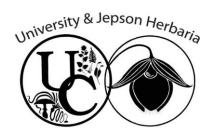
# Conserving the evolutionary diversity of the California flora

Matthew Kling, Brent Mishler, Bruce Baldwin, Andrew Thornhill, David Ackerly



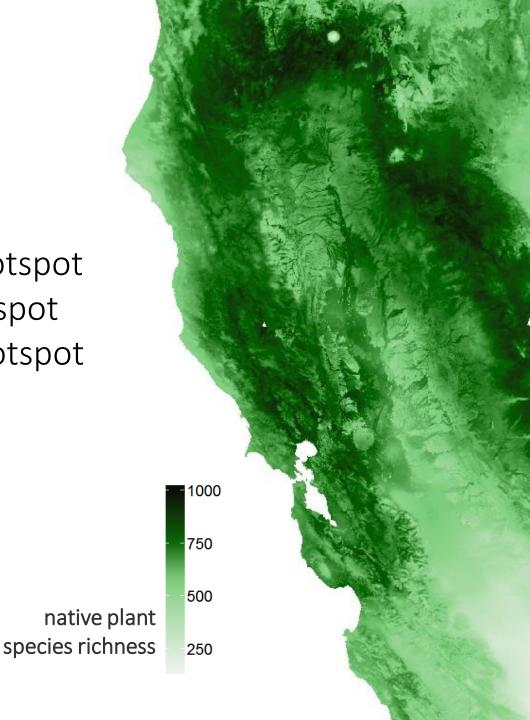






### California:

- World biodiversity hotspot
- Biodiversity data hotspot
- Land conservation hotspot



