

THE LEXICAL FREQUENCY OF LABIAL-VELAR STOPS AS A WINDOW ON THE LINGUISTIC PREHISTORY OF NORTHERN SUB-SAHARAN AFRICA

Dmitry Idiatov & Mark Van de Velde

LLACAN, CNRS, Sorbonne-Paris Cité, INALCO

dmitry.idiatov@cnrs.fr

mark.vandavelde@cnrs.fr



- Northern sub-Saharan Africa is obviously a spread zone with a **marked areal distribution** of various linguistic features
 - Macro-Sudan belt (Güldemann 2008)
 - Sudanic zone (Clements & Rialland 2008)
 - ...
- LV are **common in NSSA** languages but **typologically** they are known to be rather **rare** (e.g., Cahill 2008, Maddieson 2011)

Q₁: What can the areality of LV tell us about the history of the languages of NSSA and of the populations speaking these languages?

We **start** with the **observation** that:

- Languages with LV can **vary significantly** with respect to the **status of LV** in their phonologies and lexicons

We **proceed** by looking into the following **questions**:

- Are LV “normal” phonemes in the languages of NSSA in general?
- Are the distributions of LV within the lexicons random?
- What is the spatial pattern of the LV frequency distribution?

We **conclude** by providing an **interpretation** of our findings:

- What can our findings tell us about the spatio-temporal dynamics of the languages of NSSA and the populations speaking these languages?

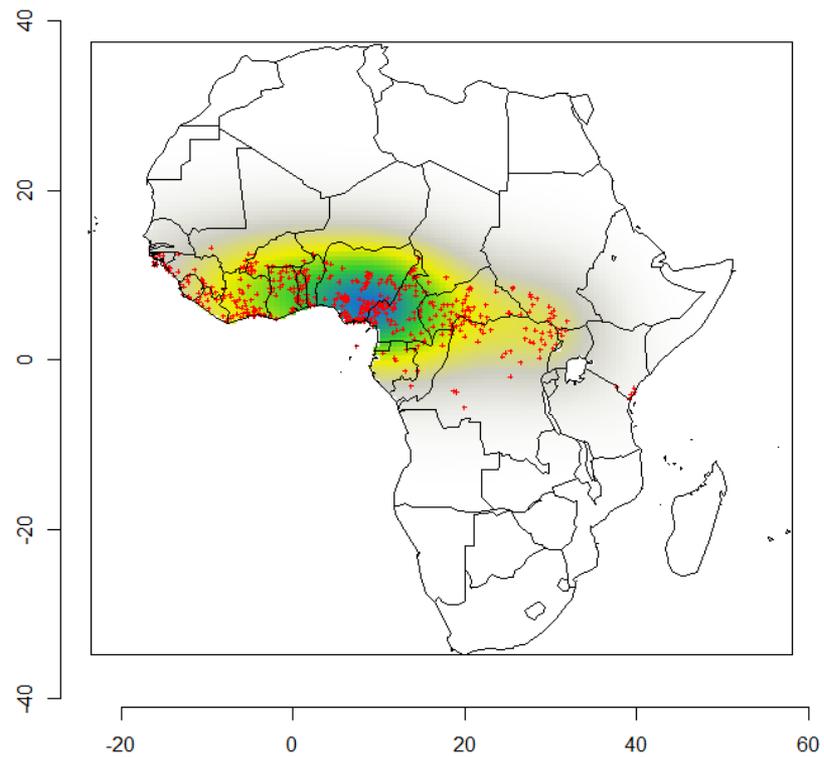
LV data sources:

- **RefLex**, www.reflex.cnrs.fr, LVFreq data
- Phoible, www.phoible.org, YN data
- Additional LVFreq data for some Mande and Bantu languages

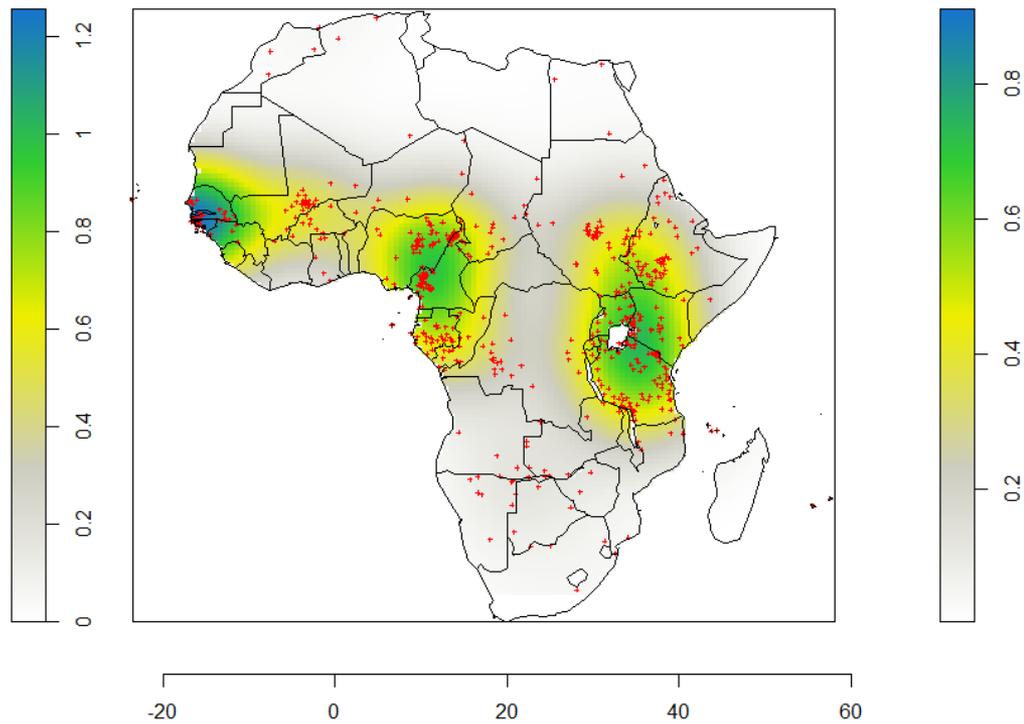
The composition of our sample (1304 languages):

- **336** languages with LV & LV frequency is known
- 230 languages with LV & LV frequency is not known
- 738 languages without LV

LVall_Y languages: geographic distribution



LVall_N languages: geographic distribution



LVFreq estimation

H_0 : In a lexicon, all C phonemes have equal frequency (have equal probability of occurrence)

$$LVFreq = \frac{LV_O}{LV_E} * 100\% = \frac{\sum T_{LV}}{\frac{\sum T_C}{\sum P_C} * \sum P_{LV}} * 100\%$$

