

1 Introduction

The word order of natural languages has been initially characterized by Behagel’s laws¹. These laws state elements that are related to each other tend to be placed close together in a sentence. Until recently, these laws have been summarized as dependency length minimization²⁻⁴. Dependency lengths are the distances between linguistic units that are related to each other in a sentence. This theory states that languages had evolved to minimize the sum of these distances so that the users can easily produce and comprehend sentences by reducing cognitive load. It has been shown 37 languages follow this principle².

One potential explanation for this phenomenon is that human have limited working memory capacity⁵. This limitation forces languages to evolve by placing related words closer to each other, allowing speakers and listeners could easily integrate words and process semantic information from memory. These ideas are known as Dependency length theory (DLT)⁶. One key prediction of DLT is the locality effects: longer dependencies leads to longer reading times and vice versa, which has been supported by various psycholinguistic studies^{7;8}.

Recently, information-theoretic approaches has been proposed to formalize the relationship between incremental working memory usages in languages processing and sequential order, known as memory-surprisal tradeoff⁹. This theory has shown the ease of comprehension is determined by the resources allocated to store elements (e.g., words) in working memory, hence there exists a tradeoff between memory usage and comprehension difficulty in a corpora of 54 languages by using surprisal (shannon entropy conditioned on limited memory system). However, it is still unclear how the architecture of different language model model, tokenization methods and the syntatic structures of different languages affect surprisal measures and thus memory-surprisal tradeoff. Here, we aim to investigate these factors by implementing a naive hidden markov model and a simple LSTM network and different tokenization methods on corpora of 10 chosen languages.

2 Method

2.1 Architecture and the tokenizer of Language Models

Here we implement two different language models: a simple markovian feed-foward neural network language model (NNLM) and a long-short term memory (LSTM) network.

Neural Network Language Model (NNLM)

The NNLM recieves a sequeence of tokens $x = \{w_0, x_1, \dots, x_t\}$ with embeddings $e = \{e_0, e_1, \dots, e_t\}$ as input and compute the probability of the next word given a context window with size n :

$$P(w_t|w_{t-n+1}, \dots, w_{t-1}) \quad (1)$$

The embedding is then passed to a hidden layer:

$$h_t = \tanh(W_{inh}e_t + b_{ih}) \quad (2)$$

The resulting hidden state is then passed to an output layer with softmax activation to compute the probability distribution of the next token:

$$P(w_t|w_{t-n+1}, \dots, w_{t-1}) = \text{softmax}(W_{out}h_t + b_{out}) \quad (3)$$

The tokenizer used for NNLM is a simple whitespace tokenizer that splits the text into words based on spaces. However, this method is problematic for languages without explicit word boundaries (e.g., Chinese, Japanese). **(Answer for Exercise1)** In addition, given that NNLM is a next-token prediction model, only last few tokens are predicted due to the limited size of the sentence. For example, a test sentence with a fixed length T were passed to the model, only the last $T - N + 1$ tokens were predicted because the first $N - 1$ tokens do not have enough context for a memory window of length N . To solve this problem, we padded the begining of each sentence with $N - 1$ special tokens `<pad>` so that all tokens in the sentence could be predicted. Our modification is maded in the `NgramsLanguageModelDataSet(Dataset)` class in the `dataloader.py` file.

Long Short-Term Memory (LSTM) Network

(Answer for Exercise 3)

A normal Recurrent Neural Network would predict the next word w_t given its preceding context (w_1, \dots, w_{t-1}) by maintaining a hidden state h_t that acts as a memory. However, such a vanilla RNN struggles with long-term dependencies due to instability in the gradient propagation (vanishing/exploding gradients).

So, we rather prefer Long Short-Term Memory networks to address this limitation. LSTM introduces an additional hidden state: a cell state c_t and a system of gates that regulate information flow. This gating mechanism allows LSTMs to selectively retain and discard information, effectively learning long-range dependencies through the cell state.

In PyTorch, implementing an LSTM-based RNNLM involves an ‘nn.Embedding’ layer to convert token IDs to dense vectors (x_t) , an ‘nn.LSTM’ module for the recurrent processing (handling h_t and c_t updates), and an ‘nn.Linear’ layer to project the LSTM’s final hidden state to a vocabulary-sized output.

