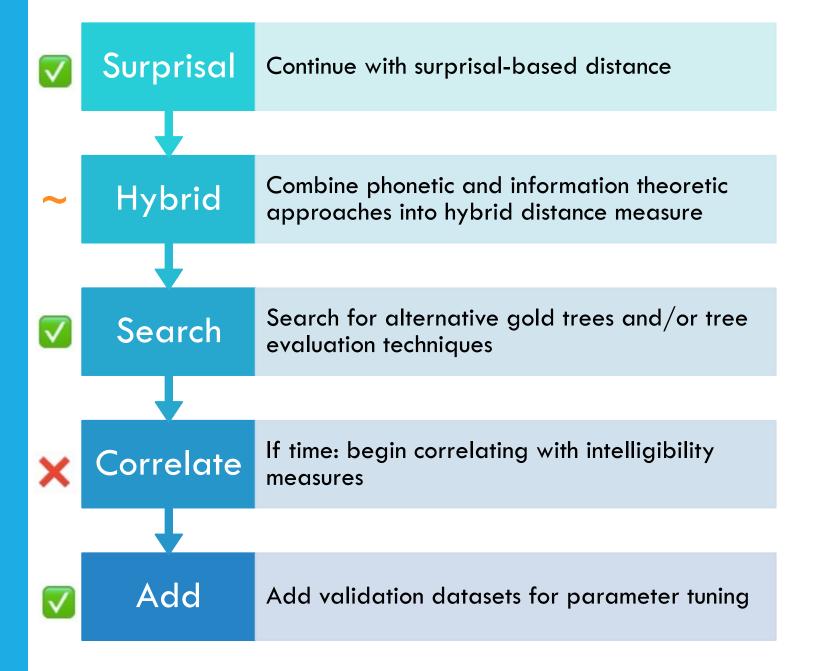


# MASTER'S THESIS MEETING

Philip Georgis
October 13, 2021

## CURRENT TASKS



## **VALIDATION DATASET**

#### Bantu

- Sample of some of the largest among the >300 doculects from dataset
- Hellenic
  - 4 modern Hellenic doculects + 2 varieties of Ancient Greek
- Japonic
  - Japanese and Ryukyuan doculects + Old Japanese; omitted Middle Japanese
- Quechuan
  - Omitted doculects without glottocodes, and non-Quechuan languages from dataset
- Uto-Aztecan
  - Omitted non-Uto-Aztecan languages from dataset
- Vietic
  - Omitted non-Vietic languages, Proto-Vietic, and ambiguous doculects

# DATASETS OVERVIEW

#### **Validation Set**

Family	Number of Languages	Time Depth	Macroarea
Bantu	37	5000	Africa
Hellenic	6	2400	Eurasia
Japonic	58	2400	Eurasia
Quechuan	33	1850	South America
Uto-Aztecan	34	7000	North America
Vietic	17	2750	Eurasia
Average	31	3567	

Family	Number of Languages	Time Depth	Macroarea
Arabic	16	1600	Africa & Eurasia
Balto-Slavic	16	3000	Eurasia
Dravidian	20	6000	Eurasia
Hokan	20	5000	North America
Italic	58	2100	Eurasia
Polynesian	30	3000	Papunesia
Sinitic	19	2500	Eurasia
Turkic	32	2100	Eurasia
Uralic	23	8000	Eurasia
Average	26	3700	

## DATASETS OVERVIEW



- Word pairs aligned using typical alignment algorithm (costs based on phonetic similarity + PMI)
  - Ancient Attic Greek  $<\alpha \tilde{i}\mu\alpha>/h\hat{a}jma/$  Modern Demotic Greek  $<\alpha i\mu\alpha>/ema/$  'BLOOD'

AAG	h	a	j	m	a
MDG	-	-	e	m	a

- Word pairs aligned using typical alignment algorithm (costs based on phonetic similarity + PMI)
  - Ancient Attic Greek <αἷμα> /hâjma/
     Modern Demotic Greek <αίμα> /ˈema/

Adaptation surprisal calculated for each pair in alignment in both directions

AAG	h	a	j	m	a
MDG	-	-	e	m	a
Surprisal	1.41	2.61	3.17	1.32	0.61

**TOTAL:** 9.12

'BLOOD'

