Abstract. This report describes two methods implemented for the CLEF eHealth 2016 Task 1 challenge. They consist of: a) a feed forward neural network; and b) a random forest for classification and a feed forward neural net, applied to automatically fill in medical handover forms using synthetic medical records as inputs. Both approaches are interesting because they rely on word embeddings, are domain independent, and are feature engineering free. We discuss the complexity of the task, and the impact in our models, having too many output classes and a limited amount of training data. The performance of the methods are based on traditional classification metrics (e.g. precision, recall, and f1-score) on the macro and micro averaged level, and focus on two sets of labels: a) the ”NA” tag, which recognize data that is irrelevant and therefore should be excluded from the form; and b) all other tags, which account for the different fields of the form. The neural network achieved an F1-score of 0.8 (for the ”NA” tag) and a macro-averaged F1-score of 0.308 and a micro-averaged result of 0.514 (for the remaining categories), while the ensemble pipeline got 0.813 (for the ”NA” tag) and 0.345 and 0.503 for the macro- and micro-averaged rates on the rest of the labels.

1 Introduction

This year’s CLEF eHealth challenge consists of 3 tasks[1], that cover different ongoing research topics, briefly summarized as: 1 - Information extraction; 2 - Multilingual Information extraction; 3 - Information retrieval. Due to several factors, such as the overloading amount of information and the lack of standardization procedures when documenting cases, the information flow in the clinical
field results hindered. In consequence, not only non-medical personnel but clinicians have problems in processing this information. Ultimately, these tasks would help in the way in which medical records are handled, processed, and shared, leading to a better understanding overall.

This report describes two statistical approaches to solve the first of these tasks: Task 1: Handover information extraction. In this first assignment, we are presented with plain text records which are the result of automated speech recognition translations from nurses’ shifts verbal information exchange, and we are asked to identify relevant chunks in order to complete a clinical handover form in a fully-automated fashion.

The outline of this report is set as follows: In section 2, a description and analysis of the datasets and the methods is presented; in section 3, the experiments results are shown and explained; section 4 includes conclusions drawn from this work, as well as encountered issues and future work; finally, there is an appendix section that includes additional information referenced in the report.

2 Methodology

2.1 Datasets

Three datasets were released for the purposes of this challenge. Though it was not compulsory, there were meant to be used as independent: a) training; b) validation; and c) testing sets. This was the case for all experiments described in this report, and that is how they are going to be referenced from now on.

Table 1 presents an overview of the datasets. These numbers account for tokens found in the data, with punctuation removal as the only pre-processing step applied. As it is shown in the table, the datasets are roughly the same in terms of size, namely: number of records included, number of tokens, and number of word types. But nearly half of the word types present in the validation and testing set are not seen in the training group (50.04% and 56.55%, respectively, when considering stopwords, and 56.62% and 62.65% excluding them). This is, definitely, an obstacle to overcome; hopefully, the vector representations are going to capture enough semantic meaning to deal with it.

Note that, with no prior handling, constructions like: ‘forty-eight’, ‘self-caring’, ‘self-inflammatory’; numbers, such as: ‘81’, ‘220’; symbols and punctuation: ‘@’, ‘,’; misspellings and wrongly spaced words: ‘bed2’, ‘1pm’, ‘gout.’, ‘litres/nasal’, ‘urine.and’, ‘gastroscopy/colonoscopy’ are treated as single tokens. This criterion follows the way in which the dataset was originally tokenized and listed with the given features.

Already at this stage it can be pointed out, one of the most important limitation to this report’s approaches, and to other techniques of the same nature: there is not enough training data in order to effectively train the neural networks. Considering the number of parameters to be learnt, these models might result too complex to train.

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3 The following symbols were left, as there is a vector representation for them: ‘,’ ‘\^’, ‘@’, ‘\_’, ‘>’, ‘\#’
Table 1: Datasets overview

<table>
<thead>
<tr>
<th>Dataset</th>
<th># docs</th>
<th># tokens</th>
<th># word types w/stopwords</th>
<th>Token overlap w/stopwords</th>
<th>Token overlap w/o stopwords</th>
</tr>
</thead>
<tbody>
<tr>
<td>Training</td>
<td>101</td>
<td>7451</td>
<td>1347</td>
<td>-</td>
<td>-</td>
</tr>
<tr>
<td>Validation</td>
<td>100</td>
<td>6798</td>
<td>1291</td>
<td>645 (49.96%)</td>
<td>560 (43.38%)</td>
</tr>
<tr>
<td>Testing</td>
<td>100</td>
<td>5741</td>
<td>1213</td>
<td>527 (43.45%)</td>
<td>453 (37.35%)</td>
</tr>
</tbody>
</table>

