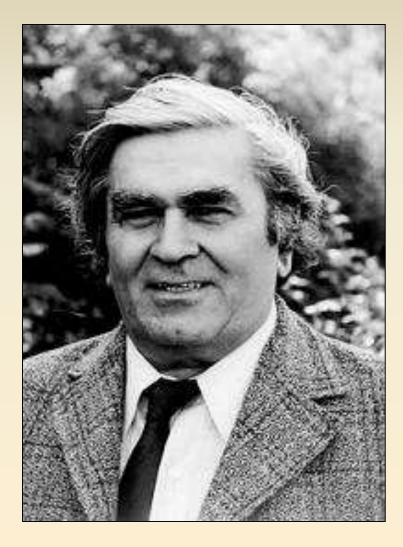
Woody Magnoliids primitive Aquatic origin of monocots

Magnoliopsida 6 subclasses, 55 orders, 352 families

Liliopsida 5 subclasses, 18 orders, 61 families

A "Lumper", did not agree with "cladists" about monophyletic groups

Armen Tahktajan (1910-2009)



Diversity and Classification of Flowering Plants, 1997

System similar to Cronquist

Magnoliopsida - Dicots 7 Subclasses, 20 Superorders Liliopsida - Monocots 3 Subclasses, 8 Superorders Woody Magnoliids primitive Monocots derived from aquatic herbs

A "Splitter" Lilliidae split into many families Agavaceae included *Hosta*

Cronquist's System

Liliales

Phylydraceae Pontederiaceae Haemodoraceae Cyanastraceae Liliaceae Iridaceae Velloziaceae Agavaceae Aloeaceae Xanthorrhoeacea Hanguanaceae Taccaceae Stemonaceae Smilacaceae Dioscoreaceae **Orchidales** Geosiridaceae Burmanniaceae

Orchidaceae

Melanthiaceae Japonoliriaceae Xerophyllaceae Nartheciaceae Heloniadaceae Chionographidaceae

Melanthiales

Tofieldiaceae

Colchicales

Tricyrtidaceae Burchardiaceae Uvulariaceae Campynemataceae Scoliopaceae Colchicaceae Calochortaceae Trilliales Trillaceae

Liliales Liliaceae Medeolaceae

Alstroemeriales Alstroemeriaceae

Takhtajan's System

Iridales

Isophysidaceae Geosiridaceae Iridaceae Tecophilaeales Ixioliriaceae Lanariaceae Walleriaceae Tecophilaeaceae Cyanastraceae Eriospermaceae

Burmanniales

Burmanniaceae Thismiaceae Corsiaceae

Hypoxidales

Hypoxidaceae

Orchidales Orchidaceae

Amaryllidales Hemerocallidaceae

Hyacinthaceae Alliaceae Hesperocallidaceae Hostaceae Agavaceae Amaryllidaceae

Asparagales

Convallariaceae Ophiopogonaceae Ruscaceae Asparagaceae Dracaenaceae Nolinaceae Blandfordiaceae Herreriaceae Phormiaceae Dianellaceae Doryanthaceae Asteliaceae Asphodelaceae Aloaceae Anthericaceae Aphyllanthaceae

Xanthorroeales

Baxteriaceae Lomandraceae Dasypogonaceae Calectasiaceae Xanthorrhoeaceae

Hanguanales Hanguanaceae

Stemonales

Stemonaceae Croomiaceae Pentastemonaceae

Smilacales

Luzuriagaceae Philesiaceae Ripogonaceae Smilacaceae Petermanniaceae

Dioscoreales

Stenomeridaceae Trichopodiaceae Avetraceae Dioscoreaceae

Rolf Dahlgren (1932-1987)

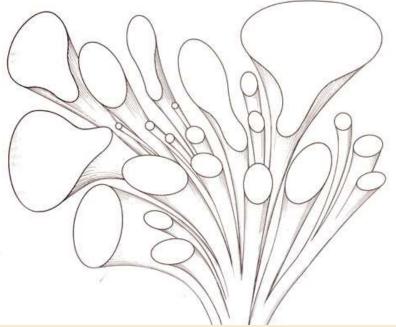


The Families of the Monocotyledons, Dahlgren, Clifford, and Yeo, 1985 System based on work of Huber, 1969 Examined micro-characters of seed coat, cuticle, endosperm, embryo etc **Monocots derived from Dioscoreales-like** dicot ancestor Liliiflorae divided into major groups **Dioscoreales - 7 families** Asparagales - 31 families Liliales - 10 families Melanthiales - 2 families

Rolf Dahlgren 1980s Microcharacters "Lacrymograms"

Helped define Asparagales and families

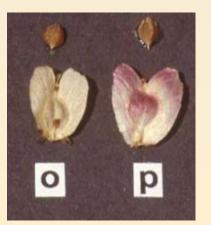


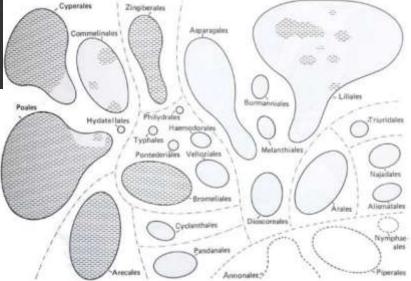


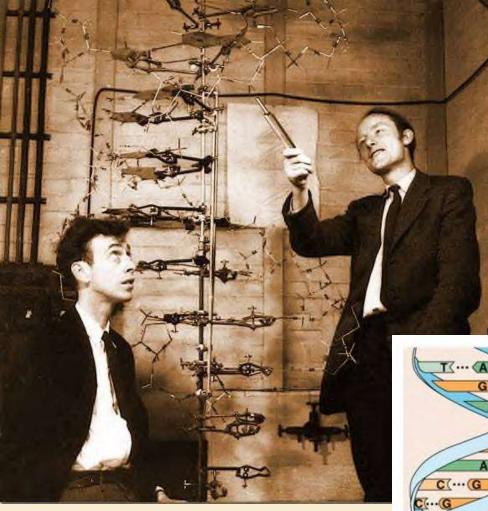




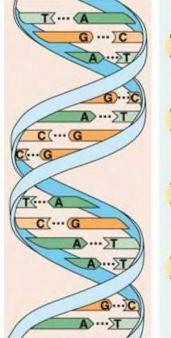


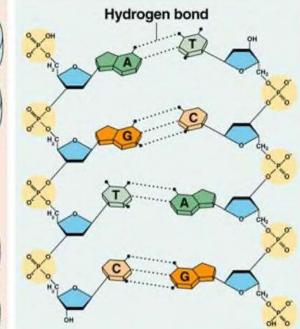






Watson and Crick 1953 Structure of DNA

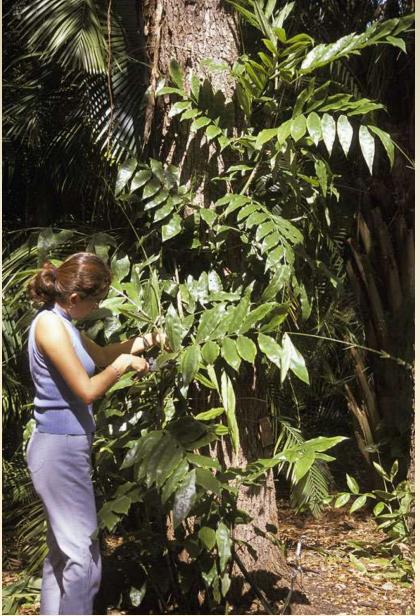






6Addison Wesley Longman, Inc.

Working with Plant DNA







Voucher Specimens

Working with Plant DNA



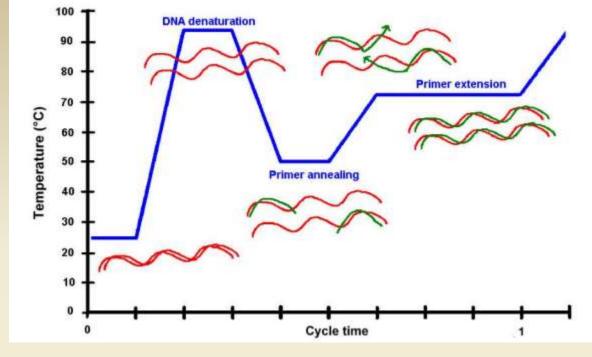


Grinding Tissue

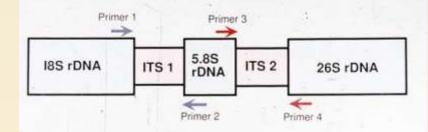
Extracting DNA

DNA Amplification





Internal Transcribed Spacer (ITS)

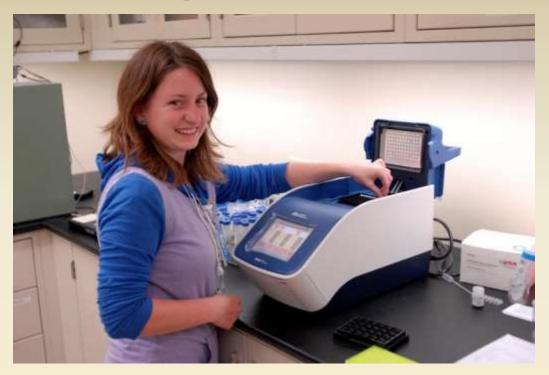


Polymerase Chain Reaction (PCR)