LV_O - observed LV count

LV_E - expected LV count

T_{LV} - LV token

T_C - any C token

P_{LV} - LV phoneme

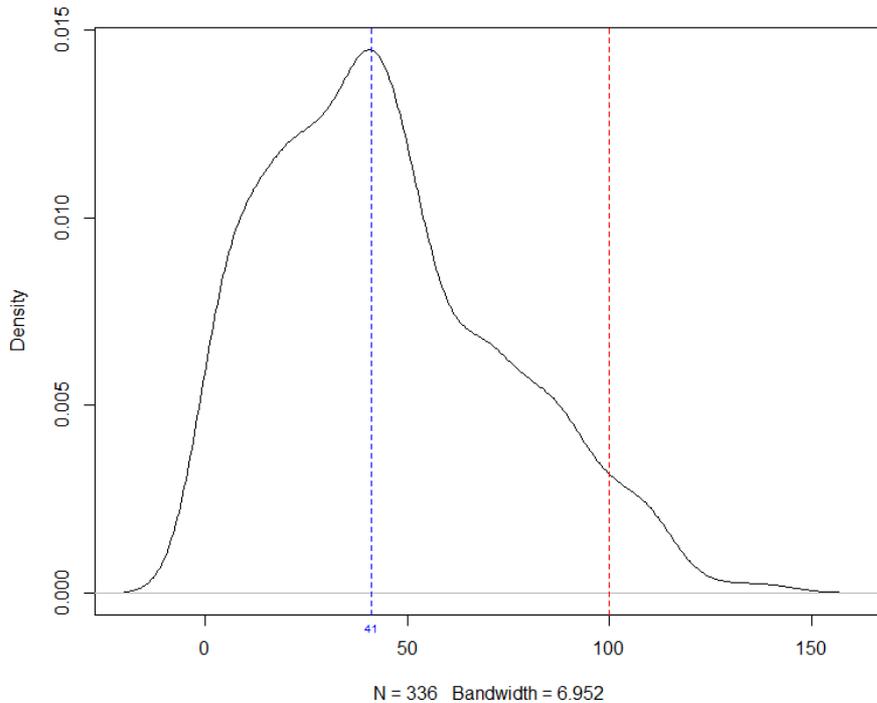
P_C - any C phoneme

LVFreq estimation

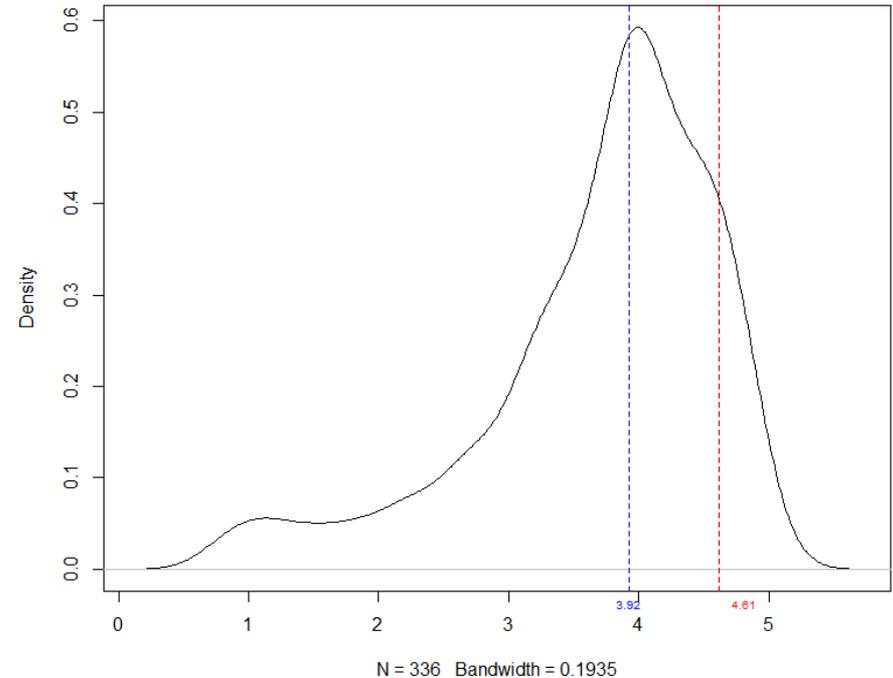
LVFreq = **0%** no LV

LVFreq = **100%** “reference LVFreq” - LV are “normal” phonemes, i.e. the observed number of occurrences of LV is the same as would be expected under the H_0

Non-zero LVFreq probability density



Log-transformed non-zero LVFreq probability density (scaled)



--- median

--- reference LVFreq

- LV are relatively **rare phonemes** in most languages that have them, which is in accordance with their typological rarity



Are the distributions of LV within the lexicons random?

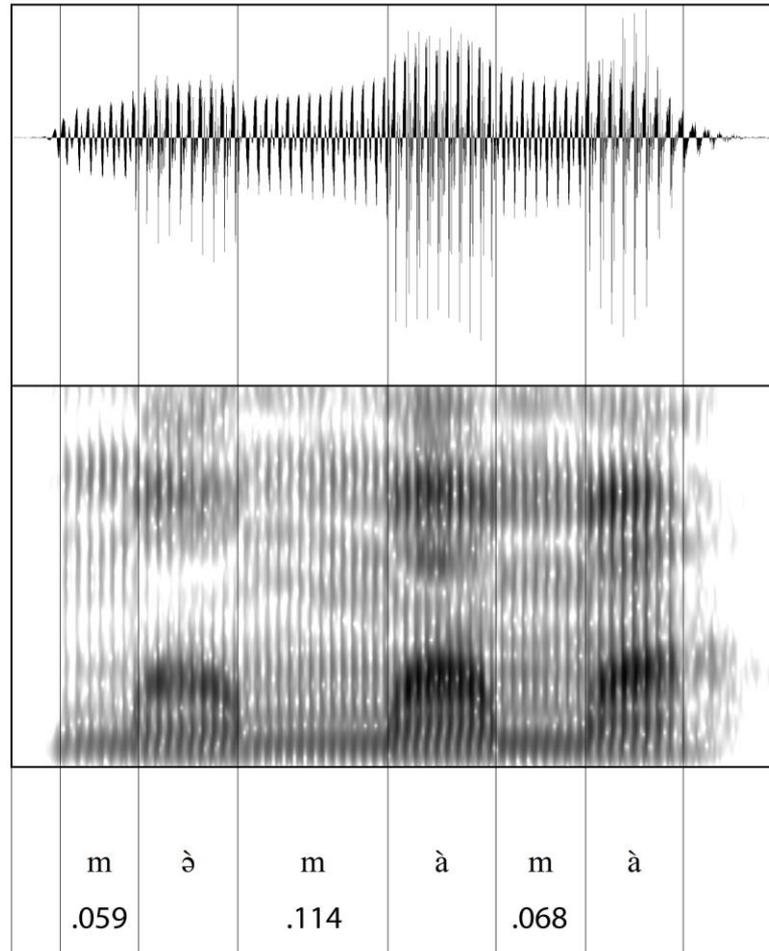
- LV tend to be less common in “basic vocabulary”
- **{H}**: LV are more common in the “**expressive**” parts of the **lexicon**, such as ideophones or property words, rather than referring expressions, such as nouns and verbs
- LV are largely restricted to the **stem-initial position**



- The correlation [LV ~ “expressive” vocabulary] is not independent of the correlation [LV ~ stem-initial position]
- **SI C-accent** (as a manifestation of a more general phenomenon of **C-emphasis prosody**) is a very important factor behind the emergence of LV in NSSA

