

Complexity clines out of Africa

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joint work with Christian Fröhlich

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WORDS BONES GENES TOOLS
Tracking Linguistic, Cultural, and Biological Trajectories of the Human Past

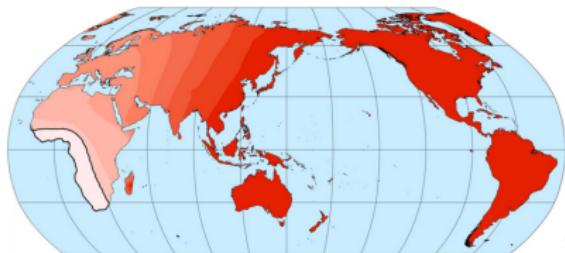
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Atkinson's observation

- Quentin Atkinson 2011
(Science, Phonemic diversity supports a serial founder effect model of language expansion from Africa):

- phonemic diversity of languages decreases with distance from Africa
- consistent with a **serial founder effect**
- evidence for origin of modern humans from Africa



(Atkinson, 2011, Figure S7)

- correlations are only informative about real-world processes if there is a plausible causal story
- Atkinson:

The number of phonemes—perceptually distinct units of sound that differentiate words—in a language is positively correlated with the size of its speaker population (*I*) in such a way that small populations have fewer phonemes. Languages continually gain and lose phonemes because of stochastic processes (2, 3). If phoneme distinctions are more likely to be lost in small founder populations, then a succession of founder events during range expansion should progressively reduce phonemic diversity with increasing distance from the point of origin, paralleling the serial founder effect observed in population genetics (4–9). A founder effect has already been

- causal mechanism rests on assumed correlation of inventory size with population size
- latter assumption debunked by Moran et al. (2012)

Looking at the data

- Phoible (Moran and McCloy, 2019)
 - phoneme inventories of 2,186 doculects
- ASJP (Wichmann et al., 2018)
 - 40-item Swadesh lists of 7,655 doculects
- intersection (linked via ISO code; only one doculect per ISO):
1,455 languages

Inventory Scots (Northern) (EA 2505)²

Segment list IPK chart

Showing 1 to 37 of 37 entries

Segment class	Segment	Marginal	Allphones	Representation
-any-		Search	~ants~	Search
consonant	m	False		2914 (96%)
vowel	i	False		2779 (92%)
consonant	j	False		2716 (90%)
vowel	u	False		2646 (89%)
vowel	a	False		2600 (86%)
consonant	w	False		2483 (82%)
consonant	n	False		2349 (78%)
consonant	s	False		2020 (67%)
consonant	b	False		1906 (63%)
consonant	ŋ	False		1897 (63%)
vowel	e	False		1841 (61%)
vowel	o	False		1826 (60%)
consonant	g	False		1712 (57%)
consonant	h	False		1703 (56%)
consonant	d	False		1376 (46%)
consonant	f	False		1329 (44%)
consonant	ʃ	False		1218 (40%)
vowel	c	False		1129 (37%)
consonant	ʒ	False		1104 (37%)
vowel	ɔ	False		1070 (36%)
consonant	z	False		893 (30%)
consonant	v	False		816 (27%)
consonant	r	False		774 (26%)
consonant	ʍ	False		605 (20%)
consonant	ɒ	False		592 (20%)
consonant	x	False		576 (19%)
consonant	ɛ	False		478 (16%)
vowel	ɪ	False		444 (15%)
consonant	p	False		403 (13%)
vowel	ʌ	False		133 (4%)
consonant	ə	False		129 (4%)
consonant	M	False		40 (1%)
consonant	t	False		31 (1%)
vowel	ai	False		19 (1%)
vowel	au	False		17 (1%)
vowel	æ	False		8 (0%)
vowel	œ	False		3 (0%)

Showing 1 to 37 of 37 entries

← Previous Next →



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Wordlist Scots

Compiled by André Müller and Helen Geyer
Showing 1 to 50 of 50 entries

No.	Meaning	Ward	Loan
1	a		False
1	an		False
2	you	ȝu	False
2	you	ȝu	False
3	we	ȝe	False
3	we	ȝe	False
11	one	an	False
11	one	e	False
12	her	her-e	False
16	parrot	bodl	False
18	falk	tȝ	False
21	dog	dg	False
21	dog	dug	False
22	knee	lus	False
22	knee	pul	False
23	tree	tr	False
26	leaf	ȝif	False
27	bird	heid	False
28	star	aleo	False
30	blood	blid	False
31	bone	ben	False
34	horn	horn	False
36	ear	ȝir	False
40	eye	i	False
40	eye	ȝldr	False
41	nose	nas	False
41	nose	rhȝ	False
43	tooth	teil	False
43	tooth	tei	False
44	fork	ȝyN	False
47	knob	ri	False
47	knob	ri	False
48	hand	hand	False
48	hand	han	False
51	breast	bretst	False
51	breast	breast	False
54	drink	emh-	False
54	drink	ȝdu	False
57	sun	ȝir	False
58	heat	he	False
61	die	di	False
66	come	kuem	False
66	come	kem	False
70	sun	ȝon	False
70	sun	ȝin	False
74	star	starn	False
74	star	bȝdN-ȝr	False
75	water	watrh	False
77	stone	sten	False
82	fire	ȝis	False
82	fire	is	False
85	path	ped	False
85	path	pȝ	False
86	moorland	ȝen	False
86	moorland	men	False
87	right	ri	False
88	full	lu	False
89	new	ynnu	False
90	name	nam	False

Showing 1 to 50 of 50 entries



Coordinates: WGS84
56.162° N, 4.018° W
55.96° - 56.26° N
0° - 4.26° E

number of speakers: 98.205

status: alive

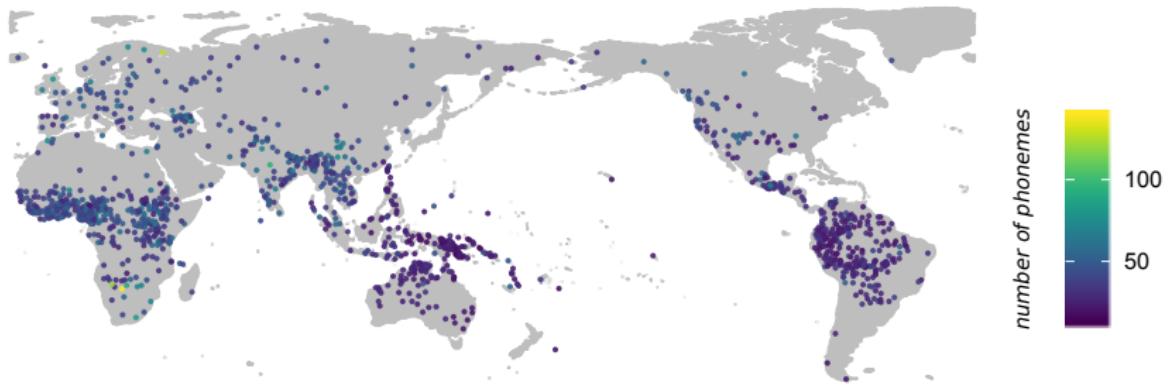
Classification

WALS
L1: Germanic
Glossing:
Indo-European > Germanic > Northwestgermanic > Westgermanic > Northgermanic >
Anglo-Frisian > English
Ethnologue
Indo-European > Germanic > West > English