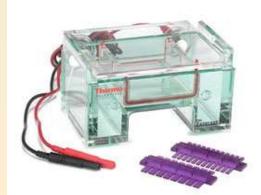


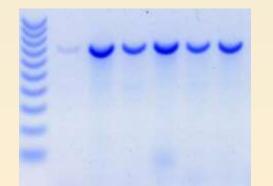
Taq Polymerase 35 Thermal Cycles Amplify DNA

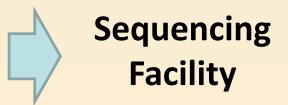
DNA Amplification: PCR



Kelsey Huisman, 2013 REU







PCR Product Gel Electrophoresis – check size

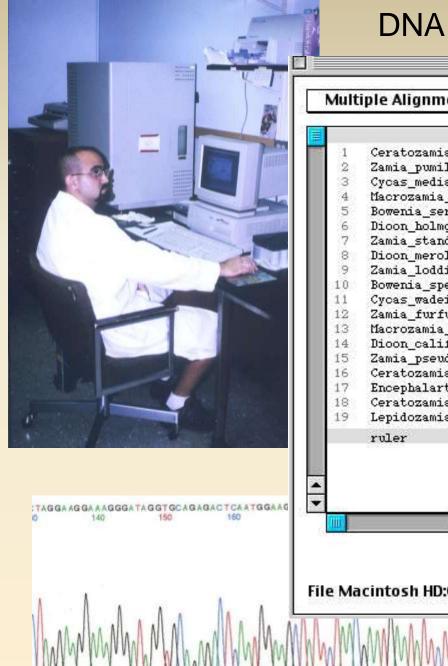
Working with Plant DNA



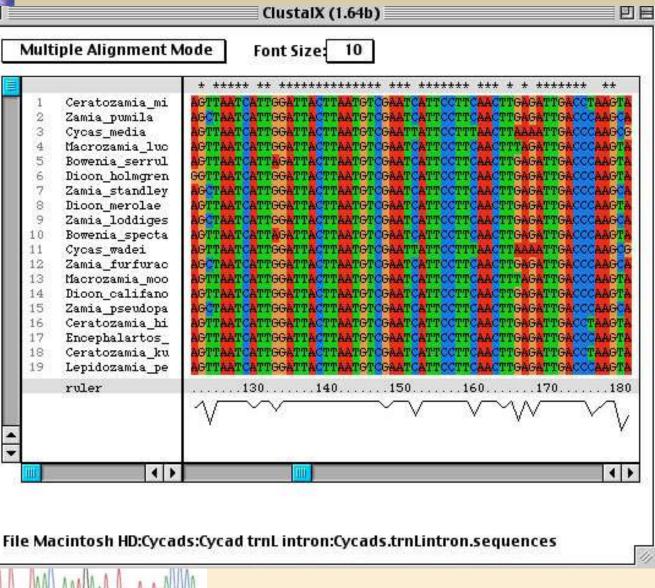
Loading DNA on Agarose Gel



Visualizing DNA

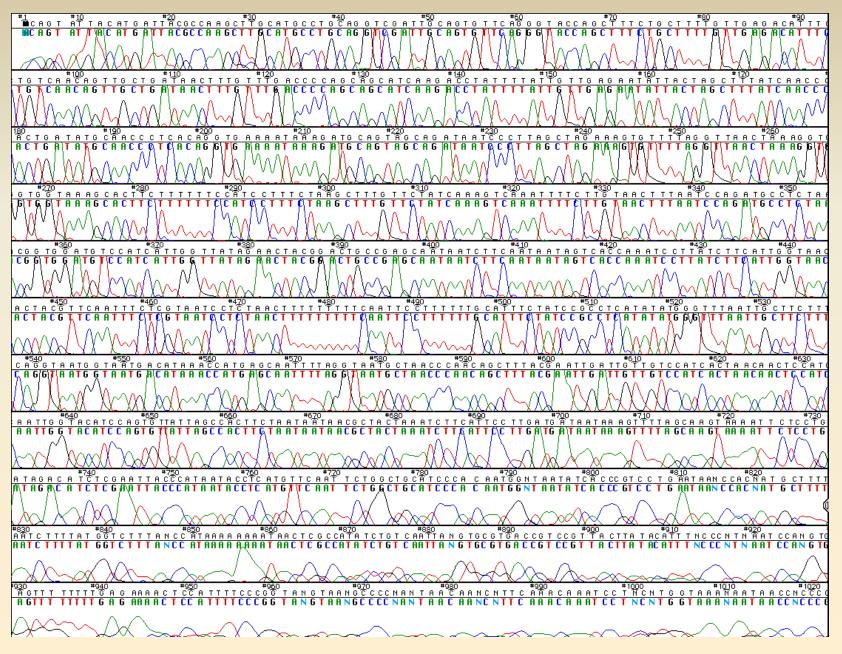


DNA Sequencing



Sequence Alignment

Chromatogram for One Sequence



🔍 DNA Baser v2.61.1 - License: Registered to Heracle - [Assembly window - Contig samples - Mismatches: 10]							
<u>F</u> ile <u>V</u> iew Search <u>E</u> dit Contig Chromatogram <u>W</u> indow <u>I</u> nfo	_ 2 ×						
	>						
Sequence B T G C A G A A G A G G A A T G T G C A A C A T T C T G C C C A C G T C A T C G C T G G C C C C T T G C							
E Sequence A T G C A G A A G A G G A A T G T G C C A C A T T C T G C C C A C G T C A T C G C C T G A C Ruler (x10) 58 59 60 61 62	<u>ี (NNNCN</u> ธ						
CONTIG T G C A G A A G A G A A T G T G C A A C A T T C T G C C C A C G T C A T C G C T G G C C C T G G C							
<u><</u>							
Stop Prev Stop Next Finish Mew contig	Map						
T C C A G G A A A T G C A G A A G A G A G A A T G T G C C A A C A T T C T C T G C C C A C G T C A T C G C T G G C C C C T T G C							
•							
M = M = M = M = M = M = M = M = M = M =	٨						
TC CAG G A A T G C A G A A G A G G A A T G T G C A A C A T T C I C T G C C C A C G T C A T C G C T G G C C C C T G C	ß						
A A T C C A G G A A A T G C A G A A G A G A A T G T G C A A C A T T C T C T G C C A C G T C A T C G C T G C C C T G A C G							
	٨						
$\left(\frac{1}{2} \right) \left(\frac{1}{2} \right) $							
A A T C C A G G A A A T G C A G A G A G A G A A T G T G C A A C A T T C I C T G C C C A C G T C A T C G C T G C G C C C T G A C G							
Contig saved as 'Contig - samples'							

Finally, the Sequence Data for alignment and analysis...

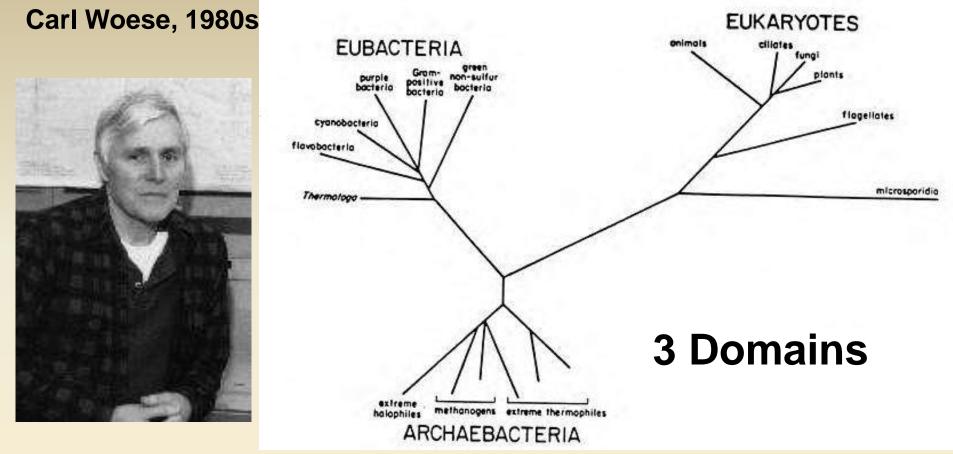
🔍 C:\DNA Baser 3.1\samples\Reference.Fasta	
>Reference	
ACCACAGTCGGAGATAATAGGACGAAGTAANACTGACGNGATACTTTCCCCGAGCTGA ACCTGGTTCTTTTACTAAGTGTTCAAATACCAGTGAACTTAAAGAATTTGTCAATCC GAAGAAAAAGAAGAGAAACTAGAAACAGTTAAAGTGTCTAATAATGCTGAAGACCCC TAAGTGGAGAAAGGGTTTTGCAAACTGAAAGATCTGTAGAGAGTAGCAGTATTTCAT TGATTATGGCACTCAGGGAAAGTATCTCGTTACTGGAAGTTAGCACTCTAGGGAAGGC AATAAATGTGTGAGTCAGTGTGCAGCATTTGAAAACCCCCAAGGGACTAATTCATGGT ATAGAAATGGACACAGAAGGCTTTAAGTATCCATTGGGACATGAAGTTAACCACACAGTC AGAAATGGAAGAAAGTGAACTTGATGCTCAGTATTTGCAGAATACATTCAAGGTTTC TTTGCTCTGTTTTCAAATCCAGGAAATGCAGAAGAGGGAATGTGCAACATTCATGCC GCCCCTTGCGAAGAGGGATATTCTACGGATCGTAATCG	TAGCCTTCCAAGA AAAGATCTCATGT TGGTACCTGGTAC AAAAAACAGAACCA TGTTCCAAAGATA GGGGAAACAAGCAT AAAGCGCCCAGTCA
	-
Copy Paste Load sample Wrap V Co	ount bases Save Close
667 bases	///