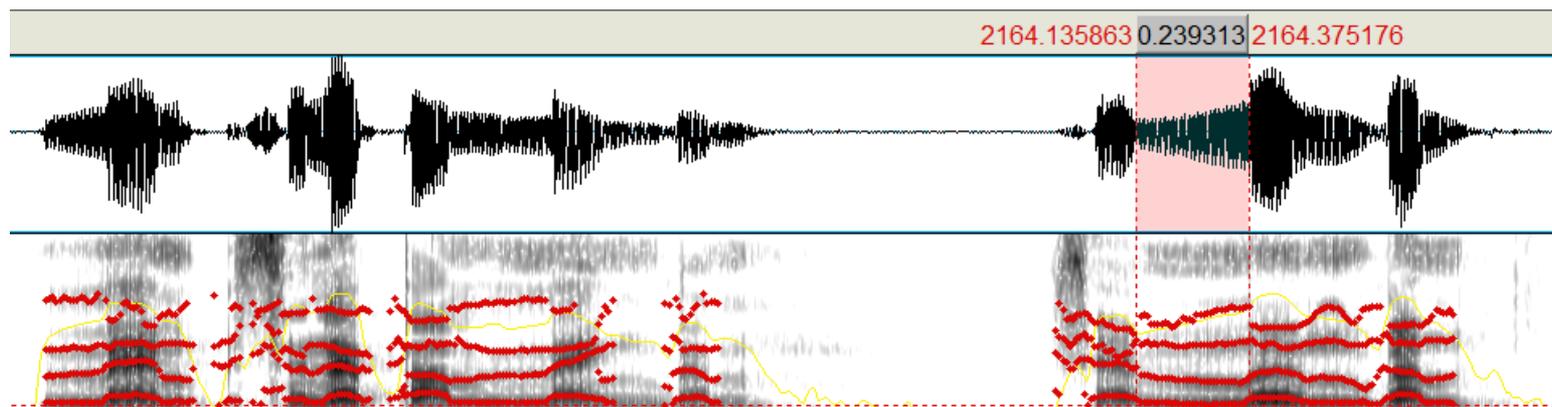
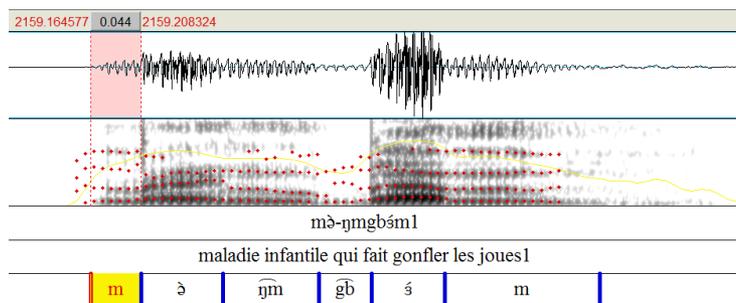


Consonant length in the nonsense word *mè-màmà* (Eton, Bantu A70)



- Corrective focus on the prefix V realized with prefix C-emphasis

Eton (A70)



FR+ET: Mais, ce n'est pas mè-ηmgbám (FOC), c'est mè-ηmgbám (FOC)

FR+ET: Mais, ce n'est pas mè-ηmgbám (FOC), c'est mè-ηmgbám (FOC)

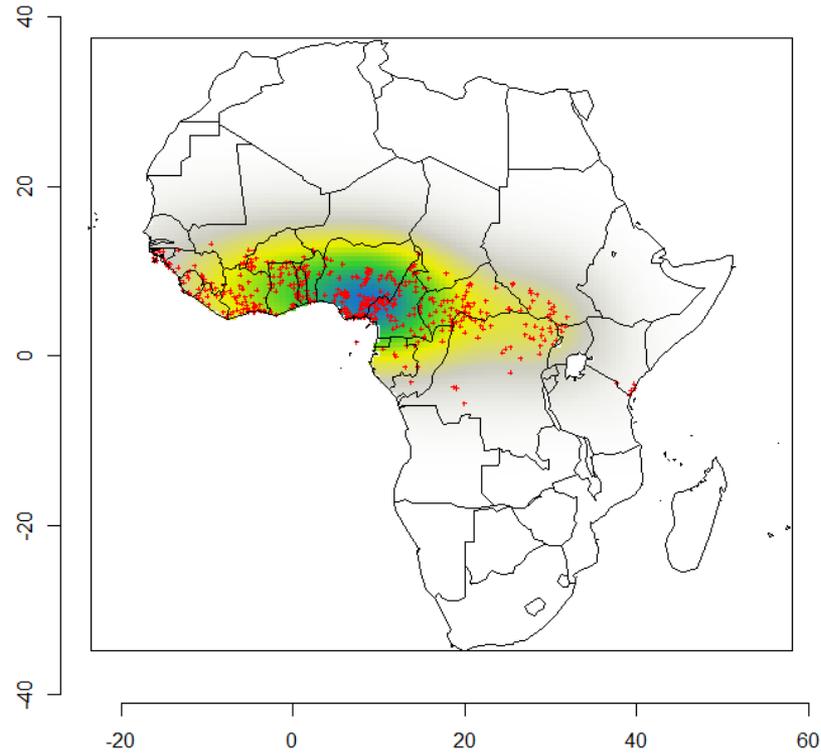
m | è | ηm | gb | s | m

m | è | η | gb | s | m

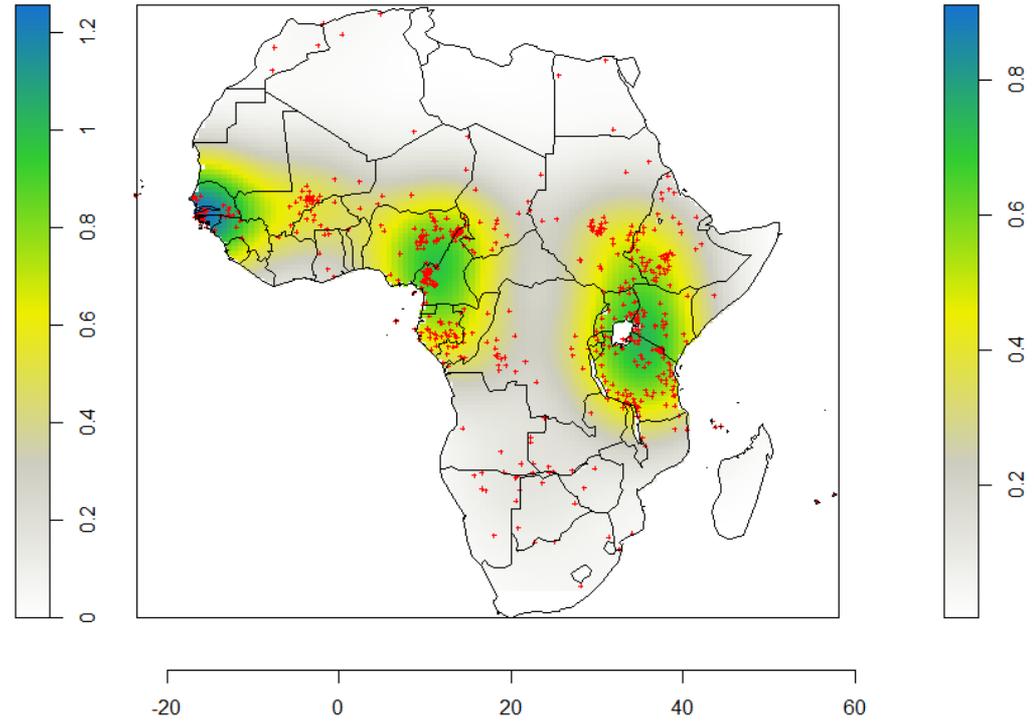


- In a broader perspective, **C-emphasis prosody** is a very good candidate for the role of a **major driving force** behind the emergence of several other types of sounds, such as labial flaps, bilabial trills, and clicks

