Improved Neural Protoform Reconstruction via Reflex Prediction

Liang Lu, Jingzhi Wang, David R. Mortensen

Language Technologies Institute, Carnegie Mellon University

{lianglu, jingzhi3, dmortens}@cs.cmu.edu

Abstract

Protolanguage reconstruction is central to historical linguistics. The comparative method, one of the most influential theoretical and methodological frameworks in the history of the language sciences, allows linguists to infer protoforms (reconstructed ancestral words) from their reflexes (related modern words) based on the assumption of regular sound change. Not surprisingly, numerous computational linguists have attempted to operationalize comparative reconstruction through various computational models, the most successful of which have been supervised encoder-decoder models, which treat the problem of predicting protoforms given sets of reflexes as a sequence-to-sequence problem. We argue that this framework ignores one of the most important aspects of the comparative method: not only should protoforms be inferable from cognate sets (sets of related reflexes) but the reflexes should also be inferable from the protoforms. Leveraging another line of research—reflex prediction—we propose a system in which candidate protoforms from a reconstruction model are reranked by a reflex prediction model¹. We show that this more complete implementation of the comparative method allows us to surpass state-of-the-art protoform reconstruction methods on three of four Chinese and Romance datasets.

Keywords: historical reconstruction, historical linguistics, phonology, reranking

1. Introduction

Historical linguistics provides a window into the human past, the diversification of and interactions between human populations, as well as the mechanisms through which languages change over time. Perhaps the most enduring theoretical and methodological contribution of historical linguistics is the comparative method, by which protolanguages-putative ancestors of families of languages—can be reconstructed (Anttila, 1989; Campbell, 2021). In the comparative method, cognate sets-groups of words believed to have descended from the same ancestral word-are compared in order to infer the corresponding ancestral words (protoforms). These reconstructions are chosen to maximize the regularity of the mapping from reconstructions to reflexes (daughter forms) and minimize the phonetic distance between reconstructions and their reflexes. The assumption is that the historical changes that affect the sounds in words are largely regular such that almost all reflexes in a language can be derived deterministically from the protoforms given a series of sound changes.

This method is difficult to employ in practice, in no small part because datasets can be very large. To deal with the cognitive burden of historical comparison, computational methods have been proposed to assist linguists in this endeavor. However, with a few exceptions (Bouchard-Côté et al., 2013; He et al., 2023; Arora et al., 2023), the most

¹Our code is available at https://github.com/ cmu-llab/reranked-reconstruction. successful comparative reconstruction models have treated this task as a fairly generic sequenceto-sequence transduction task, in essence translating sets of reflexes (cognate sets) into protoforms (Meloni et al., 2021; Chang et al., 2022; Fourrier, 2022; Kim et al., 2023; Cui et al., 2022). This ignores an important aspect of the comparative method in that it does not constrain the protoforms so that they can be deterministically translated back into each of the reflexes.

In this paper, we propose a multi-model reconstruction system that improves its reconstructions via reflex prediction-the task of predicting the reflexes given a protoform. Our system consists of a beam search-enabled sequence-to-sequence reconstruction model and a sequence-to-sequence reflex prediction model that serves as a reranker. The reflex prediction component can often provide valuable information that may not have been captured by the reconstruction model's probability distribution, thereby addressing certain reconstruction errors. We find that our linguisticallymotivated method can address some errors made by existing techniques. Figure 1 shows an example where reranking reconstruction candidates according to reflex prediction accuracy compensates for the erroneous probability ranking of the reconstruction model, leading to a correct reconstruction.

As reflex prediction prior art on our datasets of interest is limited, we test various neural reflex prediction models. We then combine pre-trained reconstruction models and reflex prediction models into reconstruction systems. We perform ablation studies and post hoc error analysis to examine

Bean	m Search Reflex Prediction (based on protoform candidates)						Reranking Result			esult						
rank	\hat{p}_i^{bs}	mi		Cantonese	Gan	Hakka	Jin	Mandarin	Hokkien	Wu	Xiang	ri		rank	$\hat{p}_i^{\mathbf{rk}}$	Si
0	pjet入	-0.1114		pi:t+	pįɛt٦	pi̯et⊣	pįə?⊣	рі́ε∖	pi̯ɛt⊦	pjɪ? ٦	pie₁	0.2500	\searrow 7	0	pit入	0.5995
1	pet入	-0.2711		pi:t+	pįɛt٦	pi̯et⊣	pįə?∔	pjel	pi̯ɛt⊣	p <u>i</u> 171	pię≀	0.2500	\checkmark	1	pjet入	0.2036
2	pit入	-0.5030		pet1	pit1	pit⊦	pįə?∔	pi∖	pit⊦	p <u>i</u> 171	pi≀	0.8750		2	pet入	0.0439
3	рер入	-1.5533		pi:p+	pįɛt٦	pi̯ap⊣	pįə?∔	pjel	рі́ар⊣	p <u>i</u> 171	pię≀	0.2500	\longrightarrow	3	рер入	-1.2383
4	pij去	-1.6329		peí⊦	pi1	pi٦	pi11	pi∖	pi≀	pi1	pi11	0.1250	\longrightarrow	4	pij去	-1.4754
				pi:t1	pit1	pit⊦	pįə?⊣	piN	piti	pj1?1	pi≀	-				

Figure 1: A scenario in which using beam search on Meloni et al. (2021)'s sequence-to-sequence GRU reconstruction model incorrectly predicts $pjet\lambda$ as the most likely reconstruction for the $\not{a} pit\lambda$ 'must' cognate set (in the WikiHan test set). The reflex prediction model could only infer 2 (bold) of the 8 reflexes from the incorrect reconstruction $pjet\lambda$, but correctly infers 7 of the 8 reflexes from the third candidate $pit\lambda$. Our reflex prediction-based reranked reconstruction system makes score adjustments that lead to the correct reranked protoform prediction $pit\lambda$. The last row in the reflex prediction table provides reference reflexes. Bold: correct protoform or reflexe; *i*: ranking index; \hat{p}_i^{bs} : beam search protoform; r_i : reranker score; \hat{p}_i^{rk} : reranker protoform candidate; s_i : adjusted score.

the effectiveness of such systems. Our reranked reconstruction system outperforms state-of-theart neural reconstruction approaches on Meloni et al. (2021)'s Romance datasets and Chang et al. (2022)'s Sinitic dataset WikiHan. Our contributions include:

- Proposing a multi-model, reranking-driven reconstruction system that achieves state-ofthe-art reconstruction results on both Romance and Sinitic datasets
- 2. Adapting and examining existing architectures, as well as modified variants, for reflex prediction on Romance and Sinitic languages
- 3. Performing phonologically-informed analysis of the reflex prediction model and its interactions with the reconstruction model in a reranking system
- 4. Providing a fast implementation of the reconstruction system with vectorized beam search and reranking

2. Related Work

Word form-related tasks in computational historical linguistics include reconstruction, reflex prediction, and cognate prediction², as summarized in Figure 2.

2.1. Reconstruction

Computational reconstruction of proto-languages was proposed as early as the 1960s (Durham and Rogers, 1969). Bouchard-Côté et al. (2013) used sound change probabilistic models along with a Monte Carlo inference algorithm to automate



Figure 2: Three word-form-related tasks in historical linguistics, exemplified by the $emtiliamediate{mtext} luk\lambda$ 'wheel' cognate set from WikiHan.

protoform reconstruction, but their method relies on a phylogenetic tree. List et al. (2022a) proposed sequence comparison and phonetic alignment for reconstruction, but this did not perform well on either the WikiHan or Romance datasets (Cui et al., 2022; Kim et al., 2023). Ciobanu and Dinu (2018) and Ciobanu et al. (2020) used conditional random fields to automate reconstruction by labeling each position in the daughter sequence with a protoform token. Meloni et al. (2021) formulated protoform reconstruction as a sequence-to-sequence task and used an encoder-decoder GRU model to perform Latin reconstruction, setting a baseline for future neuralbased reconstruction methods. Fourrier (2022) compared RNNs and Transformers for protoform reconstruction, noting that encoder-decoder architectures can encode phonetic features in the reflexes into an informative latent space from which the decoder can derive the protoform. Extending Meloni et al. (2021)'s work, Kim et al. (2023) proposed using a Transformer-based encoderdecoder architecture with language embedding for protoform reconstruction, achieving state-ofthe-art on Meloni et al. (2021)'s Romance dataset and Hou (2004)'s Sinitic dataset. Very recently, Akavarapu and Bhattacharya (2023) used an MSA Transformer (originally proposed as a protein language model with multiple sequence alignments

²These terms are sometimes confused. Because we need a distinction here, we categorize them using Arora et al. (2023)'s definitions. When only relatedness but not ancestry is concerned, the protoform is sometimes treated as part of the cognate set in the literature.

as inputs (Rao et al., 2021)) pretrained for cognate prediction to perform protoform reconstruction on automatically aligned cognate sets.

2.2. Reflex prediction

Reflex prediction involves modeling the phonological or morphological changes needed to derive reflexes from protoforms. It corresponds to recreating the evolutionary process of languages in historical linguistic studies. Marr and Mortensen (2020, 2023) developed a rule-based Latin-to-French reflex prediction model that predicted reflexes at five different stages in the history of French. Bodt and List (2022) used a semiautomatic method to predict reflexes in Western Kho-Bwa via automatic alignment and identification of sound correspondences on manually annotated cognate sets. Paralleling the use of sequence-to-sequence techniques in reconstruction, Cathcart and Rama (2020) pioneered neural reflex prediction with an LSTM encoder-decoder model that predicts Indo-Aryan languages from Old Indo-Aryan. Recently, Arora et al. (2023) introduced a new South Asian languages dataset and replicated Cathcart and Rama (2020)'s reflex prediction experiments on the dataset with both GRU and Transformer encoder-decoder models. To the best of our knowledge, no work has examined neural reflex prediction with Romance and Sinitic languages.

Protoform with prepended target language token	Predicted reflex in target language			
<cantonese>luk入 -</cantonese>		lʊk1	(Cantonese)	
<mandarin>luk入 —</mandarin>		lu۱	(Mandarin)	
<hokkien>luk入 -</hokkien>		lok1	(Hokkien)	
<wu>luk入 一</wu>		۲,20	(Wu)	

Figure 3: A reflex prediction model aims to derive the correct reflexes based on a protoform sequence tagged by the target daughter languages. A forward pass on the model involves only one daughter language.

2.3. Cognate prediction

Nitschke (2021) used neural machine translation techniques to predict missing reflexes in Ro-

mance cognate sets. The SIGTYP 2022 shared task on the prediction of conjugate reflexes called for efforts to develop cognate prediction systems and evaluated submissions on numerous language families (List et al., 2022b). Interestingly, a CNN model by Kirov et al. (2022) resembling image-inpainting (Mockingbird-I1) performed the best overall (List et al., 2022b). By treating phonemes as pixels, reflexes as rows of pixels, and cognate sets as stacked rows forming an image, Mockingbird-I1 recovers the missing rows with convolution and deconvolution networks. Cui et al. (2022) found that Mockingbird-I1 can be used to augment a reconstruction dataset, improving the model's stability while training. Although we do not perform cognate prediction in this paper, we test our methods on Cui et al. (2022)'s augmented WikiHan dataset (WikiHanaug), which will help answer the question of how well reflex prediction, cognate prediction, and reconstruction can combine to form a more effective reconstruction workflow.

3. Methods

3.1. Datasets

Romance Datasets: We use Meloni et al. (2021)'s dataset consisting of both IPA (International Phonetic Alphabet) and orthographic forms. The IPA form (Rom-phon) represents words' pronunciation in phonemes, while the orthographic form (Rom-orth) represents the words as they are spelled out in writing. To compare with the state-of-the-art reconstruction model on the Romance datasets, we match Kim et al. (2023)'s preprocessing and splits.

Sinitic Datasets: We use both Hóu (2004)'s dataset compiled by Kim et al. (2023) and Chang et al. (2022)'s WikiHan dataset³. Both datasets contain phonetic representations of Middle Chinese and its descendants. We also test an augmented version of WikiHan (WikiHan-aug) created by Cui et al. (2022), which uses the cognate prediction model Mockingbird-I1 to fill in missing

³Although the full dataset consists of 21,227 cognate sets, cognate sets with less than 3 reflexes are ignored to match previous work.

Dataset	Cognate sets	# varieties	Ancestor language
WikiHan (Chang et al., 2022)	5,165	8	Middle Chinese
WikiHan-aug (Cui et al., 2022)	8,780	8	Middle Chinese
Hóu (Hóu, 2004)	804	39	Middle Chinese
Rom-phon (Meloni et al., 2021; Ciobanu and Dinu, 2018)	8,703	5	Latin
Rom-orth (Meloni et al., 2021; Ciobanu and Dinu, 2018)	8,631	5	Latin

Table 1: Overview of the datasets with their respective number of daughter languages (# varieties) and the number of cognate sets used in the experiments.

daughter entries in the train set. Protoform labels for the Sinitic datasets are based on Baxter (2014)'s reconstructions of Middle Chinese. We match Chang et al. (2022)'s splits for WikiHan and Kim et al. (2023)'s splits for Hóu (2004).

3.2. Reflex Prediction Models

As there is limited prior work on neural reflex prediction with our datasets of interest, we adapt reflex prediction models previously used for other datasets, along with reconstruction models previously used for the current datasets. Figure 3 shows the reflex prediction task as a sequence-tosequence transduction task from the protoform with target language tokens prepended to the beginning—to the reflex in each specified language.

One notable difference between reflex prediction and translation is that, instead of decoding to one target language, the reflex prediction model needs to decode into multiple possible target languages. As a baseline, we attempt various architectural modifications to Meloni et al. (2021)'s unidirectional encoder-decoder GRU model to accommodate multiple target languages-including multi-layer bidirectional encoding, target language embedding during decode similar to Meloni et al. (2021)'s encoder, onehot vector target language prompting to the decoder's classifier, target-language-specific connections in the decoder's classifier network, and support for VAE-style latent space used by Cui et al. (2022)⁴ to decode from the same source to multiple daughters-all of which are tuned as hyperparameters.

Additionally, we adapt Kim et al. (2023)'s Transformer reconstruction model⁵ for reflex prediction and test Arora et al. (2023)'s GRU and Transformer reflex prediction models on our datasets of interest⁶. We implement batched training and inference for all the adapted models but retain the original architecture.

⁵The major difference between Kim et al. (2023)'s model and a standard Transformer encoder-decoder model is the addition of language embeddings for input daughter sequences. In the adapted version for reflex prediction, input language embedding serves little purpose (due to a singular input language) and is thus disabled, making it technically very similar to Arora et al. (2023)'s Transformer model in architecture.

⁶Since Cathcart and Rama (2020)'s model requires additional data such as part of speech for semantic embedding, their reflex prediction model is not fully replicable on our datasets of interest.

3.3. Beam Search Reconstruction Model

Since beam search is needed in the reranked reconstruction process and no prior work uses beam search in neural reconstruction, we implement beam search on top of Meloni et al. (2021)'s GRU model (GRU-BS). To isolate the effects of reranking, the architecture of the GRU is kept the same as Meloni et al. (2021), consisting of language and token embeddings, a single-layer unidirectional encoder-decoder GRU model, and a multi-layer perceptron classifier⁷. We tune GRU-BS separately to optimize for performance with beam search.

Before being passed into the reconstruction model, reflexes in a cognate set are concatenated into one long sequence, with separators between the reflexes and language tokens to identify each reflex. An example input is as follows:

*Cantonese:meil*Mandarin:meil*Wu:mel*

3.4. Reranked Reconstruction System

In a simple beam search system, the candidate sequences are ranked by their length-normalized log probability, and the candidate with the highest normalized log probability is returned. We propose to enhance this ranking using a reflex prediction model that estimates phonetic naturalness when inferring the reflexes from each candidate protoform, as detailed in Algorithm 1. Given protoform candidates predicted by GRU-BS, we compute the proportion of reflexes correctly derived from each candidate as score adjustment. The candidates are rescored by summing the normalized log probability and the score adjustment, scaled by a score adjustment weight λ . The candidate with the highest adjusted score is chosen as the final prediction.

