# A Bayesian non-Linear Method for Feature Selection in Machine Translation Quality Estimation

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Abstract We perform a systematic analysis on the effectiveness of features for the problem of predicting the quality of machine translation at the sentencelevel. Starting from a comprehensive feature set, we apply a technique based on Gaussian Processes, a Bayesian non-linear learning method, to automatically identify features leading to accurate model performance. We consider application to several datasets across different language pairs and text domains, with translations produced by various machine translation systems and scored for quality according to different evaluation criteria. We show that selecting features with this technique leads to significantly better performance in most datasets, as compared to using the complete feature sets or a state-of-theart feature selection approach. In addition, we identify a small set of features which seem to perform well across most datasets.

## **1** Introduction

Machine Translation (MT) systems have been increasingly adopted in recent years for different purposes, including gisting and aiding humans to produce professional quality translations. Since the quality of automatic translations tends to vary significantly across text segments, methods to predict translation quality become more and more relevant. This problem is referred to as Quality

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Estimation (QE). Different from standard MT evaluation metrics, QE metrics do not have access to reference (human) translations; they are aimed at MT systems in use. Applications of QE include:

- Decide which segments need revision by a human translator;
- Decide whether a reader gets a reliable gist of the text;
- Estimate how much effort will be needed to post-edit a segment;
- Select among alternative translations produced by different MT systems.

Work in QE started with the goal of estimating automatic metrics such as BLEU [14] and WER [3]. However, these metrics are difficult to interpret, particularly at the sentence-level, added to which automatic estimation proved highly inaccurate. A new surge of interest in the field started recently, motivated by the widespread use of MT systems in the translation industry, as a consequence of better translation quality, more user-friendly tools, and higher demand for translation. In order to make MT maximally useful in this scenario, a quantification of the quality of translated segments similar to "fuzzy match scores" from translation memory systems is needed.<sup>1</sup> QE work addresses this problem by using more complex metrics that go beyond matching the source segment against previously translated data. QE can also be useful for endusers reading translations for gisting, particularly those who cannot read the source language. Recent work focuses on estimating more interpretable metrics, where "quality" is defined according to the task at hand: post-editing, gisting, etc. A number of positive results have been reported (Section 2).

QE is generally addressed as a supervised machine learning task using algorithms to induce models from examples of translations described through a number of **features** and annotated for quality. One of most challenging aspects of the task is the design of feature extractors to capture relevant aspects of quality.

A wide range of features from source and translation texts and external resources and tools have been used. These go from simple, language-independent features, to advanced, linguistically motivated features. They include features that rely on information from the MT system that generated the translations, and features that are oblivious to the way translations were produced. This variety of features plays a key role in QE, but it also introduces a few challenges. Datasets for QE are usually small because of the cost of human annotation. Therefore, large feature sets bring sparsity issues. In addition, some of these features are more costly to extract as they depend on external resources or require time-consuming computations. Finally, it is generally believed that different datasets (i.e. language pair, MT system or specific quality annotation such as post-editing time vs translation adequacy) can benefit from different features.

**Feature selection** techniques can help not only select the best features for a given dataset, but also understand which features are in general effective.

 $<sup>^1</sup>$  A fuzzy match score represents the percentage of common words between a segment to translate and segments previously translated in a database, and thus for which a correct translation is available and can be used directly.

While recent work has exploited selection techniques to some extent, the focus has been on improving QE performance on individual datasets (Section 2). As a result, no general conclusions can be made about the effectiveness of features across language pairs, text domains, MT systems and quality labels.

In this paper we propose to use Gaussian Processes for feature selection, a technique that has proven effective in ranking features according to their discriminative power [27]. We benchmark with this technique on two settings: (i) 13 datasets for four language pairs, various Statistical MT (SMT) and rulebased MT (RBMT) systems and four types of quality scores with the same feature sets; (i) one dataset (same language pair and quality scores) with seven feature sets produced in a completely independent fashion (by participants in a shared task on the topic) (Section 4). The experiments showed the potential of feature selection to improve overall regression results, often outperforming published results even on feature sets that had already been previously selected using other methods. They also allowed us to identify a small number of wellperforming features across datasets (Section 5). We discuss the feasibility of extracting these features based on their dependence on external resources or specific languages.

## 2 Related work

Examples of successful cases of QE include improving post-editing efficiency by filtering out low quality segments which would require more effort or time to correct than translating from scratch [28,24], selecting high quality segments to be published as they are, without post-editing [23], selecting a translation from either an MT system or a translation memory for post-editing [10], selecting the best translation from multiple MT systems [26], and highlighting subsegments that need revision [2]. For an overview of various algorithms and features we refer the reader to the WMT12-13 shared tasks on QE [6,4].

Most previous work on QE use machine learning algorithms such as Support Vector Machines (SVM), which are fairly robust to redundant/noisy features, and therefore feature selection is often neglected. Work using explicit feature selection methods rely mostly on forward/backward selection approaches, but the order in which features are added/removed is not informed by any prior knowledge. In what follows we summarise recent work using explicit feature selection methods.

[8] performed feature selection on a set of 475 sentence- and sub-sentence level features. Principal Component Analysis and a greedy selection algorithm to iteratively create subsets of increasing size with the best-scoring individual features were exploited. Both selection methods yielded better performance than all features, with greedy selection achieving the best MAE scores with 254 features.

[11] reported positive results with a greedy backward selection algorithm that removes 21 poor features from an initial set of 66 features based on error minimisation on a development set. In an oracle-like experiment, [7] use a sequential forward selection method, which starts from an empty set and adds one feature at a time as long as it decreases the model's error, evaluating the performance of the feature subsets on the test set directly. 37 features out of 147 are selected, and these significantly improved the overall performance.

[1] tested a few feature selection methods using both greedy stepwise and best first search to select among their 266 features with 10-fold cross-validation on the training set. These resulted in sets of 30-80 features, all outperforming the complete feature set. Correlation-based selection with best first search strategy was reported to perform the best. Conversely, [13] reported no improvements in performance in experiments with several selection methods.

Finally, [22], the winning system in the WMT12 QE shared task, used a computationally-intensive method on a development set. For each of the official evaluation metrics (e.g. MAE), from an initial set of 24 features, all  $2^{24}$ possible combinations were tested, followed by an exhaustive search to find the best combinations. The 15 features belonging to most of the top combinations were selected. Other rounds were added to deal with POS features, but the final feature sets included 14-15 features depending on the evaluation metric. This technique outperformed the complete feature set by a large margin.

### **3** Gaussian Processes

Gaussian Processes (GPs; [18]) are an advanced machine learning framework incorporating Bayesian non-parametrics and kernels, and are widely regarded as state of the art for many regression tasks. Despite that, GPs have been under-exploited for language applications. Most of the previous work on QE uses kernel-based Support Vector Machines for regression (SVR), based on experimental findings that non-linear models significantly outperform linear models. This is perhaps unsuprising, given the relatively small numbers of input features used and the complexity of the response variable.