In this task, we are going to learn patterns from the data so as to fill in a fixed handover form. The description of this form can be found in the dataset paper, for the purposes of this explanation, it is relevant to know that it consists of 36 tags/labels; one of these is the 'NA' label, which accounts for information that shouldn’t be included in the form. Naturally, this tag covers the most number of tokens, and as we are dealing with a multiclass classification task, this difference in label-group sizes will play a significant role at prediction time, specifically when computing the resulting averaged metrics.

Looking at the training and validation sets, we can find out that there is a mismatch between the tags that are included in one and the other. Namely, the training dataset includes 36 labels, 3 of which are not found in the validation set (tags: 'Appointment/Procedure_ClinicianGivenNames/Initials', 'Appointment/Procedure_Ward', 'Appointment/Procedure_City'); and the validation set includes 36 tags as well, 3 of which are not seen in the training set (tags: 'PatientIntroduction_Title', 'Appointment/Procedure_ClinicianTitle', 'Appointment/Procedure_Hospital'). Finally, the testing set is in agreement with the training label data. This difference will have an impact on the validation scores, but not on the testing ones.

Conceptually, the statistical models to be trained will try to fit the conditional probability of a label given certain word tokens. It is worth analysing the data at the tag level so to get an idea of how complex the task is. Table 5 and Table 6, included in the Appendix, show the empirical distribution shaped from the training data, and a word type count overlap breakdown. The overlap summary was done considering the validation and the training set (at the moment of writing this report, the testing set labels were not released), and it includes stopwords in the counts. As it can be seen in the empirical distribution, there is one category which concentrates the majority of the tokens (the "NA" tag, leaving a very low probability mass for the rest of the labels. Clearly, some categories are easier than others to predict; but, in the end, this is going to be dependant on: a) the number of samples; and b) the token overlap of the category. These two factors will affect the quality of the trained word embeddings.

In the next subsections, the details of the two models are provided.

2.2 Method 1: a simple neural network approach

The first approach consists of a one hidden layer context window feed forward neural network. Following the idea of constructing a pipeline that is domain
independent, there are no features derived from medical data enrichment, and the neural network makes use of semantic features only, i.e. word embeddings.

From a Natural Language Processing point of view, we know that languages present a high level of ambiguity, and if we, furthermore, take into account the overlap schema presented in the previous section, it seems like a good idea to include a context window on the token to be predicted.

As mentioned before, even this simple architecture might have too many parameters to be trained with the amount of data we have. As a way to help this situation, the word embeddings are initialized using the pretrained Googlenews Word2Vec representations\textsuperscript{4}[5].

Figure 1 shows a graphical description of the implemented neural network\textsuperscript{9}.

![Neural network architecture](image)

Fig. 1: Neural network architecture

The network has two weight matrix and two bias parameters. The matrix shown as $W1$ corresponds to the word embeddings, initialised uniformly and intersected with the pretrained embeddings, while the matrix $W2$ and the bias vectors $b1$ and $b2$ are uniformly initialised. There is a non-linear behaviour in the net, introduced by the hyperbolic tangent function on the hidden layer, and the output layer is a classification step achieved by using a softmax activation function. The output layer has a dimension of 39, because of the 36 tags in the training/test set and the 3 added tags found in the validation set, as explained in the previous section.

The neural network is trained with Stochastic gradient descent, and backpropagation\textsuperscript{6}, using Adagrad\textsuperscript{7} as gradient update optimizer. Concerning the overfitting behaviour of neural nets, an L2-regularization is applied to the weight matrices $W1$ and $W2$.

\textsuperscript{4}https://code.google.com/archive/p/word2vec/
2.3 Method 2: an ensemble approach

In this second method, the pipeline is based on a layered prediction. First, a random forest[8] is set up with the purpose of predicting a subset of the tags, and then a neural network, with the same architecture as described in the previous subsection, is implemented to further discriminate between the remaining labels.

Again, both, the random forest as well as the neural network only make use of semantic features. It could be the case that the two models use different or extended types of features in order to increase their performance. As an advantage, these two methods could possibly learn different patterns from the data, helping the final prediction quality. But, at the same time, the errors that are made in the first step, are further passed to the second model; because false-positive predictions by the random forest are taken out from the sample set to be predicted later by the neural net, and false-negatives are going to be included by the neural net in one of the remaining categories as false-positives values.

