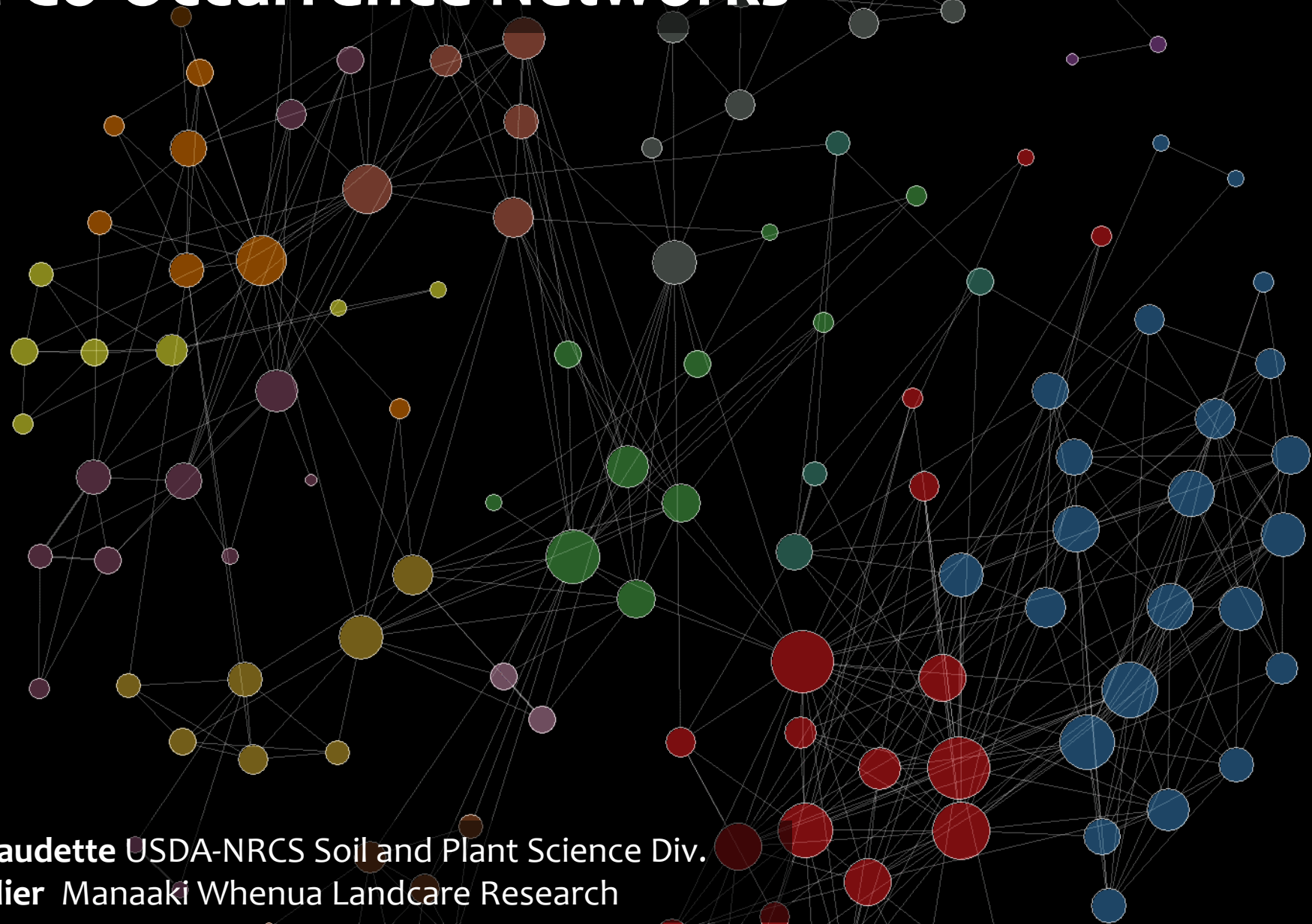




Mapping Soilscares Using Soil Co-Occurrence Networks



D.E. Beaudette USDA-NRCS Soil and Plant Science Div.
P. Roudier Manaaki Whenua Landcare Research

Acknowledgments

- Pierre Roudier and Dion O’Neale
- A.T. O’Geen, C. Stiles, Drew Kinney
- 100+ years of soil survey effort
- R graph libraries (`igraph`, `ape`)
- R soil science related libraries (`soilDB`, `sharpshootR`)

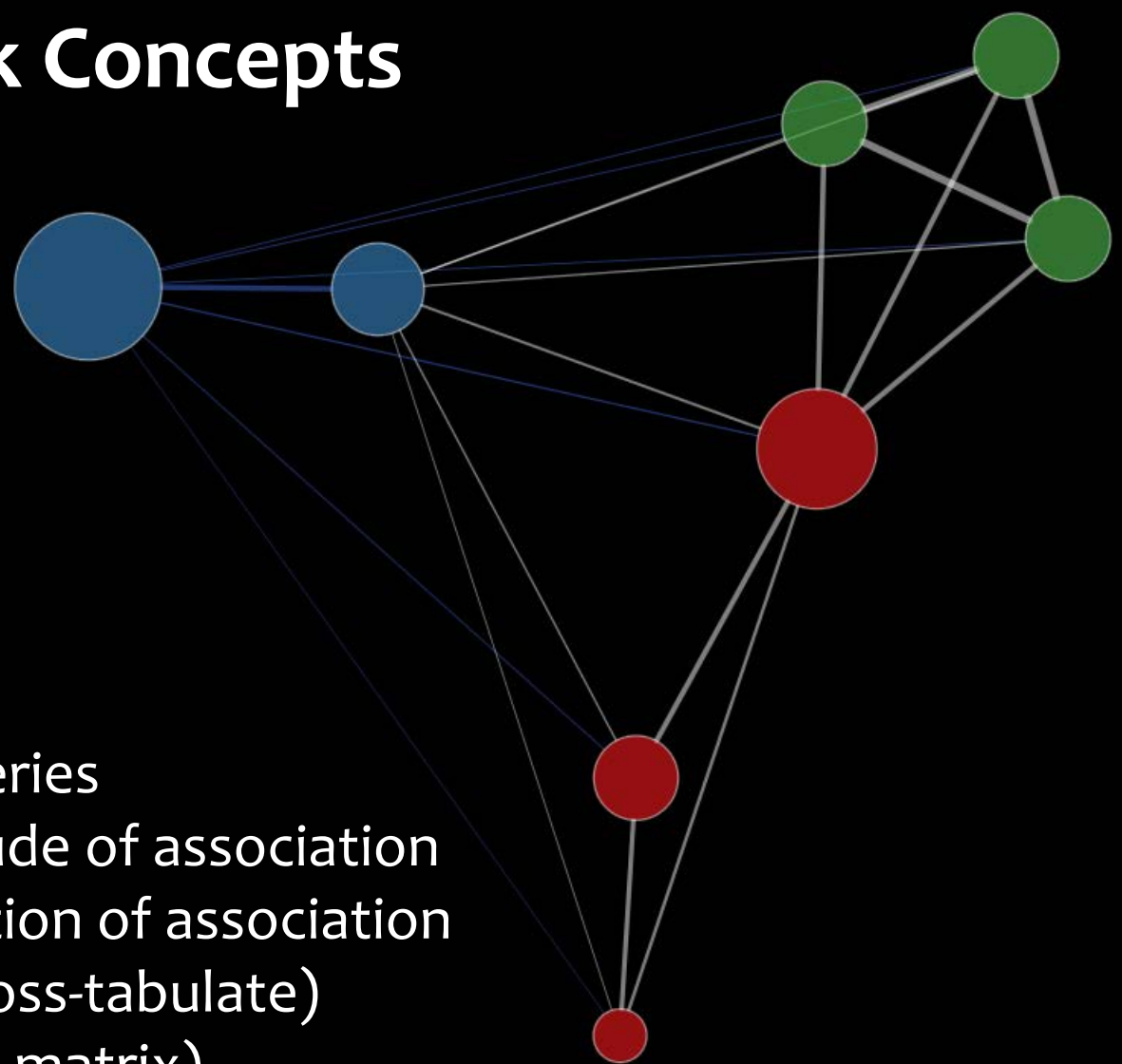
Prior Work

- 1977, 1985: F.D. Hole → soilsclapes defines patterns of association
- 2001: P. Lagacherie et al. → soilsclapes as landscape signature
- 2010: A.E. Hewitt et al. → soilsclapes as modeling domains
- 2013/2016: J.D. Phillips → spatial adjacency of mapping units
- 2013—current: SoilWeb Series Data Explorer
- 2014/2017: D.E. Beaudette and P. Roudier → mapping co-occurrence
- 2018/2019: Odgers, Roudier, Thompson, Beaudette → soilsclapes in NZ/US

