"CDG LAB": a Toolbox for Dependency Grammars and Dependency Treebanks Development

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Abstract

We present "CDG LAB", a toolkit for development of dependency grammars and treebanks. It uses the Categorial Dependency Grammars (CDG) as a formal model of dependency grammars. CDG are very expressive. They generate unlimited dependency structures, are analyzed in polynomial time and are conservatively extendable by regular type expressions without loss of parsing efficiency. Due to these features, they are well adapted to definition of large scale grammars. CDG LAB supports the analysis of correctness of treebanks developed in parallel with evolving grammars.

1 Introduction

There are two main technologies of automatic syntactic analysis of natural language:

1. grammatical parsing i.e (symbolic or statistical or mixed) parsing of a hand-crafted grammar belonging to a family of formal grammars disposing of a general purpose parser;

2. data-driven parsing, i.e. parsing with statistical parsers trained over annotated data.

Both technologies need a large amount of expensive expert linguistic data. The hand-crafted wide coverage grammars are notoriously expensive and only very few of them have been successfully realized and applied to unrestricted material (cf. (Bouma et al., 2000; Riezler et al., 2002)). Besides this, they are prone to explosion of spurious ambiguity when parsed with general purpose parsers. On the other hand, training of statistical parsers needs voluminous high quality treebanks such as the Penn Treebank (Marcus et al., 1993). Training data of this size and quality are in fact as expensive as the hand-crafted grammars and also need a long-term hand work. Even if the results obtained in the statistical parsing during the last fifteen years are very encouraging, their quality and adequacy depends on those of the hand-crafted annotated data. This is a vital issue for dependency grammars which suffer from the shortage of high quality training data. The several existing dependency treebanks (DTB) such as the Prague Dependency Treebank of Czech (Hajicova et al., 1998), the TIGER treebank of German (Brants and Hansen, 2002) or the Russian treebank (Boguslavsky et al., 2000) only partially solve the problem. First of all, they serve for particular languages. Secondly, even for these languages, the DTB use a particular inventory of dependency relations. At the same time, there is no consensus on such inventories. So the DTB are dependent on the choice of underlying syntactic theories, which makes their reuse problematic. The translation technologies (cf. (Hockenmaier and Steedman, 2007)) consisting in acquisition of dependency structures from high quality constituent structure treebanks also do not resolve the problem because, for technical reasons, they often flatten the genuine dependency structures and introduce into them multiple distortions. For all these reasons, there is a need in efficient and inexpensive methods and tools of development of wide coverage grammars and of training corpora.

Below we present "CDG LAB", a toolkit supporting parallel development of wide coverage dependency grammars and of DTB. It uses Categorial Dependency Grammars (CDG) as a formal model of dependency grammars.

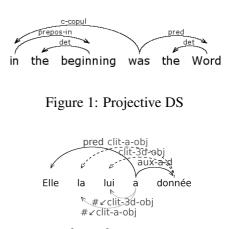
The CDG, a class of first-order type categorial grammars generating unlimited dependency structures (DS), were introduced in (Dikovsky, 2004). Since then, they were intensively studied (e.g., see (Dekhtyar and Dikovsky, 2004; Béchet et al., 2004; Dekhtyar and Dikovsky, 2008; Dekhtyar et al., 2010; Béchet et al., 2010)). CDG are very expressive. In particular, very simple CDG generate such non-CF languages as $L^{(m)} = \{a_1^n a_2^n ... a_m^n \mid n \ge 1\}$ for all m > 0 and $MIX = \{w \in \{a, b, c\}^+ \mid |w|_a = |w|_b = |w|_c\}$. They are equivalent to real time pushdown automata with independent counters (Karlov, 2008). Interesting sufficient conditions of learning CDG in the limit were found recently (Béchet et al., 2004; Béchet et al., 2010; Béchet et al., 2011).