### Los Angeles Times

CALIFORNIA

LOCAL

Couple donates \$165 million to preserve 24,000 acres at Point Conception



By THOMAS CURWEN DEC 22, 2017 | 3:00 AM



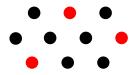
### Los Angeles Times

CALIFORNIA

Couple donates \$165 rQUESTION: 24,000 acres at Point Conception



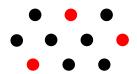
Species richness

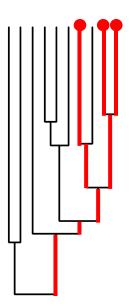


3 species

Species richness

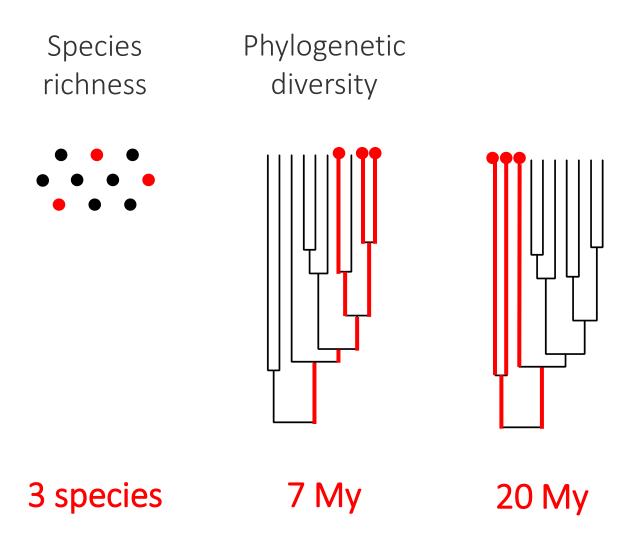
Phylogenetic diversity

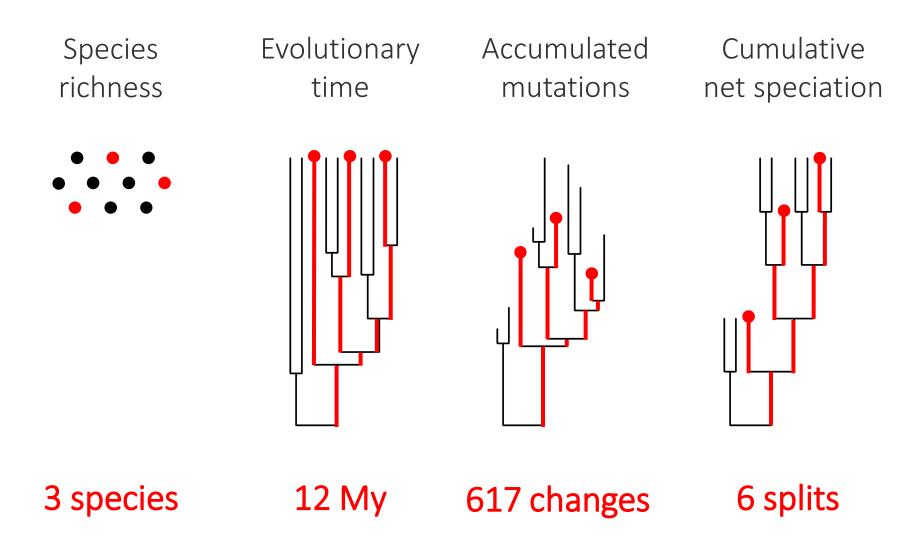




3 species

7 My





# onservation values

Why protect phylodiversity?

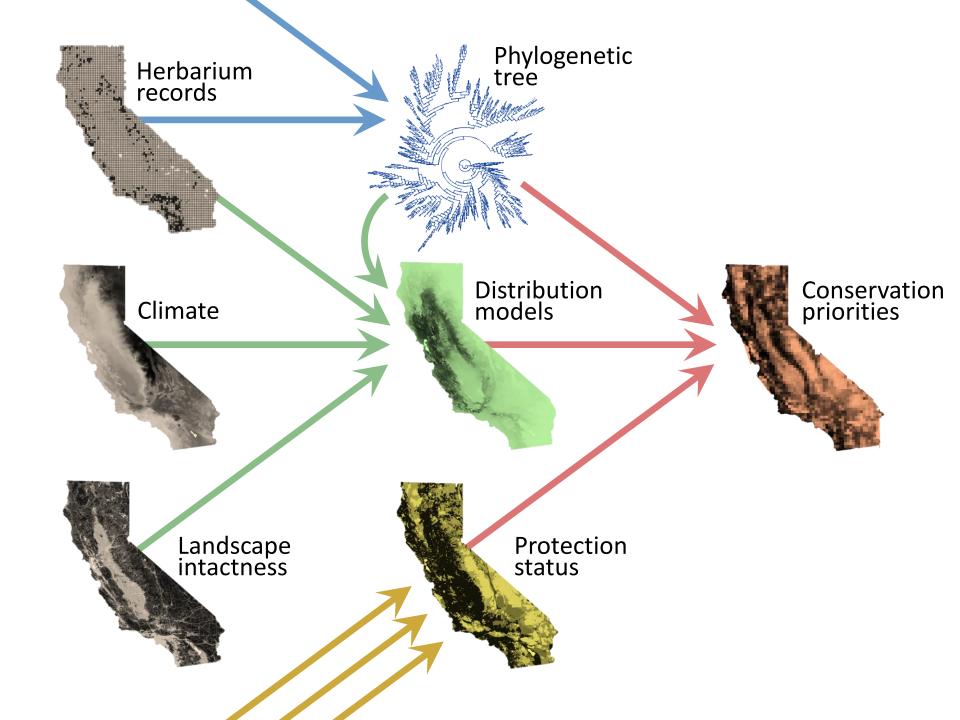
### **Phylodiversity facets**

ohylodiversity?	Mutations	Time	Speciation
Functional diversity			
Information content	<b>/</b>		
Evolutionary potential	<b>✓</b>		
Geographic properties			

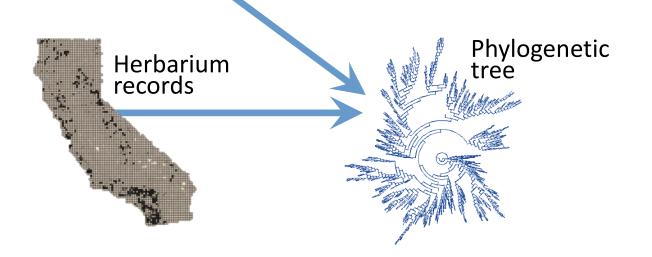
Why protect phylodiversity?

