# The Automatic Characterization of Grammars from Small Wordlists

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### **Vowel Harmony Overview**

An unsupervised algorithm for detecting and describing vowel harmony systems in small wordlists. It answers the following questions about an unknown language:

- Does the language have harmony?
- What are its harmonizing sets?
- Does it have neutral (transparent or opaque) vowels?
- Does it have secondary harmony?

### **Vowel Harmony Algorithm**

The algorithm is designed to work on short wordlists (down to about 500 types) without frequency counts. If the standard orthography roughly approximates a phonemic representation, no transcription is needed. If available, token frequencies may be used to improve results. Furthermore, the algorithm can provide a mapping between harmonizing sets if the researcher provides vowel features as input.

if frequencies provided then
Trim tail off <i>wordlist</i>
while <i>True</i> do
Calculate tier-adjacent V-V co-occurrence matrix
Calculate MI between each vowel pair
Identify vowels whose MI distributions uniform
within threshold.
Assign these to the neutral vowel set and remove
from consideration
if number of non-neutral vowels $\leq 1$ then
return
Run k-means ( $k = 2$ ) clustering on the remaining
vowels' MI vectors
if no <i>features</i> provided <b>then</b>
return
else
Map vowels between harmonizing sets by find-
ing pairs that share the most features in common.
<i>vowel list</i> $\leftarrow$ Collapse vowels along the harmo-
nizing feature
rerun for secondary harmony
return

### Results

Lang. Turkish 3 Finnish 3 Hungar. Uyghur 3 Warlpiri German 2 English <sup>2</sup>

 
 Table 1: Vowel co-occurrences are taken from corpus
orthographies. Marginal vowels (e.g. Finnish å and German y) are automatically detected and removed. Corpora are from MorphoChallenge Kurimo et al. (2010) when available. Uyghur and Hungarian were provided for the DARPA LORELEI project. Warlpiri is from Swartz (1997).

# **Next Steps**

We are continuing to develop this algorithm.

- Leveraging paradigms from our morphological segmentation will allow it to map harmonizing vowels with explicitly provided features.
- The same distributional processes can discover other typological features: whether a language exhibits stem alternations, has agglutinative morphology, tends towards prefixation or suffixation, reduplication, etc.

### References

Types	1ary H?	Correct	2ary H?	Correct
03,013	$\checkmark$	8/8	$\checkmark$	4/4
96,770	$\checkmark$	8/8	—	—
53,839	$\checkmark$	11/15	—	—
92,403	$\checkmark$	7/8	_	_
28,885	$\checkmark$	3/3	_	—
25,327	_	5/5	_	
01.438	_	6/6	_	_

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- Narasimhan, K., Barzilay, R., and Jaakkola, T. (2015). An unsupervised method for uncovering morphological chains. *arXiv preprint arXiv:1503.02335*.
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# **Segmentation Overview**

An unsupervised morphological segmentation algorithm designed with small wordlists in mind. Our algorithm is built around the concept of **paradigms**. Each root is attached to a paradigm containing all the proposed suffixes with which it is attested.

This algorithm achieves state-of-the-art results on English and Turkish. We are preparing gold standards for testing on other languages as well.

# **Segmentation Algorithm Summary**

The morphological segmentation algorithm combines three processes: segmentation, paradigm construction, and pruning.

• Segmentation - A Bayesian model estimates probability P(r, s, t|w) over candidate roots, affixes, and transformations for each word

$$P(r, s, t | w) = \frac{P(r) z}{\sum_{k \in V}}$$

• Paradigm Construction - Affix appearing with each root are grouped together into paradigms. The more common its paradigm, the greater its *support*.

Paradigm	Support
(-ed, -ing, -s)	772
(-ed, -ing)	331
(-ed, -er, -ing, -s)	219
(-ly, -ness)	208
(-ed, -ing, -ion, -s)	154

• Pruning - Affixes which do not appear in enough well-supported paradigms are pruned. For example, if *closet* is incorrectly segmented as close-t, the *close* paradigm becomes {*-er*, *-est*, *-ed*, *-ing*, *-s*, *-t*}. Pruning corrects the *-t*.

### $\times P(s) \times P(t|f(r,s))$ $\sum_{(r',s',t')\in w} P(r',s',t')$

### Results

Lang.	Model	Prec.	Recall	l
	Morfessor-Base	0.740	0.623	C
	AGMorph	0.696	0.604	C
English	MorphChain-C	0.555	0.792	C
	MorphChain-All	0.807	0.722	C
	Our model	0.804	0.764	C
	Morfessor-Base	0.827	0.362	C
	AGMorph	0.878	0.466	C
Turkish	MorphChain-C	0.516	0.652	C
	MorphChain-All	0.743	0.520	C
	Our model	0.589	0.726	C

Table 2: All numbers except for ours are reported in Narasimhan et al. (2015). Best results are reported.

# **Next Steps**

We are still developing this tool. We expect improvements to come from integration with the vowel harmony analyzer as well as more theoretically involved morphological transformations.

- Have run it on other languages: Tagalog, Navajo, Yoruba, Somali; but cannot score the outputs yet
- Are designing a **segmentation annotation scheme** to create more gold standards
- Will leverage the vowel harmony tool to create more coherent paradigms
- Will enrich transformations for languages with non-concatenative morphology

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