GraphAnno

An annotation and query tool for graph-based linguistic annotations

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1 The annotation graph

1.1 Graph format

A graph in GraphAnno consists of a set of nodes and directed edges. Nodes and edges bear attributes in the form of key-value pairs, that are used for the linguistic annotation as well as the structuring and visualization of the graph. There are some attributes – like the attribute type, that differentiates the types of nodes and edges (annotation nodes, tokens, sentence nodes; annotation edges, ordering edges, section edges) – that serve the internal representation of the graph and are not edited directly. Under the key attr, however, there is a group of key-value pairs which serve the linguistic annotations and the representation of layers and which are edited by the user directly. They will be called *annotation attributes* in the following.

The graph in GraphAnno is segmented in sentences – units that that are used for structuring and displaying the corpus, but that do not necessarily correspond to real sentences (how ever these may be defined). There is a sentence node (bearing the attribute type:s) for every sentence, that bears information concerning the whole sentence. This includes, e.g., source, medium, speaker or the like. The annotations of this node is displayed in the light grey area below the sentence graph and under the text of the sentence (blue font) in black font. The information to which sentence a node of the graph belongs is represented by an (invisible) edge with type:s that links the node in question to the sentence node. Edges are not linked to a sentence node; their affiliation follows from the affiliation of their start and end nodes.

Token nodes are characterized by the attribute type:t and they carry the annotation attribute token with the token text as value. The order of the tokens of a sentence (as well as the sentence nodes themselves) is defined by ordering edges (with type:o) that link each node to its successor. These edges are needed for the correct visualization and for traversal; the are, however, not displayed and cannot be manipulated directly.

Among the annotation attributes, nodes and edges may bear the cat attribute, that has no special meaning for the graph, but which is desplayed prominently on top of the other annotations and without the key. Nodes and edges show numbers like t23 for tokens, n23 for other nodes and e23 for edges, that are used for referencing the elements in annotation commands. These numbers are not stored in the graph model but generated dynamically each time the displayed graph is rendered.

GraphAnno also provides the possibility to assign nodes and edges to different layers. The fact that a node or edge is affiliated to a certain layer is represented by that element

bearing an annotation (among the annotation attributes) with the key corresponding to the layer and the value t (for *true*). This representation allows for elements to belong to more than one layer, with the consequence that layers my overlap in an arbitrary manner. Nodes and edges of different layers may be displayed in different colors and aligned in different ways (hierachically or horizontally). Tokens do not belong to any layer; they are displayed in black by default.

1.2 Configuration

1.2.1 Layers and visualization

The window for configuration of the layers of the currently loaded graph and its visualization is opened with the command line command config.

In the section *general settings*, you can configure the settings for nodes and edges that do not belong to any layer.¹ *Default color* applies to all nodes and edges that are not tokens and that do not belong to any layer, *token color* applies to tokens, *found color* is used for the highlighting of nodes and edges found in a search, and *filtered color* for elements that are filtered out by the filter function. The setting for *edge weight* affects the layout of the displayed graph and shows its effect only when edges with different weight are present (details will follow in the next paragraph).

In the section layers, you can configure the layers of the graph. The name is an arbitrary label for the layer, that will be shown in the dropdown field for the layer selection. Attribute is the attribute that will be set to t for elements that belong to the layer in question. The shortcut is an identifier that can be used for the annotation of elements (cf. 3.2.1 or ??); this shortcut may only consist of alphanumeric characters and underscores and must not have the same structure as element references (i.e., t, n or e followed by a number, or m; cf. 3.2.3). Color means the color that is used for displaying the elements of the layer; edge weight is the weight of the layer's edges. The heigher this value (integer values), the shorter the rendering algorithm will try to make the edges. When you create two layers, one with a high edge weight and one with a low edge weight, the graph will be rendered such that the graph of the first layer is as compact as possible; the elements of the second layer will be placed in a way that they distort the first layer only to a low degree. If you enter 0 as edge weight, the edges will not enforce a hierarchy between the nodes of that layer (otherwise, the start node of an edge is always placed higher than the end node). If a negative value is entered, all nodes of that layer are displayed horizontally on one tier.

The section *layer combinations* contains the settings for elements that belong to multiple layers. Via the checkboxes under *attributes* you determine, to which layers an

¹Depending on the browser you use, the color fields are displayed as color picker or as simple text field. In the latter case, you have to provide the color as hexadecimal RGB value: A hash (#) followed by three two-digit hexadecimal numbers for red, green and blue, respectively. #000000, e.g., stands for black, #ffffff for white, #ff0000 for light red etc.

element has to belong in order to be subject to the definition of the combination in question. The other settings work like those described for the *layers*, and for those elements that belong to the layer combination, they override the values given for the single layers.

Under *search makros* you can enter predefined graph-specific search makros as described in Section ??. These makros are then available for search queries in the graph. In the search makros text area, the definitions (def ...) have to be enterd each on its own line.

