

Analyzing Correlated Evolution of Multiple Features Using Latent Representations

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Kyoto University



Two Types of Data

1. Modern languages represented by typological features

152 discrete features

Tetum

1	2	...	1
---	---	-----	---

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Feature 26A: Prefixing vs. Suffixing in Inflectional Morphology

1. Little affixation
2. Strongly suffixing
3. Weakly suffixing
4. Equal prefixing and suffixing
5. Weakly prefixing
6. Strong prefixing

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Feature 81A: Prefixing vs. Suffixing in Inflectional Morphology

1. Feature 81A: Order of Subject, Object and Verb
2. 1. SOV
3. 2. **SVO**
4. 3. VSO
5. 4. VOS
6. 5. OVS
6. OSV:
7. No dominant Order

Two Types of Data

1. Modern languages represented by typological features

152 discrete features

2,557 languages

1	2	...	1
2	4	...	0
1	1	...	1
⋮	⋮	⋮	⋮
3	1	...	1

Two Types of Data

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2,557 languages

1	2	...	1
2	4	...	?
1	?	...	1
⋮	⋮	⋮	⋮
?	1	...	?

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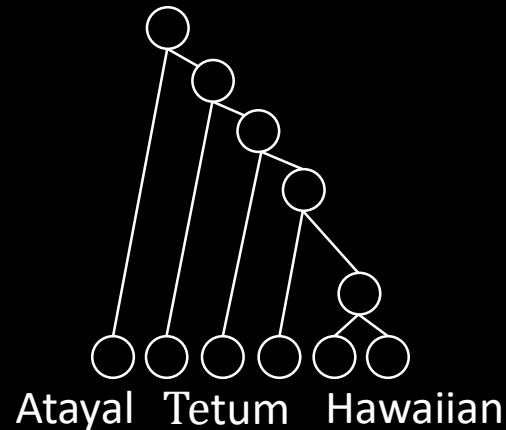
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2. Phylogenetic trees relating these modern languages



Two Types of Data

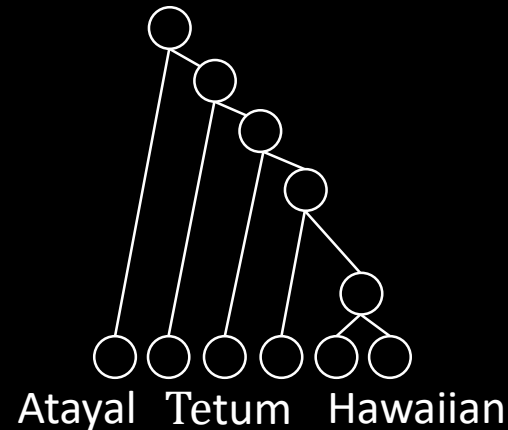
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2. Phylogenetic trees relating these modern languages

Proto-Austronesian



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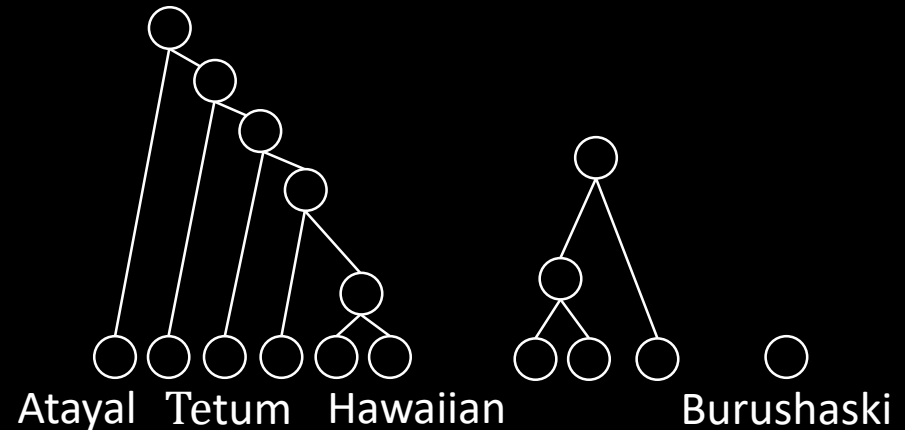
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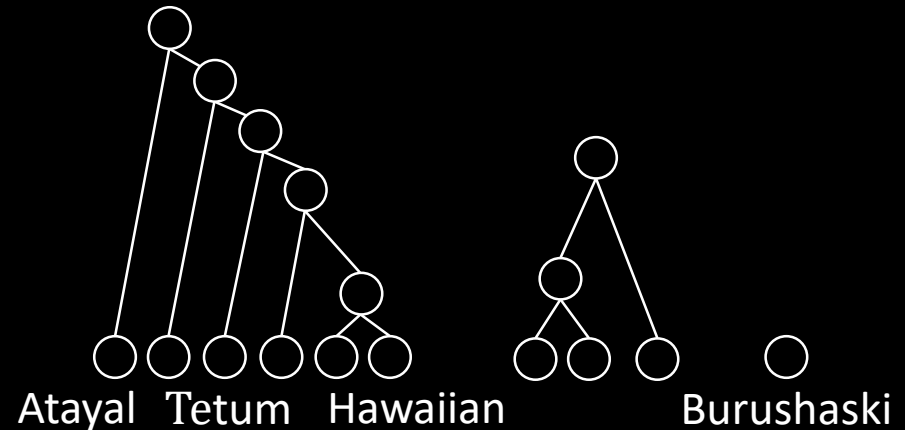
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2. Phylogenetic trees relating these modern languages

309 language families including 154 language isolates

Proto-Austronesian

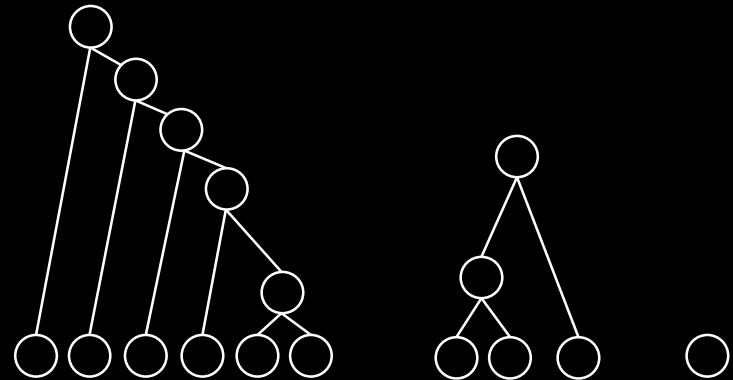


How have languages changed in the past?
How are they likely to change in the future?

