

Democratizing Advanced Attribution Analyses of Generative Language Models with the Inseq Toolkit

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Abstract

Inseq¹ is a recent toolkit providing an intuitive and optimized interface to conduct feature attribution analyses of generative language models. In this work, we present the latest improvements to the library, including efforts to simplify the attribution of large language models on consumer hardware, additional attribution approaches, and a new client command to detect and attribute context usage in language model generations. We showcase an online demo using Inseq as an attribution backbone for context reliance analysis, and we highlight interesting contextual patterns in language model generations. Ultimately, this release furthers Inseq's mission of centralizing good interpretability practices and enabling fair and reproducible model evaluations.

Keywords

Natural Language Processing, Generative Language Models, Feature Attribution, Python Toolkit

1. Introduction

Feature attribution methods have been widely adopted in NLP to quantify the importance of input tokens in driving language models' (LMs) predictions [1]. While some works used feature attribution to analyze generative NLP models, focusing mainly on machine translation [2, 3, 4, i.a.], most analyses in this area focused on classification due to the initial popularity of BERT-based encoders [5] and the challenges of autoregressive generation [6]. Although several post-hoc interpretability tools are available, few support generative LMs [7, 8, 9], often requiring ad-hoc wrappers to enable interoperability with the popular Transformers library [10] commonly used by NLP practitioners.

Inseq [11] is a Python library offering native compatibility with Transformers and supporting advanced methods and customizations. Inseq centralizes access to a broad set of feature attribution methods, sourced in part from the Captum [12] framework, enabling fair comparisons

¹Library: <https://github.com/inseq-team/inseq>, Docs: <https://inseq.org>. This paper refers to release v0.6.0. *Late-breaking work, Demos and Doctoral Consortium, colocated with The 2nd World Conference on eXplainable Artificial Intelligence: July 17–19, 2024, Valletta, Malta*

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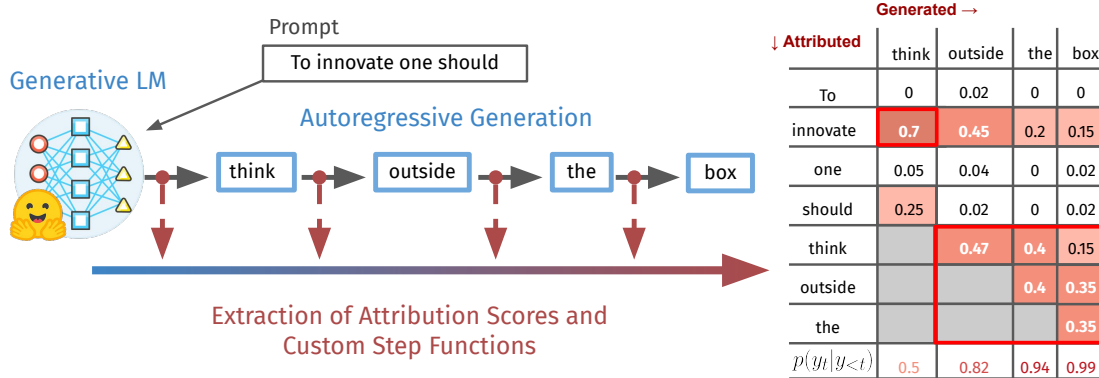


Figure 1: Example of Inseq usage with a generative LM. Given a prompt, attribution scores and next-step probabilities are extracted from the model at every generation step, with a final visualization aggregating values at the token level. Highlighted areas in the output show that the model relies on the keyword “innovate” to begin the idiomatic expression “think outside the box” at relatively low confidence ($p = \emptyset.5$). However, importance shifts to previous tokens in the idiom throughout the generation.

across various techniques for all encoder-decoder and decoder-only models supported by the Transformers library. The toolkit aims to democratize access to interpretability analyses of generative LMs with minimal setup, enabling reproducible evaluations. An example is provided in Figure 1. Thanks to its intuitive interface, users can easily integrate interpretability analyses into their text generation pipelines with just a few lines of code. Moreover, a command-line interface (CLI) and various utility methods to visualize, serialize, and reload attribution outcomes are provided to facilitate analysis at scale. Inseq is also highly flexible, including cutting-edge attribution methods with built-in post-processing features (Section 2), supporting customizable attribution targets and enabling the attribution of arbitrary sequences produced via forced decoding (Section 2.1).