MDG	-	-	e	m	a
AAG	h	a	j	m	a
Surprisal	3.10	3.31	4.21	0.90	0.87

**TOTAL:** 12.39

•Self-surprisal based on trigram probabilities calculated for each word

AAG	h	a	j	m	a
Self-Surprisal	3.29	2.00	1.91	0.92	2.98

**TOTAL:** 11.10

MDG	e	m	a
Self-Surprisal	3.35	2.14	2.10

**TOTAL:** 7.59

•Self-surprisal based on trigram probabilities calculated for each word

AAG	h	a	j	m	a
Self-Surprisal	3.29	2.00	1.91	0.92	2.98

**TOTAL:** 11.10

MDG	e	m	a
Self-Surprisal	3.35	2.14	2.10

**TOTAL:** 7.59

- Adaptation surprisal normalized by self-surprisal
  - $S(MDG \mid AAG) / S(MDG) = 9.12 / 7.59 = 1.20$
  - S(AAG | MDG) / S(AAG) = 12.39 / 11.10 = 1.12
- Take mean of normalized adaptation surprisal in each direction
  - → 1.16

•Self-surprisal based on trigram probabilities calculated for each word

AAG	h	a	j	m	a
Self-Surprisal	3.29	2.00	1.91	0.92	2.98

**TOTAL:** 11.10

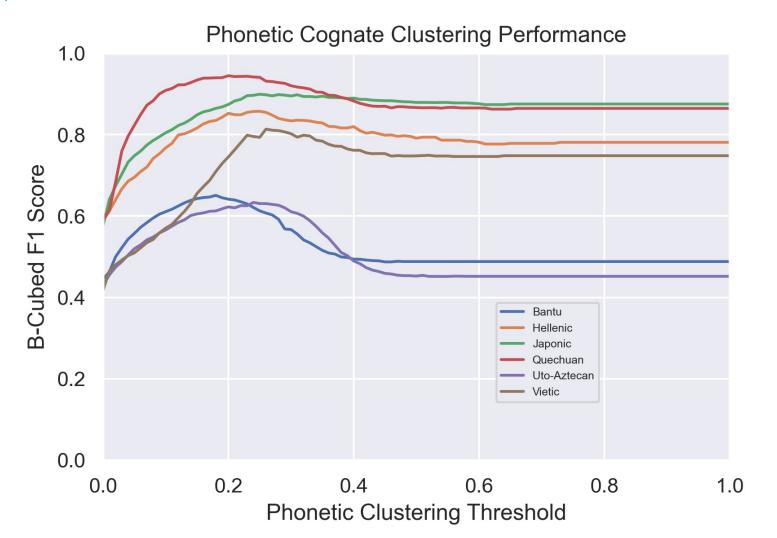
MDG	e	m	a
Self-Surprisal	3.35	2.14	2.10

**TOTAL:** 7.59

- Adaptation surprisal normalized by self-surprisal
  - $S(MDG \mid AAG) / S(MDG) = 9.12 / 7.59 = 1.20$
  - S(AAG | MDG) / S(AAG) = 12.39 / 11.10 = 1.12
- Take mean of normalized adaptation surprisal in each direction
  - $\rightarrow$  1.16  $\rightarrow$  turn into similarity: e^^-1.16 = 0.31

