

Semantic Tag Medical Concept using Word2Vec representation

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Where we are?





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Final Goal: (Semantic Search E.H.R)

To develop a Semantic Search engine, we need that the clinical Information, from free text data, it'll be map with a clinical terminology, like (Snomed-CT, ICD-10-MC, UMLS, etc)

Our Goal: (Semantic Tag Clinical Concepts)





Background and Related Work:

Semantic Tag: It is a process of associating an element from a ontology with some document.

S. T. Medical:

It is to map clinical concepts from free text clinical reports with a clinical ontology.



Supervised machine learning methods like CRF (Condition Random Fields), SSVM (Structural support Vector Machines), and UMLS MetaThesaurus, like Clinical Terminology.

Our approach is to use an unsupervised M.L. Neural Network to discover Word Embedding (Word2Vec) with algorithm rules and Snomed-CT like clinical terminology



Semantic Tag Medical Concepts (STMC):

- We proposed a mapping tool to discover from free text to clinical concepts using the ontology clinical terminology, Snomed-CT.
- We use word embedding model (Word2Vec) to represents the word in the texts by vectors and identify the semantic relation between there.
- We use Named-Based techniques combined with a query expansion system, and the Space vector Model, generate with Word2Vec, to find alternative search terms.



What is Word Embedding?

• In Spanish there is a proverb:

"Dime con quien andas y te diré quien eres". [El Quijote II 10 y 23].

"Tell me who are your friends and I'll tell you who you are".

To identify the semantic meaning of a word, it depend of the words around it.

What is Word2Vec? Created by Tomas Mikolov et al. at Google.

Word2vec is a group of related models that are used to produce word embeddings.

These models are shallow, two-layer neural networks that are trained to reconstruct linguistic contexts of words. Word2vec takes as its input a large corpus of text and produces a vector space, with hundred dimensions each unique word in the corpus.



Characteristics Word2Vec - Structure:

The neural network structure of word2vec is a feedforward network with **one hidden** layer.



The **training** method of word2vec is backpropagation with **stochastic gradient descent**.

SoftMax Function:

$$p(w_O|w_I) = \frac{\exp\left(v'_{w_O} {^{\top}v_{w_I}}\right)}{\sum_{w=1}^{W} \exp\left(v'_w {^{\top}v_{w_I}}\right)}$$

Training can be made feasible by using either *hierarchical softmax* or *negative sampling* (Mikolov et al.).



CBOW

Skip-gram



Word2Vec Skip-Gram Model (1/3):

McCormick, C. (2016, April 19). Word2Vec Tutorial - The Skip-Gram Model. Retrieved from http://www.mccormickml.com

Word2Vec uses a trick: Param(00): SG= Size (0 –CBOW, 1-Skip-Gram) We don't train a simple neural network with a single hidden layer to perform a certain task, the goal is just to **learn the weights** of the **hidden layer**.

These **weights** are the "word vectors" that we're trying to learn.

Param(01): Size of Vector = Size Hidden layer

Given a specific word (the input word). The network is going to tell us the **probability for every word** in our vocabulary (SoftMax function) of being the "nearby word" that we chose.

Param(02): Nearby word = Window size



Word2Vec Skip-Gram Model (2/3):

McCormick, C. (2016, April 19). Word2Vec Tutorial - The Skip-Gram Model. Retrieved from http://www.mccormickml.com

Model Details:

W2V build a vocabulary of words from our training documents. We have a **vocabulary of 98,103** unique words.

Param(03): min_count(n)= Ignore frequency(w) < n

We're going to represent an input word like "fracture" as a one-hot vector.





Word2Vec Skip-Gram Model (3/3):

McCormick, C. (2016, April 19). Word2Vec Tutorial - The Skip-Gram Model. Retrieved from http://www.mccormickml.com

There is no activation function on the hidden layer neurons, but the output neurons use **softmax** like clasification method to build a probability distribution.

$$p(w_O|w_I) = \frac{\exp\left(v'_{w_O}{}^{\top} v_{w_I}\right)}{\sum_{w=1}^{W} \exp\left(v'_w{}^{\top} v_{w_I}\right)}$$

Training de model:

Input: one-hot vector for a word

Output: probability distribution vector

Param(04): hs= (1 hierarchical softmax, 0 negative sampling)

We're learning word vectors with 300 features. So the hidden layer is going to be represented by a weight matrix with 98103 rows (one for every word in our vocabulary) and 300 columns (one for every hidden neuron)

We use gensim python library (Parameters):

Param(00): SG= 1

Param(01): Size=300

Param(02): window= 5

Param(03): min_count(n)= 2

Param(04): hs= 0

Param(05): negative_sampling= 5



How identify the meaning between two words? Similarity Distance between Words: Cosine Distance Word Vectors.





Snomed-CT Design and Structure





Snomed-CT Components. Logical Model





Implementation S.T.M.C.:

Big Data Corpus:



We use 615,513 emergency discharge reports. **Emergency Discharge Records**

Emergency Electronic Health Records:

• Emergency Discharge report:

- Administrative Data (anonymised)
- Reason Medical Consultation
- Personal Background:
 - Known Allergies.
 - Medicals
 - Surgeries.
 - Treatment Background.
- Actual illness.
- Exploration.
- Complementary Evidence.
- Evolution.
- Diagnostic.
- Treatment and Recommendations.

