Abstract

The use of parameters in the description of natural language syntax has to balance between the need to discriminate among (sometimes subtly different) languages, which can be seen as a cross-linguistic version of Chomsky’s (1964) descriptive adequacy, and the complexity of the acquisition task that a large number of parameters would imply, which is a problem for explanatory adequacy. Here we present a novel approach in which a machine learning algorithm is used to find dependencies in a table of parameters. The result is a dependency graph in which some of the parameters can be fully predicted from others. These empirical findings can be then subjected to linguistic analysis, which may either refute them by providing typological counter-examples of languages not included in the original dataset, dismiss them on theoretical grounds, or uphold them as tentative empirical laws worth of further study.

1 Introduction

Parametric theories of generative grammar focus on the problem of a formal and principled theory of grammatical diversity (Chomsky, 1981; Baker, 2001; Roberts, 2012). The basic intuition of parametric approaches is that the majority of observable syntactic differences among languages are derived, usually through complex deductive chains, from a smaller number of more abstract contrasts, drawn from a universal list of discrete, and normally binary, options, called parameters. The relation between observable patterns and the actual syntactic parameters which vary across languages is quite indirect: syntactic parameters are regarded as abstract differences often responsible for wider typological clusters of surface co-variation, often through an intricate deductive structure. In this sense, the concept of parametric data is not to be simplistically identified with that of syntactic pattern: co-varying syntactic properties/patterns are in fact the empirical manifestations of much more abstract cognitive structures.

Syntactic parameters are conceived as definable by UG (i.e. universally comparable) and set by each learner on the basis of her/his linguistic environment. Open parameters, or any set of more primitive concepts they can derive from (Longobardi, 2005; Lightfoot, 2017), define a variation space for biologically acquirable grammars, set (a.k.a. closed) parameters specify each of these grammars. Thus, the core grammar of every natural language can in principle be represented by a string of binary symbols (Clark and Roberts, 1993), each coding the value of a parameter of UG.

The Parametric Comparison Method (PCM, (Longobardi and Guardiano, 2009)) uses syntactic parameters to study historical
relationships among languages. An important aspect of parametric systems that is particularly relevant to the present research is that parameters form a pervasive network of partial implications (Guardiano and Longobardi, 2005; Longobardi and Guardiano, 2009; Longobardi et al., 2013): one particular value of some parameter A, but not the other, often entails the irrelevance of parameter B, whose consequences, i.e. the corresponding surface patterns, become predictable. Under such conditions, B becomes redundant and will not be set at all by the learner. The PCM system makes such interdependencies explicit: in our notation, he symbols + and − are used to represent the binary value of each parameter; the symbol ‘0’, instead, encodes the neutralising effect of implicational cross-parametric dependencies, i.e. cases in which the content of a parameter is either entirely predictable, or irrelevant altogether. The conditions which must hold for each parameter not to be neutralised are expressed in a Boolean form, i.e., either as simple states of another parameter (or negation thereof), or as conjunctions or disjunctions of values of other parameters.

The PCM has shown that an important effect of the pervasiveness of parameter interdependencies is a noticeable downsizing of the space of grammatical variation: according to some preliminary experiments (Bortolussi et al., 2011), the number of possible languages generated from a given set of independent binary parameters is reduced from $10^{18}$ to $10^{11}$ when their interdependencies are taken into account. This also crucially implies a noticeable reduction of the space of possible languages that a learner has to navigate when acquiring a language.

Here we adopt an empirical, data-driven approach to the task of identifying parameter dependencies, which has been implemented on a database of 71 languages described through the values of 91 syntactic parameters (see Appendix A) expressing the internal syntax of nominal structures. Our results show that applying machine learning techniques to the data reveals previously unknown dependencies between parameters, which could potentially lead to a further significant reduction of the search space of possible languages.

This paper sets out to identify parameters whose entire range of values can be fully predicted from the values of other parameters. There is an important difference between previously published work on parameter dependencies and this paper’s contribution, which needs to be emphasised: rather than state that, for example, any language in which $P_1 = +$ will have a fully predictable value of $P_2$ (a fact which we encode as $P_2 = 0$), we seek parameters whose value can be deduced in all cases from the values of certain other parameters, e.g. as shown in the hypothetical example in Figure 1. Should such a rule prove to have universal validity, then parameter $P_3$ would never offer any advantage in separating any two languages, yet it could clearly still play a useful role in describing them.

