GAATME: A Genetic Algorithm for Adversarial Translation Metrics Evaluation

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Abstract

Building on a recent method for decoding translation candidates from a Machine Translation (MT) model via a genetic algorithm, we modify it to generate adversarial translations to test and challenge MT evaluation metrics. The produced translations score very well in an arbitrary MT evaluation metric selected beforehand, despite containing serious, deliberately introduced errors. The method can be used to create adversarial test sets to analyze the biases and shortcomings of the metrics. We publish various such test sets for the Czech to English language pair, as well as the code to convert any parallel data into a similar adversarial test set.

1. Introduction

One of the crucial aspects of developing and deploying machine translation is automatic evaluation. The evaluation metrics introduced in recent years follow the trend of using pre-trained large language models as the core of a task-specific system. These novel metrics correlate better with human evaluation than the previous generation of metrics based on a rather shallow similarity of the proposed translation and human reference. However, many shortcomings, weaknesses and blind spots of these new metrics were already described in the literature, like insensitivity to errors in the translation of named entities, numbers and others (Hanna and Bojar, 2021; Amrhein and Sennrich, 2022).

In this paper, we modify a recently introduced genetic algorithm-based technique (Jon and Bojar, 2023) to automatically construct adversarial examples for specific metrics. Starting with an initial set of translation hypotheses generated by an MT model, we stochastically modify and combine them. The objective is to craft translations that excel in one specific metric but perform poorly across others. The main contribution of this paper is the release of metric-specific adversarial test sets and the accompanying code for creating new test sets, enabling researchers to probe various metrics' robustness, biases, and weaknesses. The code and the test sets can be found at: https://github.com/cepin19/GAATME

2. Related work

Many new MT evaluation metrics were introduced recently (Zhang et al., 2020; Yuan et al., 2021; Thompson and Post, 2020; Sellam et al., 2020; Rei et al., 2020, 2021, 2022b; Lo, 2019; Wan et al., 2021, 2022; Freitag et al., 2022; Rei et al., 2022a; Kocmi and Federmann, 2023; Guerreiro et al., 2023). They are based on representing the source, MT, and reference sentences in a (sometimes shared) high-dimensional space, computing the similarity between the representations, and (in most cases) predicting human evaluation scores. This allows more flexibility than traditional metrics based on shallow text similarities (e.g. for lexical overlap metrics like BLEU, synonyms vs. completely unrelated mistranslations are indistinguishable, while neural metrics should account for this by scoring synonyms similarly). Overall, they correlate better with human evaluation (Freitag et al., 2022; Kocmi et al., 2021). The downside of this increased flexibility is that the models are prone to be insensitive to some kinds of errors, especially in rare words and named entities, since such expressions often have similar embeddings.

Existing literature extensively probes the behavior and weaknesses of these contemporary metrics.

Moghe et al. (2023) show that neural metrics do not provide reliable results on the segment level. Amrhein and Sennrich (2022) try to find high-scoring incorrect translations, similar to our approach, to show that the analyzed metrics are not sensitive to errors in named entities and numbers. Alves et al. (2022) and Kanojia et al. (2021) further show that meaning-changing errors are hard to detect for QE. Rei et al. (2023); Leiter et al. (2022); Treviso et al. (2021); Guerreiro et al. (2023), and Fomicheva et al. (2021) explore the interpretability of the neural metrics.

Lastly, we directly build on Jon and Bojar (2023) and use a slight modification of this approach (using negative weights for "control" metrics) to build our adversarial test sets.

3. Method

We adapt the method presented by Jon and Bojar (2023). This approach is based on the genetic algorithm, which is described in the following paragraphs.



Figure 1: One iteration of the GA algorithm for a population of 4 individuals s1,..., s4. The steps with the yellow background are equivalent to simple reranking, the steps with the blue background introduce the operations of the genetic algorithm. Figure from Jon and Bojar (2023).

Genetic algorithm Our approach is the same as Jon and Bojar (2023) and we encourage the reader to find a more detailed description there. The whole process is illustrated in Figure 1.

