



Manual annotation complexity grid

Karën Fort

karen.fort@univ-lorraine.fr / <https://members.loria.fr/KFort>

Dealing with the complexity of manual annotation

Analysing the complexity of an annotation campaign

What to annotate?

How to annotate?

Weight of the context

Synthesis: using the right tool, at the right place

WYMR: What You Must Remember

Manual annotation

Why is it easy/difficult? What part should be helped/automatized?

Parts of speech [Marcus et al., 1993] :

I/PRP do/VBP n't/RB feel/VB very/RB ferocious/JJ ./.

Manual annotation

Why is it easy/difficult? What part should be helped/automatized?

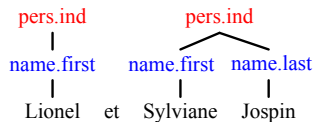
Gene renaming [Fort et al., 2012] :

The yppB gene complemented the defect of the recG40 strain. yppB and ypbC and their respective null alleles were termed recU and "recU1" (recU:cat) and recS and "recS1" (recS:cat), respectively. The recU and recS mutations were introduced into rec-deficient strains representative of the alpha (recF), beta (addA5 addB72), gamma (recH342), and epsilon (recG40) epistatic groups.

Manual annotation

Why is it easy/difficult? What part should be helped/automatized?

Structured named entities [Grouin et al., 2011] :



What is complex?



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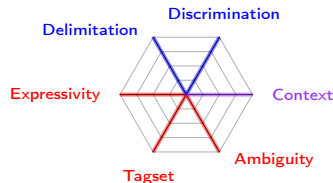
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Complexity dimensions

- ▶ 5 independent dimensions:
 - ▶ 2 related to the **localisation** of annotations
 - ▶ 3 related to the **characterisation** of annotations
- ▶ 1 not independent: the **context**



- ▶ Scale from **0** (null complexity) to **1** (maximal complexity) to allow for the comparison between campaigns
- ▶ Independent from the volume to annotate and the number of annotators

Example: gene renaming

1. Identification of gene names in the source signal:
*The **yppB** gene complemented the defect of the recG40 strain. **yppB** and **ypbC** and their respective null alleles were termed “**recU**” and “**recU1**” (recU:cat) and “**recS**” and “**recS1**” (recS:cat), respectively.*
2. Identification of gene couples expressing a renaming relation:
*The **yppB** gene complemented the defect of the recG40 strain. **yppB** and **ypbC** and their respective null alleles were termed “**recU**” and “**recU1**” (recU:cat) and “**recS**” and “**recS1**” (recS:cat), respectively.*

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Discrimination

Parts-of-speech [Marcus et al., 1993], pre-annotated :

I/PRP do/VBP n't/RB feel/VB very/RB ferocious/JJ ./.

Gene renaming [Fort et al., 2012], no pre-annotation:

The yppB:cat and ypbC:cat null alleles rendered cells sensitive to DNA-damaging agents, impaired plasmid transformation (25- and 100-fold), and moderately affected chromosomal transformation when present in an otherwise Rec⁺ B. subtilis strain. The yppB gene complemented the defect of the recG40 strain. yppB and ypbC and their respective null alleles were termed recU and “recU1” (recU:cat) and recS and “recS1” (recS:cat), respectively. The recU and recS mutations were introduced into rec-deficient strains representative of the alpha (recF), beta (addA5 addB72), gamma (recH342), and epsilon (recG40) epistatic groups.

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⇒ **more difficult** if the units to annotate are scattered, in particular if the segmentation is not obvious.

Discrimination

The discrimination weight is all the more high as the proportion of what *should* be annotated as compared to what *could* be annotated is low.

Definition

$$Discrimination(Flow) = 1 - \frac{|Annotations(Flow)|}{\sum_{i=1}^{LevelSeg} |UnitsObtainedBySeg_i(Flow)|}$$

⇒ Need for a reference segmentation

Parts-of-speech[Marcus et al., 1993] :

I/PRP do/VBP n't/RB feel/VB very/RB ferocious/JJ ./.

$$Discrimination_{PTB_{POS}} = 0$$

Gene renaming[Fort et al., 2012] :

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$$Discrimination_{Identification} = 0,9$$

$$Discrimination_{Renaming} = 0,95$$

Boundaries delimitation

- ▶ **extending** or **shrinking** the discriminated unit:
Madame Chirac → *Monsieur et Madame Chirac*

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- ▶ **extending** or **shrinking** the discriminated unit:
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- ▶ **decompose** a discriminated unit into several elements:
le préfet Érignac → *le **préfet** **Érignac***