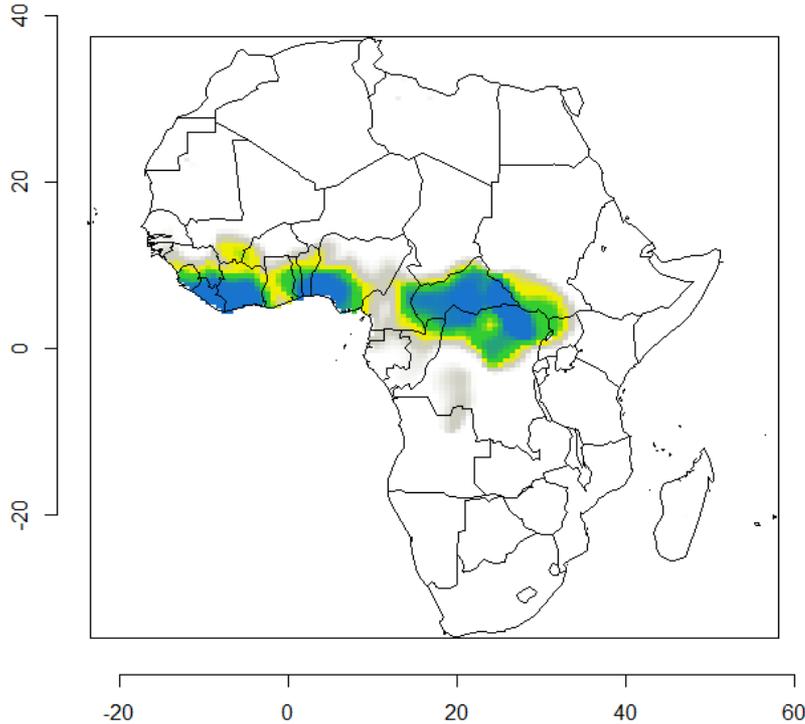
LVall_Y languages: geographic distribution



LVall_N languages: geographic distribution

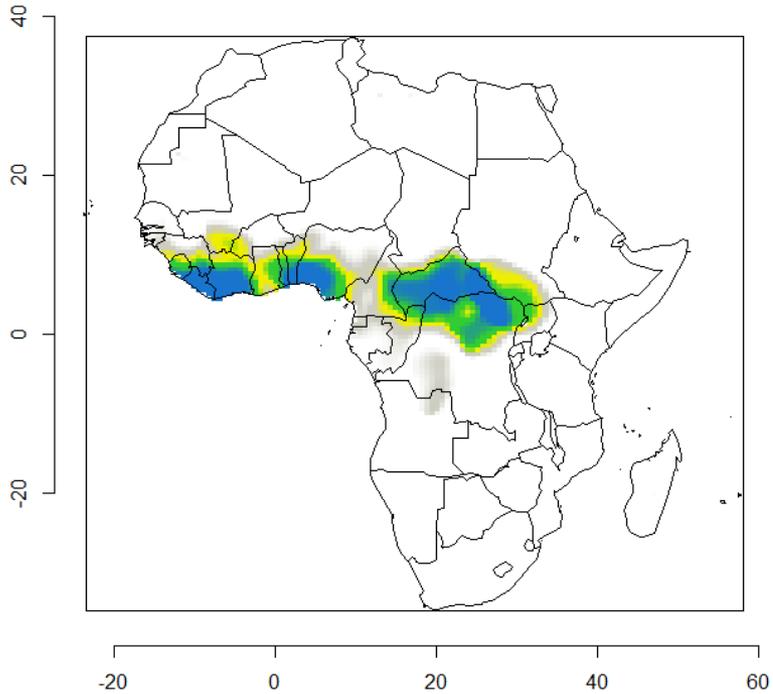


Spatially interpolated log-LVFreq (for LVall)

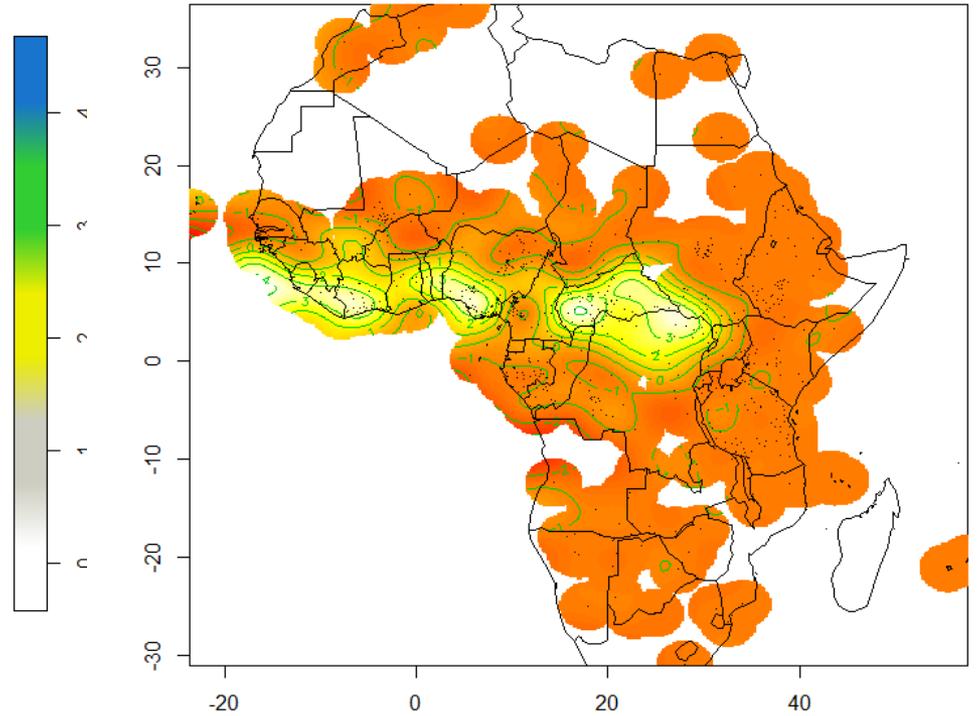


- 2 clearly separated clusters
 - Coastal West Africa (possibly itself composed of 2 sub-clusters)
 - Central Africa
- possibly, + 1 less prominent cluster
 - SE Mali & SW Burkina-Faso
- 1 major spatial discontinuity
 - NE Nigeria & Cameroon
- 1 minor spatial discontinuity
 - Ghana

Spatially interpolated log-LVFreq (for LVall)