There is little reason to expect that measures of quality, such as postediting effort, will be a linear function of the input features. To illustrate, consider how quality varies with the length of the source sentence, one of the most important features in our arsenal. Most short inputs are easy to translate, and therefore we expect the translation quality to be high. In contrast medium length sentences can be much more syntactically complex, leading to much worse translations. However we do not expect that very long sentences are much worse again: in fact they may be simpler, as these sentences are often lists or other highly structured sentences which can be more easily translated. Consequently, positing a linear relationship between input length and quality is not a reasonable proposition. For this feature and many other important features, a non-linear approach is more appropriate. Fig 1 shows three examples of features known to perform very well for translation quality prediction (source sentence length in 1(a), source sentence language model score in 1(b), and target sentence language model score in 1(c)) and their relationship with



Fig. 1 Features known to perform well for translation quality prediction (source sentence length in 1(a), source sentence language model score in 1(b), and target sentence language model score in 1(c)) and their relationship with post-editing distance – HTER, with a fitted GP to highlight the non linearity of the data.



Fig. 2 The Bayesian posterior under a GP prior, illustrated on synthetic one-dimensional data. Figs a and b show samples from the posterior (curves) over functions given several training observations (blue dots). These differ in the values of the hyperparameters, Fig a uses  $\sigma_f^2 = \sigma_n^2 = l = 1$ , while Fig b has learned the MLE hyper-parameters,  $\sigma_f^2 = 0.7$ ,  $\sigma_n^2 = 0.7$ ,

 $\pm$  one confidence interval.

post-editing distance – HTER [21], with a fitted GP to highlight the non linearity of the data. The same behaviour is observed with other features and other quality labels.

0.02, l = 0.3. Fig c shows the full posterior for the model in b, with the shaded area denoting

Like SVMs for regression – SVRs, GPs can describe non-linear functions using kernels such as the radial basis function (RBF). However in contrast, inference in GP regression can be expressed analytically and the kernel hyperparameters optimised directly using gradient descent. This avoids the need for costly cross validation, while also allowing the use of much richer kernel functions with many more parameters. Further differences between the two techniques are that GPs are probabilistic models and thus can be incorporated into larger graphical models. Moreover, GPs take a fully Bayesian approach by integrating out the model parameters to support posterior inference. Unlike most other Bayesian methods, GP regression supports exact posterior inference and learning (see Fig 2), without the need to resort to approximation (e.g., Markov Chain Monte Carlo sampling or variational approximations).



Fig. 3 Samples from a Gaussian Process prior with different length scales in an RBF kernel, showing the effect of this parameter on the smoothness of the function. We consider one dimensional input (D = 1), set  $\sigma_f = 0.7$  and omit the white-noise term.

Formulation We now present GP regression, following the formulation of Rasmussen and Williams [18, chapter 2]. GP regression assumes the presence of a latent function,  $f : \mathbb{R}^F \to \mathbb{R}$ , which maps from the input space of feature vectors **x** to a scalar. Each response value is then generated from the function evaluated at the corresponding data point,  $y_i = f(\mathbf{x}_i) + \eta$ , where  $\eta \sim \mathcal{N}(0, \sigma_n^2)$ is added white-noise. Formally f is drawn from a GP prior,

$$f(\mathbf{x}) \sim \mathcal{GP}(\mathbf{0}, k(\mathbf{x}, \mathbf{x}')),$$

which is parameterised by a mean (here, 0) and a covariance kernel function  $k(\mathbf{x}, \mathbf{x}')$ . The kernel function represents the covariance (i.e., similarities in the response) between pairs of data points. Several draws of f from a GP prior are illustrated in Fig 3.

*Kernel (covariance) function* **GPs** allow for many different kernels. Here we consider the RBF with automatic relevance determination,

$$k(\mathbf{x}, \mathbf{x}') = \sigma_f^2 \exp\left(-\frac{1}{2} \sum_{i}^{D} \frac{(x_i - x'_i)^2}{l_i}\right),$$

where the  $k(\mathbf{x}, \mathbf{x}')$  is the kernel function between two data points  $\mathbf{x}, \mathbf{x}'$ ; D is the number of features; and  $\sigma_f$  and  $l_i \geq 0$  are the kernel hyper-parameters which control the covariance magnitude and the *length scales* of variation in each dimension, respectively. This is closely related to the RBF kernel used with SVR, except that each feature is scaled independently from the others, i.e.,  $l_i = l$  for SVR, while **GPs** allow for a vector of independent values. The model hyper-parameters ( $\sigma_n, \sigma_f, \mathbf{l}$ ) are learned from data using a maximum likelihood estimation.

The learned length scale hyper-parameters can be interpreted as encoding the importance of a feature: the narrower the RBF (the smaller is  $l_i$ ) the more important a change in the feature value is to the model prediction. This is illustrated in Fig 3, which shows samples from a GP prior with different settings of the length scale: clearly the value of the input will be of less important for Fig 3(c), where the curves are mostly flat, versus Fig 3(a) which allows for rapid fluctuations. This effect can also be seen in the posterior, comparing Figs 2(a) and 2(b), where the latter's shorter length scale allows more accurate fitting of the training points.

Feature selection A model trained using GPs can be viewed as a list of features ranked by relevance, and this information can be used for feature selection by discarding the lowest ranked (least useful) features. GPs on their own do not provide a cut-off point on this ranked list of features, instead this needs to be determined in another way, e.g., by evaluating loss on a separate set to determine the optimal number of features. We experiment with variants on how to determine this cut off point in Section 4.3.

*Bayesian inference* Given the generative process defined above, prediction can be formulated as Bayesian inference under the posterior,

$$p(y_*|\mathbf{x}_*, \mathcal{D}) = \int_f p(y_*|\mathbf{x}_*, f) p(f|\mathcal{D}),$$

where  $\mathbf{x}_*$  is a test input and  $y_*$  is its response value. The posterior  $p(f|\mathcal{D})$  reflects our updated belief over possible functions after observing the training set  $\mathcal{D}$ , i.e., f should pass close to the response values for each training instance (but need not fit exactly due to additive noise). This is balanced against the biases (e.g., smoothness constraints) that arise from the GP prior. The posterior is illustrated in Fig 2 (a, b), which shows several samples of f from the posterior in (a,b), which are consistent with the observed training data,  $\mathcal{D}$ . The predictive posterior can be solved analytically, resulting in

$$y_* \sim \mathcal{N} \left( \mathbf{k}_*^T (K + \sigma_n^2 I)^{-1} \mathbf{y}, \ k(\mathbf{x}_*, \mathbf{x}_*) - \mathbf{k}_*^T (K + \sigma_n^2 I)^{-1} \mathbf{k}_* \right),$$
 (1)

where  $\mathbf{k}_* = [k(\mathbf{x}_*, \mathbf{x}_1) \ k(\mathbf{x}_*, \mathbf{x}_2) \cdots k(\mathbf{x}_*, \mathbf{x}_n)]^T$  are the kernel evaluations between the test point and the training set, and  $\{K_{ij} = k(\mathbf{x}_i, \mathbf{x}_j)\}$  is the kernel (gram) matrix over the training points. Note that the posterior in Eq. 1 includes not only the expected response (the mean) but also the variance, encoding the model's uncertainty, which is important for integration into subsequent processing, e.g., as part of a larger probabilistic model. The posterior is illustrated in figure 2(c). Note the increasing amounts of uncertainty in the centre of the graph (wider confidence interval) where there are few nearby training instances compared to the edges where there is dense training data. This nuanced modelling of uncertainty is of great importance when combining the model into a larger probabilistic graphical model, such that uncertainty can be preserved in task based inferences.

