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► **To cite this version:**

David Langlois, Sylvain Raybaud, Kamel Smaïli. LORIA System for the WMT12 Quality Estimation Shared Task. NAACL 2012 - The Seventh Workshop on Statistical Machine Translation, Jun 2012, Montréal, Canada. pp.114–119. hal-00726372

HAL Id: hal-00726372

<https://inria.hal.science/hal-00726372>

Submitted on 15 Nov 2017

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LORIA System for the WMT12 Quality Estimation Shared Task

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Abstract

In this paper we present the system we submitted to the WMT12 shared task on Quality Estimation. Each translated sentence is given a score between 1 and 5. The score is obtained using several numerical or boolean features calculated according to the source and target sentences. We perform a linear regression of the feature space against scores in the range [1:5]. To this end, we use a Support Vector Machine. We experiment with two kernels: linear and radial basis function. In our submission we use the features from the shared task baseline system and our own features. This leads to 66 features. To deal with this large number of features, we propose an in-house feature selection algorithm. Our results show that a lot of information is already present in baseline features, and that our feature selection algorithm discards features which are linearly correlated.

1 Introduction

Machine translation systems are not reliable enough to be used directly. They can only be used to grasp the general meaning of texts or help human translators. Confidence measures detect erroneous words or sentences. Such information could be useful for users to decide whether or not to post-edit translated sentences (Specia, 2011; Specia et al., 2010) or select documents mostly correctly translated (Soricut and Echiabi, 2010). Moreover, it is possible to use confidence measures to compare outputs from different systems and to recommend the best one (He et al., 2010). One can also imagine that confidence

measures at word-level could be also useful for a machine translation system to automatically correct parts of output: for example, a translation system translates the source sentence, then, this output is translated with another translation system (Simard et al., 2007). This last step could be driven by confidence measures.

In previous works (Raybaud et al., 2011; Raybaud et al., 2009a; Raybaud et al., 2009b) we used state-of-the-art features to predict the quality of a translation at sentence- and word-level. Moreover, we proposed our own features based on previous works on cross-lingual triggers (Lavecchia et al., 2008; Latiri et al., 2011). We evaluated our work in terms of Discrimination Error Trade-off, Equal Error Rate and Normalised Mutual Information.

In this article, we compare the features used in the shared task baseline system and our own features. This leads to 66 features which will be detailed in sections 3 and 4. We therefore deal with many features. We used a machine learning approach to perform regression of the feature space against scores given by humans. Machine learning algorithms may not efficiently deal with high dimensional spaces. Moreover, some features may be less discriminant descriptors and then in some cases could add more noise than information. That is why, in this article we propose an in-house feature selection algorithm to remove useless features.

The article is structured as follows. In Section 2, we give an overview of our quality estimation system. Then, in Sections 3 and 4, we describe the features we experimented with. In section 6, we describe the algorithm we propose for feature selec-

tion. Then we give the results of several configurations in Section 7.

2 Overview of our quality estimation submission

Each translated sentence is assigned a score between 1 and 5. 5 means that the machine translation output is perfectly clear and intelligible and 1 means that it is incomprehensible. The score is calculated using several numerical or boolean features extracted according to the source and target sentences. We perform a regression of the feature space against [1 : 5].

3 The baseline features

The quality estimation shared task organizers provided a baseline system including several interesting features. Among them, several are yet used in (Raybaud et al., 2011) but we give below a brief review of the whole baseline features set¹:

- Source and target sentences lengths: there is a correlation between the sizes of source and target sentences.
- Average source token length: this is the average number of letters of the words in the sentence. We guess that this feature can be useful because short words have more chance to be tool words.
- Language model likelihood of source and target sentences: a source sentence with low likelihood is certainly far from training corpus statistics. There is a risk it is badly translated. A target sentence with low likelihood is not suitable in terms of target language.
- Average number of occurrences of the words within the target sentence: too many occurrences of the same word in the target sentence may indicate a bad translation.
- Average number of translations per source word in the sentence: for each word in the source sentence, the feature indicates how many words of the target sentence are indeed translations of this word in the IBM1 table (with probability higher than 0.2).

¹Indeed, our system takes into input a set of features, and is able to discard redundant features (see Section 6).

