

Twist Decoding: Diverse Generators Guide Each Other

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Abstract

Natural language generation technology has recently seen remarkable progress with large-scale training, and many natural language applications are now built upon a wide range of generation models. Combining diverse models may lead to further progress, but conventional ensembling (e.g., shallow fusion) requires that they share vocabulary/tokenization schemes. We introduce TWIST decoding, a simple and general inference algorithm that generates text while benefiting from diverse models. Our method does not assume the vocabulary, tokenization or even generation order is shared. Our extensive evaluations on machine translation and scientific paper summarization demonstrate that TWIST decoding substantially outperforms each model decoded in isolation over various scenarios, including cases where domain-specific and general-purpose models are both available. TWIST decoding also consistently outperforms the popular reranking heuristic where output candidates from one model is rescored by another. We hope that our work will encourage researchers and practitioners to examine generation models collectively, not just independently, and to seek out models with complementary strengths to the currently available models.¹

1 Introduction

Natural language generation has now become an important building block for many applications, such as machine translation, summarization, and question answering (Ng et al., 2019; Lewis et al., 2020; Raffel et al., 2020; Brown et al., 2020; Asai et al., 2021, *inter alia*). Researchers have recently explored and advanced models for generation in various aspects, including model architecture (Bahdanau et al., 2015; Vaswani et al., 2017), domain

¹Our code is available at https://github.com/jungokasai/twist_decoding.

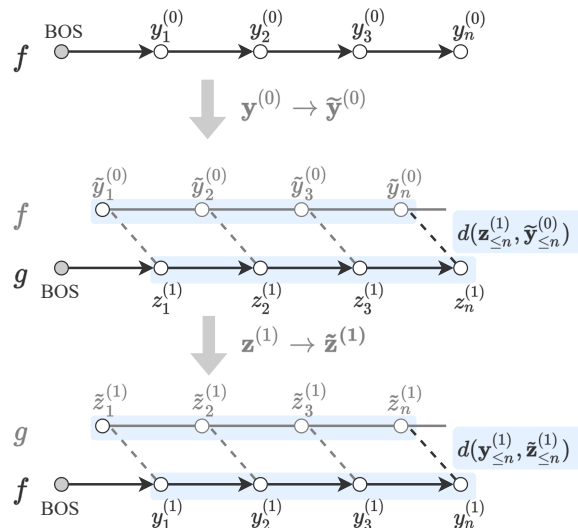


Figure 1: TWIST decoding of two generation models, f and g , that does not assume a shared vocabulary, tokenization, or generation order. Beam search is first applied to f to generate $y^{(0)}$, followed by output mapping to $\tilde{y}^{(0)}$ (e.g., f 's detokenization and g 's tokenization). g is then decoded with beam search augmented with distances from the set of previously-generated outputs (here only one sequence y is shown): $d(z_{\leq n}^{(1)}, \tilde{y}_{\leq n}^{(0)})$. Subsequently, f is similarly decoded with g 's guidance. Here we show one iteration that already achieves substantial improvements (§4), but additional iterations can be made.

adaptation (Chu and Wang, 2018; Bapna and Firat, 2019), prompting (Brown et al., 2020), and even generation order (Gu et al., 2018). The resulting generation models are diverse, trained on different data, with different assumptions, at different times. We hypothesize that diverse generation models may achieve better results through ensembling, if the various approaches have complementary strengths. Given the high cost of unifying approaches during training time (Strubell et al., 2019; Schwartz et al., 2019), inference-time combination of existing models is an attractive alternative.

One well established ensembling technique is

“shallow fusion” (Sutskever et al., 2014; Gulcehre et al., 2015; Firat et al., 2016, *inter alia*), which simply aggregates models’ scores during beam search. This approach requires, however, that the models use the same vocabulary/tokenization scheme and organize the search in the same way (e.g., autoregressive, left-to-right factorization).

We introduce a new inference algorithm, TWIST decoding (Fig. 1), that enables more diverse generators to guide each other. TWIST decoding can combine generators with different vocabularies, (de)tokenization, and even generation order without any additional training or finetuning. Our method decodes a model by standard beam search, but the scores at every step incorporate a simple function that measures the distance from outputs of the other model. We run this procedure on each generation model in turn, so that both can benefit from each other.

We present extensive experiments on machine translation and scientific paper summarization and show that TWIST decoding can improve performance over each model decoded in isolation across several scenarios: combining 1) generic and domain-specific models, 2) left-to-right and right-to-left generation models, and 3) models that generate using different conditioning inputs. Our results show consistent performance gains from combining generic and domain-specific translation models over a wide range of domains, including medical and legal translation. Applications in these domains require particularly high accuracy, and TWIST decoding is a desirable alternative to standard beam search on a single model. Interestingly, we find that TWIST decoding between generic and domain models is effective even when parallel data from the domain are scarce and the domain model yields poor performance by itself, suggesting complementary strengths of diverse generators (§3.4).

TWIST decoding can be seen as a generalization of reranking heuristics that have proven effective in syntactic parsing (Shen and Joshi, 2003; Collins and Koo, 2005), speech recognition (Collins et al., 2005), and machine translation (Shen et al., 2004; Och et al., 2004): one model generates candidate sequences, followed by rescoring from another model. We present extensive comparisons with reranking baselines and demonstrate that TWIST decoding outperforms reranking consistently. We also observe that since the encoder computations on two models can be parallelized, the inference time re-

TWIST Decoding

g generates \mathbf{z} with guidance from f at iteration t

k : beam size. M : maximum length.