The remaining question is how to determine the kernel hyperparameters. As stated above, these modulate the effect and strength of the GP prior, including in our case the relative importance of each feature. The GP framework allows for kernel hyper-parameters to be learned using a maximum likelihood estimate (type II). The marginal likelihood,  $p(\mathbf{y}|X) = \int_f p(\mathbf{y}|X, f)p(f)$ , can be expressed analytically for GP regression (X are the training inputs), which

can be optimised with respect to the hyper-parameters for model selection. Specifically, we can derive the gradient of the (log) marginal likelihood with respect to the model hyperparameters (i.e.,  $\sigma_n, \sigma_f, \mathbf{l}$  etc.) and thereby find the type II maximum likelihood estimate using gradient ascent. Note that in general the marginal likelihood is non-convex in the hyperparameter values, and consequently the solutions may only be locally optimal. Here we bootstrap the learning of complex models with many hyperparameters by initialising with the (good) solutions found for simpler models, thereby avoiding poor local optima. We refer the reader to [18, chapters 2, 4–5] for further details on Gaussian process regression, kernel formulations and model selection in this framework.

#### 4 Experimental settings

In our experiments, model learning is performed with an open source implementation of  $GPs^2$  for regression. This is used for our proposed feature selection method. In what follows we describe two groups of quality estimation datasets: the first group contains various datasets for which we have extracted common feature sets using the QuEst framework [27,19] (Section 4.1); the second group contains a single dataset with various feature sets provided as part of the WMT12 quality estimation shared task (Section 4.2).

## 4.1 Datasets with QuEst features

The following datasets have a common feature set and are available for down-load.<sup>3</sup> The statistics of these datasets are shown in Table 1.

WMT12 English-Spanish news sentence translations produced by a phrasebased (PB) Moses "baseline" SMT system,<sup>4</sup> and judged for post-editing effort in 1–5 (highest-lowest), taking a weighted average of three annotators [6].

*EAMT11* English-Spanish (EAMT11-en-es) and French-English (EAMT11fr-en) news sentence translations produced by a PB-SMT Moses baseline system and judged for post-editing effort in 1–4 (highest-lowest) [24].

*EAMT09* English sentences from the European Parliament corpus translated by four SMT systems (two Moses-like PB-SMT systems and two fully discriminative training systems) into Spanish and scored for post-editing effort in 1-4(highest-lowest). Systems are denoted by  $s_1-s_4$  [28].

<sup>&</sup>lt;sup>2</sup> http://sheffieldml.github.io/GPy/

<sup>&</sup>lt;sup>3</sup> http://www.dcs.shef.ac.uk/~lucia/resources.html

<sup>&</sup>lt;sup>4</sup> http://www.statmt.org/moses/?n=Moses.Baseline

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Dataset	Languages	Training	Test	Label
WMT12	en-es	1,832	422	PE effort 1-5
EAMT11	en-es	900	64	PE effort 1-4
EAMT11	fr-en	2,300	225	PE effort 1-4
$EAMT09-s_1-s_4$	en-es	3,095	906	PE effort 1-4
$GALE11-s_1-s_2$	ar-en	2,198	387	Adequacy 1-4
LIG	fr-en	9,000	1,881	<b>HTER 0-1</b>
TRACE	en-fr	7,599	1,323	HTER 0-1
TRACE	fr-en	7,400	1,295	HTER 0-1
WMT13	en-es	2,254	500	HTER 0-1

Table 1 Number of sentences in our datasets. Figures for EAMT09- $s_1$ - $s_4$  and GALE11- $s_1$ - $s_2$  indicate number of sentences per MT system.

*GALE11* Arabic newswire sentences translated by two Moses-like PB-SMT systems into English and scored for adequacy in 1-4 (worst-best). Systems are denoted by  $s_1-s_2$  [25].

TRACE English-French (en-fr) and French-English (fr-en) sentence translations produced by two MT systems: a rule-based system (Reverso) and LMSI's statistical MT system [29]. English-French contains a mixture of data from Ted Talks, WMT news, SemEval-2 Cross-Lingual Word Sense Disambiguation, and translation requests from Softissimo's online translation portal (the Reverso system), which can be thought of as user-generated content. The French-English data contains sentences from the OWNI – a free French online newspaper, Ted Talks and translation requests from Softissimo's online translation portal. All translations have been post-edited and the HTER scores are used as quality labels. For each language pair, 1,000 translations have been post-edited by two translators independently. We simply concatenated these in our datasets.

*LIG* French-English sentence translations of various editions of WMT news test sets, produced by a customised version of a PB-SMT Moses system by the LIG group [16]. These sentences have been post-edited by a human translator, and labelled for HTER.

*WMT13* English-Spanish sentence translations of news texts produced by a PB-SMT Moses baseline system. These were then post-edited by a professional translator and labelled for post-editing effort using HTER. This is a superset of the WMT12 dataset, with 500 additional sentences for test, and a different quality label [4].

#### 4.1.1 Feature sets

The features for these datasets are extracted using the open source toolkit QuEst.<sup>5</sup> We differentiate between *black-box* (BB) and *glass-box* (GB) features,

<sup>&</sup>lt;sup>5</sup> http://www.quest.dcs.shef.ac.uk

as only BB are available for all datasets (we did not have access to all MT systems that produced the other datasets). For the WMT12 and GALE11 datasets, we experimented with both BB and GB features. The BB feature sets are the same for all datasets, except for the Arabic-English datasets, where language-specific features supplement the initial set of features, and for the WMT13 dataset, where advanced features dependent on external resources like parsers have been exploited.

We also distinguish one special feature: the *pseudo-reference* (PR), as this is not a standard feature in that its computation requires decoding with another MT system. This feature consists in translating the source sentence using another MT system (in our case, Google Translate) to obtain a *pseudo-reference*. The geometric mean of smoothed 1-to-4-gram precision scores<sup>6</sup> is then computed between the original MT and this pseudo-reference. We note that the better the external MT system, the closer the pseudo-reference translation is to a human translation, and therefore the more reliable this feature becomes.

For each dataset we built four QE systems, each with a feature set:

- Baseline (BL): 17 BB features that performed well across languages in previous work and were used as baseline in the WMT12-13 QE shared tasks.
  - number of tokens in the source & target sentences
  - average source token length
  - average number of occurrences of the target words within the target sentence
  - number of punctuation marks in source & target sentences
  - n-gram language model (LM) probability of source & target sentences using 3-gram LMs built from the source/target corpora
  - average number of translations per source word given by IBM 1 model thresholded such that P(t|s) > 0.2
  - same as above threshold such that P(t|s) > 0.01, where the number of translations for each source word is weighted by the inverse frequency of the word in the source language corpus
  - percentage of unigrams, bigrams and trigrams in frequency quartiles 1 (lower frequency words) and 4 (higher frequency words) in the source language corpus
  - percentage of unigrams in the source sentence seen in source language corpus
- All features (AF) available for the dataset. This is a superset of the above, where:
  - Common set of 80 BB features for all datasets, and additional 43 language specific features for the Arabic-English datasets, or additional advanced features for the WMT13 dataset.