1.2.2 Permitted annotations / tagset

In the window open by the command tagset, you can specify which annotation are permitted for the nodes and edges of the graph, i.e., you can define the graph-specific tagset. Only the keys and values defined here are permitted for the annotation of elements, illicit input is answered with an error message. Already existing annotations are not affected by changes of the permitted keys and values.

If you do not specify any keys and values, all annotations are permitted. If you specify keys, only these keys are allowed for annotations. If you leave the value field for a key empty, then all values are permitted for this key. If you want to restrict the possible values for a key, you may enter these into the value field of the key in question, separated by spaces.

The notation of the values is subject to the same rules as in the query language (cf. Section ??, p. ??): Simple values are entered without further markup; if a value contains special characters (siehe p. ??) it has to be enclosed in double quotes ("..."). Additionally – just like in the query language (p. ??) – you may use regular expressions, which are enclosed in slashes (/.../). In this case the regular expressions are anchored, i.e. an annotation value has to match the whole regular expression in order to be permitted.

1.2.3 Metadata

Additionally to the configuration of layers, visualization, search makros and permitted annotations, you may save metadata for a graph as key-value pairs. The command metadata opens the window, where you can enter an arbitrary number of keys with a text as corresponding value.

2 Keyboard shortcuts

In GraphAnno most of the functions related to navigation and display are controlled by keybord shortcuts. The following is a table of the available shortcuts:

Shortcut	Function
Navigation	
Alt + ←/→	previous/next sentence
Alt + Home/End	first/last sentence
Graph	
Ctrl + Shift + -/+	scale down/up graph
Ctrl + Shift + 0	zoom to fit (wrt. height)
Ctrl + Shift + arrows	move graph
Ctrl + Shift + Home/End	go to left/right edge of graph
Ctrl + Shift + Page up/Page down	go to upper/lower edge of graph
F4	toggle element references
Window	
F1	show/hide help window
F2	show/hide text and sentence annotations
F6	show/hide filter window
F7	show/hide search window

3 Command line commands

3.1 Data and navigation

3.1.1 Load file: load

With the command load, followed by the file name, you load a graph file into the work space. Provide the file name without the extension .json; it has to enclosed in double quotes ("...") if it contains spaces. Files are loaded from the data directory located in the GraphAnno main directory. Before loading, the work space is cleared from all data. So, changes that were not saved explicitly (using the command save) are lost. The name of the loaded file is shown next to the input line on the bottom of the user interface.

3.1.2 Load file: add

Just like the command load, the command add loads a file into the work space. The difference is, that the work space is not cleared – the new file is added and the new sentences are appended to the existing ones. After adding the file no file is shown as loaded next to the input line and the files cannot be saved separately anymore.

3.1.3 Save file: save

The file name has to be entered in the same way as with the command load. If there is a filename indicated next to the input line, the work space can be saved to this file without specifying the file name. Attention: no warning is issued when an existing file will be overwritten.

3.1.4 Clear work space: clear

The command clear clears the work space from all data. Changes that were not saved are lost. Next to the input line no file will be indicated anymore.

3.1.5 Create new sentence: ns

With the command ns – followed by one or more sentence names separated by spaces – you can create a new sentence. The command creates sentence nodes with the corresponding name attribute; afterwards you are directed to the first newly created sentence.

3.1.6 Delete sentence: del

The command del deletes the current sentence including the sentence node.

3.1.7 Set sentence: s

In order to navigate from sentence to sentence, you may use (alternatively to keyboard shortcuts or the dropdown field) the command s, followed by the name of the sentence you want to change to. You are then directed to the first sentence with the given name.

3.1.8 Export graphics: image

With the command image, you can export the graphic that GraphAnno is displaying for the current sentence. The first argument is the desired format. All formats supported by Graphviz are available, e.g. dot, eps, pdf, png or svg. See under http://www.graphviz.org/content/output-formats for the complete list. The second argument is the name of the new image file (without extension; put the name in double quotes if it contains spaces). The image will be saved in GraphAnno's images directory.

3.1.9 Export corpus: export

Using the command export, you can export the contents of the work space as a corpus in another format. The first argument is the format (at present the only fully functional format is sql for the import in GraphInspect; the format paula is theoretically available, but it is heavily restricted with respect to layers), the second argument is the

name of the corpus to be saved. For paula you can optionally specify the name of the corpus document to be created as third argument. The exported corpus will be saved in GraphAnno's exports directory.

3.1.10 Import text: import text

You can import texts with the command import text. After issuing the command, a window opens where you can enter the text and set the preferences for its processing. The work space will be cleared before the text is imported (but not as soon as the window opens). Changes that hadn't been saved are lost and no file is indicated next to the input line anymore.

In the import window you can choose between two methods of entering your text: You can upload a text file or you can paste it in the text area. For the processing of the text there are two methods available, too. For unedited text you may use the method "punkt segmenter". This method uses an automatic segmenter to split the text into sentences and tokens. In order to process abbreviation etc. correctly, you need to specify the language of the text.