General Process Diagram:





Algorithm STMC: Preprocessing Text

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04. Algorithm STMC: Word2Vec Model

Corpus. DocNorm-EReports





04. Algorithm STMC: Word2Vec Cosine distance

Similarity: Cosine distance





05. Algorithm STMC: (Process)

Step 01

Step 02





06. Algorithm STMC: (Process)

Step 04

It will denote **v(w)**, the vector of the word **w** ib the Model **M**

Given a n-gram $\boldsymbol{g} = \boldsymbol{w}_1 \, \boldsymbol{w}_2 \, \dots \, \boldsymbol{w}_n$

Given a n-gram **v(g)** = **v(w**₁) + **v(w**₂) + ... + **v(w**_n)

We define Similarity between 2 n-gram $g_{1,} g_{2}$: Sim() = CosineSimilarity ($v(g_{1}), v(g_{2})$)

We use the *similarity* between *n-grams* to identify if a **concept** is named in a sentence *S*.

Given a concept *c* in a ontology *O*:

Degree in witch a n-gram **g** names the concept **c** as the maximum similarity between **g** and one of the label of **c**:

Names(g,c) = Max{Sim(g,l); I label of c}

Degree in witch *c* is named in a sentence *S*:

Names(g,S) = Max{Names(g,c); g n-gram of c}

If *Names(c,S)=1*, then *c* is standardized named in *S*



07. Algorithm STMC: (Process)

Step 05: Algorithm two passes

Pass 1: We filter n-gram and concepts to possible candidates in standard way.

For every n-gram **g** of **S** and every concept **c** of **O**:

- If *Names(g,c)=1*, then *c* is named in *S* by expression *g*.
- If Names(g,c)>alpha(0.9), then g is added to the list og GC (Grams Candidates), and c to the list of CC(Concepts Candidates) to be named in S.

Pass 2: We check if some n-gram candidate names a concept in a non standard way.

For every n-gram **g** of **GC**, we get a set of **variants** of **g**.

This variants are generated from *g* by replacing some words of *g* by one of a list of 5 *g'=Most_Similar(w',5)* For every *variants g'* of a n-gram *g* of *G*, and any concept *c* of *CC*.

• If Names(g',c)=1, then it is identified that c is named in S by the expression g.



08. Bag of Clinical Concepts (BOCC):

We can represents a Medical reports as a **bag of concepts**, similar way like bag of Words.

Given a concept *c* and a document *d*, we define the *frequency of a concept* in the document: CF(c,d) = |g in d; c named by the expression g|

We represent a document **d**, by the frequency of concepts of **O** in **d**:

CF(d) = {(*c*, *CF*(*c*,*d*)); *c* in *O*}

Or simplifying, Concept Frequency reduced Representation:

- If *c* is not named in *d*, then *c* is not considered in the reduced representation
- if c₁ and c₂ are named in d, and c₁ is more detailed than c₂ in the ontology hierarchy, then we say that c₂ is subsumed by c₁. Subsumed(c₂, c₁), and not considered in reduced representations.

C(d) = {c in O; CF(c,d) > 0}

 $MaxC(d) = \{ c \in C(d); \forall c' \in C(d) \neg Subsumed(c, c') \}$

Then we have the reduced representation by:

 $CFR(d) = \{(c, CF(c, d)); c \text{ in } MaxC(d)\}$



Example: How Algorithm identify a conceptID named In no normalized way

```
In [20]: CN=[]
         GCN=[]
         CC=[]
         GC=[]
         CN, GCN, CC, GC = paso1(Sent5, 0.9, 300)
         Paso.1:
         01. Crear ngramas: [['rodilla'], ['dcha'], ['rodilla', 'dcha']]
         03. Total conceptos de snomed-CT donde aparece alguna etiqueta "l": 1073
         05. (CN) -> Conceptos Nombrados, umbral alfa(1.0): [(1.0, ['rodilla'], ['rodilla'], '72696002')]
         06. (GCN) -> Gramas Candidatas Nombradas, umbral alfa(1.0): [['rodilla']]
         07. (GC) -> Gramas Candidatas, umbral alfa(0.9):
         id: 0 - ['rodilla', 'derecha']
         08. (CC) -> Conceptos Candidatos, superan el umbral alfa(0.9):
         id: 0 - (0.905699999999999995, ['rodilla', 'dcha'], ['rodilla', 'derecha'], '6757004')
         id: 1 - (0.905699999999999995, ['rodilla', 'dcha'], ['rodilla', 'derecha'], '210562007')
         id: 2 - (0.905699999999999995, ['rodilla', 'dcha'], ['rodilla', 'derecha'], '210562007')
```