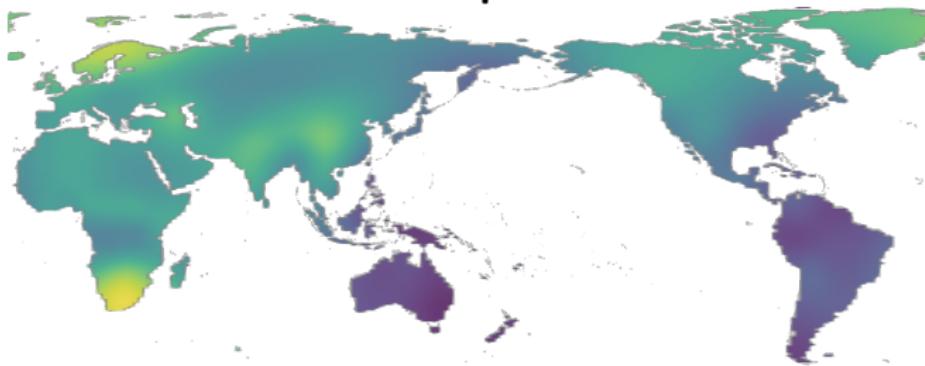
Sources

n.a.n.d.
<http://www.scots-online.org/dictionary/engscots.asp>

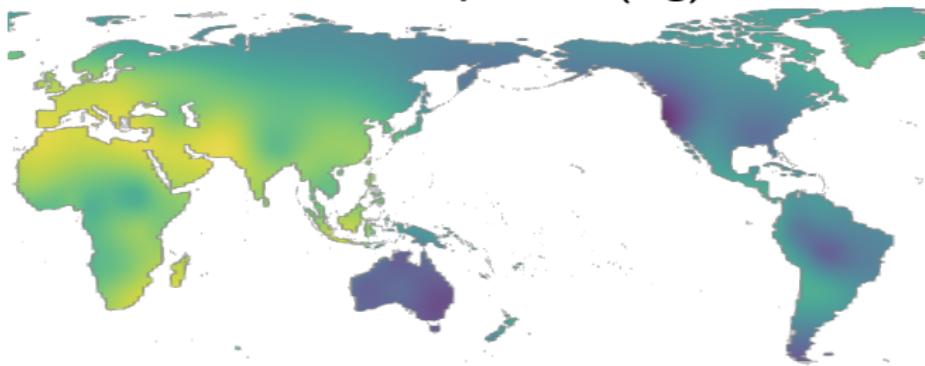




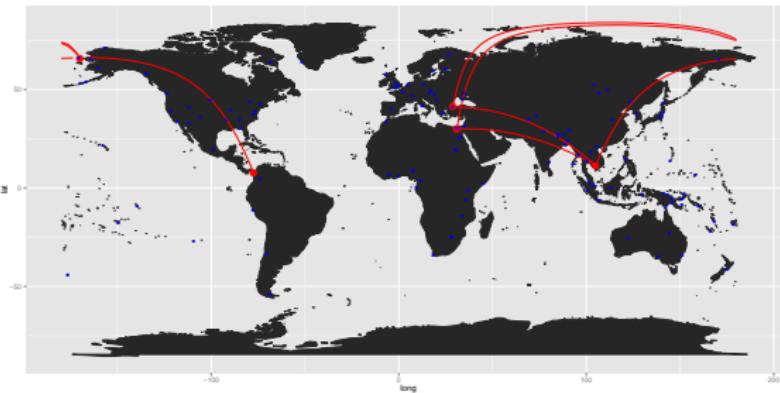
number of phonemes



number of speakers (log)

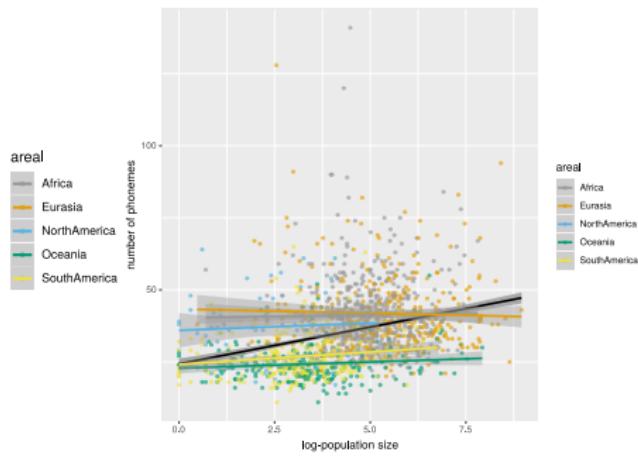
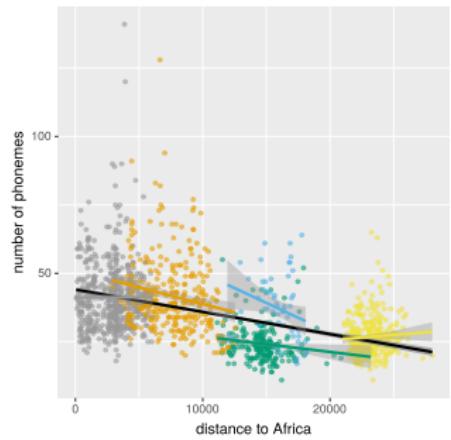


Land-based distances



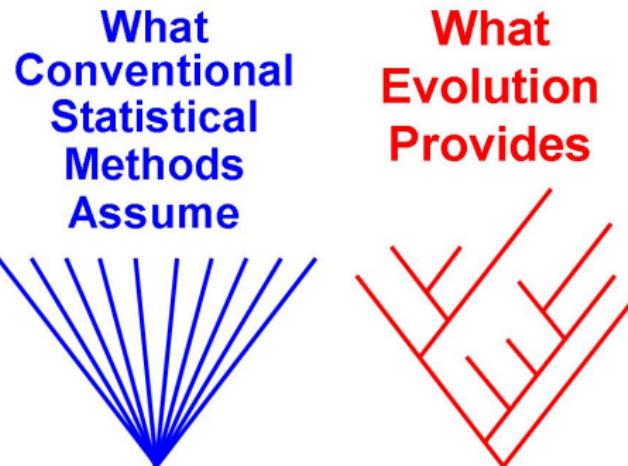
- following Atkinson 2011:
 - Africa/Asia: *Cairo*
 - Asia/Europe: *Istanbul*
 - Asia/Oceania: *Phnom Phen*
 - Asia/North America: *Bering Strait*
 - North America/South America: *Panama*
- following Wichmann et al. (2011), I use Addis Ababa as putative place of origin here





Controlling for phylogenetic non-independence

Spurious correlations in cross-linguistic data



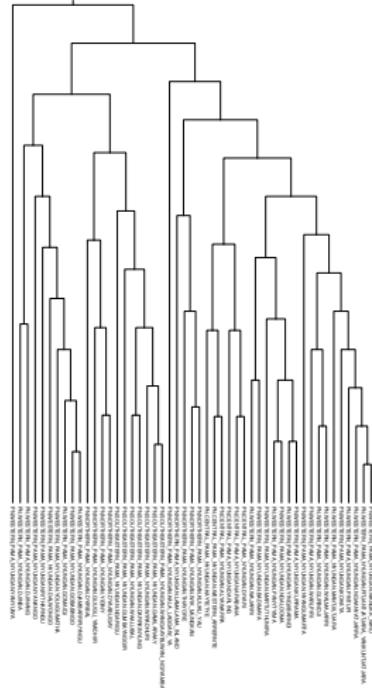
Copyright Theodore Garland, Jr. Original creation Modified from Fig. 3 of Garland, T., Jr., & P. A. Carter. 1994. Evolutionary physiology. Annu. Rev. Physiol. 55:579-621.