2 Learning Dependencies

We process our table of dimensions ($\#\text{lang} \times \#\text{param}$) with the data mining package WEKA (v.3.6.13) (Hall et al., 2009). More specifically, we take the values of all parameters but one for all languages (i.e. a dataset of size ($\#\text{lang} \times \#\text{param} - 1$), and learn a decision tree that predicts the value of the remaining parameter from the values of the other parameters. (Typically, only a few are necessary in each case.) This is repeated to produce a decision tree for each of the parameters. The machine learning algorithm used was ID3 (Quinlan, 1986). The algorithm produces a decision tree, in which each leaf corresponds to the value of the modelled parameter for the combination of parameter values listed on the way from the root to that leaf, e.g.: if $FGN = -$ and $FGP = +$ then $GCO = +$ (see Table 1). Unlike some of the more sophisticated decision tree learning algorithms, such as C4.5 (Quinlan, 1993), no postprocessing of the tree learnt

Figure 1: Parameter dependency model example

\[
\text{if } P_1 = + \text{ and } P_2 = - \text{ then } P_3 = + \\
\text{else } P_3 = -
\]
(such as pruning (Mitchell, 1997)) takes place, and the tree remains an accurate, exact reflection of the training data. If the combination of parameter values corresponding to one of the leaves of the tree is not observed in the data, the leaf contains the special label ‘null’ (see the tree predicting GCO in Table 1). In all other cases, that is, whenever the leaf label is ‘+’, ‘-’ or ‘0’, this is supported by one or more examples (languages) in the data.

Table 1: Examples of decision trees for parameters FGN and GCO

| FGN | GCO |
|-----|-----|
| if GCO = 0 then FGN = + | if GCO = 0 then GCO = null (never occurs) |
| if GCO = + then FGN = - | if GCO = + then GCO = 0 |
| if GCO = - then FGN = - | if GCO = - then GCO = + |

3 Results

The decision trees for all parameters were used to produce a dependency graph in which each vertex represents a parameter, and directed edges link the parameters, whose values are needed to predict a given parameter, with the node representing that parameter. For instance, there are edges from both FGN and FGP to GCO, as the decision tree for GCO refers to the values of FGN and FGP. Some of the decision trees are more complex, making use of up to nine separate parameters. The resulting graph is very complex (see Fig. 2). Therefore, we also present a subset of the graph (see Fig. 3), which only visualises those trees predicting one parameter from the value of one (as in the case of FGN) or two other parameters (e.g. GCO). The fact that some of the rules are missing from this graph is not an issue: for each listed node, all of the incoming edges are present, so that if we know those parameters, we are guaranteed to know the parameter they point to.

The interpretation of the graph is straightforward. For instance, looking at its top right corner, one can deduce that for any language in the dataset, it is enough to know the values of parameters EZ3 and PLS in order to know the value of EZ2, and therefore, of EZ1, too. Knowing (the value of) FVP means one also knows DMG and NSD; if one knows both FVP and DNN, the values of DNG, NSD, DSN, DMP and DMG are fully predictable for the given data set. In other words, 7 parameters (FVP, DNN, DNG, NSD, DSN, DMP and DMG) can be reduced to just 2 without any loss of information.

Some of the rules identified by the algorithm are not new, and are already contained in the dataset, as encoded by the implicational system described in Section 1. For instance, the parameter RHM is encoded as 0 when FGP = −, as the value of RHM is fully predictable in those cases. When a decision tree predicting FGP is learned, the result is as follows: if RHM = 0 then FGP = − else FGP = +.

Even the rest of the rules learned are still just empirical findings that may change with the addition of other examples of languages or their validity may be questioned by linguists on theoretical grounds.

Linguistic analysis of the results is ongoing, and while no part of the results has been accepted as sufficient evidence to dispose of a parameter, implication rules may be revised on the basis of the decision trees learned, as in the case of the parameter PLS. According to its definition, the parameter “asks if in a language without grammaticalized Number, a plural marker can also appear outside a nominal phrase, marking a distributive relation between the plural subject and the constituent bearing it.” (E.g. PLS = + for Korean, but PLS = − for Japanese.)

Prior to this research, there was an implication rule stating that PLS is neutralised (that is, its value is predictable) for all combinations of CGO and FGN values other than CGO = − and FGN = −. This rule has now been replaced with a new rule stating that PLS is neutralised for all combinations of values of FGM and FGN, except when FGM = + and FGN = −, and the evidence showing that the new rule is consistent with the data came from the tree learned for PLS.
Figure 2: Full dependency graph
Figure 3: Partial dependency graph constructed from implications with up to two antecedents
4 Discussion

The results reported here show that applying machine learning techniques to the data can reveal previously unknown dependencies between parameters, leading to a potentially significant reduction in the search space of possible languages. The data contains more features than data points, which can make for the generation of spurious rules. The most obvious way to counteract this, adding more languages, comes at a very high cost, as it requires well-trained linguists. One can also use Occam’s Razor and limit the search space of possible rules by limiting the number of antecedents in the rule, e.g. to two as we did here. Yet another approach is to collect data selectively for rules of interest, as only a small number of parameters, e.g. 2–3 per language, will be needed to test each rule.

