

## Transformational Priors Over Grammars

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### Abstract

This paper proposes a novel class of PCFG parameterizations that support linguistically reasonable priors over PCFGs. To estimate the parameters is to discover a notion of relatedness among context-free rules such that related rules tend to have related probabilities. The prior favors grammars in which the relationships are simple to describe and have few major exceptions. A basic version that bases relatedness on weighted edit distance yields superior smoothing of grammars learned from the Penn Treebank (20% reduction of rule perplexity over the best previous method).

### 1 A Sketch of the Concrete Problem

This paper uses a new kind of statistical model to smooth the probabilities of PCFG rules. It focuses on “flat” or “dependency-style” rules. These resemble subcategorization frames, but include adjuncts as well as arguments.

The verb `put` typically generates 3 dependents—a subject NP at left, and an object NP and goal PP at right:

- $S \rightarrow NP \text{ put } NP \text{ PP}$ : Jim put [the pizza] [in the oven]

But `put` may also take other dependents, in other rules:

- $S \rightarrow NP \text{ Adv } \text{put } NP \text{ PP}$ : Jim often put [a pizza] [in the oven]
- $S \rightarrow NP \text{ put } NP \text{ PP } PP$ : Jim put soup [in an oven] [at home]
- $S \rightarrow NP \text{ put } NP$ : Jim put [some shares of IBM stock]
- $S \rightarrow NP \text{ put } \text{Prt } NP$ : Jim put away [the sauce]
- $S \rightarrow \text{TO } \text{put } NP \text{ PP}$ : to put [the pizza] [in the oven]
- $S \rightarrow NP \text{ put } NP \text{ PP } \text{SBAR}$ : Jim put it [to me] [that . . . ]

These other rules arise if `put` can add, drop, reorder, or retype its dependents. These **edit operations** on rules are semantically motivated and quite common (Table 1).

We wish to learn contextual probabilities for the edit operations, based on an observed sample of flat rules. In English we should discover, for example, that it is quite common to add or delete PP at the right edge of a rule. These contextual edit probabilities will help us guess the true probabilities of novel or little-observed rules.

However, rules are often idiosyncratic. Our smoothing method should not keep us from noticing (given enough evidence) that `put` takes a PP more often than most verbs. Hence this paper’s proposal is a Bayesian smoothing method that allows idiosyncrasy in the grammar while presuming regularity to be more likely *a priori*.

The model will assign a positive probability to each of the infinitely many formally possible rules. The following bizarre rule is not observed in training, and seems very unlikely. But there is no formal reason to rule it out, and it *might* help us parse an unlikely test sentence. So the model will allow it some tiny probability:

- $S \rightarrow NP \text{ Adv } PP \text{ put } PP \text{ PP } NP \text{ AdjP } S$

### 2 Background and Other Approaches

A PCFG is a conditional probability function  $p(\text{RHS} \mid \text{LHS})$ .<sup>1</sup> For example,  $p(V \text{ NP } PP \mid VP)$  gives the probability of the rule  $VP \rightarrow V \text{ NP } PP$ . With lexicalized nonterminals, it has form  $p(V_{\text{put}} \text{ NP}_{\text{pizza}} \text{ PP}_{\text{in}} \mid VP_{\text{put}})$ .

Usually one makes an independence assumption and defines this as  $p(V_{\text{put}} \text{ NP } PP \mid VP_{\text{put}})$  times factors that choose dependent headwords `pizza` and `in` according to the selectional preferences of `put`. This paper is about estimating the first factor,  $p(V_{\text{put}} \text{ NP } PP \mid VP_{\text{put}})$ .

In supervised learning, it is simplest to use a maximum likelihood estimate (perhaps with backoff from `put`). Charniak (1997) calls this a “Treebank grammar” and gambles that assigning 0 probability to rules unseen in training data will not hurt parsing accuracy too much.

However, there are four reasons not to use a Treebank grammar. First, ignoring unseen rules necessarily sacrifices some accuracy. Second, we will show that it improves accuracy to flatten the parse trees and use flat, dependency-style rules like  $p(NP \text{ put } NP \text{ PP} \mid S_{\text{put}})$ ; this avoids overly strong independence assumptions, but it increases the number of unseen rules and so makes Treebank grammars less tenable. Third, backing off from the word is a crude technique that does not distinguish among words.<sup>2</sup> Fourth, one would eventually like to reduce or eliminate supervision, and then generalization is important to constrain the search to reasonable grammars.

To smooth the distribution  $p(\text{RHS} \mid \text{LHS})$ , one can define it in terms of a set of parameters and then estimate those parameters. Most researchers have used an  $n$ -gram model (Eisner, 1996; Charniak, 2000) or more general Markov model (Alshawi, 1996) to model the sequence of nonterminals in the RHS. The sequence  $V_{\text{put}} \text{ NP } PP$  in our example is then assumed to be emitted by some Markov model of  $VP_{\text{put}}$  rules (again with backoff from `put`). Collins (1997, model 2) uses a more sophisticated model in which all *arguments* in this sequence are generated jointly, as in a Treebank grammar, and then a Markov process is used to insert adjuncts among the arguments.

While Treebank models overfit the training data, Markov models underfit. A simple compromise (novel to this paper) is a hybrid **Treebank/Markov** model, which backs off from a Treebank model to a Markov. Like this paper’s main proposal, it can learn well-observed idiosyncratic rules but generalizes when data are sparse.<sup>3</sup>

<sup>1</sup>Nonstandardly, this allows infinitely many rules with  $p > 0$ .

<sup>2</sup>One might do better by backing off to word clusters, which Charniak (1997) did find provided a small benefit.