First, a set of candidate sentences is produced by an NMT model, either by a beam search decoding or sampling for an increased diversity. These candidates are stochastically combined using a cross-over operation. This operation selects two individuals (i.e. translation hypotheses) from the population, splits them at a random token index swaps the split parts between the individuals. The resulting sequences are further modified using the mutation operation, which randomly removes, adds or replaces tokens in the candidate. The choices for new tokens to add or replace come from two sources: the complete wordlist in the target language and the set of words from a reference sentence.

Then, these candidate translations are scored using a fitness function, in our case a weighted sum of MT metrics' scores computed with regard to a known reference. This is a difference from Jon and Bojar (2023), who use MBR decoding with the translation candidates themselves as pseudoreferences. We use a positive weight for the metric we want to analyze (i.e., the one we want to find adversarial examples for) and a small negative weight for several other MT metrics. Our goal is to find sentences that perform exceptionally well according to the metric of interest but not as well according to other metrics. Finally, a new population of candidates is selected based on their scores via tournament selection. In most of the experiments by Jon and Bojar (2023), the authors use MBR decoding to obtain the scores, in order to improve the quality of the translation. In our experiments, we do not keep the reference secret, since we are looking to obtain adversarial examples for the MT metrics instead. Jon and Bojar (2023) also ran a similarly designed small-scale experiment but they only used the analyzed metric for the fitness function, without the negatively weighted "control" metrics. They searched for the "suspicious" final translations in the outputs after running the whole GA algorithm. We are encouraging GA to directly prefer the suspicious translation candidates, making our approach proactive in seeking out translations that may reveal weaknesses in

4. Experiments

4.1. Data, tools and model

MT metrics.

The NMT model was trained on CzEng 2.0 (Bojar et al., 2016; Kocmi et al., 2020) We obtained the English wordlist from https://github.com/ dwyl/english-words. We tokenize the text using SentencePiece (Kudo and Richardson, 2018) and FactoredSegmenter¹ for the training. For the tokenization in the GA process, we use SacreMoses.² We used the wmt22 (Kocmi et al., 2022) test set in Czech to English direction to create the adversarial translations. To produce the initial translations, we use the same model as Jon and Bojar (2023), i.e. transformer-big (Vaswani et al., 2017) using MarianNMT (Junczys-Dowmunt et al., 2018) with default hyperparameters.

4.2. GA parameters

The initial population of translation candidates is created by the NMT model described in Section 4.1. We concatenate n-best list obtained by beam search with beam size 20 and 20 sampled translations. We sample uniformly from the whole output distribution, as default in MarianNMT. We copy these 40 candidates 50 times to reach a population size of 2000. Empty token positions are added before and after each token in each candidate to support the addition of new words at these positions by mutating them to non-empty tokens. Finally, all the candidates are padded with empty token positions to the length of the longest candidate multiplied by 1.1.

The candidates are combined at a crossover rate of c = 0.1. The mutation rate for modifying nonempty genes (tokens) to other non-empty genes is $m = \frac{1}{l}$, with *l* representing chromosome length (i.e. number of positions in the translation candidate, including the empty token placeholders). Mutation rates between empty and non-empty genes (word addition/deletion) are $\frac{m}{10}$. Parents for the next generation are chosen via a tournament selection with n = 3. The GA stops after 150 generations. We combine the studied metric (the one we aim to find adversarial examples for) with other metrics in the fitness function by a weighted sum. We set the weight of the studied metric to 1.0 and we explore the following weights for all the other metrics: 0, -0.001, -0.01, -0.02, -0.03, -0.05, -0.1.We note that these settings are arbitrary, based on some previous experience. A search for better parameters could bring further improvements, but running the whole process is computationally costly. This is mainly due to a need for running a large number of evaluations by the MT metrics, many of which are deep learning-based and resource-intensive.

4.3. Metrics

We assess the translations using the metrics: BLEU (Papineni et al., 2002), ChrF (Popović, 2015), BLEURT-20 (Sellam et al., 2020), wmt20-comet-da (CMT20 in the tables), wmt22-comet-da (CMT22), wmt22-cometkiwi-da (CMT22-QE) (Rei et al., 2020, 2022a,c) and UniTE-MUP (Wan et al., 2022).