The random forest uses decision trees for classification, with 100 trees and a maximum depth of 5. The neural network is trained as described in method 1.

2.4 Pretrained word embeddings

The neural networks described above utilize pretrained word embeddings as features. Given the small amount of data for training, and considering the novelty, with respect to word types, that are included in each dataset, if a word has not been seen in the training data and has no representation in the Googlenews pretrained embeddings, then it will be assigned a randomly generated vector. In these cases, there is no semantic information to use so as to assign a class.

Table 2 shows the statistics of the previously described cases for word tokens and word types in the validation and testing set.

<table>
<thead>
<tr>
<th>Dataset</th>
<th>Tokens</th>
<th>Word types</th>
<th>Most affected tags (top 3) (tag: %)</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>#</td>
<td>#</td>
<td>PatientIntroduction_Lastname: 0.156</td>
</tr>
<tr>
<td>Validation</td>
<td>231</td>
<td>160</td>
<td>PatientIntroduction_GivenNames/Initials: 0.1</td>
</tr>
<tr>
<td></td>
<td>0.03</td>
<td>0.114</td>
<td>MyShift_OtherObservation: 0.091</td>
</tr>
<tr>
<td>Testing</td>
<td>782</td>
<td>602</td>
<td>Tags not available</td>
</tr>
<tr>
<td></td>
<td>0.12</td>
<td>0.475</td>
<td></td>
</tr>
</tbody>
</table>

3 Results and analysis

The performance of the methods measure the precision, recall, and F1-score using the conlleval evaluation script, as implemented in the CoNLL 2000 Shared Task on Chunking⁵.

⁵ http://www.cnts.ua.ac.be/conll2000/chunking/
In this section, the results for the two methods are presented and compared. At the moment of submission, an error when writing the output file produced an alteration in the order of the predicted tags, completely mixing the results. The associated scores are not included in this report.

### 3.1 Method 1

Table 3 shows a summary of the results obtained when using the neural network with a context window size of 7, after being trained for 50 epochs. A detailed, per tag, analysis can be found in the Appendix section, in Table 7.

<table>
<thead>
<tr>
<th>Dataset</th>
<th>Macro average Precision</th>
<th>Macro average Recall</th>
<th>Micro average Precision</th>
<th>Micro average Recall</th>
<th>Micro average F1</th>
<th>NA Precision</th>
<th>NA Recall</th>
<th>NA F1</th>
</tr>
</thead>
<tbody>
<tr>
<td>Training</td>
<td>0.741</td>
<td>0.591</td>
<td>0.624</td>
<td>0.908</td>
<td>0.862</td>
<td>0.884</td>
<td>0.92</td>
<td>0.979</td>
</tr>
<tr>
<td>Validation</td>
<td>0.468</td>
<td>0.344</td>
<td>0.355</td>
<td>0.636</td>
<td>0.495</td>
<td>0.557</td>
<td>0.696</td>
<td>0.92</td>
</tr>
<tr>
<td>Testing</td>
<td>0.411</td>
<td>0.307</td>
<td>0.308</td>
<td>0.563</td>
<td>0.472</td>
<td>0.514</td>
<td>0.723</td>
<td>0.894</td>
</tr>
</tbody>
</table>

While training, the word embedding vectors increase their norms, this effect is reflected in Figure 2 and Figure 3, where a PCA plot of the word representations before and after training is shown. In this graph, the 36 training tags are plotted (overlapped words appear as a separated colour). It does not seem visually clear in 2 dimensions how the data could be separated; but the further transformations of these representations and the non linearity added by the network are able to identify semantic regions, to some extent.
Figure 4 presents a confusion matrix of the neural network output when predicting on the validation set. The neural net achieves a high score for some of the easy tags, such as PatientIntroduction, CurrentBed, PatientIntroduction, CurrentRoom, PatientIntroduction, Gender, while having a poor performance in complex tags like Future, Goal, TaskToBeCompleted, ExpectedOutcome or PatientIntroduction, CarePlan, but also in other easy tags like PatientIntroduction, UnderDr, GivenNames, Initials (which are misclassified as PatientIntroduction, UnderDr, Lastname). As it can be seen in the plot, many samples are being wrongly classified as "NA". The ensemble approach of method 2 will try to tackle this misclassification behaviour.