<sup>3</sup>Carroll and Rooth (1998) used a similar hybrid technique

These models are beaten by our rather different model, **transformational smoothing**, which learns common rules and common edits to them. The comparison is a direct one, based on the perplexity or cross-entropy of the trained models on a test set of  $S \rightarrow \dots$  rules.<sup>4</sup>

A subtlety is that two annotation styles are possible. In the Penn Treebank, `put` is the head of three constituents (V, VP, and S, where underlining denotes a head child) and joins with different dependents at different levels:

- [<sub>S</sub> [<sub>NP</sub> Jim] [<sub>VP</sub> [V put] [<sub>NP</sub> pizza] [<sub>PP</sub> in the oven]]]

In the **flattened** or dependency version that we prefer, each word joins with all of its dependents at once:

- [<sub>S</sub> [<sub>NP</sub> Jim] put [<sub>NP</sub> pizza] [<sub>PP</sub> in the oven]]

A PCFG generating the flat structure must estimate  $p(\text{NP } \text{put } \text{NP } \text{PP} \mid S_{\text{put}})$ . A non-flat PCFG adds the dependents of `put` in 3 independent steps, so in effect it factors the flat rule’s probability into 3 supposedly independent “subrule probabilities,”  $p(\text{NP } \text{VP}_{\text{put}} \mid S_{\text{put}}) \cdot p(\text{VP}_{\text{put}} \text{ NP } \text{PP} \mid \text{VP}_{\text{put}}) \cdot p(\text{put} \mid \text{V}_{\text{put}})$ .

Our evaluation judges the estimates of flat-rule probabilities. Is it better to estimate these directly, or as a product of estimated subrule probabilities?<sup>5</sup> Transformational smoothing is best applied to the former, so that the edit operations can freely rearrange all of a word’s dependents. We will see that the Markov and Treebank/Markov models also work much better this way—a useful finding.

### 3 The Abstract Problem: Designing Priors

This section outlines the Bayesian approach to learning probabilistic grammars (for us, estimating a distribution over flat CFG rules). By choosing among the many grammars that *could* have generated the training data, the learner is choosing how to generalize to novel sentences.

To guide the learner’s choice, one can explicitly specify a prior probability distribution  $p(\theta)$  over possible grammars  $\theta$ , which themselves specify probability distributions over strings, rules, or trees. A learner should seek  $\theta$  that maximizes  $p(\theta) \cdot p(D \mid \theta)$ , where  $D$  is the set of strings, rules, or trees observed by the learner. The first factor favors regularity (“pick an *a priori* plausible grammar”), while the second favors fitting the idiosyncrasies of the data, especially the commonest data.<sup>6</sup>

to *evaluate* rule distributions that they acquired from an automatically-parsed treebank.

<sup>4</sup>All the methods evaluated here apply also to full PCFGs, but verb-headed rules  $S \rightarrow \dots$  present the most varied, interesting cases. Many researchers have tried to learn verb subcategorization, though usually not probabilistic subcategorization.

<sup>5</sup>In testing the latter case, we sum over *all* possible internal bracketings of the rule. We do train this case on the true internal bracketing, but it loses even with this unfair advantage.

<sup>6</sup>This approach is called semi-Bayesian or Maximum A Pos-

Priors can help both unsupervised and supervised learning. (In the semi-supervised experiments here, training data is not raw text but a sparse sample of flat rules.)

Indeed a good deal of syntax induction work has been carried out in just this framework (Stolcke and Omohundro, 1994; Chen, 1996; De Marcken, 1996; Grünwald, 1996; Osborne and Briscoe, 1997). However, all such work to date has adopted rather simple prior distributions. Typically, it has defined  $p(\theta)$  to favor PCFGs whose rules are few, short, nearly equiprobable, and defined over a small set of nonterminals. Such definitions are convenient, especially when specifying an encoding for MDL, but since they treat all rules alike, they may not be good descriptions of linguistic plausibility. For example, they will never penalize the *absence* of a predictable rule.

A prior distribution can, however, be used to encode various kinds of *linguistic* notions. After all, a prior is really a soft form of Universal Grammar: it gives the learner enough prior knowledge of grammar to overcome Chomsky’s “poverty of the stimulus” (i.e., sparse data).

- A preference for small or simple grammars, as above.
- Substantive preferences, such as a preference for verbs to take 2 nominal arguments, or to allow `PP` adjuncts.
- Preferences for systematicity, such as a preference for the rules to be consistently head-initial or head-final.

This paper shows how to design a prior that favors a certain kind of systematicity. Lexicalized grammars for natural languages are very large—each word specifies a distribution over all possible dependency rules it could head—but they tend to have internal structure. The new prior prefers *grammars in which a rule’s probability can be well-predicted from the probabilities of other rules*, using linguistic transformations such as edit operations.

For example,  $p(\text{NP } \text{Adv } w \text{ put } \text{NP } \text{PP} \mid S_w)$  correlates with  $p(\text{NP } w \text{ NP } \text{PP} \mid S_w)$ . Both numbers are high for  $w = \text{put}$ , medium for  $w = \text{fund}$ , and low for  $w = \text{sleep}$ . The slope of the regression line has to do with the rate of preverbal `Adv`-insertion in English.

The correlation is not perfect (some verbs are especially prone to adverbial modification), which is why we will only model it with a prior. To just the extent that evidence about  $w$  is sparse, the prior will cause the learner to smooth the two probabilities toward the regression line.

### 4 Patterns Worth Modeling

Before spelling out our approach, let us do a sanity check.

**A frame** is a flat rule whose headword is replaced with *teriori* learning, since it is equivalent to maximizing  $p(\theta \mid D)$ . It is also equivalent to Minimum Description Length (MDL) learning, which minimizes the total number of bits  $\ell(\theta) + \ell(D \mid \theta)$  needed to encode grammar and data, because one can choose an encoding scheme where  $\ell(x) = -\log_2 p(x)$ , or conversely, define probability distributions by  $p(x) = 2^{-\ell(x)}$ .

