# Non-Exchangeable Conformal Language Generation with Nearest Neighbors

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#### Abstract

Quantifying uncertainty in automatically generated text is important for letting humans check potential hallucinations and making systems more reliable. Conformal prediction is an attractive framework to provide predictions imbued with statistical guarantees, however, its application to text generation is challenging 007 since any i.i.d. assumptions are not realistic. In this paper, we bridge this gap by leveraging recent results on non-exchangeable conformal prediction, which still ensures bounds on cov-011 erage. The result is a novel extension of the 013 conformal prediction framework to generation based on nearest neighbors. Our method can be used post-hoc for an arbitrary model without extra training and supplies token-level, calibrated prediction sets equipped with statistical guaran-017 tees. Experiments in machine translation and language modeling show encouraging results 019 in word coverage and generation quality.

#### 1 Introduction

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Natural language generation (NLG) is a multifaceted field spanning applications such as machine translation (MT), language modeling (LM), summarization, question answering and dialogue generation. Owing to the recent success of large language models (LLMs) such as GPT-4 (OpenAI, 2023), BLOOM (Scao et al., 2022) or LLaMA (Touvron et al., 2023), natural language modeling with stochastic decoding (sampling) is increasingly used as an interface with end users. While sampling allows for more fluent and varied text, few methods exist to evaluate the reliability of generated text and adequacy of the underlying sampling method. This is particularly relevant for generation scenarios where pre-trained models are applied to new data with potentially different distribution to the training data, increasing the risk of generating erroneous, misleading, and potentially harmful text (Ji et al., 2023; Guerreiro



Figure 1: Schematic representation of our approach. A decoder hidden representation  $\mathbf{z}_t$  is used during inference to retrieve the nearest neighbors and their nonconformity scores  $s_k$ . Their relevance is determined by using their distance to compute weights  $w_k$ , resulting in the quantile  $\hat{q}$  that forms conformal prediction sets.

et al., 2023; Pan et al., 2023; Alkaissi and McFarlane, 2023; Azamfirei et al., 2023).

Conformal prediction (Vovk et al., 2005; Papadopoulos et al., 2002; Angelopoulos and Bates, 2021) has recently gained popularity by providing calibrated prediction sets that are imbued with statistical guarantees about containing the correct solution. Nevertheless, applying conformal prediction to NLG is not trivial and comes with a major obstacle: The conditional generation process breaks the independence and identical distribution (i.i.d.) assumption underlying conformal prediction techniques. We tackle this problem by drawing inspiration from recent advances in nearest neighbor language modeling (Khandelwal et al., 2020b; He et al., 2021a; Xu et al., 2023) and machine translation (Khandelwal et al., 2020a; Zheng et al., 2021; Meng et al., 2022; Martins et al., 2022). This way, we are able to dynamically generate calibration sets during inference that are able to maintain statistical guarantees. We schematically illustrate our method in Figure 1.

**Contributions.** We present a general-purpose extension of the conformal framework to NLG by

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tackling the problems above. Our contributions are 067 as follows: (1) To the best of our knowledge, we are the first to present a novel technique based on non-exchangeable conformal prediction and to apply it to language generation to produce calibrated prediction sets. (2) We validate the effectiveness of the method in a Language Modeling and Machine Translation context, evaluating the coverage of the calibrated prediction sets and showing that 075 our method is on par with or even outperforms other 076 sampling-based techniques in terms of generation 077 quality, all while maintaining tighter prediction sets and better coverage. (3) We finally demonstrate that 079 these properties are also maintained under distributional shift induced by corrupting the model's latent representations. (4) We publish all the code for this project in an open-source repository.<sup>1</sup>

#### 2 Related Work

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**Conformal Prediction.** Conformal prediction is a line of work that has recently regained interest in machine learning by producing prediction sets with certain statistical guarantees about containing the correct prediction (Vovk et al., 2005; Papadopoulos et al., 2002; Angelopoulos and Bates, 2021). As the size of prediction sets is calibrated to fulfill these guarantees, one can also see the size of the prediction set itself as a proxy of the uncertainty of a model-the larger the set, the more possible predictions have to be included in order to maintain the coverage guarantee. Conformal prediction has already found diverse applications in NLP for classification (Maltoudoglou et al., 2020; Fisch et al., 2021; Schuster et al., 2021; Fisch et al., 2022; Choubey et al., 2022; Kumar et al., 2023) and sequence labeling problems (Dey et al., 2021), as well as quality estimation (Giovannotti, 2023; Zerva and Martins, 2023). Unfortunately, generation problems are challenging due to their sequential nature and constant breaking of the i.i.d. assumption, so existing works operate on the sequencelevel instead (Quach et al., 2023; Ren et al., 2023; Deutschmann et al., 2023). Conformal procedures for time-series (Xu and Xie, 2021; Lin et al., 2022b; Oliveira et al., 2022; Zaffran et al., 2022) and general non-i.i.d. data (Tibshirani et al., 2019; Barber et al., 2023; Guan, 2023; Farinhas et al., 2023) have been proposed in the literature. The most related work to ours is given by Ravfogel et al. (2023), who apply the standard conformal prediction setup to

NLG, arguing that Markov chains are a type of  $\beta$ mixing processes, for which Oliveira et al. (2022) showed coverage to degrade by an only negligible amount. However, Ravfogel et al. do not investigate this claim empirically, and furthermore do not find any benefits when generating sequences. In another related work, Quach et al. (2023) propose an approach that is specifically tailored toward language modeling. However, their prediction sets contain entire sequences instead of single tokens. In contrast, our token-level prediction sets are useful for constraining the options during generation and their widths can represent model uncertainty. 116

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Uncertainty in NLP. Modeling uncertainty in NLP has already been studied in classification (Van Landeghem et al., 2022; Ulmer et al., 2022a; Holm et al., 2022) and regression settings (Beck et al., 2016; Glushkova et al., 2021; Zerva et al., 2022). However, NLG proves more challenging due to it non-i.i.d. and combinatorial nature. Some works have proposed Bayesian Deep Learning methods for NLG: Xiao et al. (2020) use Monte Carlo Dropout (Gal and Ghahramani, 2016) to produce multiple generations for the same input and measure their pair-wise BLEU scores. Malinin and Gales (2021) define extensions of mutual information for structured prediction. Other existing approaches try to account for the paraphrastic nature of language by modeling the entropy over meaning classes (Kuhn et al., 2023), investigate the use of linguistic markers to indicate uncertainty (Zhou et al., 2023) or ask the model directly for its confidence (Lin et al., 2022a; Kadavath et al., 2022). Baan et al. (2023) provide an extensive overview of the theory and current state of the field.