 $<sup>^{6}</sup>$  This formulation is equivalent to sentence level BLEU without a brevity penalty, where the n-gram precision scores are smoothed: we add one to the numerator and denominator terms in order to avoid division by 0 errors.

- For the experiments with GB features we have all MT system-dependent features (varying according to the actual type of MT system) available for the GALE11-s<sub>1</sub> (39), GALE11-s<sub>2</sub> (48), and WMT12 (47) datasets.
   For a comprehensive list, we refer the reader to the QuEst project website.<sup>6</sup>
- **BL**+**PR**: 17 baseline features along with a pseudo reference feature.
- AF+PR: All features available (BB, GB or BB+GB) plus the pseudoreference feature.

#### 4.2 WMT12 datasets

We now turn to the second family of feature sets *i.e.*, for the WMT12 dataset described above. We use the feature sets provided by all but one of the participating teams in the WMT12 shared task on QE.<sup>7</sup> These very diverse feature sets, with features of many different natures, although some overlap with the QuEst features exists in most cases. We denote each of these feature set **AF**. We note however that in a few cases these are only a subset of the features actually used in the shared task, e.g. **UU**, since the participants could not provide us with the full feature sets. This explains the difference between the official scores reported in [6] and our figures. This difference can also be explained by the learning algorithms: while we used **GPs**, participants have used SVRs, M5P and other algorithms. Some of these feature sets already result from feature selection techniques.

*SDL* [22]: 15 features selected after an exhaustive search algorithm based on all possible combinations of features. This is the optimal set used by the winning submission. It includes many of the baseline features, the pseudo-reference feature, phrase table probabilities, and a few part-of-speech tag alignment features.

UU [9]: 82 features, a subset of those used in the shared-task as the parse tree features (based on tree-kernels) were not provided by the participants. These are similar to the common BL and BB features presented above and include various source and target LM features, average number of translations per source word, number of tokens matching certain patterns (hyphens, ellipsis, etc.), percentage of n-grams seen in corpus, percentage of non-aligned words, etc.

*UEdin* [5]: 56 black-box features including source translatability, named entities, LM back-off features, discriminative word-lexicon, edit distance between source sentence and the SMT source training corpus, and word-level features based on neural networks to select a subset of relevant words among all words in the corpus.

 $<sup>^7</sup>$  These feature sets were made available by the task organisers at <code>http://www.dcs.shef.ac.uk/~lucia/resources.html</code>

*Loria* [11]: 49 features including 1-5gram LM and back-off LM features, interlingual and cross-lingual mutual information features, IBM1 model average translation probability, punctuation checks, and out-of-vocabulary rate.

TCD [13]: 43 features based on the similarity between the (source or target) sentence and a reference set (the SMT training corpus or Google N-grams) with n-grams of different lengths, including the TF-IDF metric.

WLV-SHEF [7]: 147 features which are a superset of the common 80 blackbox features above. The additional features include a number of linguistically motivated features for source or target sentences (percentage) or their comparison (ratio), such as content words and function words, width and depth of constituency and dependency trees, nouns, verbs and pronouns.

UPC [15]: 56 features on top of the baseline features. Most of these features are based on different language models estimated on reference and automatic Spanish translations.

#### 4.3 Feature selection techniques

As previously described, our proposed technique for feature selection uses GPs. In each dataset, features are first normalised. Each feature is centred and scaled to have zero mean and unit standard deviation. For feature ranking, the models are trained on the full training sets, except in the FS(GP-dev) setting (below). The RBF widths, scale and noise variance are initialised by training first with an isotropic kernel (with a single length scale,  $l_i = l$ ), then untieing the length scales and continuing training. This helps to avoid local minima in the MLE. We apply a sparse approximation method for inference, known as Fully Independent Training Conditional [17], which bases parameter learning on a few inducing points in the training set instead of the entire training set. This approximation technique is used to make the learning less computationally expensive and therefore faster, which is important for large training sets. The hyper-parameters are learned using gradient descent with a maximum of 100 iterations. A forward selection approach is then used to select features ranked from top to worst and train models using GPs with increasing numbers of features and all available training data.

Four variant of selection techniques were used in our experiments, all applied on the entire set of features (AF+PR for the QuEst datasets, and AF for the WMT12 datasets):

- FS(GP-dev): Feature selection on AF+PR for automatic ranking and selection of top features with GPs on a development set and applied to test set.
- FS(GP-fixed): Feature selection on AF+PR for automatic ranking and selection of top features with GPs and a fixed number of features (threshold pre-defined).

- FS(GP-test): Feature selection on AF+PR for automatic ranking and selection of top features with GPs directly on test set (oracle selection).
- FS(RL): Feature selection with Randomised Lasso [12].

In FS(GP-dev), a development set is randomly selected from the training data for each of the dataset. In each of the datasets, we extracted the same number of sentences for development as there were for the test set. The development set was then used to choose the cutoff point in the ranked set of features generated by the GP model, i.e., where to stop selecting features. Once we performed feature selection over development set, the full training set was used to train the final models.

**FS(GP-fixed)** is based on pre-defined threshold on number of features to select for model training. The threshold was decided empirically based on previous experiments on various datasets. We observed that the optimal number of features oscillates between 10 and 30 features for different datasets. In these experiments, we selected 17 as fixed threshold for two reasons: it falls within this range and it allows for an interesting comparison with the hand engineered selection of 17 baseline features.

In FS(GP-test), the selected set is evaluated under an oracle condition, where the optimal number of features is decided based on the best performance obtained directly on the test set. This experiment aimed to study the upper bound in performance of the GPs-based method for feature selection. The subset of top ranked features that minimises error in each test set is selected.

As an alternative approach to GPs, we use Randomized Lasso, FS(RL), for feature selection. Randomised Lasso repeatedly resamples the training data and fits a Lasso regression model on each sample. A feature is selected for the final model if it is selected (i.e., assigned a non-zero weight) in at least 25% of the samples (we do this 1000 times). This strategy improves the robustness of Lasso in the presence of high dimensional and correlated inputs. It should be noted that FS(RL) automatically selects the number of features to be used for training. The final models are then trained using GPs with the selected features.<sup>8</sup>

#### 4.4 Evaluation metrics

To evaluate the prediction models we use Mean Absolute Error (MAE), its squared version – Root Mean Squared Error (RMSE), plus the Relative Absolute Error (RAE) and its squared version – Relative Squared Error (RSE). MAE is used as the main metric on the charts for a comparative analysis. RAE provides the average error relative to a simple predictor, which is just the average of the true values. We use it here to provide some form of comparison across prediction models for different datasets.