MI	$\alpha$	$\beta$	MI	$\alpha$	$\beta$	MI	$\alpha$	$\beta$
9.01	[NP — ADJP-PRD]	[NP — RB ADJP-PRD]	4.76	[TO — S]	[— S]	5.54	[TO — NP PP]	[NP TO — NP]
8.65	[NP — ADJP-PRD]	[NP — PP-LOC-PRD]	4.17	[TO — S]	[TO — NP PP]	5.25	[TO — NP PP]	[NP MD — NP.]
8.01	[NP — ADJP-PRD]	[NP — NP-PRD]	2.77	[TO — S]	[TO — NP]	4.67	[TO — NP PP]	[NP MD — NP]
7.69	[NP — ADJP-PRD]	[NP — ADJP-PRD.]	6.13	[TO — NP]	[TO — NP SBAR-TMP]	4.62	[TO — NP PP]	[TO — ]
8.49	[NP — NP-PRD]	[NP — NP-PRD.]	5.72	[TO — NP]	[TO — NP PP PP]	3.19	[TO — NP PP]	[TO — NP]
7.91	[NP — NP-PRD]	[NP — ADJP-PRD.]	5.36	[TO — NP]	[NP MD RB — NP]	2.05	[TO — NP PP]	[— NP]
7.01	[NP — NP-PRD]	[NP — ADJP-PRD]	5.16	[TO — NP]	[TO — NP PP PP-TMP]	5.08	[— NP]	[ADVP-TMP — NP]
8.45	[NP — ADJP-PRD.]	[NP — PP-LOC-PRD]	5.11	[TO — NP]	[TO — NP ADVP]	4.86	[— NP]	[ADVP — NP]
8.30	[NP — ADJP-PRD.]	[NP — NP-PRD.]	4.85	[TO — NP]	[TO — NP PP-LOC]	4.53	[— NP]	[— NP PP-LOC]
8.04	[NP — ADJP-PRD.]	[NP — NP-PRD]	4.84	[TO — NP]	[MD — NP]	3.50	[— NP]	[— NP PP]
7.01	[NP — ADJP-PRD.]	[NP — ADJP-PRD]	4.49	[TO — NP]	[NP TO — NP]	3.17	[— NP]	[— S]
7.01	[NP — SBAR]	[NP — SBAR.]	4.36	[TO — NP]	[NP MD — S]	2.28	[— NP]	[NP — NP]
4.75	[NP — SBAR]	[NP — SBAR.]	4.36	[TO — NP]	[NP TO — NP PP]	1.89	[— NP]	[NP — NP.]
6.94	[NP — SBAR.]	[“ NP — SBAR.]	4.26	[TO — NP]	[NP MD — NP PP]	2.56	[NP — NP]	[NP — NP.]
5.94	[NP — SBAR.]	[NP — SBAR.]	4.26	[TO — NP]	[TO — NP PP-TMP]	2.20	[NP — NP]	[— NP]
5.90	[NP — SBAR.]	[S, NP —.]	4.21	[TO — NP]	[TO — PRT NP]	4.89	[NP — NP.]	[NP ADVP-TMP — NP.]
5.82	[NP — SBAR.]	[NP ADVP — SBAR.]	4.20	[TO — NP]	[NP MD — NP]	4.57	[NP — NP.]	[NP ADVP — NP.]
4.68	[NP — SBAR.]	[— SBAR]	3.99	[TO — NP]	[TO — NP PP]	4.51	[NP — NP.]	[NP — NP PP-TMP]
4.50	[NP — SBAR.]	[NP — SBAR]	3.69	[TO — NP]	[NP MD — NP.]	3.35	[NP — NP.]	[NP — S.]
3.23	[NP — SBAR.]	[NP — S.]	3.60	[TO — NP]	[TO — ]	2.99	[NP — NP.]	[NP — NP]
2.07	[NP — SBAR.]	[NP — ]	3.56	[TO — NP]	[TO — PP]	2.96	[NP — NP.]	[NP — NP PP.]
1.91	[NP — SBAR.]	[NP — NP.]	2.56	[TO — NP]	[NP — NP PP]	2.25	[NP — NP.]	[— NP PP]
1.63	[NP — SBAR.]	[NP — NP]	2.04	[TO — NP]	[NP — S]	2.20	[NP — NP.]	[— NP]
4.52	[NP — S]	[NP — S.]	1.99	[TO — NP]	[NP — NP]	4.82	[NP — S.]	[— S]
4.27	[NP — S]	[— S]	1.69	[TO — NP]	[NP — NP.]	4.58	[NP — S.]	[NP — S]
3.36	[NP — S]	[NP — ]	1.68	[TO — NP]	[NP — NP PP.]	3.30	[NP — S.]	[NP — ]
2.66	[NP — S]	[NP — NP.]	1.03	[TO — NP]	[— NP]	2.93	[NP — S.]	[NP — NP.]
2.37	[NP — S]	[NP — NP]	4.75	[S, NP —.]	[NP — SBAR.]	2.28	[NP — S.]	[NP — NP]

Table 1: The most predictive pairs of sentential frames. If  $S \rightarrow \alpha$  occurs in training data at least 5 times with a given headword in the — position, then  $S \rightarrow \beta$  also tends to appear at least once with that headword. MI measures the mutual information of these two events, computed over all words. When MI is large, as here, the edit distance between  $\alpha$  and  $\beta$  tends to be strikingly small (1 or 2), and certain linguistically plausible edits are extremely common.

the variable “—” (corresponding to  $w$  above). Table 1 illustrates that in the Penn Treebank, if frequent rules with frame  $\alpha$  imply matching rules with frame  $\beta$ , there are usually edit operations (section 1) to easily turn  $\alpha$  into  $\beta$ .

How about rare rules, whose probabilities are most in need of smoothing? Are the same edit transformations that we can learn from frequent cases (Table 1) appropriate for predicting the rare cases? The very rarity of these rules makes it impossible to create a table like Table 1.