#### **3** Background

**Conformal Prediction.** Conformal prediction is an attractive method for uncertainty quantification due to its statistical coverage guarantees (Vovk et al., 2005; Papadopoulos et al., 2002; Angelopoulos and Bates, 2021). Given some predictor, a heldout calibration set  $\{(\mathbf{x}_i, y_i)\}_{i=1}^N$ , and a pre-defined miscoverage level  $\alpha$  (e.g., 0.1), the calibration set is used to obtain *prediction sets*  $C(\mathbf{x}^*)$  for a new test point  $\mathbf{x}^*$  satisfying

$$p(y^* \in \mathcal{C}(\mathbf{x}^*)) \ge 1 - \alpha,$$
 (1)

that is, the probability of the prediction set  $C(\mathbf{x}^*)$ containing the correct label  $y^*$  is at least  $1-\alpha$ . This

<sup>&</sup>lt;sup>1</sup>Made available upon acceptance.

is achieved by the following recipe: Firstly, one has 164 to define a non-conformity score, that provides an 165 estimate of the distance of the test point to the rest 166 of the data, i.e., a proxy for the uncertainty over the 167 test point predictions. In this context, the score can 168 be as simple as  $s_i = 1 - p_{\theta}(y | \mathbf{x})$ , i.e. one minus the 169 softmax probability of the true class, which will be 170 higher when the model is wrong or less confident. 171 Next, we define  $\hat{q}$  as the  $|(N+1)(1-\alpha)/N|$ -th quantile of the non-conformity scores. Then, when 173 we make a new prediction for a test point  $x^*$ , we 174 can create prediction sets defined as 175

$$\mathcal{C}(\mathbf{x}^*) = \left\{ y \mid p_{\boldsymbol{\theta}}(y \mid \mathbf{x}^*) \ge 1 - \hat{q} \right\}, \qquad (2)$$

which is guaranteed to fulfil the coverage requirement in Equation (1) for i.i.d. data (Vovk et al., 2005; Angelopoulos and Bates, 2021).

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Non-exchangeable Conformal Prediction. Bar-180 ber et al. (2023) address a major shortcoming in 181 the method above: When a test point and the calibration data are not i.i.d.,<sup>2</sup> the distributional drift causes any previously found  $\hat{q}$  to be miscalibrated, and thus the intended coverage can no longer be 185 guaranteed. However, we can still perform conformal prediction by assigning a weight  $w_i \in$ [0, 1] to every calibration data point, reflecting its 188 relevance-i.e. assigning lower weights to points far away from the test distribution. Then, by nor-190 malizing the weights with  $\tilde{w}_i = w_i/(1 + \sum_{i=1}^N w_i)$ , 191 we define the quantile as 192

$$\hat{q} = \inf \left\{ q \left| \sum_{i=1}^{N} \tilde{w}_i \mathbf{1} \{ s_i \le q \} \ge 1 - \alpha \right\}, \quad (3)$$

with  $1{\cdot}$  denoting the indicator function. The construction of the prediction sets then follows the same steps as before. Most notably, the coverage guarantee in Equation (1) now changes to

$$p\left(y^* \in \mathcal{C}(\mathbf{x}^*)\right) \ge 1 - \alpha - \sum_{i=1}^N \tilde{w}_i \varepsilon_i, \quad (4)$$

with an extra term including the *total variation* distance between the distribution of a calibration and a test point,  $\varepsilon_i = d_{\text{TV}}((\mathbf{x}_i, y_i), (\mathbf{x}^*, y^*))$ .<sup>3</sup> Unfortunately, this term is hard to estimate or bound, nevertheless, the selection of appropriate weights that can capture the relevance of calibration points to the test set should moderate both the impact of the distant data points on the estimation of the prediction set and the impact of  $d_{\rm TV}$  on the coverage bound. In other words, for large  $d_{\rm TV}$  values we expect to have smaller weights, that allow us to achieve coverage close to the desired values. We show in our experiments that the loss of coverage when using nearest neighbor weights is limited and revisit the practical implications in Section 5. 203

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## 3.1 Method: Non-exchangeable Conformal Prediction through Nearest Neighbors

We now present a novel method to apply conformal prediction in NLG by synthesizing the nonexchangeable approach of Barber et al. (2023) with k-NN search-augmented neural models (Khandelwal et al., 2020a,b). The related approach by Ravfogel et al. (2023) calibrates prediction sets within bins of similar entropies using the nonexchangeable procedure described in Section 3. However, this implies that we would use semantically unrelated (sub-)sequences to calibrate the model—in fact, we show experimentally that this approach obtains generally trivial coverage by producing extremely wide prediction sets. Instead, we propose to perform a *dynamic* calibration step during model inference, only considering the most relevant data points from the calibration set. We do this in the following way: Given a dataset  $\{(\mathbf{x}^{(i)}, y^{(i)})\}$  of sequences  $\mathbf{x}^{(i)} = (\mathbf{x}_1^{(i)}, \dots, \mathbf{x}_S^{(i)})$  and corresponding references consisting of gold tokens  $y^{(i)} = (y_1^{(i)}, \dots, y_T^{(i)})$ , we extract the model's decoder activations  $\mathbf{z}_t^{(i)} \in \mathbb{R}^d$  and conformity scores  $s_t^{(i)}$ .<sup>4</sup> We save those in a datastore allowing for fast and efficient nearest neighbor search using FAISS (Johnson et al., 2019). In the inference phase, during every decoding step, we then use the decoder hidden state  $\mathbf{z}_t^*$  to query the data store for the K nearest neighbors and their conformity scores and record their distances. We use the squared  $l_2$  distance to compute the weight  $w_k$  for a neighbor as

$$w_k = \exp\left(-\left|\left|\mathbf{z}_t^* - \mathbf{z}_k\right|\right|_2^2 / \tau\right), \qquad (5)$$

where  $\tau$  corresponds to a temperature hyperparameter. Overall, this formulation is equivalent to a

<sup>&</sup>lt;sup>2</sup>In fact, the coverage guarantee applies to the case where the data is *exchangeable*, a weaker requirement than i.i.d. Specifically, a series of random variables is exchangeable if their joint distribution is unaffected by a change of their order.

<sup>&</sup>lt;sup>3</sup>In this expression,  $(\mathbf{x}_i, y_i)$  and  $(\mathbf{x}^*, y^*)$  denote random variables and the total variation distance is between the two underlying distributions. See Barber et al. (2023) for details.