- Weighted average number of translations per source word in the sentence: this feature is similar to the previous one, but a frequent word is given a low weight in the averaging.
- n-gram frequency based features: the baseline system proposes to group the n-gram frequencies into 4 quartiles. The features indicate how many n-gram (unigram to trigram) in source sentence are in quartiles 1 and 4. These features indicate if the source sentence contains *n*-grams relevant to the training corpus.
- Punctuation based features: there may exist a correlation between punctuation of source and target sentences. The count of punctuation marks in both sentences may then be useful.

Overall, the baseline system proposes 17 features.

4 The LORIA features

In a previous work (Raybaud et al., 2011), we tested several confidence measures. The Quality Measure Task campaign constitutes a good opportunity for us to compare our approach to others. We give below a brief review of our features (we cite again features which are yet presented in baseline features because sometimes, we use a variant of them):

- lengths: three features are generated, lengths of source and target sentences (already presented in baseline features), and ratio of target over source length
- *n*-gram based features (Duchateau et al., 2002): each word in the source and target sentences is given its 5-gram probability. Then, the sentence-level score is the average of the scores across all words in the sentence. There are 4 features: one for each language (source and target) and one for each direction (left-to-right and right-to-left 5-gram).
- backoff *n*-gram based features: in the same way, a score is assigned to a word according to how many times the language model had to back off in order to assign a probability to the sequence (Uhrík and Ward, 1997). Here too, word scores are averaged and we get 4 scores.

- averaged features: a common property of all n -gram based and backoff based features is that a word can get a low score if it is actually correct but its neighbours are wrong. To compensate for this phenomenon we took into account the average score of the neighbours of the word being considered. More precisely, for every relevant feature x , defined at word level we also computed:

$$\begin{aligned} x^{left}(w_i) &= x.(w_{i-2}) * x.(w_{i-1}) * x.(w_i) \\ x^{centred}(w_i) &= x.(w_{i-1}) * x.(w_i) * x.(w_{i+1}) \\ x^{right}(w_i) &= x.(w_i) * x.(w_{i+1}) * x.(w_{i+2}) \end{aligned}$$

A sentence level feature is then calculated according to the average of each new "averaged feature".

- intra-lingual features: the intra-lingual score of a word in a sentence is the average of the mutual information between that word and the other words in that sentence. Mutual information is defined by:

$$I(w_1, w_2) = P(w_1, w_2) \times \log \left(\frac{P(w_1, w_2)}{P(w_1)P(w_2)} \right) \quad (1)$$

The intra-lingual score of a sentence is the average of the intra-lingual scores of the words in this sentence. There are two features, one for each language.

- cross-lingual features: the cross-lingual score of a word in a target sentence is the average of the mutual information between that word and the words in the source sentence. The cross-lingual score of a target sentence is the average of its constituents.
- IBM1 features: the score of the target sentence is the average translation probability provided by the IBM1 model.
- basic parser: this produces two scores, a binary flag indicating whether any bracketing inside the target sentence is correct, and one indicating if the sentence ends with an end of sentence symbol (period, colon, semi-colon, question/exclamation/quotation mark, comma, apostrophe, close parenthese)

- out-of-vocabulary: this generates two scores, the number of out-of-vocabulary words in the sentence, and the same one but normalized by the length of the sentence. These scores are used for both sides.

This leads to 49 features. A few ones are equivalent to or are strongly correlated to baseline ones. As we want to be able to integrate several sets of features without prior knowledge, our system is able to discard redundant features (see Section 6).

5 Regression

Our system predicts a score between 1 and 5 for each test sentence. For that, we used the training corpus to perform the linear regression of the input features against scores given by humans. We used SVM algorithm to perform this regression (LibSVM toolkit (Chang and Lin, 2011)). We experimented two kernels: linear, and radial basis function. For the radial basis function, we used grid search to optimise parameters.

6 Feature Selection

We experimented with many features. Some of them may be very poor predictors. Then, these features may disturb the convergence of the training algorithm of SVM. To prevent this drawback, we applied an in-house feature selection algorithm. A feature selection algorithm selects the most relevant features by maximizing a criterion. Feature selection algorithms can be divided into two classes: backward and forward (Guyon and Elisseeff, 2003). Backward algorithms remove useless features from a set. Forward algorithms start with an empty feature set and insert useful features. We implemented a greedy backward elimination algorithm for feature selection. It discards features until a quality criterion stops to decrease. The criterion used is the Mean Average Error (MAE) calculated on the development corpus:

$$MAE(s, r) = \frac{\sum_{i=1}^n |s_i - r_i|}{n} \quad (2)$$

where s is the list of scores predicted by the system, r is the list of scores given by experts, n is the size of these lists.