The biggest difference between our RNNLM and the previous NNLM we used lies in the way they handle context and memory. The NNLM uses a fixed-size context window of length n and concatenates the embeddings of the $n - 1$ preceding words as input for a single forward pass. It treats each prediction as independent, given its immediate $n - 1$ left neighbors in the sequence, without carrying any information beyond this window.

On the other hand, our RNNLM processes the input sequence recurrently, which means that it maintains the dynamic hidden state (h_t) and the cell state (c_t) that act as a “memory” of all preceding tokens in the sequence. This allows the RNNLM to capture much longer dependencies than a fixed window NNLM.

The most important challenges when implementing the RNNLM in PyTorch are :

- Because of their inherent recurrent structure, LSTMs are likely to suffer from gradients issues (exploding or vanishing), breaking calculation during training with NaN or infinite values. However we haven’t faced this issue as we used rather short sequences. In addition, in these conditions it can be meaningful to add clipping features when gradients get too big or too small.
- For independent n-grams, as we have been using with our previous model, each n-gram is a fresh sequence. But when we are dealing with continuous sequences, LSTMs have to maintain internal hidden and cell states. Thus, we need to pass these states from the end of one batch as the initial states for the next, which requires subtle handling of the variable sequence lengths and batching.
- These considerations are more real-life issues, but we’ve been able to run our model as a POC. But because of the recurrent nature of LSTMs, for long sequences, training can be much slower compared to feedforward models.

GPT-2 Model

(Answer for Exercise2) To compare the surprisal measures between different tokenization methods, we also used a simplified GPT-2 model with byte-pair encoding (BPE) tokenizer to control for vocabulary and vocabulary size of the model. The GPT-2 model is a transformer-based language model that uses self-attention mechanisms to capture long-range dependencies in text. It is a compact GPT-2 decoder-only transformer with a 256-dimensional embedding space, 4 transformer blocks and a hidden size of 256. Each block contains multi-head self-attention (implemented via Conv1D projections), a 1024-dimensional feedforward MLP with GELU activation and residual connections with layernorm and dropout. A final projection layer maps the hidden states to the vocabulary.

The BPE tokenizer splits the text into subword units, allowing the model to handle out-of-vocabulary words and capture morphological information.

Randomization Protocol of Syntactic Structures

(briefly describe how we do exercise 5 here)

Choice of corpora and languages

(Answer for Exercise 4) Regarding selection criteria for the corpora, we want to ensure certain things. First, the languages must be present in UD. Second, we want to prioritize languages whose corpora are large enough to estimate language models (though we will be limited by our compute capabilities, so this is more a real life concern). Third, we want to aim for diversity of families and geographic regions to make the results more general.

So we can go for the following UD available languages:

- **English** – typologically fairly fixed SVO order.
- **Spanish** – another SVO, fixed-order example, Romance family.
- **German** – but typologically more flexible (WALS shows it lacks a single dominant SVO vs. SOV) → good “semi-free” testing case.
- **Turkish** – typologically strongly SOV/OV (head-final) and relatively more fixed order (though topic-marking allows variation) → fixed-order side.
- **Hungarian** – typologically discourse-configurational with flexible word order → good free-order candidate.
- **Russian** – Slavic, moderately flexible word order (case marking allows alternations) → another free-order candidate.
- **Japanese** – strongly SOV, fairly fixed in canonical word order → fixed side.
- **Finnish** – Uralic language, fairly flexible word order due to rich morphology → free-order side.
- **Arabic (Modern Standard Arabic)** – typologically VSO or SVO depending on dialect; variation makes it interesting → flexible side.
- **Chinese (Mandarin)** – typologically SVO and relatively fixed in major clauses → fixed side.

Randomization Protocol of Syntactic Structures

(Answer for Exercise 5: Randomization methods)

By studying the provided script dependencies.py we learned its sophisticated approach to preserving dependency structure during randomization. The module offers two methods: `shuffle tree()` which preserves dependency relationships while randomizing word positions, and `shuffle projective()` which maintains projectivity constraints. However, for our information locality research, we developed a custom `conllu parser.py` module to have more direct control over word forms and seamless integration. We implemented two complementary randomization strategies to test distinct linguistic hypotheses.

