



# Advancing the study of endangered languages with computational tools for morphology

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# ADVANCING THE STUDY OF ENDANGERED LANGUAGES WITH COMPUTATIONAL TOOLS FOR MORPHOLOGY

## The case of Asama verb paradigms

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### BENEFITS OF USING COMPUTATIONAL TOOLS

1. Test the accuracy and coverage of the analysis of the morphology.
  2. Partially automate the glossing of texts.
  3. Enable quantitative measures of morphological complexity.
- ⇒ Advance the study of endangered languages.

### ASAMA



- Endangered Ryukuan language (Japonic family).
- Reference grammar project by D. Lévéque.
- 100 full + 400 near-full inflectional verb paradigms collected.
- Non-canonical phenomena (e.g. alternation of tones and vowel length) in verb inflection.

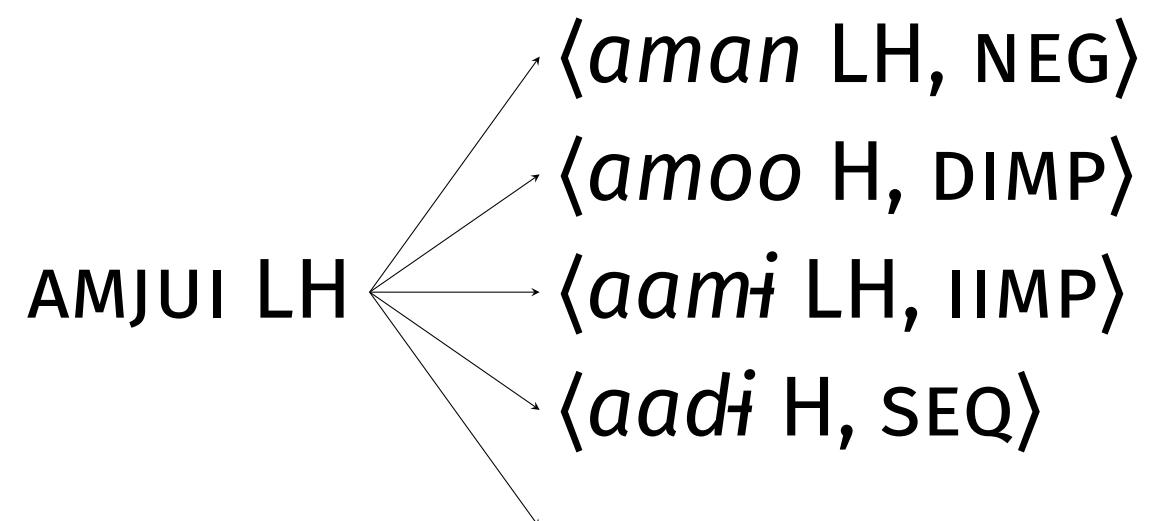
	'to sell'	'to knit'	'to go out'
NPST	ujui H	amjui H	izjijui LH
CVB2	urug99si H	amjug99si H	izjirug99si LH
NEG	uran H	aman H	izjiran LH
DIMP	uroo H	amoo H	izjiroo LH
IIMP	urii H	aami HL	izjiri LH
CVB	ui H	amii H	izjii LH
DES	uicjaahai H	amicjaahai H	izjicjaahai LH
PST	utan H	adan H	izjitan LH
SEQ	utii H	aadi HL	izjiti LH
PROG	utui LHL	aadui HL	izjuti LHL

### 1 LINGUISTIC DOCUMENTATION

**GOALS** To speed up and ease the glossing of texts.

**ISSUES** Small amount of data, limited time, usual tools (Toolbox) not well-fitted to non-concatenative phenomena.

**SOLUTION** The transducer produces complete paradigms for all existing lexemes.



tourner - IPFV - PART - PROH māwaj - u - N - na  
tourner - IPFV - PST - PART māwaj - ut - a - N

**RESULTS** A simple Python script is used to format the output, which can then be directly imported into Toolbox and used in the process of interlinearisation.

```

\lx māwajunna
\mo māwaj - u - n - na
\ge tourner - IPFV - PART - PROH

\lx māwajutaN
\mo māwaj - ut - a - N
\ge tourner - IPFV - PST - PART
  
