# A Bayesian phylogenetic study of the Siouan language family using typological data 

Edwin Ko

University of California, Berkeley
41st Siouan and Caddoan Languages Conference, 2021

## Acknowledgments

I acknowledge with respect that the University of Berkeley,
California resides on the traditional, ancestral, and unceded land of the Ohlone people.

My sincerest thanks and gratitude to John Boyle, Andrew Garrett, David Kaufman, Rory Larson, Tyler Lemon, Sarah Lundquist, Julie Marsault, Armik Mirzayan, and Corey Roberts.

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## Introduction

- Reconstructing the linguistic history of a language family involves making inferences based on available information.
- Because we do not know what the true history is, there is a degree of uncertainty associated with our inferences.
- When there are many possible hypotheses, it is important to quantify these uncertainties to determine the most likely ones.
- Sound change: ${ }^{*} c h>k$ or ${ }^{*} k>c h$ ?
- Subgrouping:



## Computational phylogenetic methods

- Determining the internal subgrouping of any language family is a non-trivial and computationally-intensive task.
- There are approximately 17 Siouan languages which amount to $2.6 \times 10^{42}$ possible trees.
- Bayesian phylogenetic methods allow us to estimate the possible trees that have the highest likelihood given the available data.
- While these tools can be very useful, the results are only as good as the data and model assumptions that are employed.


## Data used in phylolinguistic research

- Most phylolinguistic studies use lexical data for classification of the following types (Chang et al. 2015):
- Cognate: Proto-Siouan *ahpá > Crow apá 'nose’, Biloxi pá 'head' (Rankin et al. 2015)
- Root-meaning: 'nose'
(1) Crow apé, Hidatsa abá (Boyle \& Gwin 2006:70)
(2) Mandan páaxu (Kasak 2019:201, Ex.3.35e), Lakota pȟasú (Ullrich 2019)
- Other studies have incorporated typological, structural data, although this has been controversial.
- Dunn et al. (2005) use computational phylogenetic methods with typological features to argue for a shared historical association between Austronesian and Papuan languages.


## Typological data: phylogeny or geography?

- The Austronesian-Papuan controversy in a nutshell:
- Typological features can detect a geneological signal!
- Dunn et al. 2005, 2007, 2008, Dunn 2009
- Wrong, typological features detect a geographical signal!
- Donohue \& Musgrave 2007, Donohue et al. 2008, 2011
- Sicoli and Holton (2014) also used typological features to infer the true of the Dene-Yeneseian macro-family.
- It is still unclear how reliable typological features are in inferring the true phylogenetic tree.
- Typological features are often thought of as being easily diffusable across geographical space (e.g. Holman et al. 2007).


## Purpose of this talk

## Research questions

(1) Can typological/structural data be used to detect a phylogenetic signal or does it indicate a geographical signal?
(2) How do the results compare with previously proposed classifications of the Siouan language family?

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## Typological data used in previous studies

- Dunn et al. 2008: 115 binary features, selected (i) "to provide broad coverage," (ii) "to distinguish between the languages of Island Melanesia," and (iii) "on which Austronesian and Papuan languages generally diverge" (Dunn et al. 2008:730)
- Sicoli \& Holton 2014: 116 binary features, Sherzer's (1976) An areal-typological study of American Indian languages north of Mexico
- Yanovich (2020) argues against Sicoli \& Holton: there is too little data and thus the inferences are not robust


## Lessons learned

- Incorporate more data and check for robustness of inferences


## Data used in this study

- World Atlas of Language Structures (WALS; Dryer \& Haspelmath 2013) has been used in phylogenetic studies:
- Distance-based: Wichmann \& Saunders 2007, Donohue et al. 2011, 2012, Greenhill et al. 2010
- Character-based: Wichmann \& Saunders 2007, Dediu 2011, Maurits \& Griffiths 2014
- Sixteen Siouan languages: Crow, Hidatsa, Mandan, Quapaw, Osage, Omaha, Ho-chunk, loway-Oto, Chiwere, Assiboine, Lakota, Dakota, Stoney, Biloxi, Ofo, and Tutelo
- WALS has many missing information and several inaccuracies:
- Hidatsa: Incorrectly coded as having noun-demonstrative order based on Matthews's 1965 Hidatsa Syntax
- Osage: Incorrectly coded as having nominal plural citing Quintero (1997:33) - there is no mention of plurality on p .33
- Likely a typo but p.330-339 show plural api following verbs