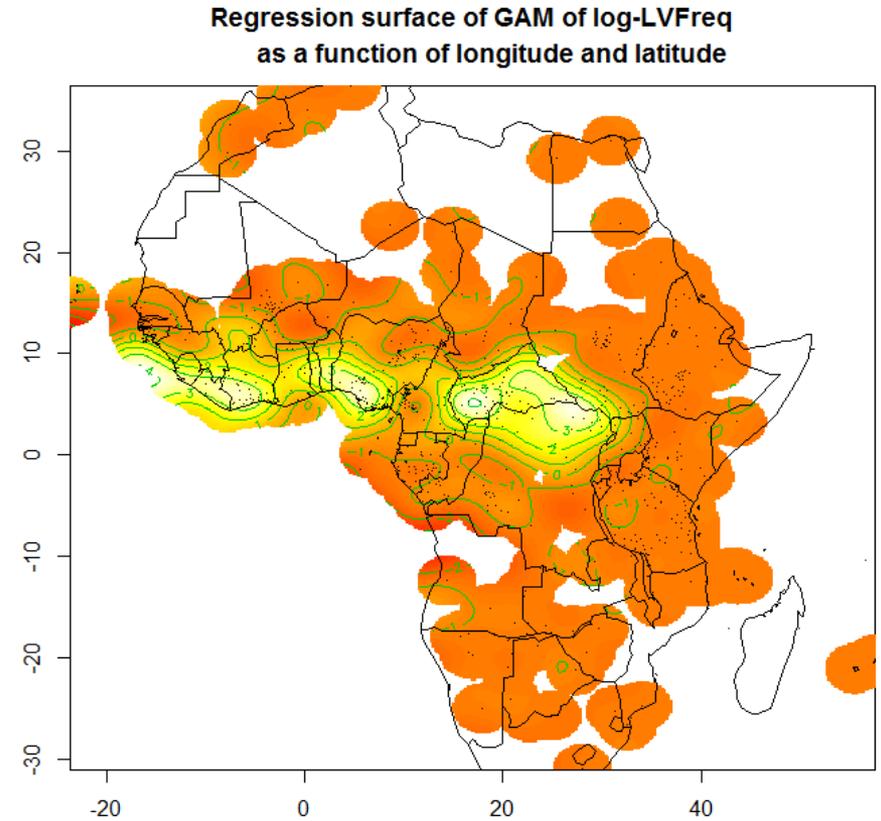
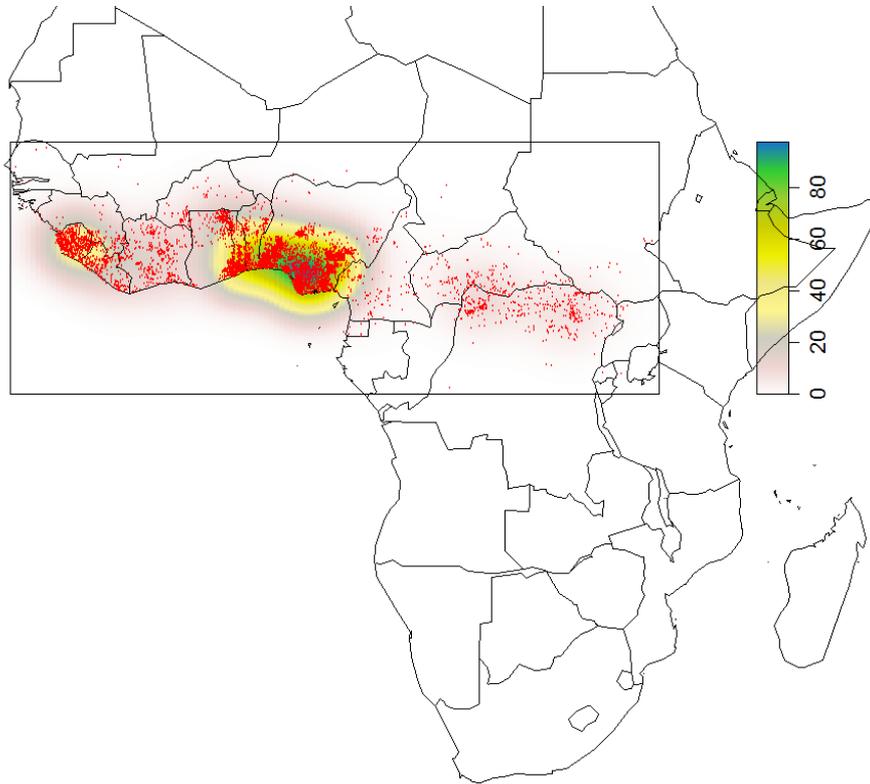
Regression surface of GAM of log-LVFreq
as a function of longitude and latitude



(thin-plate regression splines, k=16, family=Gaussian)



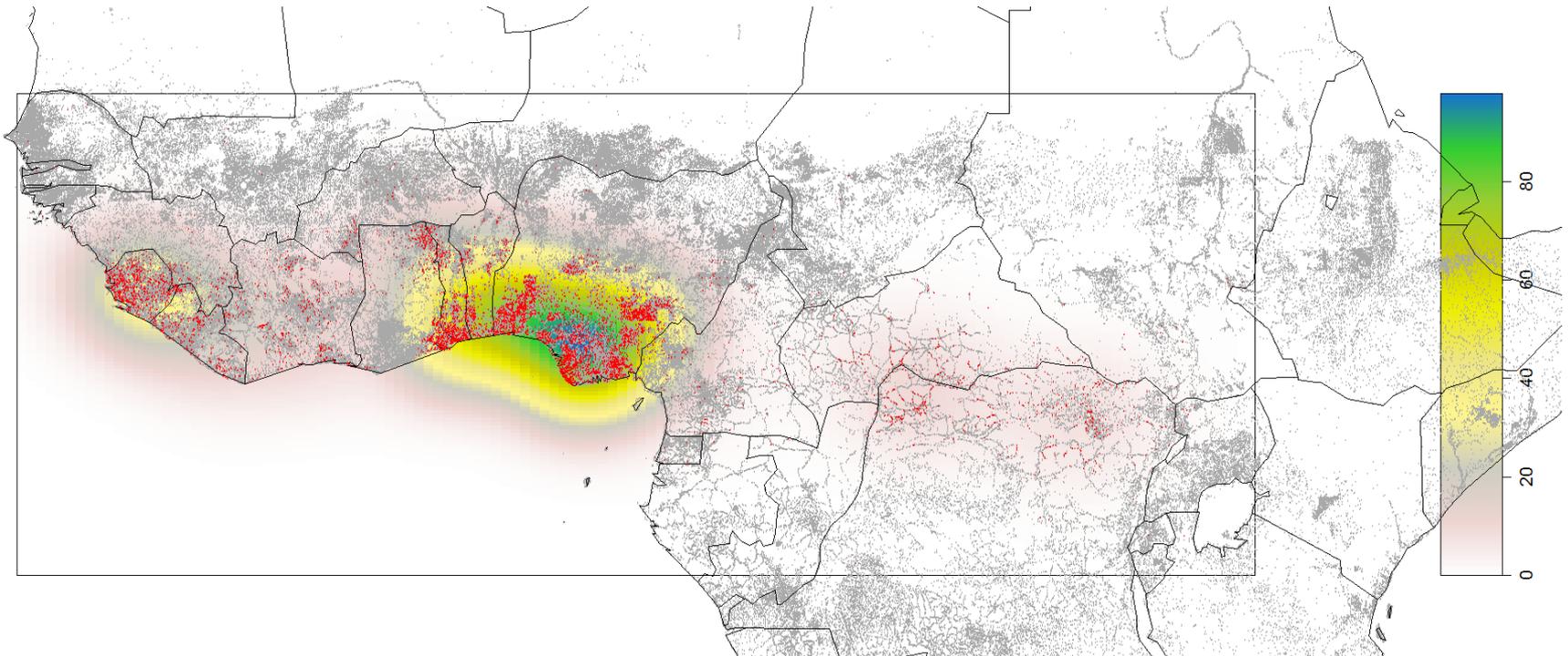
- How can we **cross-validate** our model?
- Spatial distribution of **settlement names spelled with a LV** (such as “kp”, “gb”, Yoruba “p”) on the assumption that:
 - **H₀**: Frequency of settlement names with LV in a given area should roughly correlate with (be representative of) lexical frequency of LV in the languages spoken in the area
- **Big data approach**: quantity compensates for quality
- Settlement names data source: **GeoNames.org**