3.5. Evaluation Criteria

To enable cross-task comparisons, we employ established reconstruction metrics for both reflex prediction and reranked reconstruction experiments, including token edit distance (TED), the number of token insertions, deletions, or substitutions between predictions and targets (Levenshtein et al., 1966); token error rate (TER), a length-normalized edit distance (Cui et al., 2022); accuracy (ACC), the percentage of exactly correct predictions; feature error rate (FER), a measure of phonological edit distance by PanPhon (Mortensen et al., 2016); and B-Cubed F Score (BCFS), a measure of structural similarity between predictions and targets (Amigó et al., 2009; List,

⁴Cui et al. (2022)'s report proposes a reconstruction model that learns a representation of the cognate set with a Variational Autoencoder (VAE) on the reflexes, reconstructing both the reflexes and the protoforms from the same latent space.

⁷We use Chang et al. (2022)'s PyTorch reimplementation obtained from github.com/cmu-llab/ meloni-2021-reimplementation.

Algorithm 1 Sequential representation of our reranked reconstruction algorithm **Require:** $d_1, d_2, ..., d_n$ = reflexes in daughter languages $D_1, D_2, ..., D_n$ from a cognate set with *n* reflexes **Require:** f_{θ_f} = a beam search-enabled reconstruction model with pre-trained parameters θ_f **Require:** g_{θ_q} = a reflex prediction model with pre-trained parameters θ_q **Require:** k = beam size for predicting candidate reconstructions on $f_{\theta_{\ell}}$ **Require:** a = length normalization constant **Require:** λ = score adjustment weight $D \leftarrow ``*"D_1``:"d_1``*"D_2``:"d_2``*" \cdots ``*"D_n``:"d_n``*"$ concatenate reflex sequences into a long sequence, with language labels and separators in between $C = [(\hat{p}_1, m_1), (\hat{p}_2, m_2), \dots, (\hat{p}_l, m_l)] \leftarrow f_{\theta_f}(D, k, a) \triangleright$ beam search with beam size k to obtain a list of $l \leq k$ candidate protoform predictions \hat{p}_i with their normalized log probabilities $m_i = \frac{\log P(\hat{\rho}_i | D)}{|\hat{\rho}_i|^{\alpha}}$ assigned by f_{θ_f} for $1 \le i \le I$ $C' \leftarrow []$ initialize reranked candidate list for (\hat{p}_i, m_i) in C do *a* ← 0 counter for the number of correctly derived daughters **for** *j* ← 1 to *n* **do** $\hat{p}'_i \leftarrow D_i \hat{p}_i$ > prepend the *j*-th daughter language token to the candidate protoform $\hat{d}_{ij} \leftarrow g_{ heta_g}(\hat{p}'_i)$ if $\hat{d}_{ij} = d_j$ then > predict the reflex in the *i*-th daughter language based on the *i*-th candidate $a \leftarrow a + 1$ > increment counter if predicted reflex is correct \triangleright use the accuracy of reflex predictions as the reranker score r_i $r_i \leftarrow a/n$ $s_i \leftarrow m_i + \lambda r_i$ \triangleright calculate the adjusted score s_i for the *i*-th candidate $C' \leftarrow C' + [(\hat{p}_i, \mathbf{s}_i)]$ > append entry with adjusted score to reranked candidate list $C' \leftarrow C'$ sorted by descending s_i return C'[0]return the candidate with the highest adjusted score

2019). Tokens are phonemes in all datasets with the exception of the character-level Rom-orth dataset. Consequently, FER cannot be reliably calculated for Rom-orth.

3.6. Experiments

Hyperparameters: We tune hyperparameters using WandB (Biewald, 2020) except for models already tested by Kim et al. (2023): Meloni et al. (2021)'s GRU reconstruction model and Kim et al. (2023)'s Transformer reconstruction model on Rom-phon, Rom-orth, and Hóu. We use Bayesian search with 100 total runs for the best validation phoneme edit distance, validated every 3 epochs and with early stopping. We keep a constant beam size of 5 when tuning GRU-BS to balance computation cost and effectiveness.

Reflex Prediction Experiments: First, we test the reflex prediction capability of the four aforementioned reflex prediction models. For each model on each dataset, we perform 20 runs with random seeds (same hyperparameters). We select the best-performing reflex prediction model from each architecture (GRU or Transformer) as reranker models in reranked reconstruction experiments.

Baseline Reconstruction Data: We use both Meloni et al. (2021)'s GRU and Kim et al. (2023) Transformer models as baselines. For datasets

present in Kim et al. (2023)'s work, we perform additional runs to obtain 20 runs in total (on top of Kim et al. (2023)'s 10 checkpoints).

Reranked Reconstruction Experiments: Each reranking experiment involves a pre-trained GRU-BS reconstruction model and a reflex prediction model acting as a reranker, forming a reranked reconstruction system. We select Arora et al. (2023)'s Transformer and the baseline GRU reflex prediction model as rerankers due to their higher performance among their respective architecture. For each reranked reconstruction system, we tune two additional hyperparameters: the beam size of GRU-BS k, and the score adjustment weight λ . We perform a grid search on the hyperparameters for best validation accuracy⁸. The search results across 20 pairs of pre-trained reconstruction and reflex prediction models are averaged (rounding k to the nearest integer) to obtain the final hyperparameters for evaluations on the test set.

Statistical Analysis: Considering a small sample size and unknown distribution, we use Wilcoxon Rank-Sum test (Wilcoxon, 1992) with a = 0.01 and Bootstrap test (Efron and Tibshirani, 1994) with 99% confidence interval for the mean difference between models or reconstruction systems.

⁸The search range for *k* and λ are [2, 10] with resolution 2 and [0.3, 4.2] with resolution 0.3, respectively.

We consider results to be significant if both tests indicate significance.

Ablation Studies: Our reranked reconstruction system extends Meloni et al. (2021)'s model by both beam search and reranking. To isolate the effect of reranking, we remove the reranker to obtain the performance of GRU-BS before reranking (with beam size no larger than when used in reranking) and test for differences in performance between GRU-BS with and without reranking.

Correlation Experiments: We select the worstperforming reconstruction model (by accuracy at k = 5) and rerank it with all the pre-trained reflex prediction models (20 GRU and 20 Transformer) using the same reranking hyperparameters obtained from grid search, effectively varying the reranker with controlled reconstruction and reranking hyperparameters. We then examine the correlation between reflex prediction performance and reranked reconstruction performance.

4. Results and Discussion

4.1. Reflex prediction Results

Table 2 shows the average performance of the four reflex prediction models. We found statistically significant evidence that Arora et al. (2023)'s Transformer performs the best on all metrics for WikiHan and WikiHan-aug, Arora et al. (2023)'s Transformer performs the best only on ACC for Rom-orth, and Kim et al. (2023)'s Transformer performs the best on TER, TED, and BCFS for Romphon. We find no evidence that the top-2 performing models are statistically different for the remaining metrics or datasets. Among GRU models, the

baseline performs better than Arora et al. (2023)'s GRU model across all datasets.

On all datasets except Rom-phon, we obtain overall better performance at reflex prediction than reconstruction—consistent with the hypothesis that learning regular sound changes is easier in the forward direction. It is possible that reflex prediction performs worse than reconstruction on Rom-orth due to its non-phonetic nature, obscuring the environments for sound changes.

In reranking experiments, we select the bestperforming model for each architecture: Arora et al. (2023)'s Transformer and the baseline GRU⁹.

4.2. Reranked Reconstruction Results

As shown in Table 3, our reranking system performs significantly better than both Meloni et al. (2021) and Kim et al. (2023) on all datasets except Hóu, for which it performs better on some metrics. We notice a high variance in both reflex prediction and reconstruction results for Hóu, possibly due to its small test set. Finally, we find no statistical difference between using a GRU or a Transformer as a reranker, despite evidence that Transformers outperform GRUs on reflex prediction.

Cui et al. (2022) previously found no evidence that data augmentation helps improve reconstruction. However, our result on WikiHan-aug suggests that cognate set augmentation contributes to both reflex prediction and reranked reconstruc-

⁹We also tested reranking using Kim et al. (2023)'s Transformer model, but found no statistical difference in performance.

Dataset	Model	ACC%↑	TED↓	TER↓	FER↓	BCFS ↑
WikiHan	GRU (baseline)	66.43%	0.5244	0.1547	0.0400	0.7394
	GRU (Arora et al., 2023)	64.45%	0.5558	0.1640	0.0428	0.7260
	Transformer (Kim et al., 2023)	66.39%	0.5302	0.1564	0.0406	0.7370
	Transformer (Arora et al., 2023)	67.64%	0.5128	0.1513	0.0390	0.7445
WikiHan-aug	GRU (baseline)	68.11%	0.5007	0.1477	0.0380	0.7495
	GRU (Arora et al., 2023)	66.94%	0.5159	0.1522	0.0391	0.7430
	Transformer (Kim et al., 2023)	68.96%	0.4889	0.1442	0.0371	0.7551
	Transformer (Arora et al., 2023)	69.37%	0.4826	0.1424	0.0363	0.7572
Hóu	GRU (baseline)	51.72%	0.7777	0.2037	0.0488	0.6783
	GRU (Arora et al., 2023)	49.26%	0.8266	0.2166	0.0528	0.6622
	Transformer (Kim et al., 2023)	55.46%	0.7576	0.1985	0.0494	0.6882
	Transformer (Arora et al., 2023)	55.60%	0.7520	0.1970	0.0485	0.6892
Rom-phon	GRU (baseline)	63.85%	0.7439	0.1014	0.0426	0.8361
	GRU (Arora et al., 2023)	48.28%	1.3257	0.1808	0.0930	0.7567
	Transformer (Kim et al., 2023)	64.19%	0.7349	0.1002	0.0427	0.8380
	Transformer (Arora et al., 2023)	63.96%	0.7442	0.1015	0.0428	0.8361
Rom-orth	GRU (baseline)	64.58%	0.7301	0.0967	-	0.8465
	GRU (Arora et al., 2023)	57.92%	0.8741	0.1158	-	0.8218
	Transformer (Kim et al., 2023)	64.80%	0.7258	0.0961	-	0.8478
	Transformer (Arora et al., 2023)	65.20 %	0.7247	0.0960	-	0.8476

Table 2: Average performance of the reflex prediction models across 20 runs, with bold indicating the best-performing model for each metric.

Dataset	Reconstruction System	ACC%↑	TED↓	TER↓	FER↓	BCFS ↑
WikiHan	GRU (Meloni et al., 2021)	55.58%	0.7360	0.1724	0.0686	0.7426
	Trans (Kim et al., 2023)	54.62%	0.7453	0.1746	0.0696	0.7393
	GRU-BS (<i>k</i> = 10)	54.88%	0.7507	0.1758	0.0701	0.7364
	GRU-BS ($k \le 10$) + GRU Reranker	57.14%*†	0.7045*†	0.1650*†	0.0661*†	0.7515*†
	GRU-BS ($k \le 10$) + Trans. Reranker	57.26% *†	0.7029 *†	0.1646 *†	0.0658 *†	0.7520 *†
WikiHan-aug	GRU (Meloni et al., 2021)	54.73%	0.7574	0.1774	0.0689	0.7346
	Trans (Kim et al., 2023)	55.82%	0.7317	0.1714	0.0661	0.7416
	GRU-BS (<i>k</i> = 10)	56.64%*	0.7214	0.1690	0.0658	0.7454
	GRU-BS ($k \le 10$) + GRU Reranker	58.58% *†	0.6822 *†	0.1598 *†	0.0628*†	0.7579*†
	GRU-BS ($k \le 10$) + Trans. Reranker	58.58%*†	0.6840*†	0.1602*†	0.0626 *†	0.7575*†
Hóu	GRU (Meloni et al., 2021)	34.63%	1.0916	0.2479	0.0914	0.6697
	Trans (Kim et al., 2023)	39.01%	0.9904	0.2233	0.0875	0.6955
	GRU-BS (<i>k</i> = 10)	37.36%	1.0382	0.2328	0.0917	0.6974
	GRU-BS ($k \le 10$) + GRU Reranker	40.50%†	0.9727†	0.2181†	0.0867†	0.7130*†
	GRU-BS ($k \le 10$) + Trans. Reranker	42.08% *†	0.9503 *†	0.2131 *†	0.0850†	0.7170 *†
Rom-phon	GRU (Meloni et al., 2021)	51.92%	0.9775	0.1244	0.0390	0.8275
	Trans (Kim et al., 2023)	53.04%	0.9050	0.1148	0.0377	0.8417
	GRU-BS (<i>k</i> = 10)	52.63%	0.9125	0.1018*	0.0353*	0.8402
	GRU-BS ($k \le 10$) + GRU Reranker	53.95% *†	0.8775*†	0.0979*†	0.0336*†	0.8460*†
	GRU-BS ($k \le 10$) + Trans. Reranker	53.85%*†	0.8765 *†	0.0978 *†	0.0333 *†	0.8461 *†
Rom-orth	GRU (Meloni et al., 2021)	69.41%	0.6004	0.0781	-	0.8906
	Trans (Kim et al., 2023)	71.05%	0.5636	0.0734	-	0.8981
	GRU-BS (<i>k</i> = 10)	71.09%	0.5531	0.0617*	-	0.8990
	GRU-BS ($k \le 10$) + GRU Reranker	72.60% *†	0.5237 *†	0.0584 *†	-	0.9045 *†
	GRU-BS ($k \le 10$) + Trans. Reranker	72.50%*†	0.5246*†	0.0585*†	-	0.9044*†

Table 3: Evaluation of reconstruction systems, including baselines, GRU with beam search (GRU-BS), and GRU-BS with reranking, averaged across 20 runs. Bold indicates the best-performing system for each metric, asterisks indicate statistically better performance than both baseline models (Meloni et al. (2021)'s GRU and Kim et al. (2023)'s Transformer), and daggers indicate that a reranking system performs statistically better than its beam search counterpart.

Dataset	Reranker	ACC	TED	TER	FER	BCFS
WikiHan	GRU Reranker	0.2771	0.3647	0.3647	0.1845	0.3639
	Trans. Reranker	0.3860	0.0987	0.0987	0.3132	0.0276
WikiHan-aug	GRU Reranker	0.3830	0.4179	0.4179	0.4829	0.3573
	Trans. Reranker	0.1849	-0.0435	-0.0435	-0.0655	-0.0310
Hóu	GRU Reranker	0.3735	0.1236	0.1236	0.4278	0.0173
	Trans. Reranker	0.5432	0.2742	0.2742	0.2782	0.3184
Rom-phon	GRU Reranker	-0.2207	-0.0115	-0.0115	-0.0373	0.0336
	Trans. Reranker	-0.0706	-0.0181	-0.0181	-0.0866	0.0421
Rom-orth	GRU Reranker	0.2531	0.4044	0.4044	-	0.4459
	Trans. Reranker	0.1639	0.1123	0.1123	-	0.1035

Table 4: Correlation coefficients between rerankers' reflex prediction performance and reranked reconstruction performance. The cells are color-coded by sign and strength, with red for positive correlation coefficients and blue for negative correlation coefficients.

tion performance, bringing WikiHan reconstruction accuracy to 3% above previous work.

4.3. Correlation Test Results

Correlation analysis reveals a mostly positive correlation between rerankers' reflex prediction performance and the corresponding reranking system's reconstruction performance, except on the Rom-phon dataset (see Table 4). Although statistical significance is unclear due to small sample sizes, evidence suggests that the performance of the reranker could play an important role in a reranked reconstruction system.

4.4. Ablation Studies

While GRU-BS alone (without a reranker) outperforms baseline models on some occasions, GRU-BS with a reranker performs statistically better than GRU-BS alone for all datasets and metrics, as indicated in Table 3. Even though beam search is commonly regarded as a powerful method in sequence-to-sequence tasks, its ability in a protolanguage reconstruction setting is still limited compared to reranking, where modeling reflex prediction in addition to reconstruction proves more informative.

4.5. Reranking Error Analysis

To gain insights into the reranker's behavior, we conduct error analyses on the topperforming reranking system (GRU-BS + Transformer Reranker) by randomly selecting one of the 20 runs. We denote the ranks from beam search and reranking by $r_{\rm bs}$ and $r_{\rm rk}$ respectively (better score has lower rank), and categorized the reranker's behavior into four distinct categories:

- **Improved** (*r*_{rk} < *r*_{bs}): reranker assigns a more favorable rank to the target protoform.
- **Unchanged** ($r_{rk} = r_{bs}$): reranker does not alter the rank of the target protoform.
- Worsened ($r_{\rm rk} > r_{\rm bs}$): reranker assigns a less desirable rank to the target protoform.
- Not-in: the target protoform is not predicted as a candidate by beam search and is thus not seen by the reranker. This category is not included in analyses that require the target protoform to be processed by the reranker.