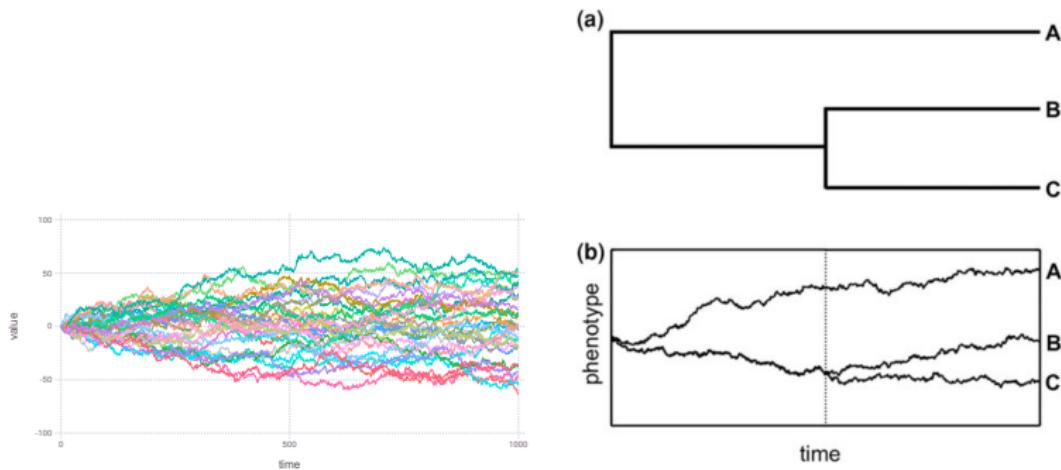
Treating different languages as independent samples creates many spurious correlations!

- mixed-effects models, with genealogical units and/or linguistic macro-areas as random factors, mitigate this problem somewhat
- genealogical non-independence is no binary distinction though; Scots is related to Hindi, but it much closer related to English
- degree of relatedness can be estimated via **phylogenetic inference** from lexical data

The phylogenetic comparative method

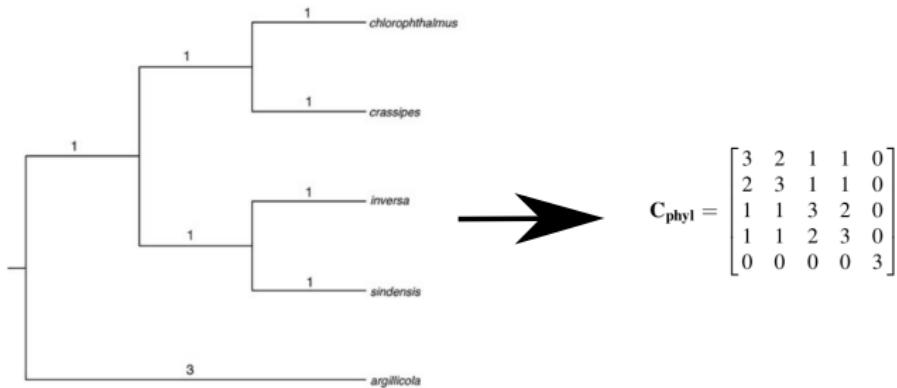
- data-generating process is governed by a **phylogenetic tree**
- simplest model for continuous variables:
 - variable evolves according to **Brownian motion** from root to leaves
 - when branch splits, to independent copies keep evolving randomly



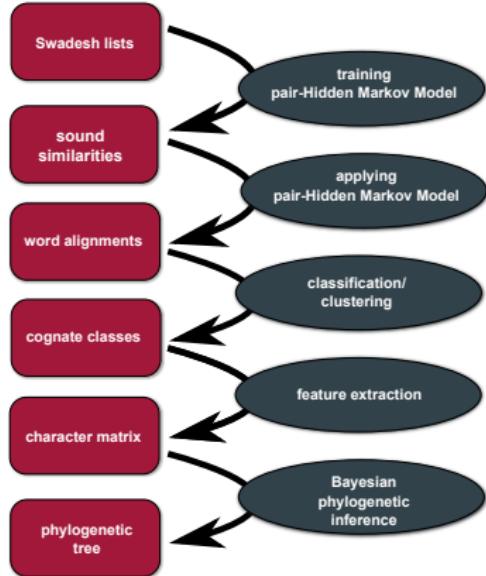


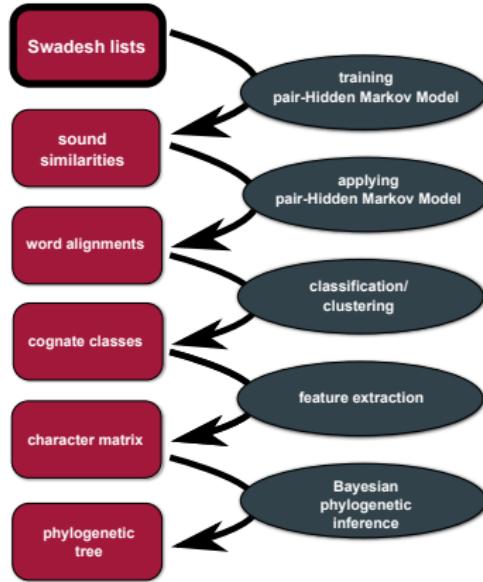
- covariance between two leaves \propto time traveled together

- measurements at the leaves are distributed according to multivariate normal distribution
- variance-covariance matrix is determined by tree

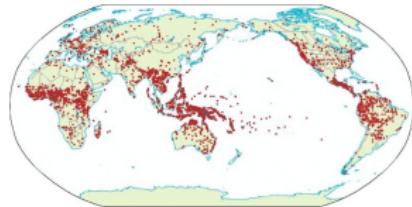


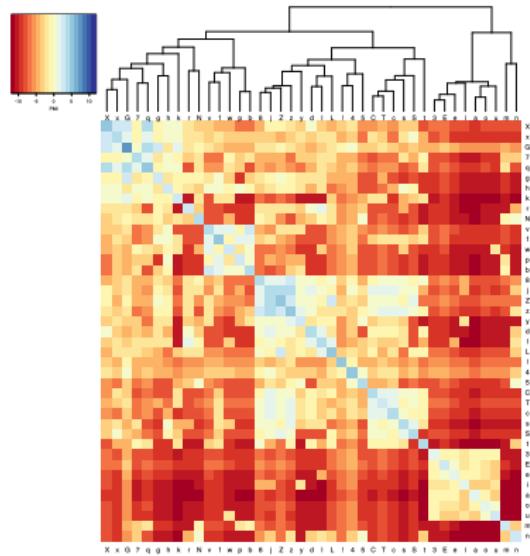
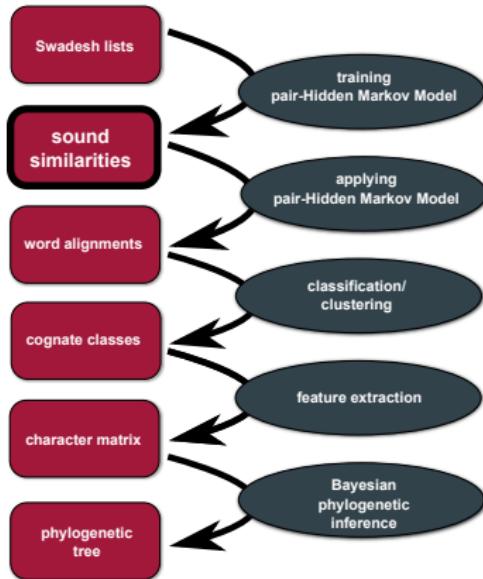
Inferring a language tree