Pairwise alignments

43.2%	2% identity; Global alignment score: 374							
	1	.0	20	30	40		50	
alpha	V-LSPADKTNVKAAWGKVGAHAGEYGAEALERMFLSFPTTKTYFPHF-DLSHGSA							
	: :.: .:.	: : ::::	: :	. : : : : :	.: ::	: :::	:.	
beta	VHLTPEEKSAVTALWGKVNVDEVGGEALGRLLVVYPWTQRFFESFGDLSTPDAVMGNP							
	10)	20	30	40	50		
	60	70	8	0	90	100	110	
alpha	QVKGHGKKVA	DALTNAVA	HVDDMPNA	LSALSDLHA	HKLRVDPVN	FKLLSHCLL	VTLAAHL	
	. : : . : : : : :	::	:.:	: : . : :	::.::::	:.:: :.	.:: :.	
beta	KVKAHGKKVLGAFSDGLAHLDNLKGTFATLSELHCDKLHVDPENFRLLGNVLVCVLAHHF							
	60	70	80	90	100	110		
				_				
	120	130	14	0				
alpha	a PAEFTPAVHASLDKFLASVSTVLTSKYR							
···· · · · · · · · · · · · · · · · · ·								
beta	beta GKEFTPPVQAAYQKVVAGVANALAHKYH							
	120 1	.30	140					

Methods of tree estimation

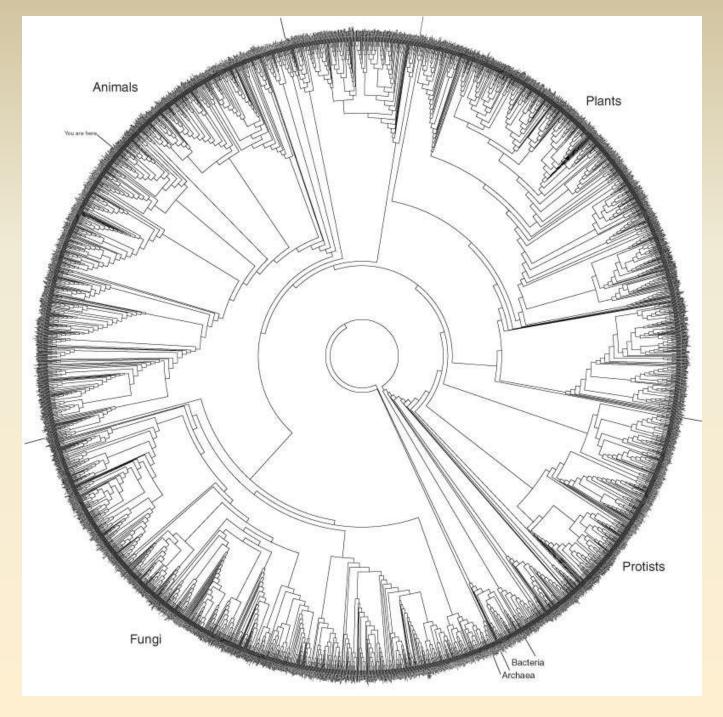
- Character based
 - Maximum parsimony (MP)
 - Fewest character changes
 - Maximum likelihood (ML)
 - Highest probability of observing data, given a model
 - Bayesian
 - Similar to ML, but incorporates prior knowledge
- Distance based
 - Minimum distance
 - Shortest summed branch lengths



Phylogeny based on ssRNA Sequences

Originally the Bacteria and Archaea were thought to be one large diverse family of prokaryotes until Carl Woese and others investigated the evolutionary tree of ribosomal RNAs and found that there were three distinct founding evolutionary domains, then named eubacteria, archaebacteria, and the eukaryotes.

ssRNA 3,000 species



Constructing Phylogenetic Trees

Characters – observations about organisms

morphological structures anatomy chromosomes chemistry DNA base sequences



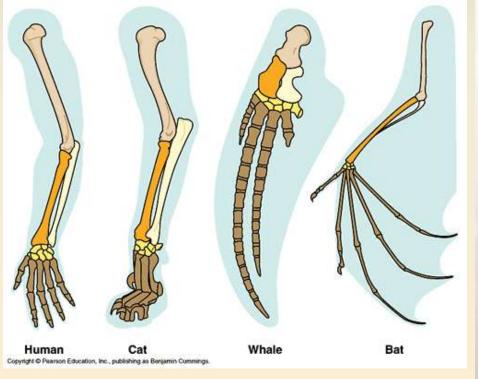






Characters

Homologous characters similarity due to common ancestry Analogous characters similarity due to convergent evolution





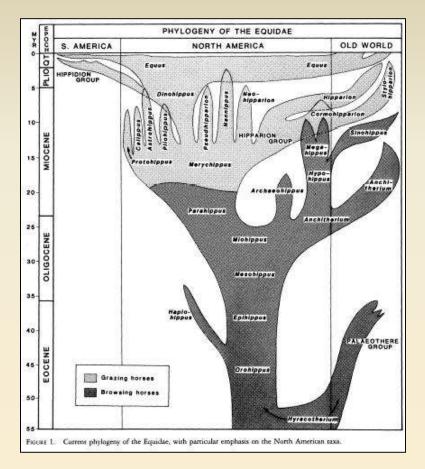
Vertebrate limbs

Euphorbs Cacti

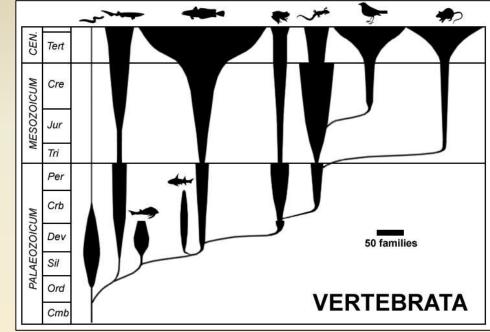
Phylogenetic Trees

- The **phylogeny** of a group of taxa (species, etc.) is its evolutionary history.
- A phylogenetic tree is a graphical summary of this history — indicating the <u>sequence</u> in which lineages appeared and how the lineages are related to one another
- Because we do not have direct knowledge of evolutionary history, every phylogenetic tree is an hypothesis about relationships
- Of course, some hypotheses are well supported by data, others are not

Evolutionary Systematics – 1950s to 1970s Linnaean system incorporated with the Darwinian modern synthesis

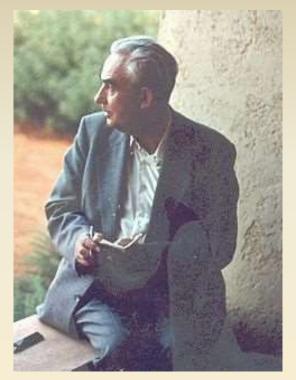


Evolution of Horses summary of data tree, width shows geographic distribution



Spindle Diagram, showing evolution of the vertebrates, showing the relative importance of the major groups through time. The vertical axis represents geological time, and the horizontal axis represents the diversity of each group, in terms of number of families.

Phylogenetic Systematics = "Cladistics"



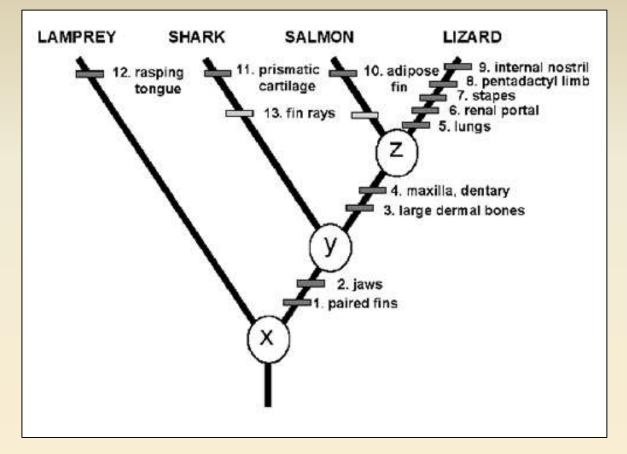
Willi Hennig about 1970

Put forward his ideas in 1950, wrote in his German, so these were completely ignored until 1966 when an English translation of a manuscript was published under the title "Phylogenetic Systematics" (Hennig 1966). Willi Hennig - 1913-1976
Germany, military entomologist, malaria prevention
Taxonomist, specialist in Dipterans (flies)
1950 - Basic outline of a theory of phylogenetic systematics

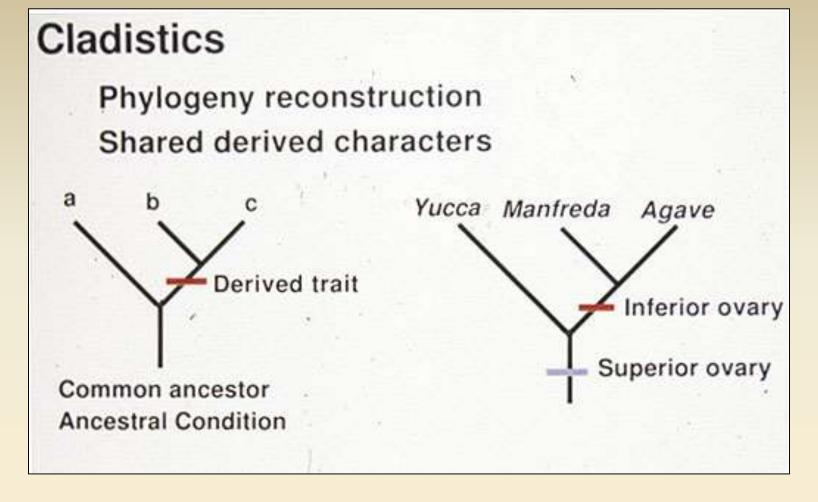
- 1. Relationships interpreted as sisterlineages (clades)
- 2. Synapomorphies determine common ancestry
- 3. Best tree determined by greatest amount of evidence
- 4. Taxonomy/classification should based on inferred pattern of historical relationships (monophyly)