 $<sup>^{8}\,</sup>$  The GP trained on the selected features consistently outperforms the linear model learned by RL.

$$MAE = \frac{\sum_{i=1}^{N} |H(s_i) - V(s_i)|}{N} \qquad RMSE = \sqrt{\frac{\sum_{i=1}^{N} (H(s_i) - V(s_i))^2}{N}}$$
$$RAE = \frac{\sum_{i=1}^{n} |H(s_i) - V(s_i)|}{\sum_{i=1}^{n} |\bar{V}(s) - V(s_i)|} \qquad RSE = \sqrt{\frac{\sum_{i=1}^{N} (H(s_i) - V(s_i))^2}{N}}$$

where:

N = |S| is the number of test instances,

 $H(s_i)$  is the predicted score for  $s_i$ ,

 $V(s_i)$  is the true (human) score for  $s_i$ ,

 $\overline{V}(s)$  is the mean true score (on the test set).

### **5** Results

We note that preliminary results for some of the datasets used here have been reported in [20]. The following results include additional datasets and further analysis.

## 5.1 Results on QuEst feature sets

The error scores for all datasets with common (QuEst) black-box (BB) features are reported in Tables 2 and 3, while Table 4 shows the results with glass-box (GB) features for a subset of these datasets, and Table 5 the results with BB and GB features together for the latter datasets. For each Table and dataset, bold-faced figures represent results that are significantly better (paired t-test with  $p \leq 0.05$ ) with respect to the following comparisons, where available:

- BL vs AF;
- BL vs BL+PR;
- AF vs AF+PR; and
- Each feature selection techniques versus AF+PR.

Underlined figures represent best results overall and double underlined figures, the second best results.

In what follows we take a closer look at some of these comparisons, as well as the difference between different types of feature selection methods. We summarise these comparisons graphically in Fig 4. This figure shows the improvements of different feature sets over the BL results. We also discuss the impact of using GB features against BB features only, showing the improvements of various feature sets over the use of black-box features in Fig 6.

Bayesian Feature Selection for Quality Estimation

Dataset	System	#feats.	MAE	RMSE	RAE	RSE
	Mean	-	0.6027	0.7314	1.0172	1.126
	BL	17	0.4857	0.6178	0.8780	1.2071
EAMIII-en-es	AF	80	0.4719	0.5418	0.8709	1.1808
	BL+PR	18	0.4490	0.5329	0.8417	0.9917
	AF+PR	81	0.4471	0.5301	0.8414	0.9905
	FS(GP-dev)	13	0.4370	0.5199	0.8424	0.9911
	FS(GP-fixed)	17	0.4397	$\overline{0.5224}$	0.8411	0.9901
	FS(GP-test)	20	0.4320	0.5260	0.8330	0.9861
	FS(RL)	69	0.4457	0.5324	0.8501	0.9937
	Mean	-	0.5411	0.6927	1.0813	1.314
	BL	17	0.4401	0.6301	0.9829	1.089
EAMT11-fr-en	AF	80	0.4292	0.6222	0.9708	1.001
	BL+PR	18	0.4183	0.6213	0.9614	0.9972
	AF+PR	81	0.4169	0.6181	0.9682	0.9913
	FS(GP-dev)	15	0.4123	0.6021	0.9627	0.9937
	FS(GP-fixed)	17	0.4166	0.6176	0.9614	0.9920
	FS(GP-test)	10	0.4110	0.6099	0.9387	0.9891
	FS(RL)	65	0.4165	0.6180	0.9553	0.9871
	Mean	_	0.5382	0.7092	1.0392	1.0036
	BL	17	0.5313	0.6655	0.8217	0.8111
EAMT09-s <sub>1</sub>	AF	80	0.5265	0.6538	0.7949	0.7535
	BL+PR	18	0.5123	0 .6492	0.7712	0.7423
	AF+PR	81	0.5109	0.6441	0.7467	0.7374
	FS(GP-dev)	16	0.5055	0.6409	0.7367	0.7166
	FS(GP-fixed)	17	0.5045	0.6392	0.7350	0.7175
	FS(GP-test)	13	$\frac{0.5025}{0.5025}$	$\frac{0.6391}{0.6391}$	$\frac{0.7317}{0.7317}$	0.7117
	FS(BL)	73	0.5195	0.6416	$\frac{0.7011}{0.7401}$	$\frac{0.7111}{0.7320}$
	Mean	-	0.6854	0.7926	1 0060	1.0016
	BL	17	0.4614	0.5816	0.8933	0.7384
EAMT09-s <sub>2</sub>	AF	80	0.4741	0.5953	0.8997	0.790
	BL+PR	18	0.4493	0.5692	0.8714	0.7211
	AF+PR	81	0.4609	0.5821	0.8701	0.7188
	FS(GP-dev)	17	0.4514	0.5735	0.8711	0.7120
	FS(GP-fixed)	17	$\frac{0.1011}{0.4514}$	0.5735	$\frac{0.0711}{0.8711}$	$\frac{0.7120}{0.7120}$
	FS(CP tost)	19	0.4014	0.5625	0.8514	$\frac{0.7120}{0.7013}$
	FS(BL)	12 50	0.4410	0.5025	0.8514	0.7013
	Moan		0.4001	0.3307	1.0013	1 0000
	BL	- 17	0.0733	0.7751	0.8114	0.7628
EAMT09-s <sub>3</sub>		80	0.5339 0.5437	0.6827	0.3114	0.7020 0.7535
	BL_PR	18	0.5457	0.0827	0.7545	0.7000
	$\Delta F \perp P R$	81	0.5115	0.6771	0.7701	0.7430
	FS(GP-dev)	15	0.5500	0.6591	0.7527	0.7324
	FS(GP-fixed)	17	0.5140	0.6572	0.7515	0.7312
	$FS(CP_{test})$	15	0.5160	$\frac{0.0012}{0.6410}$	$\frac{0.7010}{0.7471}$	$\frac{0.7012}{0.7250}$
	FS(BL)	67	0.5000	0.6727	0.7680	0.7450
	Mean	01	0.0230	0.6112	0.1000	1 0000
	BI	- 17	0.43304	0.0112	0.3331	0.0750
EAMT09-s <sub>4</sub>		80	0.3578	0.4942	0.7319	0.9739
	BL_PR	18	0.370	0.4900	0.7555	0.9712
	AF+PR	81	0.3401	0.4816	0.7224	0.0004
	$FS(GP_{dov})$	18	0 3381	0 4814	0 7112	0.0381
	FS(CP fared)	10	0.0001	0.4014	0.7113	0.3001
	FS(GF-IIXed)	10	0.3363	$\frac{0.4011}{0.4700}$	$\frac{0.7101}{0.7011}$	$\frac{0.9314}{0.0312}$
	FS(GP-test)	19	$\frac{0.3370}{0.3404}$	0.4799	$\frac{0.7011}{0.7001}$	$\frac{0.9312}{0.0571}$
	FS(KL)	40	0.3404	0.4805	0.7221	0.9571