<sup>&</sup>lt;sup>4</sup>In this phase, we do not let the model generate freely, but feed it the gold prefix during the decoding process to make sure that conformity scores can be computed correctly.

radial basis function kernel with scale parameter  $\tau$ . 249 Finally, we use the weights to compute the quantile  $\hat{q}$  as in Equation (3). The entire algorithm is given in Appendix A.4.

Adaptive Prediction Sets. The efficacy of conformal prediction hinges on the choice of non-254 conformity score, with the simple non-conformity 255 score  $s_i = 1 - p_{\theta}(y_t | \mathbf{x}, y_{\leq t})$  known to undercover hard and overcover easy subpopulations of the data. Due to the diverse nature of language, we therefore opt for *adaptive prediction sets* (Angelopoulos et al., 2021a; Romano et al., 2020). Adaptive prediction sets redefine the non-conformity score as the cumulative probability over classes necessary to reach the correct class. More formally, let  $\pi$  be a permutation function mapping all possible output tokens  $\{1, \ldots, C\}$  to the indices of a permuted version of the set, for which tokens are sorted by 266 their probability under the model, descendingly. We define the non-conformity score as

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$$s_{i} = \sum_{j=1}^{\pi(y_{t})} p_{\theta}(\pi^{-1}(j) | \mathbf{x}, y_{< t}).$$
 (6)

Since we only include the cumulative mass up until the gold label, the summation stops at  $\pi(y)$ . The prediction sets are then defined as

$$\mathcal{C}(\mathbf{x}^*, y_{< t}^*) = \left\{ \pi^{-1}(1), \dots, \pi^{-1}(\hat{c}) \right\}, \quad (7)$$

with  $\hat{c} = \sup\{c' \mid \sum_{j=1}^{c'} p_{\theta}(\pi^{-1}(j) \mid \mathbf{x}^{*}, y_{< t}^{*}) <$  $\hat{q}$  + 1. Intuitively, this means that we included all classes whose cumulative probability (after sorting descendingly) does not surpass  $\hat{q}$ , adding one extra class to avoid empty sets. Compared to the simple conformity score, this produces wider predictions sets for hard inputs, encompassing more potentially plausible continuations in a language context.

#### **Experiments** 4

In the following sections, we conduct experiments in both language modeling and machine translation. For machine translation we opt for the 400 million and 1.2 billion parameter versions of the M2M100 model (Fan et al., 2021) on the WMT-2022 shared task datasets for German to English and Japanese to English (Kocmi et al., 2022). For Language Modelling, we use the 350 million and 1.3 billion parameter versions of the OPT model (Zhang et al., 2022) and replicate the setup by Ravfogel et al. (2023): We calibrate our model on

10000 sentences from a 2022 English Wikipedia dump (Foundation, 2022) and test coverage and generation on 1000 sentences from OpenWebText (Gokaslan et al., 2019).<sup>5</sup> All models are used in a zero-shot setup without extra training or finetuning. For the datastore, we use the implementation by FAISS library (Johnson et al., 2019), computing 2048 clusters in total and probing 32 clusters per query. We also summarize the environmental impact of our experiments in Appendix A.5.

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#### 4.1 Evaluating Coverage

First of all, we demonstrate that the retrieved information from the data store enables us to successfully apply the proposed method. While it is not possible to measure coverage in a free generation setting (see next section), we can assess whether the correct class is contained in the prediction set if we feed the actual reference tokens into the decoder and check whether we include the true continuation.<sup>6</sup> For our MT task, this is reminiscent of an interactive translation prediction setup (Knowles and Koehn, 2016; Peris et al., 2017; Knowles et al., 2019), where we would like to suggest possible continuations to a translator, suggesting the next word from a set of words that (a) contains plausible options and (b) is limited in size, in order to restrict the complexity for the end user. Before we run our experiments, we need to determine  $\tau$ , which we tune on the calibration set using a stochastic hillclimbing procedure described in Appendix A.1. We compare our non-exchangeable conformal nucleus sampling (Non-Ex. CS) with nucleus sampling (Holtzman et al., 2020) and conformal nucleus sampling (Conf. Sampl.; Ravfogel et al., 2023), using 10 entropy bins and corresponding  $\hat{q}$  values.

**Evaluation.** We evaluate by measuring the total coverage using different distance metrics, namely, squared  $l_2$  distance, normalized inner product, and cosine similarity (see Tables 1 and 2),<sup>7</sup> as well as binning predictions by set size and then measuring the per-bin coverage in Figure 2 (more results given in Appendix A.2). We also summarize the plots in

<sup>&</sup>lt;sup>5</sup>Data obtained through the Hugging Face datasets package (Lhoest et al., 2021): https://huggingface. co/datasets/wikipedia and https://huggingface.co/ datasets/stas/openwebtext-10k.

<sup>&</sup>lt;sup>6</sup>We emphasize that access to gold tokens is not required by our method and only done here to measure the actual coverage.

For inner product and cosine similarity, we follow the same form as Equation (5), omitting the minus. We normalize the inner product by the square root of the latent dimension.

Table 1: Coverage results for the de  $\rightarrow$  en and ja  $\rightarrow$  en MT tasks. We report the best found temperature  $\tau$  while keeping the confidence level  $\alpha$  and number of neighbors k = 100 fixed. We also show the coverage percentage along with the avg. prediction set size as a proportion of the entire vocabulary (Ø WIDTH) as well as ECG and SSC. Tested distance metrics are inner product (IP), (squared)  $l_2$  distance, and cosine similarity (cos).

			de  ightarrow en				$ja \rightarrow en$					
	Method	Dist.	τ	% COVERAGE	$\varnothing \ Width \downarrow$	$\mathbf{SCC}\uparrow$	$\text{Ecg}\downarrow$	τ	% COVERAGE	$\varnothing  Width \downarrow$	$\mathbf{SCC}\uparrow$	Ecg↓
_	Nucleus Sampling	-	-	0.9207	0.48	0.25	0.00	-	0.9261	0.54	0.41	0.02
400M	Conf. Sampling	-	-	0.9951	0.94	0.33	0.03	-	0.9950	0.96	0.14	0.00
M2M100(	Non-Ex. CS	IP	3.93	0.8251	0.16	0.63	0.26	11.90	0.8815	0.24	0.67	0.03
		$l_2$	512.14	0.8334	0.17	0.60	0.06	419.91	0.8468	0.18	0.61	0.05
		cos	2.54	0.8371	0.17	0.63	0.06	3.53	0.8540	0.17	0.62	0.04
_	Nucleus Sampling	-	-	0.8339	0.38	0.00	0.08	-	0.7962	0.42	0.03	0.10
(1.2B	Conf. Sampling	-	-	0.9993	0.99	0.34	0.00	-	0.9998	0.99	0.60	0.00
1100	Non-Ex. CS	IP	15.79	0.8861	0.25	0.71	0.03	10.45	0.9129	0.38	0.72	0.00
M2N		$l_2$	1123.45	0.8874	0.25	0.72	0.03	605.97	0.8896	0.30	0.76	0.01
		cos	3.21	0.8858	0.25	0.72	0.03	1.48	0.8897	0.30	0.75	0.01
Coverade	0.8 0.6 0.4 0.2 0 20000 400	000 600 Se	000 80000 t Size	100000 120000	2000 <u>vi</u> 1500 <u>o</u> 1000 go 500 0		0.8 b b b b c 0.6 0.4 0.2 0	20000	40000 6000 80 Set Size	000 100000 120	- 150 - 100 - 500 - 0	00 000 Number of Points
Coverade	(a) Nucle		npling or	n de $\rightarrow$ en.	8000 0000 6000 0000 Number of Points		(b) C	Conform	al Nucleus Sa	mpling on d	$e \rightarrow en$ 25 20 15 10 50	Number of Points