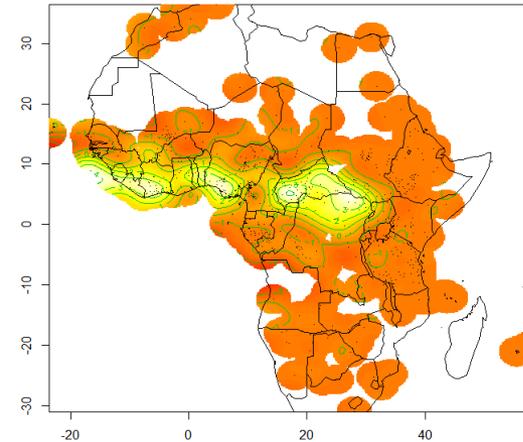
(thin-plate regression splines, k=16, family=Gaussian)

Spatial intensity of unique settlement names
with a <LV>

- The significance of the clusters should be evaluated against the general **population density** in the respective areas:
 - The seeming weakness of the E-most cluster is an artefact of the low population density in Central Africa
 - Both discontinuities are significant



Regression surface of GAM of log-LVFreq
as a function of longitude and latitude

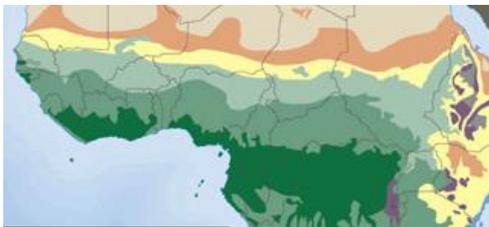
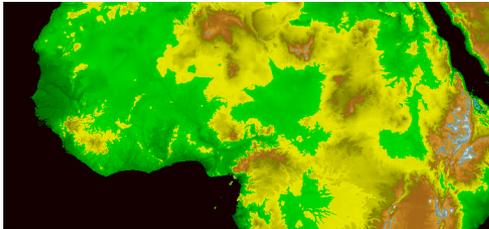
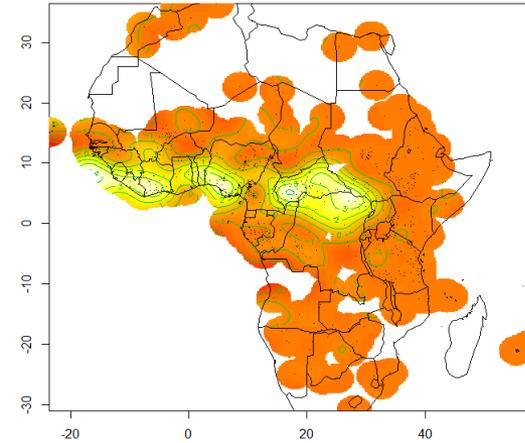


(thin-plate regression splines, k=16, family=Gaussian)

- **Logically**, the 3 major zones of high LVFreq (and the possible minor zone) are most likely to be **refuge zones**:
 - Typologically, LV are rare
 - Several emergent hotbeds of high LVFreq historically independent of each other are unlikely

HISTORICAL IMPLICATIONS: REFUGE ZONES

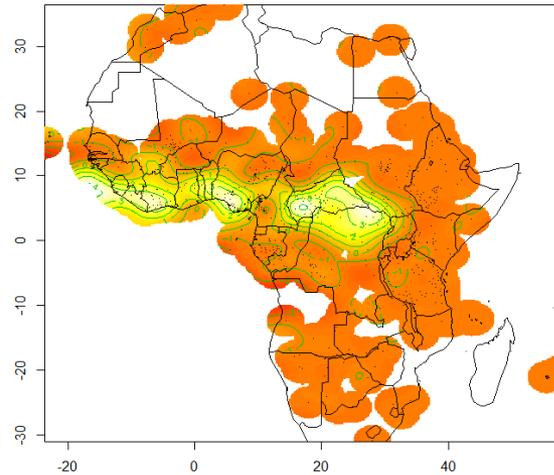
Regression surface of GAM of log-LVFreq
 as a function of longitude and latitude



- **Geographically**, the 3 major zones of high LVFreq (and the possible minor zone) also look like **refuge zones**: mostly forests delimited by **natural boundaries** (sea, savanna, mountain ranges)
- Ghana discontinuity \approx Dahomey forest gap
- NE Nigeria & Cameroon discontinuity \approx Adamawa Plateau, Cameroon mountains



Regression surface of GAM of log-LVFreq
as a function of longitude and latitude

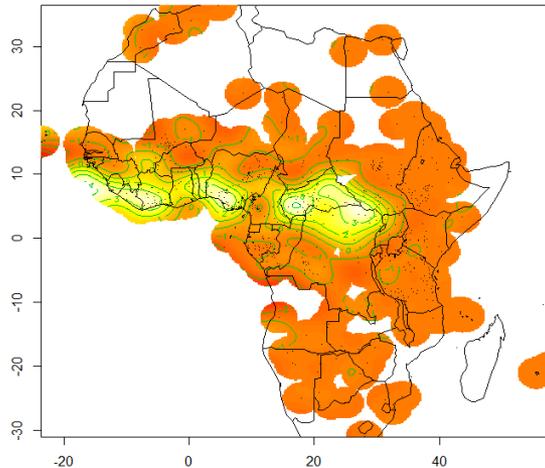


(thin-plate regression splines, k=16, family=Gaussian)

- “hotbeds” → **older presence** of LV and ultimately SIC-accent and C-emphasis prosody
- Given the refuge zone nature of the “hotbeds”, they are probably “hotbeds” not so much for propagation but for **retention** of the feature C-emphasis and derived features, including SIC-accent & LV, present in the original population



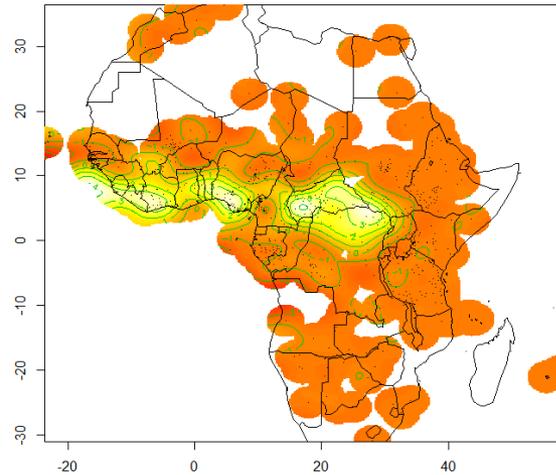
Regression surface of GAM of log-LVFreq
 as a function of longitude and latitude



(thin-plate regression splines, k=16, family=Gaussian)

- **Genetic make-up of the hotbeds:**
 - W: mostly Niger-Congo
 - E: Gbaya, Ubangian, parts of Central Sudanic
- **Genetic make-up of the periphery:**
 - W: mostly Niger-Congo
 - E: Niger-Congo, parts of Central Sudanic
- Linguistically, **the original population** with C-emphasis/SIC-accent/LV may be almost any of these (unlikely Niger-Congo or Central Sudanic) or none
- Hotbeds as refuge zones & retention:
 - hotbeds || language shift
 - periphery || change in language contact situations

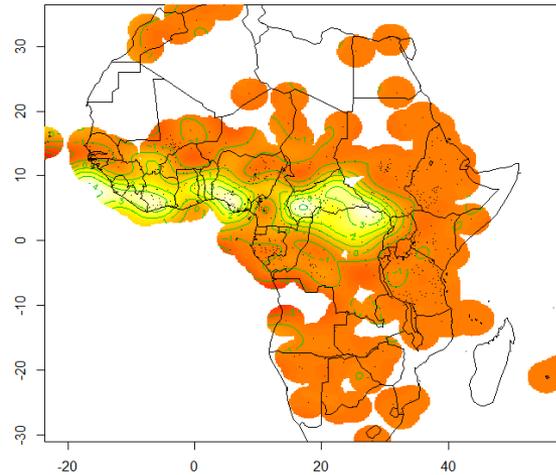
Regression surface of GAM of log-LVFreq
as a function of longitude and latitude