\mathcal{V}_g : vocabulary of g . $g(\cdot)$: scoring function.

$\mathcal{Y}^{(t-1)}$: set of output sequences from f . $\mathcal{Z}^{(t)}$: new outputs.

B_n : beam of continuing sequences.

H : expanded hypotheses before beam selection.

$d(\cdot, \cdot)$: distance between partial sequences.

λ_f : scalar coefficient for the distance.

```

1:  $\tilde{\mathcal{Y}}^{(t-1)} = \{\tilde{\mathbf{y}} = \text{map\_output}(\mathbf{y}) \mid \mathbf{y} \in \mathcal{Y}^{(t-1)}\}$ 
2:  $B_0 \leftarrow \{\text{BOS}\}$ ,  $\mathcal{Z}^{(t)} \leftarrow \emptyset$ 
3: for  $n \in \{1, \dots, M\}$  :
4:    $H \leftarrow \emptyset$ 
5:   for  $\mathbf{z} \in B_{n-1}$  : # Expansion.
6:     for  $z \in \mathcal{V}_g$  :
7:        $s \leftarrow g(\mathbf{z} \circ z) - \lambda_f \min_{\tilde{\mathbf{y}} \in \tilde{\mathcal{Y}}^{(t-1)}} d(\mathbf{z} \circ z, \tilde{\mathbf{y}}_{\leq n})$ 
8:        $H.\text{add}(\langle s, \mathbf{z} \circ z \rangle)$ 
9:    $B_n \leftarrow \text{topk}(H)$ ,  $\mathcal{Z}^{(t)}.\text{add}(\text{finished}(H))$ 
10: return  $\mathcal{Z}^{(t)}$ 

```

Figure 2: TWIST decoding when g is guided by f . Swap f and g and \mathbf{y} and \mathbf{z} to obtain $\mathcal{Y}^{(t)}$. `map_output` converts outputs from f to g ; e.g., f ’s detokenization followed by g ’s tokenization. It would also include sequence reversal if f or g is a right-to-left model. The highlighted line is the *only* modification that TWIST decoding introduces to standard beam search. The input sequence to g is omitted. See also Kasai et al. (2022b) for the stopping criterion and implementation details (the *first come, first served* heuristic).

quired for TWIST decoding is much shorter than the sum of the two models, resulting only in a 50% increase, relative to decoding of a single model in isolation (§4). TWIST decoding is therefore a viable alternative to standard beam search on a single model and the widespread reranking heuristic.

2 TWIST Decoding

We propose TWIST decoding, a general decoding algorithm that generates text from diverse models without assumptions of a shared vocabulary, tokenization, or generation order. At the core of the algorithm is a simple modification in standard beam search (highlighted in Fig. 2); we incorporate into a scoring function the distance from outputs that are previously generated by another model.

2.1 Initial Decoding

Let us assume that we have two generation models: f and g .² Both f and g assign scores to output

²The algorithm can be readily extended to three or more generators. We also abuse f or g to mean both the generator and its scoring function.

sequences.³ f is, for instance, a domain-specific translation model and g a generic translation model. f and g perform their own pre/postprocessing (e.g., (de)tokenization) and factorization (e.g., left-to-right or right-to-left factorization). Here we suppress for brevity the conditioned-upon input (e.g., machine translation input from the original language). Standard beam search with beam size k is first applied to f to produce a set of k output sequences: $\mathcal{Y}^{(0)}$. This approximately solves $\text{topk}_y f(y)$ by pruned breadth-first search, and often returns higher-quality outputs than the exact search counterpart (Stahlberg and Byrne, 2019; Meister et al., 2020a).

2.2 Mutually-Guided Decoding

Once $\mathcal{Y}^{(0)}$ is obtained, we proceed with decoding generators with mutual guidance (Fig. 2; $t \geq 1$)

Output Sequence Mapping The commonly-used technique of ensembling (Sutskever et al., 2014) or shallow fusion (Gulcehre et al., 2015; Stahlberg et al., 2018) adds scores from f and g at every step and executes the same search algorithm to approximately solve $\text{topk}_y f(y) + g(y)$. This method thus necessitates a shared vocabulary, tokenization, and generation order (Imamura and Sumita, 2017). We depart from this assumption and first map the candidates in $\mathcal{Y}^{(t-1)}$ to output sequences for g : $\tilde{\mathcal{Y}}^{(t-1)}$ (Line 1 in Fig. 2). This mapping (`map_output`) typically involves deterministic operations of f 's detokenization followed by g 's tokenization. Sequence reversal is also performed if f and g generate in the opposite order. For example, if g uses byte-pair encoding (Sennrich et al., 2016b), but f does not, we might have $y = \textit{John does n't like Mary}$ mapped to $\tilde{y} = \textit{Jo@ hn doesn't like Mar@ y}$, where `@` denotes subword separation.

Decoding with Distance Terms We then decode g with guidance from $\tilde{\mathcal{Y}}^{(t-1)}$. Specifically, we perform beam search with a simple modification in scoring (Line 7). In this work, we use a simple distance measure that adds binary distances at all positions (i.e., the Hamming distance):

$$d(\mathbf{z}_{\leq n}, \tilde{\mathbf{y}}_{\leq n}) = \sum_{i \leq n} \mathbb{1}\{z_i \neq \tilde{y}_i\}$$

We also explored using the distance between (sub)word embeddings from the model:

³They typically assign log-probabilities, but it is not necessary to assume the scores form a valid probability distribution.

