Typological Word Order Correlations with Logistic Brownian Motion

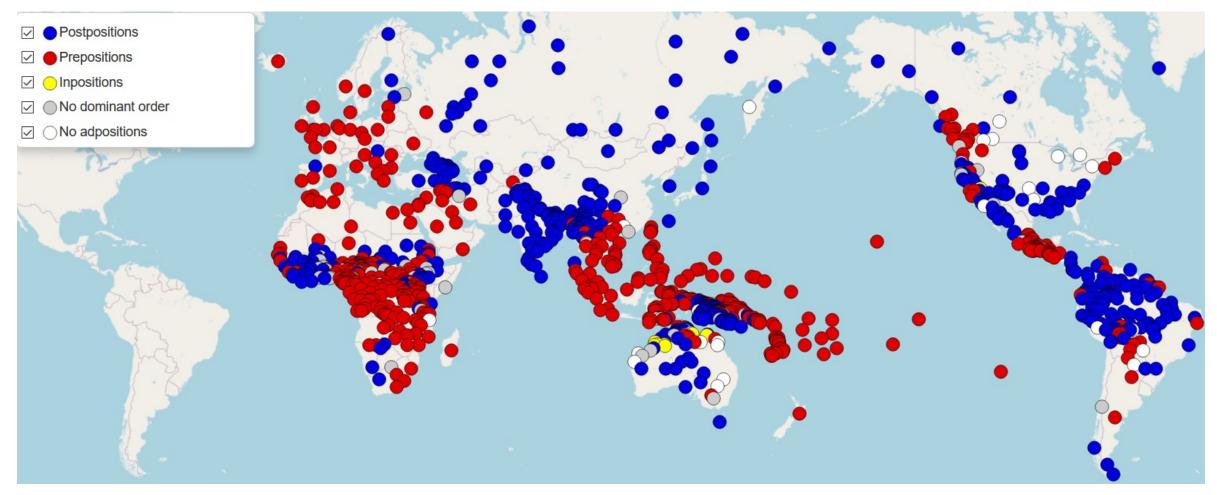


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Word-Order Traits in Languages





World Atlas of Languages (Dryer and Haspelmath, 2013)



Aim:

- Probing for universal correlation patterns in the evolution of wordorder traits
- Testing if cross-family models can capture correlation patterns not found in single-family models.

using a Logistic Brownian Motion Model

Word-Order Traits in Languages



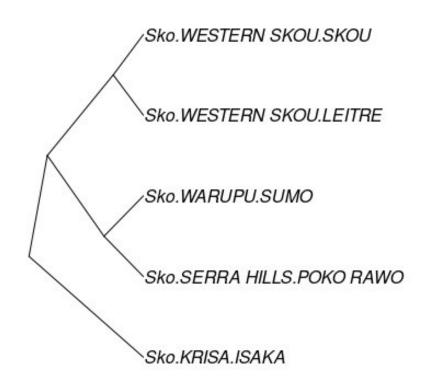
- Adjective-Noun
- Adposition-Noun
- Demonstrative-Noun
- Genitive-Noun
- Numeral-Noun
- Object-Verb
- Relative Clause-Noun
- Subject-Verb

28 binary trait pair combinations derived from WALS (Dryer and Haspelmath, 2013)

Language Families



- Evolutionary history in the form of phylogenetic trees
- 33 language families
- 768 languages in total
- Provided by Jäger (2018)