To answer these questions,
I develop statistical models that make use of

1	2	...	1
2	4	...	0
1	1	...	1
⋮	⋮	⋮	⋮
3	1	...	1

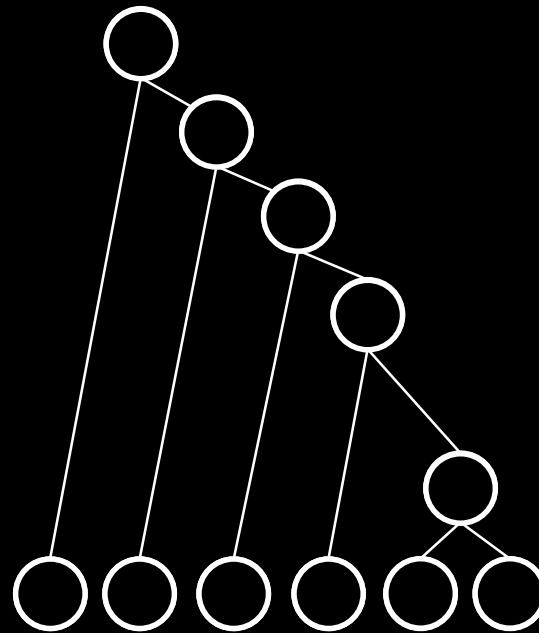
and



Phylogenetic Comparative Method

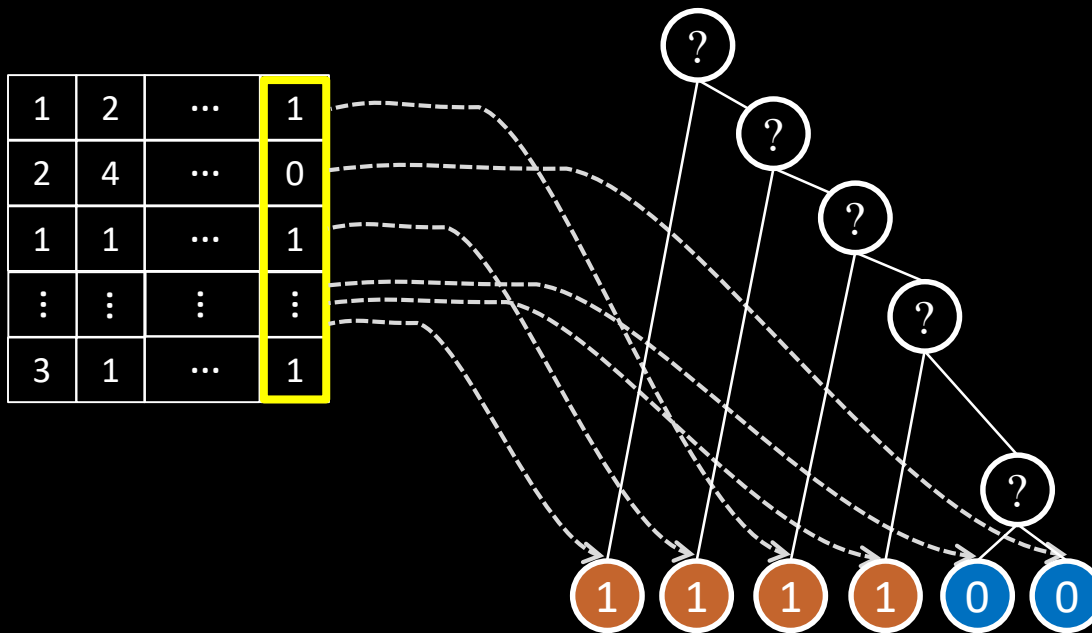
Trees allow us to infer ancestral states, with varying degrees of confidence

1	2	...	1
2	4	...	0
1	1	...	1
⋮	⋮	⋮	⋮
3	1	...	1



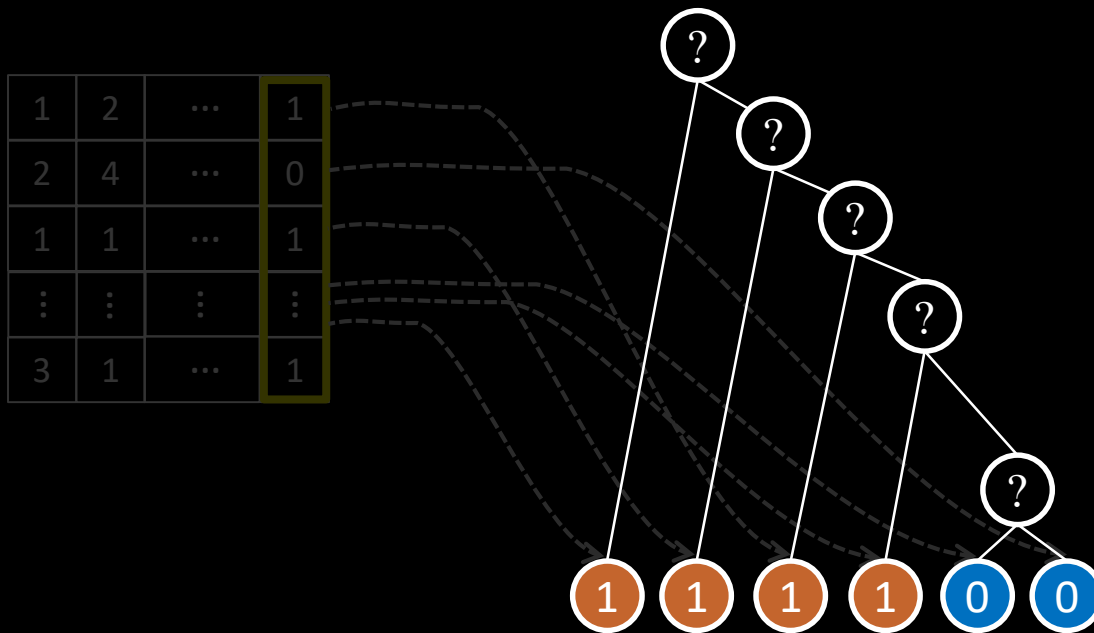
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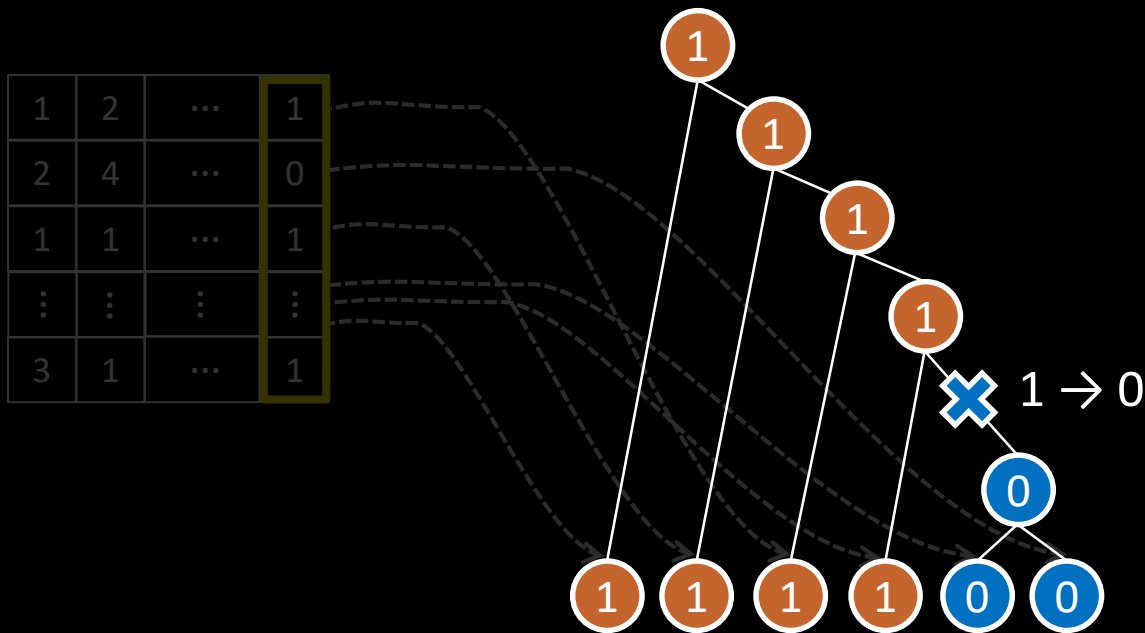
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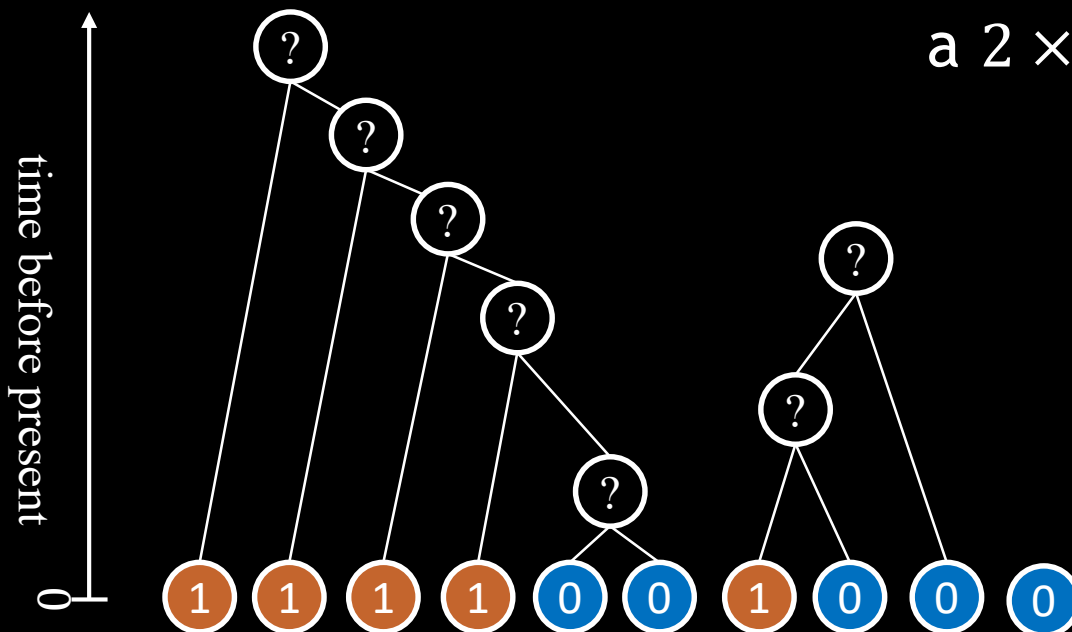
Continuous-time Markov Chains (CTMCs) for Statistical Analysis