CDG also have important advantages which make them a convenient and natural means of definition of wide coverage dependency grammars. First, they are completely lexicalized, as it is the case of all categorial, and more generally, type logical grammars (Bar-Hillel et al., 1960; Lambek, 1961; Lambek, 1999; Steedman, 1996). Second, the CDG types directly encode DS with repeatable and unlimited non-projective dependencies (see below). Third, they are parsed in a polynomial time (Dekhtyar and Dikovsky, 2004; Dekhtyar and Dikovsky, 2008). Fourth, an extension of CDG by regular type expressions (RTE) specially designed for large scale grammars is defined (Dikovsky, 2009; Dikovsky, 2011) and is implemented in the CDG parser presented below. Moreover, for this extension there is a supported by CDG LAB method of incremental bootstrapping of large scale grammars from dependency structures (Dikovsky, 2011).

The plan of this paper is as follows. Section 2 presents the basics of CDG and of their extension by RTE. Then the architecture and the main functionalities of CDG LAB are described in Section 3.

2 Categorial Dependency Grammars

CDG define projective DS (as in Fig. 1) i.e. DS in which dependencies do not cross, and also non-projective DS, as in Fig. 2, in which they may cross. In these graphs, the nodes correspond to the words of the sentence (their precedence order in the sentence is important) and the arcs represent the dependencies: named binary relations on words. Formally, a DS of a sentence x is a linearly ordered cycle-free graph with labelled arcs and the words of x as nodes. We consider connected DS with the root node. When in a DS D there is an arc $w_1 \stackrel{d}{\longrightarrow} w_2$, we say that d is a dependency between w_1 and w_2 , w_1 is the governor and w_2 is subordinate to w_1 through d. E.g., in is subordinate to was in Fig. 1 and donnée governs la through clit - a - obj and lui through clit - 3d - obj.



(fr. *she it[fem.] to him has given)

Figure 2: Non-projective DS

As all categorial grammars, the CDG are completely lexicalized and may be seen as assignments of types to words in a dictionary W. CDG types are expressions of the form

 $t = [l_1 \backslash l_2 \backslash \ldots \backslash H / \ldots / r_2 / r_1]^P.$

A type assigned to a word $w \in W$ defines its dependencies in a rather straightforward way: its subtypes $H, l_1, l_2, \ldots, \ldots, r_2, r_1$ represent the claims for w to be related to other words through projective dependencies and P, called potential of t, defines all non-projective dependencies of w. The head subtype H, claims that w should be subordinate to a word through dependency H. When w should be the root of a DS, H = S (S is a special symbol called axiom). The left subtypes l_1, l_2, \ldots define the left projective dependencies of w (i.e. the dependencies through which w governs the words occurring in the sentence on its left). The right subtypes \ldots, r_2, r_1 define the right projective dependencies of w. For instance, the projective DS in Fig. 1 is uniquely defined by the type assignment:

 $in \mapsto [c-copul/prepos-in], the \mapsto [det],$ $Word \mapsto [det \ pred], beginning \mapsto$ $[det \ prepos-in], was \mapsto [c-copul \ S/pred].$

Left and right subtypes may also be iterated. The iterated subtypes define repeatable dependencies. E.g., $l_i = d^*$ means that w may have on it left 0, 1, 2, ...occurrences of words subordinate to it through dependency d. We also use optional subtypes $l_i = d$?. Assignment of type $[d? \setminus \alpha]$ is equivalent to assignment of $[d \setminus \alpha]$ and $[\alpha]$. E.g., the DS in Fig. 3 is defined by the type assignment:

 $she \mapsto [pred], was \mapsto [pred \setminus S/a_copul^*],$

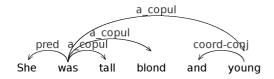


Figure 3: Iterated dependency

tall, blond, young \mapsto [coord-conj?\a_copul], and \mapsto [coord-conj].

By the way, the coordination scheme used is this analysis is more general than the traditional one, where the coordinated phrase is subordinate to the conjunction. The new scheme applies to the traditional recursive definition of coordination as well as to the iterative one, as in this sentence.