3.2 Method 2

In this case, the random forest uses a randomly initialized matrix intersected with the Googlenews pretrained embeddings as input features, and predicts whether the token belongs to the "NA" label or not. Later on, the neural network of the second step discriminates between the remaining labels. This latter model was trained using a context window of 7 for 50 epochs.

Figure 5 presents a PCA plot of the word embeddings corresponding to the tag "NA", the remaining tags, and also separates word types that belong to "NA" and, at least, some other label.

Table 4 presents a summary of the ensemble method results. A detailed, per tag, analysis can be found in Table 8.
Figure 6 shows the confusion matrix when using the ensemble model to predict on the validation set. This time, the NA false-positive results are reduced, in comparison to the first approach. While incrementing the number of true-positives, some mistakes are translated to other categories, the most prominent being: Appointment/Procedure_Description, Future_Goal/TaskToBeCompleted/ExpectedOutcome, MyShift_OtherObservation, and PatientIntroduction.AdmissionReason/Diagnosis.

The non-ensemble method achieved a macro-averaged F1-score of 0.308 on the 35 tags (all tags excluding "NA") and the ensemble system performed at 0.345. This means the ensemble method performs better overall, obtaining higher F1-results for 17 out of the 36 classes (with improvements of up to 0.3), maintaining the same scores for 8 of them, and lowering them in 11 cases (with decrements of max 0.08). Some of these improvements imply that labels that previously had an F1-score of 0.0 are now getting 0.308, like the case of PatientIntroduction_CarePlan. Considering the micro-averaged, these results are translated to 0.514 and 0.503, respectively. Analysing this updates, the F1-value goes down due to the decrease in the micro-averaged precision (the micro-averaged recall increases). In the vast majority of the cases, the "NA" false-positive classifications of method 1 are now being assigned to the remaining classes causing an increase.
in their recall (lowering the false-negatives samples), and even though negatively affecting the precision, the gain is big enough to push the F1-relation up. But for some categories this statement does not hold, and it is the sum of these misclassified tokens which causes the micro-averaged reduction. Examples of this negative behaviour can be found in labels: a) "MyShift Input/ Diet", where previously correct tokens are now assigned to other classes, while also wrongly predicting tokens as members of this tag; b) "MyShift Status", where 79 false-positives are added just for gaining 2 tokens in true predictions; and c) "PatientIntroduction, AdmissionReason/ Diagnosis", case in which 90 false-positives are included, with an advantage of only 3 new true-positive samples.

4 Conclusions and future work

In this report, a feed forward neural network, and a random forest - feed forward neural network ensemble method are presented as solutions for the CLEF eHealth Task 1 challenge. Both methods rely on semantic features and are domain independent (no medical features are used). While the neural network alone achieves a macro-averaged F1-score of 0.308 on the test set, considering 35 categories (all but "NA"), and an F1-score of 0.8 for the "NA" tag; the ensemble method produces better results with a macro F1-score of 0.345 on the 35 tags of the test dataset, and an F1-score of 0.813 for the "NA" label, increasing the precision but affecting the recall metric. This gain in F1-results suggests the second method performs better when considering the entire set of labels. On
the other hand, from the micro-averaged perspective, the former method gets an F1-score of 0.514 for the 35 tags, and the latter an F1-score of 0.503, which can be explained from the relation between the general raise of the recall and the decrement in the precision.

As explained in the analysis section, the handover form presents a large number of tags, while some appear to be easy to learn, some others are clearly a complex task. One of the main problems limitations, given the Machine Learning methods implemented, is the amount of data available for training.

In this report’s pipelines no pre-processing steps were applied, and the negative effects of this decision were explicitly pointed out in the analysis. There should be a prior stage in which abbreviations, misspellings, and errors during tokenization are treated.

While this work shows that semantic representations are able to help in this task, most likely, a higher performance could be achieved by incorporating features of other types; lexical features such as part-of-speech tags, or dependency parsing, as well as features resulting from external taggers. Moreover, a potentially useful characteristic, not exploited in this work, is the natural structure of the medical records, for instance, word or sentence locations, and tag-precedence.