The algorithm is described below:

Algorithm 1: Feature Selection algorithm

```
begin
  Start with a set  $S$  of features
  while two features in  $S$  are linearly
  correlated (more than 0.999) do
     $\perp$  discard one of them from  $S$ 
    Calculate MAE for  $S$ 
  repeat
    DecreaseMax  $\leftarrow$  0
    forall the feature  $f \in S$  do
       $S' \leftarrow S \setminus f$ 
      Calculate newMAE for  $S'$ 
      if MAE - newMAE >
      DecreaseMax then
        DecreaseMax  $\leftarrow$ 
        MAE - newMAE
         $f_{\text{chosen}} \leftarrow f$ 
      if DecreaseMax > 0 then
         $S \leftarrow S \setminus f_{\text{chosen}}$ 
        MAE  $\leftarrow$  MAE - DecreaseMax
  until DecreaseMax = 0;
```

For calculating the MAE for a feature set, several steps are necessary: performing the regression between the features and the expert scores on the training corpus, using this regression to predict the scores on the development corpus, calculate the MAE between the predicted scores and the expert scores on this development corpus.

7 Results

We used the data provided by the shared task on Quality Estimation², without additional corpus. This data is composed of a parallel English-Spanish training corpus. This corpus is made of the concatenation of europarl-v5 and news-commentary10 corpora (from WMT-2010), followed by tokenization, cleaning (sentences with more than 80 tokens removed) and truecasing. It has been used for baseline models provided in the baseline package by the shared task organizers. We used the same training corpus to train additional language models (forward and backward 5-gram with kneyser-ney discounting, obtained with the SRILM toolkit) and triggers required for our features. For feature extrac-

²<http://dl.dropbox.com/u/6447503/resources.tbz>

tion, we used the files provided by the organizers: 1832 source english sentences, their translations by the baseline translation system, and the score given by humans to these translations. We split these files into a training part (1000 sentences) and a development part (832 sentences). We used the train part to perform the regression between the features and the scores. We used the development corpus to optimise the parameters of the regression and for feature selection. We did not use additional provided information such as phrase alignment, word alignment, word graph, etc.

Table 1 presents our results in terms of MAE and Root Mean Squared Error (RMSE). MAE is described in Formula 2, and RMSE is defined by:

$$RMSE(s, r) = \sqrt{\frac{\sum_{i=1}^n (s_i - r_i)^2}{n}} \quad (3)$$

Each line of Table 1 gives the performance for a set of features. BASELINE+LORIA constitutes the union of both features BASELINE (Section 3) and LORIA (Section 4). the 'feature selection' column indicates if feature selection algorithm is applied. We experimented the SVM with two kernels: linear (LIN in Table 1) and radial basis function (RBF in Table 1). As the radial basis function uses parameters, we proposed results with default values (DEF) and with values optimised by grid search on the development corpus (OPT). MAE and RMSE are given for development corpus and for the test corpus. This test corpus (and its reference scores given by humans) is the one released for the shared 2012 task³. MAE and RMSE has been computed against the scores given by humans to the translations in this test corpus⁴.

The results show that the performance on development corpus are always confirmed by those of the test corpus. The BASELINE features alone achieve already good performance, better than ours. Although the differences are well inside the confidence interval, the fusion of both sets outperforms slightly the BASELINE. The feature selection algorithm allows to gain 0.01 point. The gain is the same for

³https://github.com/lspecia/QualityEstimation/blob/master/test_set.tar.gz

⁴available at https://github.com/lspecia/QualityEstimation/blob/master/test_set.likert

the optimisation of the radial basis function parameters. Surprisingly, the linear kernel, simpler than other kernels, yields the same performance as radial basis function.