- The first is complete randomization, where word order is entirely shuffled without preserving any structural information. This serves as the strongest baseline to test Hypothesis H1 (Information Locality): natural language word order is optimized to cluster predictively relevant information locally. If $\text{Surprisal}_{\text{natural}} < \text{Surprisal}_{\text{random}}$, this demonstrates that word order organization reduces prediction uncertainty, as proposed by Hahn et al. (2021).
- The second strategy is dependency-preserving randomization, inspired by dependencies.py’s `shuffle tree()` method. In this randomization method, we shuffle the linear order of the words in a sentence while preserving all original dependency relations. Each token in a dependency tree is associated with two indices: a linear position ID (*tid*) and the position ID of its syntactic head (*head tid*). The algorithm first applies `random.shuffle()` to the list of linear position IDs, producing a randomized target order. A mapping from old to new positions is then constructed, and each token’s head index is updated using this mapping so that all governor–dependent pairs are preserved. This procedure maintains the topology of the dependency tree while altering

the surface word order, thereby breaking grammatical word order but retaining the underlying syntactic structure. This dependency-preserving shuffle serves as a controlled baseline for dependency length minimization (DLM) experiments. Compared to fully random shuffling, it provides a more structurally informed contrast condition, as it isolates the effect of linear order independent of the dependency graph itself. This tests Hypothesis H2 (Dependency Length Minimization): the optimization in natural language specifically operates by placing syntactically related words closer together (Futrell et al., 2015).

Experimental Procedures

(briefly describe how we do exercise 7 here)

3 Results and Discussion

3.1 Effects of tokenization methods on surprisal measures

(Answer for Exercise2 Continued) The GPT-2 uses a Byte-Pair Encoding (BPE) tokenizer, which will decompose unknown words into fragments of known tokens, meaning that BPE guarantees any string is tokenizable. In addition, common patterns (e.g., the, ing) become short token sequences but for rare/unseen patterns will stay as small tokens, suggesting that an unknown sequence requires longer sequences and therefore giving higher surprisals. For example, if "rareword" is a common pattern in the training corpora, that means this pattern can be represented as a single token, hence the surprisal will be $p(\text{rareword}|\text{context}) = a$. However, if it is less likely, let's say it will be decomposed into $[t_1, t_2, t_3, t_4]$, even the model learns that these tokens are more likely (i.e., $p(t_i) = b > a$), the surprisal will still be larger due to the multiplicative nature of the tokens – $p(\text{rareword}|\text{context}) = \prod_{i=1}^n p(t_i) = \sum_{i=1}^n \log p(t_i)$, as shown in Figure 3.

The properties of BPE make it very difficult to compare two different approaches. To compare them, first an untrained GPT-2 model that has been trained on the same corpora and maintain the same vocabulary. To capture the unknown sequences that have not been shown to GPT-2 tokenizer, we have to align and concatenate the tokens so that it matches the unknown words in the context, and sum their surprisals for comparison. Then a statistical test is required to test if the KL-divergence between two distributions of the surprisals does not arise from chance, which tells us whether the two distributions are different from each other (perhaps a permutation test).

To conclude, it is not easy to compare the surprisals between the two distributions as it required sophisticated alignments and design of the GPT-2 and training procedures. From a theoretical point of view, for unknown sequences, most unknown words will receive a larger surprisal in the case of BPE tokenizer, while the model with a simple `<unk>` exhibits a more narrower range of surprisal due to weaker dependency of the context.

