MorphAGram Framework for Morphological Segmentation

Preprocessing Phase

MorphAGram uses the Pitman-Yor Adaptor-Grammar Sampler (PYAGS), developed by Mark Johnson, for training. The sampler can be downloaded at:

http://web.science.mq.edu.au/~mjohnson/code/py-cfg-2013-09-23.tgz

For complete information about how the sampler works, please see https://cocosci.princeton.edu/tom/papers/adaptornips.pdf

The sampler requires two types of inputs: a grammar and a list of training units. For the purpose of unsupervised morphological segmentation, a grammar should specify word structure, while the list of training units is a list of raw words. In addition, MorphAGram requires the text to be in the Hex format in order to meet the language-independence assumption and to escape special characters. Also the grammar should have the characters in the word list as terminals.

The first step is to provide an initial CFG (Context-Free grammar) and a list of words (one word per line) to MorpAGram, which in turn converts them into inputs to PYAGS. The initial CFG should have two parameters associated with each production rule (default values are zeros). The first number represents the value of the probability of the rule in the generator, and the second number is the value of the α parameter in the Pitman-Yor process. Below is an example CFG.

```
1 1 Word --> Prefix Stem Suffix

Prefix --> ^^
Prefix --> ^^
Prefix --> ^^
PrefixMorphs --> PrefixMorph PrefixMorphs
1 1 PrefixMorphs --> PrefixMorph PrefixMorphs
1 1 PrefixMorphs --> SubMorphs

Stem --> SubMorphs

Suffix --> $$$
Suffix --> $UffixMorph SuffixMorph SuffixMorphs --> SuffixMorph SuffixMorphs --> SuffixMorph SuffixMorphs
1 1 SuffixMorphs --> SubMorph
SuffixMorph --> SubMorph
SubMorph --> SubMorph
1 1 SubMorphs --> SubMorph
1 1 SubMorphs --> SubMorph
1 1 SubMorphs --> Char
1 1 Chars --> Char
1 1 Chars --> Char
1 1 Chars --> Char Chars
```

MorphAGram has three learning settings; Standard, Scholar-Seeded and Cascaded. For complete details, please see <Link to our paper once published>.

Here is how to preprocess the data for each setup:

```
Standard Setting
#Read the initial word list and convert it into HEX
words, encoded words, hex chars = process words(word list path)
write_encoded_words(encoded_words, word_list_output_path)
#Read the initial CFG and append the characters as terminals
grammar = read grammar(grammar path)
appended grammar = add chars to grammar(grammar, hex chars)
write grammar(appended grammar, grammar output path)
# <word list output path> and <grammar output path> then become the inputs to PYAGS.
Scholar-Seeded Setting
#Read the initial word list and convert it into HEX
words, encoded words, hex chars = process words(word list path)
write encoded words(encoded words, word list output path)
#Read the initial CFG
grammar = read_grammar(grammar_path)
#Seed scholar-seeded affixes into the grammar, where the affixes are read from a <
scholar_seeded_path> file that should have the following format:
###PREFIXES###
anti
dis
extra
im
in
###SUFFIXES###
's
ina
ed
##prefix_nonterminal> and <suffix_nonterminal> are two strings that specify which nonterminals the
prefixes and suffixes should be seeded into, respectively.
ss grammar = prepare scholar seeded grammar(scholar seeded path, prefix nonterminal,
suffix nonterminal)
write grammar(ss grammar, ss grammar path)
#Append the characters as terminals
appended ss grammar = add chars to grammar(ss grammar, hex chars)
write grammar(appended ss grammar, ss grammar output path)
# <word_list_output_path> and <ss_grammar_output_path> then become the inputs to PYAGS.
Cascaded Setting
#Read the initial word list and convert it into HEX
words, encoded words, hex chars = process words(word list path)
write encoded words(encoded words, word list output path)
#Read the initial CFG and append the characters as terminals
grammar = read grammar(grammar path)
#Build the cascaded grammar by seeding affixes whose count is <number of affixes>, from the
```

segmentation output of a first round of learning that is written into <segmentation output path>.

#<nonterminals_to_parse> a bar-separated string that denotes the nontermials that will be parsed (starting with the prefix and ending with the suffix), e.g., 'Prefix|Suffix'.

##prefix_nonterminal> and <suffix_nonterminal> are two strings that specify which nonterminals the
prefixes and suffixes should be seeded into, respectively.

cascaded_grammar = prepare_cascaded_grammar(grammar, segmentation_output_path,
number_of_affixes, nonterminals_to_parse, prefix_nonterminal, suffix_nonterminal)
write_grammar(cascaded_grammar, cascaded_grammar_path)

#Append the characters as terminals

appended_cascaded_grammar = add_chars_to_grammar(cascaded_grammar, hex_chars)
write_grammar(appended_cascaded_grammar, cascaded_grammar_output_path)
<word_list_output_path> and <cascaded_grammar_output_path> then become the inputs to
PYAGS.

Note: The appended characters and seeded affixes are non-adapted by default, where the parameters of their corresponding production rules are set to 1.

Training Phase

Download and run the Pitman-Yor Adaptor-Grammar Sampler (PYAGS), developed by Mark Johnson, where the input to the sampler is the output of the preprocessing phase. http://web.science.mq.edu.au/~mjohnson/code/py-cfg-2013-09-23.tgz

For complete information about how the sampler works, please see https://cocosci.princeton.edu/tom/papers/adaptornips.pdf

Segmentation Phase

Note: When segmenting a file, the segmentation of seen words is read from the output map generated by parsing the PYAGS segmentation output. For unseen words, our code applies MLE-based heuristics that approximate the parse of the PYAGS grammar. We strongly recommend parsing the grammar using the CKY parsing algorithm instead of using our code for segmentation.

In order to do segmentation upon running the Pitman-Yor Adaptor-Grammar Sampler (PYAGS), MorphAGram first parses the segmentation output of the sampler and converts it into words and their respective segmentations, which are then used to segment any given input text.

Parsing PYAGS Segmentation Output

#Parse the sampler's segmentation output and produce two output files: segmented words and a map of segmented words (dictionary).

<word_output_path> and <dic_output_path> are the file paths of the output segmented words and the output dictionary of words mapped to their segmentations, respectively.

<<u>nonterminals_to_parse</u>> is a bar-separated string that denotes the nontermials and the order that will be parsed and returned in the final output , e.g., 'PrefixMorph|Stem|Suffix'.

The segmented words are separated by "+", and when a "Stem" nonterminal is part of <<u>nonterminals_to_parse</u>>, the stem gets encased in parentheses, e.g. "re+(place)+d".

map = parse_PYAGS_segmentation_output(pyags_output_path,, nonterminals_to_parse,
word_output_path, dic_output_path)

Segmenting a White-Space Tokenized Plain Text File

- # Use the map output from extract all words to segment a white-space tokenized plain text file.
- # <map> is the output from extract_all_words .
- # < input_path > is the path of the input plain text file (assumed to be white-space tokenized).
- # < output path > is the path where the segmented plain text file is written.
- # < min_word_length_to_segment > is an integer that represents the minimum length of a stem in the segmentation output of unseen words
- # < multiway_segmentation > is an optional boolean that forces composite affixes to be further split into simple ones when set to True.

Segment(map, input path, output path, min word length to segment, multiway segmentation)