$\sum_{i \leq n} \|e(z_i) - e(\tilde{y}_i)\|_2$, but this did not bring improvements (§4). Note also that when i exceeds the length of $\tilde{\mathbf{y}}$, we assume $\tilde{y}_i = \text{EOS}$. The overall distance term is then

$$\min_{\tilde{\mathbf{y}} \in \tilde{\mathcal{Y}}^{(t-1)}} d(\mathbf{z}_{\leq n}, \tilde{\mathbf{y}}_{\leq n})$$

Here we minimize over the output sequences to compute the distance to the closest candidate. These candidates from $\tilde{\mathcal{Y}}^{(t-1)}$ can be equally good outputs but differ only by one word; in such cases, this minimization operation avoids overestimation of the distances. The new score at step n in beam search is now computed by:

$$g(\mathbf{z}_{\leq n}) - \lambda_f \min_{\tilde{\mathbf{y}} \in \tilde{\mathcal{Y}}^{(t-1)}} d(\mathbf{z}_{\leq n}, \tilde{\mathbf{y}}_{\leq n}),$$

where λ_f is a scalar coefficient for the distance term that controls the importance of f relative to g . We tune $\lambda_f \in \{0.1, 0.3, 1.0, 3.0\}$ during development. After this beam search, we obtain a new candidate set, $\mathcal{Z}^{(t)}$. We then run the same beam search (Fig. 2) with the roles of f , \mathcal{Y} and g , \mathcal{Z} swapped.⁴ Namely, we decode f with distance terms from $\mathcal{Z}^{(t)}$ at each step of beam search:

$$f(\mathbf{y}_{\leq n}) - \lambda_g \min_{\tilde{\mathbf{z}} \in \mathcal{Z}^{(t)}} d(\mathbf{y}_{\leq n}, \tilde{\mathbf{z}}_{\leq n})$$

Finally, the highest-scoring sequence from $\mathcal{Y}^{(t)}$ is output. This process of mutually-guided decoding can be repeated multiple times. We observe, however, that one iteration ($t = 1$) suffices to bring performance gains (§4). We also present detailed sensitivity analysis over varying λ_f and λ_g and find that TWIST decoding is particularly effective when $\lambda_g > \lambda_f$ (i.e., initial exploration by g is encouraged with relatively little guidance from f 's original outputs; see §4).

Reranking Heuristic as a Special Case Notice that as $\lambda_f \rightarrow \infty$, g 's generation falls back to a reranking heuristic: top k sequences from the initial f decoding are reranked according to g . This reranking heuristic has proven successful in a wide range of sequence generation tasks, including machine translation (Shen et al., 2004), syntactic parsing (Collins and Koo, 2005), and speech recognition (Collins et al., 2005). Reranking is performed in many strong machine translation systems to use a right-to-left model to improve a left-to-right model; e.g., top-performing systems in recent WMT competitions (Ng et al., 2019; Kiyono et al., 2020; Wang et al., 2021; Akhbardeh et al., 2021).

⁴We can stop inference with $\mathcal{Z}^{(t)}$, but we found that led to performance degradation in preliminary development.

German→English			Medicine		Law		Koran		Subtitles	
Method	f	g	COMET	BLEU	COMET	BLEU	COMET	BLEU	COMET	BLEU
Isolation	Generic	–	44.5	41.2	30.6	34.8	14.4	16.6	34.4	31.3
Isolation	Domain	–	80.7	48.3	60.7	40.9	8.7	17.0	32.3	29.0
Rerank	Generic	Domain	59.6	43.5	56.4	36.1	14.7	17.0	40.3	32.3
TWIST	Generic	Domain	71.6	47.5	61.4	40.2	16.5	18.5	41.0	32.2
Δ (TWIST– Rerank)			+12.0	+4.0	+5.0	+4.1	+1.8	+1.5	+0.7	–0.1
Rerank	Domain	Generic	75.8	48.7	59.9	40.8	12.3	18.1	36.5	30.3
TWIST	Domain	Generic	81.6	50.1	61.6	41.3	15.3	18.7	37.3	31.0
Δ (TWIST– Rerank)			+5.8	+1.4	+1.7	+0.5	+3.0	+0.6	+0.8	+0.7

Table 1: Combination of generic and domain-specific translation models. The generic model is the top-performing translation model in WMT19 (Ng et al., 2019) that is trained on a collection of parallel corpora, such as the Europarl and the UN corpora. Two settings are considered for each of the reranking baseline and our TWIST decoding: f is the generic model and g is the domain-specific model or the reverse. The best scores are in **bold**. COMET (Rei et al., 2020a,b) uses crosslingual contextual representations (Conneau et al., 2020) and achieves significantly higher correlation with expert human judgment than BLEU (Papineni et al., 2002) and other alternative metrics (Kasai et al., 2022a,c).

In our experiments, we extensively compare performances of TWIST decoding and reranking and demonstrate that the former consistently outperforms the latter.

3 Experiments

We present experiments across three scenarios: decoding domain and generic models for machine translation (§3.1), left-to-right and right-to-left machine translation models (§3.2), and scientific paper summarization models that take as input different parts of the paper (§3.3). We empirically compare TWIST decoding with decoding in isolation and the widely-adopted reranking baselines, illustrating that TWIST decoding offers performance improvements in various situations *without* any change to the trained models.