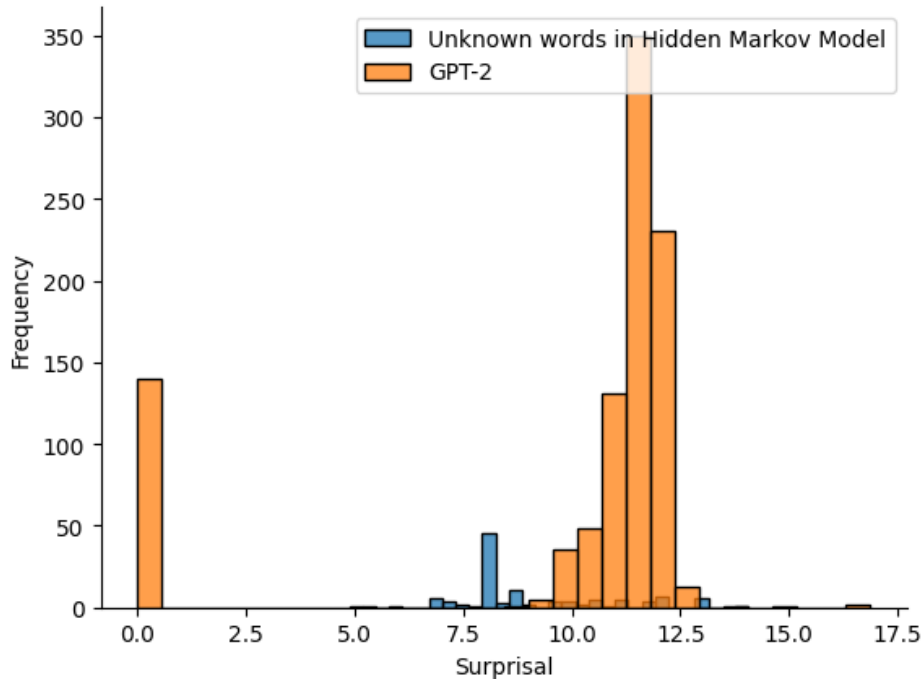


Figure 1: The distribution of surprisal for two different tokenization methods: whitespace tokenizer (blue) and byte-pair encoding tokenizer (orange) on English corpus using NNLM and GPT-2 model. The BPE tokenizer results in a more skewed distribution with a longer tail, indicating that it produces higher surprisal values for certain tokens compared to the whitespace tokenizer. This suggests that the choice of tokenization method can impact the surprisal measures obtained from language models.

3.2 Observations on artificial corpora randomized

Answer for Exercise 6

For statistical testing, we implemented a permutation test to compare the means of dependency lengths between natural corpora and randomized corpora by shuffling the labels 1000 times. The null hypothesis assumes that the average dependency length in the randomized dataset is not different from that in the natural corpora. After running the permutation test, the null hypothesis yielded a p-value of 0 across 1000 iterations, indicating that we can reject the null hypothesis and accept the alternative hypothesis: the average dependency length in the randomized dataset is significantly different from that in the natural corpora.

There is a clear and consistent difference. As shown in our results, all languages exhibit **longer** average dependency lengths after word order randomization in both methods. For example, in English: 11.01 (natural) \rightarrow 32.83 (complete shuffle) — a 198% increase. This increase is observed **across all ten languages** and **both randomization methods**. Notably, when dependency relations are preserved, the increase in dependency length is **smaller** than in full randomization, which breaks all relations. These results strongly support the hypothesis proposed in the paper: natural language word order is optimized to minimize dependency length, keeping syntactically related words closer together.

Interestingly, free word order languages do **not** systematically exhibit longer dependency lengths. For example, Turkish: 8.83 vs. English: 11.01; Russian: 60.29 vs. German: 60.15. In fact, Turkish (a free word order language) has one of the shortest dependency lengths, while Spanish (fixed SVO) has the longest.

The choice of randomization method is crucial for drawing valid conclusions. Complete randomization breaks all syntactic relations and dependencies without any constraints, which can only show that natural language is better than a completely randomized language but does not allow more nuanced observations. In contrast, dependency-preserving shuffle (shuffle tree) maintains structural constraints

and respects linguistic dependencies. This method provides a suitable baseline for controlled experiments, allowing us, for instance, to systematically manipulate the order of nouns and verbs while preserving dependency relations, isolating the effect of word order on dependency length.

Finally, comparing dependency lengths under complete randomization and dependency-preserved randomization reveals a clear cross-linguistic pattern. When a sentence is fully randomized without preserving any syntactic structure, dependency lengths increase dramatically, as expected in a system with no grammatical constraints. In contrast, shuffling only the linear order while preserving original dependency relations leads to a much smaller increase in dependency length.

This contrast suggests that natural languages follow structural regularities that constrain linearization, with dependency structure being one such principled constraint. This raises a broader question: is this locality effect a universal property of language? More specifically, beyond dependency locality, do other syntactic frameworks—such as phrase-structure grammar and constituent-based models—also exhibit some form of locality?