(thin-plate regression splines, k=16, family=Gaussian)

- LV (and correlated phonetic and phonological features) **should not be reconstructed** for Proto Niger-Congo or any of its major branches
- We should also be very **cautious** about reconstructing LV for lower-level branches (problems with “**the majority wins**” rule)

Regression surface of GAM of log-LVFreq
as a function of longitude and latitude

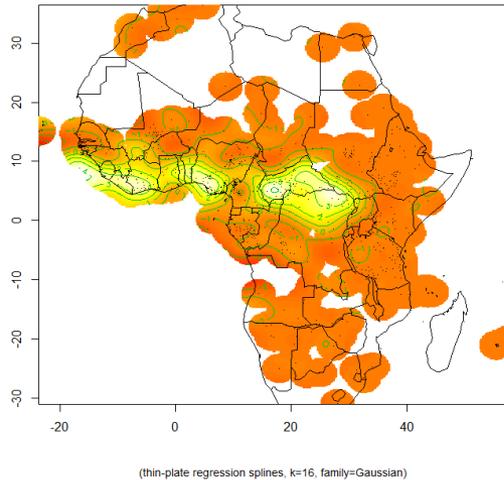


(thin-plate regression splines, k=16, family=Gaussian)

- A rather **northern** localization of the **homelands** of most **major branches** of Niger-Congo in **grassland** and **savanna** ecoregions
- The **homeland** of **Proto Niger-Congo** is then likely to have been located in the northern part of the former extent of grassland and savanna ecoregions
- Probably, somewhere in **present-day Sahel** or **southern Sahara**.

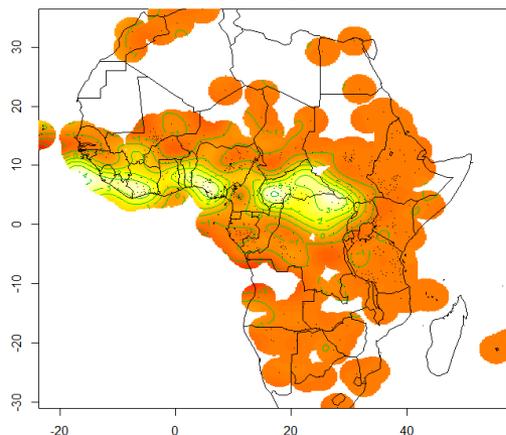


Regression surface of GAM of log-LVFreq
as a function of longitude and latitude



- Bantoid, Adamawa and Chadic are responsible for the **major discontinuity** around Cameroon & NE Nigeria
- The majority of **Bantu languages** (a major Bantoid subgroup) are spoken outside of the hotbeds of high LVFreq
- Bantoid & Adamawa are likely to have **arrived** in the discontinuity area relatively **recently**
- Bantoid may have passed it & then re-entered or just entered late

Regression surface of GAM of log-LVFreq
as a function of longitude and latitude



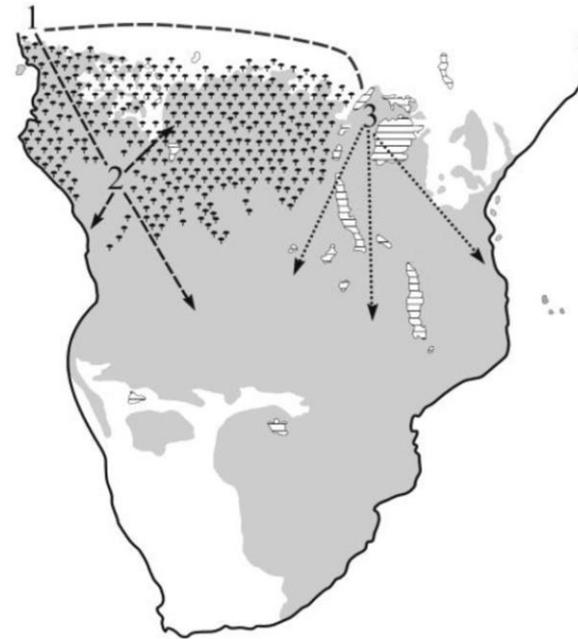
(thin-plate regression splines, k=16, family=Gaussian)

- Further spread of Bantoid, particularly **Bantu expansion**, must have happened **without much language shift** from any original “LV” populations involved
- The only Bantu group with a relatively high LVFreq are **languages in the N of DRC** which moved into the hotbed of high LVFreq from SW
- The spread of Bantu in the N or DRC is most likely to have been a **spread of languages through language shift** much more than a spread of Bantu speaking populations

Two principal models of Bantu expansion

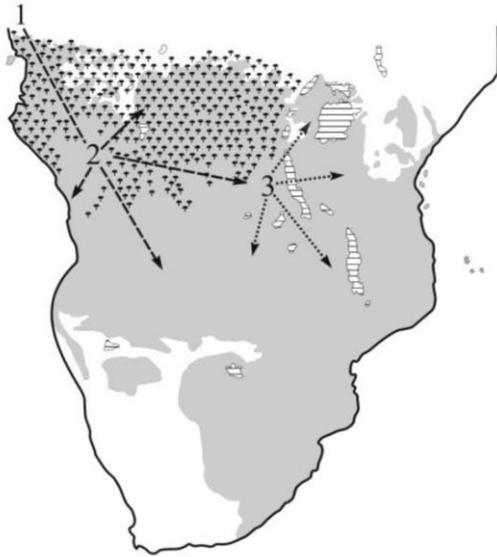


a. East out of West

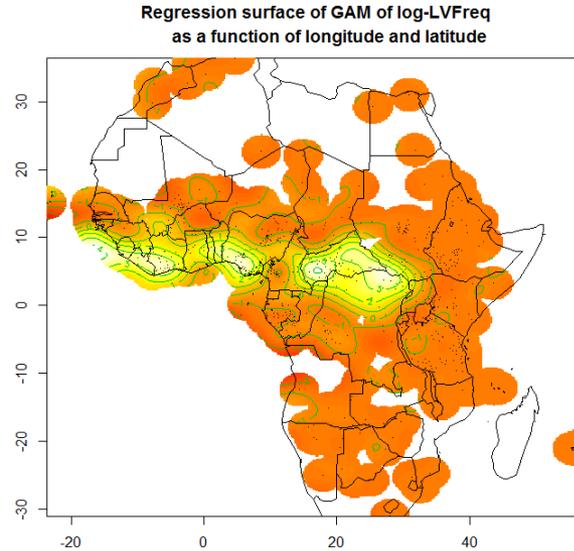


b. East separate from West

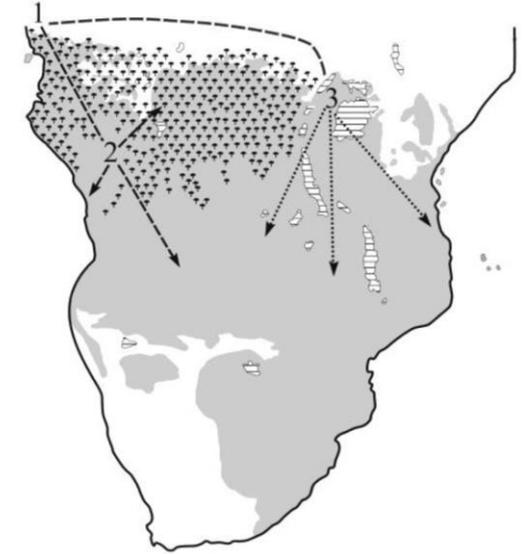
(adapted from Pakendorf et al. 2011 :8).