References

### Table 5: Training data empirical distribution

<table>
<thead>
<tr>
<th>Tag</th>
<th>Probability</th>
</tr>
</thead>
<tbody>
<tr>
<td>PatientIntroduction_GivenNames/Initials</td>
<td>0.0140</td>
</tr>
<tr>
<td>PatientIntroduction_Lastname</td>
<td>0.0117</td>
</tr>
<tr>
<td>PatientIntroduction_AgeInYears</td>
<td>0.0290</td>
</tr>
<tr>
<td>PatientIntroduction_Gender</td>
<td>0.0576</td>
</tr>
<tr>
<td>PatientIntroduction_CurrentRoom</td>
<td>0.0064</td>
</tr>
<tr>
<td>PatientIntroduction_CurrentBed</td>
<td>0.0312</td>
</tr>
<tr>
<td>PatientIntroduction_UnderDr_GivenNames/Initials</td>
<td>0.0018</td>
</tr>
<tr>
<td>PatientIntroduction_UnderDr_Lastname</td>
<td>0.0213</td>
</tr>
<tr>
<td>PatientIntroduction_AdmissionReason/Diagnosis</td>
<td>0.0488</td>
</tr>
<tr>
<td>PatientIntroduction_Allergy</td>
<td>0.0016</td>
</tr>
<tr>
<td>PatientIntroduction_ChronicCondition</td>
<td>0.0082</td>
</tr>
<tr>
<td>PatientIntroduction_Disease/ProblemHistory</td>
<td>0.0173</td>
</tr>
<tr>
<td>PatientIntroduction_CarePlan</td>
<td>0.0042</td>
</tr>
<tr>
<td>MyShift_Status</td>
<td>0.0569</td>
</tr>
<tr>
<td>MyShift_Contraption</td>
<td>0.0072</td>
</tr>
<tr>
<td>MyShift_Input/Diet</td>
<td>0.0119</td>
</tr>
<tr>
<td>MyShift_Output/Diuresis/BowelMovement</td>
<td>0.0081</td>
</tr>
<tr>
<td>MyShift_Wounds/Skin</td>
<td>0.0085</td>
</tr>
<tr>
<td>MyShift_ActivitiesOfDailyLiving</td>
<td>0.0289</td>
</tr>
<tr>
<td>MyShift_RiskManagement</td>
<td>0.0014</td>
</tr>
<tr>
<td>MyShift_OtherObservation</td>
<td>0.0425</td>
</tr>
<tr>
<td>Appointment/Procedure_Status</td>
<td>0.0187</td>
</tr>
<tr>
<td>Appointment/Procedure_Description</td>
<td>0.0185</td>
</tr>
<tr>
<td>Appointment/Procedure_ClinicianGivenNames/Initials</td>
<td>0.0002</td>
</tr>
<tr>
<td>Appointment/Procedure_Clinician_Lastname</td>
<td>0.0002</td>
</tr>
<tr>
<td>Appointment/Procedure_Day</td>
<td>0.0047</td>
</tr>
<tr>
<td>Appointment/Procedure_Time</td>
<td>0.0033</td>
</tr>
<tr>
<td>Appointment/Procedure_City</td>
<td>0.0002</td>
</tr>
<tr>
<td>Appointment/Procedure_Ward</td>
<td>0.0004</td>
</tr>
<tr>
<td>Medication_Medicine</td>
<td>0.0185</td>
</tr>
<tr>
<td>Medication_Dosage</td>
<td>0.0044</td>
</tr>
<tr>
<td>Medication_Status</td>
<td>0.0000</td>
</tr>
<tr>
<td>Future_Alert/Warning/AbnormalResult</td>
<td>0.0070</td>
</tr>
<tr>
<td>Future_Goal/TaskToBeCompleted/ExpectedOutcome</td>
<td>0.0584</td>
</tr>
<tr>
<td>Future_Discharge/TransferPlan</td>
<td>0.0105</td>
</tr>
<tr>
<td>NA</td>
<td>0.4443</td>
</tr>
</tbody>
</table>
Table 6: Validation set overlap