This talk is about quantifying co-occurrence within tabular data

This talk is not about quantifying spatial co-occurrence

Graph / Network Concepts



- node or vertex: soil series
- edge weight: magnitude of association
- adjacency: quantification of association
 - co-occurrence (cross-tabulate)
 - distance (distance matrix)
 - similarity (restatement of distance)
- degree: number of edges / node
- community detection: clustering patterns of co-occurrence

Soil Data Explorer - SIERRA

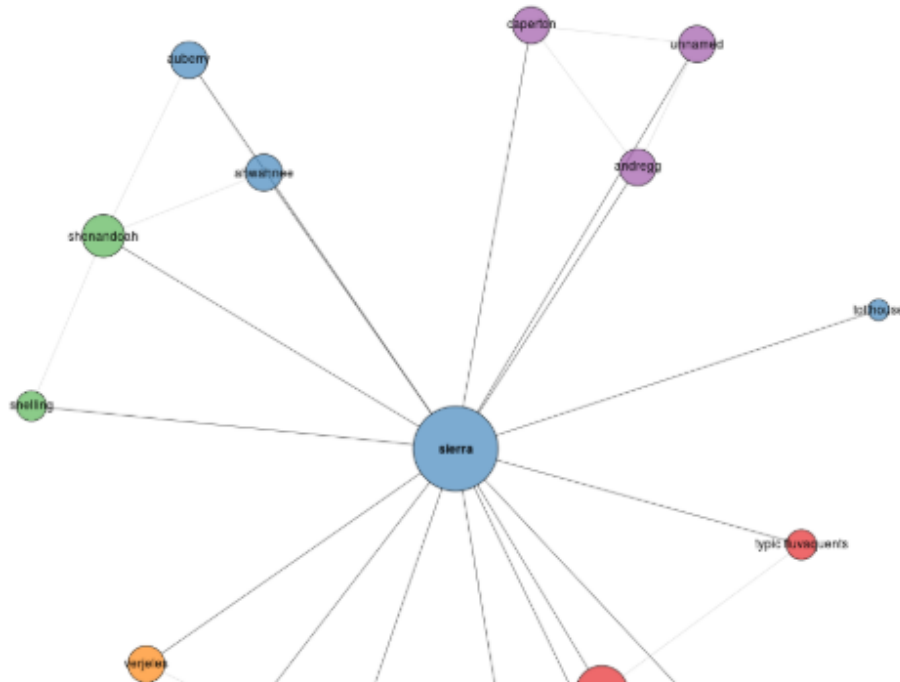
- OSD
- Lab Data
- Component Association**
- Series Association
- Block Diagrams
- Map Units
- Extent
- Competing Series

Component Association

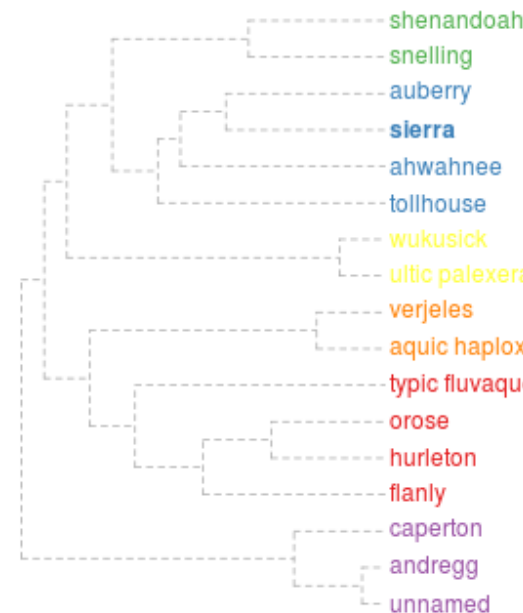
Two views depicting components that occur within map units containing the SIERRA series as a *major component*. Component association is a function of how frequently co-occurences occur. This summary *should* generate structures that are related to soil-landscape concepts that were used to construct map units containing the SIERRA series. These diagrams give only partial information on the relationships between components in the SSURGO snapshot.

Graph representation: each circle (vertex) is a component, lines (edges) connect components that have been mapped together, colors define groups of components that frequently co-occur, vertex proximity is proportional to co-occurrence as weighted by component percentage. Dark lines connect vertices to the queried soil series, grey lines connect other vertices.

Click the image to view it full size.



Dendrogram representation: branch height (relative to the left-side of the dendrogram) is proportional to co-occurrence as weighted by component percentage. Components are highlighted in this view.



Soilscapes → Mapping Units → Soil Series

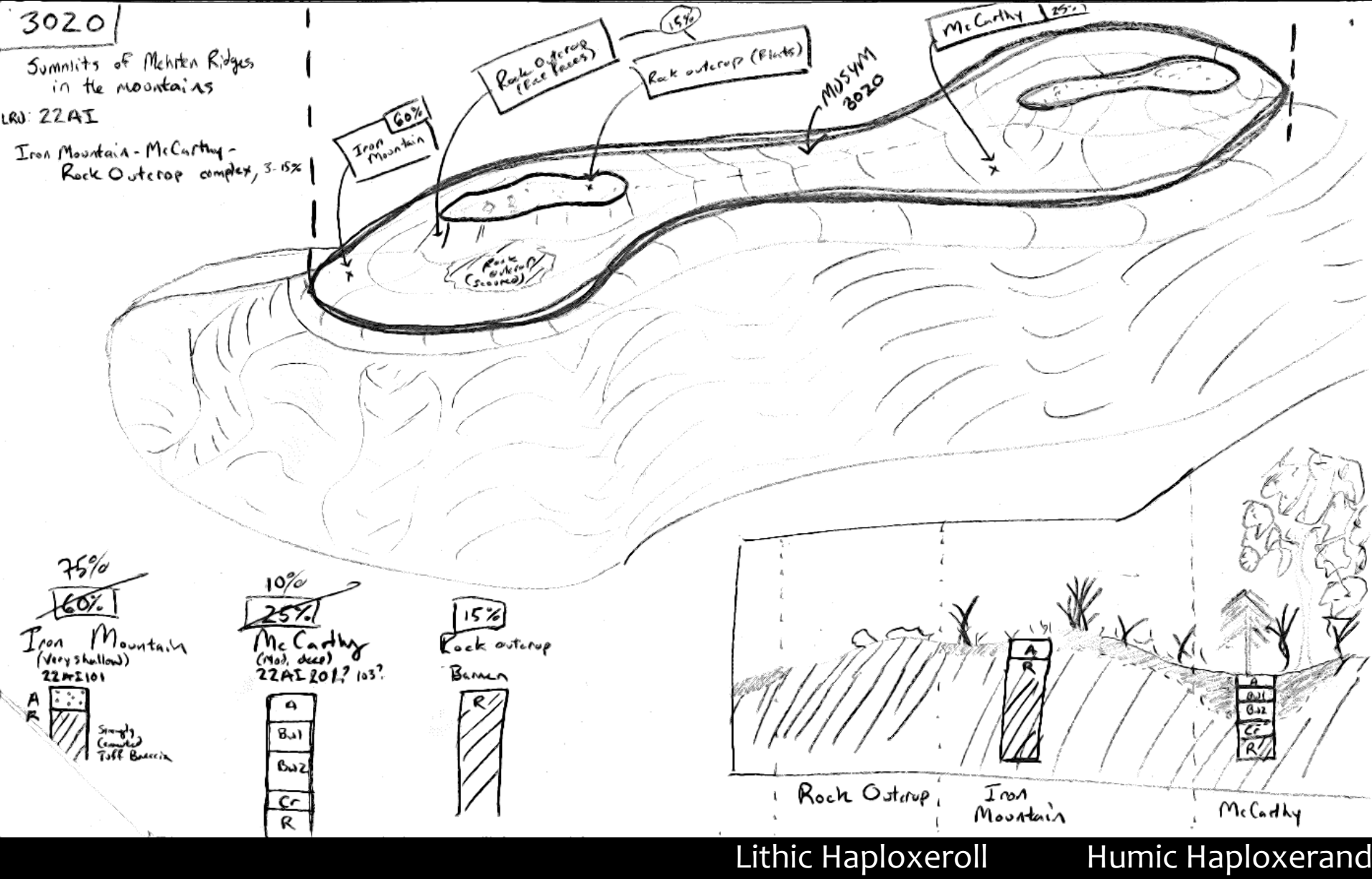
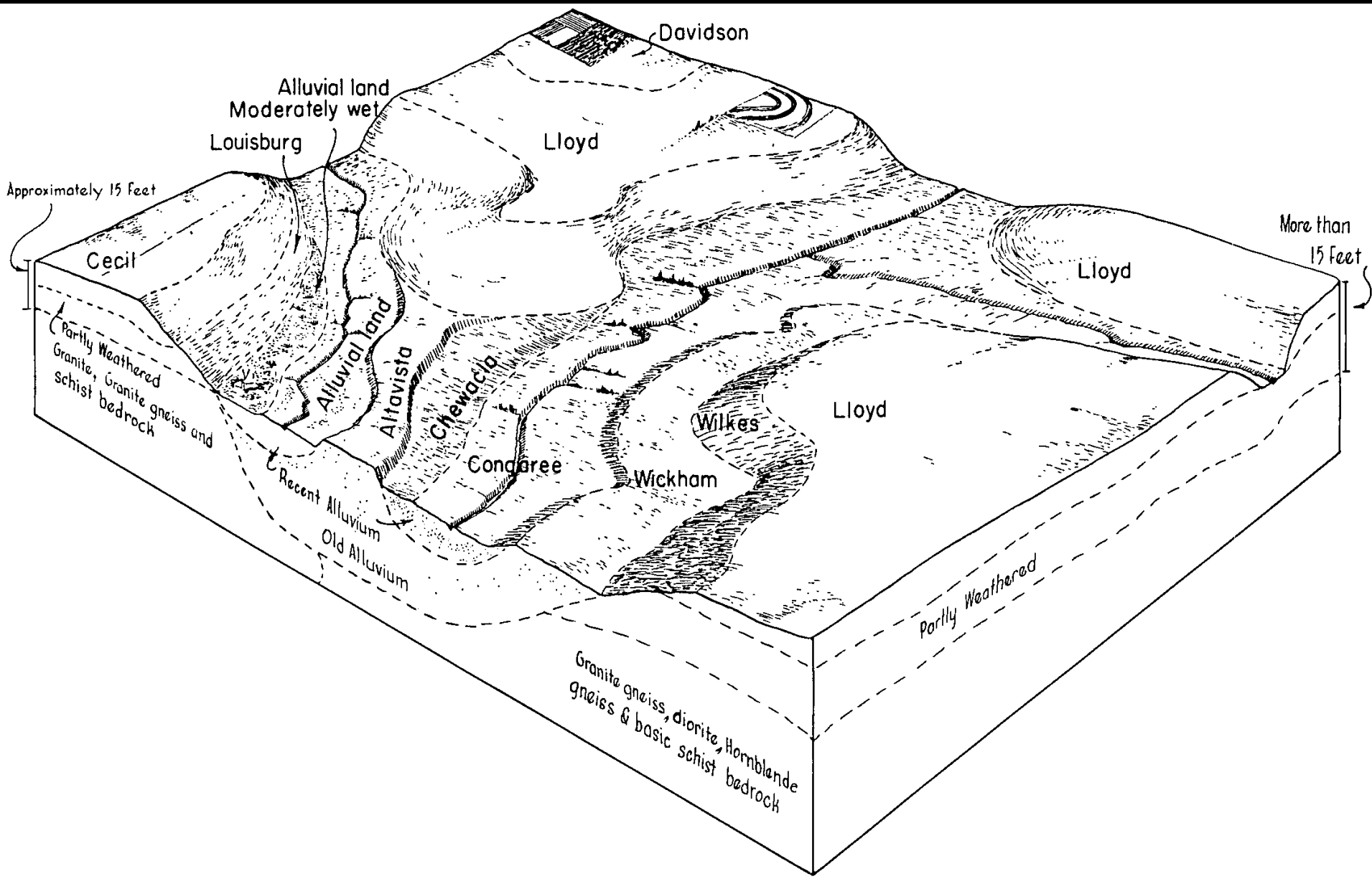