The potential P is the concatenation of so called polarized valencies of w. The polarized valencies are of four kinds: $\checkmark v, \checkmark v$ (negative) and $\nwarrow v$, $\nearrow v$ positive. E.g., if a word w has valency $\swarrow v$, this intuitively means that its governor through dependency v must occur somewhere on the right. Two polarized valencies with the same valency name v and orientation, but with the opposite signs are dual. Together they define non-projective dependency v. The order of polarized valencies in the potential P is irrelevant (so one may choose a standard lexicographic order). For instance, in the DS in Fig. 2, the potential $\leq clit - a - obj \leq clit - 3d - obj$ of the participle donnée means that it needs somewhere on its left a word subordinate through dependency clit - a - obj and also another word subordinate through dependency clit - 3d - obj. At the same time, the accusative case clitic la (it[fem.]) has potential $\sqrt{clit} - a - obj$ and the dative case clitic *lui* (*to him*) has potential $\checkmark clit - 3d - obj$. The proper pairing of their dual valencies with those of the participle defines two non-projective dependencies between the participle and its cliticized complements.

We use CDG with non-empty head subtypes. When a type $[\alpha \backslash d/\beta]^P$ has a negative valency in its potential P, say $P = \swarrow vP'$, the word w with this type has two governors: one through v, the other through d. For such cases we use special head subtypes d = #(A), called anchors, to express the adjacency of w to a host word w_0 . The anchor dependencies are displayed below the sentence for a better readability. E.g., the DS in Fig. 2 is defined by the following types assignment:

$$\begin{aligned} elle &\mapsto [pred], \\ la &\mapsto [\#(\swarrow clit - a - obj)]^{\checkmark clit - a - obj}, \\ lui &\mapsto [\#(\swarrow clit - 3d - obj)]^{\checkmark clit - 3d - obj}, \\ donnée &\mapsto [aux - a - d]^{\land clit - a - obj^{\land} clit - 3d - obj}, \\ a &\mapsto [\#(\swarrow clit - 3d - obj) \setminus \#(\swarrow clit - a - obj) \setminus pred \\ &\setminus S/aux - a - d]. \end{aligned}$$

Due to the anchor subtypes $\#(\swarrow clit - 3d - obj)$, $\#(\swarrow clit - a - obj)$ in the type of the auxiliary verb a (has), it serves as the host verb for both clitics and also defines their precedence order.

Derivability of DS in CDG is formalized through the following calculus ¹ (with C being a dependency, H being a dependency or an anchor and V being a polarized valency):

 $\mathbf{L}^{\mathbf{l}}. H^{P_1}[H \setminus \beta]^{P_2} \vdash [\beta]^{P_1P_2}$ $\mathbf{I}^{\mathbf{l}}. C^{P_1}[C^* \setminus \beta]^{P_2} \vdash [C^* \setminus \beta]^{P_1P_2}$ $\mathbf{O}^{\mathbf{l}}. [C^* \setminus \beta]^{P_1} \vdash [\beta]^{P_2}$

 $\mathbf{\Omega}^{\mathbf{l}} \cdot [C^* \backslash \beta]^{P} \vdash [\beta]^{P}$

D¹. $\alpha^{P_1(\swarrow V)P(\nwarrow V)P_2} \vdash \alpha^{P_1PP_2}$, if the potential $(\swarrow V)P(\nwarrow V)$ satisfies the following pairing rule **FA** (*first available*):²

FA: *P* has no occurrences of $\checkmark V, \diagdown V$. **L**¹ is the classical elimination rule. Eliminating the argument subtype $H \neq \#(\alpha)$ it constructs the (projective) dependency *H* and concatenates the potentials. $H = \#(\alpha)$ creates the anchor dependency. **I**¹ derives k > 0 instances of *C*. Ω^1 serves for the case k = 0. **D**¹ creates non-projective dependencies. It pairs and eliminates dual valencies with name *V* satisfying the rule **FA** to create the non-projective dependency *V*.

Let us see how DS in Fig. 2 can be derived using the type assignment shown above.