Table 2 Results for datasets with common BB features - part	Table 2	2 Results for	r datasets with	n common	BB features -	- part 1.
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Dataset	System	#feats.	MAE	RMSE	RAE	RSE
	Mean	-	0.8278	0.9898	1.0247	1.0263
	BL	17	0.6821	0.8117	0.8622	0.7152
	AF	80	0.6717	0.8103	0.8476	0.6954
WMT12	BL+PR	18	0.6290	0.7729	0.8139	0.6678
	AF+PR	81	0.6324	0.7735	0.7745	0.6699
	FS(GP-dev)	15	0.6231	0.7658	0.7719	0.6651
	FS(GP-fixed)	17	0.6224	0.7645	0.7701	0.6644
	FS(GP-test)	19	0.6131	0.7598	0.7613	0.6586
	FS(RL)	71	0.6335	0.7724	0.7755	0.6709
	Mean	-	0.1438	0.1792	1.0036	1.0009
WMT13	BL	17	0.1411	0.1812	0.9723	1.080
		114	0.1389	0.1775	0.9211 0.0711	1.062
	$\Delta F \perp P R$	10	0.1409 0.1367	0.1602 0.1750	0.9711	0.0603
	FS(GP-dev)	19	0.1217	0.1411	0.8907	0.9577
	FS(GP-fixed)	17	$\frac{0.1211}{0.1242}$	0.1489	$\frac{0.0001}{0.8911}$	$\frac{0.0011}{0.9581}$
	FS(GP-test)	19	0.1207	0.1481	0.8844	0.9531
	FS(RL)	101	$\frac{0.1201}{0.1317}$	0.1691	0.9201	0.9709
	Mean	-	0.5823	0 7214	0.8533	0.8125
	BL	17	0.5462	0.6885	0.8186	0.7715
$GALE11-s_1$	AF	123	0.5399	0.6805	0.8017	0.7619
	BL+PR	18	0.5301	0.6814	0.7929	0.7595
	AF+PR	81	0.5249	0.6766	0.7905	0.7520
	FS(GP-dev)	21	0.5220	0.6751	0.7871	0.7581
	FS(GP-fixed)	17	0.5231	0.6761	0.7856	0.7577
	FS(GP-test)	27	0.5210	0.6701	0.7811	0.7434
	FS(RL)	56	0.5258	0.6749	0.7914	0.7530
	Mean	-	0.5850	0.7527	0.8573	0.8313
GALE11-s <sub>2</sub>	BL	17	0.5540	0.7117	0.8287	0.7978
- 2	AF	123	0.5401		0.8232	0.7931
	BL+PR	18	0.5401	0.7014	0.8221	0.7901
	$FS(CP_{dev})$	01 22	0.5249 0 5107	0.0800	0.8133	0.7853
	FS(GP-fixed)	17	$\frac{0.5191}{0.5210}$	0.6794	0.8193	0.7800
	FS(CP tost)	21	0.5219	0.0734	0.0125	0.7815
	FS(BL)	54	0.5194	0.6805	0.8081	0.7813
	Maan	94	0.0200	0.0000	0.0104	1.0945
	BI	- 17	0.1520 0.1250	0.1727	.9950	1.0240 0.0223
LIG	AF	80	0.1200 0.1227	0.1038	0.9309	0.9223
	BL+PR	18	0.1159	0.1527	0.9312	0.9191
	AF+PR	81	0.1161	0.1534	0.9313	0.9183
	FS(GP-fixed)	17	0.1087	0.1504	0.9287	0.9135
	FS(GP-dev)	13	0.1077	0.1493	0.9284	0.9145
	FS(GP-test)	13	0.1054	0.1489	0.9227	0.9099
	FS(RL)	69	0.1154	0.1519	0.9321	0.9178
	Mean	-	0.1868	0.2470	1.0079	1.0030
	BL	17	0.1807	0.2429	0.9749	0.9695
RACE-fr-en	AF	80	0.1776	0.2422	0.9678	0.9611
	BL+PR	18	0.1729	0.2210	0.9611	0.9554
	AF+PR	81	0.1687	0.2127	0.9601	0.9521
	FS(GP-dev)	18	0.1564	0.2081	0.9571	0.9499
	FS(GP-fixed)	17	0.1597	0.2101	0.9565	0.9487
	FS(GP-test)	19	0.1554	0.2089	0.9515	0.9421
	FS(RL)	72	0.1654	0.2109	0.9589	0.9501
	Mean	-	0.1891	0.2460	1.0004	1.0021
FRACE-en-fr	BL	17	0.1782	0.2389	0.9315	0.9354
		80	0.1657	0.2279	0.9261	0.9292
	BL+PK	18	0.1729	0.2210	0.9292	0.9314
	AF + PK FS(CP dow)	82 15	U.1037 0 161F	0.2127	0.9211	0.9226
	FS(GP-fived)	10 17	0.1010	0.2103	0.9209 0.9201	0.9441
	FS(GP_teet)	18	0.1611	0.2121	0.9178	0.0109
	FS(RL)	71	0.1633	0.2241	0.9209	0.9220
		1 1	0.1000	0.4441	0.0400	0.0440

Table 3	Results	for	datasets	with	$\operatorname{common}$	BB	features -	part 2.
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Bayesian Feature Selection for Quality Estimation

Dataset	System	#features	MAE	RMSE	RAE	RSE
WMT19	AF	47	0.7066	0.8445	0.9114	0.7467
VV IVI I 12	FS(GP-dev)	19	0.6789	0.8318	0.8931	0.7210
	FS(GP-fixed)	17	0.6775	0.8314	0.8914	0.7196
	FS(GP-test)	21	0.6755	0.8298	0.8814	0.7119
	FS(RL)	35	0.6921	0.8388	0.9057	0.7411
CALE11 -	AF	39	0.5736	0.7402	0.8516	0.8131
GALEII-S <sub>1</sub>	FS(GP-dev)	18	0.5715	0.7385	0.8499	0.8080
	FS(GP-fixed)	17	0.5732	0.7381	0.8491	0.8098
	FS(GP-test)	19	0.5702	0.7361	0.8441	<u>0.8060</u>
	FS(RL)	33	0.5781	0.7481	0.8501	0.8111
	AF	48	0.5540	0.6979	0.8317	0.8060
GALE11-s <sub>2</sub>	FS(GP-dev)	21	0.5451	0.6998	0.8243	0.8005
	FS(GP-fixed)	13	0.5491	0.6944	0.8261	0.8011
	FS(GP-test)	13	0.5411	0.6934	0.8220	0.8001
	FS(RL)	41	0.5512	0.6950	0.8301	0.8041

Table 4 Results with GB features.

