

## I. Overview

- Models of long-distance phonology differ in whether they propose an explicit proviso for identity.
  - Gallagher & Coon (2009): IDENTITY, which requires consonants linked by a correspondence relation to be identical.
  - Rose & Walker (2004) and other work in ABC: identity not a goal; no explicit proviso for identity.
  - Zuraw (2002): identity is a goal; no explicit proviso for identity.
- A **question** that this raises: should identity preferences be analyzed using one monolithic constraint, or a set of constraints, each requiring identity for an individual feature?
- This poster:** a case study of Ngbaka vowels, which suggests that we need both.

## II. Background and research question

- Ngbaka (Ubangian; Thomas 1963, Henrix et al. 2007, Sélézilo 2008, *a.o.*) has a twelve-vowel system.

Oral vowels		Nasal vowels	
i	u	ĩ	ũ
e	o	ẽ	õ
ɛ	ɔ	ẽ̃	õ̃
ɑ		ã	

- Descriptions of Ngbaka (references above) often claim that multiple types of vowel harmony are active. Two examples:
  - ATR harmony: [e ẽ o õ] cannot precede or follow [ɛ ẽ̃ ɔ õ̃].
  - Backness harmony: [i ĩ e ẽ] cannot precede or follow [u ũ o ɔ̃].
- These requirements often coincide to ensure that, in a CV<sub>1</sub>CV<sub>2</sub> word, V<sub>1</sub> and V<sub>2</sub> are identical (noted by Sélézilo 2008). Another analytical possibility arises: Ngbaka vowels *prefer to be identical*.
- Question:** can the apparent preference for identity in Ngbaka vowels be explained through the interaction of multiple types of harmony, or must an identity preference be recognized independently?

## III. Data

- To address this question, I created a database of vowel pairs from Henrix et al.'s (2015) Ngbaka dictionary (5,571 words).
- Investigation limited to disyllabic or longer words that contain only CV syllables (3,928 words).
- For each word, I extracted each adjacent pair of vowels.

z i b ɔ<sub>1</sub> l ɔ<sub>2</sub>                      t a l ɛ k u s i  
 Pairs: [i ɔ<sub>1</sub>], [ɔ<sub>1</sub> ɔ<sub>2</sub>]                      Pairs: [a ɛ], [ɛ u], [u i]

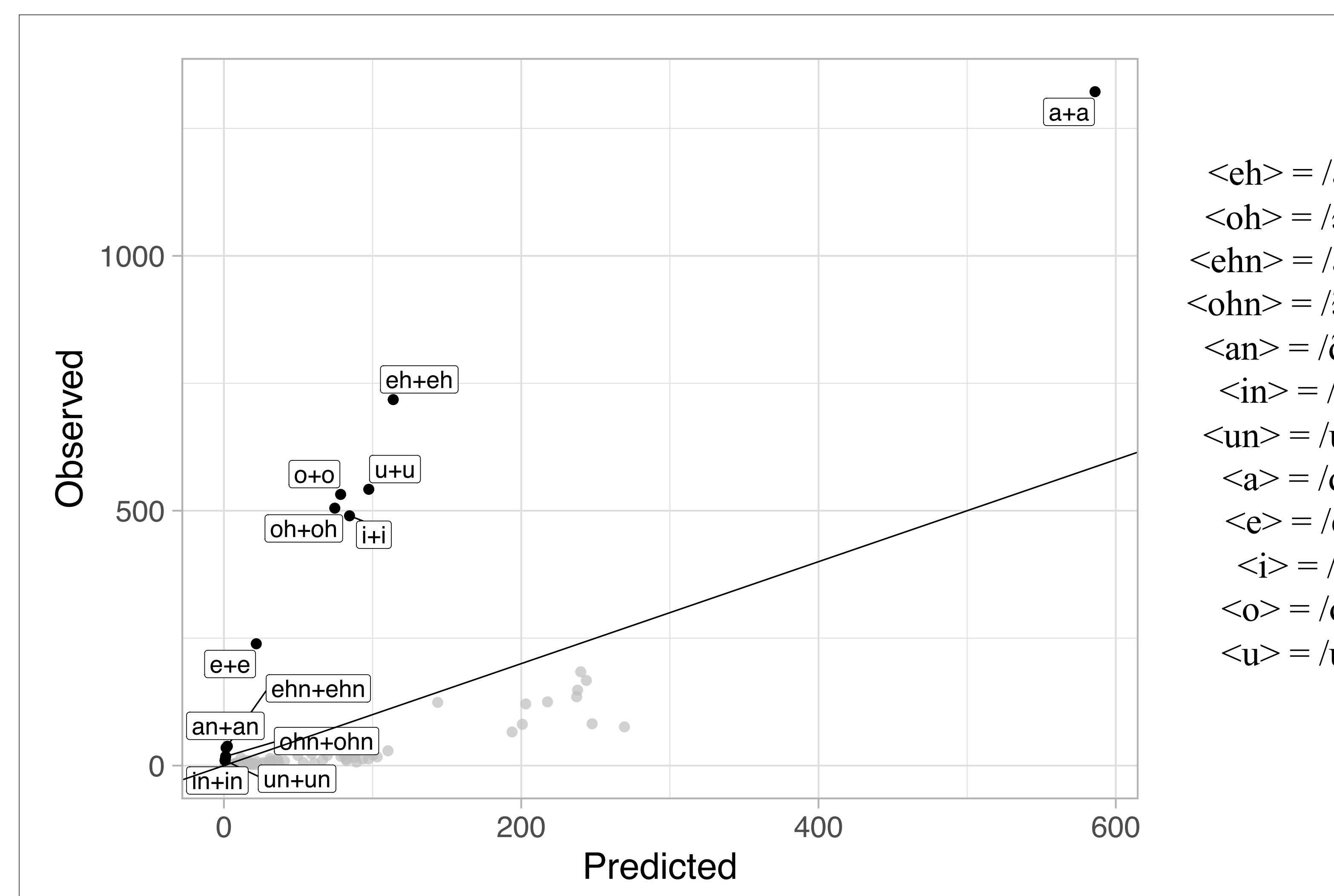
- Result:** 6,716 vowel pairs. Identical pairs (n=4,461, in white) are more common than non-identical pairs (n=2,255, in black).