The second processing method is "regular expressions"; this method is made for preformatted texts. First, you have to enter a string that will be used for segmenting the sentences. The preset is \n 2 for a file in which every sentence starts on a new line. The second string you have to enter a a regular expression that matches the tokens. The preset here is (\S+). This stands for a sequence of non-spaces, so all words that are separated by spaces are matched as tokens. The purpose of the parentheses is to save the matched string in the variable \$1, so it can be used in the next field. The next field is for an annotation command (see 3.2.3) for the tokens, that uses the string matched by the regular expression in the preceding field. The preset here is token: \$1. That means that the string matched as token is used for the annotation of the token text. Another example would be a text tagged for parts of speech, where the part of speech is appended to every word with an underscore. In this case you would enter the regular expression (\S+)_(\S+) and the annotation command token:\$1 pos:\$2. The regular expression in this case finds two strings of non-spaces that are joined by an underscore; the strings are saved to the variables \$1 (the word) and \$2 (the POS tag). In the annotation command these variables are used to annotate the token text and the pos attribute.

3.1.11 Import Toolbox data: import toolbox

Toolbox files can be imported using the command import toolbox. This command opens a window in which you can choose the file to be imported and enter the format description. The format description has to be in JSON format and consists of a list of a list of markers. The lists are sorted according to their levels – the highest (*record* level)

²\n stands for a line break, \t for a tab.

first – and contain the markers that belong to the respective level (markers are entered without backslash). The first marker of the first level (ref in the example below) will be used as record ID. The marker whose line is to be used as token text is preceded by an asterisk. Elements that lie below the token level will be joined and integrated into their respective tokens.

A format description for a toolbox file with three levels (record, word, morpheme) could look like this, e.g.:

```
[["ref", "eng"], ["*gw"], ["mph", "ge", "ps"]]
```

Like with the command load, the work space will be cleared when importing. Changes that haven't been saved will be lost; next to the input line no file will be indicated anymore.

3.1.12 Export and import configuration: export and import

The command export serves also for exporting graph configurations, that can be imported in other graphs using the command import. As first argument you enter the type of the configuration to be exported or imported: config for layers and visualization configuration (see 1.2.1), tagset for permitted annotations / tagset (see 1.2.2). The second argument is the filename for the configuration file to be saved or to be imported (without file extension). The exported file will be saved in a subdirectory – named like the configuration type – of the exports directory. Attention: when importing, the existing configuration will be replaced completely.

3.1.13 Edit configurations: config, tagset and metadata

These commands open the windows for the settings that were described in Section 1.2. config for layers and visualization, tagset for permitted annotations / tagset and metadata for metadata.

3.2 Annotation commands

GraphAnno's annotation commands are designed to be entered quickly, so their syntax is rather compact: they consist of a short command (often one letter only) and are followed by parameters separated by spaces.

The commands require to be in a sentence, i.e., you have to create a sentence first (using the command ns, see 3.1.5) if the work space is empty.

3.2.1 New node: n

The command for creating a new node is n, followed by the attributes the new node is to bear as key-value pairs in the format key: value. Key and value can be given either as

simple string (if it doesn't contain any of the special characters used in the annotation languages: "")") or as string in double quotes ("..."), that may contain any character (double quotes themselves have to be excaped with a backslash: \"). Additionally, you can specify the layer to which the new node is to belong (if you don't, it will belong to the layer set in the layer dropdown field). For this purpose you use the shortcuts defined in the layer configuration (cf. Section 1.2.1). The use of these shortcuts also has the effect of a switch, insofar as it sets the layer for the following operations (like the command \lambda, see ??).

Command n in modified BNF:

3.2.2 New edge: e

The command for creating a new edge is e, followed by start and end node of the edge to be created and the attributes, it is to bear. Like with n, the specification of a layer is possible.

Command e in modified BNF:

3.2.3 Annotieren: a

The command for annotation elements is a, followed by the elements to be annotated and the attributes with which they are to be annotated (all given elements will be

³The symbol _u stands for the space.

annotated with all given attributes; also layer shortcuts can be used). The order of the elements and attributes is free. You can also annotate sequences of elements of the same type (i.e., n, e or t) by entering the first and the last element joined by two dots. E.g., when you enter t3..t7, all tokens from t3 to t7 will be annotated (you may enter the sequence also in the inverse order, i.e., t7..t3).

At the same time you can use the command a for deleting attributes. In order to do so, enter the key to be deleted with colon, but without value.

Command a in modified BNF:

```
"a " a_parameters
command_a
                        a_parameters " " a_parameters
a parameters
                        element_reference
                        attributes
                        key
element_reference
                        node_reference
                        edge_reference
                        token_reference
                        meta node reference
                        element_sequence
edge_reference
                        "e" number
                        "m"
meta_node_reference =
element_sequence
                        node_sequence
                        token_sequence
                        edge sequence
                        node_reference ".." node_reference
node_sequence
token_sequence
                        token_reference ".." token_reference
edge_sequence
                        edge_reference ".." edge_reference
```

tbc.