Dataset	System	#features	MAE	RMSE	RAE	RSE
WMT19	AF	128	0.7185	0.8451	0.9137	0.7507
VV IVI I 12	FS(GP-dev)	25	0.6121	0.7582	0.7692	0.6629
	FS(GP-fixed)	17	0.6161	0.7591	0.7703	0.6639
	FS(GP-test)	29	0.6101	0.7561	0.7679	0.6611
	FS(RL)	99	0.6601	0.8098	0.7815	0.6779
CALE11 o	AF	163	0.5455	0.6722	0.8226	0.7755
GALEII-SI	FS(GP-dev)	27	0.5162	0.6699	0.7801	0.7411
	FS(GP-fixed)	17	0.5170	0.6701	0.7810	$\overline{0.7421}$
	FS(GP-test)	30	0.5150	0.6681	0.7785	0.7384
	FS(RL)	132	0.5310	0.6731	0.7911	0.7501
	AF	172	0.5239	0.6529	0.8177	0.7915
GALE11-s <sub>2</sub>	FS(GP-dev)	27	0.5120	0.6441	0.8094	0.7812
	FS(GP-fixed)	17	0.5119	0.6451	0.8111	0.7823
	FS(GP-test)	17	0.5109	0.6431	0.8063	0.7795
	FS(RL)	132	0.5155	0.6499	0.8144	0.7887

Table 5Results with common BB & GB features.

Baseline features vs all feature sets Adding more features (systems AF) leads to better results in most cases as compared to the baseline systems BL, except for some of the EAMT-09 datasets. However, these are small and in some cases not significant improvements. Adding more features may bring more relevant information, but at the same time it makes the representation more sparse and the learning prone to overfitting. Larger improvements over the baseline come from using either a pseudo-reference feature or performing feature selection, as we discuss in what follows.

Impact of the pseudo-reference feature Adding a single feature, the pseudo-reference (systems BL+PR) to our baseline improves results in all datasets, often by a large margin. Similar improvements are observed by adding this feature to the set with all available features (systems AF+PR).





Fig. 5 Error on (a) WMT12 and (b) EAMT11-en-es datasets with all BB features ranked by GPs.

**Impact of feature selection** Our experiments with feature selection using both GPs and RL led to significant improvements over the entire set of features. Feature selection with GPs has shown better performance over feature selection with RL.

For a more comprehensive overview of the results of feature selection using GPs, we plot the MAE error scores (on the test sets) for different cutoff points on the features in our forward selection method after features are ranked by GPs. The plots for two of our datasets are given in Fig 5. The yaxis shows the MAE scores, while the x axis shows the number of features selected. Generally, we observe an initially rapid error decrease as features are added until approximately 20 features, where the minimum (optimal) error scores are found, after which error increases to a plateau at approximately 45 features. This shows that while a tiny number of features is insufficient, adding too many low-ranked features degrades performance. Similar curves were observed for all datasets with slightly different ranges for optimal numbers of features and best score. It is interesting to note that the best performance on most datasets is observed using 10-20 top-ranked features. This explains why FS(GP-fixed) performs as well as FS(GP-dev) or FS(GP-test), in most cases. Given that FS(GP-dev) and FS(GP-fixed) perform equally well in most cases, the choice between them could be guided by the size of the dataset: if enough data is available to put a development set aside, this should be preferable, while **FS(GP-fixed)** should be used as a cheaper alternative approach if necessary.

**Black-box versus glass-box features** GB features on their own perform worse than BB features (Fig 6), but in all three datasets the combination of GB and BB followed by feature selection resulted in significantly lower error than using only BB features with feature selection, showing that the two features sets are complementary.



Fig. 6 Improvements of various feature sets over black-box features (AF+PR).

## 5.2 Results on WMT12 datasets

In order to investigate whether our feature selection results hold for other feature sets, we experimented with the feature sets provided by most teams participating in the WMT12 QE shared task. These feature sets are very diverse in terms of the types of features, resources used, and their sizes. As shown in Table 6, we observed similar results: feature selection with GPs has the potential to outperform models with all initial feature sets. For these experiments, the significance is checked between system AF trained with GPs versus each feature selection techniques. Improvements were observed even on feature sets which had already been produced as a result of some other feature selection technique. Table 6 also shows the official results from the shared task [6], which are often different from the results obtained with GPs even before feature selection, simply because of differences in the learning algorithms used. In some cases the results with GPs without feature selection (AF) are better than the official results, notably for WLV-SHEF and DCU - which both use large sets of linguistically-motivated features, showing the potential of GPs as a learning algorithm for QE.

The error curves on the test sets for different numbers of top-ranked features have a similar shape to those with the common feature sets. As an example, Fig 7 shows the Uppsala University feature set, with the lowest error score for the 15 top-ranked features.

Team	System	#features	Official	WMT12 score	Score w	ith GP
	· ·		MAE	RMSE	MAE	RMSE
CDI	AF	$15^{*}$	0.61	0.75	0.6030	0.7510
SDL	FS(GP-dev)	10	-	-	0.6021	0.7479
	FS(GP-test)	10	-	-	0.6013	0.7474
	FS(RL)	12	-	-	0.6025	0.7513
TIT	AF	82	0.64	0.79	0.6507	0.8012
00	FS(GP-dev)	13			0.6425	0.7939
	FS(GP-fixed)	17			$\overline{0.6459}$	$\overline{0.7952}$
	FS(GP-test)	10			0.6419	0.7931
	FS(RL)	67			0.6489	0.7979
Louio	AF	49	0.68	0.82	0.6866	0.8340
Loria	FS(GP-dev)	12	-	-	0.6824	0.8355
	FS(GP-fixed)	17	-	-	0.6829	0.8351
	FS(GP-test)	10	-	-	0.6824	0.8395
	FS(RL)	41	-	-	0.6861	0.8395
UEdin	AF	56	0.68	0.82	0.6949	0.8540
UEdin	FS(GP-dev)	16	-	-	0.6839	0 <u>8352</u>
	FS(GP-fixed)	17	-	-	0.6825	0 8373
	FS(GP-test)	20	-	-	0.6795	0.8323
	FS(RL)	49	-	-	0.6899	0.8478
TCD	ÂF	43	0.68	0.82	0.6906	0.8367
ICD	FS(GP-dev)	12	-	-	<u>0.6906</u>	0.8370
	FS(GP-fixed)	17	-	-	$\overline{0.6907}$	0.8372
	FS(GP-test)	10	-	-	0.6904	0.8370
	FS(RL)	37	-	-	0.6913	0.8372
WIV SHEE	AF	147	0.69	0.85	0.6665	0.8219
WLV-SIIEF	FS(GP-dev)	15	-	-	0.6611	0.8090
	FS(GP-fixed)	17	-	-	0.6633	0.8105
	FS(GP-test)	15	-	-	0.6592	0.8088
	FS(RL)	127	-	-	0.6658	0.8168
UDC	AF	57	0.84	1.01	0.8365	0.9601
UPC	FS(GP-dev)	13	-	-	0.8092	0.9304
	FS(GP-fixed)	17	-	-	0.8115	0.9368
	FS(GP-test)	15	-	-	0.8092	0.9288
	FS(RL)	46	-	-	0.8302	0.9588
DCU	AF	308	0.75	0.97	0.6782	0.8394
DCU	FS(GP-dev)	15	-	-	0.6157	0.7671
	FS(GP-fixed)	17	-	-	$\overline{0.6211}$	0.7716
	FS(GP-test)	15	-	-	0.6137	0.7602
	FS(RL)	215	-	-	0.6673	0.8250
рвніт	AF	497	0.70	0.85	0.6733	0.8297
FRIILI	FS(GP-dev)	24	-	-	0.6677	0.8201
	FS(GP-fixed)	17	-	-	0.6697	0.8219
	FS(GP-test)	30	-	-	0.6647	0.8179
	FS(RL)	337	-	-	0.6722	0.8291

Table 6 Results on WMT12 feature sets.  $\ast$  indicates that the initial feature sets already resulted from feature selection.