(c) Non-Ex. Conformal Sampling on de  $\rightarrow$  en.



Figure 2: Conditional coverage for the M2M100 on de  $\rightarrow$  en with the small 418M model (Figures 2a to 2c) and using the bigger 1.2B model (Figure 2d). We aggregate predictions by set size using 75 equally-spaced bins in total. The blue curve shows the conditional coverage per bin, whereas red bars show the number of binned predictions.

Figure 2 via the *Expected Coverage Gap*  $(ECG)^8$ that we define as

$$ECG = \sum_{b=1}^{B} \frac{|\mathcal{B}_b|}{N} \max\left(1 - \alpha - Coverage(\mathcal{B}_b), 0\right),$$
(8)

where  $\mathcal{B}_b$  denotes a single bin and N the total number of considered predictions in the dataset.<sup>9</sup> In our experiments, we use 75 bins in total. The same bins are used to also evaluate the Size-Stratified Coverage metric (SSC) proposed by Angelopoulos et al. (2021b) to assess the balance of coverage across set sizes, with a well-calibrated method resulting in a SCC close to the desired coverage  $1 - \alpha$ :

$$SCC = \min_{b \in \{1, \dots, B\}} Coverage(\mathcal{B}_b)$$
(9)

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<sup>&</sup>lt;sup>8</sup>This measure is inspired by the expected calibration error (Guo et al., 2017), but measuring the coverage against a constant target value  $1 - \alpha$ . Since Conformal Prediction provides a lower bound, overcoverage is not penalized.

<sup>&</sup>lt;sup>9</sup>Since conformal prediction produces a *lower* bound on the coverage, we do not include overcoverage in Equation (8).

Table 2: Coverage results for the LM task. We report the best found temperature  $\tau$  while keeping the confidence level  $\alpha$  and number of neighbors k = 100 fixed. We also show the coverage percentage along with the avg. prediction set size as a proportion of the entire vocabulary ( $\emptyset$  WIDTH) as well as the ECG and SSC metrics. Tested distance metrics are inner product (IP), (squared)  $l_2$  distance and cos. similarity (cos).

				OPENWEBTEXT						
	Method	Dist.	τ	% Cov.	$\varnothing  Width \downarrow$	$\mathbf{SCC}\uparrow$	Ecg↓			
	Nucl. Sampl.	-	-	0.8913	0.05	0.71	0.01			
(W)	Conf. Sampl.	-	-	0.9913	0.90	0.91	0.00			
T <sub>(35(</sub>	Non-Ex. CS	IP	4.99	0.9352	0.19	0.80	0.0			
OP		$l_2$	$0.31\times 10^4$	0.9425	0.17	0.80	0.0			
		cos	4.98	0.9370	0.15	0.83	0.0			
	Nucl. Sampl.	-	-	0.8952	0.05	0.00	0.01			
B	Conf. Sampl.	-	-	0.9905	0.88	0.95	0.0			
T <sub>(1.3</sub>	Non-Ex. CS	IP	0.48	0.9689	0.59	0.84	0.0			
OF		$l_2$	$1.55\times 10^4$	0.9539	0.20	0.83	0.0			
		cos	0.11	0.9512	0.20	0.875	0.0			

We present some additional experiments where we assess the impact of key hyperparameters in Appendix A.3.

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**Results.** We found our method to miss the desired coverage of 90% for MT by 8% or less. Beyond the reported values, we were not able to further increase coverage by varying the temperature parameter without avoiding trivial coverage (i.e., defaulting to very large set sizes), which is likely due to the impossible-to-estimate coverage in Equation (4). Most notably, our method was able to achieve better SCC scores while maintaining considerably smaller prediction sets than the baselines on average. The reason for this is illustrated in Figure 2: while standard nucleus sampling produces some prediction sets that are small, the total coverage seems to mostly be achieved by creating prediction sets between 60k-80k tokens. The behavior of conformal nucleus sampling by Ravfogel et al. (2023) is even more extreme in this regard, while our method focuses on producing smaller prediction sets, with the frequency of larger set sizes decreasing gracefully. In Figure 2d, we can see that the larger M2M100 models also tend to produce larger prediction sets, but still noticeably smaller than the baselines. Importantly, for both M2M100 models, even very small prediction sets (size  $\leq 1000$ ) achieve non-trivial coverage, unlike the baseline methods. For LM, we always found the model to slightly overcover. This does not contradict the desired lower bound on the coverage in

Equation (4) and suggests a more negligible distributional drift. While nucleus sampling produces the smallest average prediction sets, we can see that based on the SCC values some strata remain undercovered. Instead, our method is able to strike a balance between stratified coverage and prediction set size. With respect to distance measures, we find that the difference between them is minimal, indicating that the quality largely depends on the retrieved local neighborhood of the decoder encoding and that finding the right temperature can help to tune the models to approximate the desired coverage. Now we would like to find out whether this neighborhood retrieval mechanism can prove to be robust under distributional shift as well.

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#### 4.2 Coverage Under Shift

To demonstrate how the retrieval of nearest neighbors can help to maintain coverage under distributional shift, we add Gaussian noise of increasing variance-and therefore intensity-to the last decoder hidden embeddings (for MT) and the input embeddings (LM). This way, we are able to simulate distributional drift while still keeping the original sequence of input tokens intact, allowing us to measure the actual coverage. We show the achieved coverage along with the average set size (as a percentage of the total vocabulary) and the average quantile  $\hat{q}$  in Figure 3. We can see that the conformal sampling method deteriorates into returning the full vocabulary as a prediction set. Thus it behaves similarly to simple sampling as indicated by the  $\hat{q}$  values being close to 1. Nucleus sampling provides smaller prediction sets compared to conformal sampling, but they seem invariant to noise. As such, the method is not robust to noise injection in the open text generation task, and the obtained coverage deteriorates with noise variance  $\geq 0.025$ . Instead, the use of nearest neighbors allows for the estimation of prediction sets that are small but amenable to increase, such that the obtained coverage remains close to the desired one. We can specifically observe that the prediction set size increases considerably to mitigate the injected noise in the open-text generation case.