### OPTIMAL COGNATE CLUSTERING THRESHOLDS

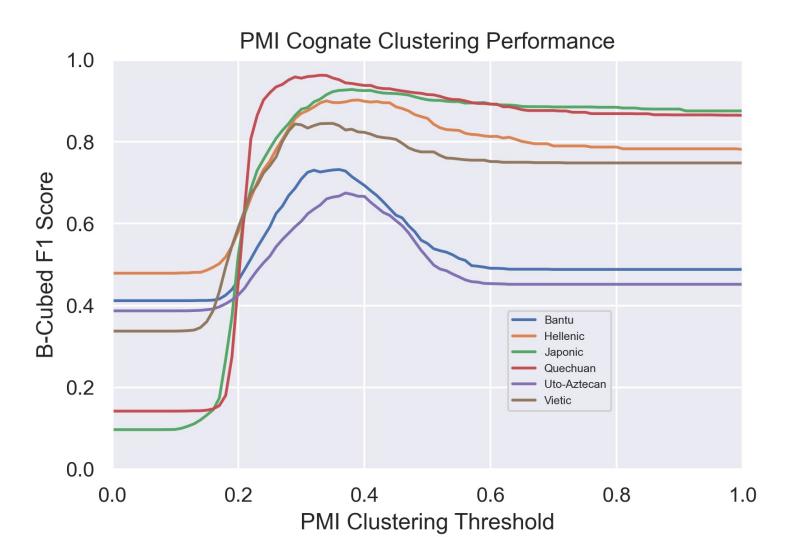
#### BASED ON VALIDATION DATASETS



Optimal threshold = 0.23

### OPTIMAL COGNATE CLUSTERING THRESHOLDS

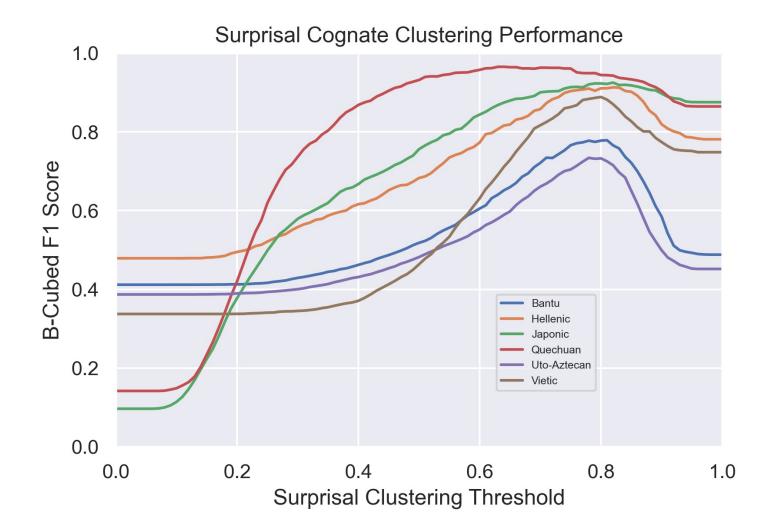
#### BASED ON VALIDATION DATASETS



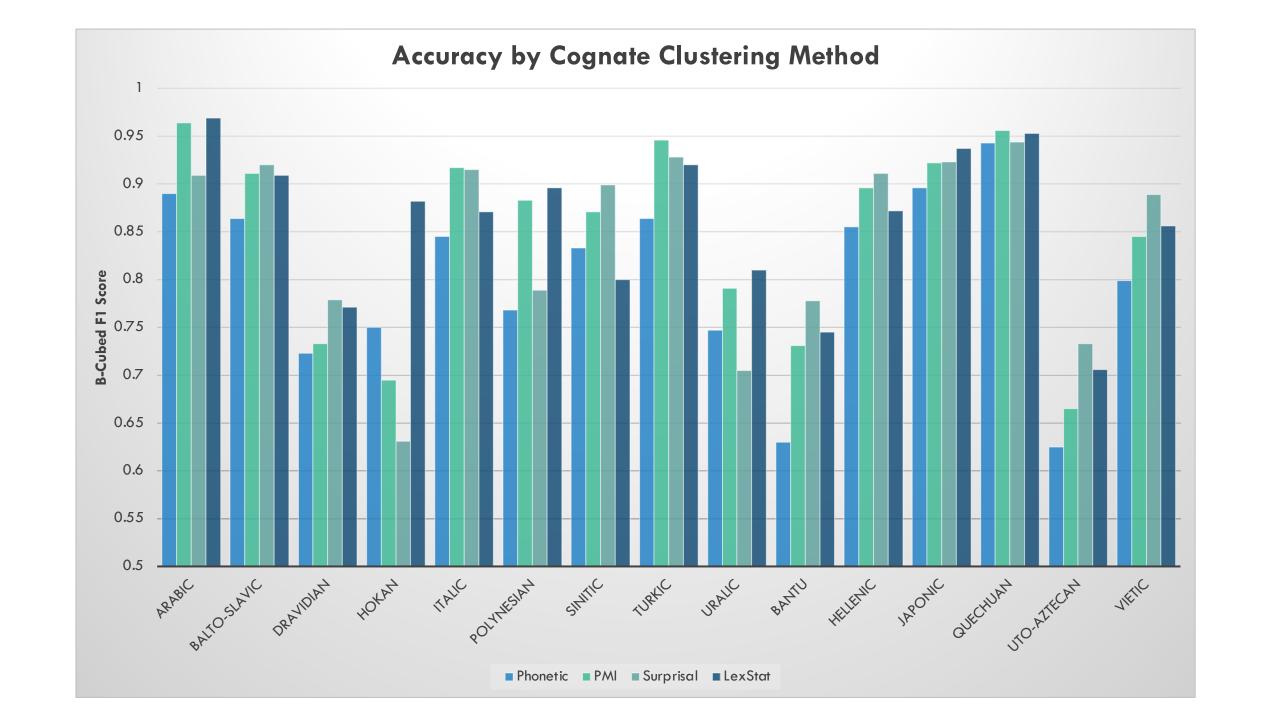
Optimal threshold = 0.35

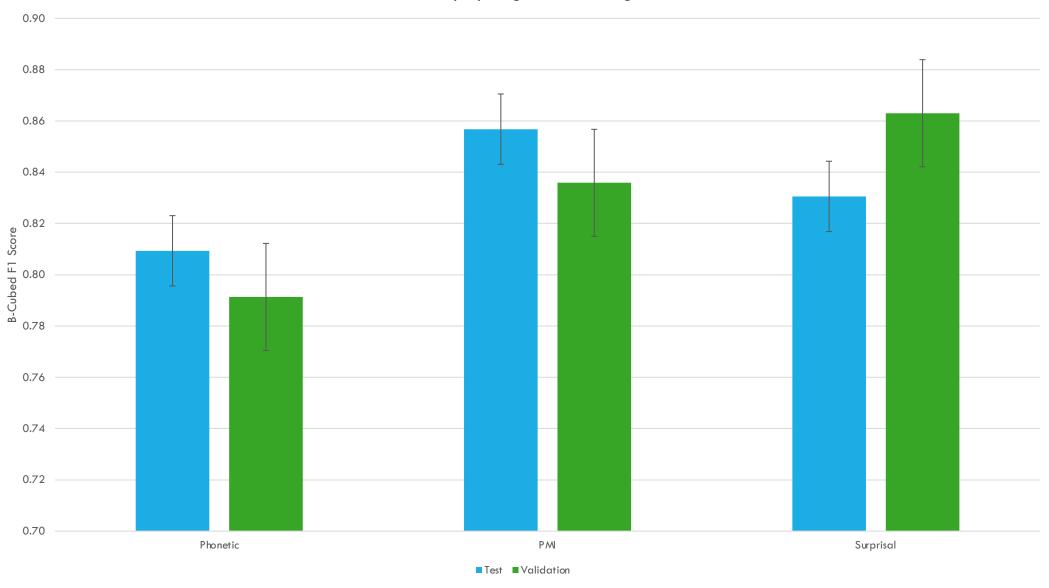