		V <sub>2</sub>											
		ɑ	ã	e	ɛ	ẽ	i	ĩ	o	ɔ	õ	u	ũ
V <sub>1</sub>	ɑ	1322	33	124	76	10	167	6	135	81	14	148	8
	ã	7	38	1	0	0	9	9	4	3	1	3	1
	e	26	0	239	5	1	5	0	13	7	1	10	0
	ɛ	82	3	9	718	6	17	0	21	24	2	33	0
	ẽ	4	4	0	1	35	1	1	2	1	1	3	4
	i	121	6	20	13	2	490	1	59	20	4	10	1
	ĩ	2	2	2	0	0	1	12	0	3	4	0	0
	o	66	2	55	7	1	20	0	532	11	1	18	0
	ɔ	125	10	6	47	3	44	5	16	505	15	25	0
	õ	2	2	0	4	0	2	5	0	4	18	1	0
	u	184	5	23	29	4	62	3	13	13	1	542	4
	ũ	4	2	0	0	5	1	4	1	0	0	5	10

- An interesting trend to note: if a vowel co-occurs with a non-identical vowel, it's usually [ɑ].
- In addition, there are trends in these data that I won't focus on (example: nasal vowels are less common than oral vowels).

## IV. Analysis

- Analysis: loglinear model of count data, using *bayesglm* from R's arm package (Gelman & Hill 2007).
- Baseline model asks: **what is the predicted frequency of each vowel pair, given the independent frequency of each vowel?**
  - Dependent variable: number of times a particular vowel-vowel pair is attested.
  - Independent variables: one predictor per vowel per position (ɑ<sub>1</sub>, ɑ<sub>2</sub>, ã<sub>1</sub>, ã<sub>2</sub>, e<sub>1</sub>, e<sub>2</sub>, etc.).
- Fitted values (obtained with R's *fitted.values* function) show that identical pairs are overattested relative to expectation.



For an interactive graph where all points are labeled, go to:

[julietstanton.github.io/files/ngbakavowels.html](http://julietstanton.github.io/files/ngbakavowels.html)



- To this baseline model, I added predictors reflecting types of vowel harmony and one predictor to look for effects of identity.

	Predictor (all binary)	Type of harmony	Assigns a 1 to...
a.	*[αATR][-αATR]	ATR	(e o)(ɛ ɔ) and (ɛ ɔ)(e o)
b.	*[αback][-αback]	Backness	(i e ɛ)(u o ɔ) and (u o ɔ)(i e ɛ)
c.	*[αhigh, -back, -low][-αhigh, -back, -low]	Height amongst front vowels	(i)(e ɛ) and (e ɛ)(i)
d.	*[αhigh, +back, -low][-αhigh, +back, -low]	Height amongst back vowels	(u)(o ɔ) and (o ɔ)(u)
e.	*[αnasal][-αnasal]	Nasal	vowels mismatching for nasality
f.	*[+syllabic] <sub>i</sub> [+syllabic] <sub>j</sub>	Identity	non-identical vowel pairs

- Question:** which combination of predictors is responsible for shaping the data?
- To answer this question, I fit the maximal model to the data and compared the goodness of fit of nested models using LRTs.

## V. Results

- The best-fit model includes predictors for **ATR harmony** (a; t = 5.51, p < .001), **backness harmony** (b; t = 4.38, p < .001); **height harmony amongst front vowels** (c; t = 5.07, p < .001), **nasal harmony** (e; t = 6.64, p < .001), and **identity** (f; t = 29.23, p < .001).
  - Further modeling suggests another possible type of harmony, between low [ɑ ã] and the [-low] vowels, is inactive.
- In addition, investigation of the coefficients shows that the identity predictor plays the largest role in shaping these data.

	Predictor (all binary)	Type of harmony	Coefficient
a.	*[αATR][-αATR]	ATR	0.40
b.	*[αback][-αback]	Backness	0.26
c.	*[αhigh, -back, -low][-αhigh, -back, -low]	Height amongst front vowels	0.53
e.	*[αnasal][-αnasal]	Nasal	0.63
f.	*[+syllabic] <sub>i</sub> [+syllabic] <sub>j</sub>	Identity	1.04

- Conclusion:** The fact that the identity predictor is significant tells us that the identity preference in Ngbaka vowels cannot be explained entirely by appealing to interacting processes of harmony; the identity predictor plays an independent role.