IIS — hw1

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1 UML Design

See Fig. 1 and Fig. 2.

2 System Description

At its core, the system uses a statistical named entity recognizer provided by Lingpipe. The recognizer is HMM based and pre–trained.

The system is rather simple. There is a collection reader called MyCollectionReader, an annotator called GeneAnnotator, and a consumer called MyConsumer. There is also a wrapper class for the Lingpipe package we used, called Chunker. Following is the description of the components.

MyCollectionReader reads the input, which is specified in the descriptor by the parameter name InputFileName. The specified file is opened and read in initialize. There is a design decision of whether the whole file should be loaded in advance or each line should be extracted from the stream on the fly. I choose the former approach since MyCollectionReader needs to implement the abstract method getProgress of the convenient base class CollectionReader_ImplBase. If the content is read from the stream, it would be hard to know the current location in the file. However, loading the whole file can be costly; so if loading a huge corpus is required, something more sophisticated should be implemented. For now the whole file is loaded into a List, and MyCollectionReader#getNext gets the content of the next line in the file through an iterator on that List. We pass on the line as a Sentence annotation, which has two self-explanatory properties id and text.

GeneAnnotator has an initialize method where it initializes the wrapper Chunker c with the model file path specified in the configuration parameter ModelPath. The constructor of Chunker in turn loads the specified model file using the API provided by Lingpipe. The process method retrieves the Sentences of the input annotation. The text property of every Sentence is fed into the named entity recognizer through the wrapper metho Chunker#chunk, which returns a Set of chunked named entites. For each recognized named

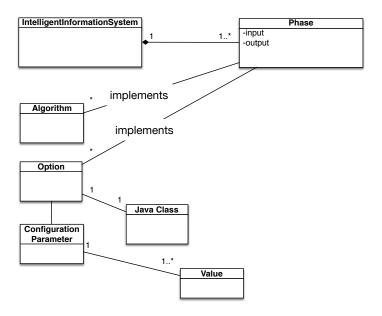


Figure 1: Task 1.1

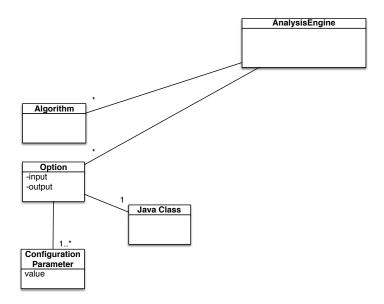


Figure 2: Task 1.2

entity, GeneAnnotator creates a new annotation GeneAnnotation, which contains the properties id. We use the inherited properties begin and end to store offsets, which we also compute in GeneAnnotator. While the homework description asks to compute the offset in the CAS consumer, I find that would introduce unnecessary dependency on MyConsumer — to compute the offset requires both the sentence and the chunked named entity. By computing the offset in GeneAnnotator, MyConsumer does not need to know about Sentence annotations, which may facilitate further maintenance.

MyConsumer gets the specified output filename from the configuration parameter OutputPath in the method initialize. It retrieves the GeneAnnotations associated with an annotation, and and writes into the output file in the specified format.