 $x \sim MultiNormal(a, V),$



 $x \sim Binomial(p)$

 $inv_logit(p) \sim MultiNormal(a, V),$

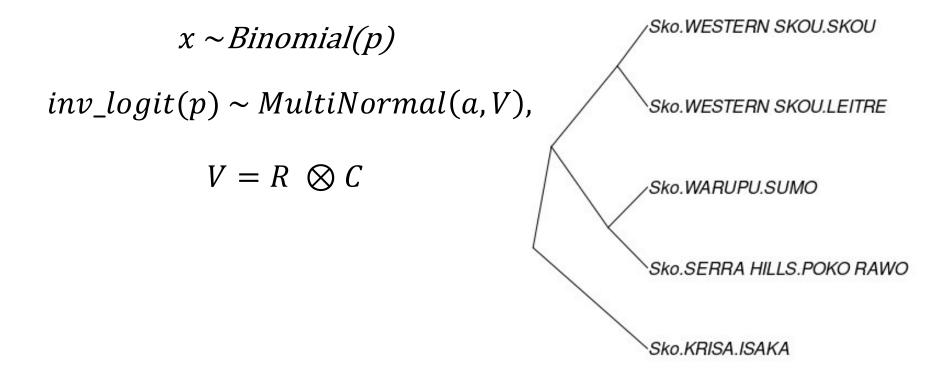
trait values x,

value probabilities p,

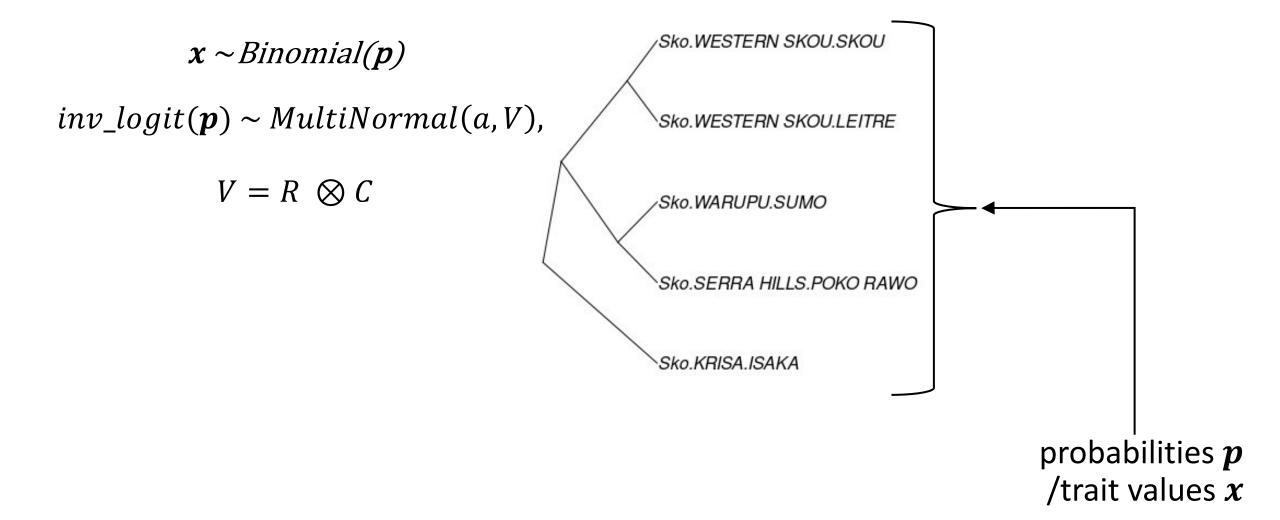
means a,

Variance-Covariance matrix $V = R \otimes C$

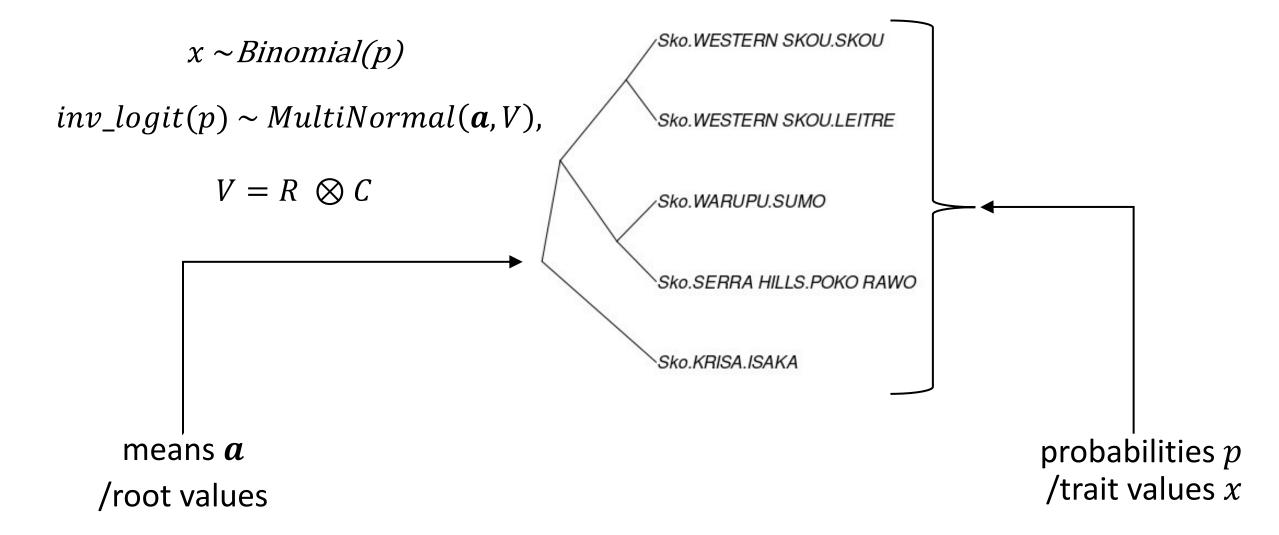




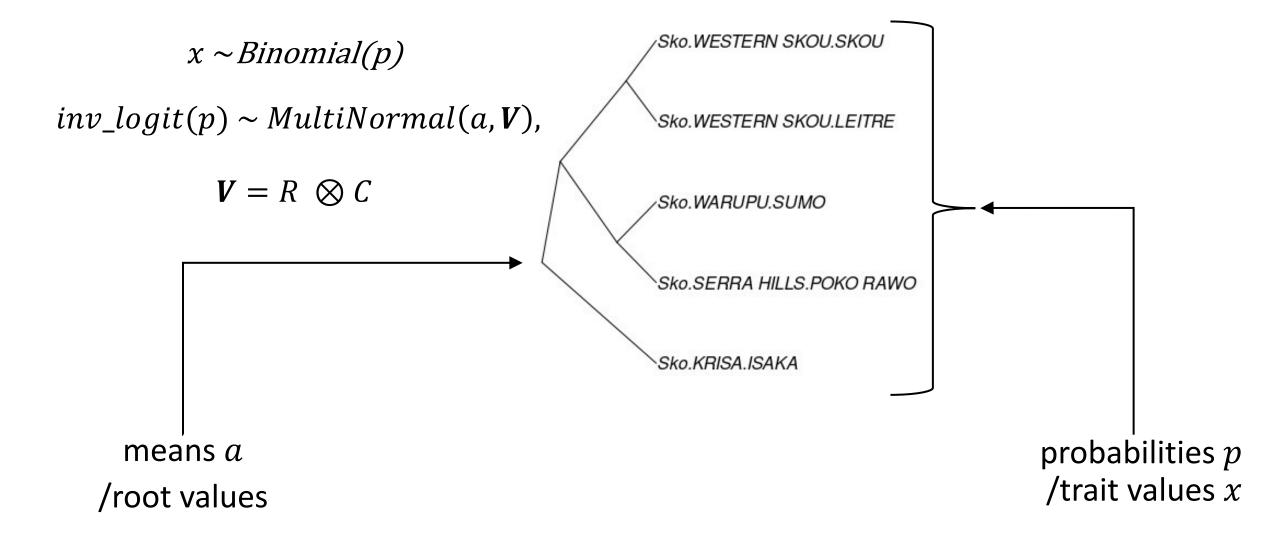




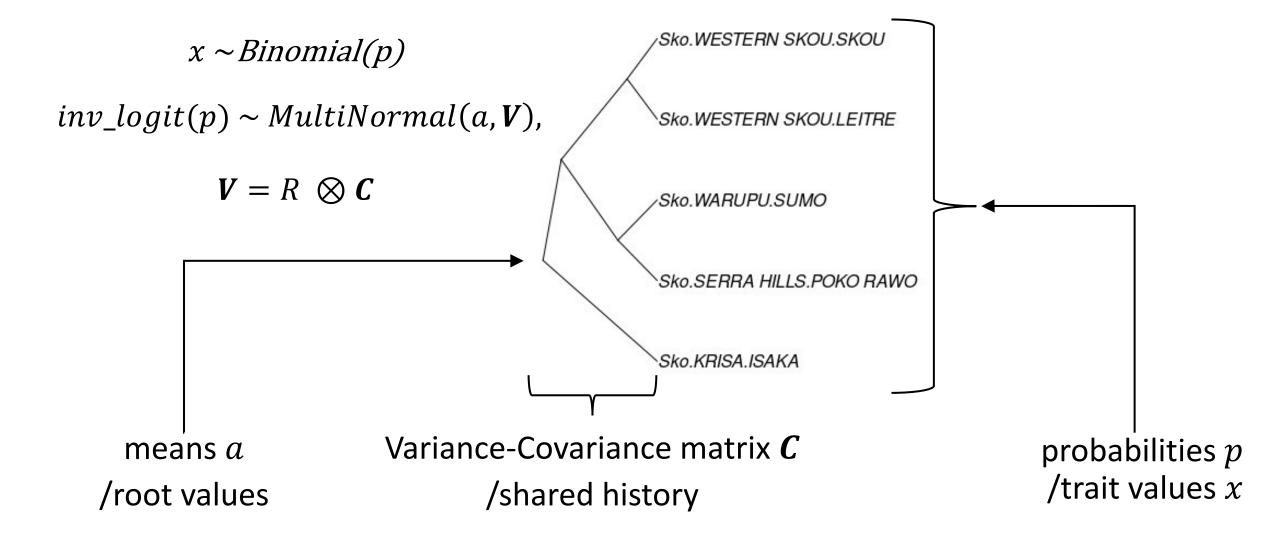
















$$x \sim Binomial(p)$$

 $inv_logit(p) \sim MultiNormal(a, V),$

$$V = R \otimes C$$

Trait correlation *R*

$$\mathbf{R} = \begin{bmatrix} \sigma^2_{11} & \sigma_{21} \\ \sigma_{12} & \sigma^2_{22} \end{bmatrix}$$





1. Models for each single family:

Correlated:

Independent:

$$R = \begin{bmatrix} \sigma^2_{11} & \sigma_{21} \\ \sigma_{12} & \sigma^2_{22} \end{bmatrix} \qquad R = \begin{bmatrix} \sigma^2_{11} & 0 \\ 0 & \sigma^2_{22} \end{bmatrix}$$

$$R = \begin{bmatrix} \sigma^2_{11} & 0 \\ 0 & \sigma^2_{22} \end{bmatrix}$$

Setup



1. Models for each single family:

Correlated:

Independent:

$$R = \begin{bmatrix} \sigma^2_{11} & \sigma_{21} \\ \sigma_{12} & \sigma^2_{22} \end{bmatrix}$$

$$R = \begin{bmatrix} \sigma^2_{11} & \sigma_{21} \\ \sigma_{12} & \sigma^2_{22} \end{bmatrix} \qquad R = \begin{bmatrix} \sigma^2_{11} & 0 \\ 0 & \sigma^2_{22} \end{bmatrix}$$

2. Models across all families:

Lineage-specific correlation:

$$R_f = \begin{bmatrix} \sigma^2_{11_f} & \sigma_{21_f} \\ \sigma_{12_f} & \sigma^2_{22_f} \end{bmatrix}, \quad R = \begin{bmatrix} \sigma^2_{11} & \sigma_{21} \\ \sigma_{12} & \sigma^2_{22} \end{bmatrix} \qquad R = \begin{bmatrix} \sigma^2_{11} & 0 \\ 0 & \sigma^2_{22} \end{bmatrix}$$
 for each family $f \in F$

$$R = \begin{bmatrix} \sigma^2_{11} & \sigma_{21} \\ \sigma_{12} & \sigma^2_{22} \end{bmatrix}$$

$$R = \begin{bmatrix} \sigma^2_{11} & 0 \\ 0 & \sigma^2_{22} \end{bmatrix}$$





- Single Family Models:
 - No trait pairs correlated consistently across language families
 - Observation consistent for Bayes Factors, WAIC, LOOIC





- Single Family Models:
 - No trait pairs correlated consistently across language families
 - Observation consistent for Bayes Factors, WAIC, LOOIC
- Cross-Family Models:
 - Bayes Factors:
 - Lineage-specific correlations valued much higher than universal models
 - Information Criteria:
 - Contrarily, universal models valued much higher than lineage-specific



Conclusions

- Single-Family models and Bayes Factors for Universal models are in favour of only lineage-specific correlations
- However, Information Criteria for Universal models are in favour of universal correlations
- => No clear evidence in favour of any universal trait correlations





Matthew S. Dryer and Martin Haspelmath, editors. 2013. WALS Online. Max Planck Institute for Evolutionary Anthropology, Leipzig.

Gerhard Jäger. 2018. A bayesian test of the lineagespecificity of word order correlations. In 12th International Conference on Language Evolution (Evolang XII), Torun