Example: Identify, relations and similar words, to discover new words meaning

In [17]:	<pre>InfAltModel.most_similar(['fx'])</pre>	In [15]:	<pre>InfAltModel.most_similar(['tce'])</pre>
Out[17]:	<pre>[('fractura', 0.8746222853660583), ('fract', 0.7706700563430786), ('fratura', 0.7633858919143677), ('frx', 0.7520125508308411), ('frac', 0.7503272294998169), ('fisrua', 0.7457991242408752), ('desplazada', 0.7253227233886719), ('conminuta', 0.7239192724227905), ('diafisis', 0.7234492897987366), ('fracrtura', 0.7221113443374634)]</pre>	Out[15]:	<pre>[('tec', 0.7197970151901245), ('craneoencefalico', 0.6779731512069702), ('conocimiento', 0.6762552857398987), ('conmocion', 0.6442399621009827), ('politraumatismo', 0.6233476400375366), ('conocimento', 0.6212176084518433), ('occipital', 0.6101372838020325), ('p/c', 0.5979176759719849), ('pdc', 0.584796667098999), ('policontusionado', 0.5799985527992249)]</pre>
In [20]:	<pre>InfAltModel.most_similar(['dcha'])</pre>	In [16]:	<pre>InfAltModel.most_similar(['hta'])</pre>
Out[20]:	<pre>[('izda', 0.8290866613388062), ('derecha', 0.7282750606536865), ('izqda', 0.7253247499465942), ('dercha', 0.7189282178878784), ('decha', 0.7020336389541626), ('decha', 0.6982229948043823), ('decho', 0.6982229948043823), ('deh', 0.680919349193573), ('izquierda', 0.6796815395355225), ('izd', 0.6736931800842285), ('deercha', 0.666525661945343)]</pre>	Out[16]:	<pre>[('hipertensiva', 0.7690305113792419), ('hipertension', 0.6535224914550781), ('captopril', 0.6313796043395996), ('hipertesiva', 0.6241427659988403), ('htva', 0.6211124658584595), ('crisisi', 0.6208800077438354), ('anisedad', 0.605313777923584), ('anisedad', 0.6037868857383728), ('resuelta', 0.596153199672699), ('hta-', 0.5947197675704956)]</pre>

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Example: Identify, relations and similar words, to discover new words meaning





Example: Visualization Examples(Tensorflow project):

Embedding projector - visualization of high-dimensional data - Google Chrome 1 \blacksquare \blacksquare $07:01$ \clubsuit \bigcirc Home × \clubsuit Embedding project × \blacksquare \blacksquare $07:01$ \clubsuit \leftarrow \rightarrow C \bigcirc							
Embedding Projector			⊘ ≇				
DATA	Points: 101 Dimension: 300 Selected 101 points	Show All Isolate 10 points	01 Clear selection				
5 tensors found Word2Vec 10K	O	Search	.* word ~				
word Color by No color map	Count 1 Count 2 Count 1 Count 1 Count 1 Count 2 Count 3 C						
Sphereize data	Cbicitopenia Paresia/apraxia Gatinfarto Construction	Nearest points in the origination of the originatio	ginal space: 0.786 0.810				
Load data Publish Checkpoint: w2v-SK-s300-w5- Graph bit, tensor tay	eisrs neurologico radiculapatia contecederos chadds2 eradiculapatia tromboembolic chadds2 eragudizadolos eradiculapatia tromboembolic eragudizadolos	tepitu diagnostic -bicitopenia	0.836 0.842 0.845				
Metadata: w2v-SK-s300-w5-Graph.txt_	parestesia incomenzationa constructiona construction cons	=180 sencitivo encefalpatia	0.855 0.862 0.875				
X Y Component #1 - Component #2 -	eroncoencertaica edemencia egeneralizada eperferico-discop	-parestesia neurologicoepisodio antececemtes neurologico-	0.879 0.880 0.890 0.895				
Z Component #3 👻 PCA is approximate.	<pre></pre>	monopatia mesenterico sentitivo	0.902 0.905 0.907				
Total variance described: 19.6%.		BOOKMARKS (0) 🔮	^				



Evaluation:

Corpus Gold:

we generate a Corpus gold from the Emergency discharge clinical reports with the help of two Expert, using the Browser ihtsdotools to codify the reports. (http://browser.ihtsdotools.org/?)

We use Precision, recall and F_Measure to analyze the performance tool

Precision: P=TP/(TP+FP) Recall: R=TP/(TP+FN) $F_Measure = (1 + \beta^2) * \frac{precision * recall}{(\beta^2 * precision) + recall}$

 $oldsymbol{eta}=0.7$ To put more emphasis on precision than recall

TABLE I. TABLE MEASURES					
Concept	Our approach				
Precision	0.8097				
Recall	0.7469				
F-Measure	0.7879				



Conclusion and Outlook

This technology can use in many practical applications:

Future applications to develop:

- Semantic Search from free text Electronic Health Records.
- A tool assistant, to help the human expert, to assign the correct clinical id concept, from clinical reports.
- Discover new local words from a closed clinical domain.
- Identify and disambiguate abbreviations from a local clinical domain.
- Identify relations between type mistakes and the correct word.
- A new kind of visualization concept, using the vector Space Model.

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