### OPTIMAL COGNATE CLUSTERING THRESHOLDS

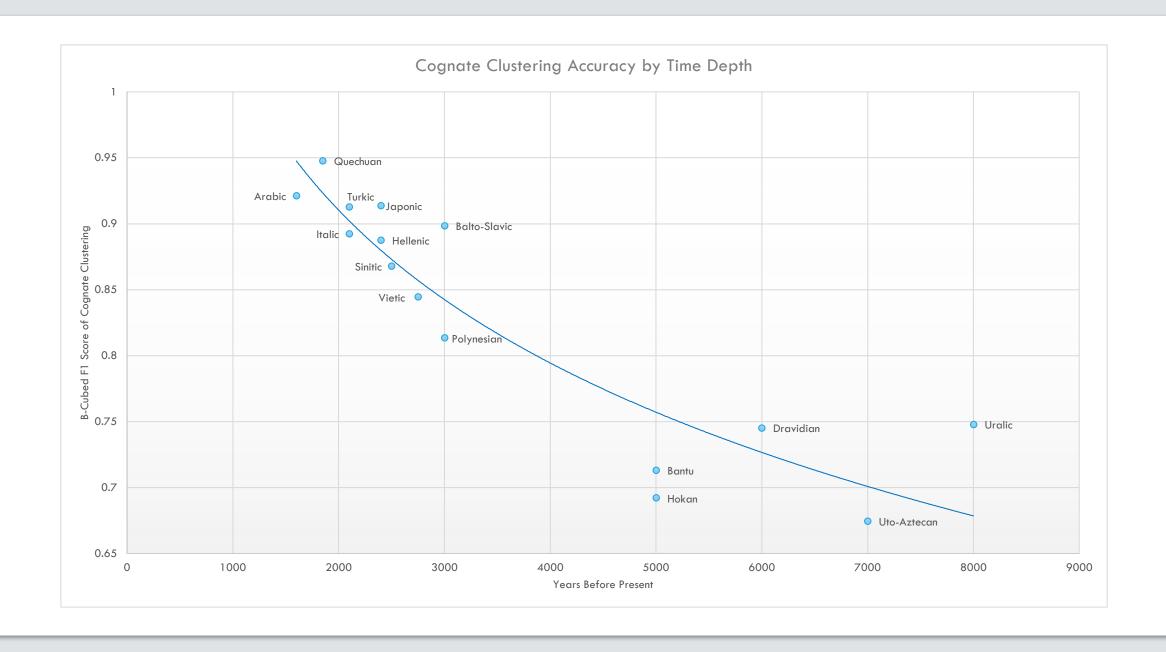
#### BASED ON VALIDATION DATASETS



Optimal threshold = 0.80

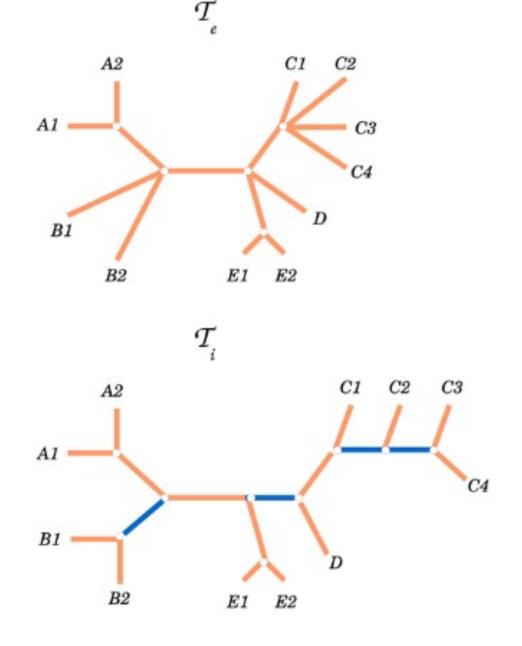






## IMPROVED TREE COMPARISON

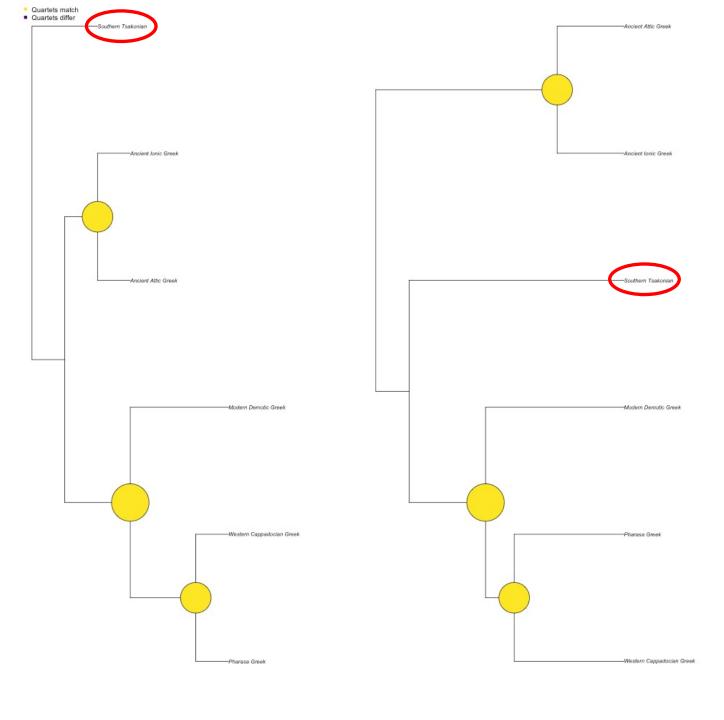
- Found paper which discusses impact of comparing resolved, binary trees with unresolved, non-binary trees on Quartet Distance and Robinson-Foulds distance
- Pompei, S., Loreto, V., & Tria, F. (2011). On the Accuracy of Language Trees. PLoS ONE, 6(6).
   <a href="https://doi.org/doi:10.1371/journal.pone.0020109">https://doi.org/doi:10.1371/journal.pone.0020109</a>
- GQD seems to be the measure which is used in other phylogenetic inference papers
- Tree Dist presumably has the same issue but found no modification → GQD as primary evaluation now



