







## . Overview



B. Phylogenetic Evidence1. Characters & Character states

b. Molecular Characters (e.g., DNA sequence characters)

> Protein coding genes (e.g., cox1 or rbcL) Structural RNA genes (e.g., 18S or 26S rDNA)

# I. Overview



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> Protein coding genes (e.g., cox1 or rbcL) Structural RNA genes (e.g., 18S or 26S rDNA)

e.g., Position 36 in rbcL gene: 0 = A; 1 = G; 2 = C; 3 = T.

or

e.g., Position 36 in rbcL gene: A; G; C; T.

# I. Overview



B. Phylogenetic Evidence1. Characters & Character states

b. Molecular Characters

(e.g., amino acid sequence characters)

e.g., Position 5 in rubisco: 0 = glutamine; 1 = proline; etc.



#### I. Overview



B. Phylogenetic Evidence

c. Primary Homology Assessment (during character coding and scoring)

e.g., PETAL COLOR: 0 = red; 1 = blue

Position 4 in rbcL A ; G; C; or T

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	d Data N						
	d. Data N	latrix					
	VERTEBRAE	LEGS	ENDOTHERMY	FUR	MAMMARY GLANDS	OPPOSABLE THUMB	TAI
Snail	0	0	0	0	0		•
Fish	1	0	0	0	0	-	1
Lizard	1	1	0	0	0	0	1
Bird	1	1	1	0	0	0	1
Cow	1	1	1	1	1	0	1
Monkey	1	1	1	1	1	1	1
Gorilla	1	1	1	1	1	1	0
Human	1	1	1	1	1	1	0
*Note: a	"-" denotes that	t the ch	aracter is inapplic	able-i.	e., not relevan	-for that spec	ies.



## I. Overview



C. Recognizing synapomorphies to resolve cladogram

1. Uses the OG to root the cladogram (cladogram is rooted Between the OG and the IG)

2. Parsimony



















































