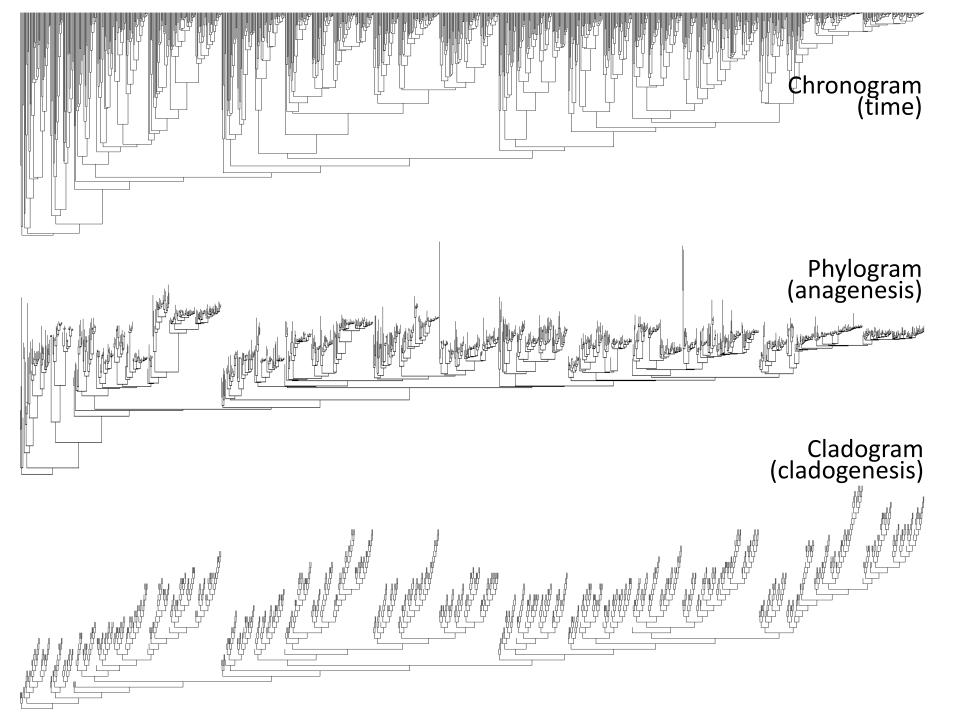
### **Phylodiversity facets**

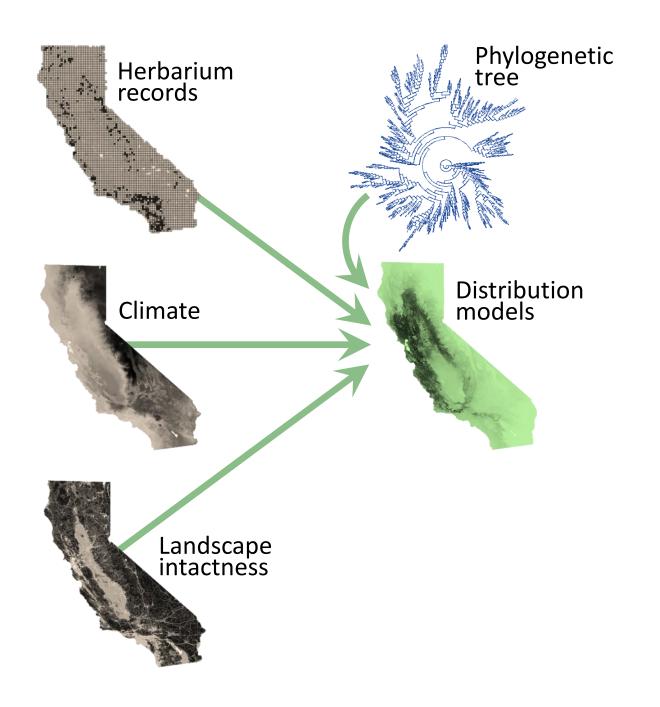
**QUESTION:** How do priority conservation sites differ by diversity facet?