Transition probability as a function of *continuous* time

$$P(x = b \mid \text{parent}(x) = a, t) = \exp(tQ)_{a,b}$$

A binary feature has
a 2×2 transition rate matrix

$$Q = \begin{pmatrix} -\alpha & \alpha \\ \beta & -\beta \end{pmatrix}$$



[Greenhill+, 2010]
[Maurits+, PNAS, 2014]

Wait, Features are not Independent Need to Model *Correlated* Evolution

1	2	⋯	1
2	4	⋯	0
1	1	⋯	1
⋮	⋮	⋮	⋮
3	1	⋯	1

Implicational universals [Greenberg, 1963]

Order of
object and verb

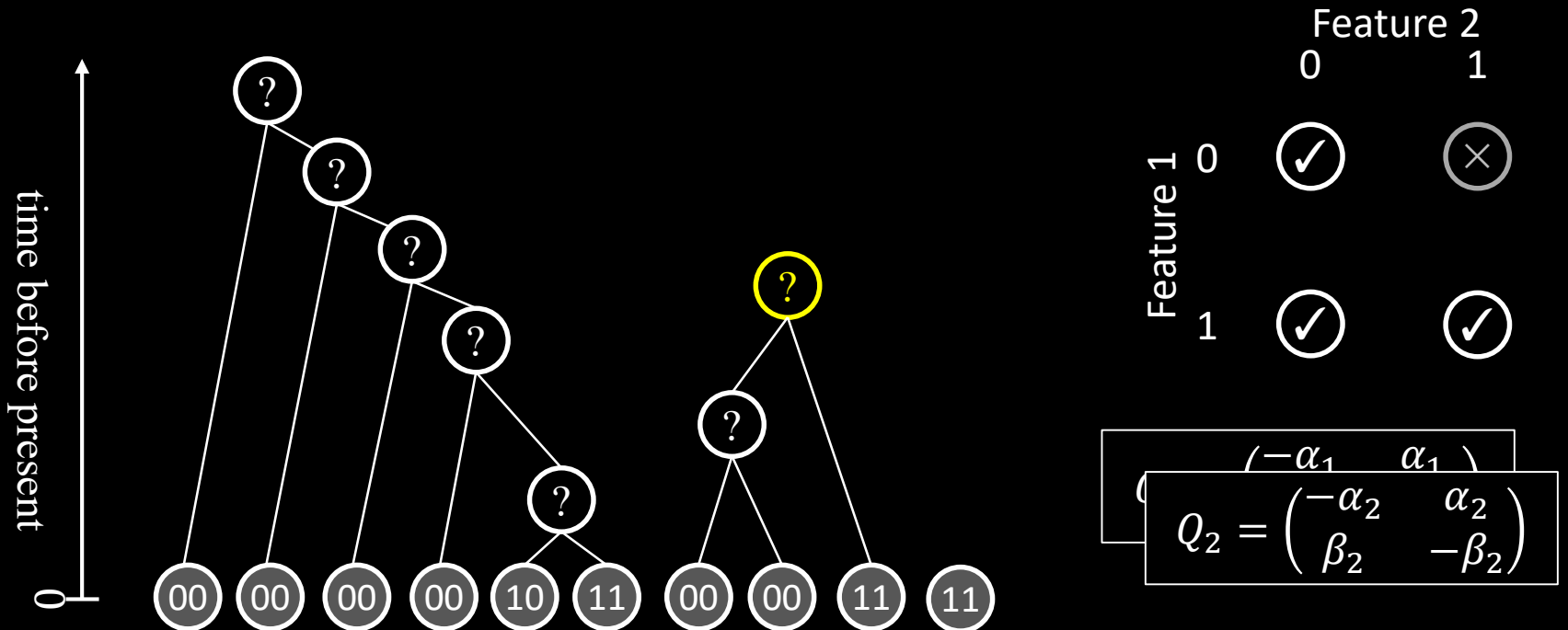
Order of noun and
relative clause

	NRel	RelN
VO	✓	×
OV	✓	✓

[Dryer, 2011]

Independent CTMCs

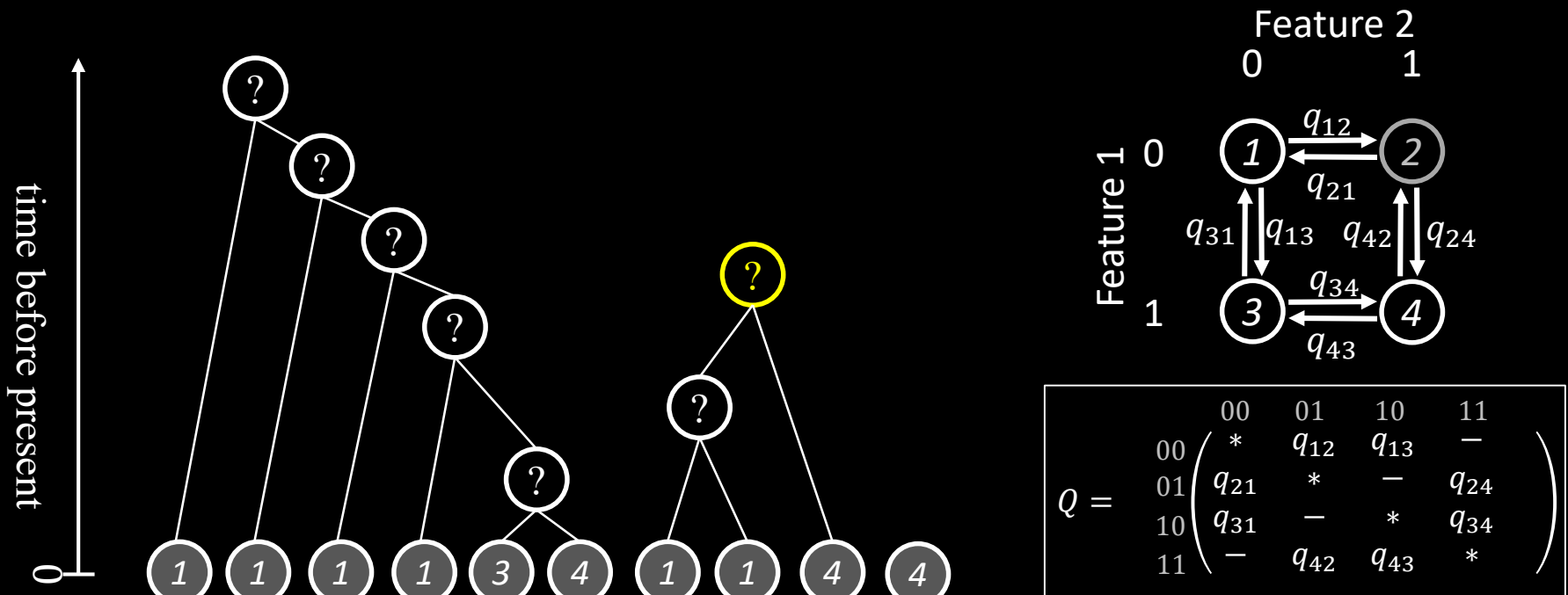
Unable to take into account the observation that the feature combination **01** is unnatural