First, we may eliminate the left anchor subtype $\#(\swarrow clit - 3d - obj)$ in the type of the auxiliary verb *a* using the type of the clitic *lui*. As a result, we generate the anchor dependency $\#(\swarrow clit - 3d - obj)$ from *a* to *lui* and the derived type of the string *lui a* becomes

 $[\#(\swarrow clit-a-obj) \ pred \ S/aux-a-d] \ clit-3d-obj$. In this type, we may eliminate the anchor subtype $\#(\swarrow clit - a - obj)$ using the type of the clitic *la*. This will generate the anchor dependency $\#(\swarrow clit - a - obj)$ from *a* to *la*. The derived type of the sequence *la lui a* is $[pred \ S/aux - a - d] \ clit-a-obj \ clit-3d-obj$. Now we may eliminate the left subtype *pred* of the derived type using the type of the subject *elle*. This generates the projective dependency from *a* to *elle* and

¹We show left-oriented rules. The right-oriented rules are symmetrical.

²Cf. a different pairing rule mentioned below.

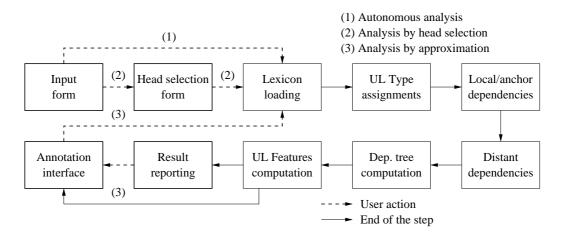


Figure 4: Architecture of the CDG parser

elle la lui a obtains the derived type [S/aux $a-d \sqrt[]{clit-a-obj}{clit-3d-obj}$. Then using the type of the participle donnée, we may eliminate the right subtype of this type. This generates the projective dependency aux - a - d from a to donnée and assigns to the sentence the derived type $[S] \checkmark clit-a-obj \checkmark clit-3d-obj \land clit-a-obj \land clit-3d-obj$. Application of the rule FA to the dual valencies $\checkmark clit - 3d - obj$ and $\nwarrow clit - 3d - obj$ generates the non-projective dependency clit-3d-obj from donnée to lui and derives for the sentence the type $[S]^{\checkmark clit-a-obj}$. Finally, applying this rule to the dual valencies $\sqrt{clit} - a - obj \wedge clit - a - obj$ we generate the DS in Fig. 2 because the nonprojective dependency clit - a - obj from $donn\acute{e}$ to *la* is generated and the derived type is S.

Extended CDG. CDG is a theoretical model not adapted to wide coverage grammars. Wide coverage grammars face a combinatorial explosion of spurious ambiguity and many other hard problems, e.g. those of compound lexical entries including complex numbers, compound terms, proper names, etc. and also that of flexible precedence order. (Dikovsky, 2009) proposes an extension of CDG adapted to wide coverage grammars.

The extended CDG use classes of words in the place of words and use restricted regular expressions defining sets of types in the place of types. I.e., the dictionary W is covered by classes:

 $W = \bigcup_{i \in I} C_i$ and the lexicon λ assigns sets of regular expressions to classes. At that:

- all words in a class C share the types defined by the expressions assigned to C,

- every word has all types of the classes to which it belongs.

The extended CDG use flat (i.e. bounded depth) regular type expressions (RTE). In these expressions, C, C_i are dependency names or anchors, B is a primitive type, i.e. a dependency name, or an anchor or an iterated or optional type, and H is a choice.

<u>Choice</u>: $(C_1|\ldots|C_k)$. (C) = C. Optional choice: $(C_1|\ldots|C_k)$?. (C)? = C?.

<u>Iteration</u>: $(C_1|...|C_k)^*$. $(C)^* = C^*$.

Dispersed subtypes expressing flexible order.

 $\underline{\text{Left}}: [\{\alpha_1, B, \alpha_2\} \setminus \alpha \setminus H/\beta]^P. \\
\underline{\text{Right}}: [\alpha \setminus H/\beta / \{\alpha_1, B, \alpha_2\}]^P. \\
\underline{\text{Two-way}}: \{\alpha_1, B, \alpha_2\} [\alpha \setminus H/\beta]^P.$