3.1 Domain and Genric Models

Machine translation has now been used over many domains, ranging from everyday conversations to medical documents. Machine translation models are often trained on large amounts of parallel data, such as the Europarl corpus (Koehn, 2005) and the OPUS data (Tiedemann, 2012). Applying these models to out-of-domain data remains a challenge (Koehn and Knowles, 2017; Chu and Wang, 2018), and some of these domains require particularly high accuracy in translation (e.g., medical and legal documents). We will demonstrate that TWIST decoding between general-purpose and domain-specific models is a viable approach to tackle this problem.

Setups We use machine translation datasets over diverse domains from prior work (Koehn and Knowles, 2017; Hu et al., 2019): German→English over medical (1.1M training sentence pairs), legal (720K pairs), Koran (religious text, 480K pairs), and subtitles (14M pairs) domains.⁵ For the domain-specific models, we train a base-sized transformer model (Vaswani et al., 2017) with a 6-layer encoder and a 6-layer decoder on the training data of each domain. The top-performing German→English system from WMT19 (Barrault et al., 2019; Ng et al., 2019)⁶ is used as the generic model. This generic model is a large-sized transformer trained on a concatenation of publicly available parallel data, including the Europarl (Koehn, 2005) and UN (Ziemski et al., 2016) corpora with the backtranslation technique (Sennrich et al., 2016a). We follow (de)tokenization (Koehn et al., 2007) and byte-pair encoding (Sennrich et al., 2016b) of previous work (Koehn and Knowles, 2017; Hu et al., 2019).⁷

For every domain, we evaluate a total of six configurations: decoding of the generic and domain models each in isolation; the reranking baseline and TWIST decoding with f being the generic model and g being the domain model, as well as the versions where f and g are swapped. In all cases, we use beam size 5 and length penalty 1

⁵We excluded the IT domain because we found significant overlap between training and dev/test data.

⁶<https://github.com/pytorch/fairseq/tree/main/examples/wmt19>.

⁷All data are available at <https://github.com/JunjieHu/dali>.

(Wu et al., 2016) and conduct all experiments using the `fairseq` library (Ott et al., 2019). All performance is measured with the COMET score (Rei et al., 2020a,b) and the SACREBLEU implementation (Post, 2018) of the BLEU score (Papineni et al., 2002). Note that COMET is based on crosslingual contextual representations (Conneau et al., 2020), and recent work showed that it achieves significantly higher correlation with expert human judgment than BLEU and other n-gram-based metrics (Kasai et al., 2022a,c). More experimental details are described in Appendix §A.1.

Results Seen in Table 1 are the results from our experiments over various domains. Firstly, given two translation models f and g , TWIST decoding outperforms the reranking baseline in all configurations (indicated in blue) with only one exception (a small drop in BLEU in the subtitles domain). Particularly noteworthy are the gains in the medical domain: TWIST decoding outperforms the reranking heuristic by 5.8 COMET and 1.4 BLEU points when f is the domain model and g is the generic model. TWIST decoding is thus an effective generalization over the reranking heuristic commonly used in the literature across domains.

Comparing the performance of decoding in isolation and TWIST decoding, we observe that the best score from TWIST decoding substantially outperforms each individual model over all domains: e.g., 81.6 vs. 80.7 (domain model) and 81.6 vs. 44.5 (generic model) in the medical domain. In both medical and legal domains, the generic model underperforms the domain model by a large margin. Nonetheless, TWIST decoding between the two improves over the domain model, suggesting that TWIST decoding makes use of their complementary strengths. Finally, we see a consistent pattern regarding f and g : both TWIST decoding and the reranking baseline perform better when the higher-performing model is chosen as f . This is expected because f is used both for initial decoding and final decoding with g 's guidance (Fig. 1).

3.2 Left-to-Right and Right-to-Left Models

Language generation models usually factorize sequences autoregressively in a left-to-right order, but previous work showed that left-to-right (L2R) models can be improved by reranking their outputs with a separate right-to-left (R2L) model (Imamura and Sumita, 2017; Ng et al., 2019; Kiyono et al., 2020, *inter alia*). TWIST decoding can be readily

applied to such scenarios since it does not assume shared generation order between models.

Setups We experiment with two language pairs from the WMT 2020 news translation task (Barrault et al., 2020): Chinese→English (WMT20 ZH-EN, 48M training sentence pairs) and English→German (WMT20 EN-DE, 48M pairs). Submissions for these language pairs to the shared task have human evaluations from professional translators (Freitag et al., 2021), and the correlation between automatic metrics and the human ratings are extensively studied in subsequent work (Kasai et al., 2022a); COMET (Rei et al., 2020b,a) achieves the highest correlation out of the 15+ metrics.

Similar to the previous experiments, we measure all performance in the COMET and BLEU scores. Note that we use two reference translations per instance for WMT20 ZH-EN and three for WMT20 EN-DE, following Kasai et al. (2022a). They both have reference translations from two different services, and WMT20 EN-DE has an additional translation created by linguists who are asked to paraphrase the two translations as much as possible. These paraphrased translations are shown to increase correlation with human judgments by mitigating the translationese effect (Graham et al., 2020) and diversifying the reference (Freitag et al., 2020). On each dataset, we follow the preprocessing and tokenization (Koehn et al., 2007; Sennrich et al., 2016b) from Kasai et al. (2022a)⁸ and train a large-sized transformer model for left-to-right and right-to-left translation, in which the output English/German sequences are reversed after tokenization. We implement all models and decoding with `fairseq` and apply beam search with beam size 5 and length penalty 1. We again consider a total of six settings: reranking and TWIST decoding with L2R as f and R2L as g or the reverse, as well as the individual models. Further details can be found in Appendix §A.2.