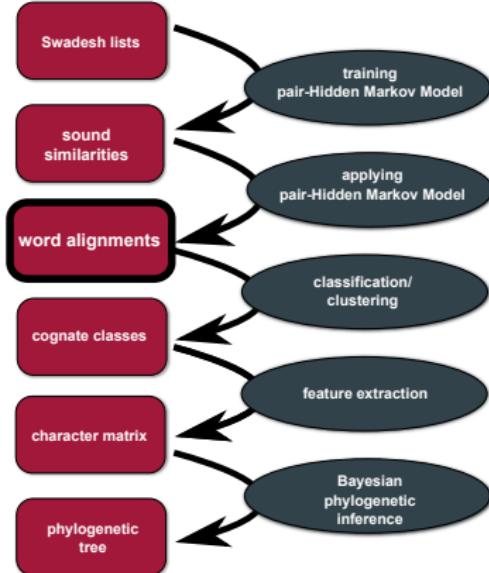




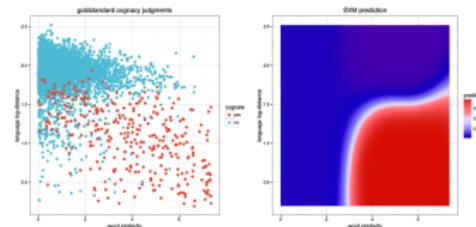
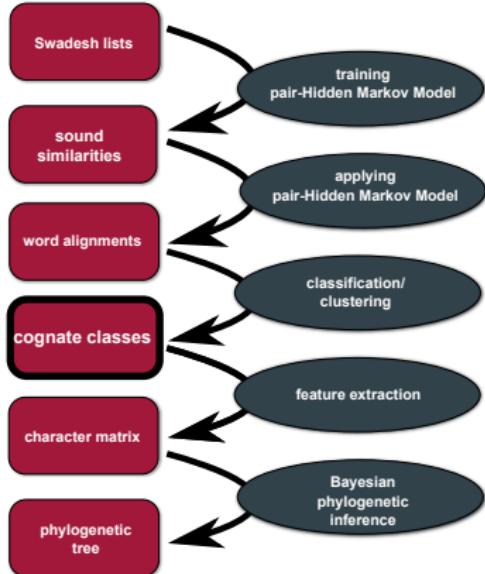
concept	Latin	English
<i>I</i>	ego	Ei
<i>you</i>	tu	yu
<i>we</i>	nos	wi
<i>one</i>	unus	w3n
<i>two</i>	duo	tu
<i>person</i>	persona, homo	pers3n
<i>fish</i>	piskis	fiS
<i>dog</i>	kanis	dag
<i>louse</i>	pedikulus	laus
<i>tree</i>	arbor	tri
<i>leaf</i>	foly~u*	lif
<i>skin</i>	kutis	skin
<i>blood</i>	saNgw~is	bl3d
<i>bone</i>	os	bon
<i>horn</i>	kornu	horn
<i>ear</i>	auris	ir
<i>eye</i>	okulus	Ei



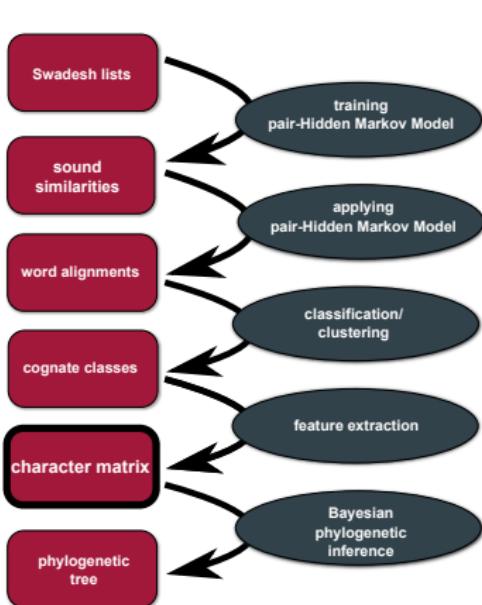




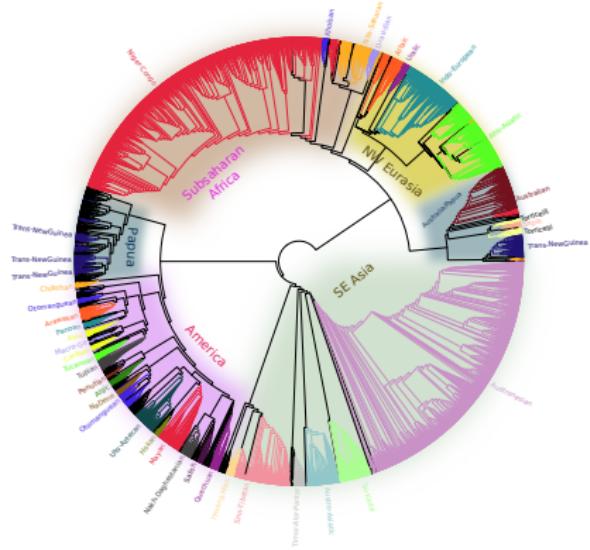
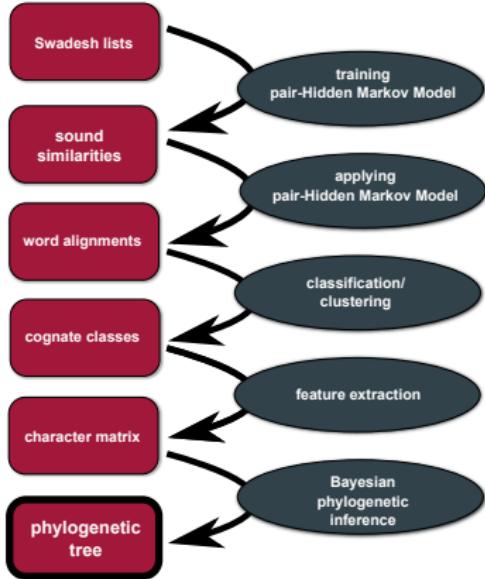
<i>Language</i>	<i>fish:z</i>	<i>tongue:1</i>	<i>smoke:1</i>
Abui-Atangmelang	-af <u>u</u>	tal-i-fii--	awai-b-a-n-o-7o-
Abui-Fuimelang	-af <u>u</u>	tal-E-b---	--ad-b-a-n-a-nka-
Adang	aab--	--j-e-bur-	-----b-e-n-a-xa-
Blagar-Bakalang	-ab--	teg-e-bur-	-----b-e-n-a-nka-
Blagar-Bama	aab--	tej-e-bur-	-----b-e-n-a-nka-
Blagar-Kulijahi	-ab--	tej-e-bur-	-----b-e-n-a-nka-
Blagar-Nule	aab--	tej-e-bur-	--ad-b-e-n-a-nka-
Blagar-Tuntuli	aab--	tel-e-bur-	a-adgeb-a-n-a-q--
Blagar-Warsalelang	-ab--	tel-e-bur-	a-ad-b-a-n-a-x--
Bunaq			-----b-o-t-o-h--
Deing	haf--		-----buu-n-
Hamap	7ab--	nar-g u N	-----b-a-n-o-7--
Kabola	hab--	tal-e-b---	aval-b-e-n-e-7o-
Kaera-Padangsul	-ab--	talee-b---	a-ad-b-e-naa-x--
Kafoa	-afUi	tal-i-p---	-----f-o-n-a---
Kamang	-ap-i	nal---pu--	-----p-u-n---a-
Kiraman	-Eb--	nal-i-bar-	--ar-b-a-n-o-kan
Klon	-eb-i	gel-E-b---	--ed-ab-o-n-----
Kui	-eb--	tal-i-ber-	--ar-b-o-n-o-k--
Kula	-ap-i	-il-I-p---	-----p---n-ekka-
Nedebang	aaf-i	gel-e-fu--	--ar-ab-u-n-----
Reta	aab--	nal-e-bul-	a-ad-b-o-n-a----
Sar-Adiabang	haf--	--p-e-fal-	--ar-buu-n-----
Sar-Nule	haf--	nal-e-faj-	-----faj-n-----
Sawila	-ap-i	gal-impuru	-----p-u-n-a-ka-
Tewa-Madar	xaf--	gel-i-vi--	-----buu-n-----
Wersing	-ap-i	nej-e-bur-	--ad-ap-u-n-a-k-
Wpantar	hap--	nal-e-bu--	-----b-unna-----