For both BLEU and ChrF metrics, Sacre-BLEU (Post, 2018) is used. Specifically, ChrF operates with a $\beta = 2$ setting, labeled ChrF2.

BLEURT is not used as the negative metric in any of the experiments, due to its 5x computational requirements compared to COMET. We only analyze it as the studied metric, with other metrics as the negative ones. We also do not use wmt20-cometda as part of the negatively weighted metrics, because we previously found that it does not correlate well with human quality assessment under these circumstances.

4.4. Results

The results from our various experimental runs are summarized in Table 1. The first column specifies the metric we are creating adversarial examples for. The second column details the negative metric weights. These "control" metrics guide the GA to produce translations with errors. The following columns provide system-level scores of the adversarial translations produced. A translation is identified as adversarial if its post-GA score in the examined metric rises, while the translation manifests serious translation mistakes introduced by the GA process, as we manuall annotated, see below. The first row shows the results of the baseline MT model that was used to create the initial population of translation candidates for the GA.

We can infer some notions about the robustness of the particular metrics based on this table. By comparing the targeted metric's score with other (control) metrics' scores, we can get a gist of its resilience against adversarial inputs. If a metric can be tricked using our method, its post-GA score should remain high, whereas scores from other metrics should decrease significantly. For instance, when optimized for BLEU or CMT22-QE, we observe a decline in most other metrics compared to their baseline, even without negative weights in the fitness function. In other words, BLEU and CMT22-QE are very susceptible to overfitting towards them. Conversely, optimization for UniTE or CMT22 enhances scores in many other metrics, indicating the robustness of UniTE and CMT22. This kind of analysis assumes there are no spurious correlations or shared blind spots between the metrics - this assumption is however certainly violated in practice, since the neural metrics share large parts of the architecture and training data.

To address this, we manually examined a selection of translations to determine the true ratio of adversarial samples. We evaluated 50 samples from each metric with negative weights of 0 and -0.1, labeling them adversarial if they presented significant translation errors (consistent samples were