Results Table 2 shows the results from L2R and R2L translation models. TWIST decoding again outperforms the reranking counterpart by a considerable margin in COMET and BLEU on both language pairs; e.g., 43.1 vs. 41.2 COMET points on WMT20 ZH-EN when f is R2L and g is L2R. The best performance is achieved by TWIST decoding

⁸<https://github.com/jungokasai/billboard/tree/master/baselines>.

WMT20 Test			ZH→EN		EN→DE	
Method	f	g	COMET	BLEU	COMET	BLEU
Isolation	L2R	–	40.8	35.5	42.9	45.5
Isolation	R2L	–	40.4	35.0	43.3	44.9
Rerank	L2R	R2L	41.4	36.1	43.7	46.0
TWIST	L2R	R2L	42.8	36.8	45.4	46.7
Δ (TWIST– Rerank)			+1.4	+0.7	+1.7	+0.7
Rerank	R2L	L2R	41.2	35.4	44.7	45.2
TWIST	R2L	L2R	43.1	36.8	44.8	46.0
Δ (TWIST– Rerank)			+1.9	+1.4	+0.1	+0.8

Table 2: Combination of left-to-right (L2R) and right-to-left (R2L) transformer translation models. ZH: Chinese. DE: German. Two settings are considered for reranking and our TWIST decoding each: L2R or R2L as f . The best scores are in **bold**.

on both datasets and improves over the individual models by more than 1 BLEU point. The reranking baseline, on the other hand, does not outperform the L2R model in BLEU when f is R2L: 35.4 vs. 35.5 (ZH-EN) and 45.2 vs. 45.5 (EN-DE). This result illustrates that TWIST decoding is a more effective approach to combine models with different generation order than the popular reranking heuristic.

3.3 Summarization with Different Input

We also experiment with strong models on a highly abstractive scientific paper summarization task: **SciTLDR** (Cachola et al., 2020). Specifically, we use two BART-based models from prior work (Cachola et al., 2020) that differ in input type: one that only takes as input the paper abstract (Abst.) and the other a concatenation of the abstract, introduction, and conclusion (AIC).⁹

Setups We use the train/dev./test split from Cachola et al. (2020). Again following Cachola et al. (2020), we use all human-written summaries (written either by authors or undergraduate computer science students) as the reference and evaluate performance in terms of the ROUGE score (Lin, 2004).¹⁰ Similar to our previous experiments, we use beam size 5 and length penalty 1. See more detail in Appendix A.3.

Results Table 3 presents our results. TWIST decoding substantially outperforms the reranking baseline when f is the AIC model (e.g., +0.5

⁹Models are available at <https://github.com/allenai/scitldr>.

¹⁰We release our models and their outputs, so other metrics can be readily used as well in the future.

ROUGE-L points), but they yield (almost) the same performance when f is the Abst. model. Nonetheless, TWIST decoding achieves the best performance out of all configurations. Our small improvements might be attributed to the fact that the input to the Abst. model is a strict subset of the AIC model and there are only limited benefits from combining them.

SciTLDR Summ. Test			ROUGE		
Method	f	g	R-1	R-2	R-L
Isolation	Abst.	–	39.9	21.1	34.5
Isolation	AIC	–	40.2	21.3	34.9
Rerank	Abst.	AIC	40.5	21.7	35.1
TWIST	Abst.	AIC	40.5	21.7	35.0
Δ (TWIST– Rerank)			0.0	0.0	–0.1
Rerank	AIC	Abst.	40.1	21.2	34.8
TWIST	AIC	Abst.	40.7	22.1	35.3
Δ (TWIST– Rerank)			+0.6	+0.9	+0.5

Table 3: Combination of scientific paper summarization models with different input types. Both models are BART-based models from prior work (Cachola et al., 2020) with different input: abstract only (Abst.) or abstract, introduction, and conclusion (AIC). The best scores are in **bold**.

3.4 Low-Resource Scenarios

In our experiments over four diverse domains (§3.1), we assumed that plenty of parallel data is available in every domain, and the domain model generally outperformed the generic model. Concretely, we used 1.1M and 720K training sentence pairs for the medical and legal domains, based on the data splits from previous work (Koehn and Knowles, 2017; Hu et al., 2019). In real-world applications, however, these domain-specific translation data are often scarce since they need to be annotated by bilingual speakers with expertise in those domains. The question arises: *can a domain model trained on small parallel data still help the generic model by its complementary strengths?* To simulate such realistic, low-resource scenarios, we randomly sample [10k, 20k, 40k, 80k] sentence pairs and conduct the same evaluations with the generic and domain models as f and g , respectively.

Fig. 3 plots COMET scores of various decoding methods on the medical and legal domains. The score from the generic model is constant because we only change the domain training data. There is a striking trend: even though the domain model

performs poorly by itself, it improves the generic model through TWIST decoding over varying sizes. Reranking also helps the generic model as the data size increases, but the improvement is less pronounced than that of TWIST decoding.

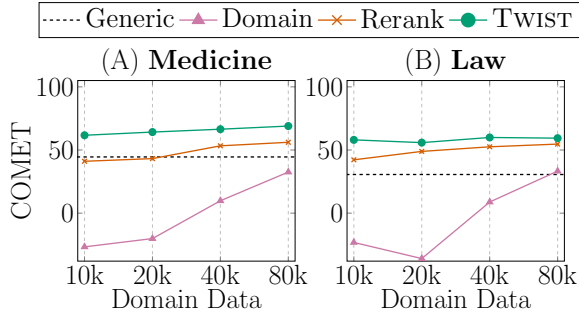


Figure 3: Results when parallel data are scarce in the target domain. Both TWIST decoding and reranking use the generic model as f and the domain model as g . COMET (Rei et al., 2020a) is a regression-based metric that can take negative values.