**Neighbor Retrieval.** We further analyze how the retrieval enables this flexibility by relating it to the entropy of the output distribution of the 400M parameters M2M100 on German to English. Intuitively, the baseline methods, faced by high-entropy output distributions, need to produce wide predic-



Figure 3: Coverage, average set size and  $\hat{q}$  based on the noise level on the de  $\rightarrow$  en MT task (top) and open text generation task (bottom). Error bars show one standard deviation.

Table 3: Average entropy of 400M M2M100 model on de  $\rightarrow$  en per noise level as well as the Spearman's  $\rho$  correlation coefficients between the predictive entropy and the prediction set size of the different methods. All results are significant with p < 0.0001.

	NOISE LEVEL							
	None	0.025	0.05	0.075	0.1			
Ø Entropy	8.46	8.71	9.20	9.71	10.08			
Nucl. Sampl. $(\rho)$	0.87	0.86	0.84	0.82	0.81			
Conf. Sampl. $(\rho)$	0.60	0.60	0.60	0.57	0.55			
Non-Ex. CS $(\rho)$	-0.14	-0.18	-0.27	-0.37	-0.45			

tion sets in order to maintain coverage. In fact, we report such results by correlating entropy levels and prediction set sizes using Spearman's  $\rho$  in Table 3, showing strong positive correlations. Our method in contrast shows consistently an *anti*correlation between these two quantities, enabled by decoupling the creation of prediction sets from statistics of the output distribution to instead considering the non-conformity scores of similar subsequences. The fact that the prediction set size is not just dependent on the entropy of the predictions while maintaining coverage demonstrates the value of the nearest neighbors: In this way, model uncertainty becomes more flexible and is corroborated by evidence gained from similar inputs.

#### 4.3 Generation Quality

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Crucially, our method should not degrade and potentially even improve generation quality. Thus, we evaluate generation quality for the same tasks without supplying the gold prefix. For language modeling, we follow Ravfogel et al. (2023) and use the first 35 tokens from the original sentence as input. We compare against a set of generation strategies including top-k sampling (Fan et al., 2018; Holtzman et al., 2018; Radford et al., 2019), nucleus sampling and conformal nucleus sampling. We also test a variant of our method using constant weights  $w_k = 1$  for retrieved neighbors (Const. Weight CS) to assess the impact of the weighted neighbor retrieval procedure. We further compare with beam search (Medress et al., 1977; Graves, 2012) with a softmax temperature of 0.1, and greedy decoding. Evaluation is performed using BLEU (Papineni et al., 2002), COMET-22 (Rei et al., 2020, 2022) and chrF (Popović, 2017) for MT as well as MAUVE (Pillutla et al., 2021) and BERTscore (Zhang et al., 2020) for text generation.<sup>10</sup>

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**Results.** We show the results for the different methods in Table 4. We see that beam search outperforms all sampling methods for MT. This corroborates previous work by Shaham and Levy (2022) who argue that (nucleus) sampling methods, by pruning only the bottom percentile of the token distribution, introduce some degree of randomness that is beneficial for open text generation but may be less optimal for conditional language generation, where the desired output is constrained and exact matching generations are preferred (which is the case for MT). Among sampling methods, we find nucleus sampling and conformal sampling to perform similarly (being in agreement with the findings of Ravfogel et al., 2023) but are

<sup>&</sup>lt;sup>10</sup>All metrics except for COMET were used through Hugging Face evaluate. MAUVE uses gpt2 as a featurizer.

			$de \to en$			$ja \to en$				Ope	NWEBTEXT
	Method	BLEU $\uparrow$	$Comet\uparrow$	ChrF↑	Bleu↑	$Comet\uparrow$	$ChrF\uparrow$		Method	MAUVE ↑	BERTSCORE $F_1 \uparrow$
	Beam search	28.53	0.88	55.58	11.37	0.63	37.74		Beam search	0.12	0.79
-	Greedy	27.81	0.9	54.9	10.73	0.58	36.5	-	Greedy	0.02	0.79
)(400m	Nucleus Sampling	$27.63{\scriptstyle~\pm 0.03}$	$0.89{\scriptstyle~\pm 0.01}$	$54.80 \pm 0.07$	$10.61{\scriptstyle~\pm 0.15}$	$0.59 \pm 0.01$	$36.52 \pm 0.19$	(350M	Nucleus Sampling	$0.91 \pm 0.02$	$0.80 \pm 0.00$
M100	Top- $k$ Sampling	$27.63{\scriptstyle~\pm 0.03}$	$0.89{\scriptstyle~\pm 0.01}$	$54.79 \pm 0.07$	$10.61{\scriptstyle~\pm 0.15}$	$0.59{\scriptstyle~\pm 0.01}$	$36.52{\scriptstyle~\pm 0.19}$	ЦО	Top-k Sampling	$0.90{\scriptstyle~\pm 0.03}$	$0.80 \pm 0.00$
M2N	Conf. Sampling	$27.63{\scriptstyle~\pm 0.03}$	$0.89{\scriptstyle~\pm 0.01}$	$54.80{\scriptstyle~\pm 0.07}$	$10.61{\scriptstyle~\pm 0.15}$	$0.59{\scriptstyle~\pm 0.01}$	$36.52{\scriptstyle~\pm 0.19}$		Conf. Sampling	$0.91{\scriptstyle~\pm 0.02}$	$0.80 \pm 0.00$
	Const. Weight CS*	$27.63{\scriptstyle~\pm 0.03}$	$0.89{\scriptstyle~\pm 0.01}$	$54.80{\scriptstyle~\pm 0.07}$	$10.61{\scriptstyle~\pm 0.15}$	$0.59{\scriptstyle~\pm 0.01}$	$36.52{\scriptstyle~\pm 0.19}$		Const. Weight CS*	$0.91{\scriptstyle~\pm 0.02}$	$0.80 \pm 0.00$
	Non-Ex. CS*	$27.65{\scriptstyle~\pm 0.10}$	$0.90{\scriptstyle~\pm 0.01}$	$54.82 \pm 0.14$	$\underline{10.74} \pm 0.11$	$0.59{\scriptstyle~\pm 0.01}$	$36.61 \pm 0.08$		Non-Ex. CS*	$0.92 \pm 0.01$	$0.80 \pm 0.00$
	Beam search	30.89	0.9	56.8	13.76	0.63	40.43		Beam search	0.17	0.80
~	Greedy	29.52	0.9	55.67	12.94	0.6	39.91	-	Greedy	0.05	0.79
0 <sub>(1.2B</sub>	Nucleus Sampling	$29.37{\scriptstyle~\pm0.12}$	$0.90{\scriptstyle~\pm 0.00}$	$55.55 \pm 0.11$	$10.61{\scriptstyle~\pm 0.15}$	$0.59 \pm 0.01$	$36.52 \pm 0.19$	C(1.3B	Nucleus Sampling	$0.91 \pm 0.02$	$0.80 \pm 0.00$
M10	Top-k Sampling	$29.53{\scriptstyle~\pm 0.00}$	$0.90{\scriptstyle~\pm 0.00}$	$55.67{\scriptstyle~\pm0.00}$	$12.91{\scriptstyle~\pm 0.08}$	$0.60{\scriptstyle~\pm 0.01}$	$39.95 \pm 0.00$	LdO	Top-k Sampling	$0.93{\scriptstyle~\pm 0.01}$	$0.81 \pm 0.00$
M2N	Conf. Sampling	$29.37{\scriptstyle~\pm0.12}$	$0.90{\scriptstyle~\pm 0.00}$	$55.55 \pm 0.11$	$12.91{\scriptstyle~\pm 0.08}$	$0.60{\scriptstyle~\pm 0.00}$	$39.95 \pm 0.08$		Conf. Sampling	$0.93{\scriptstyle~\pm 0.01}$	$0.80 \pm 0.00$
	Const. Weight CS*	$29.37{\scriptstyle~\pm0.12}$	$0.90{\scriptstyle~\pm 0.00}$	$55.55 \pm 0.11$	$12.91{\scriptstyle~\pm 0.08}$	$0.60{\scriptstyle~\pm 0.01}$	$39.95 \pm 0.08$		Const. Weight CS*	$0.91 \pm 0.02$	$0.80 \pm 0.00$
	Non-Ex. CS*	$29.37{\scriptstyle~\pm 0.12}$	$0.90{\scriptstyle~\pm 0.00}$	$55.55 \pm 0.11$	$12.91 \pm 0.08$	$0.60{\scriptstyle~\pm 0.01}$	$39.95 \pm 0.08$		Non-Ex. CS*	$0.92{\scriptstyle~\pm 0.01}$	$0.81 \pm 0.00$