However, rare rules can be measured in the aggregate, and the result suggests that the same kinds of transformations are indeed useful—perhaps even more useful—in predicting them. Let us consider the set  $R$  of 2,809,545 possible flat rules that stand at edit distance 1 from the set of  $S \rightarrow \dots$  rules observed in our English training data. That is, a rule such as  $S_{\text{put}} \rightarrow \text{NP put NP}$  is in  $R$  if it did not appear in training data itself, but could be derived by a single edit from some rule that did appear.

A bigram Markov model (section 2) was used to identify 2,714,763 rare rules in  $R$ —those that were predicted to occur with probability  $< 0.0001$  given their headwords. 79 of these rare rules actually appeared in a development-data set of 1423 rules. The bigram model would have expected only 26.2 appearances, given the lexical headwords in the test data set. The difference is statistically significant ( $p < 0.001$ , bootstrap test).

In other words, the bigram model underpredicts the edit-distance “neighbors” of observed rules by a factor of 3.<sup>7</sup> One can therefore hope to use the edit transformations to improve on the bigram model. For example, the

<sup>7</sup>Similar results are obtained when we examine just one particular kind of edit operation, or rules of one particular length.

Delete Y transformation recognizes that if  $\dots X Y Z \dots$  has been observed, then  $\dots X Z \dots$  is plausible even if the bigram  $X Z$  has not previously been observed.

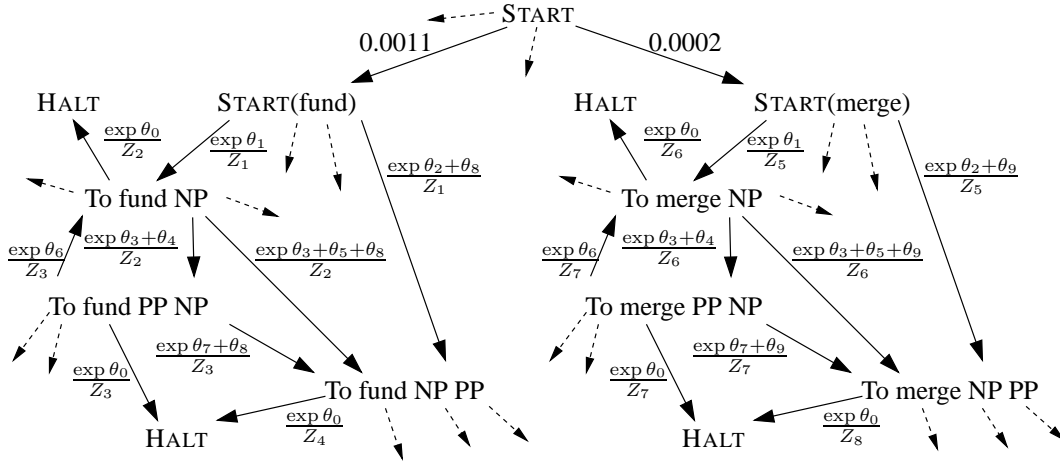
Presumably, edit operations are common because they modify a rule in semantically useful ways, allowing the filler of a semantic role to be expressed (Insert), suppressed (Delete), retyped (Substitute), or heavy-shifted (Swap). Such “valency-affecting operations” have repeatedly been invoked by linguists; they are not confined to English.<sup>8</sup> So a learner of an unknown language can reasonably expect *a priori* that flat rules related by edit operations may have related probabilities.

However, *which* edit operations varies by language. Each language defines its own weighted, contextual, asymmetric edit distance. So the learner will have to discover how likely *particular* edits are in particular contexts. For example, it must learn the rates of preverbal Adv-insertion and right-edge PP-insertion. Evidence about these rates comes mainly from the frequent rules.

## 5 A Transformation Model

The form of our new model is shown in Figure 1. The vertices are flat context-free rules, and the arcs between them represent edit transformations. The set of arcs leav-

<sup>8</sup>Carpenter (1991) writes that whenever linguists run into the problem of systematic redundancy in the syntactic lexicon, they design a scheme in which lexical entries can be derived from one another by just these operations. We are doing the same thing. The only twist that the lexical entries (in our case, flat PCFG rules) have probabilities that must also be derived, so we will assume that the speaker applies these operations (randomly from the hearer’s viewpoint) at various rates to be learned.



$\theta_0$  halts                                       $\theta_3$  inserts PP                                       $\theta_6$  deletes PP                                       $\theta_8$  yields To fund NP PP  
 $\theta_1$  chooses To — NP                                       $\theta_4$  inserts PP before NP                                       $\theta_7$  moves NP right past PP                                       $\theta_9$  yields To merge NP PP  
 $\theta_2$  chooses To — NP PP                                       $\theta_5$  inserts PP before right edge

Figure 1: A fragment of a transformation model. Vertices are possible context-free rules (their left-hand sides,  $S_{\text{fund}} \rightarrow$  and  $S_{\text{merge}} \rightarrow$ , are omitted to avoid visual clutter). Arc probabilities are determined log-linearly, as shown, from a real-valued vector  $\theta$  of feature weights. The  $Z$  values are chosen so that the arcs leaving each vertex have total probability 1. Dashed arrows represent arcs not shown here (there are hundreds from each vertex, mainly insertions). Also, not all features are shown (see Table 2).

ing any given vertex has total probability 1. The learner’s job is to discover the probabilities.

Fortunately, the learner does not have to learn a separate probability for each of the (infinitely) many arcs, since many of the arcs represent identical or similar edits. As shown in Figure 1, an arc’s probability is determined from meaningful features of the arc, using a conditional log-linear model of  $p(\text{arc} \mid \text{source vertex})$ . The learner only has to learn the finite vector  $\theta$  of feature weights. Arcs that represent similar transformations have similar features, so they tend to have similar probabilities.

This **transformation model** is really a PCFG with unusual parameterization. That is, for any value of  $\theta$ , it defines a language-specific probability distribution over all possible context-free rules (graph vertices). To sample from this distribution, take a random walk from the special vertex  $\text{START}$  to the special vertex  $\text{HALT}$ . The rule at the last vertex reached before  $\text{HALT}$  is the sample.