Table 5 shows the distribution of the reranker's behavior among the four categories. On every dataset, the reranker improves the ranking of the target protoform more often than worsens it.

We observe that, compared to the target protoform, the incorrectly predicted protoform often exhibits greater phonetic similarity, measured by both token edit distance and feature edit distance, to the reflexes¹⁰ (see Table 6). It is likely that the reflex prediction models find it easier to derive correct reflexes from predicted protoforms that are more similar to the reflexes, potentially making it challenging for the reranker to improve the ranking of the target protoforms less similar to the reflexes.

Furthermore, certain sound combinations in Wiki-Han's Middle Chinese forms, such as *ju*, *je*, and *xwo*, are absent in the daughter languages included in the dataset (see Table 7 for some examples). This highlights a notable challenge of computational reconstruction—recovering phonemes lost during language evolution—which likely require solutions other than reranking.

Finally, we observe that the reranker model has the highest overall error rate when predicting Hokkien reflexes compared to other daughter languages on WikiHan (see Table 8), despite Hokkien having the third most training examples. A possible explanation is Karlgren (1974)'s hypothetical subgrouping of Sinitic in which Hokkien is a descendent of a sister of Middle Chinese rather than Middle Chinese itself.

¹⁰Because the phonetic values of Middle Chinese tones are unknown (WikiHan represents them tones with the four abstract tone characters from Tang Dynasty rhyme books), we exclude tones when calculating D_T and D_F in Table 6 and in the case studies in Table 7.

Dataset	Improved	Worsened	Unchanged	Not-in	Total	Improved/Changed (%)
WikiHan	84 (8.13%)	32 (3.10%)	755 (73.09%)	162 (15.68%)	1033	72.41%
WikiHan-aug	88 (8.52%)	23 (2.23%)	791 (76.57%)	131 (12.68%)	1033	79.28%
Hóu	26 (16.15%)	15 (9.32%)	88 (54.66%)	32 (19.88%)	161	63.41%
Rom-phon	109 (6.21%)	61 (3.48%)	1198 (68.30%)	386 (22.01%)	1754	64.12%
Rom-orth	75 (4.29%)	23 (1.32%)	1367 (78.16%)	284 (16.24%)	1749	76.53%

Table 5: The distribution of reranker behavior categorization on the test set (left), based on a randomly sampled run for each dataset, as well as the corresponding rate of ranking improvement among instances with changed (i.e. improved or worsened) ranking (right).

Dataset	Category	$D_T(\hat{\rho}, R) < D_T(\rho, R)$	$D_F(\hat{p},R) < D_F(p,R)$
		(<i>R</i> more similar to \hat{p} than p by D_T)	(<i>R</i> more similar to \hat{p} than <i>p</i> by D_F)
WikiHan	Worsened	37.50%	53.12%
	Unchanged	35.56%	43.33%
	Improved	28.30%	32.08%
Rom-phon	Worsened	60.66%	62.30%
	Unchanged	47.21%	51.48%
	Improved	47.17%	49.06%

Table 6: Comparison between the phonetic similarity between the reflexes *R* and the predicted protoform \hat{p} versus the target protoform *p* for each category of the reranker's behavior among reconstruction errors. Similarity are measured by normalized token edit distance (*D*_T) and normalized feature edit distance (*D*_F). The table presents percentages of entries in each category where the predicted protoforms exhibit greater phonetic similarity to their modern reflexes than the target protoforms according to each similarity metric, with bold indicating the highest percentage.

	Category	Worsene	d	Unchanged		Improved	
Dataset		Proto	Prôto	Proto	Prôto	Proto	Prôto
WikiHan	Middle Chinese	mjuk™	muk™	tʰ ja ŋ	te⁺a ŋ	ts¹je k	ts e k
	Cantonese	m_ʊk	mʊk	ts ^h ₋ɔːŋ	fs ^h ɔːŋ	t̂s ₋ı k	ts I k
	Hakka	m∟uk	mu <mark>k</mark>			ts _a k	fs <mark>a</mark> k
	Mandarin	m_u_	mu	<code>fʂʰ_a ŋ</code>	<mark>f</mark> şʰa ŋ	fa ^h li L	fa ^h i 👅
	Hokkien	b <mark>u</mark> ok	bok	fe ^h _ioŋ	fɕʰi̯ɔŋ	ta ^h _iək	fehiək
Rom-phon	Latin	ast ^h ma	as_ma	f_ɛrɪtatɛm	f_ɛritam	teksere	tısserε
	Romanian	as <mark>t</mark> mə	as <mark>t</mark> mə			tsese	tsese
	French	as. m.	as_m_	fjɛʁ <mark>uteuuu</mark>	fjɛʁ <mark>_te</mark> _	ti_se	t <mark>i</mark> _se
	Italian	a <mark>z</mark> . ma	az∟ma	f_erita	f_erita_	tɛssere	t <mark>e</mark> ssere
	Spanish	as <mark>.</mark> ma	as_ma			tex_er_	t <mark>ex</mark> lerl
	Portuguese	az <mark>.</mark> me	a <mark>z</mark> _me		_	t <mark>i_seı</mark>	t <mark>i</mark> _se <mark>ı</mark>
			Color	key: 🔳 substitu	ution 🔳 ins	ertion	(_) deletion

Table 7: Instances in each category where the predicted protoform is phonetically closer to its reflexes than the target protoform by both D_T and D_F , selected from the WikiHan (top) and Rom-phon (bottom) test sets. The **Proto** and **Proto** columns show edits from the target protoform and the predicted protoform to the reflexes, respectively. Words in each column are manually aligned to reflect edits, with '_' indicating an empty position in the multi-sequence alignment in the case of deletion or insertion. Unavailable reflexes are not shown on the table, and languages without available reflexes (Gan, Jin, Wu, and Xiang) are omitted. Differences in edits between the predicted and target protoforms are shaded. The selected entries are 睦 *mjuk*^w 'friendly', 昶 *t^hjaŋ* 'long daytime', 磧, *tshjek* 'gravel' (WikiHan), *asthma* 'asthma', *feritatem* 'ferocity', and *texere* 'to weave' (Rom-phon).

Category	Cantonese	Gan	Hakka	Jin	Mandarin	Hokkien	Wu	Xiang
Not-in	75.31%	72.73%	70.83%	83.33%	72.22%	82.05%	57.69%	82.14%
Worsened	59.38%	75.00%	40.00%	71.43%	68.75%	68.75%	33.33%	60.00%
Unchanged	24.44%	21.21%	42.16%	15.79%	27.22%	46.86%	15.66%	41.67%
Improved	38.46%	25.00%	54.05%	13.33%	18.87%	50.00%	37.93%	41.18%
Overall	48.12%	40.00%	53.10%	42.22%	46.37%	62.29%	32.95%	55.81%

Table 8: Reranker's reflex prediction error rates among reranked reconstruction error entries (when predicting reflexes from the target protoform) for each daughter language in the WikiHan dataset given each reranker behavior category, obtained from a randomly selected run.

5. Conclusion

Ironically, many efforts to automate protolanguage reconstruction with neural models have thus far treated reconstruction as a sequenceto-sequence task, disregarding the comparative method's constraint that reflexes should be inferable from the reconstructions. Our reranked reconstruction system provides an elegant way to replicate the synergy between reconstruction and reflex prediction in the comparative method, yielding results that surpass existing methods—a vindication of the idea that designing reconstruction systems with the comparative method in mind can be more powerful than relying solely on sequenceto-sequence techniques.

Though our approach yields better reconstruction performance, it is left to future work to address some of the challenges identified in the present work, such as a reconstruction system's tendency to produce reconstructions relatively similar to the reflexes. In the bigger picture, reranking is but one way to bring together multiple tasks in historical linguistics, and arguably a complicated one due to its multi-step training and tuning process. Future research, therefore, could also explore approaches to integrate reconstruction and reflex prediction into one seamless model.

6. Acknowledgements

This work is supported by Carnegie Mellon University's SURF grant. We thank Kalvin Chang and Chenxuan Cui for ideas and discussions that informed this work, Kalvin Chang for guidance on using their preprocessed reconstruction datasets and their PyTorch reimplementation of Meloni et al. (2021), Aryaman Arora for guidance regarding their reflex prediction models, and Kalvin Chang, Anna Cai, and Ting Chen for help with revision and proofreading.

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A. Hyperparameters

We tune hyperparameters for all models on each dataset using WandB (Biewald, 2020) except for those tested by Kim et al. (2023) (which includes Meloni et al. (2021)'s GRU reconstruction model and Transformer reconstruction model on Romphon, Rom-orth, and Hóu). We use Bayesian search with 100 total runs (with early stopping) for the best validation phoneme edit distance (validated every 3 epochs). We keep a constant beam

size of 5 for GRU-BS reconstruction models during tuning. Tables 10, 11, 14, 13, 15, and 16 report our hyperparameter search results¹¹.

Adam optimizer's $\beta_1 = 0.9$, $\beta_2 = 0.999$, and $\varepsilon = 1e-8$ are obtained from Chang et al. (2022)'s experiments, while $\beta_1 = 0.9$, $\beta_2 = 0.98$, and $\varepsilon = 1e-9$ are used to consistently replicate Arora et al. (2023)'s experiments. We do not observe a noticeable effect β_2 and ε have on the models' performance.

B. Dataset Source and Splits

The WikiHan dataset can be obtained from Chang et al. (2022), and Hóu (2004)'s dataset can be obtained through Kim et al. (2023). WikiHan-aug is obtained from Cui et al. (2022). The Romance datasets by Meloni et al. (2021) is not licensed for redistribution and thus not included in our repository.

All the datasets are split by 70%, 10%, and 20% into train, validation, and test sets. The splits for WikiHan Chang et al. (2022) match the original work, and the splits for Meloni et al. (2021) and Hóu (2004) match Kim et al. (2023). WikiHan-aug Cui et al. (2022) has the same validation and test sets as Chang et al. (2022) but with augmented reflexes in the train set. Because Chang et al. (2022) only included cognate sets with at least 3 daughters in the train set, the train set in WikiHan-aug includes additional cognate sets that fulfill the 3-daughter requirement after augmentation.

Daughter languages included in WikiHan are Cantonese, Gan, Hakka, Jin, Mandarin, Hokkien, Wu, and Xiang. Daughter languages included in Romphon and Rom-orth are French, Italian, Spanish, Romanian, and Portuguese. Daughter languages included in Hóu are Beijing, Harbin, Tianjin, Jinan, Qingdao, Zhengzhou, Xian, Xining, Yinchuan, Lanzhou, Urumqi, Wuhan, Chengdu, Guiyang, Kunming, Nanjing, Hefei, Taiyuan, Pingyao, Hohhot, Shanghai, Suzhou, Hangzhou, Wenzhou, Shexian, Tunxi, Changsha, Xiangtan, Nanchang, Meixian, Taoyuan, Guangzhou, Nanning, Hong Kong, Xiamen, Fuzhou, Jianou, Shantou, and Haikou.

C. Training

All models are trained on NVIDIA GeForce RTX 2080 Ti or RTX A6000 GPUs. Each run takes about 1–3 hours of compute time. Our total GPU compute time is 237 days.

D. Reranking Hyperparameters

The optimal beam size *k* and score adjustment weight λ can be dataset-dependent. We use grid search with ranges and resolutions detailed in Table 9 to optimize *k* and λ on the validation set. The search results are shown in Table 17. We observe a preference for higher λ on Sinitic datasets.

Hyperparameter	Search range (inclusive)	Resolution
Beam size k	[2, 10]	2
Score adjustment	[0.3, 4.2]	0.3
weight λ		

Table 9:Grid search range and resolution forreranking hyperparameters.

E. Results with standard deviation

Table 18 shows reflex prediction performance with standard deviations, and Table 19 shows reconstruction performance with standard deviations.

F. Additional Reflex Error Analysis

Table 20 shows the reflex prediction error rate for each daughter among all test entries. Similar to Table 8, we observe an overall highest error rate on Hokkien.

G. Statistical Tests Results

We obtain *p*-values from the Wilcoxon Rank-Sum test and confidence intervals (Cl) from the Bootstrap test. Tables 21, 22, 23, 24, and 25 show *p*-values and 99% confidence intervals for reflex prediction performance. Tables 26, 27, 28, 29, and 30 show *p*-values and 99% confidence intervals for reconstruction performance.

H. Additional Reranking Case Studies

We provide additional reranking examples similar to Figure 1. Figures 4 and 5 show two additional reranking successes on WikiHan, Figures 6 and 7 show two failures on WikiHan, Figures 8 and 9 show two successes on Rom-phon, and Figures 10 and 11 show two failures on Rom-phon.

¹¹We use batch size to refer to the number of cognate sets in a batch, meaning that the number of reflex prediction training examples in each batch may vary if cognate sets have missing daughters.

	WikiHan	WikiHan-aug	Hóu	Rom-phon	Rom-orth
batch size k	128	256	32	256	256
beam search a	0.912598	0.600524	0.638660	0.825868	0.707860
dropout	0.405044	0.496428	0.497715	0.430556	0.489005
embedding size	509	148	265	154	283
feedforward size	218	471	232	310	311
hidden size	81	216	36	115	255
number of layers	1	1	1	1	1
learning rate	0.000629980	0.000550343	0.000691970	0.000762067	0.000568855
max epochs	576	204	194	285	304
warmup epochs	19	3	24	50	50
β_1	0.9	0.9	0.9	0.9	0.9
β_2	0.999	0.999	0.999	0.999	0.999
ε	1e-8	1e-8	1e-8	1e-8	1e-8

Table 10: Hyperparameters for GRU reconstruction model with beam search (GRU-BS), tuned with fixed beam size k = 5. Beam search *a* is the length normalization constant. The number of GRU layers is set to 1 to match Meloni et al. (2021).

	WikiHan	WikiHan-aug	Hóu	Rom-phon	Rom-orth
batch size	128	256	128	64	256
learning rate	0.000610810	0.00128592	0.00208360	0.000153890	0.000931776
max epochs	280	202	485	487	371
dropout	0.422406	0.411611	0.402412	0.467993	0.481404
embedding size	328	286	46	324	41
feedforward size	421	183	500	275	96
target-gated classifier	True	False	False	False	True
decode with language embedding	False	False	True	False	False
hidden size	46	33	110	177	194
number of encoder layers	2	4	1	2	1
one-hot target encoding	True	True	True	True	False
bidirectional encoder	True	True	True	True	True
use VAE latent	False	True	False	False	False
warmup epochs	0	42	28	41	6
β_1	0.9	0.9	0.9	0.9	0.9
β_2	0.999	0.999	0.999	0.999	0.999
ε	1e-8	1e-8	1e-8	1e-8	1e-8

Table 11: Hyperparameters for our GRU reflex prediction model. Target-gated classifier refers to whether each specific target language enables a specific subset of the token classifier, decode with language embedding refers to whether to embed the target sequences with language embedding similar to Meloni et al. (2021), one-hot target encoding refers to whether the classifier is prompted with an additional one-hot vector concatenated to its input to indicate the target daughter language, and use VAE latent refers to whether the decoder takes a sampled and reparametrized latent similar to Cui et al. (2022)'s VAE reconstruction model.

	WikiHan	WikiHan-aug	Hóu	Rom-phon	Rom-orth
batch size	512	32	32	128	64
learning rate	0.00133100	0.000275041	0.00223748	0.00103299	0.00117782
max epochs	413	186	177	514	383
dropout	0.109371	0.352477	0.239702	0.159863	0.250876
embedding size	128	128	64	64	64
feedforward size	429	1002	962	275	677
nhead	1	16	16	2	8
number of decoder layers	2	8	7	3	2
number of encoder layers	5	2	4	6	3
warmup epochs	20	17	40	5	37
weight_decay	6.29736e-07	5.34183e-07	9.50859e-07	4.76354e-07	9.89944e-07
β_1	0.9	0.9	0.9	0.9	0.9
β_2	0.999	0.999	0.999	0.999	0.999
ε	1e–8	1e–8	1e-8	1e–8	1e–8

Table 12: Hyperparameter for Kim et al. (2023)'s Transformer model adapted for reflex prediction.