## Data used in this study

- Using most of the features from Sherzer and WALS, I coded from scratch employing more recent and reliable sources, and cross-checking with other Siouanists (thanks again!).
- 258 binary features (153 WALS, 105 Sherzer):
- 189 features are (potentially) parsimony-informative (i.e. have different values for at least two languages).
- Only Crow and Hidatsa lack nasal vowels.
- Only Quapaw lacks /u/ (Rankin 2005:463).
- 69 features known to have uniform values across all languages
- All Siouan languages have a mid or mid-high vowel.
- No Siouan languages employ plural particles on nominals.

| Feature type | Percent |
| :--- | :--- |
| Morphological | $\sim 50 \%$ |
| Phonological | $\sim 38 \%$ |
| (Morpho)syntactic | $\sim 8 \%$ |
| (Lexico)semantic | $\sim 4 \%$ |

## Coding the data

- Binary coding for presence ('1') or absence ('0') of features
- Missing data is coded as missing ('?').
- Features that have multiple values become separate features
- Negative Morphemes has four values (affix/clitic, particle, double, auxiliary word) is converted to four distinct features
- If a language has a negative affix, then it likely does not also have a negative auxiliary word.
- Issue of interdependent data:
- Unfortunately, this is common practice even with lexical cognacy data that use binary coding which violates an assumption of independence with Bayesian methods.
- It would be ideal to use mutli-state characters.


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## NeighborNet: How tree-like is the data?



Figure: Splits graph using NeighborNet (Brant \& Moulton 2004). Boxes and reticulations (i.e. web-like patterns) indicate conflicting signals.

## NeighborNet: How tree-like is the data?



| LANGUAGE | $\delta$-SCORE | Q-RESIDUAL |
| :--- | :--- | :--- |
| Crow | 0.227 | 0.0185 |
| Hidatsa | 0.234 | 0.0204 |
| $\Rightarrow$ Mandan | $\mathbf{0 . 2 7 4}$ | $\mathbf{0 . 0 2 3 2}$ |
| Lakota | 0.221 | 0.0165 |
| Dakota | 0.232 | 0.0154 |
| Assiniboine | 0.242 | 0.0193 |
| Stoney | 0.281 | 0.0219 |
| $\Rightarrow$ Chiwere | $\mathbf{0 . 2 7 3}$ | $\mathbf{0 . 0 2 3 2}$ |
| loway-Oto | 0.271 | 0.0193 |
| $\Rightarrow$ Hochunk | $\mathbf{0 . 3 2 9}$ | $\mathbf{0 . 0 2 2 6}$ |
| $\Rightarrow$ Osage | $\mathbf{0 . 3 0 6}$ | $\mathbf{0 . 0 3 3 4}$ |
| Omaha | 0.252 | 0.0194 |
| Quapaw | 0.257 | 0.0185 |
| Tutelo | 0.248 | 0.0228 |
| $\Rightarrow$ Biloxi | $\mathbf{0 . 2 7 3}$ | $\mathbf{0 . 0 2 4 8}$ |
| $\Rightarrow$ Ofo | $\mathbf{0 . 3 0 4}$ | $\mathbf{0 . 0 3 8 7}$ |
| Average | 0.264 | 0.0224 |

- $\delta$-scores and Q-residuals: 0 (less conflict) to 1 (more conflict), where conflict represents more sharing of traits with other languages.