Boundaries delimitation

- ▶ **extending** or **shrinking** the discriminated unit:
Madame Chirac → *Monsieur et Madame Chirac*
- ▶ **decompose** a discriminated unit into several elements:
le préfet Érignac → *le **préfet** **Érignac***
- ▶ or **group** together several discriminated units into one unique annotation:
Sa Majesté
le roi Mohamed VI → ***Sa Majesté le roi Mohamed VI***

Boundaries delimitation

Definition

$$Delimitation(Flow) = \min \left(\frac{Substitutions + Additions + Deletions}{|Annotations(Flow)|}, 1 \right)$$

$$Delimitation_{Identification} = 0$$

$$Delimitation_{Renaming} = 0$$

$$Delimitation_{PTB_{POS}} = 0$$

$$Delimitation_{EN_{TypesSubtypes}} = 1$$

$$Delimitation_{EN_{Components}} = 0,3$$

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Expressiveness of the annotation language

Definition

The degrees of expressiveness of the annotation language are the following:

- ▶ 0.25: type languages
- ▶ 0.5: relational languages of arity 2
- ▶ 0.75: relational languages of arity higher than 2
- ▶ 1: higher-order languages

$$Expressiveness_{Identification} = 0.25$$

$$Expressiveness_{Renaming} = 0.25$$

Dimension of the tagset

Person			Function		
<i>pers.ind</i> (individual person)	<i>pers.coll</i> (group of persons)		<i>func.ind</i> (individual function)	<i>func.coll</i> (collectivity of functions)	
Location			Production		
administrative (<i>loc.adm.town</i> , <i>loc.adm.reg</i> , <i>loc.adm.nat</i> , <i>loc.adm.sup</i>)	physical (<i>loc.phys.geo</i> , <i>loc.phys.hydro</i> , <i>loc.phys.astro</i>)	facilities (<i>loc.fac</i>), oronyms (<i>loc.oro</i>),	<i>prod.object</i> (manufactured object)	<i>prod.serv</i> (transportation route)	<i>prod.fin</i> (financial products)
		address (<i>loc.add.phys</i> , <i>loc.add.elec</i>)	<i>prod.doctr</i> (doctrine)	<i>prod.rule</i> (law)	<i>prod.soft</i> (software)
			<i>prod.art</i>	<i>prod.media</i>	<i>prod.award</i>
Organization			Time		
<i>org.adm</i> (administration)	<i>org.ent</i> (services)		<i>time.date.abs</i> (absolute date), <i>time.date.rel</i> (relative date)		<i>time.hour.abs</i> (absolute hour),
Amount		<i>time.hour.rel</i> (relative hour)			
<i>amount</i> (with unit or general object), including duration					

Types and sub-types used for structured NE annotation [Grouin et al., 2011]

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Level 1: *pers*, *func*, *loc*, *prod*, *org*, *time*, *amount* → 7 possibilities (degree of freedom = 6).

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Amount					
<i>amount</i> (with unit or general object), including duration					

Level 1: *pers*, *func*, *loc*, *prod*, *org*, *time*, *amount* → 7 possibilities (degree of freedom = 6).

Level 2: *prod.object*, *prod.serv*, *prod.fin*, *prod.soft*, *prod.doctr*, *prod.rule*, *prod.art*, *prod.media*, *prod.award* → 9 possibilities (degree of freedom = 8).

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Level 1: *pers*, *func*, *loc*, *prod*, *org*, *time*, *amount* → 7 possibilities (degree of freedom = 6).

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Level 3: *loc.adm.town*, *loc.adm.reg*, *loc.adm.nat*, *loc.adm.sup* → 4 possibilities (degree of freedom = 3).

Dimension of the tagset

Degree of freedom

$$\nu = \nu_1 + \nu_2 + \dots + \nu_m$$

where ν_i is the maximal degree of freedom the annotator has when choosing the i^{th} sub-type ($\nu_i = n_i - 1$).

Dimension (size) of the tagset

$$Dimension(Flow) = \min\left(\frac{\nu}{\tau}, 1\right)$$

where τ is the threshold from which we consider the tagset to be very large (experimentally determined).

$$Dimension_{Identification} = 0$$

$$Dimension_{Renaming} = 0.04$$

$$Dimension_{NE_{TypesSubtypes}} = 0.34$$

Degree of ambiguity: residual ambiguity

Using the traces left by the annotators:



[...] *<EukVirus>3CDproM</EukVirus> can process both structural and nonstructural precursors of the <EukVirus **uncertainty-type** = "too-generic"><taxon>poliovirus</taxon> polyprotein</EukVirus> [...].*

Définition

$$AmbiguityRes(Flow) = \frac{|Annotations_{amb}|}{|Annotations|}$$

$$AmbiguityRes_{Identification} = 0.04$$

$$AmbiguityRes_{Renaming} = 0.02$$

Degree of ambiguity: theoretical ambiguity

Proportion of the units to annotate that corresponds to ambiguous vocables.