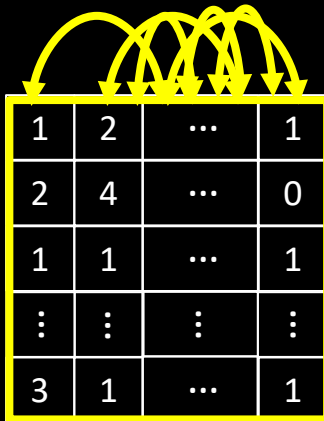
Expanding Feature Combinations Does not Scale

[Dunn+, Nature, 2011]

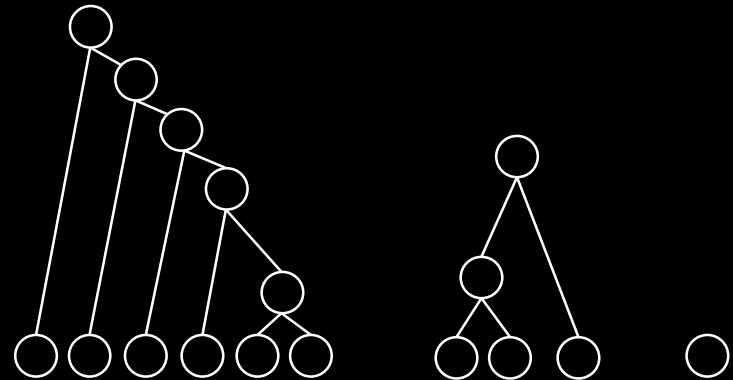
Combinatorial explosion prevents us from modeling interactions involving *multiple* features



My Goal: Model Correlated Evolution Covering All Possible Dependencies



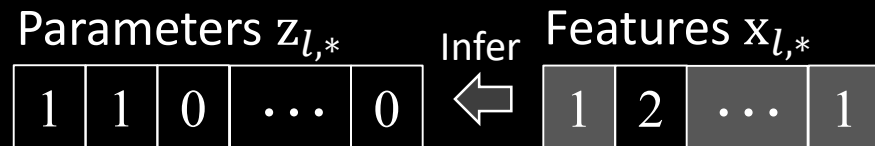
1	2	...	1
2	4	...	0
1	1	...	1
⋮	⋮	⋮	⋮
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My Solution: Latent Representations

Idea originally presented in [Murawaki, NAACL2015]

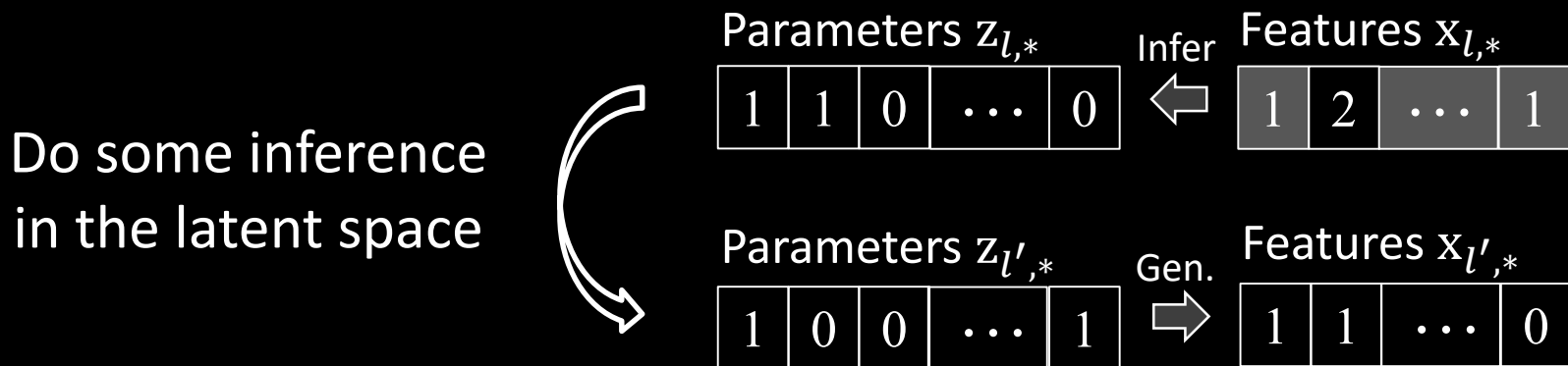
- Reorganize 152 discrete surface features into 100 binary latent *parameters*
 - Parameters are **independent** by assumption



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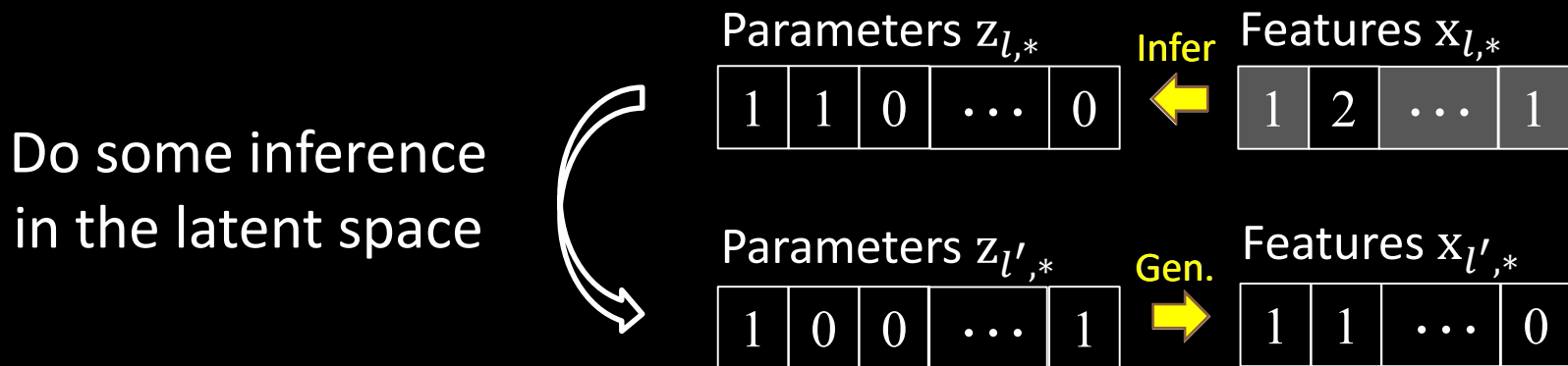
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- Inference in the latent space implicitly captures correlated evolution



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Latent Representations Capture Inter-Feature Dependencies

[Murawaki, IJCNLP2017]

Parameters $z_{l,*}$

1	0	1	...	0
---	---	---	-----	---

Features $x_{l',*}$

1	2	...	1
---	---	-----	---

Latent Representations Capture Inter-Feature Dependencies

[Murawaki, IJCNLP2017]

Parameters $z_{l,*}$

1	0	1	...	0
---	---	---	-----	---

×

Weight matrix W

2.9	0.4	-0.3	...	-0.2
6.3	-4.3	-5.7	...	5.9
8.2	-0.2	-2.5	...	0.3
⋮	⋮	⋮	⋮	⋮
0.2	0.3	1.2	...	-2.4