<table>
<thead>
<tr>
<th>Tag</th>
<th># Word types</th>
<th>Overlap</th>
<th># Unique</th>
<th>Overlapping tags (top 3)</th>
</tr>
</thead>
<tbody>
<tr>
<td>PatientIntroduction_GivenNames/Initials</td>
<td>4</td>
<td>0</td>
<td>1</td>
<td>PatientIntroduction_Lastname: 0.714</td>
</tr>
<tr>
<td>PatientIntroduction_Lastname</td>
<td>101</td>
<td>7</td>
<td>0.005</td>
<td>PatientIntroduction_GivenNames/Initials: 0.714</td>
</tr>
<tr>
<td>PatientIntroduction_Lastname</td>
<td>96</td>
<td>7</td>
<td>0.071</td>
<td>PatientIntroduction_GivenNames/Initials: 0.714</td>
</tr>
<tr>
<td>PatientIntroduction_Lastname</td>
<td>53</td>
<td>7</td>
<td>0.137</td>
<td>PatientIntroduction_GivenNames/Initials: 0.714</td>
</tr>
<tr>
<td>PatientIntroduction_Gender</td>
<td>9</td>
<td>5</td>
<td>0.556</td>
<td>MyShift_OtherObservation: 0.444</td>
</tr>
<tr>
<td>PatientIntroduction_CurrentBed</td>
<td>18</td>
<td>17</td>
<td>0.944</td>
<td>Future_Goal/TasksCompleted/ExpectedOutcome: 0.539</td>
</tr>
<tr>
<td>PatientIntroduction_CurrentBed</td>
<td>25</td>
<td>22</td>
<td>0.886</td>
<td>PatientIntroduction_AdmisionReason/Diagnosis: 0.539</td>
</tr>
<tr>
<td>PatientIntroduction_CurrentBed</td>
<td>29</td>
<td>3</td>
<td>0.103</td>
<td>Appointment/Procedure/OtherObservation: 0.400</td>
</tr>
<tr>
<td>PatientIntroduction_Lastname</td>
<td>55</td>
<td>7</td>
<td>0.127</td>
<td>PatientIntroduction_GivenNames/Initials: 0.333</td>
</tr>
<tr>
<td>PatientIntroduction_AdmisionReason/Diagnosis</td>
<td>256</td>
<td>143</td>
<td>0.416</td>
<td>Future_Goal/TasksCompleted/ExpectedOutcome: 0.400</td>
</tr>
<tr>
<td>PatientIntroduction_AdmisionReason/Diagnosis</td>
<td>8</td>
<td>7</td>
<td>0.7</td>
<td>MyShift_OtherObservation: 0.110</td>
</tr>
<tr>
<td>PatientIntroduction_AdmisionReason/Diagnosis</td>
<td>3</td>
<td>1</td>
<td>0.333</td>
<td>MyShift_OtherObservation: 0.110</td>
</tr>
<tr>
<td>PatientIntroduction_D/ProblemHistory</td>
<td>157</td>
<td>69</td>
<td>0.567</td>
<td>Future_Goal/TasksCompleted/ExpectedOutcome: 0.165</td>
</tr>
<tr>
<td>PatientIntroduction_D/ProblemHistory</td>
<td>78</td>
<td>64</td>
<td>0.803</td>
<td>Future_Goal/TasksCompleted/ExpectedOutcome: 0.168</td>
</tr>
<tr>
<td>MyShift_Status</td>
<td>91</td>
<td>73</td>
<td>0.802</td>
<td>Future_Goal/TasksCompleted/ExpectedOutcome: 0.100</td>
</tr>
<tr>
<td>MyShift_CarePlan</td>
<td>39</td>
<td>19</td>
<td>0.462</td>
<td>MyShift_OtherObservation: 0.136</td>
</tr>
<tr>
<td>MyShift_CarePlan</td>
<td>36</td>
<td>25</td>
<td>0.579</td>
<td>Future_Goal/TasksCompleted/ExpectedOutcome: 0.100</td>
</tr>
<tr>
<td>MyShift_Output/Diagnosis/BedMovement</td>
<td>26</td>
<td>13</td>
<td>0.53</td>
<td>Future_Goal/TasksCompleted/ExpectedOutcome: 0.138</td>
</tr>
<tr>
<td>MyShift_Wounds/Skin</td>
<td>18</td>
<td>14</td>
<td>0.778</td>
<td>Future_Goal/TasksCompleted/ExpectedOutcome: 0.167</td>
</tr>
<tr>
<td>MyShift_ActivityOfDailyLiving</td>
<td>48</td>
<td>20</td>
<td>0.417</td>
<td>PatientIntroduction_AdmisionReason/Diagnosis: 0.098</td>
</tr>
<tr>
<td>MyShift_RiskManagement</td>
<td>37</td>
<td>21</td>
<td>0.568</td>
<td>MyShift_ActivityOfDailyLiving: 0.111</td>
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**Table Notes:**
- **PatientIntroduction** is the main category associated with each tag.
- Overlap indicates the percentage of overlap between tags.
- # Word types refers to the number of unique word types for each tag.
- # Unique denotes the number of unique tags.
- Overlapping tags (top 3) list the most common tags overlapping with the current tag.
### Table 7: Method 1 per tag detailed scores

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Table 8: Method 2 per tag detailed scores