	WikiHan	WikiHan-aug	Hóu	Rom-phon	Rom-orth
batch size	256	512	32	256	32
bidirectional encoder	True	True	True	True	True
dropout	0.380055	0.434051	0.170343	0.278587	0.337712
embedding size	319	284	169	508	473
hidden size	353	397	367	448	422
learning rate	0.000286969	0.000321132	0.00143399	0.000153972	0.00146615
max epochs	506	434	542	436	179
number of layers	2	2	4	4	2
warmup epochs	37	43	14	16	10
β_1	0.9	0.9	0.9	0.9	0.9
β_2	0.98	0.98	0.98	0.98	0.98
ε	1e-9	1e–9	1e–9	1e-9	1e–9

Table 13: Hyperparameters for Arora et al. (2023)'s GRU reflex prediction model tuned on our datasets of interest.

	WikiHan	WikiHan-aug	Hóu	Rom-phon	Rom-orth
batch size	32	256	32	64	32
feedforward size	295	832	535	786	110
model size	64	64	256	128	64
dropout	0.199505	0.447951	0.443183	0.263578	0.251537
learning rate	0.000561076	0.00234191	0.00260176	0.00161177	0.00164335
max epochs	361	437	536	207	264
nhead	4	2	4	1	4
number of layers	4	5	3	5	7
warmup epochs	9	49	11	2	9
β_1	0.9	0.9	0.9	0.9	0.9
β_2	0.98	0.98	0.98	0.98	0.98
ε	1e–9	1e-9	1e–9	1e–9	1e–9

Table 14: Hyperparameters for Arora et al. (2023)'s Transformer reflex prediction model tuned on our datasets of interest.

	WikiHan	WikiHan-aug
batch size	512	256
dropout	0.431211	0.409475
embedding size	248	196
feedforward size	375	421
decode with language embedding	True	True
hidden size	78	278
number of layers	1	1
bidirectional encoder	False	False
use VAE latent	False	False
learning rate	0.000935879	0.000964557
max epochs	472	298
warmup epochs	18	26
β_1	0.9	0.9
β_2	0.999	0.999
ε	1e-8	1e-8

Table 15: Hyperparameters for Meloni et al. (2021)'s GRU reconstruction model on WikiHan and WikiHanaug. For the hyperparameters used to train the same model on Hóu, Rom-phon, and Rom-orth, refer to Kim et al. (2023).

	WikiHan	WikiHan-aug
batch size	512	64
dropout	0.170582	0.293413
embedding size	256	64
feedforward size	133	857
nhead	1	2
number of decoder layers	5	3
number of encoder layers	4	6
learning rate	0.000556150	0.000595262
max epochs	194	209
warmup epochs	1	4
weight decay	8.48140e-07	8.26112e-07
β_1	0.9	0.9
β_2	0.999	0.999
ε	1e–8	1e-8

Table 16: Hyperparameters for Kim et al. (2023)'s Transformer reconstruction model on WikiHan and WikiHan-aug. For the hyperparameters used to train the same model on Hóu, Rom-phon, and Rom-orth, refer to Kim et al. (2023).

Dataset	Reranking System	k (beam size)	λ (score adjustment weight)
WikiHan	GRU-BS + GRU Reranker	6	1.395
	GRU-BS + Trans Reranker 1	6	1.275
	GRU-BS + Trans Reranker 2	7	1.260
WikiHan-aug	GRU-BS + GRU Reranker	7	1.620
	GRU-BS + Trans Reranker 1	8	1.755
	GRU-BS + Trans Reranker 2	7	1.575
Hóu	GRU-BS + GRU Reranker	7	1.755
	GRU-BS + Trans Reranker 1	7	2.430
	GRU-BS + Trans Reranker 2	7	2.415
Rom-phon	GRU-BS + GRU Reranker	5	0.420
	GRU-BS + Trans Reranker 1	6	0.555
	GRU-BS + Trans Reranker 2	6	0.585
Rom-orth	GRU-BS + GRU Reranker	6	0.870
	GRU-BS + Trans Reranker 1	6	0.990
	GRU-BS + Trans Reranker 2	5	0.915

Table 17: Reranking hyperparameter search results.

Dataset	Model	ACC%↑	TED↓	TER↓	FER↓	BCFS
WikiHan	GRU (baseline)	66.43% ±	0.5244 ±	0.1547 ±	0.0400 ±	0.7394 ±
		0.29%	0.0049	0.0014	0.0006	0.0022
	Transformer (Kim et al., 2023)	66.39% ±	0.5302 ±	0.1564 ±	0.0406 ±	0.7370 ±
		0.53%	0.0089	0.0026	0.0007	0.0040
	GRU (Arora et al., 2023)	64.45% ±	0.5558 ±	0.1640 ±	0.0428 ±	0.7260 ±
		0.34%	0.0060	0.0018	0.0007	0.0027
	Transformer (Arora et al., 2023)	67.64% ±	0.5128 ±	0.1513 ±	0.0390 ±	0.7445 ±
		0.35%	0.0072	0.0021	0.0006	0.0031
WikiHan-aug	GRU (baseline)	68.11% ±	0.5007 ±	0.1477 ±	0.0380 ±	0.7495 ±
		0.44%	0.0083	0.0024	0.0007	0.0036
	Transformer (Kim et al., 2023)	68.96% ±	0.4889 ±	0.1442 ±	0.0371 ±	0.7551 ±
		0.36%	0.0055	0.0016	0.0006	0.0022
	GRU (Arora et al., 2023)	66.94% ±	0.5159 ±	0.1522 ±	0.0391 ±	0.7430 ±
		0.68%	0.0101	0.0030	0.0011	0.0043
	Transformer (Arora et al., 2023)	69.37% ±	0.4826 ±	0.1424 ±	0.0363 ±	0.7572 ±
		0.18%	0.0028	0.0008	0.0004	0.0013
Hóu	GRU (baseline)	51.72% ±	0.7777 ±	0.2037 ±	0.0488 ±	0.6783 ±
		0.70%	0.0132	0.0035	0.0010	0.0046
	Transformer (Kim et al., 2023)	55.46% ±	0.7576 ±	0.1985 ±	0.0494 ±	0.6882 ±
		1.23%	0.0243	0.0064	0.0018	0.0078
	GRU (Arora et al., 2023)	49.26% ±	0.8266 ±	0.2166 ±	0.0528 ±	0.6622 ±
		1.57%	0.0370	0.0097	0.0030	0.0120
	Transformer (Arora et al., 2023)	55.60% ±	0.7520 ±	0.1970 ±	0.0485 ±	0.6892 ±
		1.30%	0.0243	0.0064	0.0018	0.0081
Rom-phon	GRU (baseline)	63.85% ±	0.7439 ±	0.1014 ±	0.0426 ±	0.8361 ±
		0.37%	0.0068	0.0009	0.0005	0.0014
	Transformer (Kim et al., 2023)	64.19% ±	0.7349 ±	0.1002 ±	0.0427 ±	0.8380 ±
		0.64%	0.0096	0.0013	0.0006	0.0019
	GRU (Arora et al., 2023)	48.28% ±	1.3257 ±	0.1808 ±	0.0930 ±	0.7567 ±
		14.82%	0.8784	0.1198	0.0748	0.1035
	Transformer (Arora et al., 2023)	63.96% ±	0.7442 ±	0.1015 ±	0.0428 ±	0.8361 ±
		0.65%	0.0087	0.0012	0.0005	0.0018
Rom-orth	GRU (baseline)	64.58% ±	0.7301 ±	0.0967 ±	-	0.8465 ±
		0.34%	0.0069	0.0009		0.0014
	Transformer (Kim et al., 2023)	64.80% ±	0.7258 ±	0.0961 ±	-	0.8478 ±
		0.50%	0.0061	0.0008		0.0011
	GRU (Arora et al., 2023)	57.92% ±	0.8741 ±	0.1158 ±	-	0.8218 ±
		2.31%	0.0346	0.0046		0.0054
	Iransformer (Arora et al., 2023)	65.20% ±	0.7247 ±	0.0960 ±	-	0.8476 ±
		0.46%	0.0069	0.0009		0.0012

Table 18: Performance means and standard deviations of the reflex prediction models across 20 runs, with the best-performing model for each metric in bold.

Dataset	Reconstruction System	ACC% ↑	TED↓	TER↓	FER↓	BCFS ↑
WikiHan	GRU (Meloni et al., 2021)	55.58% ±	0.7360 ±	0.1724 ±	0.0686 ±	0.7426 ±
		0.86%	0.0137	0.0032	0.0026	0.0038
	Trans (Kim et al., 2023)	54.62% ±	0.7453 ±	0.1746 ±	0.0696 ±	0.7393 ±
		1.22%	0.0165	0.0039	0.0029	0.0048
	GRU-BS (<i>k</i> = 10)	54.88% ±	0.7507 ±	0.1758 ±	0.0701 ±	0.7364 ±
		1.07%	0.0186	0.0043	0.0022	0.0064
	GRU-BS (<i>k</i> ≤ 10) + GRU	57.14% ±	0.7045 ±	0.1650 ±	0.0661 ±	0.7515 ±
	Reranker	0.80%*†	0.0146*†	0.0034*†	0.0018*†	0.0048*†
	GRU-BS ($k \le 10$) + Trans	57.26% ±	0.7029 ±	0.1646 ±	0.0658 ±	0.7520 ±
	Reranker 2	0.83% *†	0.0161*†	0.0038*†	0.0021 *†	0.0052*†
WikiHan-aug	GRU (Meloni et al., 2021)	54.73% ±	0.7574 ±	0.1774 ±	0.0689 ±	0.7346 ±
Ū		0.84%	0.0127	0.0030	0.0017	0.0048
	Trans (Kim et al., 2023)	55.82% ±	0.7317 ±	0.1714 ±	0.0661 ±	0.7416 ±
	, , , , , , , , , , , , , , , , , , ,	0.97%	0.0165	0.0039	0.0020	0.0053
	GRU-BS (<i>k</i> = 10)	56.64% ±	0.7214 ±	0.1690 ±	0.0658 ±	0.7454 ±
		0.66%*	0.0113	0.0026	0.0014	0.0035
	GRU-BS (<i>k</i> < 10) + GRU	58.58% ±	0.6822 ±	0.1598 ±	0.0628 ±	0.7579 ±
	Reranker	0.70%*†	0.0143*†	0.0033*†	0.0017*†	0.0040*†
	GRU-BS ($k < 10$) + Trans	58.58% ±	0.6840 ±	0.1602 ±	0.0626 ±	0.7575 ±
	Reranker 2	0.75%*†	0.0129*†	0.0030*†	0.0017*†	0.0038*†
Hóu	GRU (Meloni et al., 2021)	34.63% ±	1.0916 ±	0.2479 ±	0.0914 ±	0.6697 ±
		2.37%	0.0629	0.0147	0.0049	0.0167
	Trans (Kim et al., 2023)	39.01% ±	$0.9904 \pm$	0.2233 ±	0.0875 ±	0.6955 ±
		2.89%	0.0443	0.0108	0.0069	0.0103
	GBU-BS (k = 10)	37.36% +	1.0382 +	0.2328 +	0.0917 +	0.6974 +
		3.25%	0.0662	0.0148	0.0065	0.0176
	GRU-BS (k < 10) + GRU	40.50% +	0.9727 +	0.2181 +	0.0867 +	0.7130 +
	Beranker	3 09%+	0.0486+	0.0109+	0.0058+	0.0132*+
	GRU-BS ($k < 10$) + Trans	42.08% ±	$0.9503 \pm$	0.2131 ±	0.0850 ±	0.7170 ±
	Reranker 2	2.96%*†	0.0525*†	0.0118*†	0.0063+	0.0137*†
Rom-phon	GRU (Meloni et al., 2021)	51.92% ±	0.9775 ±	0.1244 ±	0.0390 ±	0.8275 ±
		0.65%	0.0216	0.0028	0.0012	0.0033
	Trans (Kim et al., 2023)	53.04% ±	$0.9050 \pm$	0.1148 ±	$0.0377 \pm$	0.8417 ±
		0.80%	0.0166	0.0018	0.0008	0.0024
	GBU-BS (k = 10)	52.63% +	0.9125 +	0.1018 +	0.0353 +	0.8402 +
		0.68%	0.0174	0.0019*	0.0009*	0.0032
	GRU-BS (k < 10) + GRU	53.95% ±	0.8775 +	0.0979 +	0.0336 +	0.8460 +
	Beranker	0.77%*+	0.0165*†	0.0018*+	0.0007*+	0.0028*+
	GBU-BS ($k < 10$) + Trans	53 85% +	0.8765 +	0.0978 +	0.0333 +	0.8461 +
	Reranker 2	0.79%*†	0.0177*+	0.0020*+	0.0008*+	0.0030*+
Rom-orth	GRU (Meloni et al., 2021)	69.41% +	0.6004 +	0.0781 +	-	0.8906 +
		0.53%	0.0130	0.0018		0.0023
	Trans (Kim et al., 2023)	71.05% +	0.5636 +	0.0734 +	-	0.8981 +
		0.50%	0.0163	0.0022		0.0028
	GBU-BS ($k = 10$)	71.09% +	0.5531 +	0.0617 +	-	0.8990 +
		0.51%	0.0127	0.0014*		0.0023
	GBU-BS (k < 10) + GBU	72.60% +	0.5237 +	0.0584 +	-	0.9045 +
	Beranker	0 41%*+	0 0109*+	0 0012*+		0.0019*+
	GBU-BS(k < 10) + Trans	72 50% +	0.5246 +	0.0585 +	_	0 9044 +
	Beranker 2	0 45%*+	0.0111*+	0.0012*+		0.0020*+
		0.1070	0.0111	0.0012		5.5520 I

Table 19: Performance means and standard deviations of reconstruction systems across 20 runs. Reconstruction systems include baselines, GRU with beam search (GRU-BS), and GRU-BS with reranking. Bold indicates the best-performing system for each metric, asterisks indicate statistically better performance than both baseline models (Meloni et al. (2021)'s GRU and Kim et al. (2023)'s Transformer), and daggers indicate that a reranking system performs statistically better than its beam search counterpart.

	Cantonese	Gan	Hakka	Jin	Mandarin	Hokkien	Wu	Xiang
Not-in	75.3%	72.7%	70.8%	83.3%	72.2%	82.1%	57.7%	82.1%
Worsened	59.4%	75.0%	40.0%	71.4%	68.8%	68.8%	33.3%	60.0%
Unchanged	16.6%	19.1%	32.2%	14.5%	15.1%	39.5%	14.1%	29.7%
Improved	34.9%	30.0%	51.0%	10.0%	21.4%	48.7%	31.6%	38.1%
Overall	28.6%	26.5%	38.8%	23.7%	26.2%	47.8%	20.4%	37.3%

Table 20: Transformer reranker error rates (when predicting reflexes from the gold protoform) for each daughter language in the WikiHan dataset given each reranker behavior category, obtained from a randomly selected run.