## Comparison of $\delta$-scores and Q-residuals

- How do the $\delta$-score and Q-residual compare with those reported for other language groups?

| LANGUAGE GROUP | $\delta$-SCORE | Q-RESIDUAL | Data TYPE | Source |
| :--- | :--- | :--- | :--- | :--- |
| Siouan | 0.264 | 0.024 | Typological/Structural | - |
| Dene-Yeneseian | 0.367 | 0.0492 | Typological/Structural | Sicoli \& Holton 2014 |
| Austronesian | 0.44 | 0.0354 | Typological/Structural | Greenhill et al. 2017 |
| Indo-European | 0.23 | 0.003 | Lexical | Gray et al. 2010 |
| Polynesian | 0.41 | 0.020 | Lexical | Gray et al. 2010 |
| Ainu | 0.25 | 0.01 | Lexical | Lee \& Hasegawa 2013 |
| (Mainland) Japanese | 0.39 | 0.002 | Lexical | Lee \& Hasegawa 2014 |
| Ryukyuan | 0.23 | 0.004 | Lexical | Lee \& Hasegawa 2014 |
| Chapacuran | 0.262 | 0.016 | Lexical | Birchall et al. 2016 |
| Austronesian | 0.38 | 0.0062 | Lexical | Greenhill et al. 2017 |
| Dravidian | 0.30 | 0.0069 | Lexical | Kolipakam et al. 2018 |
| Tai | 0.2808 | 0.04088 | Lexical | Dockum 2018 |
| Turkic | 0.34 | 0.001 | Lexical | Savelyev \& Robbeets 2020 |

## Key takeaway

The splits graph, $\delta$-score, and Q-residual for the Siouan data is well within the range of what is considered tree-like.

## Estimating the true tree

- The main goal is to obtain a sample of trees (not just one tree) that explains the data relatively well.
- To do this, the algorithm (Monte Carlo Markov Chain) searches the space of all possible trees step-by-step locating the trees that best fit the data.
- I ran the analysis in BEAST 2.6.3 (Bouckaert et al. 2019) using 10 million steps (generations) with a 1,000 sampling frequency and $25 \%$ burn-in resulting in a total of 7,500 trees.
- This process was repeated two additional times to checked to ensure the results are similar across the three independent runs.


## Assessing performance: Effective sample size (ESS)

- We want to have a good sample of trees, but how can we tell if the sample is sufficient?
- Using Tracer (Rambaut et al. 2018), ESS values over 625 are considered to indicate sampling independence (Fabreti \& Hoehna 2021).

| StATISTIC | ESS |
| :--- | :--- |
| Posterior | 5114 |
| Likelihood | 3403 |
| Prior | 2265 |
| treeLikelihood.wals-sherzer | 3403 |
| TreeHeight.t:tree | 3832 |
| gammashape.s:wals-sherzer | 709 |

## Assessing performance: "Fuzzy caterpillars"

- We also want to know if the model converged; that is, did it end in a state of equilibrium?
- We can also look for "fuzzy caterpillars" in the trace.


Figure: Woolly Bear Caterpillar (from Herald Times Reporter)

## Assessing performance: "Fuzzy caterpillars"

- We also want to know if the model converged; that is, did it end in a state of equilibrium?
- We can also look for "fuzzy caterpillars" in the trace.


Figure: Non-fuzzy caterpillar trace (Source: Taming the Beast)


Figure: Fuzzy caterpillar trace of posterior probability

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## Summary tree: Maximum clade credibility tree



Figure: Maximum clade credibility tree. Values indicate relative frequency of the sampled trees that contain the particular branching.

## Visualizing all trees: DensiTree



Figure: Maximum clade credibility


Figure: DensiTree (Bouckaert 2010, Bouckaert \& Heled 2014)

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## Comparison with Rankin's (2010) proposed tree



Figure: Comparison between current analysis (left) and the tree proposed by Rankin (2010; right). Dashed lines indicate sites of divergence.

## Evidence of a geographical signal?



Figure: Map of selected Siouan languages (adapted from Wikimedia Commons). Disclaimer: This map is a rough approximation of the language communities and their geographical locations.

## A qualitative assessment: Non-contiguous languages



Figure: Map of selected Siouan languages (adapted from Wikimedia Commons). Disclaimer: This map is a rough approximation of the language communities and their geographical locations.