In summary, natural language universally shows **optimized word order** for dependency length minimization, **regardless** of whether the language is classified as "free" or "fixed" word order. The choice of randomization procedure is essential for isolating and identifying this specific optimization principle.

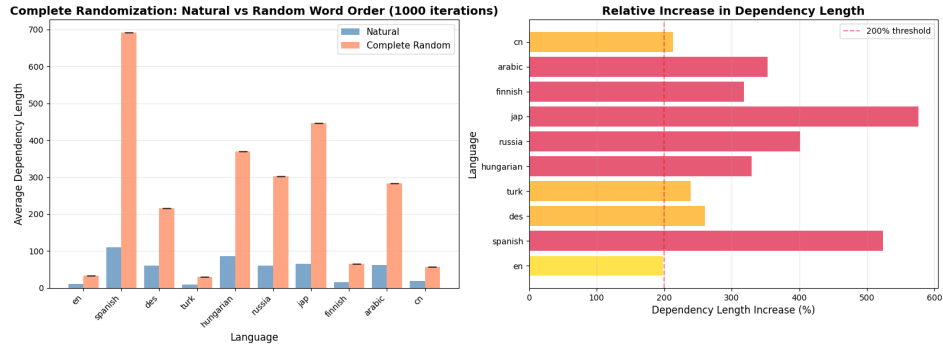


Figure 2: Average increase: 341.4%. Minimum increase: 198.0% (en). Maximum increase: 576.6% (jap). P-value range: 0.000 - 0.000 All p-values ≤ 0.05

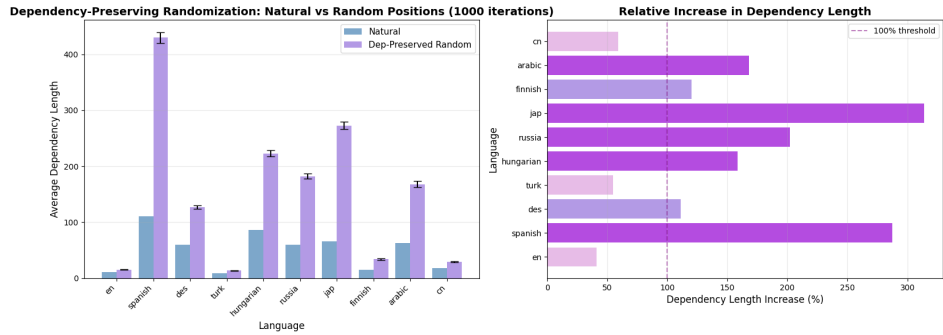


Figure 3: Average increase: 151.7%. Minimum increase: 41.1% (en) Maximum increase: 313.9% (jap). P-value range: 0.000 - 0.000 All p-values ≤ 0.05

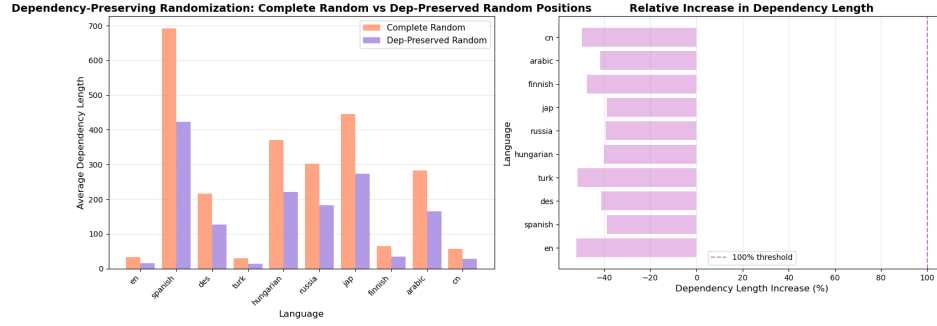


Figure 4: Statistical Summary: Dependency-Preserving Randomization relative to Complete Random. Dep-preserved increase compared to complete random: -44.1%. Minimum increase: -52.0% (en). Maximum increase: -38.7% (jap). All p-values < 0.05.

4 References

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