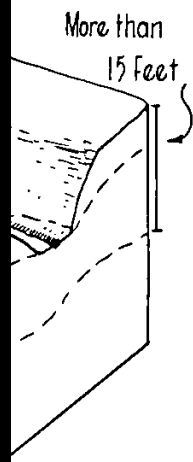
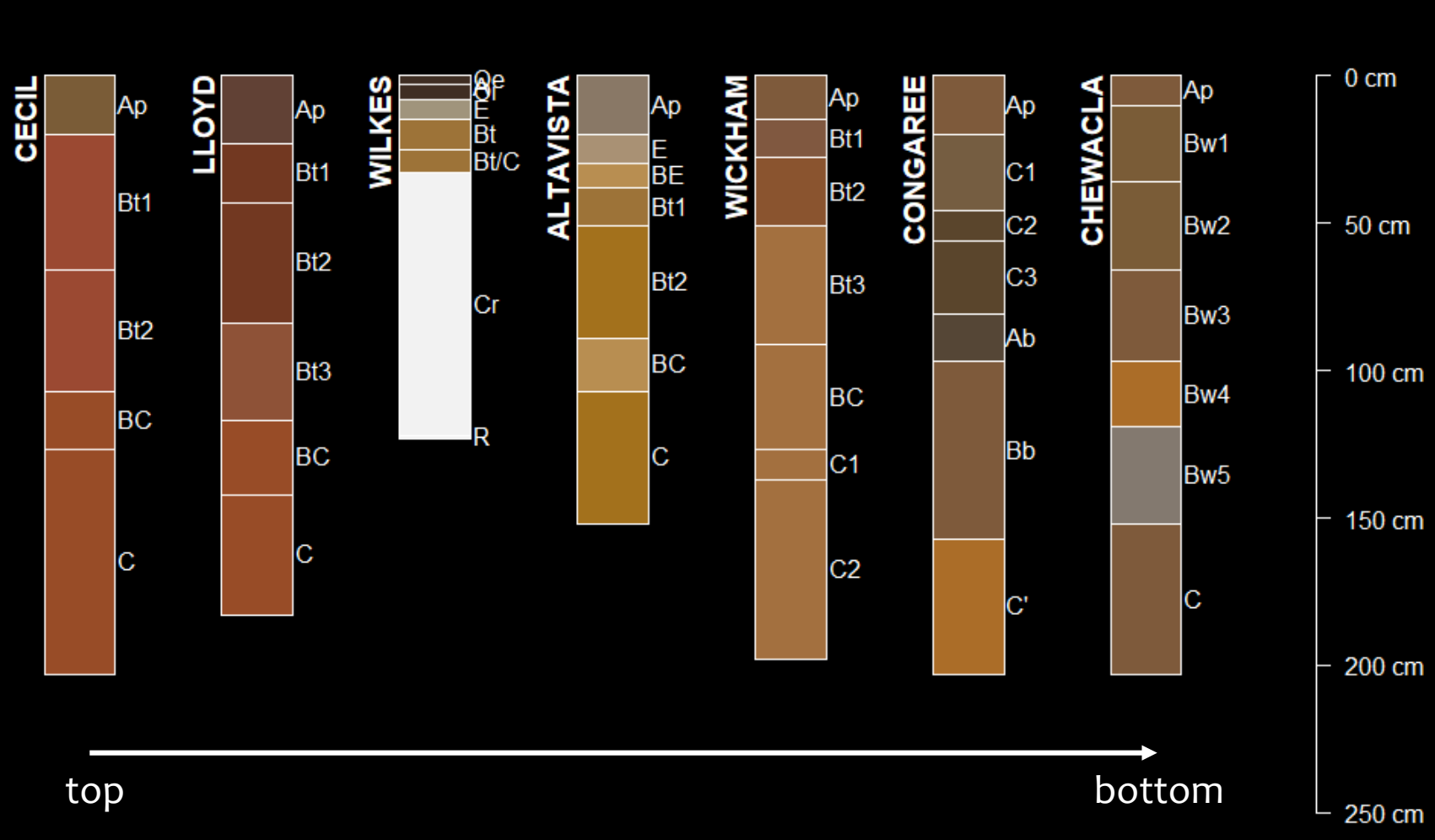
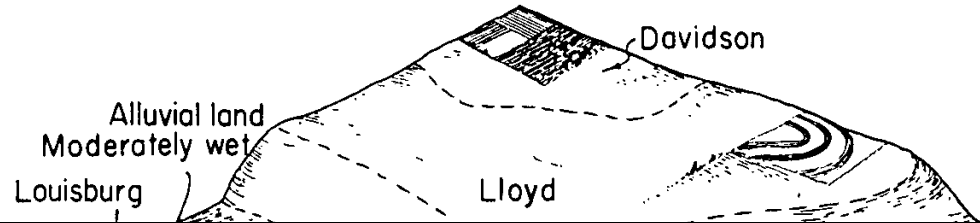
Figure by Andrew Brown, USDA-NRCS, Sonora, CA