This sampling procedure models a process where the speaker chooses an initial rule and edits it repeatedly. The random walk might reach  $S_{\text{fund}} \rightarrow \text{To fund NP}$  in two steps and simply halt there. This happens with probability  $0.0011 \cdot \frac{\exp \theta_1}{Z_1} \cdot \frac{\exp \theta_0}{Z_2}$ . Or, having arrived at  $S_{\text{fund}} \rightarrow \text{To fund NP}$ , it might transform it into  $S_{\text{fund}} \rightarrow \text{To fund PP NP}$  and then further to  $S_{\text{fund}} \rightarrow \text{To fund NP PP}$  before halting.

Thus,  $p_\theta(S_{\text{fund}} \rightarrow \text{To fund NP PP})$  denotes the probability that the random walk somehow reaches  $S_{\text{fund}} \rightarrow \text{To fund NP PP}$  and halts there. Conditionalizing this probability gives  $p_\theta(\text{To — NP PP} \mid S_{\text{fund}})$ , as needed for the PCFG.<sup>9</sup>

<sup>9</sup>The experiments of this paper do not allow transformations

Given  $\theta$ , it is nontrivial to solve for the probability distribution over grammar rules  $e$ . Let  $I_\theta(e)$  denote the **flow** to vertex  $e$ . This is defined to be the total probability of all paths from  $\text{START}$  to  $e$ . Equivalently, it is the expected number of times  $e$  would be visited by a random walk from  $\text{START}$ . The following recurrence defines  $p_\theta(e)$ :<sup>10</sup>

$$I_\theta(e) = \delta_{e, \text{START}} + \sum_{e'} I_\theta(e') \cdot p(e' \rightarrow e) \quad (1)$$

$$p_\theta(e) = I_\theta(e) \cdot p(e \rightarrow \text{HALT}) \quad (2)$$

Since solving the large linear system (1) would be prohibitively expensive, in practice we use an approximate relaxation algorithm (Eisner, 2001) that propagates flow through the graph until near-convergence. In general this may underestimate the true probabilities somewhat.

Now consider how the parameter vector  $\theta$  affects the distribution over rules,  $p_\theta(e)$ , in Figure 1:

- By raising the **initial weight**  $\theta_1$ , one can increase the flow to  $S_{\text{fund}} \rightarrow \text{To fund NP}$ ,  $S_{\text{merge}} \rightarrow \text{To merge NP}$ , and the like. By equation (2), this also increases the probability of these rules. But the effect *also* feeds through the graph to increase the flow and probability at those rules’ *descendants* in the graph, such as  $S_{\text{merge}} \rightarrow \text{To merge NP PP}$ .

So a single parameter  $\theta_1$  controls a whole complex of rule probabilities (roughly speaking, the infinitival transitives). The model thereby captures the fact that, although

that change the LHS or headword of a rule, so it is trivial to find the divisor  $p_\theta(S_{\text{fund}})$ : in Figure 1 it is 0.0011. But in general, LHS-changing transformations can be useful (Eisner, 2001).

<sup>10</sup>Where  $\delta_{x,y} = 1$  if  $x = y$ , else  $\delta_{x,y} = 0$ .

rules are mutually exclusive events whose probabilities sum to 1, transformationally related rules have positively correlated probabilities that rise and fall together.

- The **exception weight**  $\theta_9$  appears on all and only the arcs to  $S_{\text{merge}} \rightarrow \text{To merge NP PP}$ . That rule has even higher probability than predicted by PP-insertion as above (since `merge`, unlike `fund`, actually tends to subcategorize for  $\text{PP}_{\text{with}}$ ). To model its idiosyncratic probability, one can raise  $\theta_9$ . This “lists” the rule specially in the grammar. Rules derived from it also increase in probability (e.g.,  $S_{\text{merge}} \rightarrow \text{To Adv merge NP PP}$ ), since again the effect feeds through the graph.
- The **generalization weight**  $\theta_3$  models the strength of the PP-insertion relationship. Equations (1) and (2) imply that  $p_{\theta}(S_{\text{fund}} \rightarrow \text{To fund NP PP})$  is modeled as a linear combination of the probabilities of that rule’s parents in the graph.  $\theta_3$  controls the coefficient of  $p_{\theta}(S_{\text{fund}} \rightarrow \text{To fund NP})$  in this linear combination, with the coefficient approaching zero as  $\theta_3 \rightarrow -\infty$ .
- Narrower generalization weights such as  $\theta_4$  and  $\theta_5$  control *where* PP is likely to be inserted. To learn the feature weights is to learn which features of a transformation make it probable or improbable in the language.

Note that the vertex labels, graph topology, and arc parameters are language independent. That is, Figure 1 is supposed to represent Universal Grammar: it tells a learner what kinds of generalizations to look for. The language-specific part is  $\theta$ , which specifies which generalizations and exceptions help to model the data.

## 6 The Prior

The model has more parameters than data. Why? Beyond the initial weights and generalization weights, in practice we allow one exception weight (e.g.,  $\theta_8, \theta_9$ ) for each rule that appeared in training data. (This makes it possible to learn arbitrary exceptions, as in a Treebank grammar.)

Parameter estimation is nonetheless possible, using a prior to help choose among the many values of  $\theta$  that do a reasonable job of explaining the training data. The prior constrains the degrees of freedom: while many parameters are available in principle, the prior will ensure that the data are described using as few of them as possible.

The point of reparameterizing a PCFG in terms of  $\theta$ , as in Figure 1, is precisely that only one parameter is needed per linguistically salient property of the PCFG. Making  $\theta_3 > 0$  creates a broadly targeted transformation. Making  $\theta_9 \neq 0$  or  $\theta_1 \neq 0$  lists an idiosyncratic rule, or class of rules, together with other rules derived from them. But it takes more parameters to encode less systematic properties, such as narrowly targeted edit transformations ( $\theta_4, \theta_5$ ) or families of unrelated exceptions.