This research could have important implications for the understanding of processes underlyng the faculty of language (potentially strengthening the case for UG), with implications ranging from models of language acquisition to historical linguistics, where the syntactic relatedness between two languages may be more adequately measured. However, the approach requires a close collaboration between a machine learning expert, discovering empirical laws in the data, and a linguist who can test their plausibility and theoretical consequences. There is also an open theoretical computational learning challenge here presented by the need to estimate the significance of empirical findings from a given number of examples (languages) with respect to the available range of discriminative features in the dataset.

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## Appendix A: List of Parameters

| Abbreviation | Description |
|--------------|-------------|
| FGP          | gramm. person |
| FGM          | gramm. Case |
| FPC          | gramm. perception |
| FGT          | gramm. temporality |
| FGN          | gramm. number |
| GCO          | gramm. collective number |
| PLS          | plurality spreading |
| FND          | number in D |
| NOD          | NP over D |
| FSN          | feature spread to N |
| FNN          | number on N |
| SGE          | semantic gender |
| FGG          | gramm. gender |
| CGB          | unbounded sg N |
| DGR          | gramm. amount |
| DGP          | gramm. text anaphora |
| CGR          | strong amount |
| NSD          | strong person |
| FVP          | variable person |
| DGD          | gramm. distality |
| DPQ          | free null partitive Q |
| DCN          | article-checking N |
| DNN          | null-N-licensing art |
| DIN          | D-controlled infl. on N |
| FGC          | gramm. classifier |
| DBC          | strong classifier |
| GSC          | c-selection |
| NOE          | N over ext. arg. |
| DMP          | def matching pronominal possessives |
| DMG          | def matching genitives |
| GCN          | Poss*-checking N |
| GFN          | Gen-feature spread to Poss* |
| GAL          | Dependent Case in NP |
| GUN          | uniform Gen |
| EZ1          | generalized linker |
| EZ2          | non-clausal linker |
| EZ3          | non-genitive linker |
| GAD          | adpositional Gen |
| GFO          | GenO |
| PGO          | partial GenO |
| GFS          | GenS |
| GIT          | Genitive-licensing iterator |
| GSI          | grammaticalised inalienability |
| ALP          | alienable possession |
| GST          | grammaticalised Genitive |
| GEI          | Genitive inversion |
| GNR          | non-referential head marking |
| HMP          | NP-heading modifier |
| AST          | structured APs |
| STC          | structured cardinals |
| GPC          | gender polarity cardinals |
| PMN          | personal marking on numerals |
| CQU          | cardinal quantifiers |
| PCA          | number spread through cardinal adjectives |
| FFS          | feature spread to structured APs |
| ADI          | D-controlled infl. on A |
| PSC          | number spread from cardinal quantifiers |
| RHM          | Head-marking on Rel |
| FRC          | verbal relative clauses |
| NRC          | nominalised relative clause |
| NOR          | NP over verbal relative clause/ adpositional genitives |
| AER          | relative extrap. |
| ARR          | free reduced rel |
| DOR          | def on relatives |
| NOP          | NP over non-genitive arguments |
| PNP          | P over complement |
| NPP          | N-raising with obl. pied-piping |
| NGO          | N over GenO |
| NOA          | N over As |
| NM2          | N over M2 As |
| NM1          | N over M1 As |
| EAF          | fronted high As |
| NON          | N over numerals |
| FPO          | feature spread to genitive postpositions |
| ACM          | class MOD |
| DOA          | def on all +N |
| NEX          | gramm. expletive article |
| NCL          | clitic poss. |
| PDC          | article-checking poss. |
| ACL          | enclitic poss. on As |
| APO          | adjectival poss. |
| WAP          | wackernagel adjectival poss. |
| AGE          | adjectival Gen |
| OPK          | obligatory possessive with kinship nouns |
| TSP          | split deictic demonstratives |
| TSD          | split demonstratives |
| TAD          | adjectival demonstratives |
| TDC          | article-checking demonstratives |
| TLC          | Loc-checking demonstratives |
| TNL          | NP over Loc |
| XCN          | conjugated nouns |