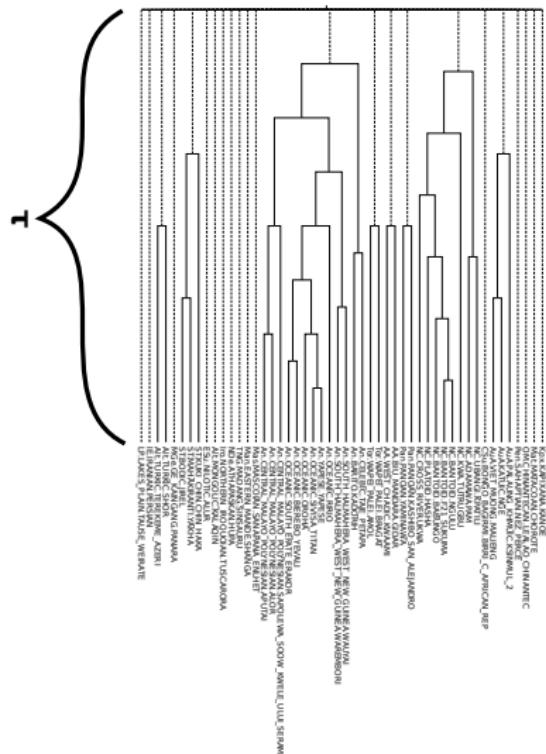
	English	Spanish	Modern Greek	Standard German
I	Ei:A	yo:B	exo:C	iX:D
you	yu:A	usted:B, tu:C	esi:D	du:E
we	wi:A	nosotros:B	enis:C	vir:Ä
one	w3n:A	uno:B	enas:C, ena:C	ains:D
two	tu:A	dos:B	By~o:C, Si:o:D	cva:i:E
person	pers3n:A	persona:A	anB~ropos:B	nBn3l:C
fish	fis:A	pezakdo:A, pes:A	parsi:B	fis:A
dog	dag:A	pero:B	sTili:C, sTiloe:C	hund:D
come	kdm:A	venir:B	erx~o:C	kh~om3n:A
sun	s3n:A	sol:B	ily~oe:C, iIos:C	zon3:A
star	star:A	estreyaa:A	asteri:A, astre:A	StErn:A
water	wat3r:A	agu~a:B	nero:C	vas3r:A
stone	ston:A	piedra:B	petra:B	Stein:A
fire	f3ir:A	fuego:B	foty~a:C	foia:D
path	pE8:A	senda:B	8ronos:C	pf~at:A, vek:D
mountain	m3unt3n:A	sero:B, montaña:A	vuno:C, oros:D	bErk:E
full	ful:A	yeno:B	yematos:C, pliria:D	fol:A
new	nu:A	nuevo:A	neos:A, Tenury~os:B	nei:A
name	nen:A	nombre:A	onoma:A	nam3:A



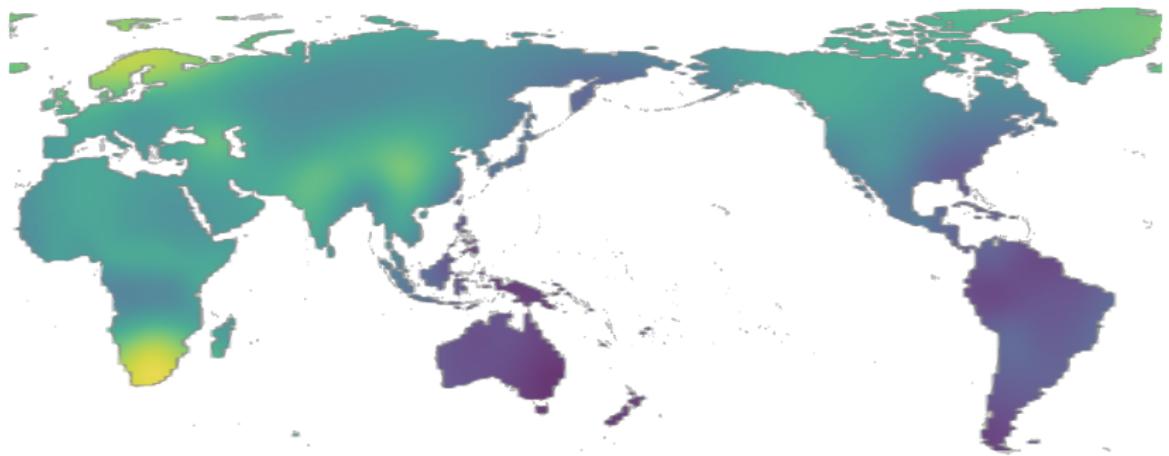
TNG, ENGAN, MAIBI
TNG, ENGAN, POLE
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TNG, ENGAN, YARIBA
TNG, FASU, FASU
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TNG, FINISTERRE, HUON, AWARA
TNG, FINISTERRE, HUON, BORONG
TNG, FINISTERRE, HUON, BURUM
TNG, FINISTERRE, HUON, BURUM MIND
TNG, FINISTERRE, HUON, DEDUA
TNG, FINISTERRE, HUON, HUBE
TNG, FINISTERRE, HUON, KATE
TNG, FINISTERRE, HUON, KOMBA
TNG, FINISTERRE, HUON, KOSORONG
TNG, FINISTERRE, HUON, MAPE
TNG, FINISTERRE, HUON, MAPE_2
TNG, FINISTERRE, HUON, MIGABAC
TNG, FINISTERRE, HUON, MINDIK
TNG, FINISTERRE, HUON, MOOLILI
TNG, FINISTERRE, HUON, HARABOG
TNG, FINISTERRE, HUON, HANKINA
TNG, FINISTERRE, HUON, HEK
TNG, FINISTERRE, HUON, HUKNA
TNG, FINISTERRE, HUON, ONO
TNG, FINISTERRE, HUON, SELEPET
TNG, FINISTERRE, HUON, TIMBE
TNG, FINISTERRE, HUON, TOBO
TNG, FINISTERRE, HUON, WANTOAT
TNG, FINISTERRE, HUON, YOPNO
TNG, GOILALAN, AFQA
TNG, GOILALAN, KINIMAIAPA
TNG, GOILALAN, MAFULUJU

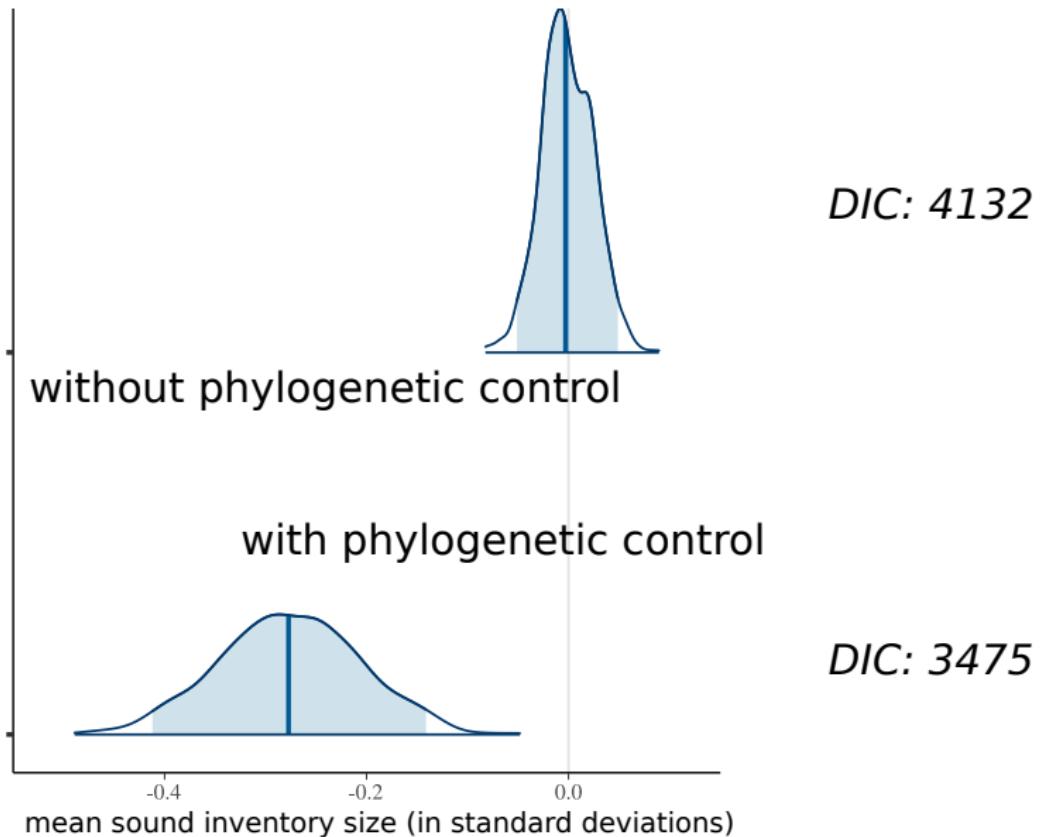


- tree estimation above established language families unreliable
 - very high phylogenetic uncertainty
 - branches tend to be much too short
- compromise used here:
 - infer trees for individual language families
 - connect them with a rake-shaped “proto-world” root with unknown depth
 - total tree depth τ is estimated from data
 - isolates are connected directly to the root