Intuitively, the choice unites several alternative types into one. When iterated, it represents all sequences of the elements of the choice occurring in the same argument position. On the contrary, assignment of the type $[\{\alpha_1, B, \alpha_2\} \setminus \alpha \setminus H/\beta]^P$ to a word w means that a word subordinate to w through projective dependency B is somewhere on its left. E.g. the assignments $w_0 \mapsto [\{d\} \setminus b \setminus a \setminus S], w_1 \mapsto [a], w_2 \mapsto [b], w_3 \mapsto [d]$ define DS of sentences: $w_3w_1w_2w_0, w_1w_3w_2w_0, w_1w_2w_3w_0$ in which $w_0 \stackrel{d}{\longrightarrow} w_3, w_0 \stackrel{a}{\longrightarrow} w_1$ and $w_0 \stackrel{b}{\longrightarrow} w_2$. The right dispersed RTE is similar. The two-way dispersed RTE claims that an element of type B were found in some left or right position.

As the original CDG, the extended CDG are formalized by a calculus wich has special rules for every kind of RTE (see (Dikovsky, 2009; Béchet et al., 2010; Béchet et al., 2011)). A fragment of this calculus may be seen in (Dikovsky, 2011) (see this vomume).

Classes and RTE do not extend the expressive power of CDG. At the same time, they dramatically reduce the grammar size. Due to the extended type calculus they can be parsed directly, without unfolding. In fact, the polynomial time parsing algorithm of (Dekhtyar and Dikovsky, 2008) can be adapted to the extended CDG.

3 CDG LAB

CDG LAB is a kit of tools supporting parsing with extended CDG, development and maintainance of dependency treebanks (DTB) and development and test of large scale extended CDG. The core element of CDG LAB is the parser of the extended CDG implemented in Steel Bank Common Lisp. Recently was issued its version 3.1 (below we will call this parser Parser-3.1).

All input and output data of Parser-3.1 are XML-structures. It may analyse sentences, text corpora and DTB either with an extended CDG integrated with an external morpho-syntactic dictionary (lexical base grammar) or only with the internal grammar lexicon (text grammar). For instance, for French is developed a large scale extended CDG (Dikovsky, 2011). Its version 3.2 (called below "French Text CDG") is integrated with the open MS-dictionary of French Lefff 3.0 (Sagot, 2010) containing 536,375 lexical units (LU). Lefff is kept in object-relational database PostgreSQL and a correspondence between the classes of the French Text CDG and the categories of Lefff is implemented through several hundreds of SQL queries. The integral grammar is called below "French LB CDG".

Modes of Analysis. Parser-3.1 is multi-purpose. It is used for semi-automatic analysis by consecutive approximations and also starting from head type/class selection, for estimation of compatibilty of analyses with an updated grammar and for expert annotation of DS needed for this estimation, for automatic re-analysis of sentences when the annotation is changed and also for autonomous syntactic analysis. Respectively, the parser is used in different modes:

- analysis by head selection,
- analysis by approximations,
- DS analysis,
- autonomous analysis.

Fig. 4 shows a scheme of functioning of Parser-3.1 in these modes. Sentences are introduced through the input form (see Fig. 5). Through this form, the User may set various parameters, e.g. the maximal parsing time, the maximal number of DS to re-



Figure 5: Query Form

turn, a graphical representation of DS, a language register (corresponding to specific choices of nonprojective dependencies common to official documents or to scientific or literary prose, to periodicals or to the spoken language), etc. The input sentence is lexically analysed. Composite forms are decomposed into separate tokens, as in the case of l'homme (the man), which is segmented into three tokens: l, ' and homme, for which are found in the lexicon all possible lemmas. In this example, the association is ambiguous: l' may be a clitic or a determiner. All possible variants of composite LU are detected (in particular, complex numbers and names recognized through regular expressions, multi-word LU, such as à la (of the kind), à travers (through) etc.) and unknown terms are identified. The transitions to and from head selection form are followed only in the mode of Analysis by head selection. Functioning in the mode of Analysis by approximations is iterative. It goes round result reporting and passes from annotation interface directly to lexicon loading. Other transitions are common for all modes. So we comment them in the head selection mode.