4 Analysis

Iterations So far, we have only applied one iteration of TWIST decoding, but Fig. 4 plots performance over multiple iterations. Iteration 0 signifies f ’s initial decoding ($y^{(0)}$ in Fig. 1), and every iteration involves g ’s decoding with f ’s guidance ($z^{(t)}$) and its reverse ($y^{(t)}$). We observe that the first iteration brings most of the performance gains. This makes TWIST decoding practically appealing, as it improves performance without much increase in the overall computation or inference time (see below).

Inference Time Table 4 reports the runtime of each decoding method, relative to f ’s decoding in isolation. We use batch size 1 on the same single A100-SXM GPU and measure the wall-clock time from when all models are loaded until all outputs are obtained. As expected, TWIST decoding results in a slowdown compared to decoding in isolation, but the increase in time is only 50%. The inference time for TWIST decoding is much shorter than the sum of f and g in isolation ($1.4\times$ vs. $2.1\times$ on medical translation) because 1) the encoder computation for f and g can be parallelized and 2) the encoder computation for f is done only once while we need two runs of f ’s decoder. We leave it to future work to further speed up TWIST decoding; since the slowdown of TWIST decoding primarily comes from the decoder, it can be sped

up by best-first beam search (Meister et al., 2020b), a deep-encoder, shallow-decoder strategy (Kasai et al., 2021a), or a fast, linear-complexity variant of the transformer decoder (Peng et al., 2021; Kasai et al., 2021b) that is shown to retain the performance of the standard encoder-decoder transformer. Another approach could be sequence-level knowledge distillation (Kim and Rush, 2016), which has proven successful in speeding up an ensemble translation model (Freitag et al., 2017).

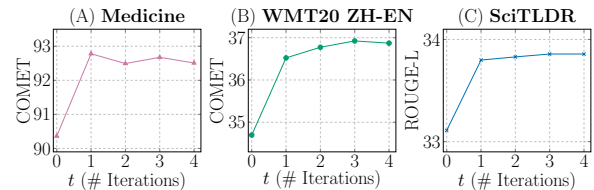


Figure 4: Effects of iterations on dev. performance. Iteration 0 refers to the initial decoding from f . Every iteration consists of g ’s decoding with f ’s guidance followed by f ’s decoding with g ’s guidance.

Inference Method	Medicine		Time	WMT20 ZH→EN		Time
	f	g		f	g	
Isolation Domain	–	–	1.0×	R2L	–	1.0×
Isolation Generic	–	–	1.1×	L2R	–	1.0×
Rerank Domain	Generic	–	1.0×	R2L	L2R	1.0×
TWIST Domain	Generic	–	1.4×	R2L	L2R	1.5×

Table 4: Inference time relative to a single model decoded in isolation. It is measured on the same single Nvidia A100-SXM GPU with batch size 1. We measure the wall-clock time from when the models are loaded until the last sentence is translated on the test data.

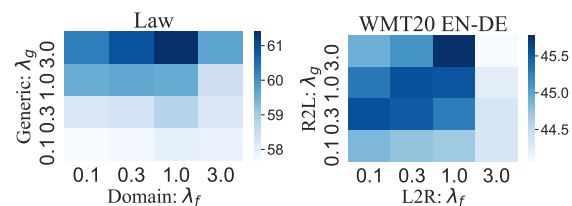


Figure 5: Dev. set performance measured in the COMET score (Rei et al., 2020a,b) with varying λ_f and λ_g . See Appendix §B for other configurations.

Sensitivity Analysis on Distance Coefficients

As discussed in §2.2, λ_f and λ_g weight the distance terms from f and g respectively. We tuned λ_f and λ_g on the dev. set from the range of $\{0.1, 0.3, 1.0, 3.0\}$. Fig. 5 visualizes how they affect the overall performance on the dev. sets. We

Dev. Set Results	Medicine		WMT20 ZH-EN	
	COMET	BLEU	COMET	BLEU
Original	92.8	58.1	36.5	26.4
One Candidate	93.2	58.2	35.4	25.8
Embed. Distance	92.8	57.9	36.5	26.2

Table 5: Variants of the distance function in TWIST decoding. f is the domain translator and g is the generic translator for medical translation (German-to-English). f is an R2L model and g is an L2R model for WMT20 Chinese \rightarrow English.

find that $\lambda_g > \lambda_f$ generally yields good performance, suggesting the effectiveness of the initial exploration by g with relatively weaker guidance from f .

Variants of Distance Functions We experiment with two variants of distance terms (Table 5): 1) *one candidate*, which measures the distance from the 1-best candidate from the other model (vs. minimization over multiple candidates; §2.2) and 2) *embed. distance*, which calculates the distance based on the Euclidean distance between the embeddings. Here the embeddings are taken from the output layer of the decoder. Overall, both two variants yield similar performance to the original distance function, but the *one candidate* method has a substantial performance drop on WMT20 ZH-EN. Note also that the *embed. distance* method necessitates additional distance computations between the token embeddings. This result illustrates that our original distance function is a simple yet effective design choice.