```

### WHY COMPUTATIONAL TOOLS?

- “The insufficiency of paper-and-pencil linguistics” (Karttunen 2006).
  - Finite-state transducers (Beesley & Karttunen 2003).
  - Bidirectional:
    - PRODUCTION** lexeme/root → inflected form;
    - RECOGNITION** inflected form → lexeme/root.
  - Syntax similar to that of phonological rewrite rules familiar to many linguists.
- ```

    regex V -> θ || V V _ ;
    regex i -> ɨ || r _ ;
    regex ə -> j || s _ a ;
  
```
- FSTs have sufficient power to handle (almost) any morphological phenomena.
  - Provide a maximally precise and explicit description.
  - Free software implementations (Foma).

### 2 LINGUISTIC ANALYSIS

- GOALS** Check the accuracy of the description and test hypotheses on new lexemes.
- ISSUES** Non-canonical verb morphology, multiple classes.
- DESCRIPTION** Two orthogonal sets of rules associated with two sets of classes.

| T <sub>1</sub> | T <sub>2</sub> | T <sub>3</sub> | T <sub>4</sub> | T <sub>5</sub> |
|----------------|----------------|----------------|----------------|----------------|
| Xbj            | -              | Xb             | Xb             | Xd             |
| Xmj            | -              | Xm             | Xm             | Xd             |
| Xkj            | -              | Xk             | Xk             | Xcj            |
| Xcj            | -              | Xc             | Xt             | Xccj           |
| Xj             | X              | Xr             | Xr             | Xcj            |
| Xj             | X              | Xr             | Xr             | Xccj           |
| Xj             | X              | X              | Xr             | Xtt            |

| Class | NPST                    | IMP.DIR                 | SEQ                       |
|-------|-------------------------|-------------------------|---------------------------|
| I1    | RT <sub>1</sub> ui (H)  | R:T <sub>4</sub> i (H)  | RT <sub>5</sub> ui (H)    |
| I2    | RT <sub>1</sub> ui (H)  | R:T <sub>4</sub> i (H)  | RT <sub>5</sub> ui (H)    |
| II1   | RT <sub>1</sub> ui (H)  | RT <sub>4</sub> ii (H)  | RT <sub>5</sub> ui (LHL)  |
| II2   | RT <sub>1</sub> ui (H)  | R:T <sub>4</sub> i (H)  | RT <sub>5</sub> ui (LHL)  |
| III   | RT <sub>1</sub> ui (LH) | R:T <sub>4</sub> i (HL) | R:T <sub>5</sub> ui (HL)  |
| IV    | RT <sub>1</sub> ui (LH) | R:T <sub>4</sub> i (LH) | R:T <sub>5</sub> ui (LHL) |
| V     | RT <sub>1</sub> ui (HL) | RT <sub>4</sub> i (HL)  | RT <sub>5</sub> ui (HL)   |

### SOLUTIONS

- FSTs allow to:
- Check the formal description on lexemes and highlight the problems (comparison between data and output);
  - Test the results with the speaker (computation of the whole paradigm for lexemes with incomplete data and check with the speaker).

**RESULTS** Robustness of formal analysis is confirmed by the computational study, and lacks of data in the description are highlighted.