Soilscapes → Mapping Units → Soil Series

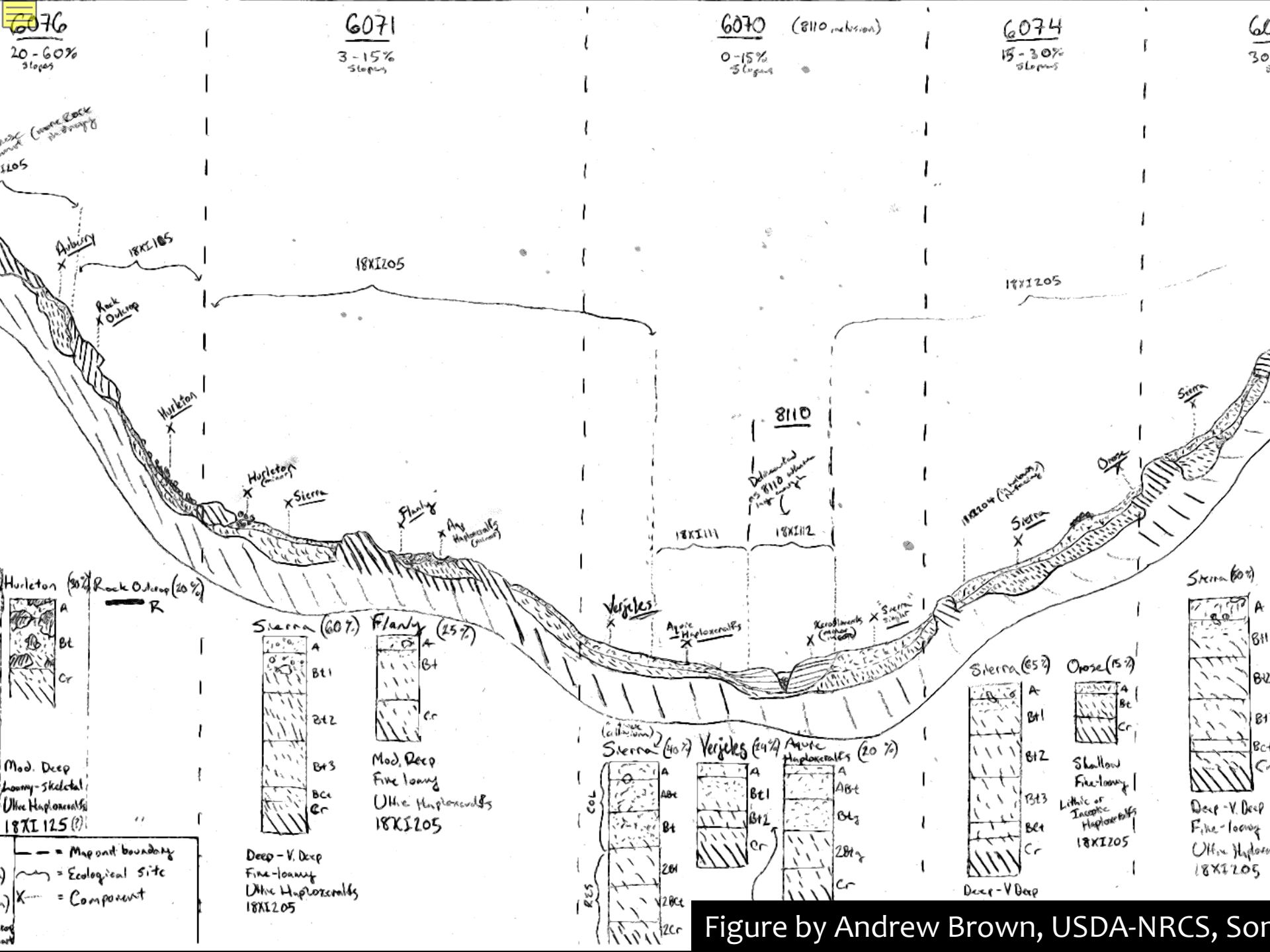


Lloyd-Davidson association (Soil Survey of Morgan County, Georgia; 1965)

Soilscapes → Mapping Units → Soil Series



Lloyd-Davidson association (Soil Survey of Morgan County, Georgia; 1965)



6076
20-60% Slopes

6071
3-15% Slopes

6070 (8110, reclassification)
0-15% Slopes

6074
15-30% Slopes

6075
30% Slopes

Rock Outcrop (same Rock as 6076)
18X1205

18X1105

18X1205

18X1205

8110

18X111

18X112

18X1204 (Alluvial)

Sierra (60%)

Hurleton (90%) Rock Outcrop (20%)
A
Bt
Cr

Sierra (60%) Flanky (25%)
A
Bt1
Bt2
Bt3
Bc1
Cr
Mod. Deep Fine-loamy Udic Haploxeralfs
18X1205

Sierra (40%) Verjules (40%) Anitic Haploxeralfs (20%)
A
ABc
Bt
2Bt
2Bt2
Cr
A
Bt1
Bt2
Cr
A
ABt
Bt3
2Bt2
Cr

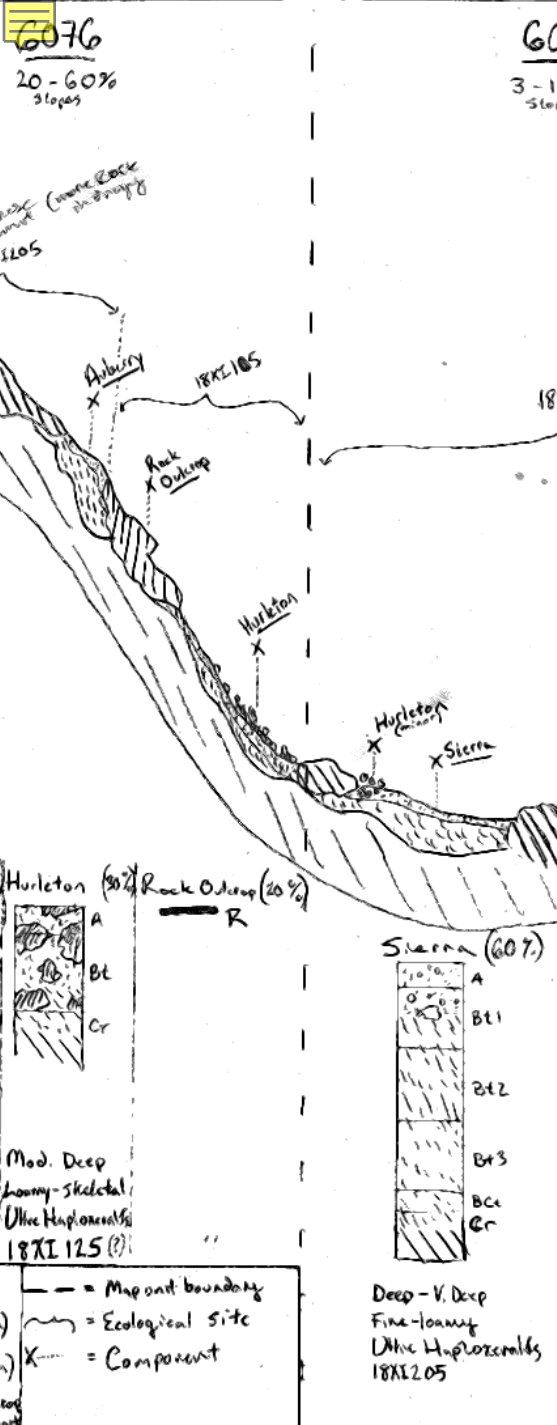
Sierra (65%) Chase (15%)
A
Bt1
Bt2
Bt3
Bc1
Cr
Shallow Fine-loamy Litic or Inceptic Haploxeralfs
18X1205
Deep-V Deep Fine-loamy Udic Haploxeralfs
18X1205

Sierra (60%)
A
Bt1
Bt2
Bt3
Bc1
Cr
Deep-V Deep Fine-loamy Udic Haploxeralfs
18X1205

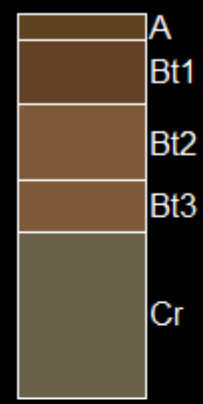
Mod. Deep Loamy-Skeletal Udic Haploxeralfs
18X1205 (?)

-- = Map unit boundary
[] = Ecological site
X = Component

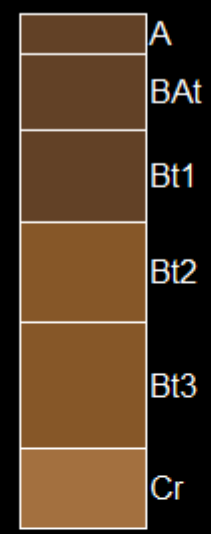
Figure by Andrew Brown, USDA-NRCS, Sor...