Model 1	Model 2	ACC%↑	TED↓	TER↓	FER↓	BCFS↑
GRU (baseline)	Trans (Kim et al., 2023)	p = 0.4196	p = 0.0093*	p = 0.0093*	p = 0.0021*	p = 0.0242
		(-0.0029, 0.0038)	(-0.0112, 0.0001)	(-0.0033, 0.0000)	(-0.0011, -0.0001)	*(-0.0003, 0.0047)
	GRU (Arora et al., 2023)	p < 0.0001*	p < 0.0001*	p < 0.0001*	p < 0.0001*	p < 0.0001*
		(0.0173, 0.0225)*	(-0.0355, -0.0268)	*(-0.0105, -0.0079)	*(-0.0033, -0.0022)	* (0.0113, 0.0152)*
	Trans (Arora et al., 2023)	p = 1.0000	p = 1.0000	p = 1.0000	p = 1.0000	p = 1.0000
		(-0.0149, -0.0095)	(0.0060, 0.0166)	(0.0018, 0.0049)	(0.0005, 0.0015)	(-0.0073, -0.0027)
Trans (Kim et al., 2023)	GRU (baseline)	p = 0.5804	p = 0.9907	p = 0.9907	p = 0.9979	p = 0.9758
		(-0.0040, 0.0031)	(-0.0003, 0.0115)	(-0.0001, 0.0034)	(0.0001, 0.0011)	(-0.0049, 0.0004)
	GRU (Arora et al., 2023)	p < 0.0001*	p < 0.0001*	p < 0.0001*	p < 0.0001*	p < 0.0001*
		$(0.0158, 0.0230)^*$	(-0.0318, -0.0195)	*(-0.0094, -0.0058)	*(-0.0027, -0.0016)	* (0.0083, 0.0138)*
	Trans (Arora et al., 2023)	p = 1.0000	<i>p</i> = 1.0000	p = 1.0000	p = 1.0000	<i>p</i> = 1.0000
		(-0.0162, -0.0089)	(0.0105, 0.0236)	(0.0031, 0.0070)	(0.0011, 0.0021)	(-0.0102, -0.0044)
GRU (Arora et al., 2023)	GRU (baseline)	p = 1.0000	<i>p</i> = 1.0000	p = 1.0000	p = 1.0000	<i>p</i> = 1.0000
		(-0.0223, -0.0173)	(0.0265, 0.0355)	(0.0078, 0.0105)	(0.0022, 0.0033)	(-0.0152, -0.0112)
	Trans (Kim et al., 2023)	p = 1.0000	<i>p</i> = 1.0000	p = 1.0000	p = 1.0000	<i>p</i> = 1.0000
		(-0.0229, -0.0158)	(0.0196, 0.0317)	(0.0058, 0.0093)	(0.0016, 0.0027)	(-0.0138, -0.0084)
	Trans (Arora et al., 2023)	p = 1.0000	<i>p</i> = 1.0000	p = 1.0000	p = 1.0000	<i>p</i> = 1.0000
		(-0.0348, -0.0291)	(0.0372, 0.0482)	(0.0110, 0.0142)	(0.0032, 0.0043)	(-0.0208, -0.0160)
Trans (Arora et al., 2023)	GRU (baseline)	p < 0.0001*	p < 0.0001*	p < 0.0001*	p < 0.0001*	p < 0.0001*
		(0.0096, 0.0148)*	(-0.0165, -0.0063)	*(-0.0049, -0.0019)	*(-0.0015, -0.0005)	*(0.0029, 0.0073)*
	Trans (Kim et al., 2023)	p < 0.0001*	p < 0.0001*	p < 0.0001*	p < 0.0001*	p < 0.0001*
		(0.0090, 0.0161)*	(-0.0236, -0.0105)	*(-0.0070, -0.0031)	*(-0.0021, -0.0011)	* (0.0044, 0.0102)*
	GRU (Arora et al., 2023)	p < 0.0001*	<i>p</i> < 0.0001*	<i>p</i> < 0.0001*	<i>p</i> < 0.0001*	<i>p</i> < 0.0001*
		(0.0293, 0.0349)*	(-0.0482, -0.0371)	*(-0.0142,-0.0110)	*(-0.0043, -0.0032)	*(0.0159, 0.0208)*

Table 21: Reflex prediction significance test results for WikiHan. Asterisks indicates that Model 1 performs better than Model 2 with the corresponding test (*p*-value or CI).

Model 1	Model 2	ACC%	TED	TER⊥	FER⊥	BCFS 1
GRU (baseline)	Trans (Kim et al., 2023)	p = 1.0000	p = 1.0000	p = 1.0000	p = 1.0000	p = 1.0000
,	, , , , , , , , , , , , , , , , , , ,	(-0.0120, -0.0053)	(0.0066, 0.0182)	(0.0020, 0.0054)	(0.0005, 0.0015)	(-0.0083, -0.0034)
	GRU (Arora et al., 2023)	p < 0.0001*	p < 0.0001*	p < 0.0001*	p < 0.0010*	p < 0.0001*
		(0.0074, 0.0166)*	(-0.0224, -0.0075)	*(-0.0066, -0.0022)	*(-0.0018, -0.0003)	* (0.0033, 0.0096)*
	Trans (Arora et al., 2023)	p = 1.0000	p = 1.0000	p = 1.0000	p = 1.0000	p = 1.0000
		(-0.0158, -0.0101)	(0.0139, 0.0245)	(0.0041, 0.0072)	(0.0013, 0.0022)	(-0.0104, -0.0058)
Trans (Kim et al., 2023)	GRU (baseline)	p < 0.0001*	p < 0.0001*	p < 0.0001*	p < 0.0001*	p < 0.0001*
		(0.0053, 0.0121)*	(-0.0183, -0.0066)	*(-0.0054, -0.0019)	*(-0.0015, -0.0005)	* (0.0034, 0.0083)*
	GRU (Arora et al., 2023)	p < 0.0001*	p < 0.0001*	p < 0.0001*	p < 0.0001*	p < 0.0001*
		(0.0162, 0.0252)*	(-0.0342, -0.0207)	*(-0.0101, -0.0061)	*(-0.0028, -0.0013)	* (0.0094, 0.0151)*
	Trans (Arora et al., 2023)	p = 0.9999	p = 1.0000	p = 1.0000	p = 1.0000	p = 0.9994
		(-0.0067, -0.0020)	(0.0030, 0.0102)	(0.0009, 0.0030)	(0.0004, 0.0012)	(-0.0037, -0.0007)
GRU (Arora et al., 2023)	GRU (baseline)	p = 1.0000	p = 1.0000	p = 1.0000	p = 0.9997	p = 1.0000
		(-0.0169, -0.0073)	(0.0075, 0.0230)	(0.0022, 0.0068)	(0.0003, 0.0019)	(-0.0098, -0.0032)
	Trans (Kim et al., 2023)	p = 1.0000	p = 1.0000	p = 1.0000	p = 1.0000	p = 1.0000
		(-0.0250, -0.0162)	(0.0208, 0.0339)	(0.0061, 0.0100)	(0.0013, 0.0028)	(-0.0150, -0.0095)
	Trans (Arora et al., 2023)	p = 1.0000	p = 1.0000	p = 1.0000	p = 1.0000	p = 1.0000
		(-0.0290, -0.0209)	(0.0279, 0.0400)	(0.0082, 0.0118)	(0.0022, 0.0035)	(-0.0170, -0.0118)
Trans (Arora et al., 2023)	GRU (baseline)	p < 0.0001*	p < 0.0001*	p < 0.0001*	p < 0.0001*	p < 0.0001*
		(0.0102, 0.0158)*	(-0.0244, -0.0140)	*(-0.0072, -0.0041)	*(-0.0022, -0.0013)	* (0.0058, 0.0104)*
	Trans (Kim et al., 2023)	p < 0.0001*	p < 0.0001*	p < 0.0001*	p < 0.0001*	p < 0.0010*
		$(0.0019, 0.0065)^*$	(-0.0102, -0.0029)	*(-0.0030, -0.0009)	*(-0.0012, -0.0004)	* (0.0007, 0.0037)*
	GRU (Arora et al., 2023)	p < 0.0001*	p < 0.0001*	p < 0.0001*	p < 0.0001*	p < 0.0001*
		$(0.0208, 0.0290)^{\star}$	(-0.0398, -0.0277)	*(-0.0117, -0.0082)	*(-0.0035, -0.0022)	* (0.0118, 0.0169)*

Table 22: Reflex prediction significance test results for WikiHan-aug. Asterisks indicates that Model 1 performs better than Model 2 with the corresponding test (*p*-value or CI).

Model 1	Model 2	ACC%↑	TED↓	TER↓	FER↓	BCFS ↑
GRU (baseline)	Trans (Kim et al., 2023)	p = 1.0000	p = 0.9989	p = 0.9989	p = 0.1337	p = 1.0000
		(-0.0453, -0.0292)	(0.0037, 0.0360)	(0.0010, 0.0094)	(-0.0017, 0.0006)	(-0.0149, -0.0044)
	GRU (Arora et al., 2023)	$p < 0.0001^{\star}$	p < 0.0001*	p < 0.0001*	p < 0.0001*	p < 0.0001*
		(0.0149, 0.0342)*	(-0.0709, -0.0276)	*(-0.0186, -0.0072)	*(-0.0057, -0.0022)	* (0.0091, 0.0235)*
	Trans (Arora et al., 2023)	p = 1.0000	p = 0.9998	p = 0.9998	p = 0.9075	p = 1.0000
		(-0.0474, -0.0305)	(0.0097, 0.0414)	(0.0025, 0.0109)	(-0.0009, 0.0014)	(-0.0161, -0.0053)
Trans (Kim et al., 2023)	GRU (baseline)	p < 0.0001*	p = 0.0011*	p = 0.0011*	p = 0.8663	p < 0.0001*
		(0.0294, 0.0455)*	(-0.0355, -0.0042)	*(-0.0093, -0.0011)	*(-0.0006, 0.0017)	(0.0046, 0.0150)*
	GRU (Arora et al., 2023)	p < 0.0001*	p < 0.0001*	p < 0.0001*	p < 0.0010*	p < 0.0001*
		(0.0507, 0.0732)*	(-0.0949, -0.0441)	*(-0.0249, -0.0116)	*(-0.0054, -0.0015)	* (0.0180, 0.0343)*
	Trans (Arora et al., 2023)	p = 0.6221	p = 0.8066	p = 0.8066	p = 0.9384	p = 0.6964
		(-0.0118, 0.0086)	(-0.0137, 0.0254)	(-0.0036, 0.0067)	(-0.0006, 0.0023)	(-0.0076, 0.0053)
GRU (Arora et al., 2023)	GRU (baseline)	p = 1.0000	p = 1.0000	p = 1.0000	p = 1.0000	p = 1.0000
		(-0.0340, -0.0147)	(0.0272, 0.0711)	(0.0071, 0.0186)	(0.0022, 0.0058)	(-0.0234, -0.0090)
	Trans (Kim et al., 2023)	p = 1.0000	p = 1.0000	p = 1.0000	p = 0.9998	p = 1.0000
		(-0.0733, -0.0505)	(0.0444, 0.0942)	(0.0116, 0.0247)	(0.0014, 0.0054)	(-0.0341, -0.0181)
	Trans (Arora et al., 2023)	p = 1.0000	p = 1.0000	p = 1.0000	p = 1.0000	p = 1.0000
		(-0.0750, -0.0518)	(0.0495, 0.0999)	(0.0130, 0.0262)	(0.0023, 0.0063)	(-0.0353, -0.0187)
Trans (Arora et al., 2023)	GRU (baseline)	p < 0.0001*	p < 0.0010*	p < 0.0010*	p = 0.0925	p < 0.0001*
		(0.0301, 0.0472)*	(-0.0416, -0.0098)	*(-0.0109, -0.0026)	*(-0.0014,0.0010)	(0.0054, 0.0162)*
	Trans (Kim et al., 2023)	p = 0.3779	p = 0.1934	p = 0.1934	p = 0.0616	p = 0.3036
		(-0.0087, 0.0120)	(-0.0256, 0.0147)	(-0.0067, 0.0039)	(-0.0023, 0.0007)	(-0.0056, 0.0076)
	GRU (Arora et al., 2023)	p < 0.0001*	p < 0.0001*	p < 0.0001*	p < 0.0001*	p < 0.0001*
		(0.0518, 0.0749)*	(-0.0995, -0.0494)	*(-0.0261, -0.0129)	*(-0.0062, -0.0024)	* (0.0188, 0.0351)*

Table 23: Reflex prediction significance test results for Hóu. Asterisks indicates that Model 1 performs better than Model 2 with the corresponding test (*p*-value or Cl).

Model 1	Model 2	ACC%↑	TED↓	TER↓	FER↓	BCFS 1
GRU (baseline)	Trans (Kim et al., 2023)	p = 0.9848	p = 0.9983	p = 0.9983	p = 0.4143	p = 0.9983
		(-0.0076, 0.0010)	(0.0022, 0.0158)	(0.0003, 0.0022)	(-0.0005, 0.0004)	(-0.0032, -0.0005)
	GRU (Arora et al., 2023)	p < 0.0001*	p < 0.0001*	p < 0.0001*	p < 0.0001*	p < 0.0001*
		(0.0956, 0.2815)*	(-1.3739, -0.2318)	*(-0.1874, -0.0316)	*(-0.1226, -0.0206)	* (0.0392, 0.1708)*
	Trans (Arora et al., 2023)	p = 0.8066	p = 0.3132	p = 0.3132	p = 0.0616	p = 0.3727
		(-0.0051, 0.0034)	(-0.0064, 0.0064)	(-0.0009, 0.0009)	(-0.0005, 0.0002)	(-0.0014, 0.0012)
Trans (Kim et al., 2023)	GRU (baseline)	p = 0.0152	p = 0.0017*	p = 0.0017*	p = 0.5857	p = 0.0017*
		(-0.0010, 0.0075)	(-0.0160, -0.0022)	*(-0.0022, -0.0003)	*(-0.0004, 0.0005)	(0.0005, 0.0033)*
	GRU (Arora et al., 2023)	p < 0.0001*	p < 0.0001*	p < 0.0001*	p < 0.0001*	p < 0.0001*
		(0.0989, 0.2843)*	(-1.3808, -0.2415)	*(-0.1883, -0.0329)	*(-0.1225, -0.0206)	* (0.0411, 0.1726)*
	Trans (Arora et al., 2023)	p = 0.2085	$p = 0.0027^*$	$p = 0.0027^*$	p = 0.1789	$p = 0.0043^*$
		(-0.0030, 0.0076)	(-0.0167, -0.0017)	*(-0.0023, -0.0002)	*(-0.0006, 0.0003)	$(0.0003, 0.0033)^*$
GRU (Arora et al., 2023)	GRU (baseline)	p = 1.0000	p = 1.0000	p = 1.0000	p = 1.0000	p = 1.0000
		(-0.2892, -0.0959)	(0.2447, 1.3977)	(0.0334, 0.1906)	(0.0214, 0.1162)	(-0.1743, -0.0405)
	Trans (Kim et al., 2023)	p = 1.0000	p = 1.0000	p = 1.0000	p = 1.0000	p = 1.0000
		(-0.2924, -0.0998)	(0.2544, 1.4058)	(0.0347, 0.1917)	(0.0213, 0.1160)	(-0.1760, -0.0423)
	Trans (Arora et al., 2023)	p = 1.0000	p = 1.0000	p = 1.0000	p = 1.0000	p = 1.0000
		(-0.2885, -0.0971)	(0.2443, 1.3977)	(0.0333, 0.1906)	(0.0211, 0.1160)	(-0.1744, -0.0404)
Trans (Arora et al., 2023)	GRU (baseline)	p = 0.1934	p = 0.6868	p = 0.6868	p = 0.9384	p = 0.6273
		(-0.0035, 0.0053)	(-0.0064, 0.0065)	(-0.0009, 0.0009)	(-0.0002, 0.0005)	(-0.0013, 0.0014)
	Trans (Kim et al., 2023)	p = 0.7915	p = 0.9973	p = 0.9973	p = 0.8211	p = 0.9957
		(-0.0077, 0.0031)	(0.0015, 0.0167)	(0.0002, 0.0023)	(-0.0003, 0.0006)	(-0.0033, -0.0002)
	GRU (Arora et al., 2023)	p < 0.0001*	p < 0.0001*	p < 0.0001*	p < 0.0001*	p < 0.0001*
		(0.0963, 0.2810)*	(-1.3713, -0.2328)	*(-0.1870, -0.0317)	*(-0.1225, -0.0204)	* (0.0392, 0.1710)*

Table 24: Reflex prediction significance test results for Rom-phon. Asterisks indicates that Model 1 performs better than Model 2 with the corresponding test (*p*-value or CI).