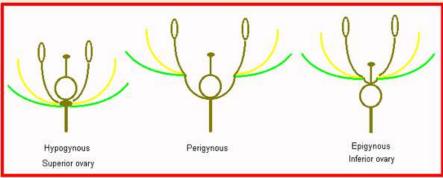
*population genetics, natural selection and adaptation have little to do with the discovery of genealogical relationships

Hennig's Method Phylogenetic Tree based on shared derived characters = "Cladogram"

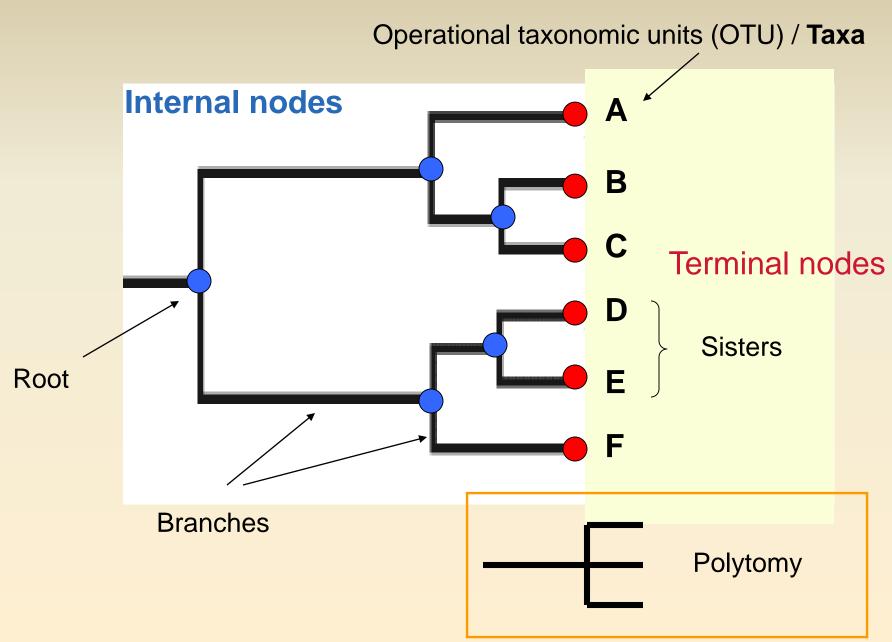


An example of a phylogeny showing characters by which taxa are recognized. Characters 1 – 4 are synapomorphies, 5 – 12 are autapomorphies and 13 is an attribute seen in the salmon and the shark.





Tree Terminology



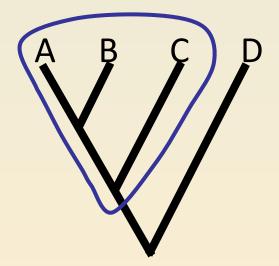
Dendrogram is a broad term for the diagrammatic representation of a phylogenetic tree.

- **Cladogram** is a phylogenetic tree formed using cladistic methods. This type of tree only represents a *branching pattern*; i.e., its branch spans do not represent time or relative amount of character change.
- **Phylogram** is a phylogenetic tree that has branch *spans proportional to the amount of character* change.
- **Chronogram** is a phylogenetic tree that explicitly represents *evolutionary time* through its branch spans.

Phylogeny and Classification

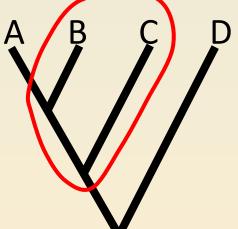
Monophyletic group

Includes an ancestor and all of its descendants



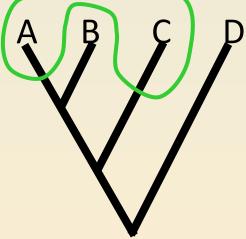
Paraphyletic group

Includes ancestor and some, but not all of its descendants



Polyphyletic group

Includes two convergent descendants but not their common ancestor



How could this happen?

Taxon A is highly derived and looks very different from B, C, and ancestor Taxon A and C share similar traits through convergent evolution

Only <u>monophyletic</u> groups (clades) are recognized in cladistic classification

cp DNA Restriction Site Analysis

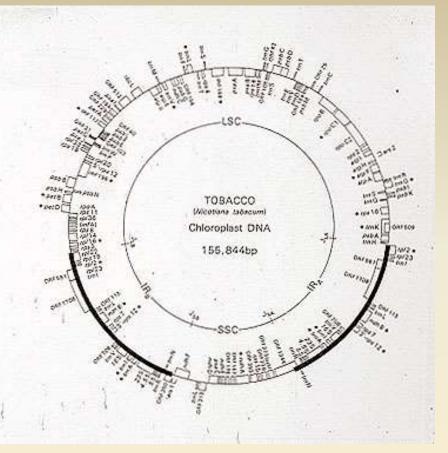
Restriction Enzymes Used:

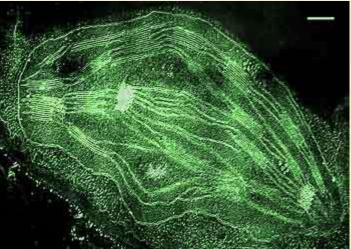
Bam HI Bcl I BstN I Dra I Eco RI Eco RV Hae II Hae III Hha I Hind III Msp I Xho I

Total # Tobacco cpDNA Probes - 40 Subset of Probes Used for this Analysis - 20

100 Restriction Sites Surveyed

18 Autapomorphies 82 sites Shared by Two or More Taxa





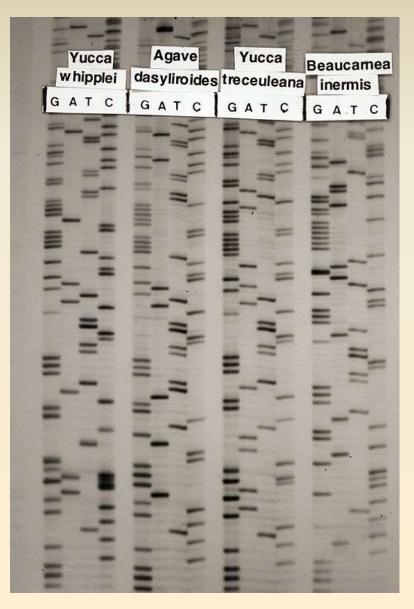
Restriction fragment analysis Digoxigenin-labeled cpDNA probe

rotricht e tophyll ong Iss. mermis I indhe ine alfbanus hocker margina ucca rupicola VIES tolina texana Agave stricts ambdan Marker Sansevieria dichoto anthorrhoez lesperaloe asylition deaucharnea Irlope s sparagas Dasyltrio Manfreda Oracaena Dasylir. Dasylfr. solina . Aloe

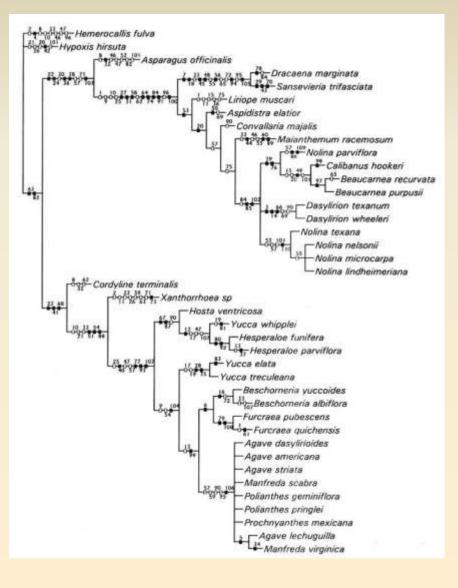


Eco RI Probe 29

Sanger Sequencing P32-labeled dNTP X-ray film



cpDNA Restriction Sites Bogler and Simpson. 1995. Syst. Bot. 20: 191

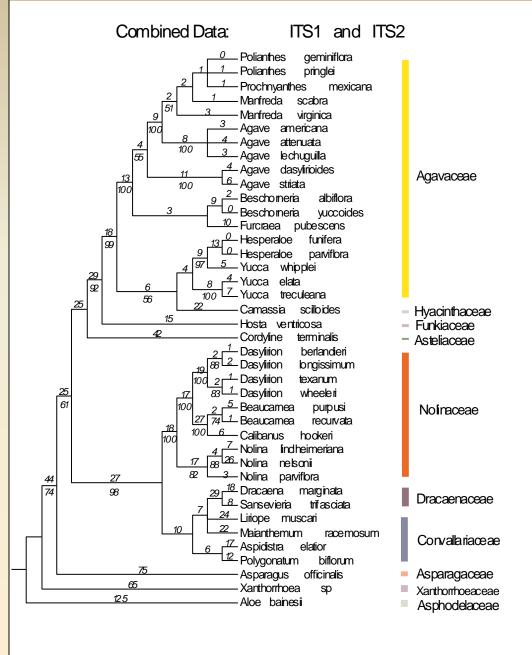


Dracaenaceae Convallariaceae

Nolinaceae

Agavaceae s.s.