(a) Generation results for the de  $\rightarrow$  en and ja  $\rightarrow$  en translation tasks.

(b) Results for the open text generation.

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Table 4: Generation results for the two tasks. We report performance using 5 beams for beam-search, top-k sampling with k = 10, and nucleus sampling with p = 0.9. Conformal methods all use  $\alpha = 0.1$ , with non-exchangeable variants retrieving 100 neighbors. MT results for sampling use a softmax temperature of 0.1. Our methods are marked with \*. Results using 5 different seeds that are stat. significant according to the ASO test (Del Barrio et al., 2018; Dror et al., 2019; Ulmer et al., 2022b) with a confidence level of 0.95 and threshold  $\varepsilon_{\min} \leq 0.3$  are underlined.

sometimes on par or even outperformed by our non-exchangeable conformal sampling for MT. For text generation, our method performs best for the smaller OPT model but is slightly beaten by conformal nucleus sampling in terms of MAUVE. When using constant weights, performance deteriorates to the conformal sampling setup, emphasizing the importance of not considering all conformity scores equally when computing  $\hat{q}$ , even though the effect seems to be less pronounced for larger models. This illustrates the benefit of creating flexible prediction sets that are adapted on token-basis, suggesting that both the latent space neighborhoods induced by the model as well as the conformity scores are informative.

#### 5 Discussion

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Our experiments have shown that despite the absence of i.i.d. data in NLG and the loss in coverage induced by using dynamic calibration sets, the resulting coverage is still close to the pre-specified desired level for both LM and MT. Additionally, even though the coverage gap predicted by the method of Barber et al. (2023) is infeasible to quantify for us, we did not observe any critical degradation in practice. Further, we demonstrated how sampling from these calibrated prediction sets performs similarly or better than other sampling methods. Even though our method is still outperformed by beam search in the MT setting, previous work such as minimum bayes risk decoding has shown how multiple samples can be re-ranked to produce better outputs (Kumar and Byrne, 2004; Eikema and Aziz, 2020; Freitag et al., 2023; Fernandes et al., 2022). Additionally, recent dialogue systems based on LLMs use sampling instead of beam search for generation. Since our prediction sets are more flexible and generally tighter, our results serve as a starting point for future work. For instance, our technique could be used with new non-conformity scores that do not consider token probabilities alone (e.g. Meister et al., 2023) or using prediction set widths as a proxy for model uncertainty (Angelopoulos et al., 2021a).

#### 6 Conclusion

We successfully demonstrated the application of a non-exchangeable variant of conformal prediction to machine translation and language modeling with the help of k-NN retrieval. We showed our method to be able to maintain the desired coverage best across different dataset strata while keeping prediction sets smaller than other sampling methods. We validated our method to produce encouraging results for generation tasks. Lastly, we analyzed the behavior under distributional drift, showing how the k-NN retrieval maintains desirable properties for the estimated prediction sets.

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Limitations

We highlight two main limitations of our work 538 here: Potential issues arising from different kinds 539 of dataset shift as well as efficiency concerns. 540

**Distributional Drifts.** Even though any loss of 541 coverage due to the term quantifying distributional 542 drift in Equation (4) was limited in our experiments (see Sections 4.1 and 4.2), this might not hold across all possible setups. As long as we 545 cannot feasibly approximate the shift penalty, it is impossible to determine a priori whether the loss of 547 coverage might prove to be detrimental, and would 548 have to be checked in a similar way as in our ex-549 periments. Furthermore, we only consider shifts between the models' training distributions and test data distributions here, while many other, uncon-552 sidered kinds of shifts exist (Moreno-Torres et al., 553 2012; Hupkes et al., 2022). 554

Computational Efficiency. Even using opti-555 mized tools such as FAISS (Johnson et al., 2019), 556 moving the conformal prediction calibration step to inference incurs additional computational cost during generation. Nevertheless, works such as He et al. (2021b); Martins et al. (2022) show that there are several ways to improve the efficiency of k-NN approaches, and we leave such explorations to future work.