Model 1	Model 2	ACC%↑	TED↓	TER↓	FER↓	BCFS ↑
GRU (baseline)	Trans (Kim et al., 2023)	p = 0.9837	p = 0.9661	p = 0.9661	-	p = 0.9971
		(-0.0052, 0.0020)	(-0.0012, 0.0094)	(-0.0002, 0.0012)		(-0.0022, -0.0003)
	GRU (Arora et al., 2023)	p < 0.0001*	p < 0.0001*	p < 0.0001*	-	p < 0.0001*
		(0.0559, 0.0829)*	(-0.1649, -0.1257)	*(-0.0218, -0.0166)	*	(0.0218, 0.0281)*
	Trans (Arora et al., 2023)	p = 1.0000	p = 0.9848	p = 0.9848	-	p = 0.9893
		(-0.0095, -0.0028)	(-0.0003, 0.0110)	(-0.0000, 0.0015)		(-0.0022, -0.0000)
Trans (Kim et al., 2023)	GRU (baseline)	p = 0.0163	p = 0.0339	p = 0.0339	-	p = 0.0029*
		(-0.0021, 0.0052)	(-0.0092, 0.0012)	(-0.0012, 0.0002)		(0.0003, 0.0022)*
	GRU (Arora et al., 2023)	p < 0.0001*	p < 0.0001*	p < 0.0001*	-	p < 0.0001*
		(0.0577, 0.0850)*	(-0.1690, -0.1299))* (-0.0224, -0.0172)	*	(0.0230, 0.0292)*
	Trans (Arora et al., 2023)	p = 0.9951	p = 0.6118	p = 0.6118	-	p = 0.3132
		(-0.0083, -0.0005)	(-0.0039, 0.0068)	(-0.0005, 0.0009)		(-0.0008, 0.0011)
GRU (Arora et al., 2023)	GRU (baseline)	p = 1.0000	p = 1.0000	p = 1.0000	-	p = 1.0000
		(-0.0826, -0.0555)	(0.1246, 0.1651)	(0.0165, 0.0219)		(-0.0280, -0.0216)
	Trans (Kim et al., 2023)	p = 1.0000	p = 1.0000	p = 1.0000	-	p = 1.0000
		(-0.0850, -0.0576)	(0.1294, 0.1689)	(0.0171, 0.0224)		(-0.0292, -0.0230)
	Trans (Arora et al., 2023)	p = 1.0000	p = 1.0000	p = 1.0000	-	p = 1.0000
		(-0.0890, -0.0617)	(0.1301, 0.1702)	(0.0172, 0.0225)		(-0.0291, -0.0228)
Trans (Arora et al., 2023)	GRU (baseline)	p < 0.0001*	p = 0.0152	p = 0.0152	-	p = 0.0107
		(0.0027, 0.0094)*	(-0.0109, 0.0002)	(-0.0014, 0.0000)		(0.0001, 0.0022)*
	Trans (Kim et al., 2023)	$p = 0.0049^*$	p = 0.3882	p = 0.3882	-	p = 0.6868
		(0.0003, 0.0085)*	(-0.0067, 0.0041)	(-0.0009, 0.0005)		(-0.0011, 0.0008)
	GRU (Arora et al., 2023)	p < 0.0001*	p < 0.0001*	p < 0.0001*	-	p < 0.0001*
		$(0.0619, 0.0894)^{\star}$	(-0.1702, -0.1308)	*(-0.0225, -0.0173)	*	(0.0229, 0.0291)*

Table 25: Reflex prediction significance test results for Rom-orth. Asterisks indicates that Model 1 performs better than Model 2 with the corresponding test (*p*-value or Cl).

Reconstruction System 1	Reconstruction System 2	ACC%↑	TED↓	TER↓	FER↓	BCFS ↑
GRU (Meloni et al., 2021)	Trans (Kim et al., 2023)	p = 0.0075*	p = 0.0283	p = 0.0283	p = 0.1719	$p = 0.0080^*$
		(0.0011, 0.0185)*	(-0.0217, 0.0029)	(-0.0051, 0.0007)	(-0.0034, 0.0011)	(-0.0004, 0.0067)
	GRU-BS	p = 0.0405	$p = 0.0040^*$	$p = 0.0040^*$	p = 0.0442	p < 0.0010*
		(-0.0009, 0.0152)	(-0.0288, -0.0018)	*(-0.0068, -0.0004)	*(-0.0035, 0.0004)	(0.0023, 0.0111)*
	GRU-BS + GRU Reranker	p = 1.0000	p = 1.0000	p = 1.0000	p = 0.9992	p = 1.0000
		(-0.0223, -0.0090)	(0.0196, 0.0428)	(0.0046, 0.0100)	(0.0006, 0.0042)	(-0.0122, -0.0050)
	GRU-BS + Trans. Reranker	p = 1.0000	p = 1.0000	p = 1.0000	p = 0.9991	p = 1.0000
		(-0.0240, -0.0101)	(0.0216, 0.0464)	(0.0051, 0.0109)	(0.0009, 0.0047)	(-0.0134, -0.0057)
Trans (Kim et al., 2023)	GRU (Meloni et al., 2021)	p = 0.9925	p = 0.9717	p = 0.9717	p = 0.8281	p = 0.9920
		(-0.0182, -0.0014)	(-0.0023, 0.0222)	(-0.0005, 0.0052)	(-0.0011, 0.0034)	(-0.0067, 0.0002)
	GRU-BS	p = 0.8066	p = 0.1427	p = 0.1427	p = 0.2581	p = 0.0684
		(-0.0124, 0.0063)	(-0.0197, 0.0092)	(-0.0046, 0.0022)	(-0.0025, 0.0016)	(-0.0015, 0.0077)
	GRU-BS + GRU Reranker	p = 1.0000	p = 1.0000	p = 1.0000	p = 0.9999	p = 1.0000
		(-0.0339, -0.0173)	(0.0280, 0.0533)	(0.0066, 0.0125)	(0.0017, 0.0056)	(-0.0158, -0.0080)
	GRU-BS + Trans. Reranker	p = 1.0000	p = 1.0000	p = 1.0000	p = 1.0000	p = 1.0000
		(-0.0354, -0.0183)	(0.0302, 0.0570)	(0.0071, 0.0134)	(0.0019, 0.0060)	(-0.0169, -0.0087)
GRU-BS	GRU (Meloni et al., 2021)	p = 0.9595	p = 0.9960	p = 0.9960	p = 0.9558	p = 0.9996
		(-0.0150, 0.0007)	(0.0016, 0.0286)	(0.0004, 0.0067)	(-0.0004, 0.0035)	(-0.0109, -0.0022)
	Trans (Kim et al., 2023)	p = 0.1934	p = 0.8573	p = 0.8573	p = 0.7419	p = 0.9316
		(-0.0068, 0.0122)	(-0.0085, 0.0197)	(-0.0020, 0.0046)	(-0.0016, 0.0025)	(-0.0080, 0.0014)
	GRU-BS + GRU Reranker	p = 1.0000	p = 1.0000	p = 1.0000	p = 1.0000	p = 1.0000
		(-0.0304, -0.0150)	(0.0327, 0.0607)	(0.0077, 0.0142)	(0.0024, 0.0056)	(-0.0200, -0.0106)
	GRU-BS + Trans. Reranker	p = 1.0000	p = 1.0000	p = 1.0000	p = 1.0000	p = 1.0000
		(-0.0317, -0.0162)	(0.0344, 0.0626)	(0.0081, 0.0147)	(0.0026, 0.0061)	(-0.0208, -0.0111)
GRU-BS + GRU Reranker	GRU (Meloni et al., 2021)	p < 0.0001*	p < 0.0001*	p < 0.0001*	p < 0.0010*	p < 0.0001*
		(0.0090, 0.0224)*	(-0.0426, -0.0193)	*(-0.0100, -0.0045)	*(-0.0042, -0.0006)	* (0.0051, 0.0121)*
	Trans (Kim et al., 2023)	p < 0.0001*	p < 0.0001*	p < 0.0001*	p < 0.0001*	p < 0.0001*
		(0.0172, 0.0339)*	(-0.0537, -0.0281)	*(-0.0126, -0.0066)	*(-0.0056, -0.0018)	* (0.0078, 0.0158)*
	GRU-BS	p < 0.0001*	p < 0.0001*	p < 0.0001*	p < 0.0001*	p < 0.0001*
		(0.0151, 0.0303)*	(-0.0598, -0.0321)	*(-0.0140, -0.0075)	*(-0.0056, -0.0024)	* (0.0105, 0.0197)*
	GRU-BS + Trans. Reranker	p = 0.6723	p = 0.5962	p = 0.5962	p = 0.6425	p = 0.5216
		(-0.0080, 0.0053)	(-0.0102, 0.0158)	(-0.0024, 0.0037)	(-0.0012, 0.0020)	(-0.0051, 0.0032)
GRU-BS + Trans. Reranker	GRU (Meloni et al., 2021)	p < 0.0001*	p < 0.0001*	p < 0.0001*	p < 0.0010*	p < 0.0001*
		(0.0099, 0.0237)*	(-0.0456, -0.0213)	*(-0.0107, -0.0050)	*(-0.0047, -0.0009)	* (0.0058, 0.0132)*
	Trans (Kim et al., 2023)	p < 0.0001*	p < 0.0001*	p < 0.0001*	p < 0.0001*	p < 0.0001*
		(0.0183, 0.0356)*	(-0.0564, -0.0297)	*(-0.0132, -0.0070)	*(-0.0060, -0.0020)	* (0.0085, 0.0169)*
	GRU-BS	p < 0.0001*	p < 0.0001*	p < 0.0001*	p < 0.0001*	p < 0.0001*
		$(0.0160, 0.0314)^{\star}$	(-0.0625, -0.0338)	*(-0.0146, -0.0079)	*(-0.0061, -0.0026)	* (0.0110, 0.0207)*
	GRU-BS + GRU Reranker	p = 0.3277	p = 0.4038	p = 0.4038	p = 0.3575	p = 0.4784
		(-0.0054, 0.0080)	(-0.0150, 0.0100)	(-0.0035, 0.0023)	(-0.0019, 0.0012)	(-0.0034, 0.0050)

Table 26: Reconstruction significance test results for WikiHan. Asterisks indicates that Reconstruction System 1 performs better than Reconstruction System 2 with the corresponding test (*p*-value or CI).

Reconstruction System 1	Reconstruction System 2	ACC%↑	TED↓	TER↓	FER↓	BCFS ↑
GRU (Meloni et al., 2021)	Trans (Kim et al., 2023)	p = 0.9998	p = 1.0000	p = 1.0000	p = 1.0000	p = 0.9998
		(-0.0171, -0.0025)	(0.0120, 0.0362)	(0.0028, 0.0085)	(0.0012, 0.0042)	(-0.0108, -0.0025)
	GRU-BS	<i>p</i> = 1.0000	<i>p</i> = 1.0000	<i>p</i> = 1.0000	p = 1.0000	<i>p</i> = 1.0000
		(-0.0245, -0.0121)	(0.0253, 0.0449)	(0.0059, 0.0105)	(0.0018, 0.0044)	(-0.0141, -0.0074)
	GRU-BS + GRU Reranker	p = 1.0000	p = 1.0000	p = 1.0000	p = 1.0000	p = 1.0000
		(-0.0442, -0.0315)	(0.0633, 0.0854)	(0.0148, 0.0200)	(0.0047, 0.0074)	(-0.0268, -0.0196)
	GRU-BS + Trans. Reranker	p = 1.0000	p = 1.0000	p = 1.0000	p = 1.0000	<i>p</i> = 1.0000
		(-0.0444, -0.0313)	(0.0623, 0.0829)	(0.0146, 0.0194)	(0.0050, 0.0077)	(-0.0263, -0.0194)
Trans (Kim et al., 2023)	GRU (Meloni et al., 2021)	p < 0.0010*	p < 0.0001*	p < 0.0001*	p < 0.0001*	<i>p</i> < 0.0010*
		(0.0022, 0.0171)*	(-0.0361, -0.0125)	*(-0.0084, -0.0029)	*(-0.0042, -0.0012)	* (0.0025, 0.0107)*
	GRU-BS	p = 0.9971	p = 0.9595	p = 0.9595	p = 0.6723	p = 0.9801
		(-0.0160, -0.0023)	(0.0001, 0.0226)	(0.0000, 0.0053)	(-0.0010, 0.0019)	(-0.0079, -0.0005)
	GRU-BS + GRU Reranker	<i>p</i> = 1.0000	<i>p</i> = 1.0000	<i>p</i> = 1.0000	p = 1.0000	<i>p</i> = 1.0000
		(-0.0358, -0.0216)	(0.0375, 0.0629)	(0.0088, 0.0147)	(0.0018, 0.0049)	(-0.0205, -0.0127)
	GRU-BS + Trans. Reranker	<i>p</i> = 1.0000	<i>p</i> = 1.0000	p = 1.0000	p = 1.0000	<i>p</i> = 1.0000
		(-0.0360, -0.0212)	(0.0367, 0.0607)	(0.0086, 0.0142)	(0.0021, 0.0052)	(-0.0200, -0.0126)
GRU-BS	GRU (Meloni et al., 2021)	p < 0.0001*	p < 0.0001*	p < 0.0001*	p < 0.0001*	p < 0.0001*
		(0.0118, 0.0247)*	(-0.0450, -0.0251)	*(-0.0105, -0.0059)	*(-0.0043, -0.0019)	* (0.0073, 0.0142)*
	Trans (Kim et al., 2023)	$p = 0.0029^*$	p = 0.0405	p = 0.0405	p = 0.3277	p = 0.0199
		(0.0022, 0.0158)*	(-0.0230, -0.0001)	*(-0.0054, -0.0000)	*(-0.0019, 0.0010)	(0.0005, 0.0079)*
	GRU-BS + GRU Reranker	<i>p</i> = 1.0000	<i>p</i> = 1.0000	p = 1.0000	p = 1.0000	<i>p</i> = 1.0000
		(-0.0251, -0.0141)	(0.0285, 0.0496)	(0.0067, 0.0116)	(0.0017, 0.0043)	(-0.0155, -0.0094)
	GRU-BS + Trans. Reranker	p = 1.0000	<i>p</i> = 1.0000	p = 1.0000	p = 1.0000	<i>p</i> = 1.0000
		(-0.0253, -0.0139)	(0.0276, 0.0475)	(0.0065, 0.0111)	(0.0020, 0.0045)	(-0.0152, -0.0091)
GRU-BS + GRU Reranker	GRU (Meloni et al., 2021)	p < 0.0001*	p < 0.0001*	p < 0.0001*	p < 0.0001*	p < 0.0001*
		(0.0312, 0.0442)*	(-0.0853, -0.0631)	*(-0.0200, -0.0148)	*(-0.0074, -0.0047)	* (0.0196, 0.0268)*
	Trans (Kim et al., 2023)	p < 0.0001*	p < 0.0001*	p < 0.0001*	p < 0.0001*	p < 0.0001*
		(0.0216, 0.0354)*	(-0.0629, -0.0380)	*(-0.0147, -0.0089)	*(-0.0049, -0.0018)	* (0.0128, 0.0204)*
	GRU-BS	p < 0.0001*	p < 0.0001*	p < 0.0001*	p < 0.0001*	p < 0.0001*
		(0.0141, 0.0249)*	(-0.0489, -0.0282)	*(-0.0114, -0.0066)	*(-0.0042, -0.0017)	* (0.0093, 0.0153)*
	GRU-BS + Trans. Reranker	p = 0.3882	p = 0.3625	p = 0.3625	p = 0.5804	p = 0.3727
		(-0.0058, 0.0058)	(-0.0122, 0.0097)	(-0.0029, 0.0023)	(-0.0011, 0.0017)	(-0.0030, 0.0035)
GRU-BS + Trans. Reranker	GRU (Meloni et al., 2021)	p < 0.0001*	p < 0.0001*	p < 0.0001*	p < 0.0001*	p < 0.0001*
		(0.0311, 0.0444)*	(-0.0830, -0.0620)	*(-0.0194, -0.0145)	*(-0.0077, -0.0050)	* (0.0193, 0.0264)*
	Trans (Kim et al., 2023)	<i>p</i> < 0.0001*	<i>p</i> < 0.0001*	<i>p</i> < 0.0001*	<i>p</i> < 0.0001*	<i>p</i> < 0.0001*
		(0.0214, 0.0355)*	(-0.0604, -0.0366)	*(-0.0142, -0.0086)	*(-0.0052, -0.0021)	* (0.0125, 0.0200)*
	GRU-BS	$p < 0.0001^*$	<i>p</i> < 0.0001*	<i>p</i> < 0.0001*	<i>p</i> < 0.0001*	<i>p</i> < 0.0001*
		(0.0137, 0.0252)*	(-0.0468, -0.0275)	^(-0.0110, -0.0064)	^ (-0.0045, -0.0020)	^ (U.U091, 0.0150)*
	GRU-BS + GRU Reranker	p = 0.6118	p = 0.6375	p = 0.6375	p = 0.4196	p = 0.6273
		(-0.0058, 0.0058)	(-0.0092, 0.0130)	(-0.0022, 0.0030)	(-0.0017, 0.0011)	(-0.0036, 0.0028)

Table 27: Reconstruction significance test results for WikiHan-aug. Asterisks indicates that Reconstruction System 1 performs better than Reconstruction System 2 with the corresponding test (*p*-value or CI).