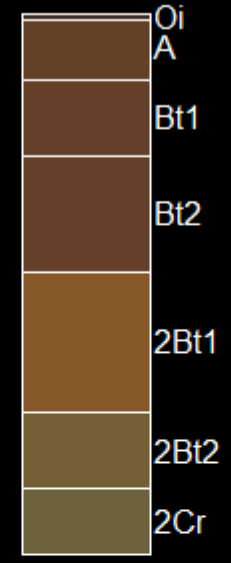
OROSE



FLANLY



VERJELES

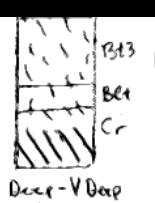
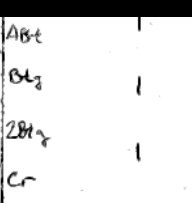
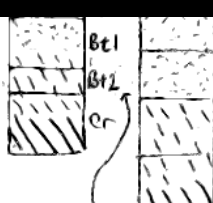
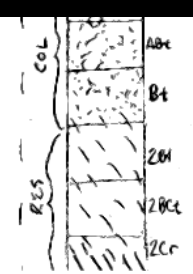


SIERRA



Ultic Haploxeralfs, complex arrangement

Fine loamy Ultic Haploxeralfs 18XI205



Fine-loamy Lithic or Inceptic Haploxeralfs 18XI205

Deep-V. Deep Fine-loamy Ultic Haploxeralfs 18XI205

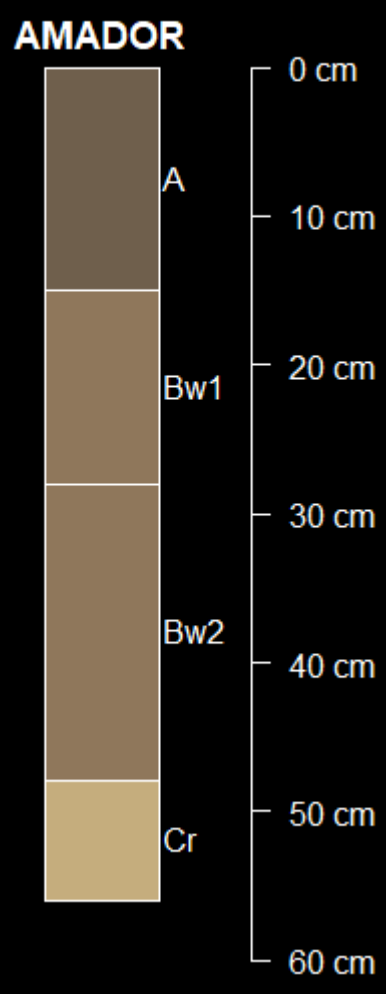
Figure by Andrew Brown, USDA-NRCS, Sor

Mapping Units in SSURGO

mukey	compname	comppct_r
1612048	Amador	45
1612048	Gillender	40
1612048	Ranchoseco	3
1612048	Vleck	3
1612048	Pardee	3
1612048	Peters	3
2766838	Amador	76
2766838	Gillender	9
2766838	Pardee	5
2766838	Miltonhills	5
2766838	Redding	2
	...	

complex

consociation



Amador: Loamy, mixed, superactive, thermic, shallow Typic Haploxerepts

Mapping Units in SSURGO



mukey	compname	compct_r
1612048	Amador	45
1612048	Gillender	40

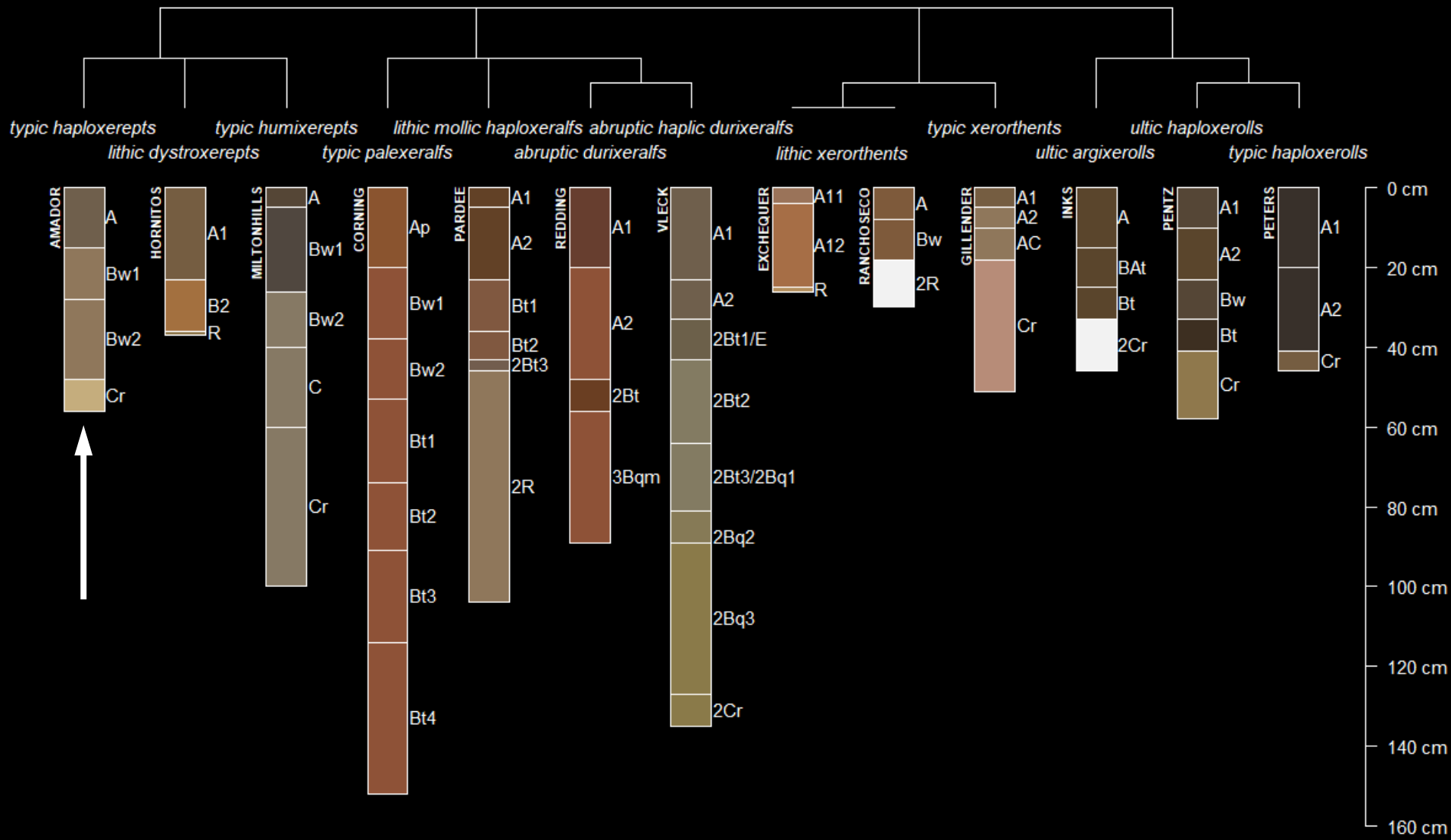
Mapping Units in SSURGO



mukey	compname	comppct_r
2766838	Amador	76
2766838	Gillender	9
2766838	Pardee	5
2766838	Miltonhills	5
2766838	Redding	2