In this paper, we summarize recent efforts in the development of the Inseq toolkit, focusing specifically on newly added usability features to support the attribution of large LMs (LLMs) (Section 2.2), and a new command to contrastively attribute context usage in LMs generations (Section 3). Finally, we present various applications of Inseq in recent research (Section 4).

2. The Inseq Toolkit

Inseq provides an easy-to-use interface to apply feature attribution methods, extending Captum [12] as attribution back-end to generative models from the Transformers library [10].

Table 1 (left) presents an updated list of supported attribution methods, categorized into three groups, *gradient-based*, *internals-based* and *perturbation-based*, depending on their underlying approach to importance quantification. Aside from popular model-agnostic methods, Inseq notably provides built-in support for attention weight attribution and a range of cutting-edge methods not supported in any other toolkit, such as Discretized Integrated Gradients [17], Sequential Integrated Gradients [18], Value Zeroing [22], and ReAGent [23], with many of those allowing for the importance attribution of custom intermediate model layers.

Among its notable features, Inseq offers flexible **source and target-side attribution** for

Method	Source	$f(l)$	Method	Source
(Input \times) Gradient	Simonyan et al.	yes	(Log) Probability	-
DeepLIFT	Shrikumar et al.	yes	Softmax Entropy	-
G GradientSHAP	Lundberg and Lee	no	Target Cross-entropy	-
Integrated Gradients	Sundararajan et al.	yes	Perplexity	-
Discretized IG	Sanyal and Ren	no	Contrastive logits Δ	Yin and Neubig
Sequential IG	Enguehard	no	Contrastive prob. Δ	-
I Attention Weights	Bahdanau et al.	yes	μ MC Dropout Prob.	Gal and Ghahramani
Occlusion (Blank-out)	Zeiler and Fergus	no	P-CXMI	Fernandes et al.
P LIME	Ribeiro et al.	no	KL divergence	-
Value Zeroing	Mohebbi et al.	yes	In-context PVI	Lu et al.
ReAGent	Zhao and Shan	no	Top-p tokens	-

Table 1

Gradient (**G**), internals (**I**) and perturbation-based (**P**) attribution methods and built-in step functions (**S**) in Inseq. $f(l)$: supports intermediate layers attribution. New methods are **bolded**.

encoder-decoder systems, alongside several Aggregator classes to aggregate attribution scores across various dimensions (e.g. at the token level), and AggregatorPipeline for chaining various aggregation steps (e.g. extract the weight of the i -th attention head at the n -th layer).

2.1. Customizing generation and attribution

At every generation step, in addition to computing attribution scores, Inseq can also use models’ information to compute functions of the output distributions or intermediate representations, which we collectively refer to as **step functions** (Table 1, **S**). For example, the resulting scores can provide additional insights into the generation process for uncertainty quantification or outlier detection. Inseq provides access to several built-in step functions and allows users to create and register custom ones. Step scores are computed alongside attribution and visualized in the same matrix of attribution scores (e.g. $p(y_t|y_{<t})$ in Figure 1).

Various attribution methods rely on model outputs to predict input importance, using functions of the model’s output logits or token probabilities [27]. Yin and Neubig [6] propose contrastive metrics to help disentangle how various factors contribute to the prediction. For example, the gradient $\nabla(p(\text{barking}) - p(\text{crying}))$ given the prompt “Can you stop the dog from ___” will highlight the role of the entity *dog* in selecting *barking*, disentangling the semantic component from grammatical correctness by providing a *crying* as grammatically valid choice. Figure 2 illustrates an example. Inseq users can leverage any built-in or custom-defined step function as an **attribution target**, enabling advanced use cases like contrastive comparisons.