Analysis by Head Selection. In this mode a selection form is proposed (see Fig. 6), in which the User may select the proper composite LU (if and when several possibilities are detected) and for every LU, to select one of possible classes and one of possible dependency relation groups

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	obj	O aux-A O aux-a O aux-a-A O aux-a-d O aux-a-g O aux-a-l

Figure 6: Selection Form

(or of their elements). The selected dependency is nothing but the head subtype of the possible types of the LU. The latter selection is in fact decisive. It corresponds to the strong constraint that the LU is subordinate through the selected dependency (or limits the choice of such dependencies to the elements of the selected group).

This selection drastically limits the searchspace. As it concerns the Parser-3.1 and the French LB CDG, it reduces the number of possible analyses by two-three orders of magnitude, i.e. in the place of a thousand of possible DS only about ten are found, most often differing between them in positions of repeatable dependencies (such as *modif* (modifier), *attr* (attribute) or *circ* (circumstantial). E.g., in this example the genuine DS will be immediately found due to this selection (see Fig. 7).

Then is created the sentence's workspace (WS), an XML-structure representing the subgrammar corresponding to all detected LU to which are affected the classes, the types and the features' values compatible with the pre-selection. After this follow the steps common to all modes. First, all possible projective (and anchor) dependencies are computed and registered in the triangular matrix

. This computation is done by an CKY-like al-

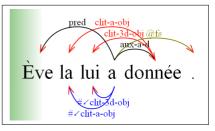


Figure 7: Resulting DS

gorithm adapted to extended CDG. Then all possible pairings of dual valencies providing nonprojective dependencies are independently computed and registered in the resulting matrix (more precisely, in the submatrix in which the axiom Scan be proved). It should be noted that the pairing principle may be chosen for a CDG and for a particular non-projective dependency. By default, this is the **FA** rule. But in rare exceptional situations, such as that of unlimited cross-serial dependencies in subordinate clauses in Dutch, one may use a different rule FC (first cross) defined in (Dikovsky, 2007). This independence of computations of projective and non-projective dependencies is founded on the fundamental projection independence property of CDG proved for the rule FA in (Dekhtyar and Dikovsky, 2004; Dekhtyar and Dikovsky, 2008) and for rule FC in

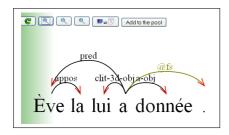


Figure 8: Incorrect DS

(Dikovsky, 2007). Till the end of this step the parsing algorithm is polynomial. The resulting triangular matrix is in fact a packed chart from which it is possible to enumerate all possible DS of the sentence. Given that the number of these DS may be exponential with respect to the size of the matrix, the next step is exponential in space in the worst case. In this step, the DS are generated from the matrix in a certain order and the feature values are assigned to LU in every generated DS. Finally, the parser generates the HTML report page, which includes various useful statistics. An XML structure representation of every DS including all necessary information, in particular the CDG classes and the feature values is also generated and saved to be used by other programs.

Analysis by Approximations. This important mode represents another User-guided strategy of parsing. It allows to find the needed DS starting from any obtained DS by consecutive approximations computed from User's annotations in the DS. There are three possible annotations of dependency relations: positive, negative and neutral. The positively annotated dependencies are those adequate. They will be kept during the whole sequence of approximations (if not discarded). The neutrally annotated dependencies are kept till they are compatible with the positively annotated ones. The negatively annotated dependencies are to be eliminated from the DS. When used in this mode, the Parser computes for every DS the total number of positively annotated dependencies and that of negatively annotated dependencies. The obtained DS are sorted first by the negative annotations' weight (the less negative annotations the better) then by the positive annotations' weight (the more positive annotations the better).

Suppose, that the approximations start from the (partially incorrect) DS of the sentence Eve la lui a donnée

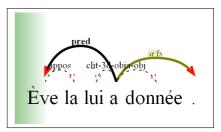


Figure 9: First annotated DS

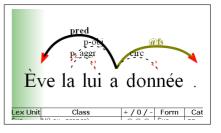


Figure 10: Next approximation

(Eve it[fem.] to him has given) shown in Fig. 8. There are only two correct dependencies in this DS: the predicative one: pred and the punctuation dependency @fs. We annotate both positively (this annotation being displayed by boldface arcs). The other three dependencies appos, clit - 3d - obj and a - obj are erroneous. We annotate them as negative (which is displayed by broken arcs). So we obtain the first annotated DS shown in Fig. 9.