ITS1 and ITS2 Strict Consensus 4 Trees 979 Steps CI = 0.659 RI = 0.815

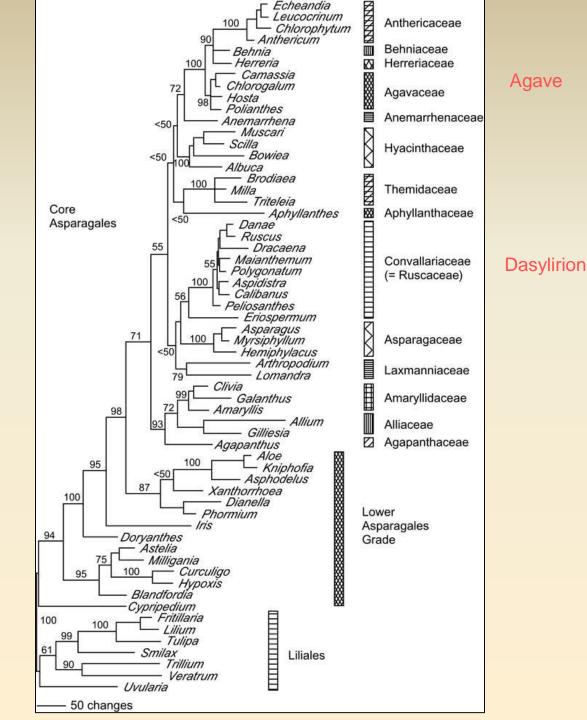


Bogler and Simpson. 1996. AJB 83: 1225-1235.

Asparagales

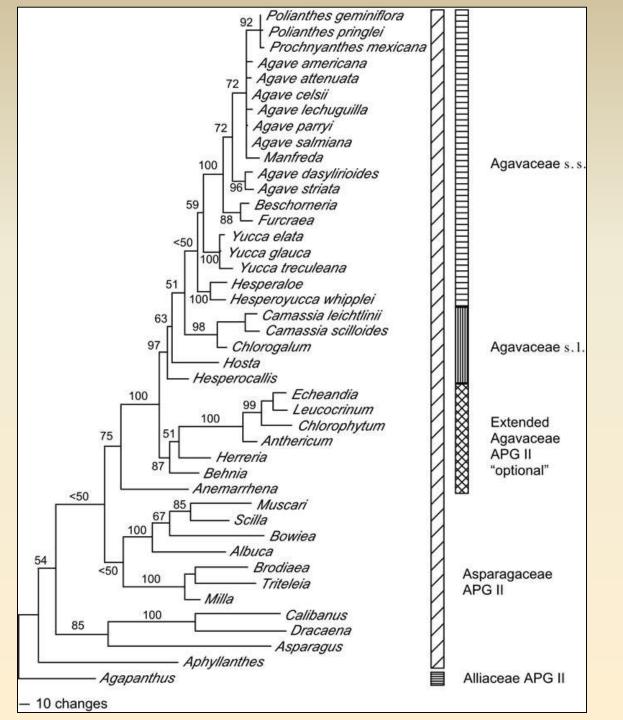
Combined rbcL and ndhF Sequence data

Bogler et al. 2006. Aliso 22: 313–328



Combined rbcL, ndhF, and ITS sequence data

Bogler et al. 2006 Aliso **22**: 313–328



121 Taxa Asparagales three nuclear and plastid DNA coding genes, 18S rDNA (1796 bp), *rbcL* (1338 bp) and *matK* (1668 bp), approx. 4-8 kb

Kim, Joo-Hwan et al. "Molecular Phylogenetics of Ruscaceae *sensu Lato* and Related Families (Asparagales) Based on Plastid and Nuclear DNA Sequences." *Annals of Botany* 106.5 (2010): 775–790. *PMC*. Web. 29 June 2015.

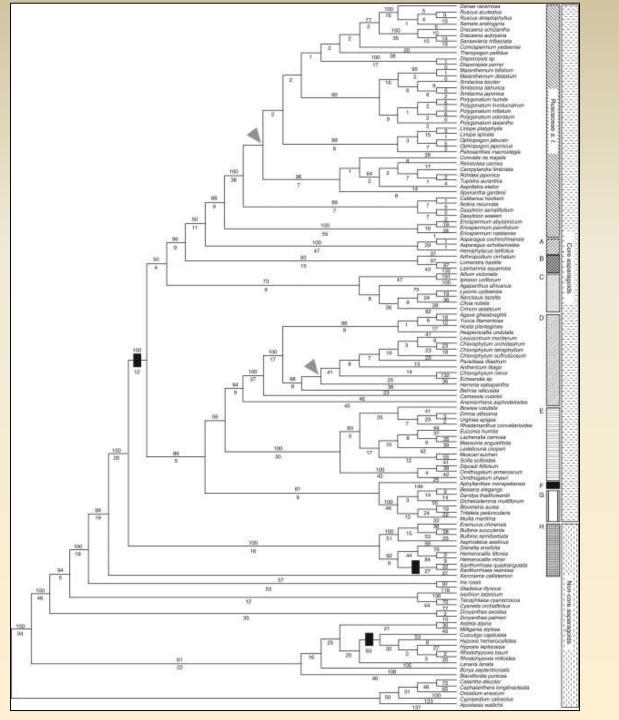
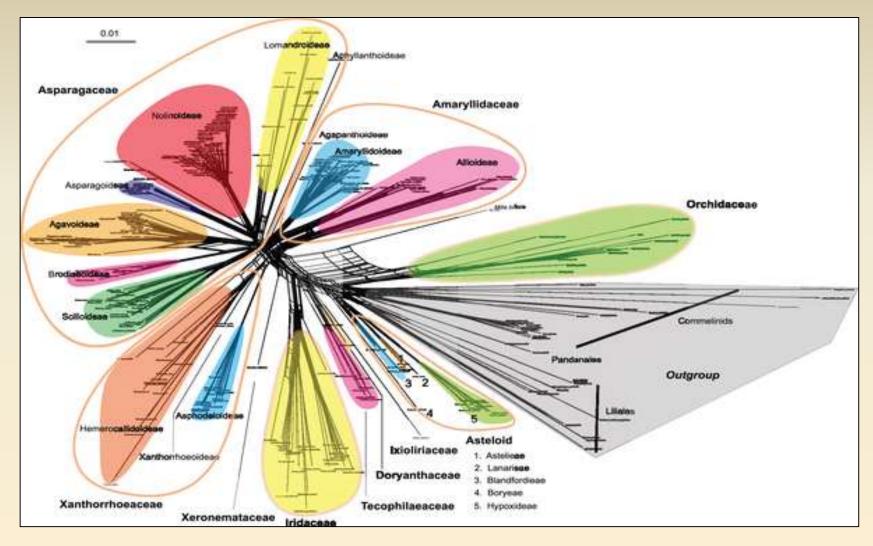


Figure 1. Neighbour net for Asparagales and outgroups.



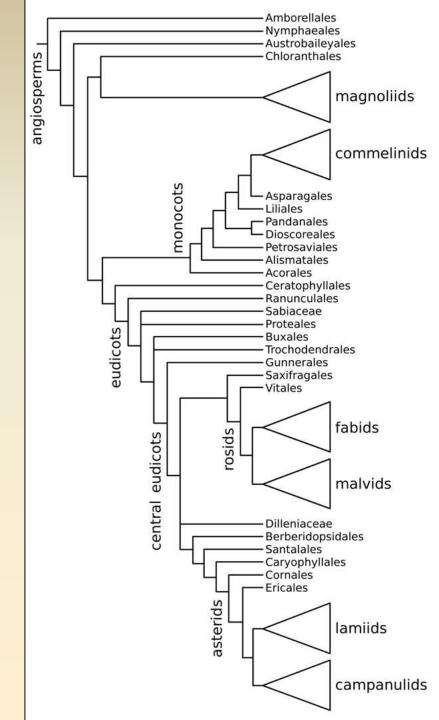
Chen S, Kim DK, Chase MW, Kim JH (2013) Networks in a Large-Scale Phylogenetic Analysis: Reconstructing Evolutionary History of Asparagales (Lilianae) Based on Four Plastid Genes. PLoS ONE 8(3): e59472. doi:10.1371/journal.pone.0059472 http://127.0.0.1:8081/plosone/article?id=info:doi/10.1371/journal.pone.0059472

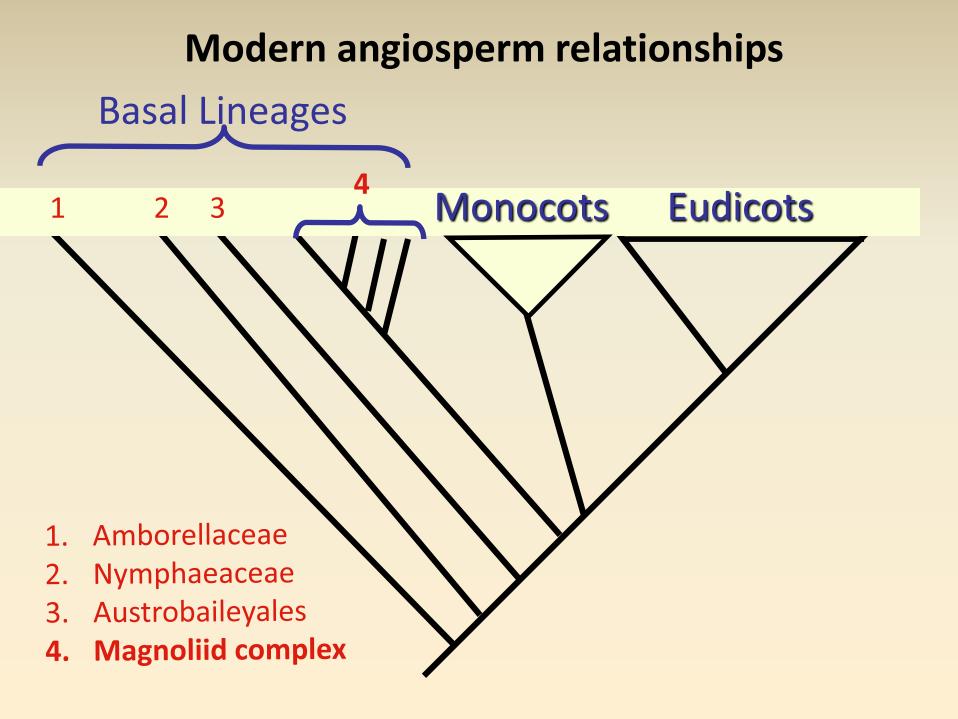


Angiosperm Phylogeny Group APG I1(998, 2003, 2009

Main Features:

- Anonymous, with contributions from many people
- No subclasses, only informal higher groups
- 40 orders
- 462 families
- Based largely on molecular phylogenies
- Taxa are monophyletic





"Agavaceae" APG3 Asparagaceae – Agavoideae



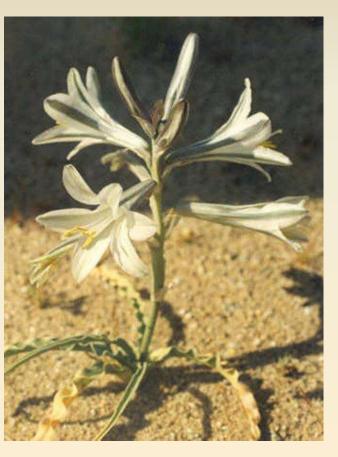
<u>Core Genera:</u> Yucca Hesperoyucca Hesperaloe Beschorneria Furcraea Agave Manfreda Polianthes Prochnyanthes

Basal Genera: Hosta Camassia Chlorogalum Hesperocallis

Agave salmiana

Basal Genera in Agavoideae

Hesperocallis



H. undulata SW U.S., Mex. n = 24, bimodal

Camassia

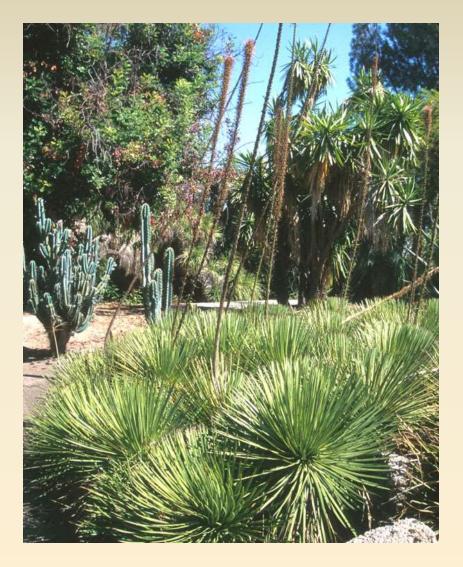
Hosta



Camassia 5 spp. n= 30. bimodal Chlorogalum - 5 spp. n=15, 30 Schoenolirion - 3 spp. Hastingsia - 5 spp.

Hosta 25-many spp., Korea, China n = 30, bimodal

Genus Agave - Group Striatae



Characters thought by Gentry (1982) to be ancestral

- Perennial habit
- Hard, serrulate leaves
- Simple spikes
- Flowers geminate
- Non-dimorphic tepals
- Ovary incompletely inferior
- Reproduce only by seed

Agave striata

All Agave? Name New Genera?

Polianthes

Prochnyanthes

Manfreda

Subgenus Littaea

Subgenus Agave

Agave bracteata etc.

Agave striata etc.

Yucca and the Yucca Moth





Tegiticula – Yucca moth larvae inside fruits



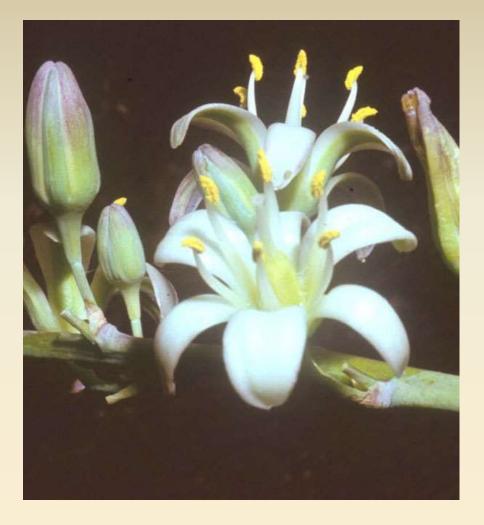
Hesperoyucca whipplei



Monocarpic Capitate Stigma Pollinated by *Tegiticula* moths





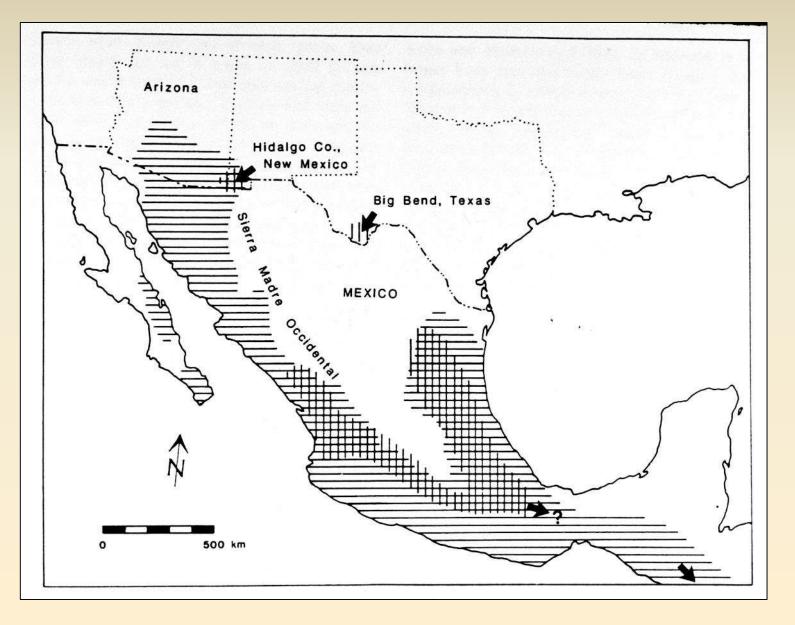


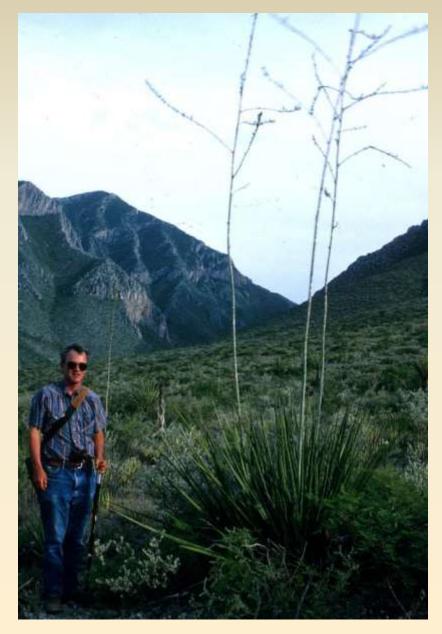


Hesperaloe funifera

Leptonycteris

Nectar-Feeding Bat Range and Migration







Phylogram of Agavaceae based on cpDNA Restriction Sites

Chronogram of the Agavaceae based on cpDNA intergenic spacer sequences.

100

100

940

 (90.33 ± 5.30)