The new version of Inseq supports customizable **word alignments**, i.e. indices aligning tokens in the original and contrastive generated texts, to support contrastive comparisons between texts of different lengths, including automatic alignments using the multilingual LaBSE encoder [28] to streamline their application.

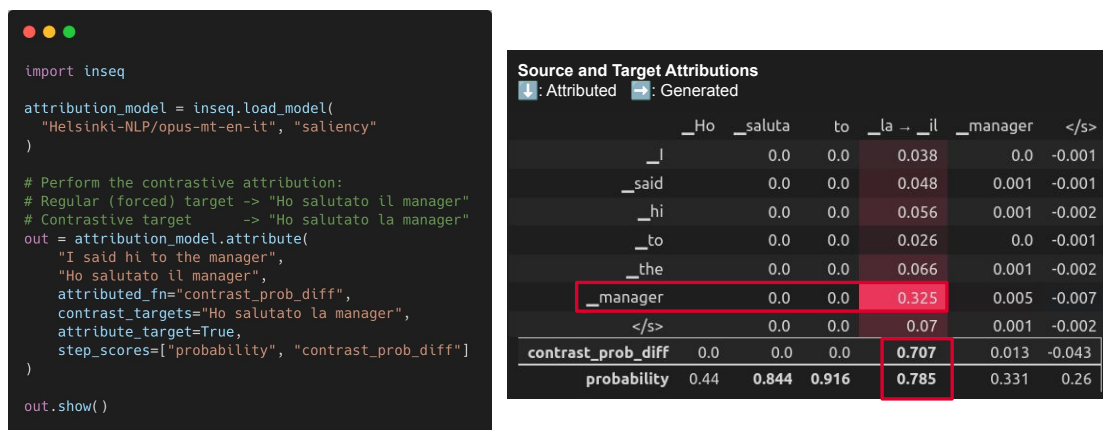


Figure 2: Left: Code using Inseq to compute contrastive attributions for an English-to-Italian machine translation model using raw gradient attribution. **Right:** Source-to-target attributions aggregated at token-level, indicating the importance of the stereotypical noun “manager” to generate the Italian masculine pronoun “il” (original) over the feminine “la” (contrastive case).

2.2. Usability Features

Inseq supports batching to simplify analysis at scale and customizable start/end positions to accelerate the attribution process for studies on localized phenomena (e.g., pronoun coreference). Moreover, it offers a CLI to attribute single examples or entire Datasets from the command line, storing resulting outputs and visualizations. Attributions can be saved in JSON format with metadata to identify their provenance, allowing for easy reloading and visualization.

Quantization and distributed attribution All models allowing for **quantization** using bitsandbytes [29] can be loaded in 4-bit or 8-bit precision directly from Transformers, and their attributions can be computed normally using Inseq at a fraction of the cost. Similarly, Inseq is compatible with the Petals library [30], supporting gradient-based attribution across language models whose computation is distributed across several machines. This can alleviate the need for high-end GPUs to run LLMs, enabling the distributed computation of attribution scores.¹

3. Case study: Attributing Context Influence using PECoRe

The PECoRe framework [31] was proposed to identify and attribute context usage in language models, and further adapted by Qi et al. [32] to produce model internals-based citations for LLM generations. First, contrastive functions such as KL divergence select generated tokens sensitive to context ablation. Then, contrastive feature attribution is used to identify context tokens driving the contextual prediction. Inseq provides an ad-hoc CLI command (`attribute-context`) for PECoRe usage, supporting all contrastive step functions and attribution methods. Figure 3 provides an example output in a GUI built on top of the Inseq API.² In the example, an LLM³ is

¹A tutorial for distributed attribution is available here: <https://inseq.org/en/latest/examples/petals.html>