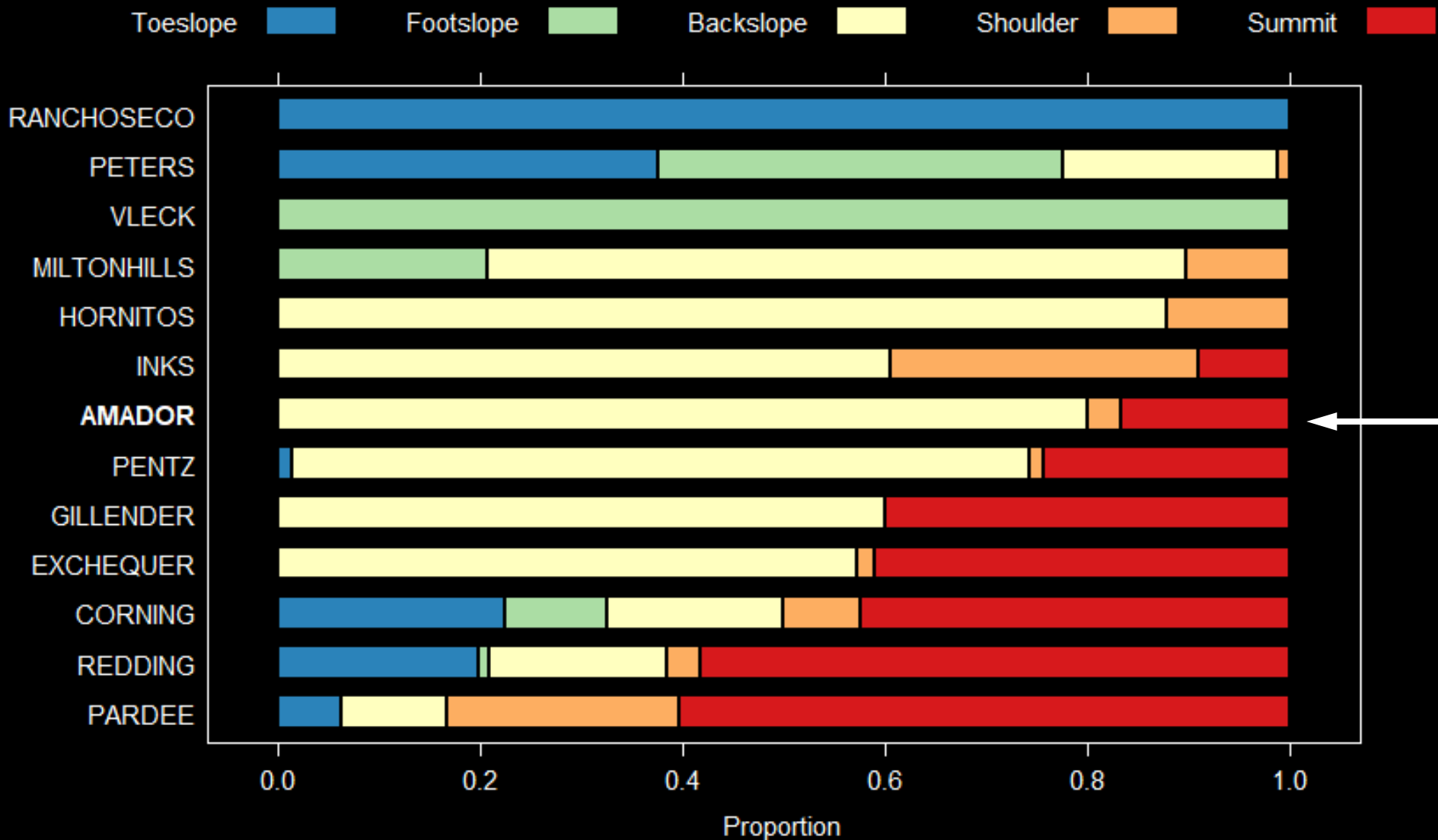
“Siblings”

Components* that co-occur with **Amador** in map units



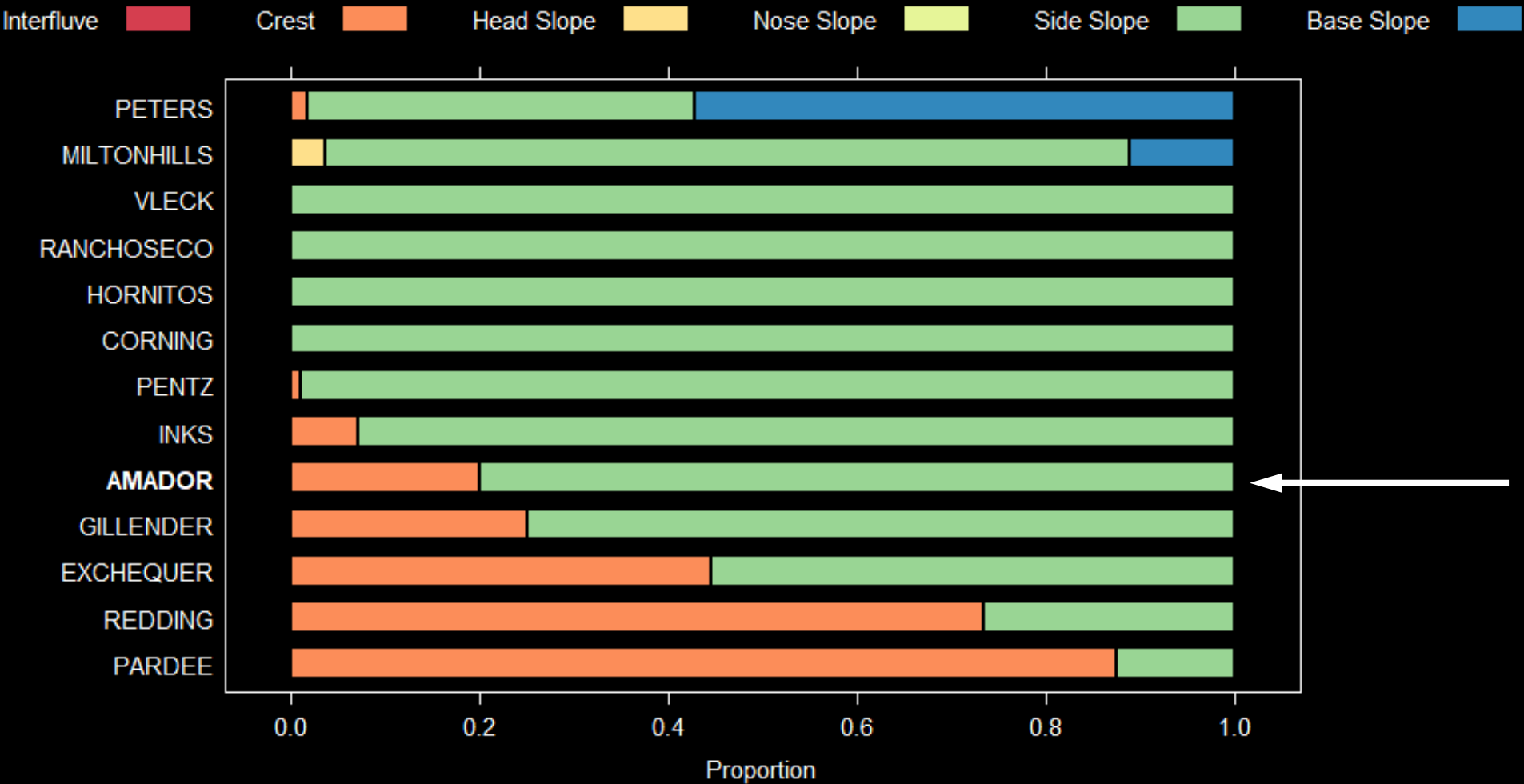
“Siblings”

Components* that co-occur with **Amador** in map units



“Siblings”

Components* that co-occur with **Amador** in map units



Quantify Co-Occurrence: Adjacency Matrix

Tabulate “co-occurrence” → *information is lost*

	Gillender	Miltonhills	Pardee	Peters	Ranchoseco	Redding	Vleck
Amador	2	1	2	1	1	1	1
Gillender		1	2	1	1	1	1
Miltonhills			1			1	
Pardee				1	1	1	1
Peters					1		1
Ranchoseco							1
Redding							
Vleck							

Compute “similarity”

1. collect mapunit / component records
2. reshape into "community matrix"
3. standardize and compute distance matrix (methods from numerical ecology)
4. convert distance matrix into similarity matrix

→ this is the adjacency matrix (details on next slide)

Why all the fuss? Component percentages (weights) matter!