Reconstruction System 1	Reconstruction System 2	ACC%↑	TED↓	TER↓	FER↓	BCFS ↑
GRU (Meloni et al., 2021)	Trans (Kim et al., 2023)	p = 1.0000	p = 1.0000	p = 1.0000	p = 0.9867	p = 1.0000
		(-0.0658, -0.0224)	(0.0571, 0.1453)	(0.0139, 0.0348)	(-0.0015, 0.0084)	(-0.0369, -0.0146)
	GRU-BS	p = 0.9991	p = 0.9914	p = 0.9975	p = 0.6772	p = 1.0000
		(-0.0516, -0.0059)	(-0.0009, 0.1053)	(0.0026, 0.0269)	(-0.0056, 0.0038)	(-0.0417, -0.0132)
	GRU-BS + GRU Reranker	p = 1.0000	p = 1.0000	p = 1.0000	p = 0.9953	p = 1.0000
		(-0.0813, -0.0370)	(0.0739, 0.1652)	(0.0193, 0.0404)	(0.0000, 0.0087)	(-0.0557, -0.0309)
	GRU-BS + Trans. Reranker	p = 1.0000	p = 1.0000	p = 1.0000	p = 0.9990	p = 1.0000
		(-0.0988, -0.0540)	(0.0935, 0.1891)	(0.0237, 0.0455)	(0.0020, 0.0111)	(-0.0600, -0.0349)
Trans (Kim et al., 2023)	GRU (Meloni et al., 2021)	p < 0.0001*	p < 0.0001*	p < 0.0001*	p = 0.0133	p < 0.0001*
		(0.0230, 0.0658)*	(-0.1463, -0.0571)	*(-0.0348, -0.0140	*(-0.0083, 0.0016)	(0.0143, 0.0370)*
	GRU-BS	p = 0.0468	p = 0.0142	p = 0.0199	p = 0.0199	p = 0.7331
		(-0.0087, 0.0405)	(-0.0935, -0.0028)	*(-0.0199, 0.0011)	(-0.0099, 0.0012)	(-0.0137, 0.0101)
	GRU-BS + GRU Reranker	p = 0.9119	p = 0.8416	p = 0.9029	p = 0.5751	p = 0.9998
		(-0.0398, 0.0096)	(-0.0186, 0.0562)	(-0.0032, 0.0143)	(-0.0043, 0.0063)	(-0.0272, -0.0081)
	GRU-BS + Trans. Reranker	p = 0.9969	p = 0.9920	p = 0.9907	p = 0.8543	p = 1.0000
		(-0.0562, -0.0086)	(0.0031, 0.0804)	(0.0017, 0.0197)	(-0.0024, 0.0083)	(-0.0320, -0.0126)
GRU-BS	GRU (Meloni et al., 2021)	p < 0.0010*	$p = 0.0086^*$	$p = 0.0025^*$	p = 0.3228	p < 0.0001*
		(0.0056, 0.0516)*	(-0.1048, -0.0027)	*(-0.0268, -0.0034)* (-0.0039, 0.0054)	(0.0141, 0.0413)*
	Trans (Kim et al., 2023)	p = 0.9532	p = 0.9858	p = 0.9801	p = 0.9801	p = 0.2669
		(-0.0410, 0.0084)	(0.0025, 0.0944)	(-0.0011, 0.0202)	(-0.0014, 0.0096)	(-0.0097, 0.0135)
	GRU-BS + GRU Reranker	p = 0.9963	p = 0.9985	p = 0.9985	p = 0.9923	p = 0.9945
		(-0.0571, -0.0059)	(0.0211, 0.1130)	(0.0047, 0.0253)	(0.0002, 0.0103)	(-0.0284, -0.0034)
	GRU-BS + Trans. Reranker	p = 1.0000	p = 0.9999	p = 0.9999	p = 0.9976	p = 0.9996
		(-0.0736, -0.0236)	(0.0416, 0.1381)	(0.0094, 0.0310)	(0.0019, 0.0124)	(-0.0330, -0.0076)
GRU-BS + GRU Reranker	GRU (Meloni et al., 2021)	p < 0.0001*	p < 0.0001*	p < 0.0001*	p = 0.0047*	p < 0.0001*
		$(0.0370, 0.0817)^{\star}$	(-0.1638, -0.0743)	*(-0.0402, -0.0194	*(-0.0087, 0.0002)	(0.0313, 0.0553)*
	Trans (Kim et al., 2023)	p = 0.0881	p = 0.1584	p = 0.0971	p = 0.4249	p < 0.0010*
		(-0.0093, 0.0385)	(-0.0553, 0.0186)	(-0.0141, 0.0032)	(-0.0063, 0.0042)	(0.0080, 0.0272)*
	GRU-BS	$p = 0.0037^*$	$p = 0.0015^*$	p = 0.0015*	$p = 0.0077^*$	$p = 0.0055^*$
		(0.0042, 0.0556)*	(-0.1130, -0.0189))*(-0.0253, -0.0042))* (-0.0104, -0.0000)	* (0.0028, 0.0282)*
	GRU-BS + Trans. Reranker	p = 0.9029	p = 0.8663	p = 0.8663	p = 0.8029	p = 0.7915
		(-0.0432, 0.0067)	(-0.0169, 0.0639)	(-0.0038, 0.0144)	(-0.0028, 0.0071)	(-0.0158, 0.0059)
GRU-BS + Trans. Reranker	GRU (Meloni et al., 2021)	p < 0.0001*	p < 0.0001*	p < 0.0001*	p = 0.0010*	p < 0.0001*
		(0.0548, 0.0978)*	(-0.1883, -0.0950)	*(-0.0455, -0.0241))* (-0.0110, -0.0018)	* (0.0354, 0.0597)*
	Trans (Kim et al., 2023)	p = 0.0031*	$p = 0.0080^*$	$p = 0.0093^*$	p = 0.1457	p < 0.0001*
		$(0.0081, 0.0559)^*$	(-0.0801, -0.0021))* (-0.0199, -0.0015)* (-0.0083, 0.0025)	(0.0123, 0.0321)*
	GRU-BS	p < 0.0001*	p < 0.0010*	p < 0.0010*	$p = 0.0024^*$	p < 0.0010*
		$(0.0224, 0.0724)^{*}$	(-0.1373, -0.0404))* (-0.0308, -0.0091)* (-0.0125, -0.0019)	* (0.0067, 0.0328)*
	GRU-BS + GRU Reranker	p = 0.0971	p = 0.1337	p = 0.1337	p = 0.1971	p = 0.2085
		(-0.0084, 0.0404)	(-0.0630, 0.0185)	(-0.0141, 0.0041)	(-0.0068, 0.0029)	(-0.0066, 0.0152)

Table 28: Reconstruction significance test results for Hóu. Asterisks indicates that Reconstruction System 1 performs better than Reconstruction System 2 with the corresponding test (*p*-value or CI).

Reconstruction System 1	Reconstruction System 2	ACC%↑	TED↓	TER↓	FER↓	BCFS ↑
GRU (Meloni et al., 2021)	Trans (Kim et al., 2023)	p = 0.9999	p = 1.0000	<i>p</i> = 1.0000	p = 0.9993	p = 1.0000
		(-0.0174, -0.0052)	(0.0567, 0.0886)	(0.0077, 0.0116)	(0.0005, 0.0021)	(-0.0167, -0.0119)
	GRU-BS	p = 0.9979	p = 1.0000	p = 1.0000	p = 1.0000	p = 1.0000
		(-0.0124, -0.0013)	(0.0487, 0.0810)	(0.0206, 0.0246)	(0.0028, 0.0045)	(-0.0154, -0.0100)
	GRU-BS + GRU Reranker	p = 1.0000	p = 1.0000	p = 1.0000	p = 1.0000	<i>p</i> = 1.0000
		(-0.0260, -0.0142)	(0.0846, 0.1159)	(0.0246, 0.0285)	(0.0046, 0.0062)	(-0.0210, -0.0160)
	GRU-BS + Trans. Reranker	p = 1.0000	p = 1.0000	p = 1.0000	p = 1.0000	p = 1.0000
		(-0.0252, -0.0131)	(0.0843, 0.1167)	(0.0246, 0.0286)	(0.0049, 0.0065)	(-0.0212, -0.0160)
Trans (Kim et al., 2023)	GRU (Meloni et al., 2021)	p < 0.0001*	p < 0.0001*	p < 0.0001*	p < 0.0010*	p < 0.0001*
		(0.0053, 0.0174)*	(-0.0890, -0.0570)	*(-0.0116, -0.0077))* (-0.0021, -0.0004)	* (0.0119, 0.0167)*
	GRU-BS	p = 0.0881	p = 0.1280	p = 1.0000	p = 1.0000	p = 0.0925
		(-0.0017, 0.0103)	(-0.0222, 0.0055)	(0.0114, 0.0145)	(0.0017, 0.0031)	(-0.0007, 0.0039)
	GRU-BS + GRU Reranker	p = 0.9992	p = 1.0000	p = 1.0000	p = 1.0000	p = 1.0000
		(-0.0153, -0.0025)	(0.0141, 0.0408)	(0.0155, 0.0184)	(0.0035, 0.0048)	(-0.0063, -0.0021)
	GRU-BS + Trans. Reranker	p = 0.9973	p = 1.0000	p = 1.0000	p = 1.0000	p = 1.0000
		(-0.0144, -0.0015)	(0.0136, 0.0414)	(0.0154, 0.0185)	(0.0038, 0.0051)	(-0.0065, -0.0020)
GRU-BS	GRU (Meloni et al., 2021)	p = 0.0021*	p < 0.0001*	p < 0.0001*	p < 0.0001*	p < 0.0001*
		(0.0017, 0.0123)*	(-0.0811, -0.0490)	*(-0.0246, -0.0206))* (-0.0045, -0.0028)	* (0.0100, 0.0154)*
	Trans (Kim et al., 2023)	p = 0.9119	p = 0.8720	p < 0.0001*	p < 0.0001*	p = 0.9075
		(-0.0102, 0.0017)	(-0.0057, 0.0216)	(-0.0145, -0.0115))* (-0.0031, -0.0017)	*(-0.0038,0.0007)
	GRU-BS + GRU Reranker	p = 1.0000	p = 1.0000	p = 1.0000	p = 1.0000	<i>p</i> = 1.0000
		(-0.0189, -0.0075)	(0.0218, 0.0485)	(0.0024, 0.0054)	(0.0011, 0.0024)	(-0.0082, -0.0034)
	GRU-BS + Trans. Reranker	p = 1.0000	p = 1.0000	p = 1.0000	p = 1.0000	<i>p</i> = 1.0000
		(-0.0182, -0.0063)	(0.0212, 0.0495)	(0.0024, 0.0055)	(0.0013, 0.0027)	(-0.0083, -0.0034)
GRU-BS + GRU Reranker	GRU (Meloni et al., 2021)	p < 0.0001*	p < 0.0001*	p < 0.0001*	p < 0.0001*	p < 0.0001*
		(0.0146, 0.0260)*	(-0.1160, -0.0845)	*(-0.0285, -0.0246))* (-0.0062, -0.0046)	* (0.0160, 0.0211)*
	Trans (Kim et al., 2023)	p < 0.0010*	p < 0.0001*	p < 0.0001*	p < 0.0001*	p < 0.0001*
		(0.0027, 0.0154)*	(-0.0410, -0.0141)	*(-0.0184, -0.0155))* (-0.0048, -0.0035)	* (0.0021, 0.0064)*
	GRU-BS	p < 0.0001*	p < 0.0001*	p < 0.0001*	p < 0.0001*	p < 0.0001*
		(0.0076, 0.0191)*	(-0.0490, -0.0219)	*(-0.0055, -0.0024))*(-0.0024,-0.0011)	* (0.0035, 0.0083)*
	GRU-BS + Trans. Reranker	p = 0.3474	p = 0.6066	p = 0.6066	p = 0.8830	<i>p</i> = 0.6066
		(-0.0053, 0.0072)	(-0.0133, 0.0137)	(-0.0015, 0.0015)	(-0.0003, 0.0009)	(-0.0024, 0.0023)
GRU-BS + Trans. Reranker	GRU (Meloni et al., 2021)	p < 0.0001*	p < 0.0001*	p < 0.0001*	ρ < 0.0001*	p < 0.0001*
		(0.0136, 0.0252)*	(-0.1165, -0.0843)	*(-0.0286, -0.0246))* (-0.0065, -0.0049)	* (0.0160, 0.0211)*
	Trans (Kim et al., 2023)	$p = 0.0027^*$	p < 0.0001*	p < 0.0001*	p < 0.0001*	p < 0.0001*
		(0.0014, 0.0146)*	(-0.0417, -0.0138)	*(-0.0185, -0.0154))* (-0.0051, -0.0038)	* (0.0021, 0.0065)*
	GRU-BS	p < 0.0001*	p < 0.0001*	p < 0.0001*	p < 0.0001*	p < 0.0001*
		$(0.0065, 0.0182)^*$	(-0.0501, -0.0218)	*(-0.0056, -0.0024))* (-0.0028, -0.0014)	* (0.0034, 0.0084)*
	GRU-BS + GRU Reranker	p = 0.6526	p = 0.3934	p = 0.3934	p = 0.1170	<i>p</i> = 0.3934
		(-0.0071, 0.0052)	(-0.0139, 0.0132)	(-0.0016, 0.0015)	(-0.0009, 0.0003)	(-0.0022, 0.0024)

Table 29: Reconstruction significance test results for Rom-phon. Asterisks indicates that Reconstruction System 1 performs better than Reconstruction System 2 with the corresponding test (*p*-value or CI).