#### **Ethical Considerations** 564

The main promise of conformal prediction lies in its correctness-i.e. producing prediction sets that contain the correct prediction and are thus reliable. In an application, this could potentially create a false sense of security. On the one hand, the conformal guarantee holds in expectation, and not necessarily on a per-sample basis. On the other hand, our experiments have demonstrated that coverage might also not hold when distributional shifts are at work or when looking at specific subpopulations. Therefore, any application should certify that coverage is maintained for potentially sensitive inputs.

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## **A** Experimental Appendix

In this appendix, we bundle more details about experiments and their results. Appendix A.1 details the procedure to determine the temperature in Equation (5). We present more results from the experiments in Section 4.1 in Appendix A.2.

We illustrate the overall algorithm in Appendix A.4 and estimate environmental impact of our work in Appendix A.5.

#### A.1 Temperature Search

In order to determine the temperature used in Equation (5) for the different distance metrics in Table 1, we adopt a variation of a simple hill-climbing procedure. Given user-defined bounds for the temperature search  $\tau_{\min}$  and  $\tau_{\max}$ , we sample an initial candidate  $\tau_0 \sim \mathcal{U}[\tau_{\min}, \tau_{\max}]$ , and then evaluate the coverage of the method given the candidate on the first 100 batches of the calibration dataset. The next candidate then is obtained via

$$\tau_{t+1} = \tau_t + \eta \cdot \varepsilon \cdot \operatorname{sgn}(1 - \alpha - \operatorname{Coverage}(\tau_t));$$

$$\varepsilon \sim \mathcal{N}(0, \tau_{\max} - \tau_{\min}),$$
(10)
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where  $\eta$  is a predefined step size (in our case 0.1) and Coverage( $\tau_t$ ) the achieved coverage given a candidate  $\tau_t$ . The final temperature is picked after a fixed number of steps (t = 20 in our work) based on the smallest difference between achieved and desired coverage.

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Overall, we found useful search ranges to differ greatly between datasets, models, and distance metrics, as illustrated by the reported values in Table 1 and Table 2. In general, the stochastic hillclimbing could also be replaced by a grid search, even though we sometimes found the best temperature to be "hidden" in a very specific value range. It also has to be noted that temperature for the  $l_2$ distance is the highest by far since FAISS returns *squared*  $l_2$  distances by default.

#### A.2 Additional Coverage Results

We show additional plots illustrating the coverage per set size-bins in Figure 4. We can see the counterparts for Figure 2 using the larger  $M2M100_{(1.2B)}$ model in Figures 4a and 4b: Instead of leveling off like for the smaller model, most prediction set sizes are either in a very small range or in a size of a few ten thousand. In Figures 4c and 4d, we show similar plots for the two different OPT model sizes. Since in both cases, most prediction set sizes are rather small, we zoom in on the the sizes from 1 to 100. Here, we can observe a similar behavior to the smaller M2M100<sub>(400m)</sub>, gradually leveling off. We do not show similar plots for other distance metrics as they show similar trends.

#### A.3 Impact of Coverage Threshold and Neighborhood Size Choice

In this section, we present experiments surrounding the two most pivotal parameters of our method: The desired confidence level  $\alpha$ , as well as the number of neighbors.

**Coverage Threshold.** In Table 5, we investigate 1115 the impact of different values on  $\alpha$  on our evalua-1116 tion metrics. We show that the increase in  $\alpha$  does 1117 indeed produce the expected decrease in coverage, 1118 however with a certain degree of overcoverage for 1119 the de  $\rightarrow$  en MT and the LM task. The loss in 1120 1121 coverage always goes hand in hand with a decrease in the average prediction set width as well, as the 1122 model can allow itself to produce tighter prediction 1123 sets at the cost of higher miscoverage. As this also 1124 produces bin in which all contained instances are 1125









(c) Conditional coverage for  $OPT_{(350M)}$  on Language Modelling.



(d) Conditional coverage for  $OPT_{(1.3B)}$  on Language Modelling.

Figure 4: Additional conditional coverage plots for the MT and LM dataset using our non-exchangeable conformal prediction method, aggregating predictions by prediction set size. The blue curve shows the conditional coverage per bin, whereas red bars show the number of predictions per bin. For Figures 4c and 4d, we zoom in on the prediction set sizes from 1 and 100.

uncovered, this produces zero values for the SCC,	
while we cannot discern clearn trends for the ECG.	

Neighborhood Size.In Table 6, we vary the ef-<br/>fect of the chosen neighborhood size (with 100112811291129

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	$\alpha$	% Cov.	$arnothing$ Width $\downarrow$	$\mathbf{SCC}\uparrow$	Ecg↓
	0.1	0.9442	0.31	0.8702	0.0011
E.	0.2	0.8767	0.18	0.7906	$8.63\times 10^{-5}$
e ↑	0.3	0.7963	0.12	0	0.0016
/ de	0.4	0.7058	0.09	0.1393	0.0082
(M004	0.5	0.6081	0.07	0.2836	0.0055
100 <sub>(4</sub>	0.6	0.5017	0.06	0.1393	0.0082
2M	0.7	0.3896	0.05	0	0.0091
Σ	0.8	0.2800	0.05	0	0.0090
	0.9	0.1762	0.04	0	0.0071
	0.1	0.7453	0.15	0.3080	0.1511
œ	0.2	0.5579	0.07	0.2728	0.2446
e ↑	0.3	0.4277	0.04	0.2770	0.2779
/ja	0.4	0.3438	0.03	0.1212	0.2438
400M)	0.5	0.2749	0.03	0.0455	0.1883
$100_{(i)}$	0.6	0.2175	0.02	0	0.1207
I2M	0.7	0.1685	0.02	0	0.0560
Σ	0.8	0.1309	0.01	0	0.0117
	0.9	0.0989	0.02	0	0.0099
	0.1	0.9460	0.26	0.8	$1.85  imes 10^{-5}$
EXT	0.2	0.8937	0.16	0.8	0
BTJ	0.3	0.8392	0.10	0.5	$8.74\times10^{-6}$
NWI	0.4	0.7782	0.08	0.6667	0
OPE	0.5	0.7171	0.06	0	$1.19\times 10^{-5}$
M) /	0.6	0.6559	0.06	0.6033	0
ľ	0.7	0.5945	0.05	0	$8.21\times 10^{-6}$
OPJ	0.8	0.5349	0.05	0.4462	0
	0.9	0.4757	0.05	0.3580	0

Table 5: Results for different values of  $\alpha$  using different models and datasets.

Table 6: Results for different neighborhood sizes K using different models and datasets.