Pseudo-community Matrix

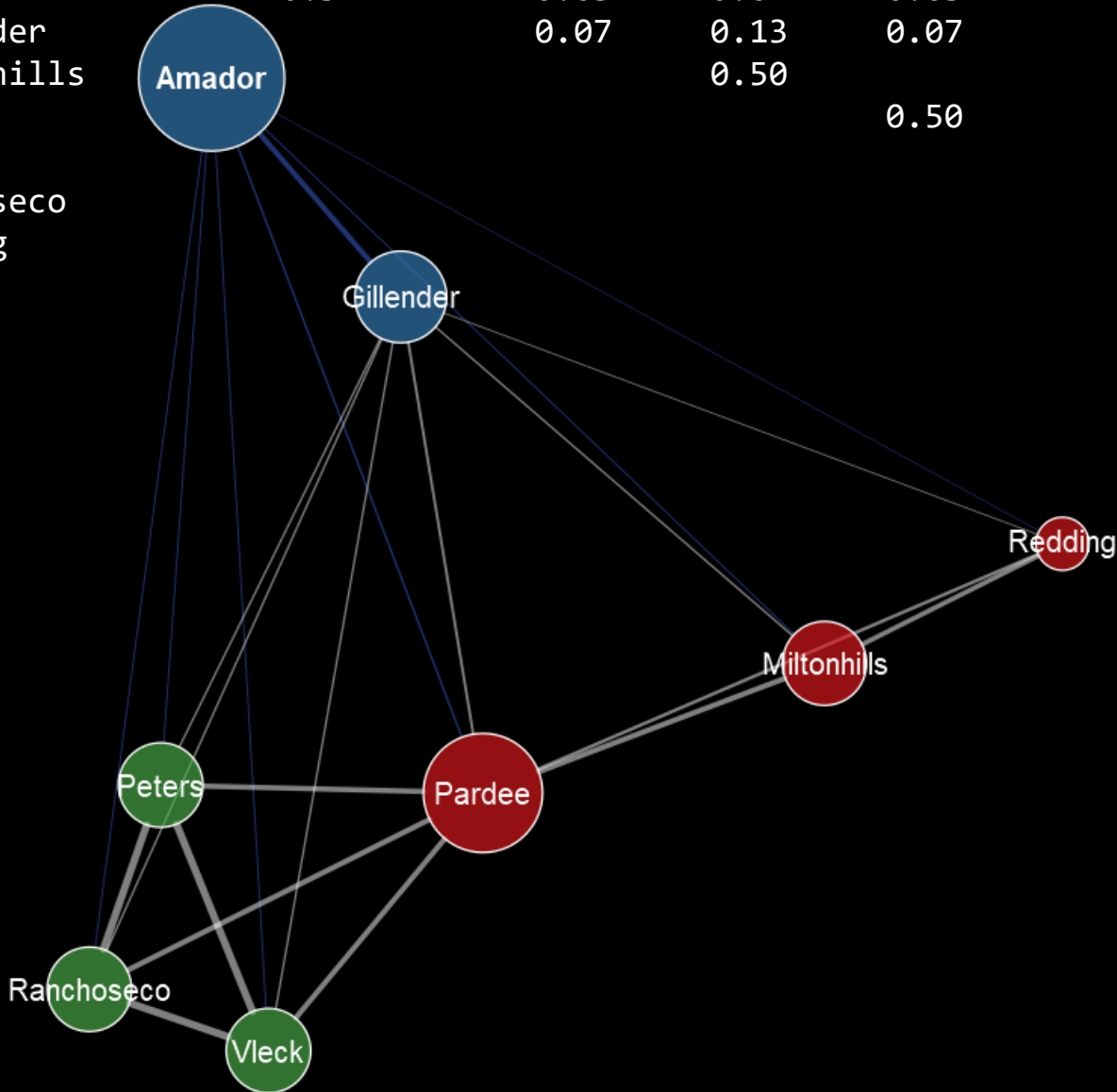
	461845	461980	462528	462529	462530	462531	462954	462955
Amador	45	25	85	85	85	85	85	85
Corning	0	5	0	0	0	0	0	0
Exchequer	0	0	5	5	5	5	0	0
Gillender	40	0	0	0	0	0	0	0
Hornitos	0	0	5	5	5	5	10	10
Inks	0	0	0	0	0	0	0	0
Miltonhills	0	0	0	0	0	0	0	0
Pardee	3	0	0	0	0	0	0	0
Pentz	0	0	5	5	5	5	5	5
Peters	3	0	0	0	0	0	0	0
Ranchoseco	3	0	0	0	0	0	0	0
Redding	0	4	0	0	0	0	0	0
Vleck	3	40	0	0	0	0	0	0

Similarity Matrix (via Jaccard index)

	Gillender	Miltonhills	Pardee	Peters	Ranchoseco	Redding	Vleck
Amador	0.5	0.03	0.07	0.03	0.03	0.01	0.03
Gillender		0.07	0.13	0.07	0.07	0.03	0.07
Miltonhills			0.50			0.40	
Pardee				0.50	0.50	0.20	0.50
Peters					1.00		1.00
Ranchoseco							1.00
Redding							
Vleck							

Adjacency Matrix → Graph

	Gillender	Miltonhills	Pardee	Peters	Ranchoseco	Redding	Vleck
Amador	0.5	0.03	0.07	0.03	0.03	0.01	0.03
Gillender		0.07	0.13	0.07	0.07	0.03	0.07
Miltonhills			0.50			0.40	
Pardee				0.50	0.50	0.20	0.50
Peters					1.00		1.00
Ranchoseco							1.00
Redding							
Vleck							



Why are we doing this?

<https://xkcd.com/1838/>

THIS IS YOUR MACHINE LEARNING SYSTEM?

YUP! YOU POUR THE DATA INTO THIS BIG PILE OF LINEAR ALGEBRA, THEN COLLECT THE ANSWERS ON THE OTHER SIDE.

WHAT IF THE ANSWERS ARE WRONG?

JUST STIR THE PILE UNTIL THEY START LOOKING RIGHT.



Why are we doing this?

Explicit, quantitative, human/machine readable encoding of historic (and future) soil knowledge.

Continuous predictions of soil properties / membership are (of course) important.

Failing to integrate hard-won knowledge (collected and synthesized in places where soils occur) into modern efforts would be a terrible tragedy.

Groups of co-occurring soils are useful strata that should be integrated into statistical models.

Co-occurrence networks are neat.

Thank You



Generate Soil Networks in R

<http://ncss-tech.github.io/AQP/>

<https://goo.gl/6HMWRR>

```
soilDB::siblings()
```

```
soilDB::fetchOSD()
```

```
sharpshootR::component.adj.matrix()
```

```
sharpshootR::polygonAgency()
```

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Help Test SoilWeb for Android Devices