### IMPLEMENTATION

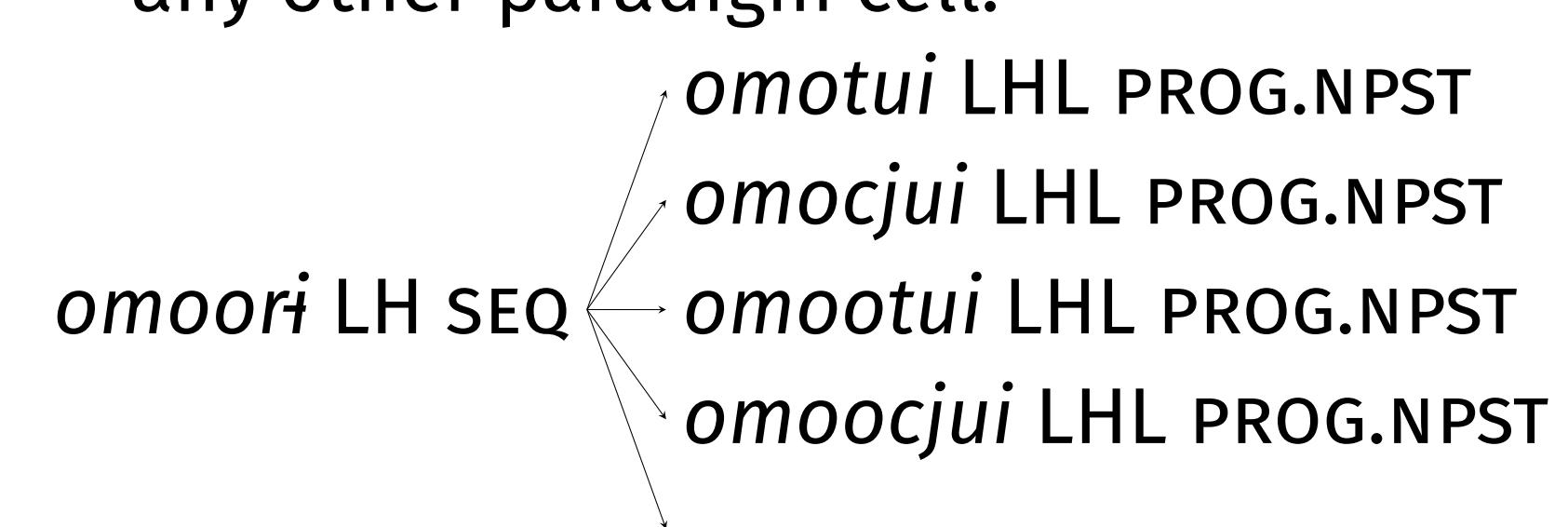
- Content paradigm cell  $\langle L, \sigma \rangle \rightarrow$  realised cell  $w \langle AMJUI LH, SEQ \rangle \rightarrow aadi LH$
- Unlabeled form  $w \rightarrow$  all possible analyses  $\langle L, \sigma \rangle$ 
  - $\langle AMJUI LH, SEQ \rangle$
  - $\langle AAMJUI H, SEQ \rangle$
  - $\langle ABJUI LH, SEQ \rangle$
  - $\langle AABJUI H, SEQ \rangle$
- $L \rightarrow$  full realised paradigm, i.e. a set of cells  $\{ \langle w, \sigma \rangle, \langle w', \sigma' \rangle, \dots \}$ .
- Any realised form  $\langle L, \sigma \rangle \rightarrow$  list of all possible realised forms of any other paradigm cell.

### 3 THEORETICAL MORPHOLOGY

**GOALS** Exploring the **implicative structure** with complexity measures and Shannon **entropy** and solve the *Paradigm Cell Filling Problem* (“What licenses reliable inferences about the inflected (and derived) surface forms of a lexical item?”, Ackerman et al. 2009: 54).

**ISSUES** Too few full paradigms, no pre-existing interactive database.

**SOLUTION** Composition of two FSTs in order to obtain for any given realized cell  $\langle w, \sigma \rangle$  the list of all possible realized forms of any other paradigm cell.



**RESULTS** Entropy measures help identifying the **sources of uncertainty** (unpredictable segmental alternations, neutralisation of vowel length and neutralisation of tone) and the **principal parts** of the system (Finkel & Stump 2007).

| H(C L)    | NPST  | CONV  | IIMP  | SEQ   | PROG.NPST |
|-----------|-------|-------|-------|-------|-----------|
| NPST      | 0.000 | 0.000 | 0.222 | 0.244 |           |
| CONV      | 0.923 | 0.082 | 0.296 | 0.550 |           |
| IIMP      | 0.950 | 0.211 | 0.222 |       | 0.525     |
| SEQ       | 1.362 | 1.012 | 0.425 |       | 0.317     |
| PROG.NPST | 1.204 | 0.910 | 0.423 | 0.000 |           |

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