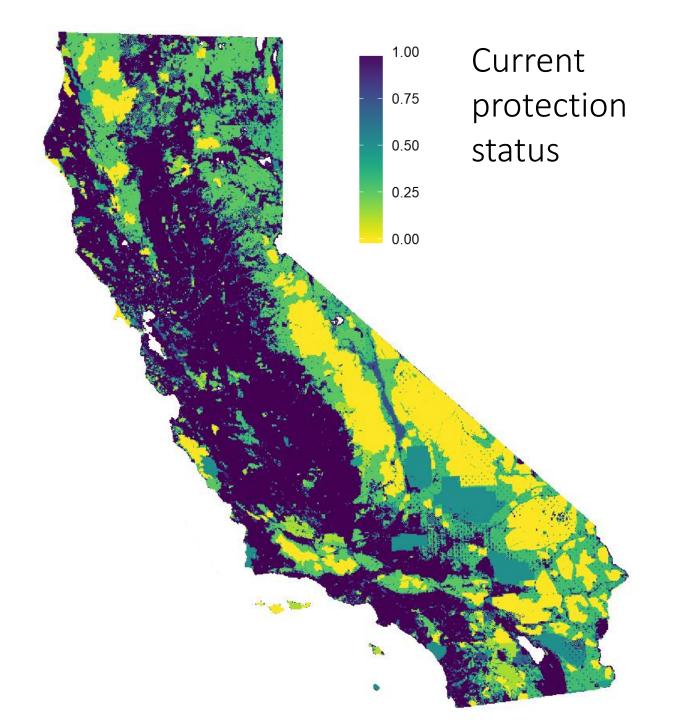


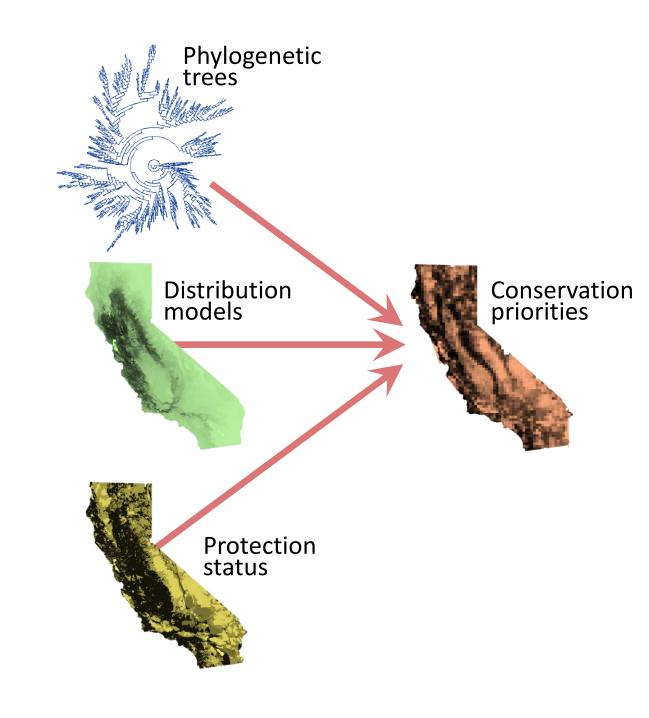