A natural prior for the parameter vector  $\theta \in \mathbb{R}^k$  is therefore specified in terms of a variance  $\sigma^2$ . We simply

say that the weights  $\theta_1, \theta_2, \dots, \theta_k$  are independent samples from the normal distribution with mean 0 and variance  $\sigma^2 > 0$  (Chen and Rosenfeld, 1999):

$$\Theta \sim N(0, \sigma^2) \times N(0, \sigma^2) \times \dots \times N(0, \sigma^2) \quad (3)$$

or equivalently, that  $\theta$  is drawn from a multivariate Gaussian with mean  $\vec{0}$  and diagonal covariance matrix  $\sigma^2 I$ , i.e.,  $\Theta \sim N(\vec{0}, \sigma^2 I)$ .

This says that *a priori*, the learner expects most features in Figure 1 to have weights close to zero, i.e., to be irrelevant. Maximizing  $p(\theta) \cdot p(D | \theta)$  means finding a relatively small set of features that adequately describe the rules and exceptions of the grammar. Reducing the variance  $\sigma^2$  strengthens this bias toward simplicity.

For example, if  $S_{\text{fund}} \rightarrow \text{To fund NP PP}$  and  $S_{\text{merge}} \rightarrow \text{To fund NP PP}$  are both observed more often than the current  $p_{\theta}$  distribution predicts, then the learner can follow either (or both) of two strategies: raise  $\theta_8$  and  $\theta_9$ , or raise  $\theta_3$ . The former strategy fits the training data only; the latter affects many disparate arcs and leads to generalization. The latter strategy may harm  $p(D | \theta)$  but is preferred by the prior  $p(\theta)$  because it uses one parameter instead of two. If *more* than two words act like `merge` and `fund`, the pressure to generalize is stronger.

## 7 Perturbation Parameters

In experiments, we have found that a slight variation on this model gets slightly better results. Let  $\theta_e$  denote the exception weight (if any) that allows one to tune the probability of rule  $e$ . We eliminate  $\theta_e$  and introduce a different parameter  $\pi_e$ , called a **perturbation**, which is used in the following replacements for equations (1) and (2):

$$I_{\theta}(e) = \delta_{e, \text{START}} + \sum_{e'} I_{\theta}(e') \cdot \exp \pi_e \cdot p(e' \rightarrow e) \quad (4)$$

$$p_{\theta}(e) = I_{\theta}(e) \cdot \exp \pi_e \cdot p(e \rightarrow \text{HALT}) / Z \quad (5)$$

where  $Z$  is a global normalizing factor chosen so that  $\sum_e p_{\theta}(e) = 1$ . The new prior on  $\pi_e$  is the same as the old prior on  $\theta_e$ .

Increasing either  $\theta_e$  or  $\pi_e$  will raise  $p_{\theta}(e)$ ; the learner may do this to account for observations of  $e$  in training data. The probabilities of other rules consequently decrease so that  $\sum_e p_{\theta}(e) = 1$ . When  $\pi_e$  is raised, *all* rules’ probabilities are scaled down slightly and equally (because  $Z$  increases). When  $\theta_e$  is raised,  $e$  steals probability from its siblings,<sup>11</sup> but these are similar to  $e$  so tend to appear in test data if  $e$  is in training data. Raising  $\theta_e$  without disproportionately harming  $e$ ’s siblings requires manipulation of many other parameters, which is discouraged by the prior and may also suffer from search error. We speculate that this is why  $\pi_e$  works better.

<sup>11</sup>Raising the probability of an arc from  $e'$  to  $e$  decreases the probabilities of arcs from  $e'$  to siblings of  $e$ , as they sum to 1.

(Insert)	(Insert, target)	If the arc inserts Adv after TO in TO fund PP, then target=Adv left=TO right=— side=left of head
(Insert, left)	(Insert, target, left)	
(Insert, right)	(Insert, target, right)	
(Insert, left, right)		
(Insert, side)	(Insert, side, target)	
(Insert, side, left)	(Insert, side, target, left)	
(Insert, side, right)	(Insert, side, target, right)	
(Insert, side, left, right)		

Table 2: Each Insert arc has 14 features. The features of any given arc are found by instantiating the tuples above, as shown. Each instantiated tuple has a weight specified in  $\theta$ .

$S \rightarrow \dots$ rules only	train	dev	test
Treebank sections	0–15	16	17
sentences	15554	1343	866
rule tokens	18836	1588	973
rule types	11565	1317	795
frame types	2722	564	365
headword types	3607	756	504
novel rule tokens		51.6%	47.8%
novel frame tokens		8.9%	6.3%
novel headword tokens		10.4%	10.2%
novel rule types		61.4%	57.5%
novel frame types		24.6%	16.4%
novel headword types		20.9%	18.8%
nonterminal types	78		
# transformations applicable to rule with RHS length = $n$	$158n - 1$	$158n - 1$	$158n - 1$

Table 3: Properties of the experimental data. “Novel” means not observed in training. “Frame” was defined in section 4.

## 8 Evaluation<sup>12</sup>

To evaluate the quality of generalization, we used reparsed training data  $D$  and testing data  $E$  (Table 3). Each dataset consisted of a collection of flat rules such as  $S_{\text{put}} \rightarrow \text{NP}_{\text{put}} \text{NP}_{\text{PP}}$  extracted from the Penn Treebank (Marcus et al., 1993). Thus,  $p(D | \theta, \pi)$  and  $p(E | \theta, \pi)$  were each defined as a product of rule probabilities of the form  $p_{\theta, \pi}(\text{NP}_{\text{put}} \text{NP}_{\text{PP}} | S_{\text{put}})$ .

