## **A** Appendices

## A.1 Features

We build two sets of features, one for finding the corresponding translation  $t_j$  and  $r_k$  for each  $s_i$ , the other for checking the equivalence if  $t_j$  and  $r_k$ .

## A.1.1 Bilingual alignment features

Bilingual alignment features assume that if  $s_i$  is aligned to any existing  $t_j$  and  $r_k$  by bilingual alignment methods. We rely on Giza++ (Och and Ney, 2000) to find bilingual alignment between S and T (R). However, as Giza++ generates noisy alignment especially for low-frequent words, we propose a bunch of features to complement Giza++ results for more accurate alignments. We apply the same types of features to both T and R, hence only the feature for the alignment between S and T are described here. Letting  $s_i$  is aligned to  $t_j^8$ .

**POS tag:** This is a feature on the source. The intuition is that functional words, indicated by POS tags, usually do not need translation. Hence it may not need to align to any words in the target language.

**NER feature:** This is a binary feature to indicate whether the NER tags of  $s_i$  and  $t_j$  are the same. A correctly aligned word pair should have the same NER tag. Also this feature helps to determine the WT and MT error class.

**Giza++ confidence**: Besides Giza++ translation probability, we also use the word frequency of  $s_i$ and  $t_j$  in our parallel corpus to penalize the alignment confidence score for low frequency words.

**Word-level similarity:** We obtain another alternate translation  $t'_j$  for  $s_i$  using a dictionary and compare the similarity between  $t_j$  and  $t'_j$  using morphology (e.g., edit distance, number of gram letters overlap, common prefix) and semantic dimension (e.g., word embedding similarity).

**Context:** We assume that  $s_i$  and  $t_j$  should be aligned if they are translation equivalent and the same applies to the words linking to  $s_i$  and  $t_j$  in the dependency tree parsed by Standford Corenlp (Manning et al., 2014). Thus, we define context as the number of aligned-pairs among words linking to  $s_i$  and  $t_j$ .

Sentence-level translation quality feature: This include the sentence-level QE shared-task baseline

features used in (Specia et al., 2018). Such features are to estimate the overall quality of the MT hypothesis. The intuition is the that better sentence level translation the less probable word-level misalignment.

## A.1.2 Monolingual equivalence checking

We leverage on monolingual alignment to compare  $t_j$  and  $r_k$  and expect that semantic equivalent words can be aligned by monolingual alignment methods. We use the state-of-the-art alignment tool proposed by Sultan et al. (2014). The tool leverages on paraphrase lexicon (Pavlick et al., 2015) and dependency relations to find equivalent expression between two sentences in the same language. The following features are used.

**Monolingual alignment feature:** A binary feature to indicate whether  $t_j$  and  $r_k$  are aligned by tool proposed by (Sultan et al., 2014). This feature is set to be 0 if either  $t_j$  or  $r_k$  does not exist.

**NER feature:** A binary feature to indicate whether the NER tag of  $t_j$  and  $r_k$  are the same.

<sup>&</sup>lt;sup>8</sup>Following Hu et al. (2018), we make adjustment based on by penalizing the frequency of both words.