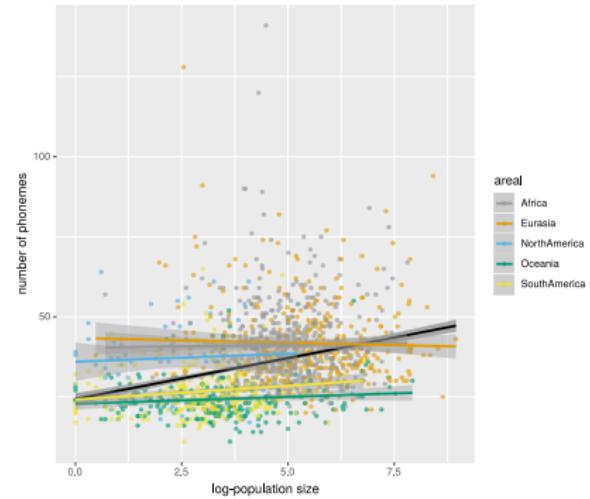
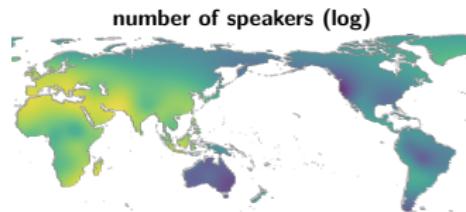
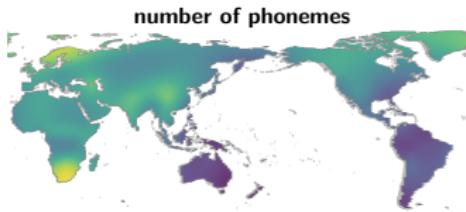


Null models for distribution of sound inventory size



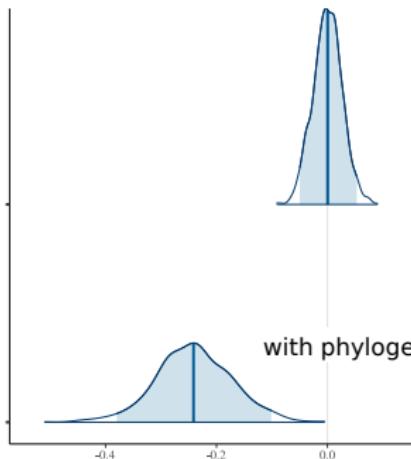


Using number of speakers as predictor



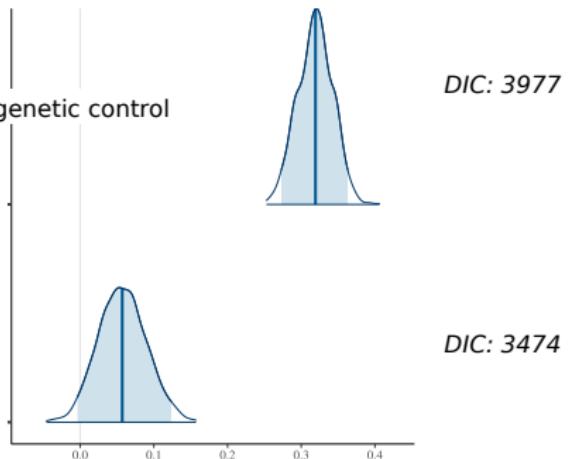
intercept

slope



without phylogenetic control

with phylogenetic control

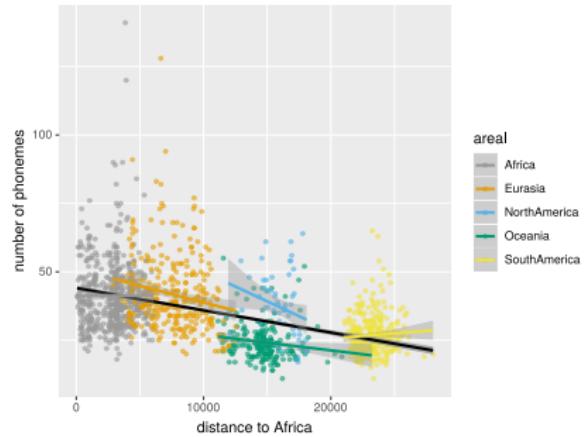
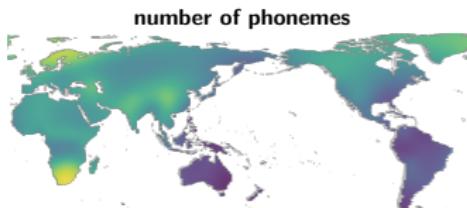


DIC: 3977

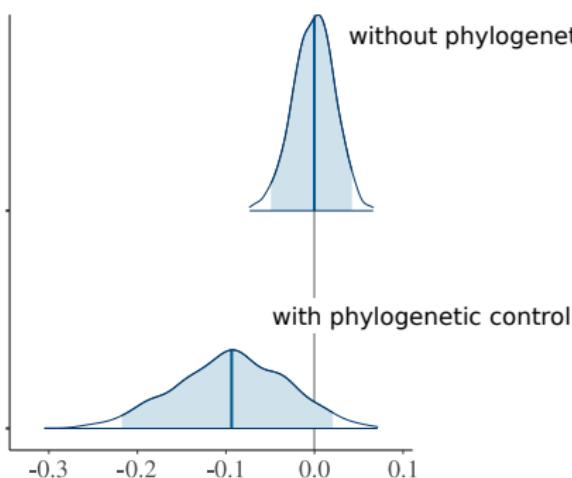
DIC: 3474

- with phylogenetic control:
 - posterior distribution for intercept includes 0 within 95% HPD (if barely)
 - model comparison shows model with and without predictor to be essentially tied