Reconstruction System 1	Reconstruction System 2	ACC%↑	TED↓	TER↓	FER↓	BCFS ↑
GRU (Meloni et al., 2021)	Trans (Kim et al., 2023)	p = 1.0000	p = 1.0000	p = 1.0000	-	p = 1.0000
		(-0.0214, -0.0128)	(0.0249, 0.0495)	(0.0030, 0.0064)		(-0.0097, -0.0055)
	GRU-BS	p = 1.0000	p = 1.0000	p = 1.0000	-	p = 1.0000
		(-0.0217, -0.0130)	(0.0379, 0.0595)	(0.0153, 0.0180)		(-0.0106, -0.0067)
	GRU-BS + GRU Reranker	p = 1.0000	p = 1.0000	p = 1.0000	-	p = 1.0000
		(-0.0366, -0.0287)	(0.0682, 0.0882)	(0.0186, 0.0212)		(-0.0158, -0.0124)
	GRU-BS + Trans. Reranker	p = 1.0000	p = 1.0000	p = 1.0000	-	p = 1.0000
		(-0.0353, -0.0274)	(0.0672, 0.0869)	(0.0185, 0.0210)		(-0.0157, -0.0122)
Trans (Kim et al., 2023)	GRU (Meloni et al., 2021)	p < 0.0001*	p < 0.0001*	p < 0.0001*	-	p < 0.0001*
		(0.0128, 0.0216)*	(-0.0493, -0.0254)	*(-0.0064, -0.0031)*	*	(0.0055, 0.0097)*
	GRU-BS	p = 0.7419	p = 0.9848	p = 1.0000	-	p = 0.7591
		(-0.0040, 0.0041)	(-0.0011, 0.0222)	(0.0103, 0.0133)		(-0.0030, 0.0011)
	GRU-BS + GRU Reranker	p = 1.0000	p = 1.0000	p = 1.0000	-	p = 1.0000
		(-0.0189, -0.0115)	(0.0293, 0.0515)	(0.0137, 0.0165)		(-0.0083, -0.0045)
	GRU-BS + Trans. Reranker	p = 1.0000	p = 1.0000	p = 1.0000	-	p = 1.0000
		(-0.0178, -0.0101)	(0.0281, 0.0506)	(0.0136, 0.0164)		(-0.0082, -0.0043)
GRU-BS	GRU (Meloni et al., 2021)	p < 0.0001*	p < 0.0001*	p < 0.0001*	-	p < 0.0001*
		(0.0130, 0.0214)*	(-0.0590, -0.0380)	*(-0.0179, -0.0152) ³	*	(0.0067, 0.0105)*
	Trans (Kim et al., 2023)	p = 0.2581	p = 0.0152	p < 0.0001*	-	p = 0.2409
		(-0.0041, 0.0043)	(-0.0226, 0.0008)	(-0.0133, -0.0103)	*	(-0.0011, 0.0030)
	GRU-BS + GRU Reranker	p = 1.0000	p = 1.0000	p = 1.0000	-	p = 1.0000
		(-0.0191, -0.0115)	(0.0201, 0.0389)	(0.0022, 0.0043)		(-0.0072, -0.0038)
	GRU-BS + Trans. Reranker	p = 1.0000	p = 1.0000	p = 1.0000	-	p = 1.0000
		(-0.0180, -0.0102)	(0.0191, 0.0380)	(0.0021, 0.0042)		(-0.0071, -0.0036)
GRU-BS + GRU Reranker	GRU (Meloni et al., 2021)	p < 0.0001*	p < 0.0001*	p < 0.0001*	-	p < 0.0001*
		(0.0286, 0.0365)*	(-0.0882, -0.0684)	*(-0.0212, -0.0186) ³	*	(0.0124, 0.0159)*
	Trans (Kim et al., 2023)	p < 0.0001*	p < 0.0001*	p < 0.0001*	-	p < 0.0001*
		(0.0116, 0.0190)*	(-0.0517, -0.0295)	*(-0.0166, -0.0137) ³	*	(0.0045, 0.0084)*
	GRU-BS	p < 0.0001*	p < 0.0001*	p < 0.0001*	-	p < 0.0001*
		(0.0115, 0.0190)*	(-0.0390, -0.0199)	*(-0.0043, -0.0022)	*	(0.0037, 0.0072)*
	GRU-BS + Trans. Reranker	p = 0.3779	p = 0.4409	p = 0.4409	-	p = 0.5431
		(-0.0022, 0.0047)	(-0.0097, 0.0079)	(-0.0011, 0.0009)		(-0.0014, 0.0017)
GRU-BS + Trans. Reranker	GRU (Meloni et al., 2021)	p < 0.0001*	p < 0.0001*	p < 0.0001*	-	p < 0.0001*
		(0.0274, 0.0353)*	(-0.0875, -0.0674)	*(-0.0211, -0.0185)	*	(0.0122, 0.0158)*
	Trans (Kim et al., 2023)	p < 0.0001*	p < 0.0001*	p < 0.0001*	-	p < 0.0001*
		(0.0103, 0.0180)*	(-0.0509, -0.0282)	*(-0.0165, -0.0136) ³	*	(0.0044, 0.0083)*
	GRU-BS	p < 0.0001*	p < 0.0001*	p < 0.0001*	-	p < 0.0001*
		(0.0102, 0.0180)*	(-0.0380, -0.0189)	· (-0.0042, -0.0021)	*	(0.0036, 0.0071)*
	GRU-BS + GRU Reranker	p = 0.6221	p = 0.5591	p = 0.5591	-	p = 0.4569
		. (-0.0047, 0.0022)	(-0.0077, 0.0100)	(-0.0009, 0.0011)		(-0.0018, 0.0014)

Table 30: Reconstruction significance test results for Rom-orth. Asterisks indicates that Reconstruction System 1 performs better than Reconstruction System 2 with the corresponding test (*p*-value or CI).

Bear	n Search		Reflex Prec	Reflex Prediction (based on protoform candidates)									nking Re	sult
rank	\hat{p}_i^{bs}	mi	Cantonese	Hakka	Jin	Mandarin	Hokkien	Wu	Xiang	ri		rank	ρ̂ ^{rk}	Si
0	kjwet入	-0.2299	 ky:t⊦	ki̯et∔	t͡ɕyə?∔	tâyx1	ku̯at⊦	t͡ɕyɪ?⊺	kşja	0.2857	$\sim z$	0	kwit入	0.3318
1	kit入	-0.2866	 ket1	kit⊦	⊧, reija	۲iat	kit⊦	۲?ıjât	t⊋i≀	0.1429	\checkmark	1	kjwet入	0.1301
2	kwit入	-0.3882	 k∾et1	ki̯ut₊	t⊊yə?∔	tayx1	kut-l	teyi?1	t͡sy∕	0.5714	$\overline{}$	2	kjwit入	-0.0567
3	kjɨt入	-0.3979	 ket1	kit⊦	۱،۲eja	fail	kit⊦	۲٩ <u>۱</u>	t€i∕l	0.1429	\checkmark	3	kit入	-0.1066
4	kjwit入	-0.5967	 k∾et1	k <u>i</u> ut⊣	tsya?-	tayx1	kut⊦	f sy171	t⊊yn√	0.4286		4	kj i t入	-0.2179
5	kjet入	-0.7380	 ki:t+	kiet∔	ا۔?eiُa	faia1	kiεt⊦	۲?ıiât	t⊊ie≀	0.1429	\longrightarrow	5	kjet入	-0.5580
6	kεt入	-0.8579	 ka:t+	kat∔	⊧?eį́a	רַפַוָּפ	kat I	ka?1	kγ́∕	0.0000	\longrightarrow	6	kεtλ	-0.8579
			kwet 1	kit⊦	t͡ɕyə?∔	tsy1	kįεt⊦	tsyı?1	t͡ɕy∕	-				

Figure 4: Successful reranking of 橘 kwit入 'mandarin orange'.

Bean	n Search		Reflex Pred	Reflex Prediction (based on protoform candidates)							nking Re	sult
rank	\hat{p}_i^{bs}	mi	Cantonese	Hakka	Mandarin	Hokkien	Wu	ri		rank	<i>̂p</i> i ^{rk}	Si
0	kwaj去	-0.2234	 kwo:1	kui1	k⁺u̯ai̯∖	kue≀	kwę1	0.0000	\ 7	0	γwaj去	0.3959
1	?waj去	-0.3368	 wu:y+	vojī	<u>ue</u> j\	ŭe≀	ωę́1	0.0000	X_{7}	1	γwoj去	0.2032
2	γwaj去	-0.3601	 wu:y₁	fi1	xueį≀	ue-I	ĥwe∤	0.6000	\sim	2	kwaj去	-0.2234
3	kwoj去	-0.3787	 k∾u:y⊣	kui⊺	ku̯ei̯∖	kue√	kuę́1	0.0000	\checkmark	3	?waj去	-0.3368
4	γwoj去	-0.5528	 wu:y4	fi1	xuei≀	hue⊦	ĥué∤	0.6000		4	kwoj去	-0.3787
5	?woj去	-0.5998	 wu:ył	ve1	uei	ue√	ue1	0.0000	\longrightarrow	5	?woj去	-0.5998
6	kʰwaj去	-0.8277	 fuːy-	kʰu̯ai̯٦	k ^h uai _l i	k⁺ue√	k ^h wę1	0.0000	\longrightarrow	6	kʰwaj去	-0.8277
			k ^h uːy1	fi1	хџеј≀	ku̯e⊣	huẹ̯́ł	-				

Figure 5: Successful reranking of 繪 ywaj去 'to draw'.

Bean	n Searcl		Reflex Prec		Rera	nking R	esult			
rank	\hat{p}_i^{bs}	mi	Cantonese	Mandarin	Hokkien	ri		rank	$\hat{p}_i^{\mathbf{rk}}$	Si
0	sew平	-0.1557	 si:u̯ๅ	۲µsia	siaul	0.0000	$\backslash \nearrow$	0	suw平	0.0196
1	suw平	-0.4004	 seyl	soʊ̯ˈ	saul	0.3333	X	1	suw上	-0.0471
2	suw上	-0.4671	 seų1	soŭ1	saų∛	0.3333	\sim	2	sew平	-0.1557
3	sju平	-0.5906	 seyl	۶y٦	۲ia	0.0000	\longrightarrow	3	sju平	-0.5906
4	sew上	-0.5909	 siːu̯1	⊾iau	۶jau	0.0000	\longrightarrow	4	sew上	-0.5909
5	sju上	-0.9313	 sey1	εγJ	۶i	0.0000	\longrightarrow	5	sju上	-0.9313
			seų1	soʊ̯]	sxV	-				

Figure 6: Unsuccessful reranking of 艘 sew平 'small boat'.

														-		
Bean	n Search			Reflex Prec	diction (b	based o	n proto	form candi	dates)				Reranking Result			sult
rank	\hat{p}_i^{bs}	mi		Cantonese	Gan	Hakka	Jin	Mandarin	Hokkien	Wu	Xiang	ri		rank	$\hat{p}_i^{\mathbf{rk}}$	Si
0	kjun平	-0.1272	·	ken⊺	t͡ɕyn√	kjun∤	t͡ɕỹŋ⅃	t͡ɕyn1	kun1	fayn	t͡ɕyn⊣	0.8750	$\overline{}$	0	kwin平	1.0834
1	kwin平	-0.1766		k∾en1	t͡ɕyn√	ki̯un∤	t͡ɕỹŋ⅃	t͡ɕyn⊺	kun1	tern 1	t͡ɕyn⊣	1.0000	\sim	1	kjun平	0.9753
2	kin平	-0.6991		ken1	t⊊in≀	kin≀	tົ⊊ĩŋ⅃	۲nia	kin1	۲ŋıaf	t͡ɕin⊣	0.0000	\longrightarrow	2	kin平	-0.6991
3	kwen平	-0.9995		ky:n1	t⊊yen√	ki̯en≀	t͡ɕye⊥	۲apan	ku̯an٦	۲øya	t͡ɕyẹ⊣	0.0000	\longrightarrow	3	kwen平	-0.9995
4	kjin平	-1.0021		ken1	tâin	kin≀	ເຣົ້າງ∟	tiin1	kin٦	۲qīat	t͡ɕin⊣	0.0000	\longrightarrow	4	kjin平	-1.0021
5	kjon平	-1.2847		ki:n1	۲nsjat	ki̯en≀	€ie」	רחsija ד	kຼiɛn٦	۲qīat	t⊊jã⊣	0.0000	\longrightarrow	5	kjon平	-1.2847
				k∾en1	t͡syn√	kjun∕	t͡ɕỹŋ⅃	t͡ɕyn⊺	kun⊺	fern	t͡ɕyn⊣	-				

Figure 7: Unsuccessful reranking of 君 kjun平 'sovereign'.

Bean	n Search		Reflex Prediction (based on protoform candidates)					Rera		
rank	\hat{p}_i^{bs}	mi	Italian	Spanish	Portuguese	ri		rank	ρ̂ ^{rk}	Si
0	dıskrɛptɪam	-0.0876	 diskrɛtsa	diskrepθja	di∫kıεpsje	0.0000	$\backslash $	0	dıskrɛpantıam	0.4777
1	dıskrɛpantıam	-0.1073	 diskrepantsa	diskrepanθja	di∫kɹɨpẽŋsje	1.0000	X	1	dıskrepantıam	0.1887
2	dıskrepantıam	-0.3963	 diskrepantsa	diskrepanθja	di∫kɹɨpẽŋsje	1.0000	\sim	2	dıskrɛptıam	-0.0876
3	dıskrɛpsɪam	-0.4401	 diskressia	diskrepsja	di∫kuipsie	0.0000	\longrightarrow	3	dıskrɛpsɪam	-0.4401
4	dıskrɛpsantıam	-0.4729	 diskressantsa	diskrepsanθja	di∫kɹɨpsẽŋsje	0.0000	\longrightarrow	4	dıskrɛpsantıam	-0.4729
			diskrepantsa	diskrepan0ja	diſkɹɨpẽŋsje	-				

Figure 8: Succssful reranking of discrepantiam 'discordance'.

Beam Search			Reflex Prediction (based on protoform candidates)					Rera	nking Result	
rank	\hat{p}_i^{bs}	mi	French	Italian	Spanish	r _i		rank	ρ̂ r k	Si
0	εksεkwikεrε	-0.2887	 εgzeke	ezekwire	ekseyir	0.0000	\ 7	0	εkssεkwi	-0.0423
1	εksεkwɪtarɛ	-0.3491	 εgzekite	ezekwitare	eksekitar	0.0000	\mathbb{N}	1	εksεkwi	-0.0641
2	εkssεkwɪtarε	-0.3907	 εgzekite	ezekwitare	eksekiðar	0.0000	XX	2	εksεkwikεrε	-0.2887
3	εksεkwirε	-0.4299	 εgzeke	ezekwire	ekseyir	0.0000	X	3	εksεkwɪtarɛ	-0.3491
4	εkssεkwi	-0.4323	 εgzekyte	ezegwire	ekseyir	0.6667	- Mγ	4	εkssεkwɪtarε	-0.3907
5	εksεkwi	-0.4541	 εgzekyte	ezegwire	ekseyir	0.6667	/ \	5	εksεkwirε	-0.4299
			εgzekyte	ezegwire	exekutar	-		-		

Figure 9: Successful reranking of exsequi 'follow after'.

Beam Search			Reflex Prediction (based on protoform candidates)						Rera	nking Result	
rank	\hat{p}_i^{bs}	mi	French	Italian	Spanish	Portuguese	ri		rank	ρ̂ ^{rk}	Si
0	prokrastinarɛ	-0.1000	 prokrastine	prokrastinare	prokrastinar	puukua∫tinau	0.7500	\searrow	0	prokrastinarɛ	0.4696
1	prokrastinarɛ	-0.1154	 psoksastine	prokrastinare	prokrastinar	pıukıe∫tinaı	1.0000	\sim	1	prokrastınarɛ	0.3388
2	prokrastınari	-0.2423	 psoksastine	prokrastinare	prokrastinar	pıukıa∫tinaı	0.7500	\searrow	2	prokrastinari	0.3221
3	prokrastinari	-0.2629	 psoksastine	prokrastinare	prokrastinar	pıukıe∫tinaı	1.0000	\sim	3	prokrastınari	0.1965
4	prokrastinare	-0.6461	 psoksastine	prokrastinare	prokrastinar	pıukıe∫tinaı	1.0000	\longrightarrow	4	prokrastinare	-0.0611
5	prokrastınare	-0.6774	 psoksastine	prokrastinare	prokrastinar	puukua∫tinau	0.7500	\longrightarrow	5	prokrastinare	-0.2386
			prokrastine	prokrastinare	prokrastinar	pıukıe∫tinaı	-				

Figure 10: Unsuccessful reranking of procrastinare 'to defer'.

Beam Search			Reflex Prediction (based on protoform candidates)							Rera	nking Result	
rank	$\hat{\rho}_i^{bs}$	mi	Romanian	French	Italian	Spanish	Portuguese	ri		rank	$\hat{\rho}_i^{\mathbf{rk}}$	Si
0	aıkwınɔktıʊm	-0.1209	 ekinokts	ekinoks	ekwinɔtso	ekinokθjo	ekinusjʊ	0.2000	$\overline{\mathbf{x}}$	0	ɛkwɪnɔktɪʊm	0.3296
1	ɛkwɪnɔktɪʊm	-0.1384	 ekinoktsiw	ekinoks	ekwinɔtsio	ekinokθjo	ekinosjʊ	0.8000	\prec_{r}	1	ɛkwɪnoktɪʊm	0.1556
2	aɪkwɪnoktɪʊm	-0.2969	 ekinokts	ekinoks	ekwinɔtso	ekinokθjo	ekinosjo	0.2000	\checkmark	2	aıkwınɔktıʊm	-0.0039
3	ɛkwɪnoktɪʊm	-0.3124	 ekinoktsiw	ekinoks	ekwinɔtsio	ekinokθjo	ekinɔsjʊ	0.8000		3	ɛkwinɔktɪʊm	-0.0256
4	aɪkwinɔktɪʊm	-0.4513	 ekinokts	ekinoks	ekwinoktsio	ekinokθjo	ekinusjʊ	0.2000	\checkmark	4	aıkwınoktıʊm	-0.1799
5	ɛkwinɔktɪʊm	-0.4936	 ekinoktsiw	ekinoks	ekwinɔtsio	ekinokθjo	ekinosjʊ	0.8000		5	aıkwinoktıʊm	-0.3343
			ekinoktsiw	ekinoks	ekwinɔtsio	ekinokθjo	ekwinɔsjʊ	-				

Figure 11: Unsuccessful reranking aequinoctium 'equinox'.