From this annotated DS, the Parser computes the next approximation shown in Fig. 10, which is also incorrect. It has the same two correct dependencies and three other incorrect: p-obj, p-aggrand *circ*. We annotate the three as negative, as it is shown in Fig. 10.

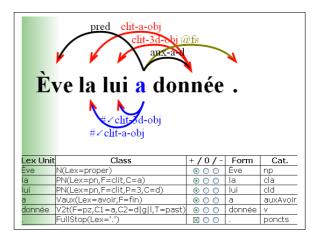


Figure 11: Final approximation

From this annotated DS, the parser finally computes the right one shown in Fig. 11. Not only this final approximation is correct, but it is also annotated as such. This difference is very important for the other mode of use of Parser-3.1, that of DS analysis.

DS, DTB and Grammar Analysis. In fact, Parser-3.1 considers every DS as annotated. The case where there are no annotations is considered as that with weight 0. Moreover, not only the dependencies, but also the LU may be annotated. The LU may have only two annotations: positive and neutral. Annotating a LU w positively is equivalent to positively annotate all dependencies in the sub-structure with the root w. This is seen in Fig 11, were the positive annotation of the root (displayed in a contrasting color) implies the positive annotations of dependencies (displayed in boldface). More than that, the class and the feature values assigned to every LU in DS may also be annotated as positive, negative or neutral. In the fragment of the class/feature table shown in Fig 11, one may see that not only the dependencies, but also the class/feature assignments for its LU are all annotated as positive. So this analysis is 100% correct. It is using this integral annotation weight, that Parser-3.1 evaluates the DS. Now, for two DS of the same sentence, it is possible to measure the difference of their weights. This simple measure turns out to be an efficient means of analysis of DTB and of the CDG used while their development. Every sentence processed by Parser-3.1 using French LB CDG obtains its status. The status includes the analysis result ('NO', when there are no parses, 'YES' otherwise) and for a parsable sentence, also the maximal number of returned DS and the difference (in percents) of annotation weights (of dependencies and of LU) between the best obtained DS and the one present in the DTB (if any) before this sentence processing. When the grammar is updated, the DS of sentences in the DTB become potentially irrelevant. For this case, Parser-3.1 has a special function of re-parsing of a DTB, which computes the difference between the DS before and after update, comparing their statuses. The User may choose between keeping or not the same head subtypes while re-parsing. Using this function, one may easily find all sentences to be revised.

0.409 DEFAULT ≥ 1 100.09 arse group 829. ≥ 1 100.09 Ok group 830 YES 0.136 ≥ 1 100.09 e aroup 831 $\geq 1\,100.0^{\circ}$ arse group 832 arse group 833. 0.20 OK YES $\geq 1|100.0$ 31.341 arse aroup 834 YES $\geq 1\,100.0$ group 83 0.22 Oł 0.23 se group 836 Ok droup 837 0.20 Ok 0.0% Ok YE 0.0% arse aroup 838 0. e group 839 0.488 Ok YES ≥ 5 77.

Figure 12: Re-parse results

grammar itself. The French Text CDG was created using the Structural Bootstrapping Method (Dikovsky, 2011), a method specific to the extended CDG and consisting in an incremental transformation of DS of a sample of sentences σ into an extended CDG $G(\sigma)$ generating these sentences. The incrementality is interpreted in the strong sense: $\Delta(G(\sigma)) \subseteq \Delta(G(\sigma \cup \{s\}))$ for every new sentence s. So such transformations represent monotone grammar updates. Basically, the bootstrapping of French Text CDG was incremental in this sense, except three important revisions which were not. Taking in mind the size and the complexity of this grammar (it consists of more than 3120 RTE distributed between 185 lexicon classes, has 84 projective and 20 non-projective dependencies), it was a very hard task to find all sentences in the sample wrongly analysed using the updated grammar. Indeed, to find them, it was necessary to look through thousands of DS of hundreds of sentences in order to find linguistically adequate DS (the simple existence of a generated DS is of course not sufficient). The situation has completely changed after the implementation of Parser-3.1. Indeed, now all sentences in the sample are initially annotated. The procedure of re-parsing of the sample finds all inconsistencies. When a DS is compatible with the grammar before update (i.e. has 100 % correct annotations) and becomes incompatible after the update, then the dependencies with changed weight of annotation correctness indicate (together with the word classes) the RTE of the grammar to be updated.