¹https://github.com/microsoft/

factored-segmenter

²https://github.com/alvations/ sacremoses

Adv metric	W_{neg}	ChrF	BLEU	CMT20	CMT22	CMT22-QE	BLEURT	UniTE	% better	% adv
Baseline MT		64.1	39.9	0.434	0.794	0.751	0.671	0.123		
	0	84.8	58.2	-0.220	0.634	0.508	0.543	-0.433	100	78 (78)
	0.001	85.2	58.6	-0.240	0.635	0.511	0.539	-0.427		
	0.01	85.0	56.4	-0.433	0.589	0.472	0.495	-0.668		
chrF	0.02	84.8	55.1	-0.640	0.544	0.435	0.444	-0.829		
	0.03	84.4	52.1	-0.815	0.517	0.415	0.408	-0.991		
	0.05	82.6	45.0	-1.078	0.460	0.381	0.363	-1.193		
	0.1	79.9	33.6	-1.257	0.406	0.341	0.353	-1.374	96	96 (100)
	0	74.8	63.4	0.122	0.730	0.607	0.588	-0.157	100	70 (70)
	0.001	71.4	63.0	-0.535	0.570	0.485	0.445	-0.789		
	0.01	69.2	62.5	-0.907	0.477	0.428	0.389	-1.099		
BLEU	0.02	68.7	62.7	-0.972	0.452	0.404	0.362	-1.149		
	0.03	67.5	62.1	-1.022	0.438	0.397	0.356	-1.187		
	0.05	65.7	61.3	-1.186	0.401	0.370	0.321	-1.321		
	0.1	61.8	59.2	-1.292	0.363	0.328	0.292	-1.427	98	98 (100)
	0	70.4	49.6	0.803	0.851	0.739	0.727	0.401	100	22 (22)
	0.001	69.9	49.0	0.800	0.850	0.733	0.719	0.357		
	0.01	70.7	49.7	0.801	0.849	0.723	0.701	0.299		
CMT20*	0.02	69.1	48.2	0.799	0.843	0.721	0.692	0.239		
	0.03	67.8	45.0	0.794	0.839	0.698	0.680	0.139		
	0.05	64.1	37.7	0.772	0.820	0.673	0.641	-0.016		
	0.1	55.5	24.0	0.716	0.779	0.599	0.561	-0.449	82	78 (96)
	0	69.4	48.4	0.667	0.879	0.742	0.710	0.311	100	26 (26)
	0.001	69.5	49.3	0.649	0.879	0.739	0.715	0.294		
	0.01	64.6	40.4	0.580	0.876	0.715	0.681	0.069		
CMT22	0.02	59.3	29.9	0.471	0.867	0.658	0.621	-0.225		
	0.03	53.3	22.6	0.259	0.854	0.612	0.569	-0.532		
	0.05	45.8	12.3	-0.071	0.828	0.535	0.490	-0.908		
	0.1	35.5	2.3	-0.518	0.788	0.449	0.401	-1.194	38	38 (100)
	0	61.2	35.3	0.400	0.809	0.824	0.674	0.080	100	20 (20)
	0.001	61.5	35.2	0.404	0.807	0.822	0.667	0.098		
	0.01	56.9	28.2	0.220	0.775	0.819	0.629	-0.157		
CMT22-QE	0.02	50.5	17.8	-0.217	0.711	0.810	0.551	-0.526		
	0.03	46.9	12.4	-0.461	0.670	0.801	0.510	-0.754		
	0.05	40.6	6.9	-0.770	0.601	0.783	0.449	-1.038		
	0.1	32.7	2.3	-1.079	0.515	0.752	0.402	-1.236	30	30 (100)
	0	65.1	40.8	0.048	0.721	0.611	0.822	-0.241	100	78 (78)
	0.001	65.0	40.4	0.076	0.732	0.638	0.819	-0.211		
	0.01	61.1	35.1	-0.374	0.633	0.515	0.819	-0.612		
BLEURI*	0.02	58.9	28.3	-0.665	0.567	0.459	0.817	-0.855		
	0.03	53.2	20.3	-0.814	0.531	0.429	0.809	-0.985		
	0.05	49.0	15.8	-1.008	0.480	0.401	0.806	-1.143		
	0.1	36.4	4.9	-1.275	0.406	0.348	0.833	-1.359	76	/6 (100)
	0	68.4	45.4	0.591	0.817	0.726	0.707	0.628	100	22 (22)
	0.001	68.3	44.8	0.555	0.808	0.719	0.707	0.622		
U	0.01	67.5	45.1	0.588	0.810	0./1/	0.706	0.643		
UNITE	0.02	67.8	45.1	0.609	0.821	0.723	0./15	0.636		
	0.03	67.3	43.5	0.548	0.808	0.723	0.702	0.622		
	0.05	66.3	41.5	0.544	0.804	0.705	0.692	0.615	100	
	0.1	62.7	33.9	0.471	0.783	0.687	0.665	0.610	100	44 (44)

Table 1: Average scores of the generated test sets. Metrics marked with * were not used in the negative component of the fitness function for analyzing the other metrics. Scores in analyzed metric (the one we are searching adversarial examples for) are bold. ChrF and BLEU scores are multiplied by 100, in the algorithm they are in the 0-1 range. Higher is better for all the metrics.

used across all settings). Significant errors are defined as omissions, misinterpretations, additions not related to the source, redundant repetitions, or severe word-order errors. Our analysis is summarized in the last two columns. The "% better" column displays cases where the post-GA metric score surpasses its pre-GA value. The final column highlights instances that meet the previous criterion but also introduce a significant error via the GA. The presence of these errors was manually assessed. The numbers in parentheses show the total percentages of examples that contain newly introduced errors, regardless of whether the GA has improved the score or not.