	K	% Cov.	$\varnothing \text{ Width } {\downarrow}$	$SCC\uparrow$	Ecg↓
	10	0.9923	0.39	0.9728	0
→ en	25	0.9563	0.37	0.8877	0.0011
de -	50	0.9504	0.32	0.8870	0.0006
/ (M(	75	0.9444	0.32	0.8641	0.0014
0(400	100	0.9442	0.31	0.8702	0.0011
MIO	200	0.9422	0.31	0.8125	0.0016
M2	300	0.9404	0.31	0.8483	0.0019
	500	0.9389	0.31	0.8214	0.0023
	10	0.8013	0.17	0.2995	0.1606
→ en	25	0.7353	0.17	0.2994	0.1438
ja –	50	0.7540	0.17	0.3023	0.1603
/ (MC	75	0.7368	0.16	0.3019	0.1603
$0_{(40)}$	100	0.7453	0.15	0.3072	0.1529
MIC	200	0.7295	0.14	0.2938	0.1787
M2	300	0.7192	0.13	0.2948	0.1788
	500	0.7110	0.13	0.2756	0.1867
Ţ	10	0.9438	0.35	0.8824	0.0019
TEX	25	0.9522	0.33	0.8333	$2.06\times 10^{-5}$
WEB	50	0.9442	0.27	0	$1.86\times 10^{-5}$
PEN	75	0.9477	0.27	0.8	$1.03\times 10^{-5}$
0	100	0.9460	0.26	0.8	$1.86\times 10^{-5}$
\$50M)	200	0.9487	0.28	0.8571	$6.20\times 10^{-5}$
PT	300	0.9500	0.28	0.8181	$1.86\times 10^{-5}$
0	500	0.9508	0.29	0.8181	$1.86\times 10^{-5}$

our experiments, we leave more principled ways to determine the neighborhood size to future work.

#### A.4 Algorithm

We show the algorithm that was schematically depicted in Figure 1 in pseudo-code in Algorithm 1. It mostly requires that we have pre-generated a datastore of latent representations of the model on a held-out set along with their non-conformity scores (in our case, using the score defined in 6 and the FAISS (Johnson et al., 2019) as the datastore architecture). Furthermore, we need to have determined an appropriate value for the temperature  $\tau$  in advance (see Appendix A.1). Then, the algorithm involves the following steps:

1. Extract the latent encoding for the current time1158step  $\mathbf{z}_t$  from the model. Even though different1159

being the value we use in our main experiments). 1130 We make the following, interesting observations: 1131 Coverage on the MT task seems to decrease with an 1132 increase in the neighborhood size as prediction set 1133 widths get smaller on average, with a neighborhood 1134 size around 100 striking a balance between cover-1135 age, width, computational cost and SCC / ECG. For 1136 LM, coverage seems to be mostly constant, with 1137 prediction set width hitting an inflection point for 1138 1139 100 neighbors. We speculate that initially there might be a benefit to considering more neighbors 1140 to calibrate  $\hat{q}$ , but that considering too large neigh-1141 borhoods might introduce extra noise. While we 1142 found 100 to be a solid choice for the purpose of 1143

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Algorithm 1 Non-exchangeable Conformal Language Generation with Nearest Neighbors

**Require:** Sequence  $\mathbf{x}^{(i)}$ , model  $f_{\theta}$ , datastore  $DS(\cdot)$  with model activations collected from held-out set, temperature  $\tau$ 

while generating do

 $\triangleright$  1. Extract latent encoding for current input  $\mathbf{z}_t^{(i)} \leftarrow f_{\boldsymbol{\theta}}(\mathbf{x}_t)$ 

 $\triangleright$  2. Retrieve K neighbors & non-conformity scores

$$\{(\mathbf{z}_1, s_1), \dots (\mathbf{z}_K, s_K)\} \leftarrow \mathsf{DS}(\mathbf{z}_t)$$

 $\triangleright 3. \text{ Compute weights } w_k \text{ and normalize} \\ w_k \leftarrow \exp(-||\mathbf{z}_t^* - \mathbf{z}_k||_2^2 / \tau) \\ \tilde{w}_k \leftarrow w_k / (1 + \sum_{k=1}^K w_k)$ 

▷ 4. Find quantile 
$$\hat{q}$$
  
 $\hat{q} \leftarrow \inf\{q \mid \sum_{i=1}^{N} \tilde{w}_i \mathbf{1}\{s_i \le q\} \ge 1 - \alpha\}$ 

▷ 5. Create prediction set  $\hat{c} \leftarrow \sup\{c' | \sum_{j=1}^{c'} p_{\theta}(y = \pi(j) | \mathbf{x}^*) < \hat{q}\} + \mathcal{C}(\mathbf{x}^*) \leftarrow \{\pi(1), \dots, \pi(\hat{c})\}$ 

 $\triangleright$  6. Generate next token  $y_t \leftarrow \text{generate}(\mathcal{C}(\mathbf{x}^*))$ end while

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options are imaginable, we utilize the activa-<br/>tions of the uppermost layer.11601161

- 2. Retrieve K neighbors and their corresponding<br/>non-conformity scores from the datastore.11621163
- 3. Compute the weights  $w_k$  based on the squared1164 $l_2$  distance between  $\mathbf{z}_t$  and its neighbors in the1165datastore and normalize the weights to obtain1166 $\tilde{w}_k$ .1167

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- 4. Use Equation (3) to find the quantile  $\hat{q}$ .
- 5. Use  $\hat{q}$  to create prediction sets, for instance the adaptive prediction sets defined in Equation (7).
- 6. Finally, generate the new token  $y_t$  by sampling from the prediction set.

The main computational bottleneck of this algorithm is the retrieval process that fetches the closest neighbors from the datastore during every generation step. However, while not explored further in this work, there are some potential avenues to reduce this load: On the one hand, works such as He et al. (2021b); Martins et al. (2022) have demonstrated ways to reduce the computational load of k-NN based approaches. On other hand, we treat the number of neighbors K fixed during every generation step. However, it seems intuitive that the number of neighbors necessary to create good prediction sets would not be the same for all tokens. Future research could explore setting Kdynamically during every time step, thus reducing the overall slowdown.

## A.5 Environmental Impact

We track the carbon emissions produced by this work using the codecarbon tracking tool (Schmidt et al., 2021; Lacoste et al., 2019; Lottick et al., 2019). The carbon efficiency was estimated to be 0.12 kgCO<sub>2</sub>eq / kWh. 159.5 hours of computation were performed on a NVIDIA RTX A6000. Total emissions are estimated to be 6.99 kgCo2eq. All of these values are upper bound including debugging as well as failed or redundant runs, and thus any replication of results will likely be shorter and incur fewer carbon emissions.