In Fig. 12 we show a fragment of the table representing the results of re-parsing applied to a DTB. In this table:

- the first column is the reference to the DS of a sentence,

- the second column shows the (folded) characteristics of parsing complexity,

Another way round, this test applies to the

- in the third column, 'OK' means that all LU of

the sentence are present in the grammar lexicon and 'DEFAULT' means that there is at least one LU absent in the lexicon and replaced by the default unit,

- in the fourth column, 'YES' means that the reparsing was successful in the sense that a successful analysis was found and 'NO' means the contrary,

- in the fifth column is given the maximal number of DS requested for re-parsing ($\geq k$ means that there are at least k requested DS),

- the sixth column shows the part (in percents) of annotation status coincidence of the DS before and after the update (so it is 0% for new sentences).

Autonomous Analysis. This is the mode of non-User-guided analysis. Parser-3.1 may be used as a general purpose parser for the extended CDG. In CDG LAB, there is a possibility to upload one's own grammar or to introduce it through the Sandbox. Parser -3.1 is not well adapted to parsing with the grammars of such size and as amiguous as French LB CDG. For instance, it passed 2373 seconds when applied to a test set of 559 French sentences of various complexity, repesenting the majority of French syntactic constructions. In so doing, it failed on 157 sentences exceeding the time limit of 10 seconds and successfully analysed the other 402 sentences. To compare: in the reparsing mode, Parser -3.1 successfully analysed in 175 seconds 1442 DS of sentences, some of which are extremely long (up to 73 words). Another problem with Parser-3.1 is that it generates the DS not in the order of their adequacy. With ambiguous CDG, such as French LB, it generates hundreds of spurious structures per sentence. So for very long and complex sentences, it is practically impossible to know whether an adequate DS was computed. This is why we consider Parser-3.1 as a tool of development of DTB using head subtype selection, approximations and re-parsing. In these modes it performs very well and doesn't impose any length limits on sentences. A higher-performing autonomous mixed stochastic-symbolic parser of extended CDG is under design.

DTB Development. The annotation based development of DTB in CDG LAB leads to a notable change in the point of view on the quality of treebanks. It is now the grammar, implementing a set of linguistic subjective expert knowledge, which will serve as the "gold standard". As to the

DTB, they should all be correct with respect to the grammar and should be tested for correctness after every non-monotone grammar update. By definition, the monotone grammar updates preserve correctness of DS.

Besides the means based on DS annotation, CDG LAB also has rather standard means for creation and updates of DTB and for search of DS by projective and non-projective dependency names and by LU in the sentences.

Grammar Development. Besides the described above general purpose means supporting nonmonotone grammar updates, CDG LAB has some means specific for CDG of French integrated with Lefff 3.0. In particular, it has several functions for completion of the lexicon of these CDG. Basically, there are two problems: the first is to automatically complete the lexicon by all forms of a missing word (this concerns mainly the verbs), the second is to compute the government pattern of a missing word from that of a present word. For the former problem, CDG LAB has several functions based on updates of the lexicon of French Text CDG. The latter problem mainly concerns the deverbals. The work on completions of this kind is in progress.

4 Conclusion

CDG LAB combines several means of incremental parallel development of wide coverage dependency grammars and of dependency treebanks provably correct with respect to the grammars. These means were successfully tested in the course of development of a wide coverage categorial dependency grammar of French and of an experimental dependency treebank. Some of these means are general purpose. E.g., the annotation weight difference test applies to any kind of structural incremental development based on expert annotations. Some other, such as head subtype selection and consecutive approximations, may be used with other classes of dependency grammars and may be implemented in tabular dependency grammar parsers. Some means are specific to the Parser-3.1 and to the French CDG integrated with Lefff. Several important means of CDG LAB are still under construction, but even this experimental version has proved its high efficiency.

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