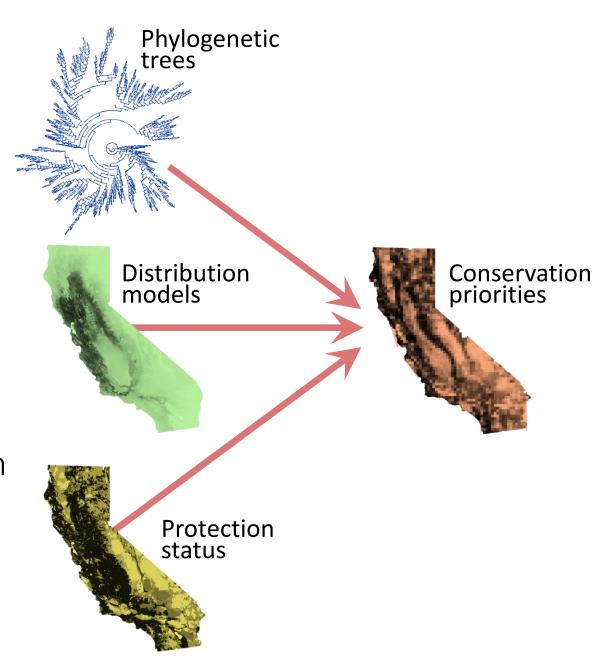


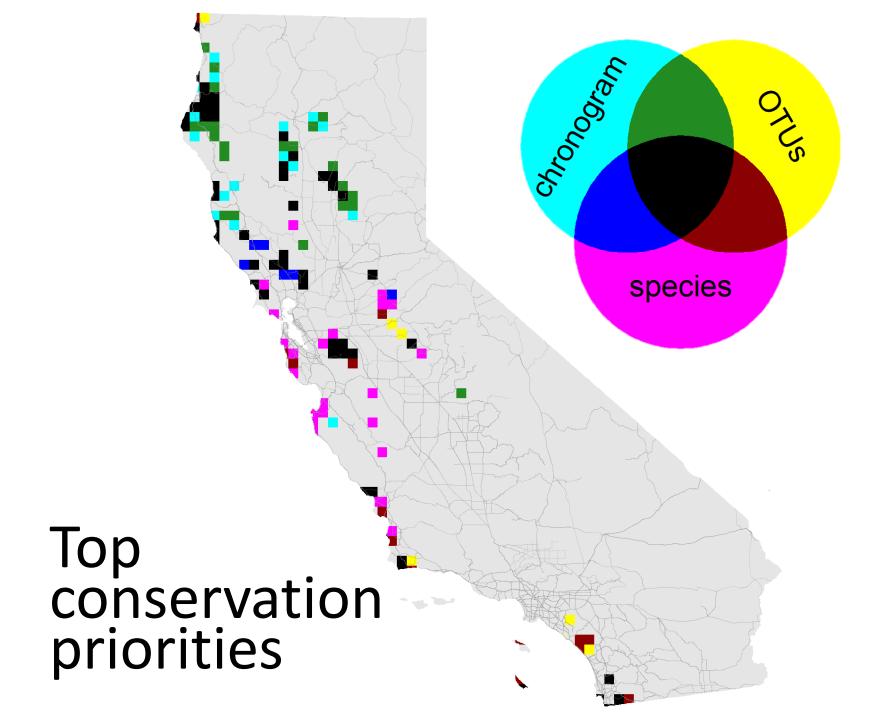