## A qualitative assessment: Contiguous languages



Figure: Map of selected Siouan languages (adapted from Wikimedia Commons). Disclaimer: This map is a rough approximation of the language communities and their geographical locations.

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## Future directions

- Identifying the features responsible for the subgroupings
- Phonological features:
- Higher-level subgroupings
- Mandan $\Rightarrow$ Crow, Hidatsa
- Morphological features:
- Lower-level subgroupings
- Mandan $\Rightarrow$ Mississippi Valley
- Quality-checking the data again (and again)
- If any linguists would be willing to take a look at (a subset of) the data, I would greatly appreciate it!
- Incorporating Catawba and Yuchi typological data for inferring deeper historical relations
- Catawba and Yuchi are grouped with the Siouan languages suggesting perhaps that deeper time depths increases the potential for conflicting signals.


## Future directions

- Comparing analyses with lexical data (e.g. Kasak, n.d.)
- Data in the Comparative Siouan Dictionary need to be checked thoroughly
- Checking the results with other linguistic (e.g. shared innovations) and historical evidence
- Thoughts on the possibility of grouping Missouri River (Crow and Hidatsa) with Ohio Valley (Biloxi, Tutelo, and Ofo)?
- Some potentially shared innovations:
(1) Loss of glottalized consonants
(2) Collapse of the distinction arrive here/there
(3) Emergence of distinct nominal and verbal conjunctions
- Impressionistically, the Missoui River and Ohio Valley Siouan subgroups appear quite distinct.


## Ahóo!

## Thank you for listening!

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## Seeing the forests for the trees



Figure: Maximum clade credibility


Figure: DensiTree (Bouckaert 2010, Bouckaert \& Heled 2014)

## Model comparison

- Eighteen models with different settings were considered and ctmc-bd-relax-gam fits the data best.


## - Marginal likelihood was estimated using the Nested Sampling algorithm (Maturana et al. 2019) using 12 particles.

| Analysis | SUBSTITUTION | Tree | Clock | Marginal | Bayes |
| :---: | :---: | :---: | :---: | :---: | :---: |
|  | MODEL | Prior | MODEL | LOG-LIKELIHOOD | FACTOR |
| $\Rightarrow$ ctmc-bd-relax-gam | CTMC $+\gamma$ | Birth-death | Relaxed | -1247.3 | - |
| ctmc-yule-relax-gam | CTMC $+\gamma$ | Yule (pure birth) | Relaxed | -1247.9 | 1.2 |
| ctmc-bdsky-relax-gam | CTMC $+\gamma$ | Birth-death skyline | Relaxed | -1249.4 | 4.2 |
| cov-yule-strict | Covarion | Yule (pure birth) | Strict | -1251.1 | 7.6 |
| ctmc-yule-strict-gam | CTMC $+\gamma$ | Yule (pure birth) | Strict | -1251.2 | 7.8 |
| ctmc-bdsky-strict-gam | CTMC $+\gamma$ | Birth-death skyline | Strict | -1251.8 | 9.0 |
| ctmc-bd-strict-gam | CTMC $+\gamma$ | Birth-death | Strict | -1253.7 | 12.8 |
| cov-yule-relax | Covarion | Yule (pure birth) | Relaxed | -1256.1 | 17.6 |
| cov-bd-relax | Covarion | Birth-death | Relaxed | -1256.1 | 17.6 |
| cov-bd-strict | Covarion | Birth-death | Strict | -1256.7 | 18.8 |
| cov-bdsky-strict | Covarion | Birth-death skyline | Strict | -1258.5 | 22.4 |
| cov-bdsky-relax | Covarion | Birth-death skyline | Relaxed | -1258.7 | 22.8 |
| ctmc-yule-strict | CTMC | Yule (pure birth) | Strict | -1284.5 | 74.4 |
| ctmc-yule-relax | CTMC | Yule (pure birth) | Relaxed | -1285.5 | 76.4 |
| ctmc-bd-relax | CTMC | Birth-death | Relaxed | -1286.0 | 77.4 |
| ctmc-bdsky-relaxed | CTMC | Birth-death skyline | Relaxed | -1288.1 | 81.6 |
| ctmc-bdsky-strict | CTMC | Birth-death skyline | Strict | -1290.9 | 87.2 |
| ctmc-bd-strict | CTMC | Birth-death | Strict | -1293.3 | 92.0 |

Note: Interpreting BF: 1-2: weak, 2-6: positive, $6-10$ : strong, $>10$ : very strong.