²The presented demo is available here: <https://huggingface.co/spaces/gsarti/pecore>

³We used StableLM 2 Zephyr 1.6B: https://huggingface.co/stabilityai/stablelm-2-zephyr-1_6b

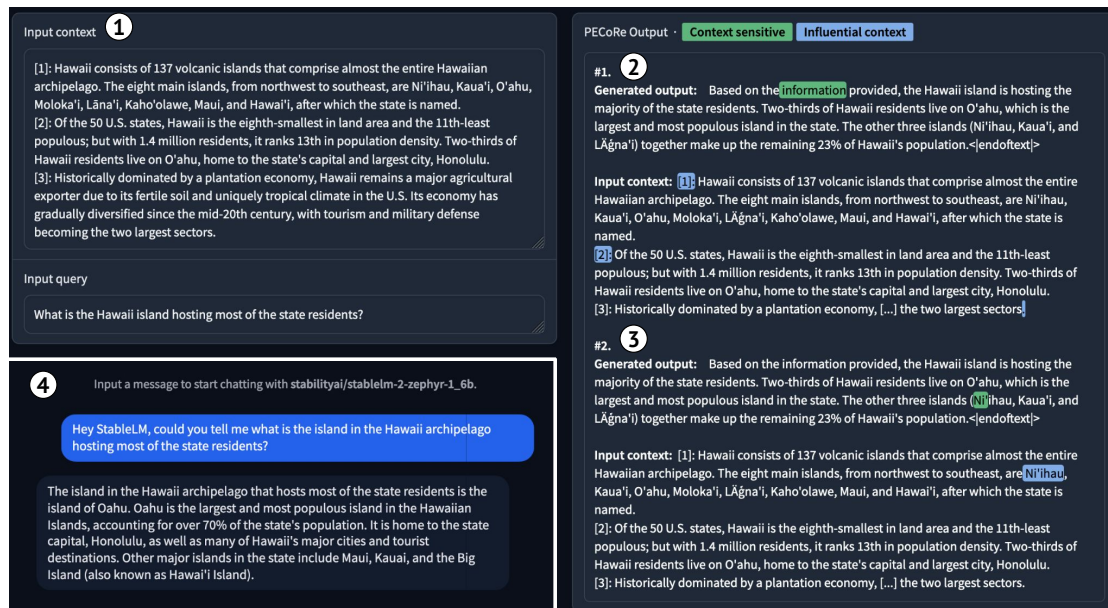


Figure 3: Context attribution for retrieval-augmented long-form QA using the Inseq-powered demo.

prompted with contexts retrieved from Wikipedia to provide a long-form answer to a query (1). When referring to context information (2), PECoRe shows that the indices of the two documents containing relevant information are salient. On the other hand, the names of other Hawaiian islands are important when the model produces an additional remark on their population (3). We observe that the context is not salient for answering the question, suggesting the model might have memorized the answer. We test this by prompting the model in a closed-book setting, finding that the model can indeed respond correctly without context (4).

4. Related Work using Inseq

Since its first release, Inseq was adopted to conduct several feature attribution analyses of generative LMs. In the conversational domain, its Integrated Gradients implementation was used to study longitudinal dialogues with conversational models for Italian [33]. Inseq was also used to measure agreement between attribution scores and a new metric of LLMs' factual reliability [34], and to analyze the context repetition in dialogues [35]. In machine translation, Inseq attribution methods were used to select salient in-context examples with the aim to mitigate gender bias in translated sentences [36] and to evaluate the usage of source and target-side information in character-level machine translation systems across several languages [37].

Inseq was integrated into several tools and methods, including the LLMCheckup interface [38], using Inseq for producing attributions for fact-checking and conversational question answering (QA), and the PECoRe framework [31] for detecting and attributing context usage in language models. Finally, Inseq methods were also used as baselines to compare proposed new feature attribution approaches [23], and to probe the contextual influence in affixal negation [39].

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