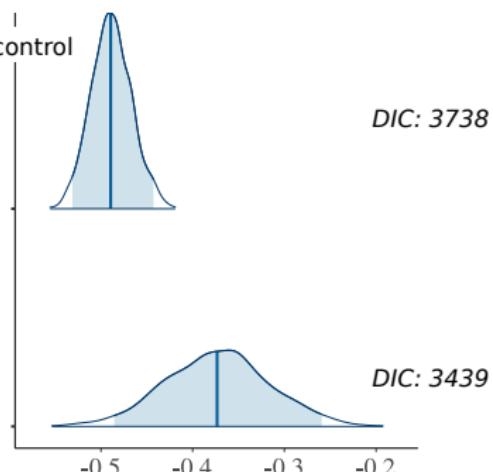
Using distance from Africa as predictor



intercept



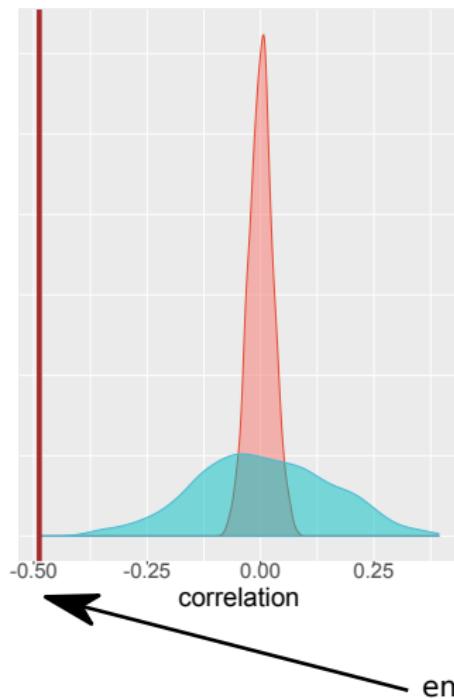
slope



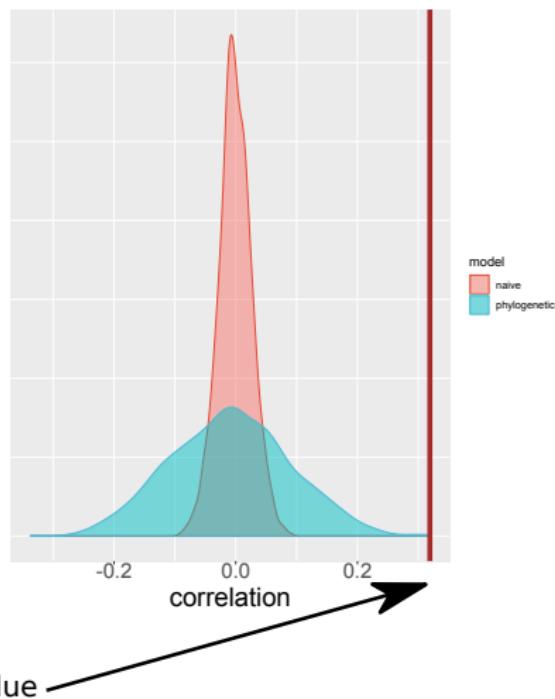
- with phylogenetic control:
 - both posterior distribution and model comparison indicate that distance to Africa is a relevant predictor for sound inventory size
 - with phylogenetic control, the effect becomes weaker, but it does not disappear

Posterior predictive check: correlations under null models

distance to Africa

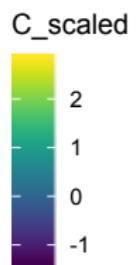
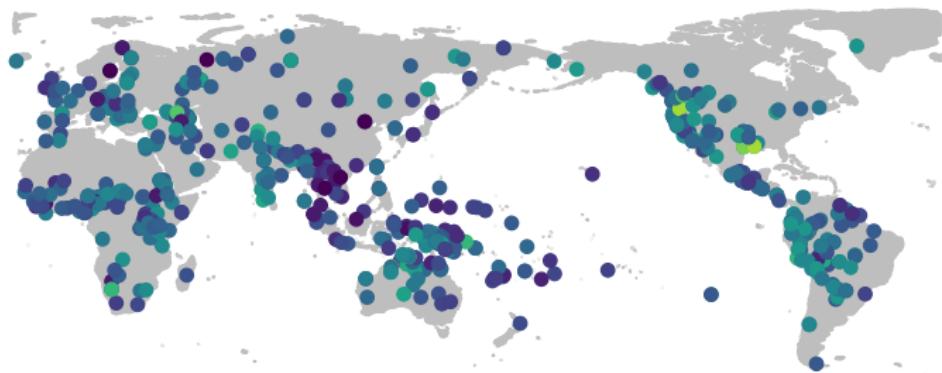
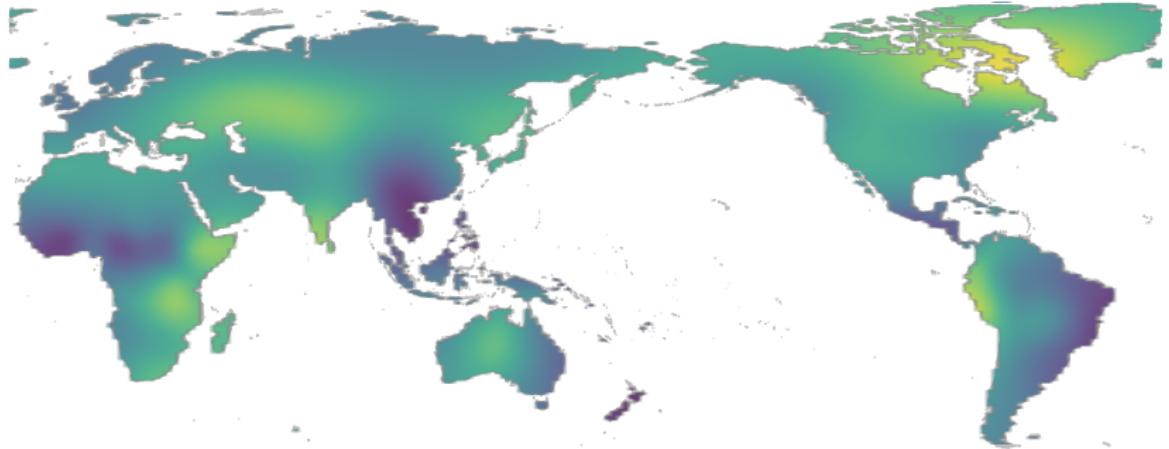


log-population size

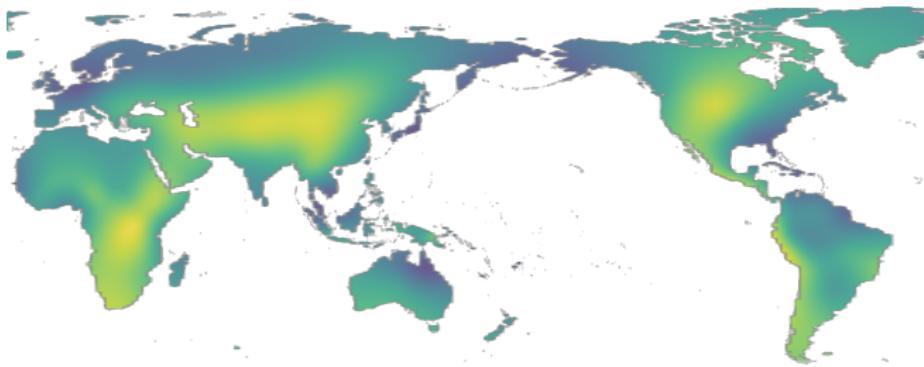


Morphological complexity

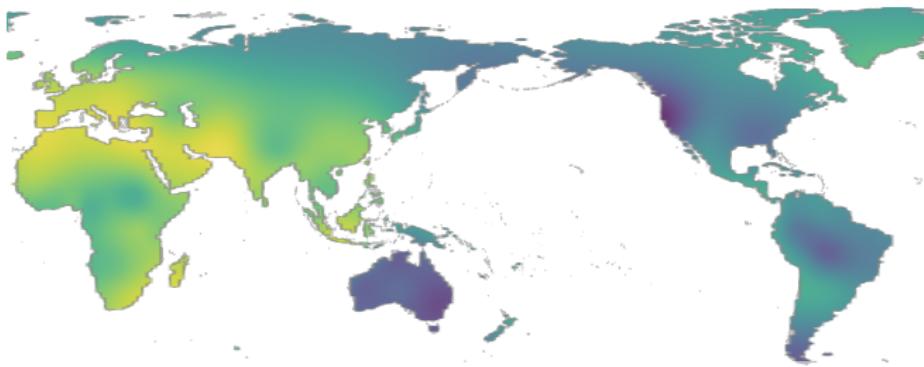
- dataset prepared by Christian Bentz and Johanna Nichols (Nichols and Bentz, 2018)
- numerical measure of morphological complexity
- aggregated from corpus unigram entropy and from number of features according to WALS (Dryer and Haspelmath, 2013) and Autotyp (Bickel et al., 2018)
- 1,060 languages in intersection with ASJP
- various hypotheses about non-linguistic factors influencing complexity, such as
 - distance to equator
 - altitude
 - population size