## Model comparison

- Four other randomly-selected models cov-yule-relax, ctmc-yule-relax-gam, ctmc-bdsky-strict-gam, and cov-bdsky-relax produced similar tree topologies suggesting that the analysis is robust to the choice of tree priors (see Yanovich 2020).

Key takeaways

- There is some evidence for ctmc-bd-relax-gam to explain the data better than other models.
- The dataset is sufficient enough in size to make robust inferences about most likely trees given the data.


## WALS features (1/3) - non-binary

1. Consonant Inventories
2. Vowel Quality Inventories
3. Consonant-Vowel Ratio
4. Voicing in Plosives and Fricatives
5. Voicing and Gaps in Plosive Systems
6. Uvular Consonants
7. Glottalized Consonants
8. Lateral Consonants
9. The Velar Nasal
10. Vowel Nasalization
11. Front Rounded Vowels
12. Syllable Structure
13. Tone
14. Absence of Common Consonants
15. Presence of Uncommon Consonants
16. Exponence of Selected Inflectional Formatives
17. Locus of Marking in the Clause
18. Locus of Marking in Possessive Noun Phrases
19. Locus of Marking: Whole-language Typology
20. Zero Marking of A and P Arguments
21. Prefixing vs. Suffixing in Inflectional Morphology
22. Case Syncretism
23. Syncretism in Verbal Person/Number Marking
24. Number of Genders
25. Sex-based and Non-sex-based Gender Systems
26. Systems of Gender Assignment
27. Coding of Nominal Plurality
28. Occurrence of Nominal Plurality
29. Plurality in Independent Personal Pronouns
30. Associative Plural
31. Definite Articles
32. Definite Affix
33. Indefinite Articles
34. Indefinite Affix
35. Inclusive/Exclusive Distinction in Independent Pronouns
36. Inclusive/Exclusive Distinction in Verbal Inflection
37. Distance Contrasts in Demonstratives
38. Pronominal and Adnominal Demonstratives

## WALS features (2/3) - non-binary

39. Third Person Pronouns and Demonstratives
40. Gender Distinctions in Independent Personal Pronouns
41. Politeness Distinctions in Pronouns
42. Indefinite Pronouns
43. Intensifiers and Reflexive Pronouns
44. Person Marking on Adpositions
45. Number of Cases
46. Position of Case Affixes
47. Comitatives and Instrumentals
48. Ordinal Numerals
49. Numeral Classifiers
50. Conjunctions and Universal Quantifiers
51. Position of Pronominal Possessive Affixes
52. Possessive Classification
53. Adjectives without Nouns
54. Noun Phrase Conjunction
55. Nominal and Verbal Conjunction
56. Perfective/Imperfective Aspect
57. The Past Tense
58. The Future Tense
59. The Perfect
60. Position of Tense-Aspect Affixes
61. The Morphological Imperative
62. The Prohibitive
63. Imperative-Hortative Systems
64. Semantic Distinctions of Evidentiality
65. Coding of Evidentiality
66. Verbal Number and Suppletion
67. Order of Subject, Object and Verb
68. Order of Adposition and Noun Phrase
69. Order of Genitive and Noun
70. Order of Adjective and Noun
71. Order of Demonstrative and Noun
72. Order of Numeral and Noun
73. Order of Relative Clause and Noun
74. Order of Degree Word and Adjective