100

100

Acorus calamus

Phalaenopsis aphrodite

Simethis mattiazzii

Agapanthus africanus

Muscari parviflorum

Convallaria majalis

Nolina microcarpa

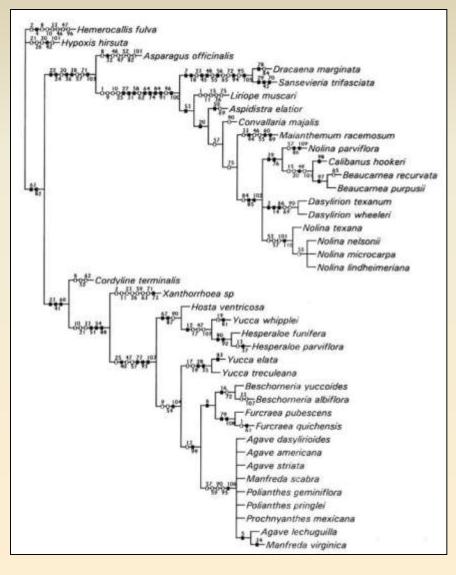
Dasylirion acrostrichum

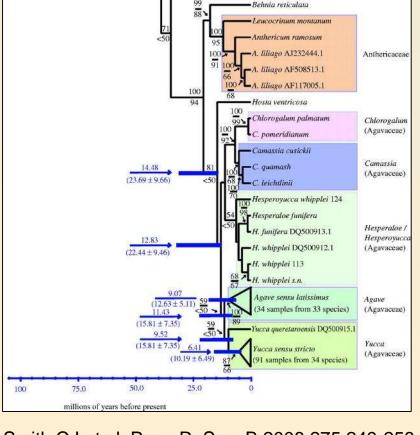
Ruscaceae

101

10

90



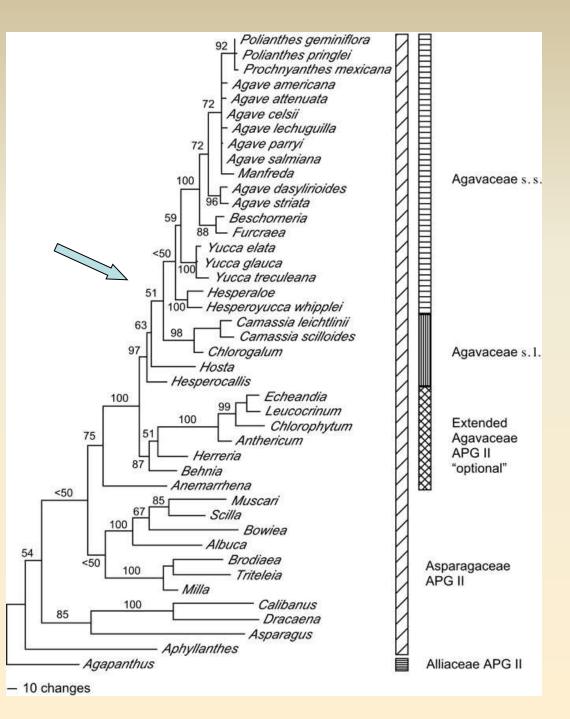


Bogler and Simpson. 1995. Syst. Bot. 20: 191

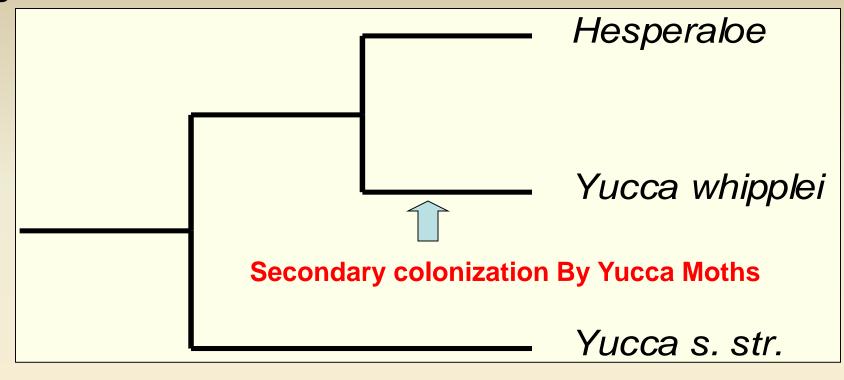
Smith C I et al. Proc. R. Soc. B 2008;275:249-258

Combined rbcL, ndhF, and ITS sequence data

Bogler et al. 2006 Aliso **22**: 313–328

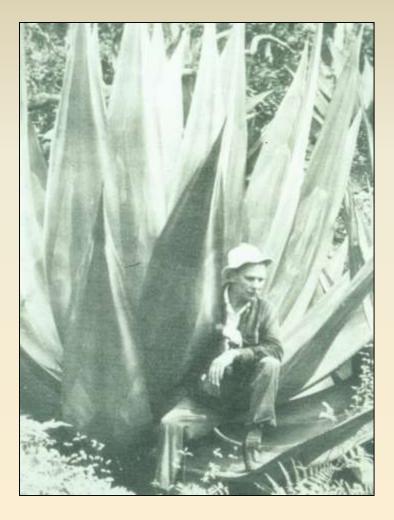


Bogler et al. 1995. PNAS 92:6864-6867





"Groups" of Agave (Gentry, 1982)



Howard Scott Gentry

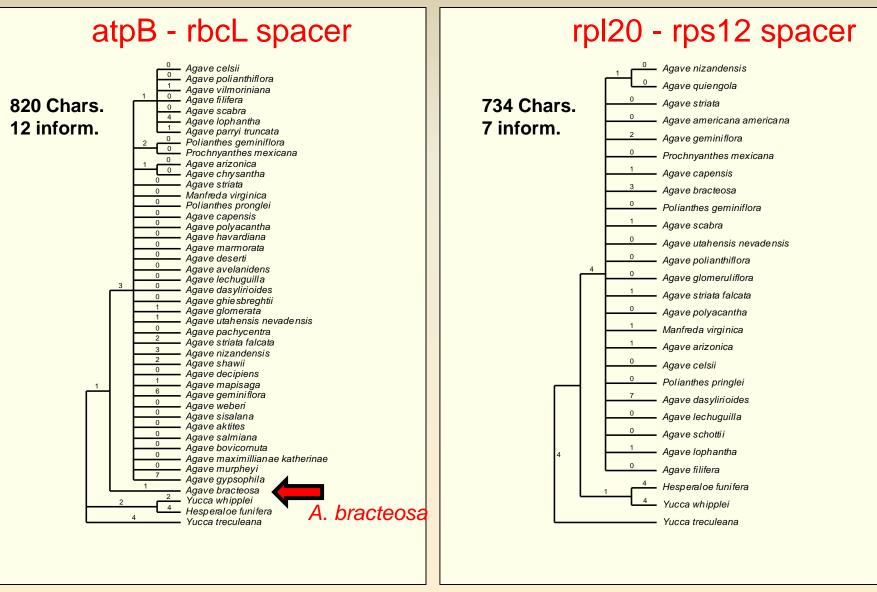
Subgenus Littaea:

Group Amolae Group Chloripetalae Group Filiferae Group Marginatae Group Parviflorae Group Polycepahalae Group Striatae Group Urceolatae

Subgenus Agave:

Group Americanae
Group Campaniflorae
Group Crenatae
Group Deserticolae
Group Ditepalae
Group Hiemeflorae
Group Marmoratae
Group Parryanae
Group Rigidae
Group Salmianae
Group Sisalanae
Group Umbelliflorae

Chloroplast Gene Spacers in Agave



What to do?

Agave bahamense – Andros Island



Agave sp. Whale Point, Eleuthera, Bahamas



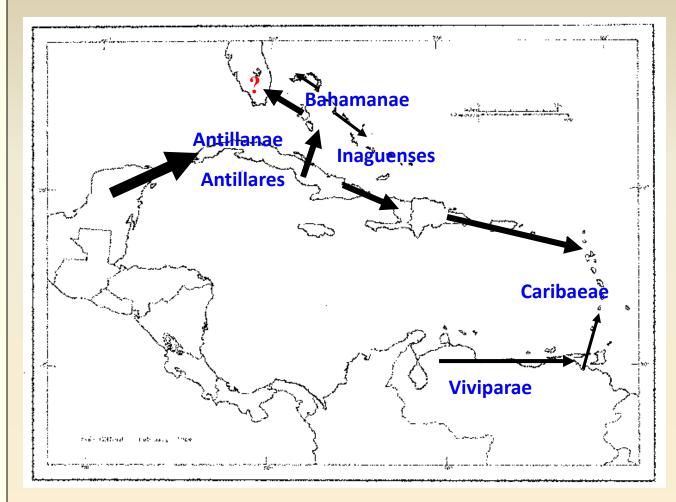
Agave in the West Indies and Florida

Agaves of the West Indies (mostly from Trelease, 1913) Group Bahamanae - 6 spp. Group Inaguenses - 2 spp. Group Antillanae - 13 spp. Group Antillares - 5 spp. Group Caribaeae - 15 spp Group Viviparae - 6 spp.

Introduced Group Rigidae A. angustifolia group Group Sisalanae A. sisalana group

Agaves in Florida

A. decipiens A. desmettiana A. neglecta



Pleistocene Sea Level Fluctuations?

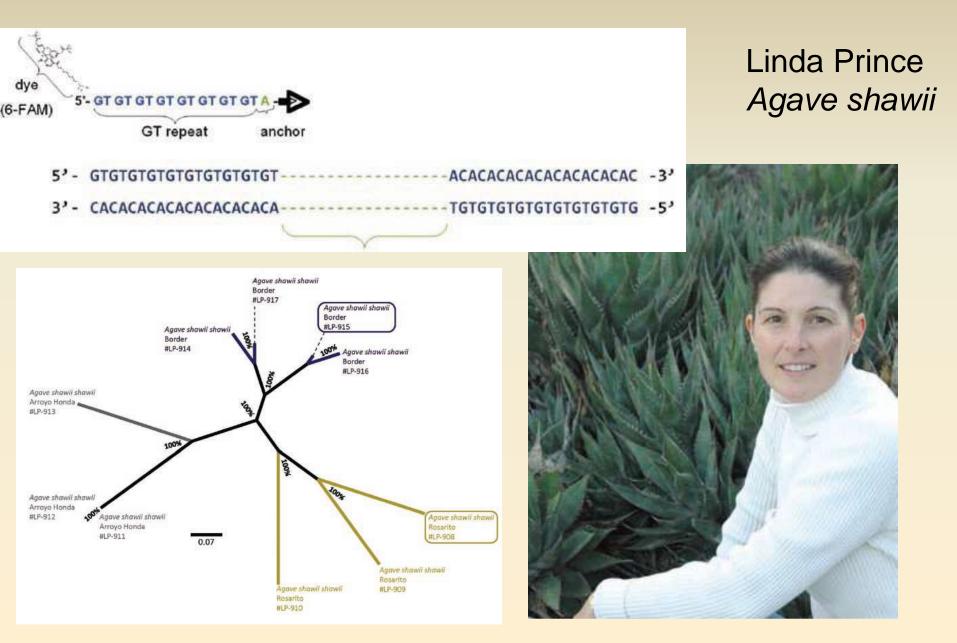
Agave eggersiana St. Croix, U.S. Virgin Islands ~200 individuals left