The learner attempted to maximize  $p(\theta, \pi) \cdot p(D | \theta, \pi)$  by gradient ascent. This amounts to learning the generalizations and exceptions that related the training rules  $D$ . The evaluation measure was then the perplexity on test data,  $-\log_2 p(E | \theta, \pi) / |E|$ . To get a good (low) perplexity score, the model had to assign reasonable probabilities to the many novel rules in  $E$  (Table 3). For many of these rules, even the frame was novel.

Note that although the training data was reparsed into rules, it was not annotated with the paths in Figure 1 that generated those rules, so estimating  $\theta$  and  $\pi$  was still an unsupervised learning problem.

The transformation graph had about 14 features per arc (Table 2). In the finite part of the transformation graph that was actually explored (including bad arcs that compete with good ones), about 70000 distinct features were encountered, though after training, only a few hundred

<sup>12</sup>See (Eisner, 2001) for full details of data preparation, model structure, parameter initialization, backoff levels for the comparison models, efficient techniques for computing the objective and its gradient, and more analysis of the results.

(a)	basic		Treebank/Markov		
	flat	non-flat <sup>b</sup>	Katz	one-count <sup>a</sup>	
			flat	flat	non-flat
Treebank	$\infty$	$\infty$			
1-gram	1774.9	86435.1	340.9	160.0	193.2
2-gram	<b>135.2</b>	199.3	127.2	<b>116.2</b>	174.7
3-gram	136.5	177.4	132.7	123.3	174.8
Collins <sup>c</sup>	363.0	494.5	197.9		
transformation	<b>108.6</b>				
averaged <sup>d</sup>	<b>102.3</b>				
(b)					
1-gram	1991.2	96318.8	455.1	194.3	233.1
2-gram	<b>162.2</b>	236.6	153.2	<b>138.8</b>	205.6
3-gram	161.9	211.0	156.8	145.7	208.1
Collins	414.5	589.4	242.0		
transformation	<b>124.8</b>				
averaged	<b>118.0</b>				

<sup>a</sup>Back off from Treebank grammar with Katz vs. one-count backoff (Chen and Goodman, 1996) (Note: One-count was always used for backoff *within* the  $n$ -gram and Collins models.)

<sup>b</sup>See section 2 for discussion

<sup>c</sup>Collins (1997, model 2)

<sup>d</sup>Average of transformation model with best other model

Table 4: Perplexity of the test set under various models. (a) Full training set. (b) Half training set (sections 0–7 only).

feature weights were substantial, and only a few thousand were even far enough from zero to affect performance. There was also a parameter  $\pi_e$  for each observed rule  $e$ .

Results are given in Table 4a, which compares the transformation model to various competing models discussed in section 2. The best (smallest) perplexities appear in boldface. The key results:

- The transformation model was the winner, reducing perplexity by 20% over the best model replicated from previous literature (a bigram model).
- Much of this improvement could be explained by the transformation model’s ability to model exceptions. Adding this ability more directly to the bigram model, using the new Treebank/Markov approach of section 2, also reduced perplexity from the bigram model, by 6% or 14% depending on whether Katz or one-count backoff was used, versus the transformation model’s 20%.
- Averaging the transformation model with the best competing model (Treebank/bigram) improved it by an additional 6%. So using transformations yields a total perplexity reduction of 12% over Treebank/bigram, and 24% over the best previous model from the literature (bigram).
- What would be the cost of achieving such a perplexity improvement by additional annotation? Training the averaged model on only the first half of the training set, with no further tuning of any options (Table 4b), yielded a test set perplexity of 118.0. So by using transformations, we can achieve about the same perplexity as the best model without transformations (Treebank/bigram, 116.2), *using only half as much training data*.
- Furthermore, comparing Tables 4a and 4b shows that the transformation model had the most graceful performance degradation when the dataset was reduced in size.

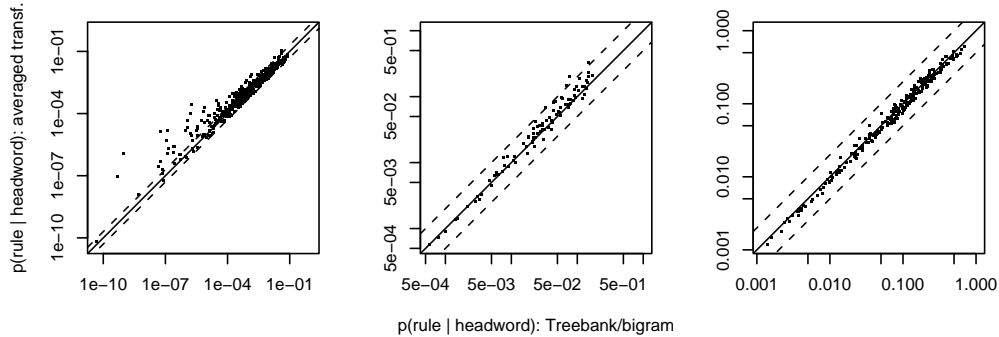


Figure 2: Probabilities of test set flat rules under the averaged model, plotted against the corresponding probabilities under the best transformation-free model. Improvements fall above the main diagonal; dashed diagonals indicate a factor of two. The three log-log plots (at different scales!) partition the rules by the number of training observations: 0 (left graph), 1 (middle),  $\geq 2$  (right).

This is an encouraging result for the use of the method in less supervised contexts (although results on a *noisy* dataset would be more convincing in this regard).

- The competing models from the literature are best used to predict flat rules directly, rather than by summing over their possible non-flat internal structures, as has been done in the past. This result is significant in itself. Extending Johnson (1998), it shows the inappropriateness of the traditional independence assumptions that build up a frame by several rule expansions (section 2).

Figure 2 shows that averaging the transformation model with the Treebank/bigram model improves the latter not merely on balance, but across the board. In other words, there is no evident class of phenomena for which incorporating transformations would be a bad idea.