=

Feature score vector

10.2	-9.8	-8.9	...	-4.9
------	------	------	-----	------

Drawn from locally normalized distributions ↓

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1	2	...	1
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Drawn from locally normalized distributions ↓

Features $x_{l',*}$

1	2	...	1
---	---	-----	---

For $z_{l,k} = 1$, $w_{k,f(i_1,j_1)} \gg 0$ and $w_{k,f(i_2,j_2)} \ll 0$ indicate that feature i_1 is likely to take value j_1 and that feature i_2 is unlikely to take value j_2

Estimate Transition Rate Matrices for Parameters and Use Them for Simulation

Do some inference
in the latent space



Parameters $z_{l,*}$

1	1	0	...	0
---	---	---	-----	---

Infer

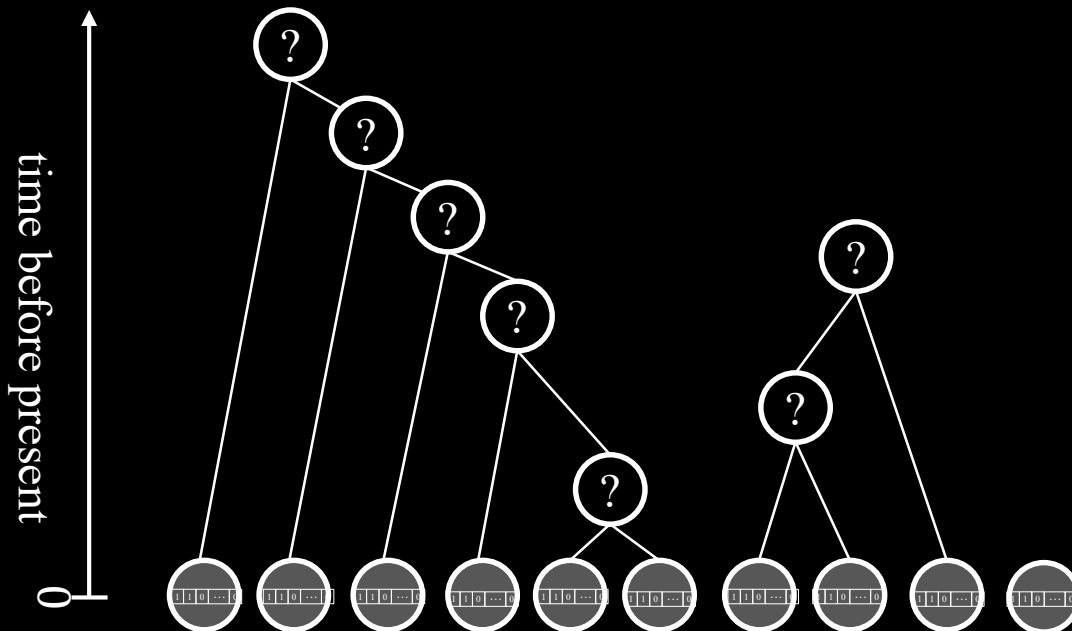
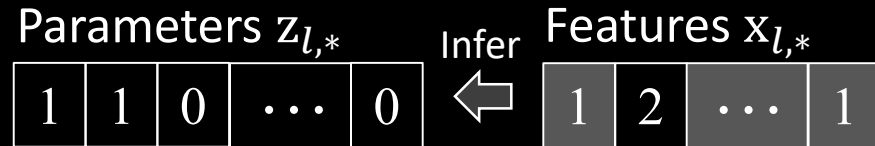


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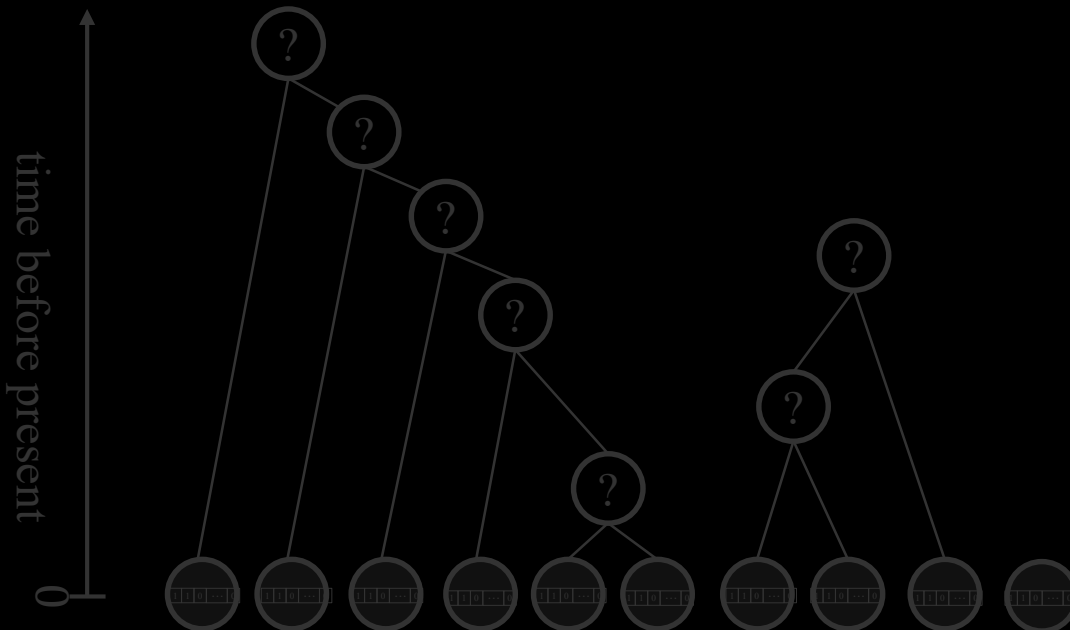
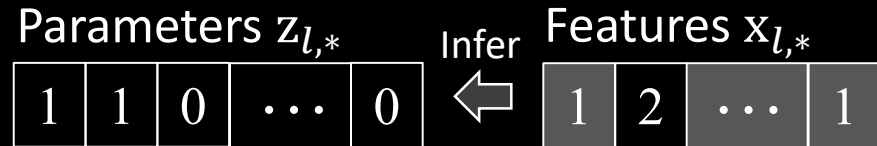
Do some inference in the latent space



$$Q_{100} = \begin{pmatrix} -\alpha_{100} & \alpha_{100} \\ \beta_{100} & -\beta_{100} \end{pmatrix}$$

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Parameters $z_{l,*}$

1	1	0	...	0
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Infer



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1	2	...	1
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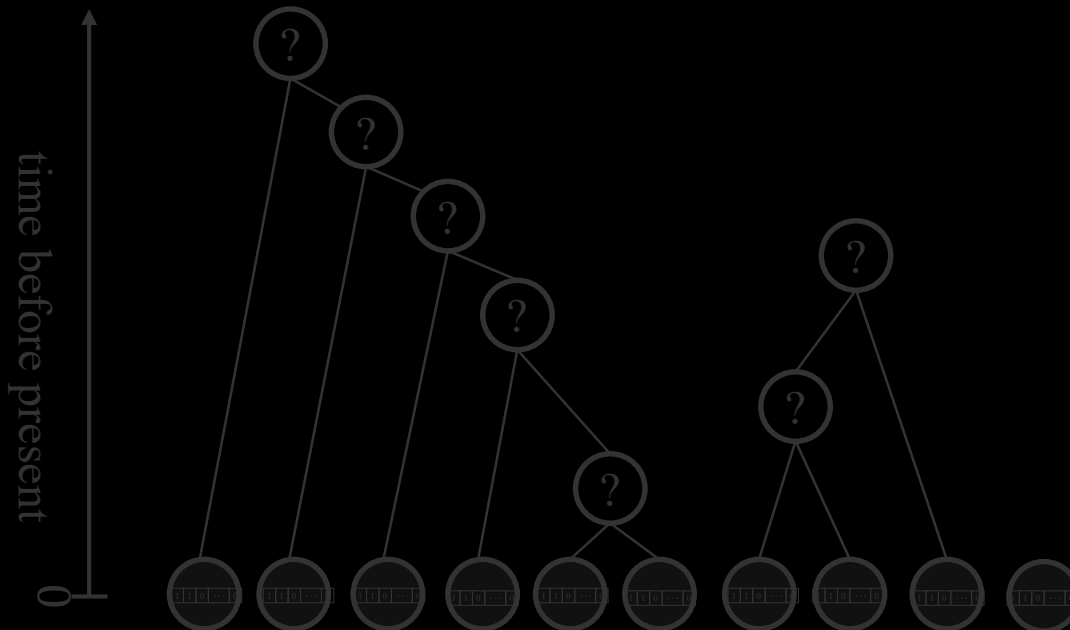
1	0	0	...	1
---	---	---	-----	---

Gen.