Definition

$$AmbiguityTh(Flow) = \frac{\sum_{voc_i=1}^{|Voc(Flow)|} (Ambig(voc_i) * freq(voc_i, Flow))}{|Units(Flow)|}$$

with

$$Ambig(voc_i) = \begin{cases} 1 & \text{if } |Tags(voc_i)| > 1 \\ 0 & \text{else} \end{cases}$$

$$AmbiguityTh_{Identification} = 0.01$$

→ Does not apply to renaming relations

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Context to take into account

- ▶ **size of the window** to take into account in the source signal:

- ▶ The sentence:

I/PRP do/VBP n't/RB feel/VB very/RB ferocious/JJ ./.

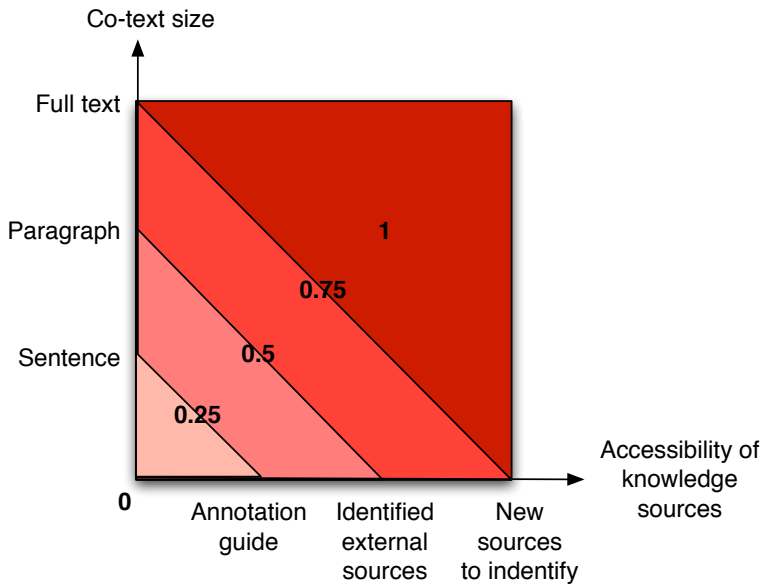
- ▶ ... or more:

Fabien Lévêque : C'est bien fait , avec **Gouffran** maintenant . **Gouffran** qui va tenter sa chance , et ça fait le but . Le but !
Xavier Gravelaine : Oh la la la la !
Fabien Lévêque : Et le but du plus breton des **Girondins** . C'est **Yoann Gourcuff** qui vient mettre un quatrième but ici au **stade de France** . Le cauchemar continue pour le **VOC** . Quatre à zéro en faveur des **Girondins** .



- ▶ number of **knowledge elements** to be rallied or degree of accessibility of the knowledge sources that are consulted:
 - ▶ annotation guidelines
 - ▶ nomenclatures (Swiss-Prot)
 - ▶ new sources to be found (Wikipedia, etc.)

Weight of the context



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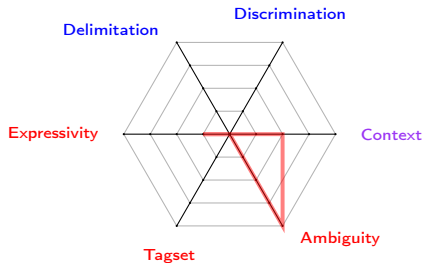
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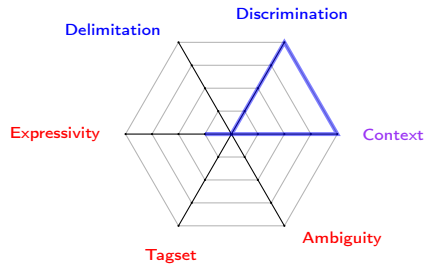
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Synthesis of the complexity dimensions



Classification of it pronouns as anaphoric or impersonal



Gene names identification



Manual annotation and NLP:

- ▶ usage
- ▶ cost

Manual annotation:

- ▶ definition
- ▶ organization
- ▶ complexity grid

Practice: Compute the complexity dimensions for your annotations

Assignment: Reading a research paper (graded)

Read carefully and email your teacher the synopsis (2p max, pdf) by next week

[Dandapat et al., 2009] :

Cheap and Fast – But is it Good?

Evaluating Non-Expert Annotations for Natural Language Tasks

<https://aclanthology.org/D08-1027.pdf>

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