- Transformations particularly helped raise the estimates of the low-probability novel rules in test data, as hoped.
- Transformations also helped on test rules that had been observed once in training with relatively infrequent words. (In other words, the transformation model does not discount singletons too much.)
- Transformations hurt slightly on balance for rules observed more than once in training, but the effect was tiny.

All these differences are slightly exaggerated if one compares the transformation model directly with the Treebank/bigram model, without averaging.

The transformation model was designed to use edit operations in order to generalize appropriately from a word’s observed frames to new frames that are likely to appear with that word in test data. To directly test the model’s success at such generalization, we compared it to the bigram model on a pseudo-disambiguation task.

Each instance of the task consisted of a pair of rules from test data, expressed as (word, frame) pairs  $(w_1, f_1)$  and  $(w_2, f_2)$ , such that  $f_1$  and  $f_2$  are “novel” frames that did *not* appear in training data (with any headword).

Each model was then asked: Does  $f_1$  go with  $w_1$  and  $f_2$  with  $w_2$ , or vice-versa? In other words, which is bigger,  $p(f_1 | w_1) \cdot p(f_2 | w_2)$  or  $p(f_2 | w_1) \cdot p(f_1 | w_2)$ ?

Since the frames were novel, the model had to make the choice according to whether  $f_1$  or  $f_2$  looked more like the frames that had actually been observed with  $w_1$  in the past, and likewise  $w_2$ . What this means depends on the model. The bigram model takes two frames to look alike if they *contain many bigrams in common*. The transformation model takes two frames to look alike if they are *connected by a path of probable transformations*.

The test data contained 62 distinct rules  $(w, f)$  in which  $f$  was a novel frame. This yielded  $\frac{62 \cdot 61}{2} = 1891$  pairs of rules, leading to 1811 task instances after obvious ties were discarded.<sup>13</sup>

Baseline performance on this difficult task is 50% (random guess). The bigram model chose correctly in 1595 of the 1811 instances (88.1%). Parameters for “memorizing” specific frames do not help on this task, which involves only novel frames, so the Treebank/bigram model had the same performance. By contrast, the transformation model got 1669 of 1811 correct (92.2%), for a more-than-34% reduction in error rate. (The development set showed similar results.) However, since the 1811 task instances were derived non-independently from just 62 novel rules, this result is based on a rather small sample.

## 9 Discussion

This paper has presented a nontrivial way to reparameterize a PCFG in terms of “deep” parameters representing transformations and exceptions. A linguistically sensible prior was natural to define over these deep parameters.

Famous examples of “deep reparameterization” are the Fourier transform in speech recognition and the SVD transform for Latent Semantic Analysis in IR. Like our technique, they are intended to reveal significant structure through the leading parameters while relegating noise and exceptions to minor parameters. Such representations

<sup>13</sup>An obvious tie is an instance where  $f_1 = f_2$ , or where both  $w_1$  and  $w_2$  were novel headwords. (The 62 rules included 11 with novel headwords.) In such cases, neither the bigram nor the transformation model has any basis for making its decision: the probabilities being compared will necessarily be equal.

make it easier to model the similarity or probability of the objects at hand (waveforms, documents, or grammars).

Beyond the fact that it shows at least a good perplexity improvement (it has not yet been applied to a real task), an exciting “big idea” aspect of this work is its flexibility in defining linguistically sensible priors over grammars. Our reparameterization is made with reference to a user-designed transformation graph (Figure 1). The graph need not be confined to edit distance transformations, or to the simple features of Table 2 (used here for comparability with the Markov models), which condition a transformation’s probability on *local* context.

In principle, the approach could be used to capture a great many linguistic phenomena. Figure 1 could be extended with more ambitious transformations, such as gapping, gap-threading, and passivization. The flat rules could be annotated with internal structure (as in TAG) and thematic roles. Finally, the arcs could bear further features. For example, the probability of unaccusative movement (*someone sank the boat* → *the boat sank*) should depend on whether the headword is a change-of-state verb.

Indeed, Figure 1 can be converted to any lexicalized theory of grammar, such as categorial grammar, TAG, LFG, HPSG, or Minimalism. The vertices represent lexical entries and the arcs represent probabilistic lexical redundancy rules or metarules (see footnote 8). The transformation model approach is therefore a full stochastic treatment of lexicalized syntax—apparently the first to treat lexical redundancy rules, although (Briscoe and Copestake, 1999) give an *ad hoc* approach. See (Eisner, 2001; Eisner, 2002a) for more discussion.

It is worthwhile to compare the statistical approach here with some other approaches:

- Transformation models are similar to graphical models: they allow similar patterns of deductive and abductive inference from observations. However, the vertices of a transformation graph do not represent different random variables, but rather mutually exclusive values of the same random variable, whose probabilities sum to 1.
- Transformation models incorporate conditional log-linear (maximum entropy) models. As an alternative, one could directly build a conditional log-linear model of  $p(\text{RHS} \mid \text{LHS})$ . However, such a model would learn probabilities, not relationships. A feature weight would not really model the strength of the relationship between two frames  $e, e'$  that share that feature. It would only influence both frames’ probabilities. If the probability of  $e$  were altered by some *unrelated* factor (e.g., an exception weight), then the probability of  $e'$  would not respond.
- A transformation model can be regarded as a probabilistic FSA that consists mostly of  $\epsilon$ -transitions. (Rules are only emitted on the arcs to HALT.) This perspective allows use of generic methods for finite-state parameter

estimation (Eisner, 2002b). We are strongly interested in improving the speed of such methods and their ability to avoid local maxima, which are currently the major difficulty with our system, as they are for many unsupervised learning techniques. We expect to further pursue transformation models (and simpler variants that are easier to estimate) within this flexible finite-state framework.

The interested reader is encouraged to look at (Eisner, 2001) for a much more careful and wide-ranging discussion of transformation models, their algorithms, and their relation to linguistic theory, statistics, and parsing. Chapter 1 provides a good overview. For a brief article highlighting the connection to linguistics, see (Eisner, 2002a).

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