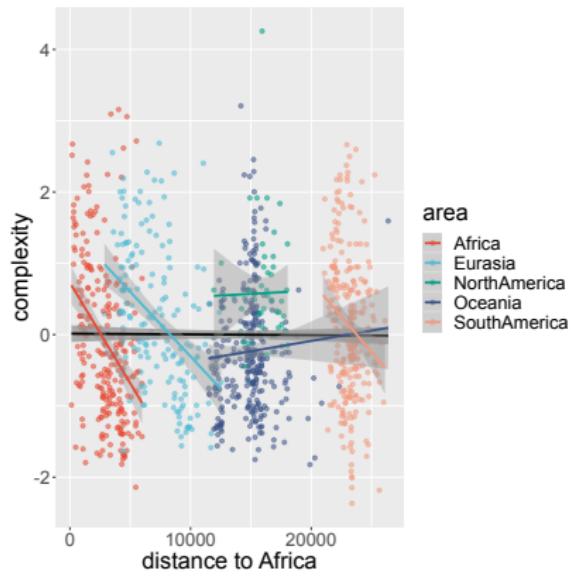
Altitude



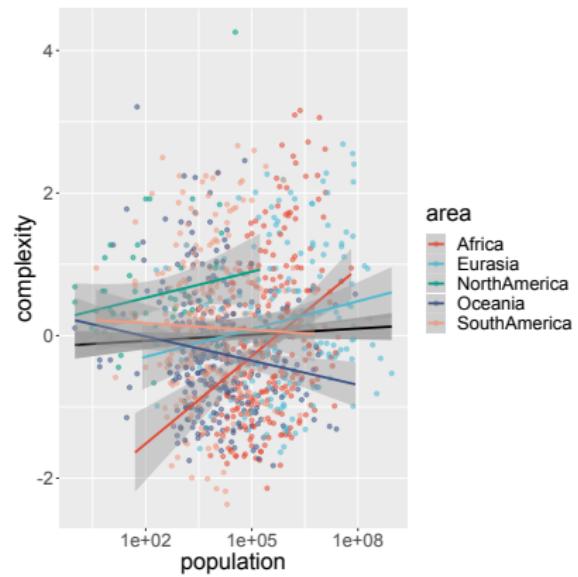
Population size



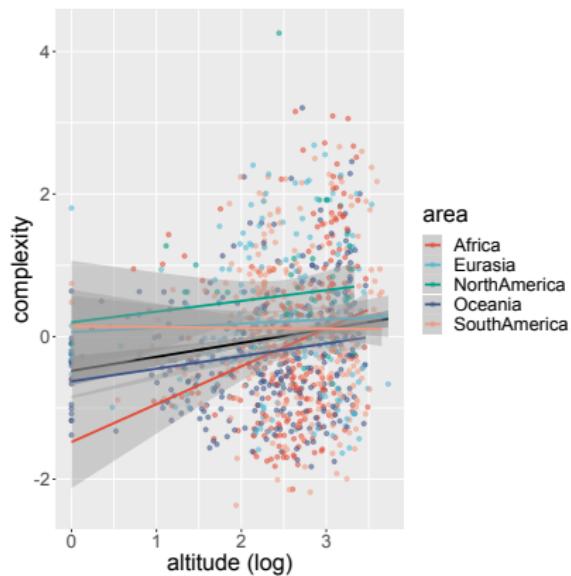
distance to Africa



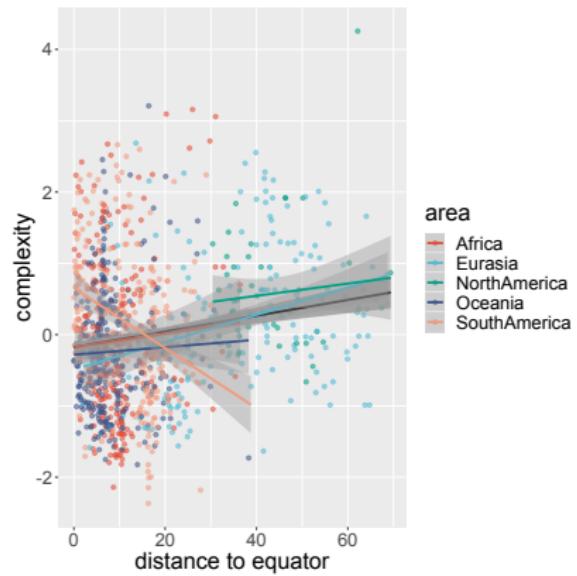
(log-)population size



(log-) altitude



distance to equator



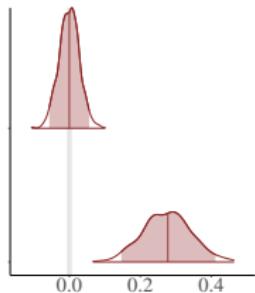
- I fitted two Bayesian regression models

$$\text{complexity} \sim \mathcal{N}(\text{intercept} + \text{distance to Africa} + \log(\text{population size}) + \log(\text{altitude}) + \text{distance from equator}, \Sigma)$$

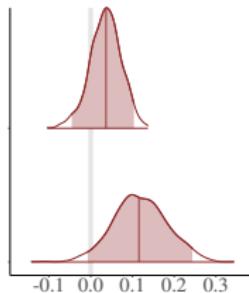
- model 1: Σ is diagonal matrix
- model 2: Σ is derived from phylogenetic trees

posterior estimates of regression parameters

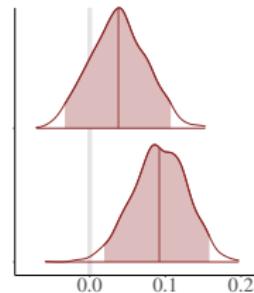
intercept



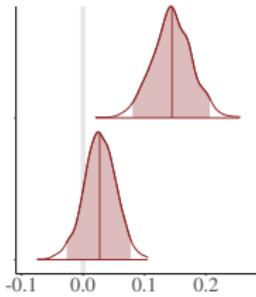
distance to Africa



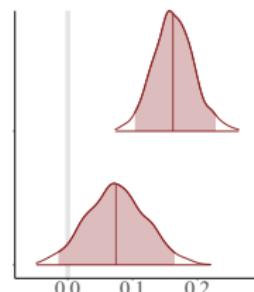
population size



altitude



distance from equator



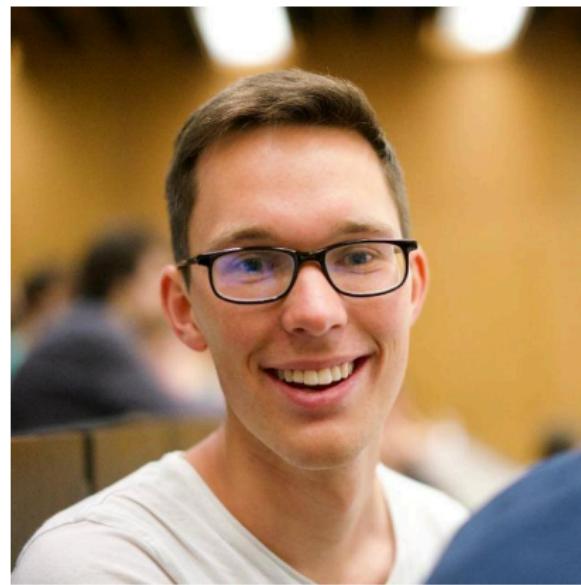
naive model (DIC: 2966)

phylogenetic model (DIC: 2494)

Conclusion

- controlling for common descent is essential when comparing populations on a global scale
- language trees, as derived via phylogenetic inference, are a suitable starting point for assessing autocorrelation
- resulting model arguably still underestimates autocorrelation since it ignores horizontal transmission
- first experiments with geographic spatial autocorrelation did not alter qualitative picture though

special thanks to Christian Fröhlich



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