Examples Seen in Table 6 are example German \rightarrow English translations from the medical domain. The left section presents a case where the domain model translates the technical term, *Spätdyskinesie*, into the corresponding English term: **tardive dyskinesia**.¹¹ The generic model, on the other hand, generates a literal translation: **late dyskinesia**. In the right section, the domain model fails to handle the coordinate structure: **12.1% and 3.2% with aripiprazole** vs. **12.1% with aripiprazole and 3.2% with placebo**. In both cases, TWIST decoding benefits from the better translation.

¹¹https://en.wikipedia.org/wiki/Tardive_dyskinesia.

5 Further Related Work

Decoding from Multiple Models Much early work proposed methods to generate text from multiple models especially for machine translation (often called *consensus-based decoding*; Bangalore et al., 2001, 2002; Matusov et al., 2006; Rosti et al., 2007; Sim et al., 2007; Hildebrand and Vogel, 2008). Most of these methods limit their search space to n-best candidates from individual translation models (Li et al., 2009), contrasting with our TWIST decoding where one model can update its translation outputs under the guidance of another model. *Collaborative decoding* (Li et al., 2009) trains a separate feature-based scorer that measures the consensus between phrase-based Chinese-to-English translation models.

Alternatives to Left-to-Right Decoding We showed that TWIST decoding can be used to benefit from models with diverging generation order. Several prior works proposed approaches for generating text in a different fashion than the standard left-to-right order. For example, much recent work explored non-autogressive generation (Gu et al., 2018; Lee et al., 2018; Mansimov et al., 2019; Ghazvininejad et al., 2019; Kasai et al., 2020, *inter alia*) primarily to parallelize and speed up inference. More specifically, several works introduced training and/or inference algorithms that combine left-to-right and right-to-left models for machine translation (Zhou et al., 2019) and commonsense inference (Zaidi et al., 2020). Qin et al. (2020) incorporated right (future) context into a left-to-right language model by iterative gradient-based updates on the output representations. Those algorithms are designed specifically for the combination of left-to-right and right-to-left generation and cannot be easily extended to more general situations, such as diverging tokenization and vocabularies where TWIST decoding has been shown effective.

6 Conclusion

We presented TWIST decoding, a general inference algorithm that generates text from diverse models without the assumption of a shared vocabulary, tokenization, or generation order. Our method enables diverse models to guide each other, thereby outperforming individual models over various scenarios, even when one of the models is much weaker because of limited data. We also demonstrated that TWIST decoding can be viewed as a generalization





Medicine	Domain (Isolation) 	Generic (Isolation) 	Domain (Isolation) 	Generic (Isolation) 
Reference	If signs and symptoms of tardive dyskinesia appear in a patient on ABILIFY, dose reduction or discontinuation should be considered.		In placebo-controlled trials, the incidence of akathisia in bipolar patients was 12.1% with aripiprazole and 3.2% with placebo.	
Domain	If signs and symptoms of tardive dyskinesia appear in one patient on ABILIFY, a dose reduction or discontinuation should be considered.		In placebo-controlled trials, the incidence of akathisia in bipolar disorder was 12.1% and 3.2% with aripiprazole.	
Generic	If a patient treated with ABILIFY shows signs and symptoms of late dyskinesia, it should be considered to reduce the dose or stop treatment.		In placebo-controlled studies, the incidence of akathisia in bipolar patients was 12.1% with aripiprazole and 3.2% with placebo.	
TWIST <i>f</i> : Domain <i>g</i> : Generic	If signs and symptoms of tardive dyskinesia appear in one patient on ABILIFY, a dose reduction or discontinuation should be considered.		In placebo-controlled trials, the incidence of akathisia in bipolar patients was 12.1% with aripiprazole and 3.2% with placebo.	

Table 6: Example outputs from machine translation on the medical domain. For TWIST decoding, f is the domain model, and g is the generic model. In the left section, the generic model fails to capture technical terminology (late dyskinesia vs. tardive dyskinesia for the German term, *Spätdyskinesie*), and TWIST decoding chooses the correct term of tardive dyskinesia from the domain model. In the right example, on the other hand, the domain model has a problem in coordination (12.1% and 3.2% with aripiprazole vs. 12.1% with aripiprazole and 3.2% with placebo), and TWIST decoding successfully benefits from the accurate translation of the generic model.

and improvement of the commonly-adopted reranking heuristic. As it only requires a small change in code, we hope that researchers and practitioners of language generation will explore complementary strengths of diverse generation models through TWIST decoding.

Acknowledgements

We thank Hila Gonen, Phillip Keung, the ARK group at UW, and the Mosaic team at the Allen Institute for AI for their helpful feedback on this work.

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Appendices

A Hyperparameters and Settings

We provide training and implementation details for easy replication of our work.

A.1 Domain Machine Translation

We generally follow the preprocessing and subword tokenization from Koehn and Knowles (2017); Hu et al. (2019). Table 7 lists the hyperparameters and setting on `fairseq` that we use for all domain-specific translation models. All embeddings are shared (Press and Wolf, 2017; Inan et al., 2017). We choose the checkpoint that achieved the best loss on the validation data.

Hyperparameter	Value
label smoothing	0.1
# max tokens	8192
dropout rate	0.1
encoder embedding dim	512
encoder ffn dim	2048
# encoder attn heads	8
decoder embedding dim	512
decoder ffn dim	2048
# decoder attn heads	8
max source positions	1024
max target positions	1024
Adam lr	5×10^{-4}
Adam β_1	0.9
Adam β_2	0.98
lr-scheduler	inverse square
warm-up lr	1×10^{-7}
# warmup updates	4000
# max updates	600K
# GPUs	8
length penalty	0.6

Table 7: Domain translation `fairseq` hyperparameters and setting. We generally follow the base-sized configuration from Vaswani et al. (2017).