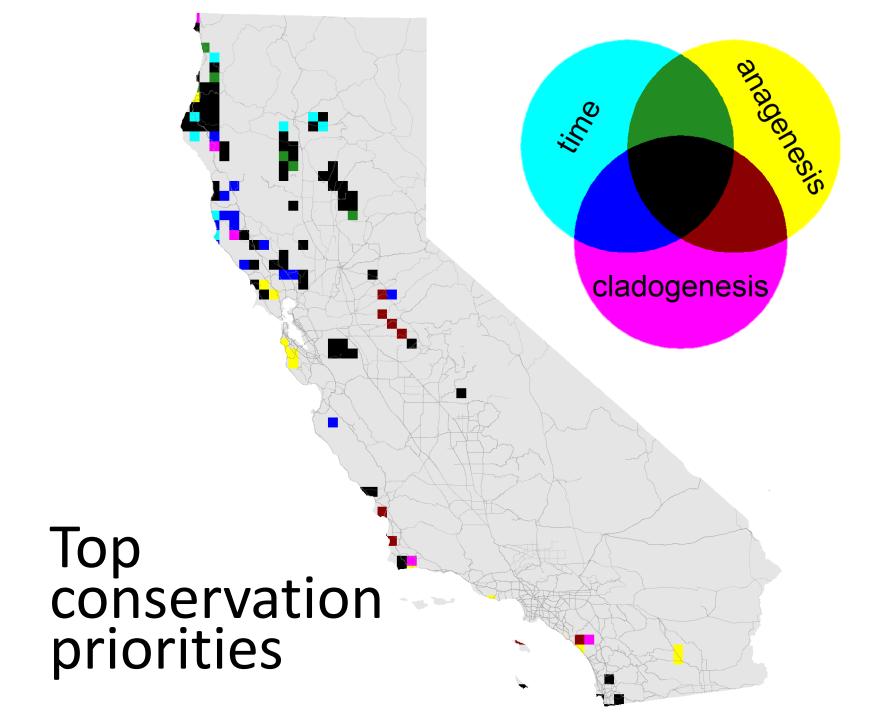


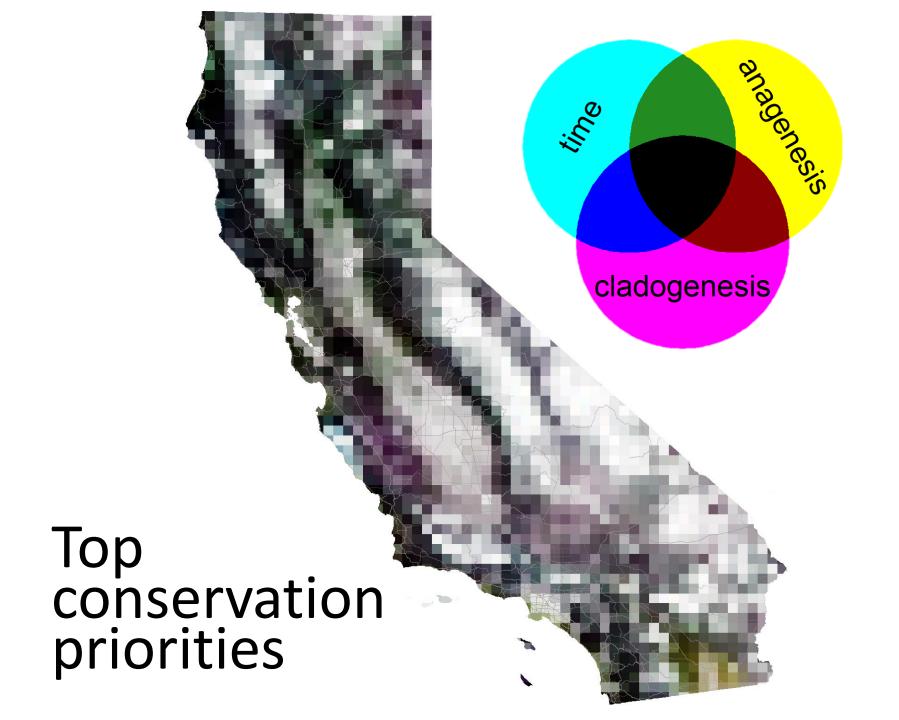
# Optimal conservation targets:

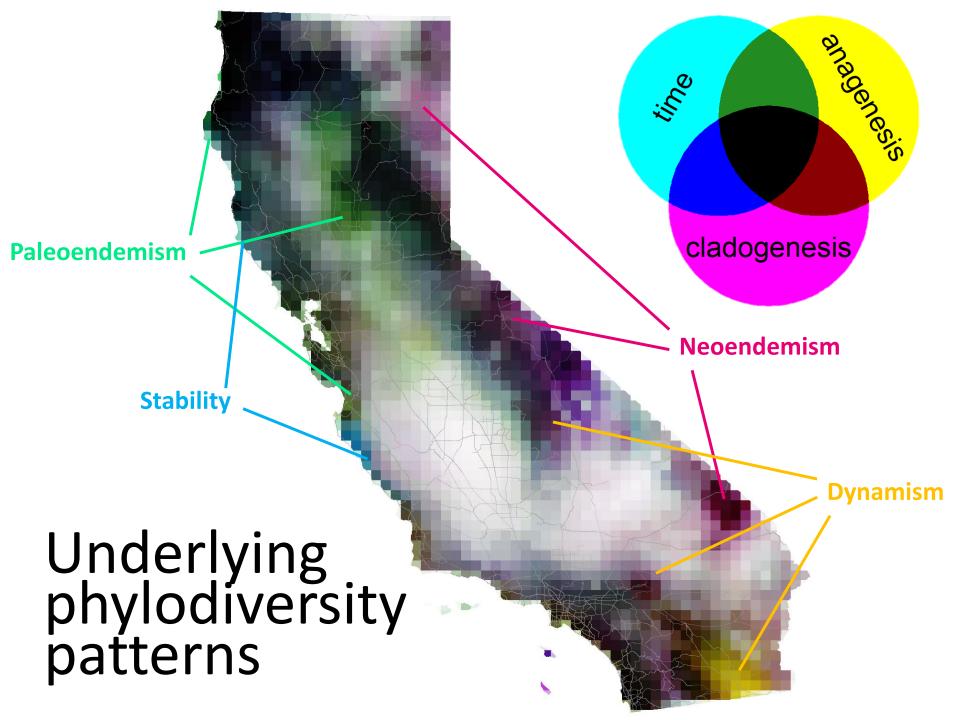
- Poorly protected
- Many resident taxa with:
  - Long branches
  - Small ranges
  - Poor protection across ranges

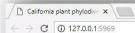










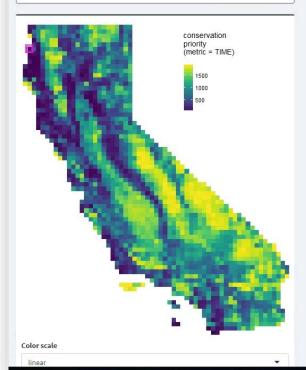




### California plant phylodiversity atlas

# Geography Map variable conservation priority Overall conservation priority for this diversity facet 1.0W values are HIGH priority sites.

Overall conservation priority for this diversity facet. LOW values are HIGH priority sites, which contain concentrations of long-branch taxa with small ranges that are largely unprotected by current preserves or other top-priority sites.



### Community

This table displays the floristic composition of the grid cell selected at left. The table is sortable and searchable.

Click to select a taxon. To map its range, change map variable to 'taxon range'. To view it in the evolutionary tree, change phylogeny variable to 'selected taxon'.

how	10 v entries		Search:		
	name #	combined benefit	presence 🖣	endemism 🛊	branch length
1	Lysichiton	1	0.3909	0.0466	0.469
2	Lycopodium	0.8587	0.3618	0.0206	0.759
3	Lycopodiella	0.549	0.4619	0.0189	0.759
4	Thuja	0.2799	0.3508	0.0121	0.544
5	Lycopodium & Lycopodiella (2- OTU clade)	0.2348	0.6565	0.0108	0.304
6	Armeria	0.1876	0.2829	0.0212	0.314
7	Empetrum	0.1646	0.2338	0.0962	0.090
8	Chrysosplenium	0.1324	0.3886	0.0338	0.100
9	Ribes_Heritiera	0.1051	0.3841	0.0218	0.12
10	Dryopteris_1	0.096	0.4875	0.0055	0.319
now	ing 1 to 10 of 2,164 er	ntries			
	Previous	1 2 3	3 4 5	217	Next

Phylogeny

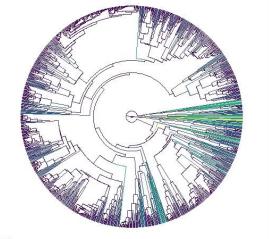
Select biodiversity facet

time

Time as a measure of biodiversity based on a 'chronogram' represents the total number of years that independent ancestral lineages presisted to give rise to taxon assemblage.

Lineages colored by

branch length



### Options

- ☐ Plot only the clade selected in table (not yet working)
- Show tip labels

### Color scale