4.5. Examples

Examples from our final test sets are presented in Table 2. Each row of the final test set displays the name of the analyzed metric, the source sentence (which is omitted in the table for conciseness), the machine translation (the first translation from the n-best list used as the initial population), the best translation post-GA, as well as the pre-GA and post-GA scores for the analyzed metric. This table offers insights into common errors associated with specific metrics. For instance, BLEURT appears to favor unfamiliar words or terms from other languages. These words were part of the noise in the English wordlist. We were unaware of their presence in the

Metric	МТ	post-GA	Ref	MT score GA score
CMT22-QE	In the NHL, "France" caught 36 games, its save rate at 92.3%. The 31-year-old full-back will be on	In yn, "Frederic" clocked up 36 or- dain, with touchdown rate at 92.3 Basilica The fullback will toilette on rota-	He has played 36 games in the NHL, where his save percentage is 92.3%. The thirty-one-year-old Pilsen na	e 0.6407 0.7679 e 0.6901 0.7242
	the scoresheet and could soon be in goal.	tion and could get into goal soon fungo	soon be in goal.	1
	The highest ranked in the affair is Berbr , who no longer features in any of the football functions.	The highest profile in glave affair is Piute Denten who no longer longer figures stanno any football func- tions	The most senior figure in the affai is Berbr, who is no longer involved in any football function.	r 0.7947 0.8104 J
CMT22	Prince William, Duke of Cam- bridge, is wearing the same as Princes George and Louis shorts and a collared T-shirt.	Prince Pippo, Duke of Gold- wyn , dressed the same as Princes Alexander and Louis in shorts and a T-shirt	Prince William, Duke of Cam bridge, and Princes George and Louis are wearing shorts and a polo shirt.	- 0.7675 0.8402 1 a
	Interior has got respirators signif- icantly cheaper than the Depart- ment of Health	Interior got respirators mushy Asch cheaper than the Ministry oie Ministry of Health natl. fur. LADT Goethe	The Ministry of the Interior got res pirators much cheaper than the Ministry of Health	- 0.5342 0.8168 9
CHRF	PVO: medium-cold war, outdated; short range - good, modern, rela- tively good number.	unpaint: medium-cold war, ob- solete; short range - good, mod- ern, relatively Orth . enterable number. fugitively favorer POS SMDF R.A.A.F. pm . SM	SHORAD: medium - cold war, ob solete; short range - good, moderr relatively favorable number.	- 67.9 82.0 I,
BLEU	The picture, which will serve as a Christmas card, was also posted by heir to the throne Prince Charles and wife Camilla.	knotty-leaved, which will be fat-shunning weeny-bopper Christmas card, was also posted by the heir to the throne, Prince Charles, dichlorodiphenyl- trichloroethane duodenochole- cystostomy cock-a-doodle-doos his wife, Camilla.	The image, which will be used for the Christmas card, was also posted by the heir to the throne Prince Charles, and his wife Camilla.	d 34.3 68.6
	By the time I got off my seat, it was gone.	Idun epicanthi got achenium tundun terebinthial off Ladakhi Morgenthaler gone ecliptically scholium mesonasal	By the time I got off the deer-stand he was gone.	l, 0.3965 0.8183
BLEURT	His return to goal in the NHL eventually extended to more than two months.	succinimid Badajoz hootchie- kootchie cheongsam NHL tao- tai meromyarian Abyla Nadean vainer tenson months	In the end, his time away from the NHL was extended by more than two months.	e 0.5695 0.8510 1

Table 2: Examples from the adversarial test set. Superfluous words in the post-GA translation and words from before GA that are missing post-GA are in bold.

wordlist before running the experiments.

5. Conclusion

We have presented a method to automatically generate adversarial test sets for arbitrary evaluation metrics. Our results demonstrate that this method is capable of producing translations that, while scoring higher than initial (correct or at least reasonable) MT outputs, contain serious translation errors. We have found that robustness against this method varies between metrics, with wmt22comet-da and UniTE being particularly robust, while BLEURT (alongside BLEU and CHRF) can be surprisingly easy to deceive. We publish the code and the created test sets to allow further use of this method.

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