### WEIRD BUG?

- Quartet Distance and Tree Distance seem not to penalize an outgroup item being grouped elsewhere in the tree
- O distance for both measures between these two trees, despite Southern Tsakonian appearing in different positions

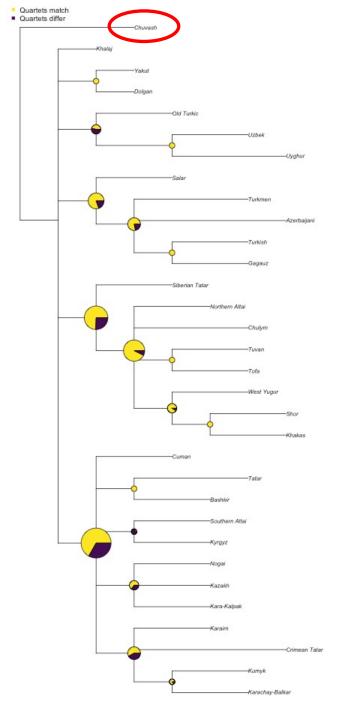
Why? How to fix?

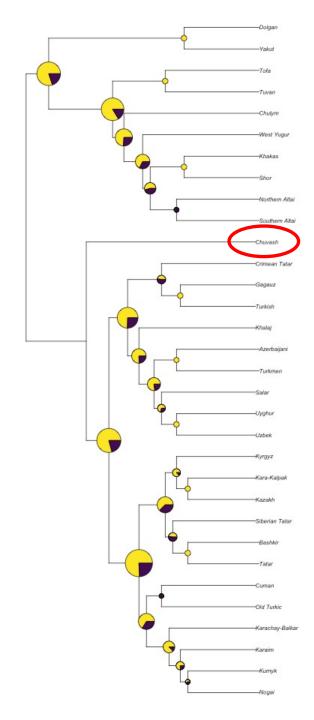


## WEIRD BUG?

- Quartet Distance and Tree Distance seem not to penalize an outgroup item being grouped elsewhere in the tree
- Similar issue in Turkic tree with placement of Chuvash

Why? How to fix?





## BEST AUTOMATIC TREES (NOT USING GOLD COGNATE SETS)

Dataset	TreeDist	Generalized Quartet Distance	Cognate Clustering Method	Word Form Evaluation Method	Linkage Method
Arabic	0.67	0.22	PMI	surprisal	Ward
Balto-Slavic	0.16	0.02	none	surprisal	complete
Dravidian	0.41	0.22	none	phonetic	average
Hokan	0.40	0.04	surprisal	phonetic	average
Italic	0.39	0.16	none	surprisal	Ward
Polynesian	0.58	0.12	PMI	phonetic	Ward
Sinitic	0.61	0.16	none	phonetic	average
Turkic	0.56	0.31	surprisal	PMI	Ward
Uralic	0.32	0.04	none	phonetic	complete

#### **NOTES:**

- Significantly lower tree distances using
   Generalized Quartet
   Distance
- No separation of cognate/non-cognates before evaluation often produces best trees
- Phonetic (5/9) and surprisal (3/9) are the most effective evaluation methods

## BEST GOLD TREES (USING GOLD COGNATE SETS)

Dataset	TreeDist	Generalized Quartet Distance	Word Form Evaluation Method	Linkage Method
Arabic	0.64	0.30	PMI	Ward
Balto-Slavic	0.16	0.02	surprisal	complete
Dravidian	0.44	0.26	surprisal	complete
Hokan	0.36	0.02	phonetic	average
Italic	0.41	0.21	surprisal	Ward
Polynesian	0.57	0.12	PMI	Ward
Sinitic	0.63	0.27	surprisal	Ward
Turkic	0.57	0.32	surprisal	weighted
Uralic	0.40	0.05	phonetic	average

#### **NOTES:**

- Most trees based on gold cognate sets are less accurate than those using either automatic cognate sets or no separation of cognates/noncognates before evaluation
- Only Hokan tree is better using gold cognate sets
- **Surprisal** seems to be the most effective word evaluation technique (5/9)

## TIME PLAN...

Finish coding distance measures: OCTOBER 19

• Finish testing on cognate detection: OCTOBER 22

• Finish testing on trees: OCTOBER 29

Finish all data collection/experiment: NOVEMBER 1

• First draft complete: NOVEMBER 15 \*

Second draft complete: DECEMBER 1

<sup>\*</sup> Will have begun writing sooner

## NEXT TASKS