Features $x_{l',*}$

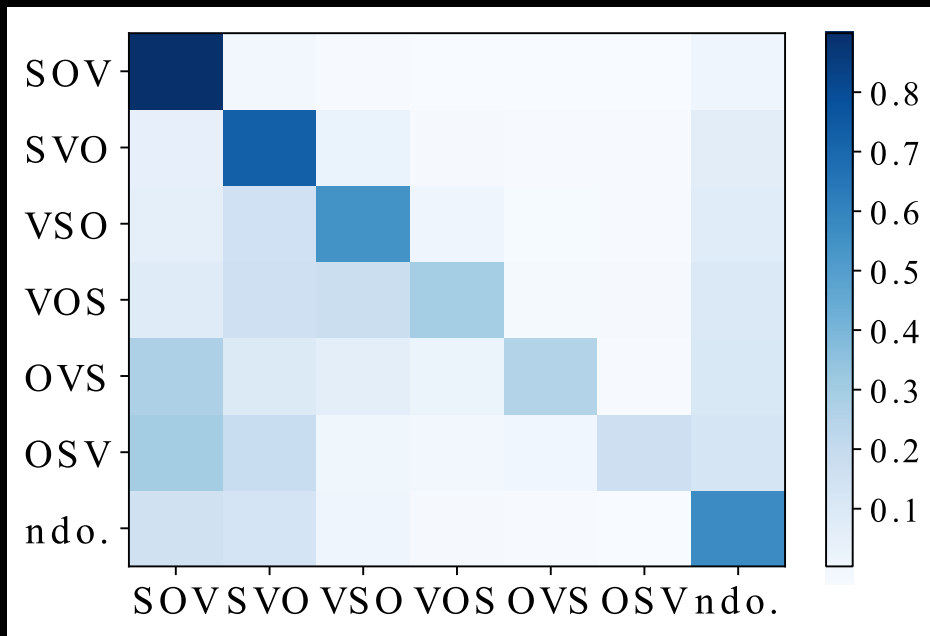
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Results: Order of Subject, Object and Verb

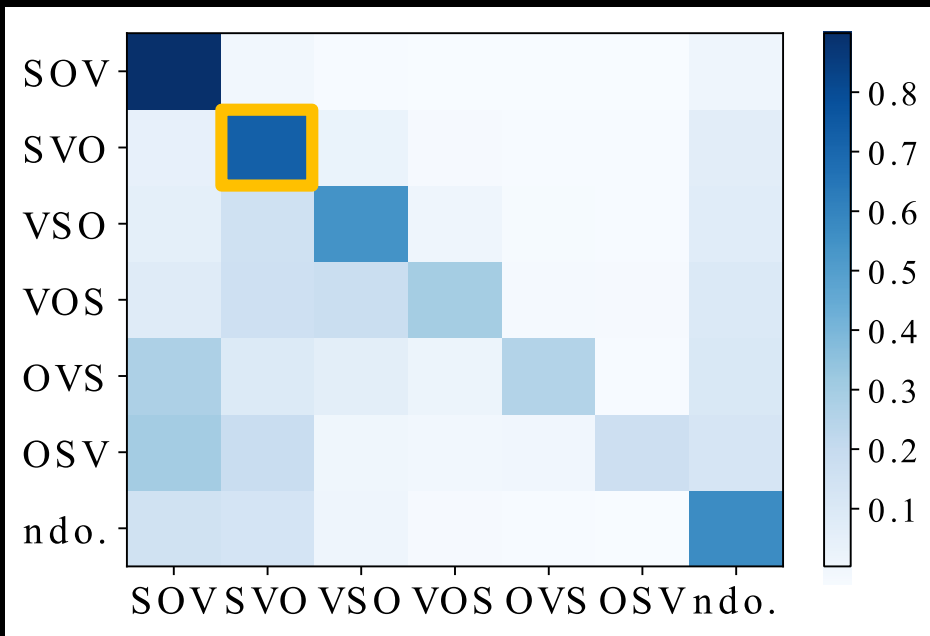
Transition prob. with $t = 2,000$



Largely agree with the results of
[Maurits+ PNAS, 2014]

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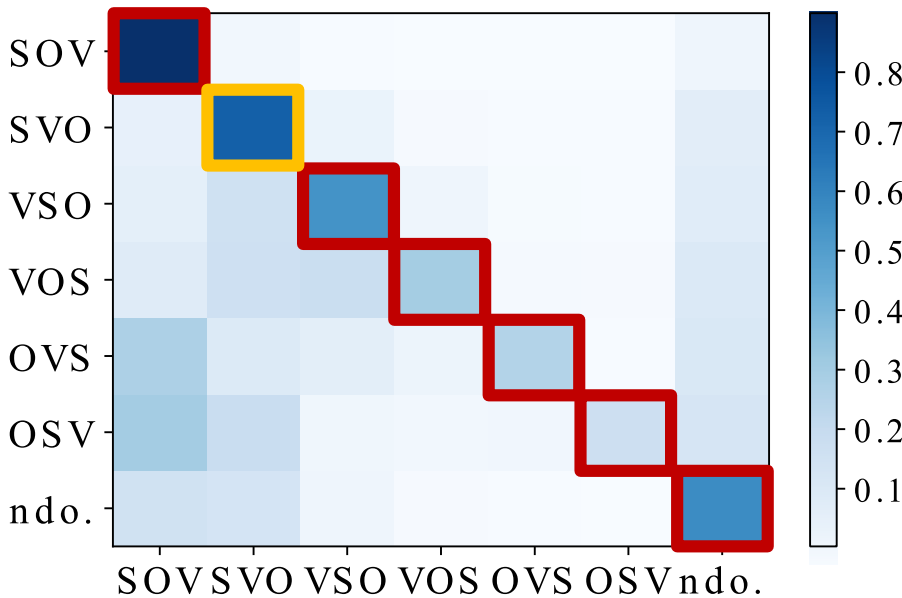
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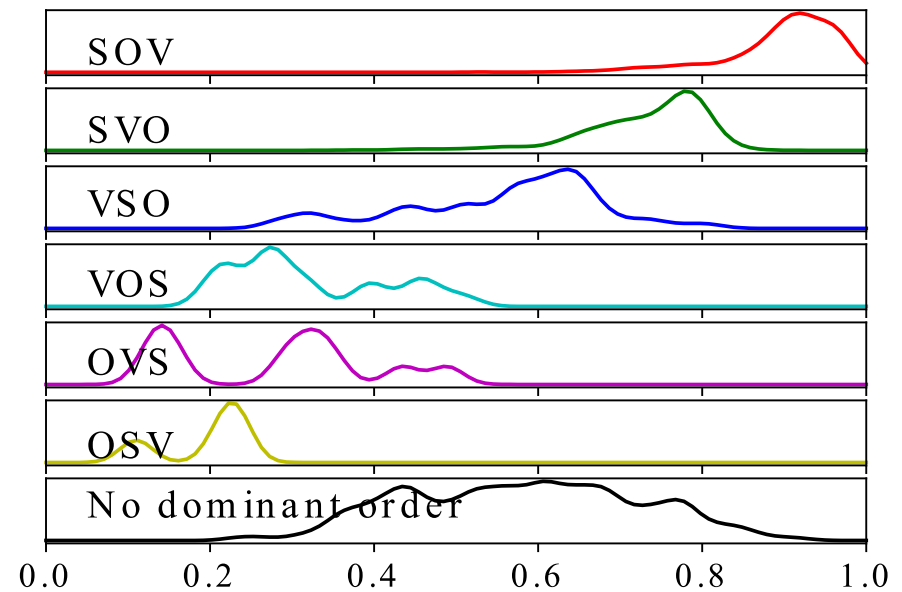
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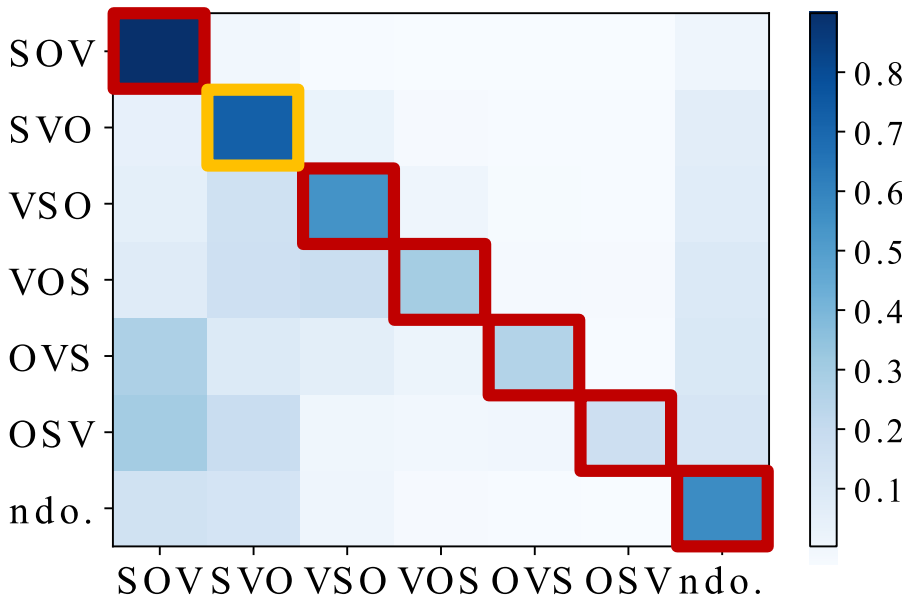
Variability in prob. of keeping the same order