a. East out of West



(thin-plate regression splines, k=16, family=Gaussian)

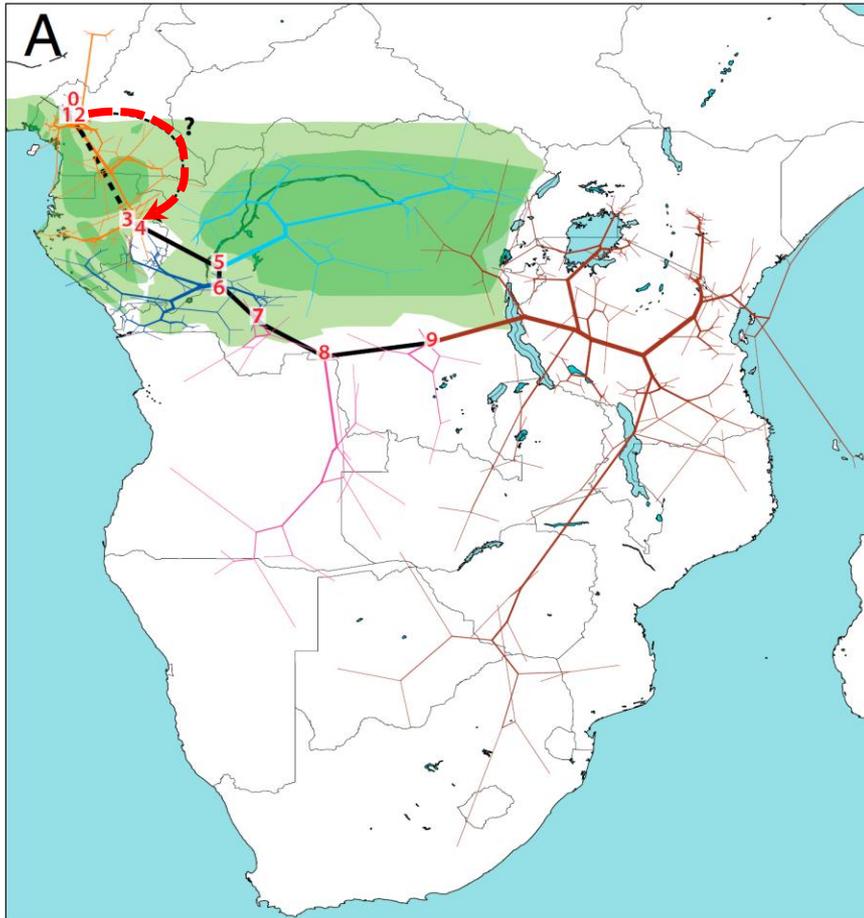


b. East separate from West

- Our model supports the “**East-out-of-West**” hypothesis of the E Bantu emergence with the E Bantu break-off point somewhere south of the rainforest



HISTORICAL IMPLICATIONS: BANTU EXPANSION



Bantu migration route reconstructed by Grollemund et al. (2015) on consensus tree by using geographical locations of contemporary languages and connecting ancestral locations by straight lines (true route will differ).

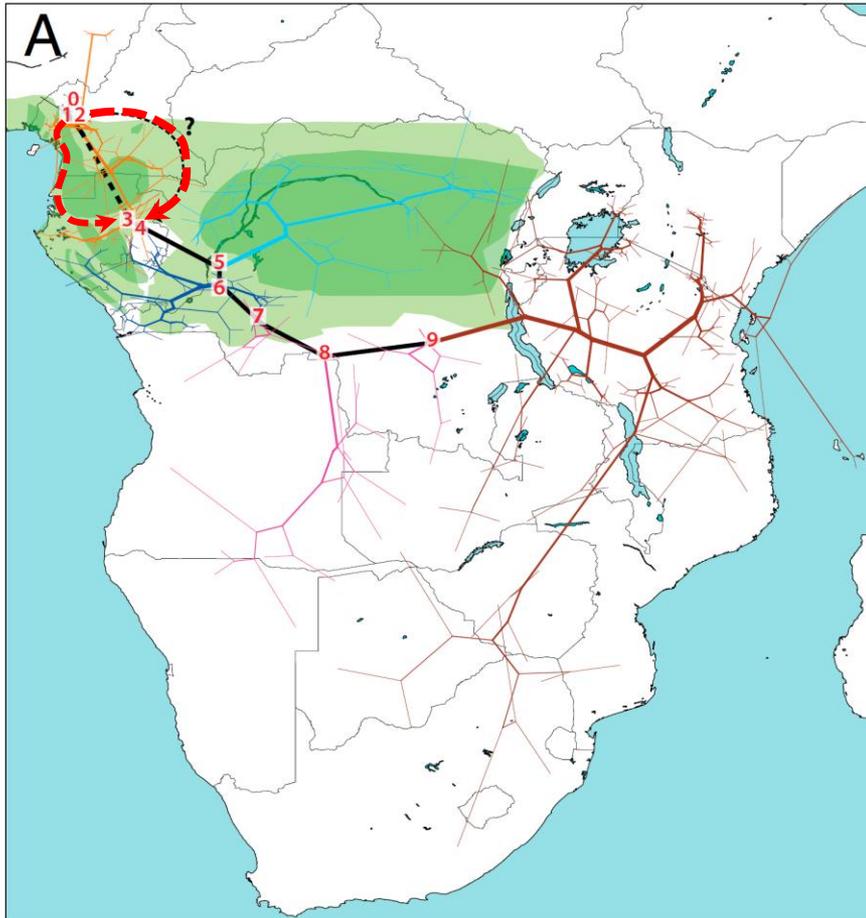
Numbered positions correspond to major diversification nodes on the consensus tree.

Curved dashed line indicates suggested **migration route through savannah corridors (Sangha River Interval)**

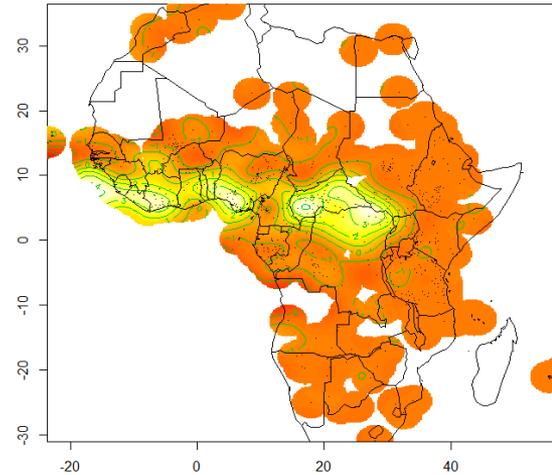
Lighter green shading corresponds to the delimitation of the rainforest at 5.000 B.P.; the darker green corresponds to the delimitation of the rainforest at 2.500 B.P.



HISTORICAL IMPLICATIONS: BANTU EXPANSION



Regression surface of GAM of log-LVFreq
as a function of longitude and latitude



(thin-plate regression splines, k=16, family=Gaussian)

- Our model suggests that the migration between nodes 2 and 3 is more likely to have happened through a **coastal route** rather than the Sangha River Interval.