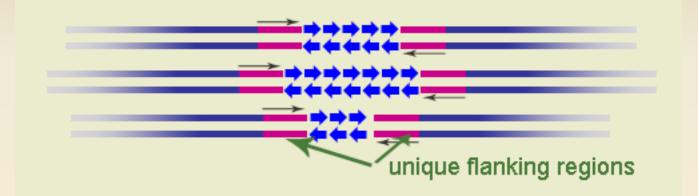
Agave shawii - Shaw's Century Plant



Inter-Simple-Sequence-Repeats - ISSRs



SSRs - Simple Sequence Repeats (= Microsatellites) Short repeating sequences scattered throughout the genome, e.g..GTGTGTGTGTGTGT, or CATCATCATCATCAT The number of SSRs is highly variable among individuals



Two Kinds of Markers Use SSRs

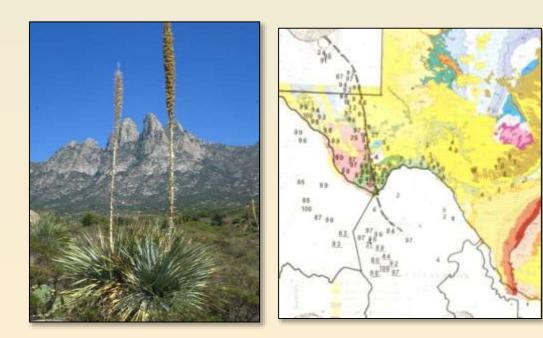
ISSRs – Inter-Simple-Sequence-Repeats Repeating unit used as a primer to amplify region in between SSRs

Microsatellites

Flanking regions used to amplify SSR repeating units

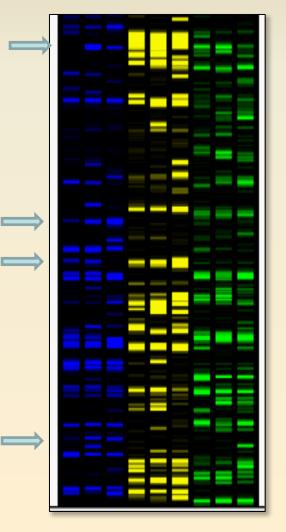
Hybrids between *Dasylirion wheeleri* and *D. leiophyllum* in west Texas?

 D. wheeleri - Organ Mtns.
 D. wheeleri/leio. - Hueco Tanks Putative hybrid
 D. leiophyllum - Chinati Mtns.

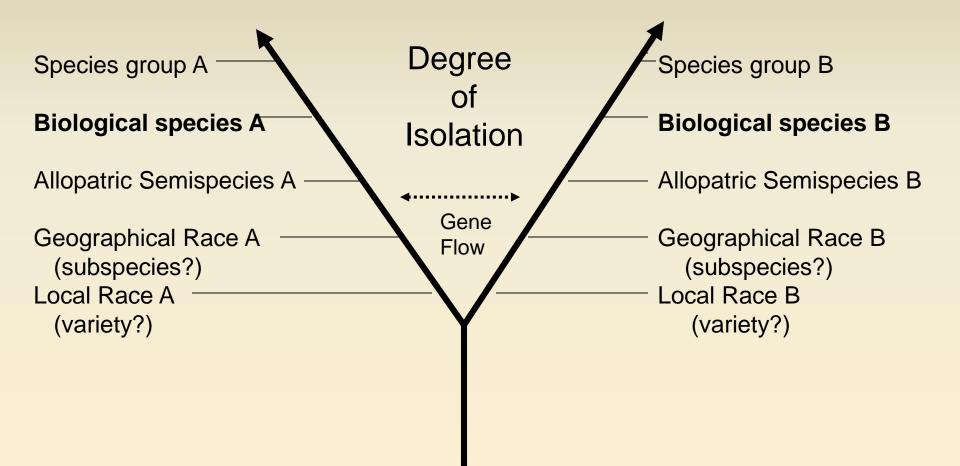


Need to look at larger sample size

123123123

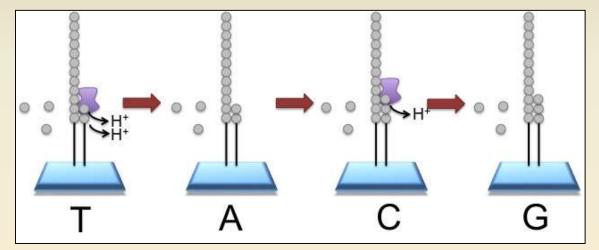


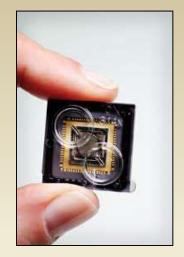
Stages in Divergence Leading to Biological Species from V. Grant, 1981



NGS in a Nutshell

DNA is fragmented. Adapters are added. One molecule is placed on a bead. Each bead is placed in a single well on a slide





Semiconductor Chip

Molecules are amplified on the bead by emulsion PCR

Slide is flooded with a single species of dNTP, along with buffers and

polymerase, one NTP at a time.

The pH is detected in each of the wells, as each H⁺ ion released will decrease the pH.

The changes in pH allow us to determine if that base, and how many thereof, was added to the sequence read. Computer keeps track.

The dNTPs are washed away, and the process is repeated cycling through the different dNTP species.

Sequence fragments are assembled into fragments by software

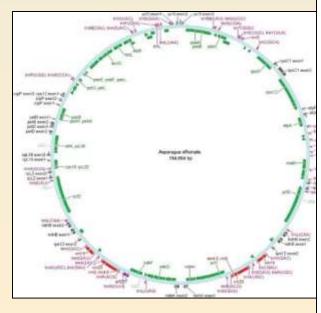


Sequence fragments can also be screened for microsatellite regions New NGS population techniques on horizon - RADSeq

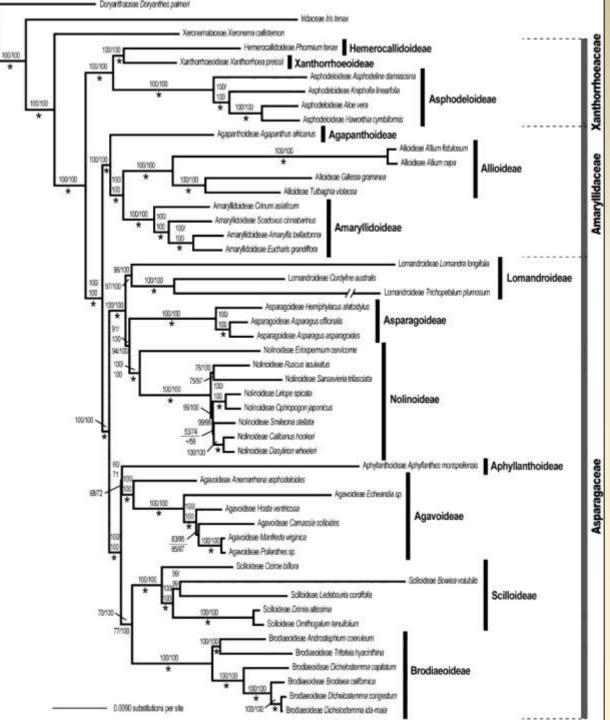
Whole Genome Sequencing

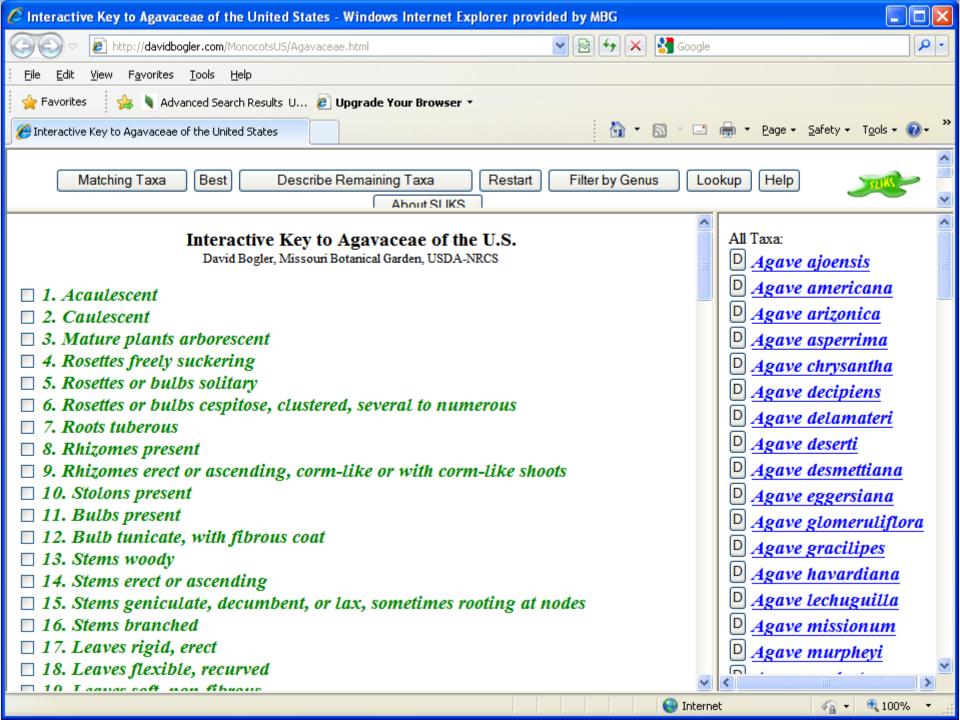
Steele et al. 2012 Am. J. Bot. 99:330-348

79 gene markers + other data sets ML tree



Asparagus cp genome







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