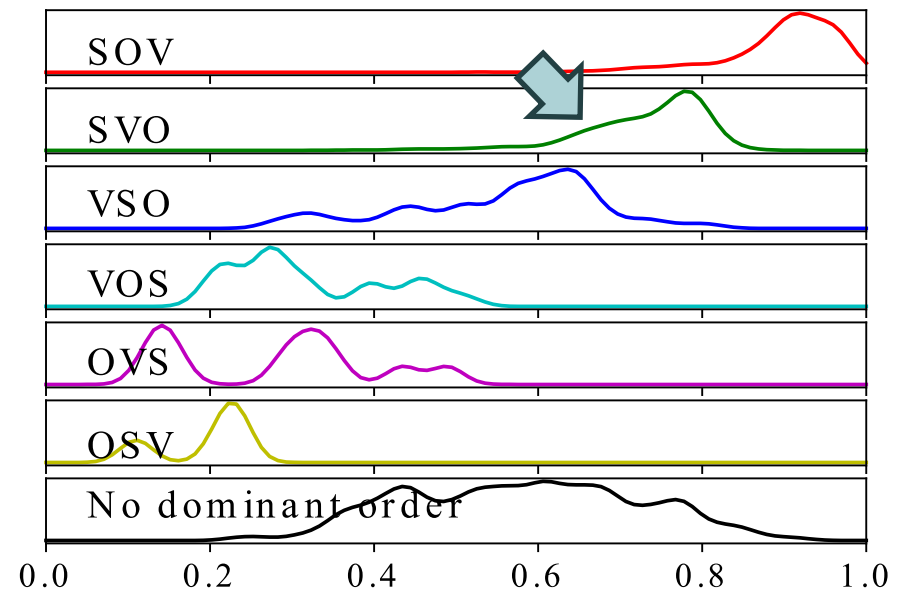
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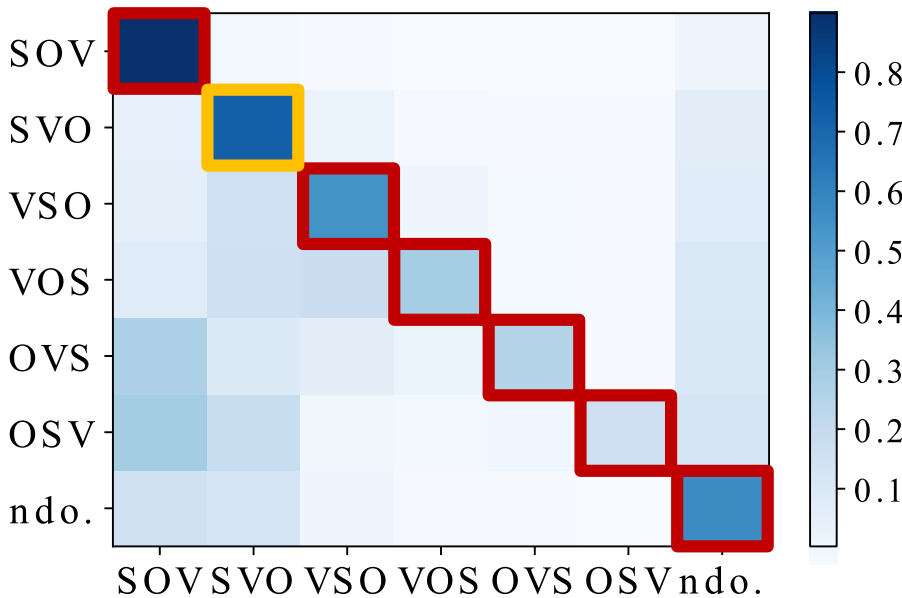
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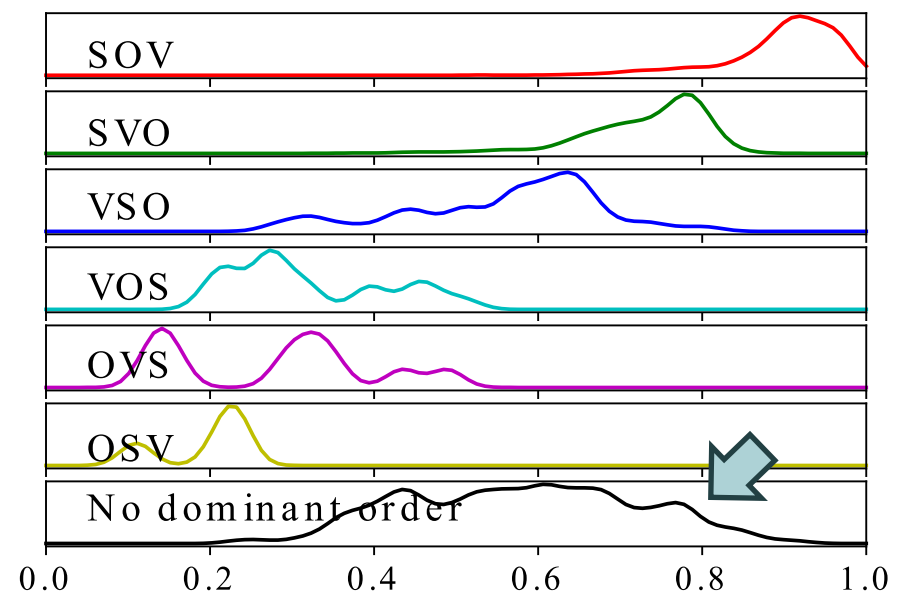
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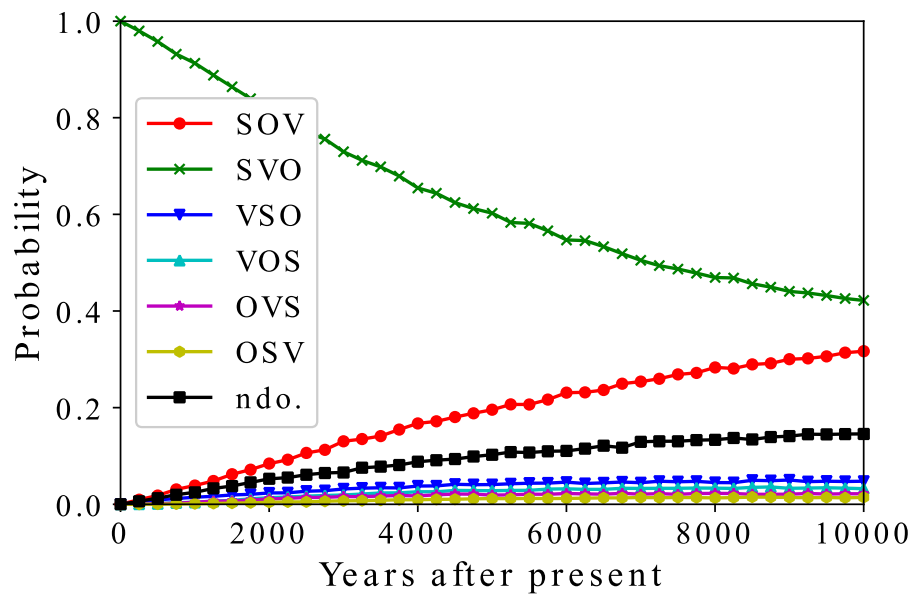
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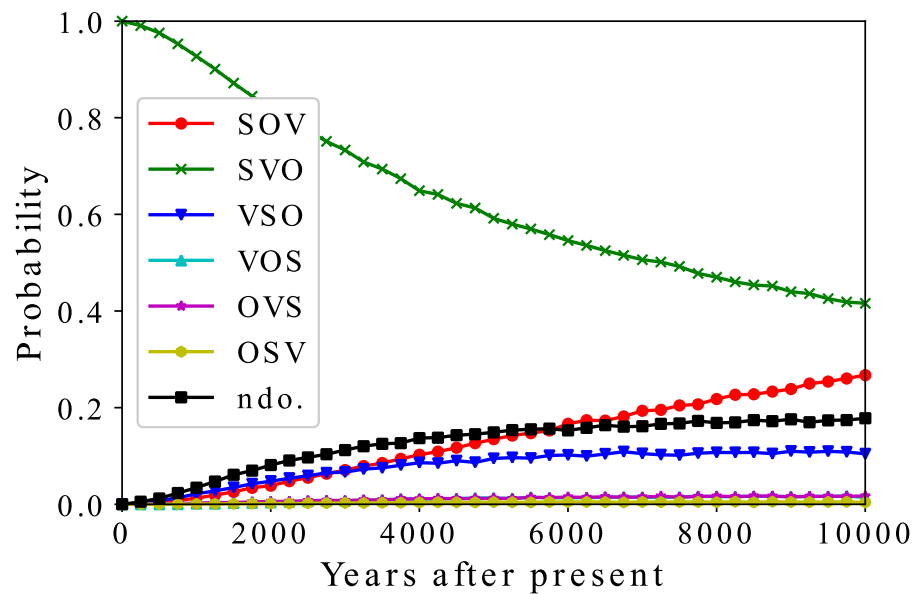
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Results: Order of Subject, Object and Verb