# 5.3 Commonly selected features

Next we investigate whether it is possible to identify a common subset of features which are selected for the optimal feature sets in most datasets. To





Top-ranked feature	Avg. rank
source sentence perplexity	9.12
number of mismatched quotation marks	11.28
source sentence perplexity without end of sentence marker	12.57
number of tokens in target	15.59
number of tokens in source	15.61
average number of translations per source word in the sentence	16.42
(GIZA++ lexical table thresholded: prob. > 0.1)	
LM log probability of POS of the target	18.33
average number of translations per source word in the sentence	19.12
(GIZA++ lexical table thresholded: prob. $> 0.2$ ) weighted by the	
inverse frequency of each word in the source corpus	
pseudo-reference	19.99
absolute difference between no tokens in source and target normalised	20.45
by source length	
Bottom-ranked features	Avg. rank
Bottom-ranked features absolute difference between number of commas in source and target	<b>Avg. rank</b> 73.10
Bottom-ranked features absolute difference between number of commas in source and target percentage of distinct unigrams seen in the corpus (in all quartiles)	Avg. rank 73.10 69.13
Bottom-ranked features absolute difference between number of commas in source and target percentage of distinct unigrams seen in the corpus (in all quartiles) average unigram frequency in quartile 3 of frequency (lower frequency	Avg. rank 73.10 69.13 65.14
Bottom-ranked features absolute difference between number of commas in source and target percentage of distinct unigrams seen in the corpus (in all quartiles) average unigram frequency in quartile 3 of frequency (lower frequency words) in the corpus of the source sentence	Avg. rank 73.10 69.13 65.14
Bottom-ranked features absolute difference between number of commas in source and target percentage of distinct unigrams seen in the corpus (in all quartiles) average unigram frequency in quartile 3 of frequency (lower frequency words) in the corpus of the source sentence absolute difference between number of : in source and target nor-	Avg. rank 73.10 69.13 65.14 64.19
Bottom-ranked features absolute difference between number of commas in source and target percentage of distinct unigrams seen in the corpus (in all quartiles) average unigram frequency in quartile 3 of frequency (lower frequency words) in the corpus of the source sentence absolute difference between number of : in source and target nor- malised by target length	Avg. rank 73.10 69.13 65.14 64.19
Bottom-ranked features absolute difference between number of commas in source and target percentage of distinct unigrams seen in the corpus (in all quartiles) average unigram frequency in quartile 3 of frequency (lower frequency words) in the corpus of the source sentence absolute difference between number of : in source and target nor- malised by target length absolute difference between number of : in source and target	Avg. rank 73.10 69.13 65.14 64.19 62.12
Bottom-ranked features absolute difference between number of commas in source and target percentage of distinct unigrams seen in the corpus (in all quartiles) average unigram frequency in quartile 3 of frequency (lower frequency words) in the corpus of the source sentence absolute difference between number of : in source and target nor- malised by target length absolute difference between number of : in source and target percentage of tokens in the target which do not contain only a-z	Avg. rank 73.10 69.13 65.14 64.19 62.12 61.16
Bottom-ranked features absolute difference between number of commas in source and target percentage of distinct unigrams seen in the corpus (in all quartiles) average unigram frequency in quartile 3 of frequency (lower frequency words) in the corpus of the source sentence absolute difference between number of : in source and target nor- malised by target length absolute difference between number of : in source and target percentage of tokens in the target which do not contain only a-z number source tokens that do not contain only a-z	Avg. rank 73.10 69.13 65.14 64.19 62.12 61.16 61.01
Bottom-ranked features absolute difference between number of commas in source and target percentage of distinct unigrams seen in the corpus (in all quartiles) average unigram frequency in quartile 3 of frequency (lower frequency words) in the corpus of the source sentence absolute difference between number of : in source and target nor- malised by target length absolute difference between number of : in source and target percentage of tokens in the target which do not contain only a-z number source tokens that do not contain only a-z average number of translations per source word in the sentence	Avg. rank 73.10 69.13 65.14 64.19 62.12 61.16 61.01 59.66
Bottom-ranked features absolute difference between number of commas in source and target percentage of distinct unigrams seen in the corpus (in all quartiles) average unigram frequency in quartile 3 of frequency (lower frequency words) in the corpus of the source sentence absolute difference between number of : in source and target nor- malised by target length absolute difference between number of : in source and target percentage of tokens in the target which do not contain only a-z number source tokens that do not contain only a-z average number of translations per source word in the sentence (threshold in giza: prob > 0.5)	Avg. rank $73.10$ $69.13$ $65.14$ $64.19$ $62.12$ $61.16$ $61.01$ $59.66$
Bottom-ranked features absolute difference between number of commas in source and target percentage of distinct unigrams seen in the corpus (in all quartiles) average unigram frequency in quartile 3 of frequency (lower frequency words) in the corpus of the source sentence absolute difference between number of : in source and target nor- malised by target length absolute difference between number of : in source and target percentage of tokens in the target which do not contain only a-z number source tokens that do not contain only a-z average number of translations per source word in the sentence (threshold in giza: prob > 0.5) percentage of punctuation marks in target	Avg. rank           73.10           69.13           65.14           64.19           62.12           61.16           61.01           59.66           59.11

 ${\bf Table \ 7 \ Top \ and \ bottom \ ranked \ features \ based \ on \ their \ average \ {\tt GP \ rank \ across \ datasets}.}$ 

do that, we took the average rank of each feature across all datasets containing common feature sets. The 10 top and bottom ranked features, along with their average rankings, are shown in Table 7.

Interestingly, not all top ranked features are among the 17 reportedly good baseline features. All of these features are language-independent. Also, most of them are simple and straightforward to extract: they either do not rely on external resources, or else use resources that are easily available, such as language models (e.g., SRILM), or word-alignment models (e.g., GIZA++).

The same analysis on the feature sets from the WMT12 shared task is not possible, given the very little overlap in the competitors' feature sets.

#### 6 Conclusions and future work

We have presented a number of experiments showing the potential of a promising feature ranking technique based on Gaussian Processes for translation quality estimation. Using an oracle approach to select the number of top-ranked features to train quality estimation models, this technique has been shown to outperform all feature sets available with only a small fraction of their features on a number of datasets with common feature sets. More important, we were able to obtain the same or comparable performance with this technique when selecting the number of features based on a development set, or even based on an empirically pre-defined threshold (17). The proposed feature selection technique has also been shown to improve the performance of all participating systems in the WMT12 shared task on quality estimation. Finally, our analysis led to the identification of a set of features which perform well on average across many datasets with different language pairs, machine translation systems, text domains and quality labels.

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