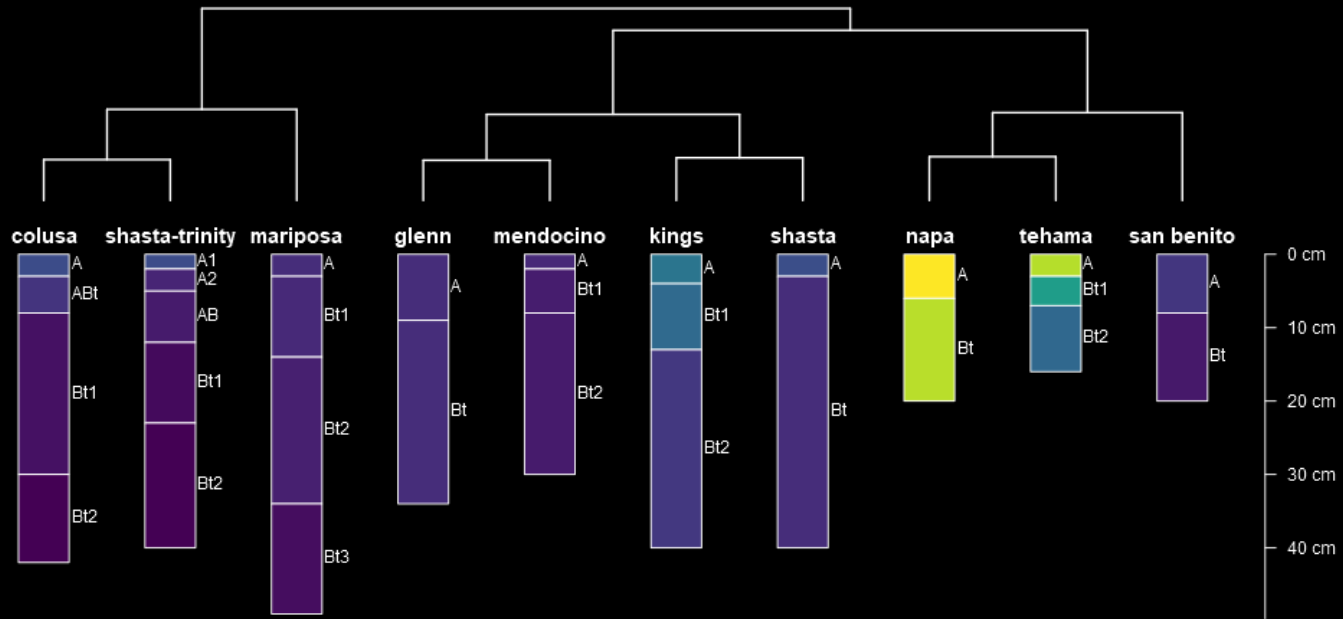


<https://goo.gl/WvrV8Y>



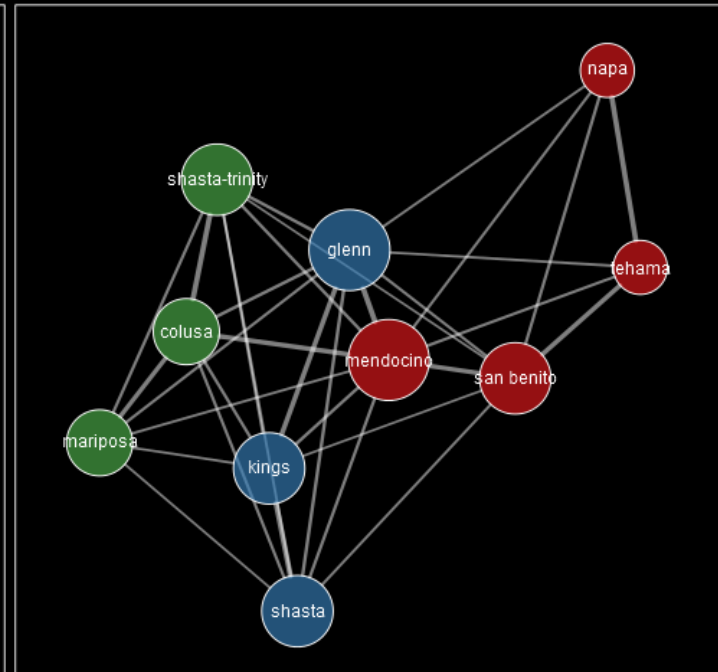
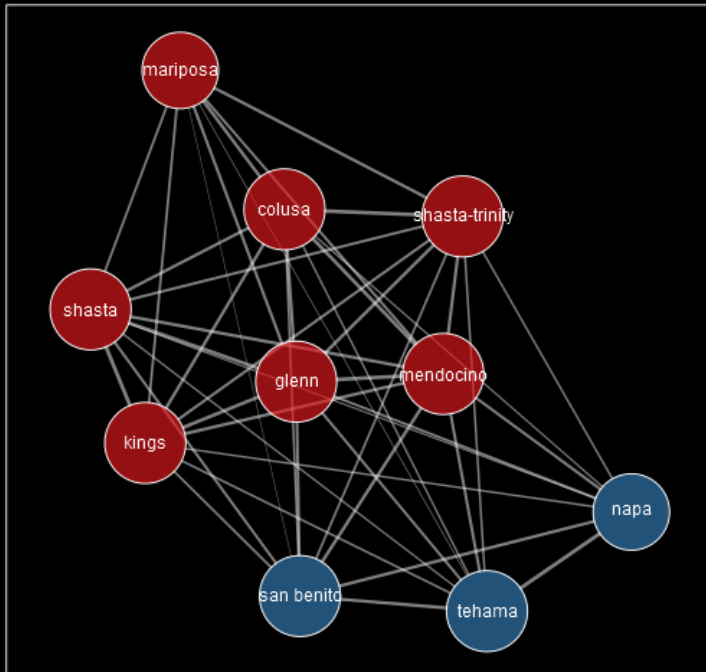
Exchangeable Ca to Mg Ratio

0 0.2 0.4 0.6 0.8 1 1.2 1.4 1.6



Full Distance Matrix

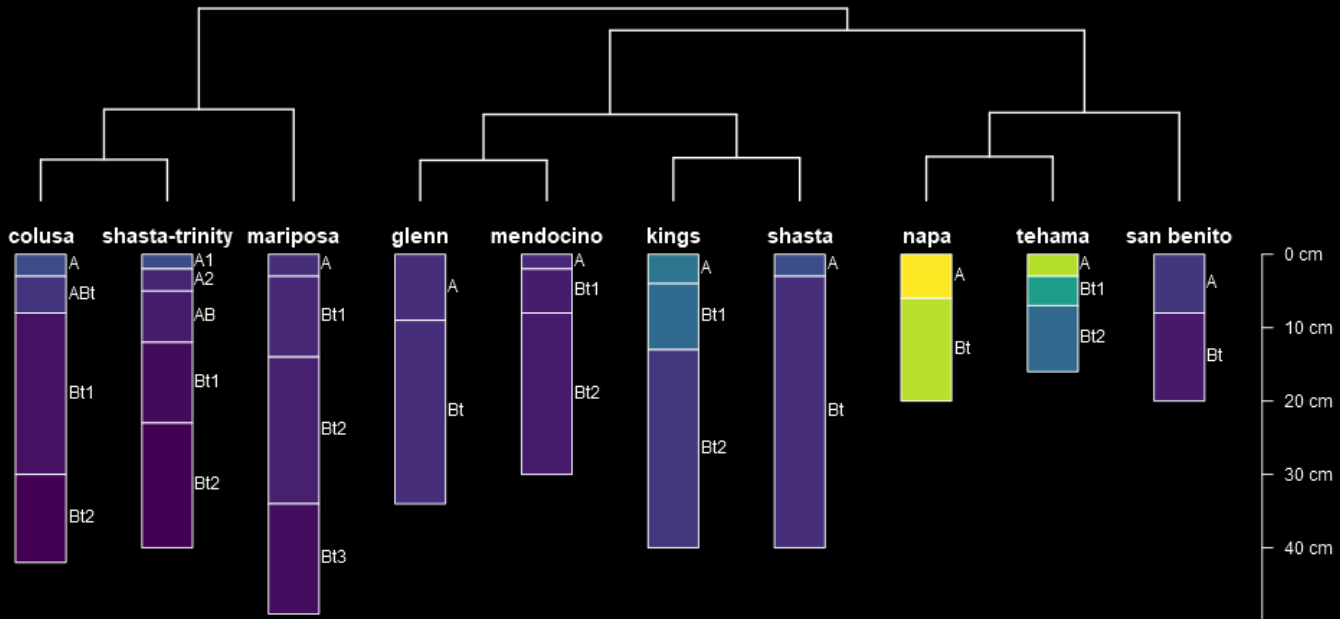
Max Span. Tree + Edges > Q25





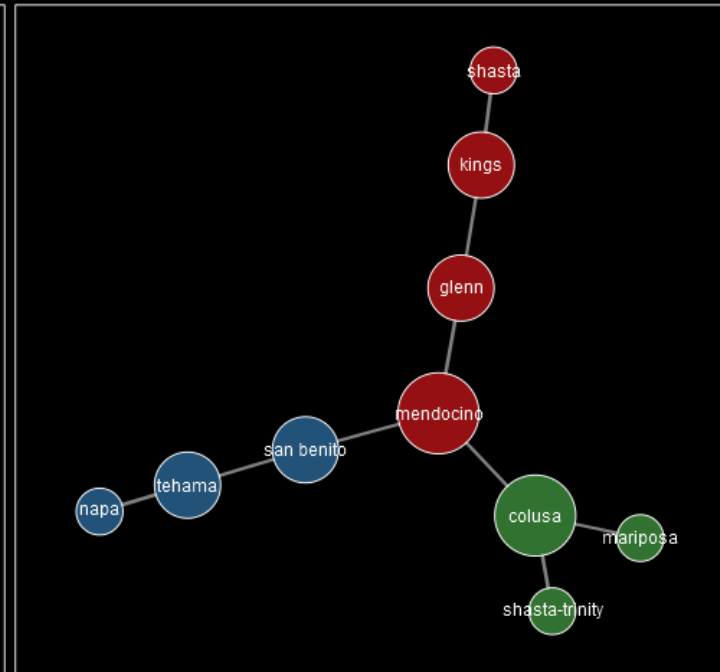
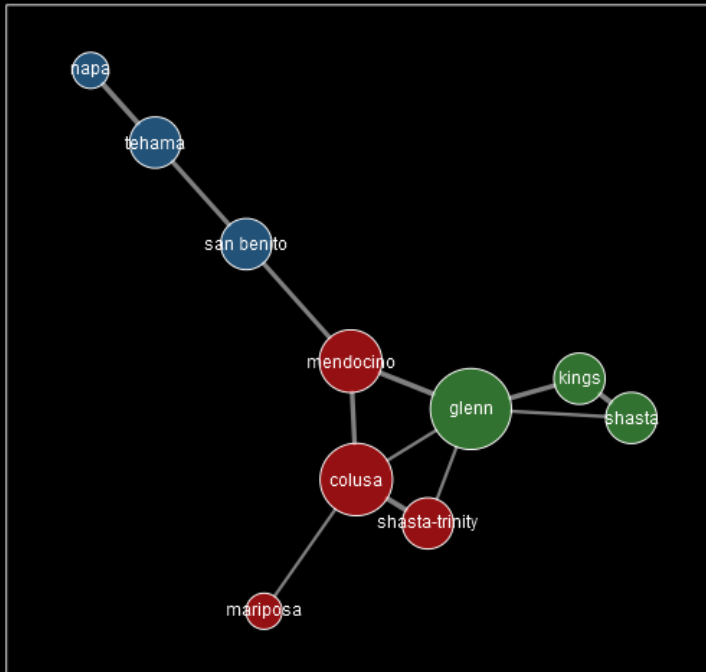
Exchangeable Ca to Mg Ratio

0 0.2 0.4 0.6 0.8 1 1.2 1.4 1.6



Max Span. Tree + Edges > Q75

Max Span. Tree



Adjacency Matrix → Graph

	Gillender	Miltonhills	Pardee	Peters	Ranchoseco	Redding	Vleck
Amador	2	1	2	1	1	1	1
Gillender		1	2	1	1	1	1
Miltonhills			1			1	
Pardee				1	1	1	1
Peters					1		1
Ranchoseco							1
Redding							
Vleck							