In addition to MAE and RMSE results, we studied the linear correlations between features: our objective is to check if BASELINE and LORIA complement each other. We computed the linear correlation between all features (BASELINE+LORIA). This leads to 2145 values. Table 2 shows in line +/- the number of features pairs which correlate with an absolute score higher than thresholds 0.9, 0.8 or 0.7. Among these pairs we give in line + the number of pairs with positive correlation, and in line - the number of pairs with negative correlation. For lines + and -, we give 4 numbers: number of pairs, number of LORIA-LORIA (e.g. the number of correlations between a LORIA feature and another LORIA feature) pairs, number of BASELINE-BASELINE pairs, number of LORIA-BASELINE pairs. We remark that only 6% of the pairs correlates (column 0.7, line +/-) and that the correlations are mostly between LORIA features. This last point is not surprising because there are more LORIA features than BASELINE ones. There are very few correlations between LORIA and BASELINE features. We studied precisely the correlated pairs. There is a strong (more than 0.9) positive correlation between n -gram and backoff based features and their averaged feature versions. Sometimes, there is also a strong correlation between 'forward' and 'backward' features. Source and target sentences lengths linearly correlate (0.98). This is the same case for source and target language model likelihoods. There is also a high correlation between forward and backward 5-gram scores (0.89). There are very few negative correlations between features. As they are not numerous, one can list these pairs with correlation between -1 and -0.7: target sentence length and target language model probability; source sentence length and source language model probability; ratio of OOV words over sentence length in source sentence and percentage of unigrams in the source sentence seen in the SMT training corpus; and number of OOV words in source sentence and percentage of unigrams in the source sentence seen in the SMT training corpus. These correlations are not surprising. First, language model probability is not normalized

	≥ 0.9	≥ 0.8	≥ 0.7
+/-	64	103	127
+	56/49/3/4	94/87/3/4	117/105/6/6
-	8/0/4/4	9/0/4/5	10/0/4/6

Table 2: Statistics on the linear correlations between LORIA+BASELINE features

by the number of tokens: the more tokens, the lower probability. Second, the more OOV in the sentence, the fewer known unigrams.

Last, we present the set of features discarded by our feature selection algorithm. We give only this description for the LORIA+BASELINE set, with linear kernel. The algorithm discards 18 LORIA features out of 49 (37%) and 3 BASELINE out of 17 (18%). The features discarded from LORIA are mostly averaged features based on n -gram and back-off. This is consistent with the fact that these features are strongly correlated with n -gram and back-off features. We remark that very few BASELINE features are discarded: lengths of source and target language because these features are yet included in LORIA features, and "average number of translations per source word in the sentence" maybe because the LORIA feature giving the average IBM1 probabilities is more precise. Last, we remark that the target length feature is discarded, and only ratio between target and source length is kept.

8 Conclusion

In this paper, we present our system to evaluate the quality of machine translated sentences. A sentence is given a score between 1 and 5. This score is predicted using a machine learning approach. We use the training data provided by the organizers to perform the regression between numerical features calculated from source and target sentences and scores given by human experts. The features are the baseline ones provided by the organizers and our own features. We proposed a feature selection algorithm to discard useless features. Our results show that baseline features contain already the main part of information for prediction. Concerning our own features, a study of the linear correlations shows that averaged features do not provide new information compared to n -gram and backoff features. This last

Set of features	feature selection	kernel	Dev		Test	
			MAE	RMSE	MAE	RMSE
BASELINE	no	RBF DEF	0.63	0.79	0.69	0.83
LORIA	no	RBF DEF	0.66	0.82	0.73	0.87
BASELINE+LORIA	no	RBF DEF	0.62	0.78	0.69	0.82
BASELINE+LORIA	yes	RBF DEF	0.61	0.77	0.69	0.83
BASELINE+LORIA	no	RBF OPT	0.62	0.77	0.68	0.82
BASELINE+LORIA	no	LIN	0.62	0.78	0.69	0.83
BASELINE+LORIA	yes	LIN	0.61	0.77	0.68	0.82

Table 1: Results of the various sets of features in terms of MAE and RMSE

remark is confirmed by our feature selection algorithm. Our feature selection algorithm seems to discard features linearly correlated with others while keeping relevant features for prediction. Last, we remark that the choice of kernel, optimisation of parameters and feature selection have not a strong effect on performance. The main effort may have to be concentrated on features in the future.

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