Tetum (Austronesian)



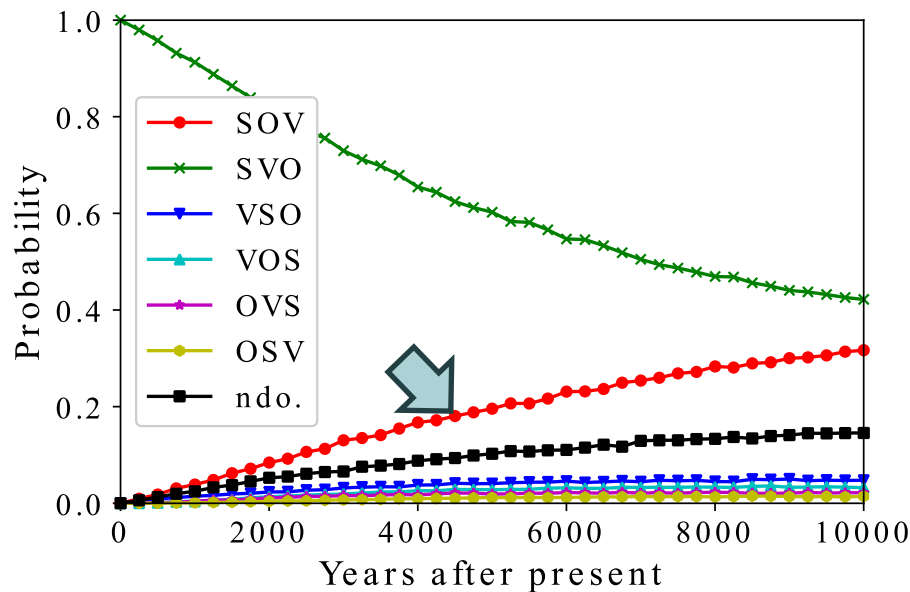
South-Central Kikongo (Atlantic-Congo)



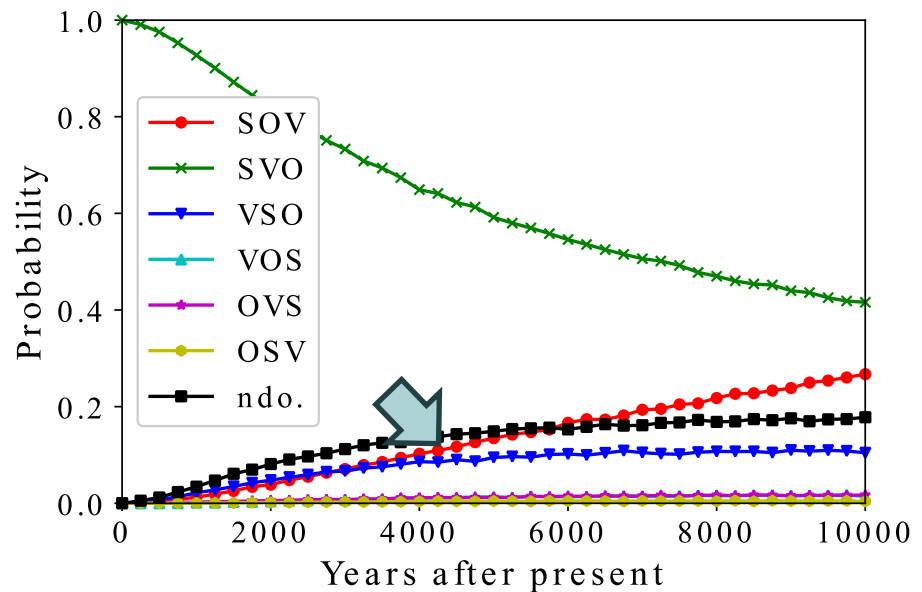
The strongly prefixing language, South-Central Kikongo, is more resistant to the change to SVO than Tetum

Results: Order of Subject, Object and Verb

Tetum (Austronesian)



South-Central Kikongo (Atlantic-Congo)



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Conclusions

- Proposed a new framework of latent representation-based analysis of diachronic typology
 - Investigate correlated evolution in an exploratory manner
- Future work
 - Analyze features other than the order of subject, object and verb
 - Inspect inferred ancestral states
 - Modeling contacts