## WALS features (3/3) - non-binary

75. Position of Polar Question Particles
76. Position of Interrogative Phrases in Content Questions
77. Relationship between the Order of Object and Verb and the Order of Adposition and Noun Phrase
78. Relationship between the Order of Object and Verb and the Order of Relative Clause and Noun
79. Relationship between the Order of Object and Verb and the Order of Adjective and Noun
80. Alignment of Case Marking of Full Noun Phrases
81. Alignment of Case Marking of Pronouns
82. Alignment of Verbal Person Marking
83. Expression of Pronominal Subjects
84. Verbal Person Marking
85. Third Person Zero of Verbal Person Marking
86. Order of Person Markers on the Verb
87. Reciprocal Constructions
88. Passive Constructions
89. Antipassive constructions
90. Applicative constructions
91. Nonperiphrastic Causative Constructions
92. Negative Morphemes
93. Polar Questions
94. Predicative Adjectives
95. Zero Copula for Predicate Nominals
96. 'Want' Complement Subjects
97. Hand and Arm
98. Finger and Hand
99. Numeral Bases
100. Green and Blue
101. Red and Yellow
102. M-T Pronouns
103. $M$ in First Person Singular
104. N-M Pronouns
105. M in Second Person Singular
106. Position of Negative Word With Respect to Subject, Object, and Verb

## Sherzer features (1/3) - binary

1. Three vowel
2. 1-1-1
3. $2-1$
4. Four vowel
5. 2-2
6. 2-1-1
7. 1-2-1
8. Five vowel
9. 3-2
10. 3-1-1
11. 2-2-1
12. Six vowel
13. 2-2-2
14. 2-3-1
15. 3-2-1
16. Seven vowel
17. 2-2-2-1
18. 3-3-1
19. Voiceless vowel
20. Nasal vowel
21. not a,e,i,o,u
22. vowel length contrast
23. mid or mid-high vowel
24. one stop series: voiceless
25. two stop series: voiceless/voiced
26. two stop series: voiceless/glottalized
27. three stop series: voiceless/voiced/glottalized
28. four stop series
29. glottalized stop series
30. labial stop present
31. $\mathrm{c} / \mathrm{t} \mathrm{t}$
32. $k / c$
33. $k / q$
34. either $k / c ̌$ or $k / q$
35. t
36. q
37. kw
38. qw

## Sherzer features (2/3) - binary

39. one fricative series: voiceless
40. two fricative series: voiceless/voiced
41. two fricative series: voiceless/glottalized
42. three fricative series:
voiceless/voiced/glottalized
43. glottalized fricatives
44. pharyngeal fricatives
45. labial fricative
46. $\theta$
47. ð
48. $\mathrm{s} / \mathrm{\int}$
49. $z$
50. x
51. $x w$
52. x .
53. x.w
54. $\mathrm{\gamma}$
55. yw
56. h
57. hw
58. I
59. $\$$
60. t 4
61. $t \$^{\prime}$
62. dl
63. I'
64. \$'
65. ly
66. $\ddagger y$
67. voiceless nasal
68. glottalized nasal
69. n
70. ท
71. $r$
72. voiceless $r$
73. glottalized $r$
74. $\mathrm{r} / \mathrm{l}$
75. voiceless semivowel

## Sherzer features (3/3) - binary

76. glottalized semivowel
77. possessive pronouns independent morpheme
78. alienable/inalienable possession?
79. reduplication $=$ distributive or plual
80. reduplication $=$ diminutive
81. augmentative-diminutive consonant symbolism
82. masculine/feminine gender
83. animate/inanimate gender
84. plural in pronouns
85. inclusive/exclusive plural in pronouns
86. dual in pronouns
87. dual in nouns
88. inclusive/exclusive dual in pronouns
89. demonstratives for visible/invisible objects
90. numerals classified by form or shape of object
91. locative prefixes
92. locative suffifxes
93. locative prepositions
94. locative postpositions
95. nominal incorporation
96. subject person marker prefixes
97. subject person marker suffixes
98. subject person markers independent pronouns
99. reduplication in verb $=$ distribution, repetition
100. reduplication in verb $=$ diminutive
101. evidential or source of information marked
102. instrumental markers
103. locative-directional markers
104. locative-directional markers prefix
105. locative-directional markers suffix