A.2 Left-to-Right and Right-to-Left

WMT20 ZH-EN Table 8 lists the hyperparameters and setting on `fairseq` that we use for left-to-right and right-to-left models on the WMT20 ZH-EN dataset. We generally follow the preprocessing and tokenization from Kasai et al. (2022a). We use `newstest-2019` as the dev. set and the official training data.¹² We apply Moses tokenization (Koehn et al., 2007) and BPE with 32K operations (Sennrich et al., 2016b) to English text. We tokenize Chinese text with the Jieba package,¹³

¹²<http://www.statmt.org/wmt20/translation-task.html>.

¹³<https://github.com/fxsjy/jieba>.

following Hassan et al. (2018). Separately from English, BPE with 32K operations is then applied to Chinese. The decoder input and output embeddings are tied.

WMT20 EN-DE The same hyperparameters are chosen as in WMT20 ZH-EN (Table 8). We again follow Kasai et al. (2022a) and preprocess both English and German text by the Moses tokenizer and *joint* BPE with 32K operations. All embeddings are shared.

Hyperparameter	Value
label smoothing	0.1
# max tokens	4096
dropout rate	0.1
encoder embedding dim	1024
encoder ffn dim	4096
# encoder attn heads	16
decoder embedding dim	1024
decoder ffn dim	4096
# decoder attn heads	16
max source positions	1024
max target positions	1024
Adam lr	5×10^{-4}
Adam β_1	0.9
Adam β_2	0.98
lr-scheduler	inverse square
warm-up lr	1×10^{-7}
# warmup updates	4000
# max updates	600K
# GPUs	8
length penalty	0.6

Table 8: L2R and R2L translation `fairseq` hyperparameters and setting. We generally follow the large-sized configuration from Vaswani et al. (2017).

A.3 SciTLDRS

We use two BART-based pretrained models from Cachola et al. (2020): the abstract-only version of BART and the AIC version of `CATTSXSUM`.¹⁴ These two models are both BART-based models; `CATTSXSUM` is obtained by finetuning BART on the XSUM dataset (Narayan et al., 2018) with multitask scaffolding (Cachola et al., 2020).

A.4 λ Tuning

We tune λ_f and λ_g from $\{0.1, 0.3, 1.0, 3.0\}$, based on the dev. BLEU/ROUGE-L score on machine translation and paper summarization, respectively. Table 9 reports the selected λ values in all scenarios.

¹⁴They are both available at <https://github.com/allenai/scitldr>.

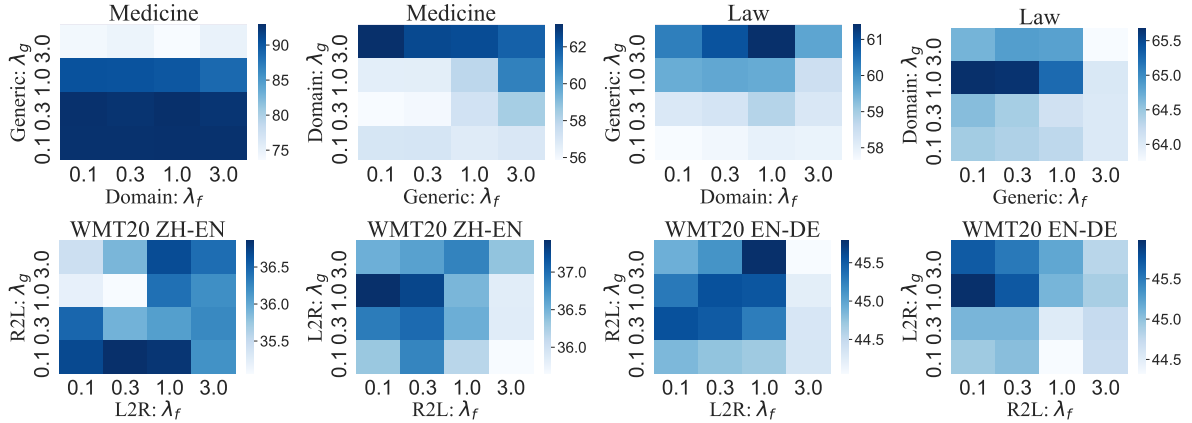


Figure 6: Dev. set performance measured in the COMET score (Rei et al., 2020a,b) with varying λ_f and λ_g .

Dataset	f	g	Tuned λ	
			λ_f	λ_g
Medicine	Domain	Generic	0.1	0.3
	Generic	Domain	0.1	3.0
Law	Domain	Generic	1.0	0.1
	Generic	Domain	0.1	3.0
Koran	Domain	Generic	1.0	3.0
	Generic	Domain	0.3	3.0
Subtitles	Domain	Generic	1.0	1.0
	Generic	Domain	1.0	1.0
WMT20 ZH-EN	L2R	R2L	1.0	3.0
	R2L	L2R	0.1	3.0
WMT20 EN-DE	L2R	R2L	0.3	0.3
	R2L	L2R	0.1	0.3
SciTLDR	Abst.	AIC	1.0	3.0
	AIC	Abst.	0.3	3.0

Table 9: Selected λ_f and λ_g values.

B Sensitivity Analysis on λ

Fig. 6 presents the sensitivity analysis in the COMET score over many scenarios. Apart from a few exceptions, $\lambda_g > \lambda_f$ tends to yield good performance, suggesting the effectiveness of the initial exploration by g with relatively weaker guidance from f .